Dossier zur Nutzenbewertung gemäß § 35a SGB V

Abrocitinib (Cibinqo®)

Pfizer Europe MA EEIG als örtlicher Vertreter des Zulassungsinhabers Pfizer Europe MA EEIG

Modul 4 A

Behandlung von mittelschwerer bis schwerer atopischer Dermatitis bei Jugendlichen ab einem Alter von 12 Jahren, die für eine systemische Therapie infrage kommen

Medizinischer Nutzen und medizinischer Zusatznutzen, Patient:innengruppen mit therapeutisch bedeutsamem Zusatznutzen

Stand: 18.04.2024

Anhang 4-G: Zusatzanalysen

| Resp | onderanalysen JADE DARE zu Woche 26 | 10 |
|------|--|----|
| Pr | oportion of Subjects Achieving EASI Response >= 75% Improvement from Baseline at Week 26 by Subgroup - (FAS, NRI) | 11 |
| Pr | oportion of Subjects Achieving EASI Response >= 90% Improvement from Baseline at Week 26 by Subgroup - (FAS, NRI) | 20 |
| Bi | nary Outcome Analysis: EASI-100 response at week 26 - Full Analysis Set Safety Population | 29 |
| | oportion of Subjects Achieving Peak Pruritus Numerical Rating Scale (PP-NRS4) Response >= 4 Points Improvement from Baseline at Week 26 by abgroup - (FAS with Baseline >= 4, NRI) | 31 |
| Bi | nary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 26 - Full Analysis Set Safety Population | 41 |
| Pr | oportion of Subjects with DLQI < 2 Response at Week 26 by Subgroup (FAS with Baseline >= 2, NRI) | 43 |
| Pr | oportion of Subjects Achieving POEM Total Score < 3 Response at Week 26 by Subgroup - (FAS with Baseline >= 3, NRI) | 52 |
| Bi | nary Outcome Analysis: Achieving 0 in POEM total score at week 26 - Full Analysis Set Safety Population | 61 |
| Pr | oportion of Subjects Achieving SCORAD Response >= 75% Improvement from Baseline at Week 26 by Subgroup - (FAS, NRI) | 63 |
| Pr | oportion of Subjects Achieving SCORAD Response >= 90% Improvement from Baseline at Week 26 by Subgroup - (FAS, NRI) | 72 |
| Bi | nary Outcome Analysis: SCORAD-100 response at week 26 - Full Analysis Set Safety Population | 81 |
| Bi | nary Outcome Analysis: EQ-5D VAS 15-point improvement at week 26 - Full Analysis Set Safety Population | 83 |
| Bi | nary Outcome Analysis: Adverse Events-Total mortality at 26 weeks (TEAE leading to death) - Safety Set | 85 |
| Pr | oportion of Subjects with Treatment-Emergent Adverse Events (Excluding Adverse Events Related to AD, All Causalities) - Safety Analysis Set | 86 |
| | oportion of Subjects with Treatment-Emergent Serious Adverse Events (Excluding Adverse Events Related to AD, All Causalities) - Safety Analyset | |
| | oportion of Subjects with Treatment-Emergent Severe Adverse Events (Excluding Adverse Events Related to AD, All Causalities) - Safety Analysi | |
| Pr | oportion of Subjects with Non-Severe Treatment-Emergent Adverse Events (All Causalities) - Safety Analysis Set | 89 |
| Pr | oportion of Subjects with Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set | 90 |
| | oportion of Subjects with Treatment-Emergent Adverse Events by Subgroup (Excluding Adverse Events Related to AD, All Causalities) - Safety | |
| Aı | nalysis Set | 95 |

| roportion of Subjects with Treatment-Emergent Serious Adverse Events by Subgroup (All Causalities) - Safety Analysis Set100 |
|--|
| roportion of Subjects with Treatment-Emergent Serious Adverse Events by Subgroup (Excluding Adverse Events Related to AD, All Causalities) - afety Analysis Set |
| roportion of Subjects with Treatment-Emergent Severe Adverse Events by Subgroup (All Causalities) - Safety Analysis Set108 |
| Proportion of Subjects with Treatment-Emergent Severe Adverse Events by Subgroup (Excluding Adverse Events Related to AD, All Causalities) - afety Analysis Set |
| Proportion of Subjects with Treatment-Emergent Adverse Events Occurring in >= 10 Subjects in any Treatment Group by Subgroup, System Organ Class and Preferred Term (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Treatment-Emergent Serious Adverse Events Occurring in >= 10 Subjects in any Treatment Group by Subgroup, System Organ Class and Preferred Term (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Treatment-Emergent Severe Adverse Events Occurring in >= 10 Subjects in any Treatment Group by Subgroup, System Organ Class and Preferred Term (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Non-Severe Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set199 |
| Proportion of Subjects with Superinfections of Infections (Treated with Antibiotics, Antiviral or Fungicidal Agents for a Duration > 2 Weeks) Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Special Interest Treatment-Emergent Adverse Events (Herpes Zoster) by Subgroup (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Special Interest Treatment-Emergent Adverse Events (Acne) by Subgroup (All Causalities) - Safety Analysis Set213 |
| Proportion of Subjects with Special Interest Treatment-Emergent Adverse Events (Folliculitis) by Subgroup (All Causalities) - Safety Analysis Set218 |
| Proportion of Subjects with Special Interest Treatment-Emergent Adverse Events (Conjonctivitis) by Subgroup (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Special Interest Treatment-Emergent Serious Adverse Events by Subgroup (All Causalities) - Safety Analysis Set228 |
| Proportion of Subjects with Special Interest Treatment-Emergent Severe Adverse Events by Subgroup (All Causalities) - Safety Analysis Set29 |
| Proportion of Subjects with Serious Superinfections of Infections (Treated with Antibiotics, Antiviral or Fungicidal Agents for a Duration > 2 Weeks) Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set |
| Proportion of Subjects with Severe Superinfections of Infections (Treated with Antibiotics, Antiviral or Fungicidal Agents for a Duration > 2 Weeks) Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set |

| Proportion of Subjects with Discontinuations from Study Drug due to Adverse Events by Subgroup (All Causalities) - Safety Ana | alysis Set232 |
|---|--------------------|
| Non-Severe Treatment Emergent Subset Flag Adverse Events, overall and by SOC/PT - Safety Population | 237 |
| Responderanalysen JADE DARE zu Woche 12 | 238 |
| Binary Outcome Analysis: SCORAD-75 response at week 12 - Full Analysis Set Safety Population | 239 |
| Binary Outcome Analysis: SCORAD-90 response at week 12 - Full Analysis Set Safety Population | 241 |
| Binary Outcome Analysis: SCORAD-100 response at week 12 - Full Analysis Set Safety Population | 243 |
| Binary Outcome Analysis: EASI-75 response at week 12 - Full Analysis Set Safety Population | 245 |
| Binary Outcome Analysis: EASI-90 response at week 12 - Full Analysis Set Safety Population | 247 |
| Binary Outcome Analysis: EASI-100 response at week 12 - Full Analysis Set Safety Population | 249 |
| Binary Outcome Analysis: Achieving 0-2 in POEM total score at week 12 - Full Analysis Set Safety Population | 251 |
| Binary Outcome Analysis: Achieving 0 in POEM total score at week 12 - Full Analysis Set Safety Population | 253 |
| Binary Outcome Analysis: Achieving 0-1 in CDLQI total score at week 12 - Full Analysis Set Safety Population | 255 |
| Binary Outcome Analysis: PP-NRS total score 4-point improvement at week 12 - Full Analysis Set Safety Population | 257 |
| Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 12 - Full Analysis Set Safety Population | 259 |
| Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 12 - Full Analysis Set Safety Population | 261 |
| Binary Outcome Analysis: Adverse Events-Total mortality at 12 weeks (TEAE leading to death) - Safety Set | 263 |
| Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events at 12 weeks - Safety Set | 265 |
| Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks - Safety Set | 267 |
| Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks - Safety Set | 269 |
| Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set | 271 |
| Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events at 12 weeks excluding progression events - | Safety Set273 |
| Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks excluding progression event | s - Safety Set275 |
| Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks excluding progression even | ts - Safety Set277 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Superinfections at 12 weeks - Safety Set | 279 |

| | Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Herpes Zoster at 12 weeks - Safety Set | 281 |
|---|--|-----|
| | Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Conjunctivitis at 12 weeks - Safety Set | 283 |
| | Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Acne (PT) at 12 weeks - Safety Set | 285 |
| | Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Folliculitis (PT) at 12 weeks - Safety Set | 287 |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Infections and infestations [SOC] – Safety Set | |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Gastrointestinal disorders [SOC] – Safety Set | |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Skin and subcutaneous tissue disorders[SOC] - Safety Set | 294 |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Investigations [SOC] - Safety Set | 297 |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nervous system disorders [SOC] – Safety Set | |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nausea [PT] - Safety Set | 302 |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Headache [PT] - Safety Set | 304 |
| | Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Acne [PT] - Safety Set | 306 |
| | Binary Outcome Analysis: Selected Adverse Events (PT)-PT Natural killer cell count decreased at 12 weeks - Safety Set | 308 |
| | Binary Outcome Analysis: Selected Adverse Events (PT)-PT Headache at 12 weeks - Safety Set | 310 |
| | Binary Outcome Analysis: Selected Adverse Events (PT)-SOC Nervous system disorders at 12 weeks - Safety Set | 312 |
| | Binary Outcome Analysis: Selected Adverse Events (SOC)-SOC Skin and subcutaneous tissue disorders at 12 weeks - Safety Set | 314 |
| N | MRM Analysen JADE DARE | 316 |
| | Descriptive Summary of Total EASI Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) | 317 |
| | Least Squares Mean of Percent Change from Baseline in Total EASI Score at Weeks 2, 4, 8, 12, 16, 20 and 26 - MMRM (FAS, OD) | 319 |
| | Plot of Least Squares Mean of Percent Change from Baseline in Total EASI Score at Weeks 2, 4, 8, 12, 16, 20 and 26 - MMRM (FAS. OD) | 324 |

| Descriptive Summary of Peak Pruritus Numerical Rating Scale (PP-NRS), Change from Baseline and Percent Change from Baseline up to (FAS, OD) | |
|--|-------------|
| Least Squares Mean of Change from Baseline in Peak Pruritus Numerical Rating Scale (PP-NRS) up to Week 26 - MMRM (FAS, OD) | 327 |
| Plot of Least Squares Mean of Change from Baseline in Peak Pruritus Numerical Rating Scale (PP-NRS) at Weeks 2, 4, 8, 12, 16, 20 and (FAS, OD) | |
| Descriptive Summary of DLQI, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) | 333 |
| Least Squares Mean of Change from Baseline in DLQI at Weeks 2, 12, 16, 20 and 26 - MMRM (FAS, OD) | 334 |
| Plot of Least Squares Mean of Change from Baseline in DLQI - MMRM (FAS, OD) | 338 |
| Descriptive Summary of EQ-5D VAS Score, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (| FAS, OD)339 |
| Least Squares Mean of Change from Baseline in EQ-5D VAS Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) | 340 |
| Plot of Least Squares Mean of Percent Change from Baseline in EQ-5D VAS Score up to Week 26 (FAS, OD) | 342 |
| Descriptive Summary of POEM, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) | 343 |
| Least Squares Mean of Change from Baseline in POEM Total Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) | 344 |
| Plot of Least Squares Mean of Change from Baseline in POEM - MMRM (FAS, OD) | 346 |
| Descriptive Summary of SCORAD Total Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) | 347 |
| Least Squares Mean of Percent Change from Baseline in SCORAD Total Score - MMRM (FAS, OD) | 349 |
| Plot of Least Squares Mean of Percent Change from Baseline in SCORAD Total Score - MMRM (FAS, OD) | 355 |
| Responderanalysen JADE TEEN zu Woche 12 | 356 |
| Binary Outcome Analysis: SCORAD-75 response at week 12 - Full Analysis Set Safety Population | 357 |
| Binary Outcome Analysis: SCORAD-90 response at week 12 - Full Analysis Set Safety Population | 359 |
| Binary Outcome Analysis: SCORAD-100 response at week 12 - Full Analysis Set Safety Population | 361 |
| Binary Outcome Analysis: EASI-75 response at week 12 - Full Analysis Set Safety Population | 363 |
| Binary Outcome Analysis: EASI-90 response at week 12 - Full Analysis Set Safety Population | 365 |
| Binary Outcome Analysis: EASI-100 response at week 12 - Full Analysis Set Safety Population | 367 |

| Binary Outcome Analysis: Achieving 0-2 in POEM total score at week 12 - Full Analysis Set Safety Population | 369 |
|--|-----|
| Binary Outcome Analysis: Achieving 0 in POEM total score at week 12 - Full Analysis Set Safety Population | 371 |
| Binary Outcome Analysis: Achieving 0-1 in CDLQI total score at week 12 - Full Analysis Set Safety Population | 373 |
| Binary Outcome Analysis: PP-NRS total score 4-point improvement at week 12 - Full Analysis Set Safety Population | 375 |
| Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 12 - Full Analysis Set Safety Population | 377 |
| Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 12 - Full Analysis Set Safety Population | 379 |
| Binary Outcome Analysis: Adverse Events-Total mortality (TEAE leading to death) - Safety Set | 381 |
| Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events - Safety Set | 382 |
| Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events - Safety Set | 384 |
| Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events - Safety Set | 386 |
| Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set | 388 |
| Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events excluding progression events - Safety Set | 390 |
| Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events excluding progression events - Safety Set | 392 |
| Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events excluding progression events - Safety Set | 394 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Superinfections - Safety Set | 396 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Herpes Zoster - Safety Set | 397 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Conjunctivitis - Safety Set | 398 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Acne (PT) - Safety Set | 400 |
| Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Folliculitis (PT) - Safety Set | 402 |
| Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Infections and infestations [SOC] - Safety Set | 404 |
| Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Gastrointestinal disorders [SOC] - Safety Set | 406 |
| Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Nervous system disorders [SOC] - Safety Set | 408 |
| Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Nausea [PT] - Safety Set | 410 |
| | |

| Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Upper respiratory tract infection [PT] - Safety Set |
|---|
| Binary Outcome Analysis: Selected Adverse Events (PT)-PT Natural killer cell count decreased - Safety Set |
| Binary Outcome Analysis: Selected Adverse Events (PT)-PT Headache - Safety Set |
| Binary Outcome Analysis: Selected Adverse Events (PT)-SOC Nervous system disorders - Safety Set |
| Binary Outcome Analysis: Selected Adverse Events (SOC)-SOC Skin and subcutaneous tissue disorders - Safety Set |
| Safety incidence analysis: Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set421 |
| Safety incidence analysis: Severe Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set422 |
| Safety incidence analysis: Serious Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set423 |
| MMRM Analysen JADE TEEN424 |
| MMRM models: EASI01-Total EASI Score (Derived) percent change from baseline, Total population - Full Analysis Set Population425 |
| MMRM models: SCA01-Total Score percent change from baseline, Total population - Full Analysis Set Population |
| MMRM models: POEM01-Total Score, Total population - Full Analysis Set Population |
| MMRM models: NRS01-Worst Itching-Atopic Dermatitis-Analysis, Total population - Full Analysis Set Population |
| MMRM models: CDLQ1-Total Score, Total population - Full Analysis Set Population |
| MMRM models: EQ5D03-EQ VAS Score, Total population - Full Analysis Set Population |
| EASI01-Total EASI Score (Derived) percent change from baseline, Least-Square Means estimates |
| SCA01-Total Score percent change from baseline, Least-Square Means estimates |
| POEM01-Total Score, Least-Square Means estimates |
| NRS01-Worst Itching-Atopic Dermatitis-Analysis, Least-Square Means estimates |
| CDLQ1-Total Score, Least-Square Means estimates |
| EQ5D03-EQ VAS Score, Least-Square Means estimates |
| Zusatzanalysen JADE DARE und JADE TEEN467 |
| Binary Outcome Analysis: SCORAD-90 response - Full Analysis Set Safety Population |

| Binary Outcome Analysis: SCORAD-100 response - Full Analysis Set Safety Population |) |
|--|----------|
| Binary Outcome Analysis: EASI-100 response - Full Analysis Set Safety Population |) |
| Binary Outcome Analysis: Achieving 0-2 in POEM total score - Full Analysis Set Safety Population | - |
| Binary Outcome Analysis: Achieving 0 in POEM total score - Full Analysis Set Safety Population | <u>)</u> |
| Binary Outcome Analysis: Achieving 0-1 in DLQI total score - Full Analysis Set Safety Population | } |
| Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score - Full Analysis Set Safety Population | ļ |
| Binary Outcome Analysis: SCORAD-90 response - Full Analysis Set Safety Population | ; |
| Binary Outcome Analysis: SCORAD-100 response - Full Analysis Set Safety Population | , |
| Binary Outcome Analysis: EASI-100 response - Full Analysis Set Safety Population | } |
| Binary Outcome Analysis: Achieving 0-2 in POEM total score - Full Analysis Set Safety Population |) |
| Binary Outcome Analysis: Achieving 0 in POEM total score - Full Analysis Set Safety Population |) |
| Binary Outcome Analysis: Achieving 0-1 in DLQI total score - Full Analysis Set Safety Population | L |
| Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score - Full Analysis Set Safety Population | <u>)</u> |

Responderanalysen JADE DARE (Woche 26)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 230 | 247 |
| | Week 26, N | 230 | 247 |
| | Number of Subjects with observed Case, N1 (%) | 185 (80.4) | 220 (89.1) |
| | Number of Subjects with NRI, N2 (%) | 45 (19.6) | 27 (10.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 154 (67.0) | 179 (72.5) |
| | 95% CI | (60.9, 73.0) | (66.9, 78.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9239 | |
| | 95% CI | (0.8203, 1.0406) | |
| | Two-sided P-value | 0.1923 | |
| | | | - |
| Age (years) group: >=40 | Baseline, N | 132 | 118 |
| | Week 26, N | 132 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 116 (87.9) | 104 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 16 (12.1) | 14 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 100 (75.8) | 82 (69.5) |
| | 95% CI | (68.4, 83.1) | (61.2, 77.8) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0902 | |
| | 95% CI | (0.9349, 1.2712) | |
| | Two-sided P-value | 0.2707 | |
| | | | |
| | P-value of interaction | 0.0951 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 193 | 204 |
| | Week 26, N | 193 | 204 |
| | Number of Subjects with observed Case, N1 (%) | 169 (87.6) | 180 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 24 (12.4) | 24 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 137 (71.0) | 134 (65.7) |
| | 95% CI | (64.6, 77.4) | (59.2, 72.2) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0807 | |
| | 95% CI | (0.9451, 1.2357) | |
| | Two-sided P-value | 0.2568 | |
| | | | |
| Sex: Female | Baseline, N | 169 | 161 |
| | Week 26, N | 169 | 161 |
| | Number of Subjects with observed Case, N1 (%) | 132 (78.1) | 144 (89.4) |
| | Number of Subjects with NRI, N2 (%) | 37 (21.9) | 17 (10.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 117 (69.2) | 127 (78.9) |
| | 95% CI | (62.3, 76.2) | (72.6, 85.2) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.8776 | |
| | 95% CI | (0.7719, 0.9979) | |
| | Two-sided P-value | 0.0464 | |
| | | | |
| | P-value of interaction | 0.0280 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| D 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 | In 11 27 | 155 | 105 |
| Region of enrollment: US/Canada/Australia | | 177 | 195 |
| | Week 26, N | 177 | 195 |
| | Number of Subjects with observed Case, N1 (%) | 142 (80.2) | 169 (86.7) |
| | Number of Subjects with NRI, N2 (%) | 35 (19.8) | 26 (13.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 118 (66.7) | 136 (69.7) |
| | 95% CI | (59.7, 73.6) | (63.3, 76.2) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate Estimate | 0.9559 | |
| | | + | |
| | 95% CI | (0.8316, 1.0987) | |
| | Two-sided P-value | 0.5255 | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| | Week 26, N | 150 | 132 |
| | Number of Subjects with observed Case, N1 (%) | 130 (86.7) | 122 (92.4) |
| | Number of Subjects with NRI, N2 (%) | 20 (13.3) | 10 (7.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 111 (74.0) | 94 (71.2) |
| | 95% CI | (67.0, 81.0) | (63.5, 78.9) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0391 | |
| | 95% CI | (0.8997, 1.2002) | |
| | Two-sided P-value | 0.6014 | |
| | Ta. 11. 12 | | |
| Region of enrollment: Asia | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (88.2) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 2 (11.8) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 11 (64.7) | 16 (84.2) |
| | 95% CI | (42.0, 87.4) | (67.8, 100.0) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.7684 | |
| | 95% CI | (0.5143, 1.1480) | |
| | Two-sided P-value | 0.1983 | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| • | Week 26, N | 18 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 14 (77.8) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (22.2) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 14 (77.8) | 15 (78.9) |
| | 95% CI | (58.6, 97.0) | (60.6, 97.3) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9852 | |
| | 95% CI | (0.7020, 1.3827) | |
| | Two-sided P-value | 0.9312 | |
| | P-value of interaction | 0.5379 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 216 | 220 |
| | Week 26, N | 216 | 220 |
| | Number of Subjects with observed Case, N1 (%) | 178 (82.4) | 195 (88.6) |
| | Number of Subjects with NRI, N2 (%) | 38 (17.6) | 25 (11.4) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 159 (73.6) | 160 (72.7) |
| | 95% CI | (67.7, 79.5) | (66.8, 78.6) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0122 | |
| | 95% CI | (0.9034, 1.1340) | |
| | Two-sided P-value | 0.8350 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 146 | 145 |
| | Week 26, N | 146 | 145 |
| | Number of Subjects with observed Case, N1 (%) | 123 (84.2) | 129 (89.0) |
| | Number of Subjects with NRI, N2 (%) | 23 (15.8) | 16 (11.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 95 (65.1) | 101 (69.7) |
| | 95% CI | (57.3, 72.8) | (62.2, 77.1) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9342 | |
| | 95% CI | (0.7959, 1.0965) | |
| | Two-sided P-value | 0.4047 | |
| | | | |
| | P-value of interaction | 0.4236 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 230 | 247 |
| | Week 26, N | 230 | 247 |
| | Number of Subjects with observed Case, N1 (%) | 185 (80.4) | 220 (89.1) |
| | Number of Subjects with NRI, N2 (%) | 45 (19.6) | 27 (10.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 114 (49.6) | 113 (45.7) |
| | 95% CI | (43.1, 56.0) | (39.5, 52.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0834 | |
| | 95% CI | (0.8975, 1.3078) | |
| | Two-sided P-value | 0.4042 | |
| | | | |
| Age (years) group: >=40 | Baseline, N | 132 | 118 |
| | Week 26, N | 132 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 116 (87.9) | 104 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 16 (12.1) | 14 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 76 (57.6) | 59 (50.0) |
| | 95% CI | (49.1, 66.0) | (41.0, 59.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1515 | |
| | 95% CI | (0.9127, 1.4527) | |
| | Two-sided P-value | 0.2341 | |
| | | | |
| | P-value of interaction | 0.6895 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 193 | 204 |
| | Week 26, N | 193 | 204 |
| | Number of Subjects with observed Case, N1 (%) | 169 (87.6) | 180 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 24 (12.4) | 24 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 95 (49.2) | 83 (40.7) |
| | 95% CI | (42.2, 56.3) | (33.9, 47.4) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2098 | |
| | 95% CI | (0.9718, 1.5061) | |
| | Two-sided P-value | 0.0883 | |
| | | | |
| Sex: Female | Baseline, N | 169 | 161 |
| | Week 26, N | 169 | 161 |
| | Number of Subjects with observed Case, N1 (%) | 132 (78.1) | 144 (89.4) |
| | Number of Subjects with NRI, N2 (%) | 37 (21.9) | 17 (10.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 95 (56.2) | 89 (55.3) |
| | 95% CI | (48.7, 63.7) | (47.6, 63.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0169 | |
| | 95% CI | (0.8389, 1.2326) | |
| | Two-sided P-value | 0.8645 | |
| | | | |
| | P-value of interaction | 0.2428 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 177 | 195 |
| | Week 26, N | 177 | 195 |
| | Number of Subjects with observed Case, N1 (%) | 142 (80.2) | 169 (86.7) |
| | Number of Subjects with NRI, N2 (%) | 35 (19.8) | 26 (13.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 88 (49.7) | 89 (45.6) |
| | 95% CI | (42.4, 57.1) | (38.6, 52.6) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0893 | |
| | 95% CI | (0.8802, 1.3480) | |
| | Two-sided P-value | 0.4314 | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| Region of enforment. Europe | Week 26, N | 150 | 132 |
| | | | - |
| | Number of Subjects with observed Case, N1 (%) | 130 (86.7) | 122 (92.4) |
| | Number of Subjects with NRI, N2 (%) | 20 (13.3) | 10 (7.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 83 (55.3) | 60 (45.5) |
| | 95% CI | (47.4, 63.3) | (37.0, 53.9) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2173 | |
| | 95% CI | (0.9616, 1.5410) | |
| | Two-sided P-value | 0.1021 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (88.2) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 2 (11.8) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 6 (35.3) | 10 (52.6) |
| | 95% CI | (12.6, 58.0) | (30.2, 75.1) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.6706 | |
| | 95% CI | (0.3098, 1.4515) | |
| | Two-sided P-value | 0.3104 | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| - | Week 26, N | 18 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 14 (77.8) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (22.2) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 13 (72.2) | 13 (68.4) |
| | 95% CI | (51.5, 92.9) | (47.5, 89.3) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0556 | |
| | 95% CI | (0.6944, 1.6046) | |
| | Two-sided P-value | 0.8002 | |
| | P-value of interaction | 0.5107 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 216 | 220 |
| | Week 26, N | 216 | 220 |
| | Number of Subjects with observed Case, N1 (%) | 178 (82.4) | 195 (88.6) |
| | Number of Subjects with NRI, N2 (%) | 38 (17.6) | 25 (11.4) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 119 (55.1) | 104 (47.3) |
| | 95% CI | (48.5, 61.7) | (40.7, 53.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1654 | |
| | 95% CI | (0.9692, 1.4013) | |
| | Two-sided P-value | 0.1036 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 146 | 145 |
| | Week 26, N | 146 | 145 |
| | Number of Subjects with observed Case, N1 (%) | 123 (84.2) | 129 (89.0) |
| | Number of Subjects with NRI, N2 (%) | 23 (15.8) | 16 (11.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 71 (48.6) | 68 (46.9) |
| | 95% CI | (40.5, 56.7) | (38.8, 55.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0370 | |
| | 95% CI | (0.8154, 1.3188) | |
| | Two-sided P-value | 0.7673 | |
| | | | |
| | P-value of interaction | 0.4499 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; EASI = eczema area and severity index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 09SEP2021 (22:33)

Binary Outcome Analysis: EASI-100 response at week 26 - Full Analysis Set Safety Population

| | | | - rop - root |
|-----------|---------------|----------|--------------|
| JADE DARE | (PF-04965842) |) - 2023 | datacut |

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|---------------------------|-----------|-----------------|-------------|------------|--------------|-----------------|---------------------------|-------------------|---------------|
| | | | | | | | | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 26 | | | | | | | | | |
| Full Analysis Set Safety | 362 | 79 (21.8%) | 365 | 50 (13.7%) | 1.76 | 1.59 | 8.1% | 0.0041+* | |
| Population | | (17.7%, | | (10.3%, | (1.19, 2.60) | (1.15, 2.20) | (2.6%, 13.7%) | | |
| | | 26.4%) | | 17.7%) | | | | | |
| Baseline Disease Severity | | | | | | | | | |
| Week 26 | | | | | | | | | |
| Moderate baseline | 216 | 52 (24.1%) | 220 | 31 (14.1%) | 1.93 | 1.71 | 10.0% | 0.0102* | 0.3916 |
| disease (IGA=3) | | (18.5%, | | (9.8%, | (1.18, 3.16) | (1.14, 2.56) | (2.7%, 17.3%) | | |
| | | 30.3%) | | 19.4%) | | | | | |
| Severe baseline disease | 146 | 27 (18.5%) | 145 | 19 (13.1%) | 1.50 | 1.41 | 5.4% | 0.2607 | |
| (IGA=4) | | (12.6%, | | (8.1%, | (0.79, 2.85) | (0.82, 2.42) | (-3.0%, 13.7%) | | |
| | | 25.8%) | | 19.7%) | | | | | |
| Gender | | | | | | | | | |
| Week 26 | | | | | | | | | |
| Male | 193 | 43 (22.3%) | 204 | 22 (10.8%) | 2.37 | 2.07 | 11.5% | 0.0026* | 0.2046 |
| | | (16.6%, | | (6.9%, | (1.36, 4.14) | (1.29, 3.32) | (4.2%, 18.7%) | | |
| | | 28.8%) | | 15.9%) | | | | | |
| Female | 169 | 36 (21.3%) | 161 | 28 (17.4%) | 1.29 | 1.22 | 3.9% | 0.4049 | |
| | | (15.4%, | | (11.9%, | (0.74, 2.23) | (0.79, 1.91) | (-4.6%, 12.4%) | | |
| | | 28.3%) | | 24.1%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

Binary Outcome Analysis: EASI-100 response at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|---------------------|-----------|-----------------|-------------|------------|---|---|---------------------------|-------------------|---------------|
| | | | | | | _ | _ | CMH or Logistic P | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 26 | | | | | | | | | |
| US/Canada/Australia | 177 | 46 (26.0%) | 195 | 18 (9.2%) | 3.45 | 2.82 | 16.8% | < 0.0001* | 0.0057* |
| | | (19.7%, | | (5.6%, | (1.91, 6.23) | (1.70, 4.67) | (9.1%, 24.4%) | | |
| | | 33.1%) | | 14.2%) | | | | | |
| Europe | 150 | 26 (17.3%) | 132 | 21 (15.9%) | 1.11 | 1.09 | 1.4% | 0.8730 | |
| 1 | | (11.6%, | | (10.1%, | (0.59, 2.08) | (0.64, 1.84) | (-7.3%, 10.1%) | | |
| | | 24.4%) | | 23.3%) | | | , | | |
| Asia | 17 | 1 (5.9%) | 19 | 5 (26.3%) | 0.18 | 0.22 | -20.4% | 0.1821 | |
| | | (0.1%, | | (9.1%, | (0.02, 1.68) | (0.03, 1.73) | (-43.2%, 2.3%) | | |
| | | 28.7%) | | 51.2%) | (, , , , , , , , , , , , , , , , , , , | (, | (==,, | | |
| Latin America | 18 | 6 (33.3%) | 19 | 6 (31.6%) | 1.08 | 1.06 | 1.8% | 1.0000 | |
| | | (13.3%, | | (12.6%, | (0.27, 4.29) | (0.42, 2.68) | (-28.4%, 31.9%) | | |
| | | 59.0%) | | 56.6%) | (, , , , , , , , , , , , , , , , , , , | (, , , , , , , , , , , , , , , , , , , | (2. 2., 2 2. 2., | | |
| Age Subgroup | | , | | , | | | | | |
| Week 26 | | | | | | | | | |
| <40 years | 230 | 41 (17.8%) | 247 | 31 (12.6%) | 1.51 | 1.42 | 5.3% | 0.1247 | 0.1947 |
| , | | (13.1%, | | (8.7%, | (0.91, 2.51) | (0.92, 2.18) | (-1.2%, 11.7%) | | |
| | | 23.4%) | | 17.3%) | | | , | | |
| >=40 years | 132 | 38 (28.8%) | 118 | 19 (16.1%) | 2.11 | 1.79 | 12.7% | 0.0231* | |
| - 3 | | (21.2%, | | (10.0%, | (1.13, 3.91) | (1.09, 2.92) | (2.5%, 22.9%) | ***-*- | |
| | | 37.3%) | | 24.0%) | (====, 5,5,1) | (====, ====) | (=15 /3, 22 15 /0) | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 227 | 246 |
| | Week 26, N | 227 | 246 |
| | Number of Subjects with observed Case, N1 (%) | 195 (85.9) | 221 (89.8) |
| | Number of Subjects with NRI, N2 (%) | 32 (14.1) | 25 (10.2) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 142 (62.6) | 152 (61.8) |
| | 95% CI | (56.3, 68.9) | (55.7, 67.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0124 | |
| | 95% CI | (0.8796, 1.1653) | |
| | Two-sided P-value | 0.8636 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| Age (years) group: >=40 | Baseline, N | 130 | 118 |
| | Week 26, N | 130 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 116 (89.2) | 106 (89.8) |
| | Number of Subjects with NRI, N2 (%) | 14 (10.8) | 12 (10.2) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 99 (76.2) | 77 (65.3) |
| | 95% CI | (68.8, 83.5) | (56.7, 73.8) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1670 | |
| | 95% CI | (0.9914, 1.3737) | |
| | Two-sided P-value | 0.0634 | |
| | | | |
| | P-value of interaction | 0.1958 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 191 | 204 |
| | Week 26, N | 191 | 204 |
| | Number of Subjects with observed Case, N1 (%) | 174 (91.1) | 183 (89.7) |
| | Number of Subjects with NRI, N2 (%) | 17 (8.9) | 21 (10.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 127 (66.5) | 119 (58.3) |
| | 95% CI | (59.8, 73.2) | (51.6, 65.1) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1399 | |
| | 95% CI | (0.9776, 1.3291) | |
| | Two-sided P-value | 0.0948 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| Sex: Female | Baseline, N | 166 | 160 |
| | Week 26, N | 166 | 160 |
| | Number of Subjects with observed Case, N1 (%) | 137 (82.5) | 144 (90.0) |
| | Number of Subjects with NRI, N2 (%) | 29 (17.5) | 16 (10.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 114 (68.7) | 110 (68.8) |
| | 95% CI | (61.6, 75.7) | (61.6, 75.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9989 | |
| | 95% CI | (0.8628, 1.1565) | |
| | Two-sided P-value | 0.9883 | |
| | | | |
| | P-value of interaction | 0.2229 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 174 | 195 |
| | Week 26, N | 174 | 195 |
| | Number of Subjects with observed Case, N1 (%) | 147 (84.5) | 172 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 27 (15.5) | 23 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 122 (70.1) | 120 (61.5) |
| | 95% CI | (63.3, 76.9) | (54.7, 68.4) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1394 | |
| | 95% CI | (0.9832, 1.3203) | |
| | Two-sided P-value | 0.0827 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Baseline, N | 148 | 131 |
| | Week 26, N | 148 | 131 |
| | Number of Subjects with observed Case, N1 (%) | 133 (89.9) | 122 (93.1) |
| | Number of Subjects with NRI, N2 (%) | 15 (10.1) | 9 (6.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 99 (66.9) | 84 (64.1) |
| | 95% CI | (59.3, 74.5) | (55.9, 72.3) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0432 | |
| | 95% CI | (0.8792, 1.2378) | |
| | Two-sided P-value | 0.6280 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 16 (94.1) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 1 (5.9) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Asia | Responders, n (%) | 7 (41.2) | 14 (73.7) |
| | 95% CI | (17.8, 64.6) | (53.9, 93.5) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.5588 | |
| | 95% CI | (0.2981, 1.0477) | |
| | Two-sided P-value | 0.0696 | |
| | | | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| | Week 26, N | 18 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (83.3) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 3 (16.7) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 13 (72.2) | 11 (57.9) |
| | 95% CI | (51.5, 92.9) | (35.7, 80.1) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Region

| | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|-------------------------|------------------------|
| Abrocitinib vs Dupilumab Response Ratio | | |
| Estimate | 1.2475 | |
| 95% CI | (0.7729, 2.0133) | |
| Two-sided P-value | 0.3653 | |
| | | |
| P-value of interaction | 0.1556 | |
| | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------------|--|-------------------------|------------------------|
| | | | |
| aseline disease severity: Moderate | Baseline, N | 213 | 219 |
| | Week 26, N | 213 | 219 |
| | Number of Subjects with observed Case, N1 (%) | 182 (85.4) | 196 (89.5) |
| | Number of Subjects with NRI, N2 (%) | 31 (14.6) | 23 (10.5) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 139 (65.3) | 132 (60.3) |
| | 95% CI | (58.9, 71.7) | (53.8, 66.8) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0827 | |
| | 95% CI | (0.9361, 1.2522) | |
| | Two-sided P-value | 0.2844 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|--|-------------------------|------------------------|
| Baseline disease severity: Severe | Baseline, N | 144 | 145 |
| | Week 26, N | 144 | 145 |
| | Number of Subjects with observed Case, N1 (%) | 129 (89.6) | 131 (90.3) |
| | Number of Subjects with NRI, N2 (%) | 15 (10.4) | 14 (9.7) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 102 (70.8) | 97 (66.9) |
| | 95% CI | (63.4, 78.3) | (59.2, 74.6) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0588 | |
| | 95% CI | (0.9066, 1.2366) | |
| | Two-sided P-value | 0.4703 | |
| | | | |
| | P-value of interaction | 0.8374 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 09SEP2021 (22:40)

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|--|-----------|---|-------------|----------------------------------|----------------------|---|---------------------------|----------------------------|--------------------------|
| | | | | | | | | CMH or Logistic I | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | , | | , , | , , | • | , | | |
| Week 26 | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 139 (38.4%) (33.4%, 43.6%) | 365 | 114 (31.2%) (26.5%, 36.3%) | 1.37 (1.01, 1.86) | 1.23 (1.01, 1.50) | 7.2% (0.3%, 14.1%) | 0.0425+* | |
| Baseline Disease Severity Week 26 | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 82 (38.0%) (31.5%, 44.8%) | 220 | 73 (33.2%) (27.0%, 39.8%) | 1.23 (0.83, 1.83) | 1.14 (0.89, 1.47) | 4.8% (-4.2%, 13.8%) | 0.3178 | 0.4227 |
| Severe baseline disease (IGA=4) | 146 | 57 (39.0%) (31.1%, 47.5%) | 145 | 41 (28.3%) (21.1%, 36.3%) | 1.62 (0.99, 2.65) | 1.38 (0.99, 1.92) | 10.8% (-0.0%, 21.6%) | 0.0628 | |
| Gender Week 26 | | ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | | , | | | | | |
| Male | 193 | 73 (37.8%) (31.0%, 45.1%) | 204 | 53 (26.0%) (20.1%, 32.6%) | 1.73 (1.13, 2.66) | 1.46 (1.09, 1.95) | 11.8% (2.7%, 21.0%) | 0.0131* | 0.1628 |
| Female | 169 | 66 (39.1%) (31.7%, 46.8%) | 161 | 61 (37.9%) (30.4%, 45.9%) | 1.05 (0.67, 1.64) | 1.03 (0.78, 1.35) | 1.2% (-9.3%, 11.7%) | 0.9099 | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|---------------------|-----------|-----------------|-------------|------------|--------------|---------------------|---------------------------|-------------------|---------------|
| | | | | | | | | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 26 | | | | | | | | | |
| US/Canada/Australia | 177 | 70 (39.5%) | 195 | 59 (30.3%) | 1.51 | 1.31 | 9.3% | 0.0644 | 0.3786 |
| | | (32.3%, | | (23.9%, | (0.98, 2.32) | (0.99, 1.73) | (-0.4%, 19.0%) | | |
| | | 47.2%) | | 37.2%) | | , | | | |
| Europe | 150 | 61 (40.7%) | 132 | 43 (32.6%) | 1.42 | 1.25 | 8.1% | 0.1749 | |
| • | | (32.7%, | | (24.7%, | (0.87, 2.31) | (0.91, 1.71) | (-3.1%, 19.3%) | | |
| | | 49.0%) | | 41.3%) | | | , | | |
| Asia | 17 | 3 (17.6%) | 19 | 6 (31.6%) | 0.46 | 0.56 | -13.9% | 0.4513 | |
| | | (3.8%, | | (12.6%, | (0.10, 2.25) | (0.16, 1.90) | (-41.6%, 13.7%) | | |
| | | 43.4%) | | 56.6%) | | ` ' ' | , | | |
| Latin America | 18 | 5 (27.8%) | 19 | 6 (31.6%) | 0.83 | 0.88 | -3.8% | 1.0000 | |
| | | (9.7%, | | (12.6%, | (0.20, 3.43) | (0.32, 2.38) | (-33.2%, 25.6%) | | |
| | | 53.5%) | | 56.6%) | | ` ' ' | , | | |
| Age Subgroup | | , | | , | | | | | |
| Week 26 | | | | | | | | | |
| <40 years | 230 | 79 (34.3%) | 247 | 80 (32.4%) | 1.09 | 1.06 | 2.0% | 0.6977 | 0.0437* |
| , | | (28.2%, | | (26.6%, | (0.75, 1.60) | (0.82, 1.37) | (-6.5%, 10.4%) | | |
| | | 40.9%) | | 38.6%) | | ` ' ' | , , , | | |
| >=40 years | 132 | 60 (45.5%) | 118 | 34 (28.8%) | 2.06 | 1.58 | 16.6% | 0.0088* | |
| - 5 | | (36.8%, | | (20.8%, | (1.22, 3.48) | (1.12, 2.22) | (4.9%, 28.4%) | ***** | |
| | | 54.3%) | | 37.9%) | (, 5110) | (===, = :==) | (112,73, 2011,70) | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 227 | 243 |
| | Week 26, N | 227 | 243 |
| | Number of Subjects with observed Case, N1 (%) | 184 (81.1) | 216 (88.9) |
| | Number of Subjects with NRI, N2 (%) | 43 (18.9) | 27 (11.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 75 (33.0) | 77 (31.7) |
| | 95% CI | (26.9, 39.2) | (25.8, 37.5) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0427 | |
| | 95% CI | (0.8028, 1.3543) | |
| | Two-sided P-value | 0.7541 | |
| | | | - |
| Age (years) group: >=40 | Baseline, N | 131 | 118 |
| | Week 26, N | 131 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 116 (88.5) | 105 (89.0) |
| | Number of Subjects with NRI, N2 (%) | 15 (11.5) | 13 (11.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 62 (47.3) | 37 (31.4) |
| | 95% CI | (38.8, 55.9) | (23.0, 39.7) |
| | Aboration of Deniloonal Barrage Batis | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.5094 | |
| | 95% CI | (1.0935, 2.0835) | |
| | Two-sided P-value | 0.0123 | |
| | | | |
| | P-value of interaction | 0.0807 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 191 | 201 |
| | Week 26, N | 191 | 201 |
| | Number of Subjects with observed Case, N1 (%) | 168 (88.0) | 178 (88.6) |
| | Number of Subjects with NRI, N2 (%) | 23 (12.0) | 23 (11.4) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 70 (36.6) | 64 (31.8) |
| | 95% CI | (29.8, 43.5) | (25.4, 38.3) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1510 | |
| | 95% CI | (0.8742, 1.5155) | |
| | Two-sided P-value | 0.3163 | |
| | | | |
| Sex: Female | Baseline, N | 167 | 160 |
| | Week 26, N | 167 | 160 |
| | Number of Subjects with observed Case, N1 (%) | 132 (79.0) | 143 (89.4) |
| | Number of Subjects with NRI, N2 (%) | 35 (21.0) | 17 (10.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 67 (40.1) | 50 (31.3) |
| | 95% CI | (32.7, 47.6) | (24.1, 38.4) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2838 | |
| | 95% CI | (0.9556, 1.7247) | |
| | Two-sided P-value | 0.0972 | |
| | | | |
| | P-value of interaction | 0.5958 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 174 | 191 |
| | Week 26, N | 174 | 191 |
| | Number of Subjects with observed Case, N1 (%) | 142 (81.6) | 166 (86.9) |
| | Number of Subjects with NRI, N2 (%) | 32 (18.4) | 25 (13.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 68 (39.1) | 62 (32.5) |
| | 95% CI | (31.8, 46.3) | (25.8, 39.1) |
| | A breeitinib ve Dunihumah Despense Batio | | |
| | Abrocitinib vs Dupilumab Response Ratio | 1 2020 | |
| | Estimate | 1.2039 | |
| | 95% CI | (0.9134, 1.5868) | |
| | Two-sided P-value | 0.1878 | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| tegion of emornion. Europe | Week 26, N | 150 | 132 |
| | Number of Subjects with observed Case, N1 (%) | 129 (86.0) | 122 (92.4) |
| | Number of Subjects with NRI, N2 (%) | 21 (14.0) | 10 (7.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 56 (37.3) | 40 (30.3) |
| | 95% CI | (29.6, 45.1) | (22.5, 38.1) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2320 | |
| | 95% CI | (0.8843, 1.7163) | |
| | Two-sided P-value | 0.2174 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 16 | 19 |
| | Week 26, N | 16 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (93.8) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 1 (6.3) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 7 (43.8) | 5 (26.3) |
| | 95% CI | (19.4, 68.1) | (6.5, 46.1) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.6625 | |
| | 95% CI | (0.6525, 4.2360) | |
| | Two-sided P-value | 0.2868 | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| 9 | Week 26, N | 18 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 14 (77.8) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (22.2) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 6 (33.3) | 7 (36.8) |
| | 95% CI | (11.6, 55.1) | (15.2, 58.5) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9048 | |
| | 95% CI | (0.3755, 2.1801) | |
| | Two-sided P-value | 0.8235 | |
| | P-value of interaction | 0.8316 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 212 | 217 |
| | Week 26, N | 212 | 217 |
| | Number of Subjects with observed Case, N1 (%) | 175 (82.5) | 192 (88.5) |
| | Number of Subjects with NRI, N2 (%) | 37 (17.5) | 25 (11.5) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 81 (38.2) | 72 (33.2) |
| | 95% CI | (31.7, 44.7) | (26.9, 39.4) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1515 | |
| | 95% CI | (0.8925, 1.4858) | |
| | Two-sided P-value | 0.2779 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 146 | 144 |
| | Week 26, N | 146 | 144 |
| | Number of Subjects with observed Case, N1 (%) | 125 (85.6) | 129 (89.6) |
| | Number of Subjects with NRI, N2 (%) | 21 (14.4) | 15 (10.4) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 56 (38.4) | 42 (29.2) |
| | 95% CI | (30.5, 46.2) | (21.7, 36.6) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.3151 | |
| | 95% CI | (0.9481, 1.8241) | |
| | Two-sided P-value | 0.1009 | |
| | | | |
| | P-value of interaction | 0.5303 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; DLQI = Dermatology Life Quality Index; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated by using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 13SEP2021 (02:03)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 226 | 246 |
| | Week 26, N | 226 | 246 |
| | Number of Subjects with observed Case, N1 (%) | 182 (80.5) | 217 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 44 (19.5) | 29 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 56 (24.8) | 47 (19.1) |
| | 95% CI | (19.2, 30.4) | (14.2, 24.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2969 | |
| | 95% CI | (0.9203, 1.8278) | |
| | Two-sided P-value | 0.1375 | |
| | | | |
| Age (years) group: >=40 | Baseline, N | 132 | 117 |
| | Week 26, N | 132 | 117 |
| | Number of Subjects with observed Case, N1 (%) | 117 (88.6) | 103 (88.0) |
| | Number of Subjects with NRI, N2 (%) | 15 (11.4) | 14 (12.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 50 (37.9) | 22 (18.8) |
| | 95% CI | (29.6, 46.2) | (11.7, 25.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 2.0145 | |
| | 95% CI | (1.3035, 3.1133) | |
| | Two-sided P-value | 0.0016 | |
| | | | |
| | P-value of interaction | 0.1194 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 190 | 202 |
| | Week 26, N | 190 | 202 |
| | Number of Subjects with observed Case, N1 (%) | 167 (87.9) | 178 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 23 (12.1) | 24 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 54 (28.4) | 34 (16.8) |
| | 95% CI | (22.0, 34.8) | (11.7, 22.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.6885 | |
| | 95% CI | (1.1540, 2.4707) | |
| | Two-sided P-value | 0.0070 | |
| | | | |
| Sex: Female | Baseline, N | 168 | 161 |
| | Week 26, N | 168 | 161 |
| | Number of Subjects with observed Case, N1 (%) | 132 (78.6) | 142 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 36 (21.4) | 19 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 52 (31.0) | 35 (21.7) |
| | 95% CI | (24.0, 37.9) | (15.4, 28.1) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.4238 | |
| | 95% CI | (0.9835, 2.0613) | |
| | Two-sided P-value | 0.0613 | |
| | | | |
| | P-value of interaction | 0.5289 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 177 | 193 |
| | Week 26, N | 177 | 193 |
| | Number of Subjects with observed Case, N1 (%) | 144 (81.4) | 166 (86.0) |
| | Number of Subjects with NRI, N2 (%) | 33 (18.6) | 27 (14.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 51 (28.8) | 37 (19.2) |
| | 95% CI | (22.1, 35.5) | (13.6, 24.7) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.5030 | |
| | 95% CI | (1.0373, 2.1778) | |
| | Two-sided P-value | 0.0313 | |
| | | | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| | Week 26, N | 150 | 132 |
| | Number of Subjects with observed Case, N1 (%) | 129 (86.0) | 121 (91.7) |
| | Number of Subjects with NRI, N2 (%) | 21 (14.0) | 11 (8.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 45 (30.0) | 23 (17.4) |
| | 95% CI | (22.7, 37.3) | (11.0, 23.9) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.7217 | |
| | 95% CI | (1.1038, 2.6857) | |
| | Two-sided P-value | 0.0166 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 14 | 19 |
| | Week 26, N | 14 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 13 (92.9) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 1 (7.1) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 4 (28.6) | 6 (31.6) |
| | 95% CI | (4.9, 52.2) | (10.7, 52.5) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9048 | |
| | 95% CI | (0.3134, 2.6120) | |
| | Two-sided P-value | 0.8532 | |
| Region of enrollment: Latin America | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 13 (76.5) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (23.5) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 6 (35.3) | 3 (15.8) |
| | 95% CI | (12.6, 58.0) | (0.0, 32.2) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 2.2353 | |
| | 95% CI | (0.6588, 7.5843) | |
| | Two-sided P-value | 0.1969 | |
| | P-value of interaction | 0.6651 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 213 | 219 |
| | Week 26, N | 213 | 219 |
| | Number of Subjects with observed Case, N1 (%) | 175 (82.2) | 193 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 38 (17.8) | 26 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 65 (30.5) | 41 (18.7) |
| | 95% CI | (24.3, 36.7) | (13.6, 23.9) |
| | | | - |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.6300 | |
| | 95% CI | (1.1575, 2.2955) | |
| | Two-sided P-value | 0.0052 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 145 | 144 |
| | Week 26, N | 145 | 144 |
| | Number of Subjects with observed Case, N1 (%) | 124 (85.5) | 127 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 21 (14.5) | 17 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 41 (28.3) | 28 (19.4) |
| | 95% CI | (20.9, 35.6) | (13.0, 25.9) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.4542 | |
| | 95% CI | (0.9540, 2.2167) | |
| | Two-sided P-value | 0.0817 | |
| | P-value of interaction | 0.6804 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; POEM = patient-oriented eczema measure; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 15SEP2021 (02:11)

Binary Outcome Analysis: Achieving 0 in POEM total score at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|--|-----------|---------------------------------|------------|-------------------------------|----------------------|----------------------|---------------------------|----------------------------|--------------------------|
| | | | | _ | | | _ | CMH or Logistic F | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | | | | , | , , | , | | |
| Week 26 | | | | | | | | | |
| Full Analysis Set Safety Population | 359 | 49 (13.6%) (10.3%, 17.6%) | 365 | 26 (7.1%) (4.7%, 10.3%) | 2.06 (1.25, 3.40) | 1.92 (1.22, 3.01) | 6.5% (2.1%, 10.9%) | 0.0040+* | |
| Baseline Disease Severity Week 26 | | , | | , | | | | | |
| Moderate baseline disease (IGA=3) | 214 | 29 (13.6%) (9.3%, 18.9%) | 220 | 15 (6.8%) (3.9%, 11.0%) | 2.14 (1.11, 4.12) | 1.99 (1.10, 3.60) | 6.7% (1.1%, 12.4%) | 0.0254* | 0.9166 |
| Severe baseline disease (IGA=4) | 145 | 20 (13.8%) (8.6%, 20.5%) | 145 | 11 (7.6%) (3.8%, 13.2%) | 1.95 (0.90, 4.23) | 1.82 (0.90, 3.66) | 6.2% (-0.9%, 13.3%) | 0.1274 | |
| Gender Week 26 | | , | | , | | | | | |
| Male | 191 | 26 (13.6%) (9.1%, 19.3%) | 204 | 11 (5.4%) (2.7%, 9.4%) | 2.76 (1.33, 5.77) | 2.52 (1.28, 4.97) | 8.2% (2.5%, 14.0%) | 0.0056* | 0.4235 |
| Female | 168 | 23 (13.7%) (8.9%, 19.8%) | 161 | 15 (9.3%) (5.3%, 14.9%) | 1.54 (0.77, 3.08) | 1.47 (0.80, 2.71) | 4.4% (-2.5%, 11.2%) | 0.2314 | |

Notes:

Number of subjects: Full Analysis Set Safety Population Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0 in POEM total score at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitini | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|---------------------|------------|-----------------|-------------|------------|---------------|-----------------|---------------------------|-------------------|---------------|
| | | • | | - | | - | | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 26 | | | | | | | | | |
| US/Canada/Australia | 177 | 27 (15.3%) | 195 | 16 (8.2%) | 2.01 | 1.86 | 7.0% | 0.0360* | 0.9211 |
| | | (10.3%, | | (4.8%, | (1.05, 3.88) | (1.04, 3.33) | (0.5%, 13.6%) | | |
| | | 21.4%) | | 13.0%) | ` ' ' | | | | |
| Europe | 150 | 17 (11.3%) | 132 | 7 (5.3%) | 2.28 | 2.14 | 6.0% | 0.0874 | |
| 1 | | (6.7%, | | (2.2%, | (0.92, 5.69) | (0.91, 4.99) | (-0.3%, 12.4%) | | |
| | | 17.5%) | | 10.6%) | ` ' ' | | , , , | | |
| Asia | 15 | 2 (13.3%) | 19 | 2 (10.5%) | 1.31 | 1.27 | 2.8% | 1.0000 | |
| | | (1.7%, | | (1.3%, | (0.16, 10.56) | (0.20, 7.97) | (-19.2%, 24.9%) | | |
| | | 40.5%) | | 33.1%) | (3, 3, 3, 3, | (, , , , , , , | (1 : 11, 11 : 11, | | |
| Latin America | 17 | 3 (17.6%) | 19 | 1 (5.3%) | 3.86 | 3.35 | 12.4% | 0.3255 | |
| | | (3.8%, | | (0.1%, | (0.36, 41.20) | (0.38, 29.26) | (-8.3%, 33.1%) | | |
| | | 43.4%) | | 26.0%) | , , | , , , | , | | |
| Age Subgroup | | , | | , | | | | | |
| Week 26 | | | | | | | | | |
| <40 years | 227 | 22 (9.7%) | 247 | 19 (7.7%) | 1.29 | 1.26 | 2.0% | 0.5137 | 0.0094* |
| | | (6.2%, | | (4.7%, | (0.68, 2.45) | (0.70, 2.27) | (-3.1%, 7.1%) | | |
| | | 14.3%) | | 11.8%) | (3323, 337) | (, | (====,===, | | |
| >=40 years | 132 | 27 (20.5%) | 118 | 7 (5.9%) | 4.08 | 3.45 | 14.5% | 0.0008* | |
| | 102 | (13.9%, | | (2.4%, | (1.70, 9.76) | (1.56, 7.62) | (6.4%, 22.6%) | 0.0000 | |
| | | 28.3%) | | 11.8%) | (10,)0) | (1.53, 7.02) | (0.173, 22.070) | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

CMH: Cochran-Mantel-Haenszel; NE: not estimable; OR: odds ratio; RD: Risk Difference; RR: Relative Risk.

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 230 | 247 |
| | Week 26, N | 230 | 247 |
| | Number of Subjects with observed Case, N1 (%) | 183 (79.6) | 219 (88.7) |
| | Number of Subjects with NRI, N2 (%) | 47 (20.4) | 28 (11.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 89 (38.7) | 91 (36.8) |
| | 95% CI | (32.4, 45.0) | (30.8, 42.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.0503 | |
| | 95% CI | (0.8341, 1.3226) | |
| | Two-sided P-value | 0.6764 | |
| | | | |
| Age (years) group: >=40 | Baseline, N | 132 | 118 |
| | Week 26, N | 132 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 117 (88.6) | 104 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 15 (11.4) | 14 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 63 (47.7) | 42 (35.6) |
| | 95% CI | (39.2, 56.2) | (27.0, 44.2) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.3409 | |
| | 95% CI | (0.9921, 1.8124) | |
| | Two-sided P-value | 0.0564 | |
| | | | |
| | P-value of interaction | 0.2069 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 193 | 204 |
| | Week 26, N | 193 | 204 |
| | Number of Subjects with observed Case, N1 (%) | 168 (87.0) | 180 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 25 (13.0) | 24 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 77 (39.9) | 61 (29.9) |
| | 95% CI | (33.0, 46.8) | (23.6, 36.2) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.3342 | |
| | 95% CI | (1.0162, 1.7518) | |
| | Two-sided P-value | 0.0379 | |
| | | | |
| Sex: Female | Baseline, N | 169 | 161 |
| | Week 26, N | 169 | 161 |
| | Number of Subjects with observed Case, N1 (%) | 132 (78.1) | 143 (88.8) |
| | Number of Subjects with NRI, N2 (%) | 37 (21.9) | 18 (11.2) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 75 (44.4) | 72 (44.7) |
| | 95% CI | (36.9, 51.9) | (37.0, 52.4) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9924 | |
| | 95% CI | (0.7800, 1.2625) | |
| | Two-sided P-value | 0.9502 | |
| | | | |
| | P-value of interaction | 0.1104 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 177 | 195 |
| | Week 26, N | 177 | 195 |
| | Number of Subjects with observed Case, N1 (%) | 141 (79.7) | 168 (86.2) |
| | Number of Subjects with NRI, N2 (%) | 36 (20.3) | 27 (13.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 75 (42.4) | 65 (33.3) |
| | 95% CI | (35.1, 49.7) | (26.7, 39.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2712 | |
| | 95% CI | (0.9777, 1.6528) | |
| | Two-sided P-value | 0.0732 | |
| | | | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| | Week 26, N | 150 | 132 |
| | Number of Subjects with observed Case, N1 (%) | 130 (86.7) | 122 (92.4) |
| | Number of Subjects with NRI, N2 (%) | 20 (13.3) | 10 (7.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 61 (40.7) | 47 (35.6) |
| | 95% CI | (32.8, 48.5) | (27.4, 43.8) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.1421 | |
| | 95% CI | (0.8461, 1.5417) | |
| | Two-sided P-value | 0.3853 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (88.2) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 2 (11.8) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 4 (23.5) | 9 (47.4) |
| | 95% CI | (3.4, 43.7) | (24.9, 69.8) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.4967 | |
| | 95% CI | (0.1866, 1.3226) | |
| | Two-sided P-value | 0.1614 | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| Region of enforment: Laun America | Week 26, N | 18 | 19 |
| | | | |
| | Number of Subjects with observed Case, N1 (%) | 14 (77.8) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (22.2) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 12 (66.7) | 12 (63.2) |
| | 95% CI | (44.9, 88.4) | (41.5, 84.8) |
| | Abrocitinib vs Dupilumab Response Ratio | I | |
| | Estimate Estimate | 1.0556 | |
| | 95% CI | (0.6571, 1.6956) | |
| | Two-sided P-value | 0.8231 | |
| | | | |
| | P-value of interaction | 0.3197 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 216 | 220 |
| | Week 26, N | 216 | 220 |
| | Number of Subjects with observed Case, N1 (%) | 176 (81.5) | 194 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 40 (18.5) | 26 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 92 (42.6) | 71 (32.3) |
| | 95% CI | (36.0, 49.2) | (26.1, 38.5) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.3198 | |
| | 95% CI | (1.0318, 1.6882) | |
| | Two-sided P-value | 0.0272 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 146 | 145 |
| | Week 26, N | 146 | 145 |
| | Number of Subjects with observed Case, N1 (%) | 124 (84.9) | 129 (89.0) |
| | Number of Subjects with NRI, N2 (%) | 22 (15.1) | 16 (11.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 60 (41.1) | 62 (42.8) |
| | 95% CI | (33.1, 49.1) | (34.7, 50.8) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.9611 | |
| | 95% CI | (0.7333, 1.2597) | |
| | Two-sided P-value | 0.7738 | |
| | | | |
| | P-value of interaction | 0.0893 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|--|-------------------------|------------------------|
| | | | |
| Age (years) group: <40 | Baseline, N | 230 | 247 |
| | Week 26, N | 230 | 247 |
| | Number of Subjects with observed Case, N1 (%) | 183 (79.6) | 219 (88.7) |
| | Number of Subjects with NRI, N2 (%) | 47 (20.4) | 28 (11.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 42 (18.3) | 32 (13.0) |
| | 95% CI | (13.3, 23.3) | (8.8, 17.1) |
| | | | _ |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.4095 | |
| | 95% CI | (0.9230, 2.1525) | |
| | Two-sided P-value | 0.1121 | |
| | | | - |
| Age (years) group: >=40 | Baseline, N | 132 | 118 |
| | Week 26, N | 132 | 118 |
| | Number of Subjects with observed Case, N1 (%) | 117 (88.6) | 104 (88.1) |
| | Number of Subjects with NRI, N2 (%) | 15 (11.4) | 14 (11.9) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Age group (<40, >=40)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|---|-------------------------|------------------------|
| Age (years) group: >=40 | Responders, n (%) | 38 (28.8) | 20 (16.9) |
| | 95% CI | (21.1, 36.5) | (10.2, 23.7) |
| | Ahmasitinih wa Dunilumah Dasmansa Datia | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.6985 | |
| | 95% CI | (1.0498, 2.7480) | |
| | Two-sided P-value | 0.0309 | |
| | | | |
| | P-value of interaction | 0.5685 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|--|-------------------------|------------------------|
| | | | |
| Sex: Male | Baseline, N | 193 | 204 |
| | Week 26, N | 193 | 204 |
| | Number of Subjects with observed Case, N1 (%) | 168 (87.0) | 180 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 25 (13.0) | 24 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 43 (22.3) | 23 (11.3) |
| | 95% CI | (16.4, 28.2) | (6.9, 15.6) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.9761 | |
| | 95% CI | (1.2394, 3.1507) | |
| | Two-sided P-value | 0.0042 | |
| | | | |
| Sex: Female | Baseline, N | 169 | 161 |
| | Week 26, N | 169 | 161 |
| | Number of Subjects with observed Case, N1 (%) | 132 (78.1) | 143 (88.8) |
| | Number of Subjects with NRI, N2 (%) | 37 (21.9) | 18 (11.2) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Sex

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------|---|-------------------------|------------------------|
| Sex: Female | Responders, n (%) | 37 (21.9) | 29 (18.0) |
| | 95% CI | (15.7, 28.1) | (12.1, 23.9) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2155 | |
| | 95% CI | (0.7863, 1.8789) | |
| | Two-sided P-value | 0.3799 | |
| | | | |
| | P-value of interaction | 0.1356 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---|--|-------------------------|------------------------|
| | | | |
| Region of enrollment: US/Canada/Australia | Baseline, N | 177 | 195 |
| | Week 26, N | 177 | 195 |
| | Number of Subjects with observed Case, N1 (%) | 141 (79.7) | 168 (86.2) |
| | Number of Subjects with NRI, N2 (%) | 36 (20.3) | 27 (13.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 44 (24.9) | 22 (11.3) |
| | 95% CI | (18.5, 31.2) | (6.8, 15.7) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 2.2034 | |
| | 95% CI | (1.3777, 3.5240) | |
| | Two-sided P-value | 0.0010 | |
| | | | |
| Region of enrollment: Europe | Baseline, N | 150 | 132 |
| | Week 26, N | 150 | 132 |
| | Number of Subjects with observed Case, N1 (%) | 130 (86.7) | 122 (92.4) |
| | Number of Subjects with NRI, N2 (%) | 20 (13.3) | 10 (7.6) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|------------------------------|--|-------------------------|------------------------|
| Region of enrollment: Europe | Responders, n (%) | 27 (18.0) | 19 (14.4) |
| | 95% CI | (11.9, 24.1) | (8.4, 20.4) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2505 | |
| | 95% CI | (0.7300, 2.1422) | |
| | Two-sided P-value | 0.4156 | |
| | | | |
| Region of enrollment: Asia | Baseline, N | 17 | 19 |
| | Week 26, N | 17 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 15 (88.2) | 18 (94.7) |
| | Number of Subjects with NRI, N2 (%) | 2 (11.8) | 1 (5.3) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 1 (5.9) | 4 (21.1) |
| | 95% CI | (0.0, 17.1) | (2.7, 39.4) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Region

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 0.2794 | |
| | 95% CI | (0.0345, 2.2620) | |
| | Two-sided P-value | 0.2321 | |
| Region of enrollment: Latin America | Baseline, N | 18 | 19 |
| | Week 26, N | 18 | 19 |
| | Number of Subjects with observed Case, N1 (%) | 14 (77.8) | 15 (78.9) |
| | Number of Subjects with NRI, N2 (%) | 4 (22.2) | 4 (21.1) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 8 (44.4) | 7 (36.8) |
| | 95% CI | (21.5, 67.4) | (15.2, 58.5) |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.2063 | |
| | 95% CI | (0.5512, 2.6400) | |
| | Two-sided P-value | 0.6387 | |
| | P-value of interaction | 0.1251 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------------------|--|-------------------------|------------------------|
| | | | |
| Baseline disease severity: Moderate | Baseline, N | 216 | 220 |
| | Week 26, N | 216 | 220 |
| | Number of Subjects with observed Case, N1 (%) | 176 (81.5) | 194 (88.2) |
| | Number of Subjects with NRI, N2 (%) | 40 (18.5) | 26 (11.8) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |
| | Responders, n (%) | 50 (23.1) | 30 (13.6) |
| | 95% CI | (17.5, 28.8) | (9.1, 18.2) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.6975 | |
| | 95% CI | (1.1245, 2.5626) | |
| | Two-sided P-value | 0.0118 | |
| | | | |
| Baseline disease severity: Severe | Baseline, N | 146 | 145 |
| | Week 26, N | 146 | 145 |
| | Number of Subjects with observed Case, N1 (%) | 124 (84.9) | 129 (89.0) |
| | Number of Subjects with NRI, N2 (%) | 22 (15.1) | 16 (11.0) |
| | Number of Subjects Missing Cases without NRI, N3 (%) | 0 | 0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Baseline disease severity

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-----------------------------------|---|-------------------------|------------------------|
| Baseline disease severity: Severe | Responders, n (%) | 30 (20.5) | 22 (15.2) |
| | 95% CI | (14.0, 27.1) | (9.3, 21.0) |
| | | | |
| | Abrocitinib vs Dupilumab Response Ratio | | |
| | Estimate | 1.3543 | |
| | 95% CI | (0.8215, 2.2325) | |
| | Two-sided P-value | 0.2344 | |
| | | | |
| | P-value of interaction | 0.4942 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; CMH = Cochran-Mantel-Haenszel; SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set at the specified visit; n (%) = number of subjects with response (percentage based on N); NRI = non-responder imputation.

CI for the response rate was based on normal approximation (or the Clopper-Pearson exact method when there were 0 or 100% responders); Response ratio was calculated using wald method without stratification.

A log regression model was used to provide two sided Wald p-value for testing subgroup and treatment interaction. 0.5 was added to the empty cells.

P-value of interaction was based on a type3 Wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment * subgroup category) are zero.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 14SEP2021 (05:07)

Binary Outcome Analysis: SCORAD-100 response at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--|-----------|--------------------------------|------------|-------------------------------|----------------------|--|-----------------------|-------------------------------|--------------------------|--|
| | | | | | | | - | CMH or Logistic P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | (7570 CI) | | (5570 CI) | (3370 CI) | (55% C1) | (3370 CI) | varae [1] | meraction | |
| Week 26 | | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 37 (10.2%) (7.3%, 13.8%) | 365 | 22 (6.0%) (3.8%, 9.0%) | 1.77 (1.03, 3.07) | 1.70 (1.02, 2.82) | 4.2% (0.2%, 8.2%) | 0.0384+* | - | |
| Baseline Disease Severity Week 26 | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 25 (11.6%) (7.6%, 16.6%) | 220 | 12 (5.5%) (2.8%, 9.3%) | 2.27 (1.11, 4.64) | 2.12 (1.09, 4.11) | 6.1% (0.9%, 11.3%) | 0.0253* | 0.2397 | |
| Severe baseline disease (IGA=4) | 146 | 12 (8.2%) (4.3%, 13.9%) | 145 | 10 (6.9%) (3.4%, 12.3%) | 1.21 (0.51, 2.89) | 1.19 (0.53, 2.67) | 1.3% (-4.7%, 7.4%) | 0.8251 | | |
| Gender Week 26 | | , | | , | | | | | | |
| Male | 193 | 21 (10.9%) (6.9%, 16.2%) | 204 | 9 (4.4%) (2.0%, 8.2%) | 2.65 (1.18, 5.93) | 2.47 (1.16, 5.25) | 6.5% (1.2%, 11.7%) | 0.0213* | 0.2219 | |
| Female | 169 | 16 (9.5%) (5.5%, 14.9%) | 161 | 13 (8.1%) (4.4%, 13.4%) | 1.19 (0.55, 2.56) | 1.17 (0.58, 2.36) | 1.4% (-4.7%, 7.5%) | 0.7007 | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-100 response at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 200mg QD Dupilumab 300mg Q2W | | | | | | Dupilumab 300mg | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | |
|--|-----|------------|-----|--------------|---------------|-----------------|--|-------------------|---------------|
| | | • | | - | | - | | CMH or Logistic I | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 26 | | | | | | | | | |
| US/Canada/Australia | 177 | 15 (8.5%) | 195 | 7 (3.6%) | 2.49 | 2.36 | 4.9% | 0.0504 | 0.2424 |
| | | (4.8%, | | (1.5%, 7.3%) | (0.99, 6.25) | (0.99, 5.66) | (0.0%, 9.7%) | | |
| | | 13.6%) | | , , , | | | , | | |
| Europe | 150 | 17 (11.3%) | 132 | 9 (6.8%) | 1.75 | 1.66 | 4.5% | 0.2199 | |
| 1 | | (6.7%, | | (3.2%, | (0.75, 4.06) | (0.77, 3.60) | (-2.1%, 11.2%) | | |
| | | 17.5%) | | 12.5%) | | , , , | , | | |
| Asia | 17 | 0 (0.0%) | 19 | 3 (15.8%) | 0.13 | 0.16 | -14.7% | 0.2310 | |
| | | ` / | | (3.4%, | (0.01, 2.81) | (0.01, 2.87) | (-33.0%, 3.6%) | | |
| | | | | 39.6%) | | , , , | ` ' ' | | |
| Latin America | 18 | 5 (27.8%) | 19 | 3 (15.8%) | 2.05 | 1.76 | 12.0% | 0.4470 | |
| | | (9.7%, | | (3.4%, | (0.41, 10.24) | (0.49, 6.31) | (-14.4%, 38.4%) | | |
| | | 53.5%) | | 39.6%) | , | , , , | , , , | | |
| Age Subgroup | | , | | , | | | | | |
| Week 26 | | | | | | | | | |
| <40 years | 230 | 19 (8.3%) | 247 | 14 (5.7%) | 1.50 | 1.46 | 2.6% | 0.2836 | 0.3246 |
| · | | (5.0%, | | (3.1%, 9.3%) | (0.73, 3.06) | (0.75, 2.84) | (-2.0%, 7.2%) | | |
| | | 12.6%) | | , , , | | , , , | | | |
| >=40 years | 132 | 18 (13.6%) | 118 | 8 (6.8%) | 2.17 | 2.01 | 6.9% | 0.0969 | |
| | | (8.3%, | | (3.0%, | (0.91, 5.20) | (0.91, 4.45) | (-0.5%, 14.3%) | 2172 | |
| | | 20.7%) | | 12.9%) | (,) | (1.1.7) | (1.1.1.) | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitinib 200mg QD Dupilumab 300mg Q2W | | | | | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | |
|--|--|----------------------------------|-----|----------------------------------|----------------------|----------------------|--|----------------------------|--------------------------|
| | | | | | | | | CMH or Logistic I | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | , , | | , | , | , , | , | | |
| Week 26 | | | | | | | | | |
| Full Analysis Set Safety Population | 302 | 119 (39.4%) (33.9%, 45.2%) | 309 | 136 (44.0%) (38.4%, 49.7%) | 0.82 (0.60, 1.14) | 0.89 (0.74, 1.08) | -4.7% (-12.5%, 3.1%) | 0.2380+ | |
| Baseline Disease Severity Week 26 | | , | | , | | | | | |
| Moderate baseline disease (IGA=3) | 177 | 59 (33.3%) (26.4%, 40.8%) | 184 | 78 (42.4%) (35.2%, 49.9%) | 0.68 (0.44, 1.04) | 0.79 (0.60, 1.03) | -9.1% (-19.0%, 0.9%) | 0.0832 | 0.2250 |
| Severe baseline disease (IGA=4) | 125 | 60 (48.0%) (39.0%, 57.1%) | 125 | 58 (46.4%) (37.4%, 55.5%) | 1.07 (0.65, 1.75) | 1.03 (0.80, 1.34) | 1.6% (-10.8%, 14.0%) | 0.8992 | |
| Gender Week 26 | | , | | , | | | | | |
| Male | 161 | 57 (35.4%) (28.0%, 43.3%) | 172 | 65 (37.8%) (30.5%, 45.5%) | 0.90 (0.58, 1.41) | 0.94 (0.71, 1.24) | -2.4% (-12.7%, 8.0%) | 0.7329 | 0.4258 |
| Female | 141 | 62 (44.0%) (35.6%, 52.6%) | 137 | 71 (51.8%) (43.1%, 60.4%) | 0.73 (0.46, 1.17) | 0.85 (0.66, 1.09) | -7.9% (-19.6%, 3.9%) | 0.2298 | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 26 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| %) OR | - nn | | CMH or Logistic | P-val [2] for |
|---------------------------------------|----------------------------|---|--|---|
| %) OR | D.D. | | | , var [2] 101 |
| | RR | RD | Regression p- | trt*subgroup |
| (I) (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| | | | | |
| | | | | |
| %) 0.75 | 0.83 | -6.7% | 0.2731 | 0.6191 |
| (0.46, 1.21) | (0.61, 1.13) | (-17.7%, 4.2%) | | |
| | , , | , , , | | |
| | 0.94 | -3.0% | 0.7040 | |
| (0.54, 1.46) | (0.73, 1.22) | (-15.4%, 9.4%) | | |
| | , | (,, | | |
| | 0.53 | -22.4% | 0.2928 | |
| * | | | | |
| | (0.20, 11.0) | (== = , = , = = , = , | | |
| , | 1.21 | 9.2% | 0.7319 | |
| * | | | | |
| · · · · · · · · · · · · · · · · · · · | (0.03, 2.17) | (2.1070, 10.270) | | |
| , | | | | |
| | | | | |
| %) 0.88 | 0.93 | -3.1% | 0.5475 | 0.6106 |
| , | | | 0.0 1.75 | 0.0100 |
| | (0.7 1, 1.17) | (12.770, 0.170) | | |
| , | 0.83 | -7 5% | 0.3100 | |
| | | | 0.3199 | |
| | (0.00, 1.10) | (-21.0%, 0.0%) | | |
| | %) 0.75 %, (0.46, 1.21) | %) 0.75 0.83 6, (0.46, 1.21) (0.61, 1.13) 6) 0.89 0.94 6, (0.54, 1.46) (0.73, 1.22) 6) 0.37 0.53 6, (0.09, 1.57) (0.20, 1.40) 6) 1.45 1.21 6, (0.37, 5.70) (0.59, 2.47) 6) %) 0.88 0.93 6, (0.59, 1.30) (0.74, 1.17) 6) %) 0.73 0.83 6, (0.42, 1.28) (0.60, 1.16) | %) 0.75 0.83 -6.7% 6, (0.46, 1.21) (0.61, 1.13) (-17.7%, 4.2%) 6) 0.89 0.94 -3.0% 6, (0.54, 1.46) (0.73, 1.22) (-15.4%, 9.4%) 6) | %) 0.75 0.83 -6.7% 0.2731 6, (0.46, 1.21) (0.61, 1.13) (-17.7%, 4.2%) 6) 0.89 0.94 -3.0% 0.7040 6, (0.54, 1.46) (0.73, 1.22) (-15.4%, 9.4%) 6) 0.37 0.53 -22.4% 0.2928 6, (0.09, 1.57) (0.20, 1.40) (-53.3%, 8.5%) 6) 1.45 1.21 9.2% 0.7319 6, (0.37, 5.70) (0.59, 2.47) (-24.8%, 43.2%) 6) 76 0.88 0.93 -3.1% 0.5475 76, (0.59, 1.30) (0.74, 1.17) (-12.7%, 6.4%) 76 0.73 0.83 -7.5% 0.3199 76, (0.42, 1.28) (0.60, 1.16) (-21.0%, 6.0%) |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Total mortality at 26 weeks (TEAE leading to death) - Safety Set JADE DARE (PF-04965842) - 2023 datacut

No adverse events of this type occurred

| | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------|-----------------------------|--|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | | | |
| Overall | Overall | N | 362 | 365 |
| | | n (%) | 264 (72.9) | 237 (64.9) |
| | | 95% CI ^a | (68.35, 77.51) | (60.04, 69.83) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.12 (1.02, 1.24) | |
| | | P-value | 0.0203 | |
| | | - | | |
| | | Odds Ratio (95% CI) ^a | 1.45 (1.06, 2.00) | |
| | | P-value | 0.0201 | |
| | | | | |
| | | Risk Difference% (95% CI) ^a | 8.00 (1.29, 14.70) | |
| | | P-value | 0.0194 | |

MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 23SEP2021 (23:43)

Output File: ./nda1_cdisc/B7451050_GBA/adae_prop_2_e

Included data up to 28 days after last dose of study.
Subjects were counted only once per treatment per event.
.5 was added to 0 count cells when RR or OR can't be calculated.
a. Results calculated with normal approximation.

| | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------|-----------------------------|--|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | | | |
| Overall | Overall | N | 362 | 365 |
| | | n (%) | 6 (1.7) | 5 (1.4) |
| | | 95% CI ^a | (0.34, 2.97) | (0.18, 2.56) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.21 (0.37, 3.93) | |
| | | P-value | 0.7512 | |
| | | | | |
| | | Odds Ratio (95% CI) ^a | 1.21 (0.37, 4.01) | |
| | | P-value | 0.7511 | |
| | | | | |
| | | Risk Difference% (95% CI) ^a | 0.29 (-1.49, 2.06) | |
| | | P-value | 0.7509 | |

MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 23SEP2021 (23:43)

Output File: ./nda1_cdisc/B7451050_GBA/adae_prop_5_e

Included data up to 28 days after last dose of study.
Subjects were counted only once per treatment per event.
.5 was added to 0 count cells when RR or OR can't be calculated.
a. Results calculated with normal approximation.

| | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------|-----------------------------|--|---------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | | | |
| Overall | Overall | N | 362 | 365 |
| | | n (%) | 10 (2.8) | 8 (2.2) |
| | | 95% CI ^a | (1.07, 4.45) | (0.69, 3.69) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.26 (0.50, 3.16) | |
| | | P-value | 0.6214 | |
| | | | | |
| | | Odds Ratio (95% CI) ^a | 1.27 (0.49, 3.25) | |
| | | P-value | 0.6213 | |
| | | | | |
| | | Risk Difference% (95% CI) ^a | 0.57 (-1.69, 2.83) | |
| | | P-value | 0.6206 | |

MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 23SEP2021 (23:43)

Output File: ./nda1_cdisc/B7451050_GBA/adae_prop_6_e

Included data up to 28 days after last dose of study.
Subjects were counted only once per treatment per event.
.5 was added to 0 count cells when RR or OR can't be calculated.
a. Results calculated with normal approximation.

| | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------|-----------------------------|--|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | | | |
| Overall | Overall | N | 362 | 365 |
| | | n (%) | 267 (73.8) | 237 (64.9) |
| | | 95% CI ^a | (69.22, 78.29) | (60.04, 69.83) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.14 (1.03, 1.25) | |
| | | P-value | 0.0102 | |
| | | | | |
| | | Odds Ratio (95% CI) ^a | 1.52 (1.10, 2.09) | |
| | | P-value | 0.0101 | |
| | | | | |
| | | Risk Difference% (95% CI) ^a | 8.83 (2.15, 15.50) | |
| | | P-value | 0.0095 | |

MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 23SEP2021 (23:43)

Output File: ./nda1_cdisc/B7451050_GBA/adae_prop_10

Included data up to 28 days after last dose of study.
Subjects were counted only once per treatment per event.
.5 was added to 0 count cells when RR or OR can't be calculated.
a. Results calculated with normal approximation.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 174 (75.7) | 162 (65.6) |
| | | | 95% CI ^a | (70.11, 81.20) | (59.66, 71.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.15 (1.03, 1.30) | |
| | | | P-value | 0.0162 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 94 (71.2) | 77 (65.3) |
| | | | 95% CI ^a | (63.49, 78.94) | (56.66, 73.85) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.09 (0.92, 1.29) | |
| | | | P-value | 0.3154 | |
| | | | | | |
| | | | Test for interaction ^b | 0.5042 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 141 (73.1) | 128 (62.7) |
| | | | 95% CI ^a | (66.80, 79.32) | (56.11, 69.38) |
| | | | Relative Risk (95% CI) ^a | 1.16 (1.02, 1.33) | |
| | | | P-value | 0.0284 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 127 (75.1) | 111 (68.9) |
| | | | 95% CI ^a | (68.63, 81.66) | (61.80, 76.09) |
| | | | Relative Risk (95% CI) ^a | 1.00 (0.05, 1.25) | |
| | | | ` ' | 1.09 (0.95, 1.25) | |
| | | | P-value | 0.2115 | |
| | | | Test for interaction ^b | 0.6631 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------|----------------------------------|---|---|--|
| System Organ | MedDRA Preferred | | | |
| Class | Term | | n (%) | n (%) |
| | | | | |
| a Overall | Overall | N | 177 | 195 |
| | | n (%) | 133 (75.1) | 130 (66.7) |
| | | 95% CI ^a | (68.77, 81.51) | (60.05, 73.28) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.13 (0.99, 1.28) | |
| | | P-value | 0.0723 | |
| | | | | |
| Overall | l Overall | N | 150 | 132 |
| | | n (%) | 111 (74.0) | 90 (68.2) |
| | | 95% CI ^a | (66.98, 81.02) | (60.24, 76.13) |
| | | D 1 .: D: 1 (050) CD2 | | |
| | | | | |
| | | P-value | 0.2855 | |
| | 0 " | | 4.5 | 10 |
| Overall | Overall | | · | 19 |
| | | II. | | 10 (52.6) |
| | | 95% CI ^a | (48.93, 92.25) | (30.18, 75.08) |
| | | | | |
| | | Relative Risk (95% CI) ^a | 1.34 (0.79, 2.27) | |
| ii | Organ Class ia Overall Overall | System Organ Class Preferred Term ia Overall Overall Overall Overall Overall Overall | Organ Class Preferred Term ia Overall N n (%) 95% CI ^a Relative Risk (95% CI) ^a P-value Overall N n (%) 95% CI ^a Relative Risk (95% CI) ^a P-value P-value | System Organ Class MedDRA Preferred Term N 177 133 (75.1) 95% CI ^a (68.77, 81.51) Relative Risk (95% CI) ^a 1.13 (0.99, 1.28) P-value 0.0723 Overall Overall N 150 111 (74.0) 95% CI ^a (66.98, 81.02) Relative Risk (95% CI) ^a 1.09 (0.93, 1.26) P-value 0.2855 Overall Overall Overall N 17 12 (70.6) 12 (70.6) 14 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) 15 (70.6) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|---------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 12 (66.7) | 9 (47.4) |
| | | | 95% CI ^a | (44.89, 88.44) | (24.92, 69.82) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.41 (0.79, 2.50) | |
| | | | P-value | 0.2446 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9036 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| Subgroup | Ciass | TCIII | | 11 (70) | 11 (70) |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 162 (75.0) | 148 (67.3) |
| | | | 95% CI ^a | (69.23, 80.77) | (61.07, 73.47) |
| | | | Relative Risk (95% CI) ^a | 1.11 (0.99, 1.26) | |
| | | | P-value | 0.0760 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 106 (72.6) | 91 (62.8) |
| | | | 95% CI ^a | (65.37, 79.84) | (54.89, 70.63) |
| | | | Relative Risk (95% CI) ^a | 1.16 (0.00.1.26) | |
| | | | ` ' | 1.16 (0.99, 1.36) | |
| | | | P-value | 0.0746 | |
| | | | Test for interaction ^b | 0.8701 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| <u> </u> | | | | . , | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 170 (73.9) | 161 (65.2) |
| | | | 95% CI ^a | (68.24, 79.59) | (59.24, 71.12) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.13 (1.01, 1.28) | |
| | | | P-value | 0.0387 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 94 (71.2) | 76 (64.4) |
| | | | 95% CI ^a | (63.49, 78.94) | (55.77, 73.05) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.11 (0.93, 1.31) | |
| | | | P-value | 0.2538 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7420 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|----------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | o retuit | O vertain | n (%) | 138 (71.5) | 126 (61.8) |
| | | | 95% CI ^a | (65.13, 77.87) | (55.10, 68.43) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.16 (1.01, 1.33) | |
| | | | P-value | 0.0404 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 126 (74.6) | 111 (68.9) |
| | | | 95% CI ^a | (67.99, 81.12) | (61.80, 76.09) |
| | | | Relative Risk (95% CI) ^a | 1.08 (0.94, 1.24) | |
| | | | P-value | 0.2595 | |
| | | | | | |
| | | | Test for interaction ^b | 0.6783 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 132 (74.6) | 129 (66.2) |
| | | | 95% CI ^a | (68.16, 80.99) | (59.51, 72.80) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.13 (0.99, 1.29) | |
| | | | P-value | 0.0756 | |
| | | | | | |
| Region of enrollment: Europe | Overall | erall Overall | N | 150 | 132 |
| | | | n (%) | 109 (72.7) | 89 (67.4) |
| | | | 95% CI ^a | (65.53, 79.80) | (59.43, 75.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.08 (0.92, 1.26) | |
| | | | P-value | 0.3404 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 11 (64.7) | 10 (52.6) |
| | | | 95% CI ^a | (41.99, 87.42) | (30.18, 75.08) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.23 (0.71, 2.14) | |
| | | | P-value | 0.4637 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 12 (66.7) | 9 (47.4) |
| | | | 95% CI ^a | (44.89, 88.44) | (24.92, 69.82) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.41 (0.79, 2.50) | |
| | | | P-value | 0.2446 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9209 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| Baseline disease severity: Moderate | Overell | Overell | N | 216 | 220 |
| Baseline disease severity. Woderate | Overan | Overall | | | <u> </u> |
| | | | n (%) | 161 (74.5) | 147 (66.8) |
| | | | 95% CI ^a | (68.73, 80.35) | (60.60, 73.04) |
| | | | Relative Risk (95% CI) ^a | 1.12 (0.99, 1.26) | |
| | | | P-value | 0.0777 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 103 (70.5) | 90 (62.1) |
| | | | 95% CI ^a | (63.15, 77.94) | (54.17, 69.97) |
| | | | D 1 (' D' 1 (050/ CT) ³ | | |
| | | | Relative Risk (95% CI) ^a | 1.14 (0.96, 1.34) | |
| | | | P-value | 0.1279 | |
| | | | Test for interaction ^b | 0.0500 | I |
| | | | 1 est for interaction | 0.9582 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 3 (1.3) | 4 (1.6) |
| | | | 95% CI ^a | (0.00, 2.77) | (0.05, 3.19) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.81 (0.18, 3.56) | |
| | | | P-value | 0.7754 | |
| | | | | | |
| Age (years) group: >=40 | Overall Overall | Overall | N | 132 | 118 |
| | | | n (%) | 3 (2.3) | 2 (1.7) |
| | | | 95% CI ^a | (0.00, 4.82) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.34 (0.23, 7.89) | |
| | | | P-value | 0.7456 | |
| | | | | | |
| | | | Test for interaction ^b | 0.6669 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|---------------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 4 (2.1) | 4 (2.0) |
| | | | 95% CI ^a | (0.06, 4.08) | (0.06, 3.86) |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.27, 4.17) | |
| | | | P-value | 0.9369 | |
| | _ | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 2 (1.2) | 2 (1.2) |
| | | | 95% CI ^a | (0.00, 2.81) | (0.00, 2.95) |
| | | | Relative Risk (95% CI) ^a | 0.95 (0.14, 6.68) | I |
| | | | P-value | · · · · · · · · · · · · · · · · · · · | |
| | | | P-value | 0.9611 | |
| | | | Test for interaction ^b | 0.9268 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 2 (1.1) | 3 (1.5) |
| | | | 95% CI ^a | (0.00, 2.69) | (0.00, 3.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.73 (0.12, 4.34) | |
| | | | P-value | 0.7336 | |
| Pasion of angellment Eugene | Oxyone 11 | Overall | N | 150 | 132 |
| Region of enrollment: Europe | Overall | Overall | | | - |
| | | | n (%) | 4 (2.7) | 3 (2.3) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 4.82) |
| | | | Relative Risk (95% CI) ^a | 1.17 (0.27, 5.15) | |
| | | | P-value | 0.8322 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| Region of emonificit. Laun America | Overall | Overall | n (%) | 0 | 0 |
| | | | 11 (70) | U | U |
| | | | | | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| • | | | | | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 5 (2.3) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.30, 4.24) |
| | | | | | - |
| | | | Relative Risk (95% CI) ^a | 1.02 (0.30, 3.47) | |
| | | | P-value | 0.9766 | |
| | | | | | - |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 1 (0.7) | 1 (0.7) |
| | | | 95% CI ^a | (0.00, 2.02) | (0.00, 2.04) |
| | 0 | | -7- | | -0 |
| | | | Relative Risk (95% CI) ^a | 0.99 (0.06, 15.73) | |
| | | | P-value | 0.9961 | |
| | | | | | |
| | | | Test for interaction ^b | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | , , | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 3 (1.3) | 4 (1.6) |
| | | | 95% CI ^a | (0.00, 2.77) | (0.05, 3.19) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.81 (0.18, 3.56) | |
| | | | P-value | 0.7754 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 3 (2.3) | 1 (0.8) |
| | | | 95% CI ^a | (0.00, 4.82) | (0.00, 2.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.68 (0.28, 25.43) | |
| | | | P-value | 0.3901 | |
| | | | | | |
| | | | Test for interaction ^b | 0.3599 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 4 (2.1) | 3 (1.5) |
| | | | 95% CI ^a | (0.06, 4.08) | (0.00, 3.12) |
| | | 1 | Relative Risk (95% CI) ^a | 1 11 (0 22 5 22) | |
| | | | ` ' | 1.41 (0.32, 6.22) | |
| | | | P-value | 0.6504 | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 2 (1.2) | 2 (1.2) |
| | | | 95% CI ^a | (0.00, 2.81) | (0.00, 2.95) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.95 (0.14, 6.68) | |
| | | | P-value | 0.9611 | |
| | | | I | | |
| | | | Test for interaction ^b | 0.7121 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|---------|---------------------|-------------------------------------|---------------------------------|--------------------------------|
| | Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 2 (1.1) | 3 (1.5) |
| | | | 95% CI ^a | (0.00, 2.69) | (0.00, 3.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.73 (0.12, 4.34) | |
| | | | P-value | 0.7336 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 2 (1.5) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 3.60) |
| | | | Relative Risk (95% CI) ^a | 1.76 (0.22, 0.45) | |
| | | | P-value | 1.76 (0.33, 9.45) | |
| | | | P-value | 0.5099 | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | I | | |
| | | | Test for interaction ^b | 0.9031 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| <u>U</u> 1 | | | | . , | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 5 (2.3) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.30, 4.24) |
| | | | Relative Risk (95% CI) ^a | 1.02 (0.20. 2.47) | |
| | | | P-value | 1.02 (0.30, 3.47) 0.9766 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 1 (0.7) | 0 |
| | | | 95% CI ^a | (0.00, 2.02) | (0.00, 2.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.99 (0.07, 58.95) | |
| | | | P-value | 0.6898 | |
| | | | Test for interaction ^b | 0.8593 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 6 (2.6) | 6 (2.4) |
| | | | 95% CI ^a | (0.55, 4.67) | (0.51, 4.35) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.07 (0.35, 3.28) | |
| | | | P-value | 0.9004 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 5 (3.8) | 2 (1.7) |
| | | | 95% CI ^a | (0.53, 7.04) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.23 (0.44, 11.30) | |
| | | | P-value | 0.3309 | |
| | R | | | | |
| | | | Test for interaction ^b | 0.4442 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 6 (3.1) | 2 (1.0) |
| | | | 95% CI ^a | (0.66, 5.56) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.17 (0.65, 15.52) | |
| | | | P-value | 0.1544 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 5 (3.0) | 6 (3.7) |
| | | | 95% CI ^a | (0.40, 5.51) | (0.80, 6.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.79 (0.25, 2.55) | |
| | | | P-value | 0.6983 | |
| | | ı | m . c . i . b | | 1 |
| | | | Test for interaction ^b | 0.2392 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) | |
|---|-----------------|---------------------|-------------------------------------|---------------------------------|-----------------------------|----------|
| | System Organ | MedDRA Preferred | | | | |
| Subgroup | Class | Term | | n (%) | n (%) | |
| | | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 | |
| | | | n (%) | 5 (2.8) | 6 (3.1) | |
| | | | 95% CI ^a | (0.38, 5.27) | (0.65, 5.50) | |
| | | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.92 (0.29, 2.96) | | |
| | | | P-value | 0.8861 | | |
| | | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 | |
| | | | | n (%) | 5 (3.3) | 2 (1.5) |
| | | | 95% CI ^a | (0.46, 6.21) | (0.00, 3.60) | |
| | | | D-1-4: D:-1- (050/ CD) | | | |
| | | | Relative Risk (95% CI) ^a | 2.20 (0.43, 11.15) | | |
| | | | P-value | 0.3410 | | |
| 5 · · · · · · · · · · · · · · · · · · · | 0 11 | 0 " | | | 10 | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 | |
| | | | n (%) | 1 (5.9) | 0 | |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 17.65) | |
| | | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.29 (0.08, 64.21) | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-----------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.8432 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Subgroup | Class | Term | | 11 (70) | 11 (70) |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 5 (2.3) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.30, 4.24) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.02 (0.30, 3.47) | |
| | | | P-value | 0.9766 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 6 (4.1) | 3 (2.1) |
| | | | 95% CI ^a | (0.89, 7.33) | (0.00, 4.39) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.99 (0.51, 7.79) | |
| | | | P-value | 0.3250 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4201 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 5 (2.2) | 6 (2.4) |
| | | | 95% CI ^a | (0.29, 4.06) | (0.51, 4.35) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.89 (0.28, 2.89) | |
| | | | P-value | 0.8529 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 5 (3.8) | 2 (1.7) |
| | | | 95% CI ^a | (0.53, 7.04) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.23 (0.44, 11.30) | |
| | | | P-value | 0.3309 | |
| | | | | | |
| | | | Test for interaction ^b | 0.3408 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 6 (3.1) | 2 (1.0) |
| | | | 95% CI ^a | (0.66, 5.56) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.17 (0.65, 15.52) | |
| | | | P-value | 0.1544 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 4 (2.4) | 6 (3.7) |
| | | | 95% CI ^a | (0.07, 4.66) | (0.80, 6.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.64 (0.18, 2.21) | |
| | | | P-value | 0.4754 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1439 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|---------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| 0.1 | Organ | MedDRA Preferred | | (01) | (01) |
| Subgroup | Class | Term | | n (%) | n (%) |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 5 (2.8) | 6 (3.1) |
| | | | 95% CI ^a | (0.38, 5.27) | (0.65, 5.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.92 (0.29, 2.96) | |
| | | | P-value | 0.8861 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 5 (3.3) | 2 (1.5) |
| | | | 95% CI ^a | (0.46, 6.21) | (0.00, 3.60) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.20 (0.43, 11.15) | |
| | | | P-value | 0.3410 | |
| Desire of succession Asia | O11 | Overall | N | 17 | 19 |
| Region of enrollment: Asia | Overan | Overali | | | |
| | | | n (%) | 0 | 0 |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.8759 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | 1 | | (,,,) | (/+/ |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 5 (2.3) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.30, 4.24) |
| | | | Relative Risk (95% CI) ^a | 1.02 (0.30, 3.47) | |
| | | | P-value | 0.9766 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 3 (2.1) |
| | | | 95% CI ^a | (0.47, 6.37) | (0.00, 4.39) |
| | | | Polotive Piels (050/ CD4 | 1.55 (0.10.500) | |
| | | | Relative Risk (95% CI) ^a | 1.66 (0.40, 6.80) | |
| | | | P-value | 0.4845 | |
| | | | Test for interaction ^b | 0.5825 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|----------------------------|-----------------------|-------------------------|---------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | | | |
| Overall | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 129 (56.1) | 87 (35.2) |
| | | | 95% CI ^a | (49.67, 62.50) | (29.27, 41.18) |
| | | | Relative Risk (95% CI) ^a | 1.50 (1.00, 1.05) | |
| | | | | 1.59 (1.30, 1.95) | |
| | | | P-value | <.0001 | |
| | | | \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ | 122 | 110 |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 68 (51.5) | 42 (35.6) |
| | | | 95% CI ^a | (42.99, 60.04) | (26.95, 44.23) |
| | | | Polotive Diele (050/ CD8 | 1.17 (1.00.1.01) | |
| | | | Relative Risk (95% CI) ^a | 1.45 (1.08, 1.94) | |
| | | | P-value | 0.0136 | |
| | | | Test for interaction ^b | 0.4642 | |
| | | | Test for interaction | 0.4042 | |
| Gastrointestinal disorders | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 53 (23.0) | 7 (2.8) |
| | | | 95% CI ^a | (17.60, 28.49) | (0.76, 4.90) |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 8.13 (3.77, 17.52) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 17 (12.9) | 4 (3.4) |
| | | | 95% CI ^a | (7.16, 18.59) | (0.12, 6.66) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.80 (1.32, 10.97) | |
| | | | P-value | 0.0136 | |
| | <u>"</u> | | | | |
| | | | Test for interaction ^b | 0.0148 | |
| | | | | | |
| | Nausea | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 53 (23.0) | 5 (2.0) |
| | | | 95% CI ^a | (17.60, 28.49) | (0.27, 3.78) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 11.38 (4.63, 27.98) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 17 (12.9) | 3 (2.5) |
| | | 95% CI ^a | (7.16, 18.59) | (0.00, 5.38) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|-------------------------|---|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 5.07 (1.52, 16.85) | |
| | | | P-value | 0.0082 | |
| | | | Test for interaction ^b | 0.0132 | |
| | | | rest for interaction | 0.0132 | |
| | Vomiting | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 9 (3.9) | 4 (1.6) |
| | | | 95% CI ^a | (1.41, 6.42) | (0.05, 3.19) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.42 (0.75, 7.74) | |
| | | | P-value | 0.1374 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 2 (1.5) | 2 (1.7) |
| | | | 95% CI ^a | (0.00, 3.60) | (0.00, 4.02) |
| | | | | | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.89 (0.13, 6.25) | |
| | | | Relative Risk (95% CI) ^a P-value | 0.89 (0.13, 6.25) 0.9100 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--|-----------------------|-------------------------|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| General disorders and administration site conditions | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 3 (1.3) | 3 (1.2) |
| | | | 95% CI ^a | (0.00, 2.77) | (0.00, 2.58) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.07 (0.22, 5.27) | |
| | | | P-value | 0.9300 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 7 (5.3) | 2 (1.7) |
| | | | 95% CI ^a | (1.48, 9.13) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.13 (0.66, 14.77) | |
| | | | P-value | 0.1497 | |
| | | | m of the h | | |
| | | | Test for interaction ^b | 0.1610 | |
| | Estima | A 22 (12222) 2021 | N | 220 | 247 |
| | Fatigue | Age (years) group: <40 | N | 230 | 241 |
| | | | n (%) | 3 (1.3) | 3 (1.2) |
| | | | 95% CI ^a | (0.00, 2.77) | (0.00, 2.58) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|----------------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.07 (0.22, 5.27) | |
| | | | P-value | 0.9300 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 7 (5.3) | 2 (1.7) |
| | | | 95% CI ^a | (1.48, 9.13) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.13 (0.66, 14.77) | |
| | | | P-value | 0.1497 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1610 | |
| | | | | | |
| nfections and infestations | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 44 (19.1) | 48 (19.4) |
| | | | 95% CI ^a | (14.05, 24.21) | (14.50, 24.37) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.98 (0.68, 1.42) | |
| | | | P-value | 0.9333 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 24 (18.2) | 28 (23.7) |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|-------------------|-----------------------|----------------------------|---|---|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 0.77 (0.47, 1.24) | |
| | | | P-value | 0.2823 | |
| | | | Test for interaction ^b | 0.4024 | 1 |
| | | | Test for interaction | 0.4034 | |
| | COVID-19 | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 9 (3.9) | 9 (3.6) |
| | | 95% CI ^a | (1.41, 6.42) | (1.31, 5.98) | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.07 (0.43, 2.66) | |
| | | | P-value | 0.8774 | |
| | | | | | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | Age (years) group: >=40 | N n (%) | 6 (4.5) | 3 (2.5) |
| | | Age (years) group: >=40 | | | |
| | | Age (years) group: >=40 | n (%) | 6 (4.5) | 3 (2.5) |
| | | Age (years) group: >=40 | n (%) | 6 (4.5) | 3 (2.5) |
| | | Age (years) group: >=40 | n (%) 95% CI ^a | 6 (4.5) (0.99, 8.10) | 3 (2.5) |
| | | Age (years) group: >=40 | n (%) 95% CI ^a Relative Risk (95% CI) ^a | 6 (4.5) (0.99, 8.10) 1.79 (0.46, 6.99) | 3 (2.5) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Conjunctivitis | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 4 (1.7) | 20 (8.1) |
| | | | 95% CI ^a | (0.05, 3.43) | (4.70, 11.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.21 (0.07, 0.62) | |
| | | | P-value | 0.0044 | |
| | | 1 | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 4 (3.0) | 15 (12.7) |
| | | | 95% CI ^a | (0.11, 5.95) | (6.70, 18.72) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.24 (0.08, 0.70) | |
| | | | P-value | 0.0089 | |
| | | | | | |
| | | | Test for interaction ^b | 0.3803 | |
| | F. 11: 12: | A () | NY . | 220 | 247 |
| | Folliculitis | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 7 (3.0) | 2 (0.8) |
| | | | 95% CI ^a | (0.82, 5.26) | (0.00, 1.93) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 3.76 (0.79, 17.91) | |
| | | | P-value | 0.0965 | |
| | | | | | |
| | | >=40 n | N | 132 | 118 |
| | | | n (%) | 5 (3.8) | 1 (0.8) |
| | | | 95% CI ^a | (0.53, 7.04) | (0.00, 2.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.47 (0.53, 37.71) | |
| | | | P-value | 0.1688 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7526 | |
| | | | | | |
| | Herpes simplex | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 9 (3.9) | 3 (1.2) |
| | | | 95% CI ^a | (1.41, 6.42) | (0.00, 2.58) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.22 (0.88, 11.75) | |
| | | | P-value | 0.0764 | |
| | | * | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 3 (2.3) | 2 (1.7) |
| | | | 95% CI ^a | (0.00, 4.82) | (0.00, 4.02) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|-------------------------|---|-----------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.34 (0.23, 7.89) | |
| | | | P-value | 0.7456 | |
| | | | Test for interaction ^b | 0.3522 | |
| | | | rest for interaction | 0.3322 | |
| | Nasopharyngitis | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 11 (4.8) | 10 (4.0) |
| | | 95% CI ^a | (2.02, 7.54) | (1.59, 6.51) | |
| | · | | | | |
| | | | Relative Risk (95% CI) ^a | 1.18 (0.51, 2.73) | |
| | | | P-value | 0.6965 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | n (| n (%) | 3 (2.3) | 2 (1.7) |
| | | | 11 (/0) | 3 (2.3) | () |
| | | | 95% CI ^a | (0.00, 4.82) | (0.00, 4.02) |
| | | | | | |
| | | | | | |
| | | | 95% CI ^a | (0.00, 4.82) | |
| | | | 95% CI ^a Relative Risk (95% CI) ^a | (0.00, 4.82) 1.34 (0.23, 7.89) | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Oral herpes | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 7 (3.0) | 7 (2.8) |
| | | | 95% CI ^a | (0.82, 5.26) | (0.76, 4.90) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.07 (0.38, 3.01) | |
| | | | P-value | 0.8923 | |
| | | - | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 2 (1.5) | 8 (6.8) |
| | | | 95% CI ^a | (0.00, 3.60) | (2.24, 11.32) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.22 (0.05, 1.03) | |
| | | | P-value | 0.0548 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0692 | |
| | | | | | |
| | Upper respiratory tract infection | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 6 (2.6) | 7 (2.8) |
| | | | 95% CI ^a | (0.55, 4.67) | (0.76, 4.90) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 0.92 (0.31, 2.70) | |
| | | | P-value | 0.8800 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 4 (3.0) | 2 (1.7) |
| | | | 95% CI ^a | (0.11, 5.95) | (0.00, 4.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.79 (0.33, 9.58) | |
| | | | P-value | 0.4976 | |
| | | | | | |
| | | | Test for interaction ^b | 0.5198 | |
| | | | | | ' |
| Investigations | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 22 (9.6) | 19 (7.7) |
| | | | 95% CI ^a | (5.76, 13.37) | (4.37, 11.02) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.24 (0.69, 2.24) | |
| | | | P-value | 0.4668 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 16 (12.1) | 7 (5.9) |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2' (N=365) |
|--------------------|--|-------------------------|---|------------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.04 (0.87, 4.79) | |
| | | | P-value | 0.1005 | |
| | | | Test for interaction ^b | 0.3293 | 1 |
| | | | rest for interaction | 0.3293 | |
| | Blood creatine phosphokinase increased | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 5 (2.2) | 10 (4.0) |
| | | | 95% CI ^a | (0.29, 4.06) | (1.59, 6.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.54 (0.19, 1.55) | |
| | | | P-value | 0.2495 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | >=40 | | | |
| | | >=40 | n (%) | 9 (6.8) | 3 (2.5) |
| | | >=40 | n (%) 95% CI ^a | 9 (6.8) (2.52, 11.12) | 3 (2.5) (0.00, 5.38) |
| | | >=40 | | | |
| | | >=40 | | | |
| | | >=40 | 95% CI ^a | (2.52, 11.12) | |
| | | >=40 | 95% CI ^a Relative Risk (95% CI) ^a | (2.52, 11.12) 2.68 (0.74, 9.67) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-------------------------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Natural killer cell count decreased | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 7 (3.0) | 0 |
| | | | 95% CI ^a | (0.82, 5.26) | (0.00, 1.48) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 15.07 (0.86, 263.98) | |
| | | | P-value | 0.0634 | |
| | · | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 3 (2.3) | 0 |
| | | | 95% CI ^a | (0.00, 4.82) | (0.00, 3.08) |
| | | | | | 200 |
| | | | Relative Risk (95% CI) ^a | 5.39 (0.27, 106.43) | |
| | | | P-value | 0.2687 | |
| | | | | | |
| | | | Test for interaction ^b | 0.5919 | |
| | | | | | |
| | SARS-CoV-2 test positive | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 10 (4.3) | 9 (3.6) |
| | | | 95% CI ^a | (1.71, 6.98) | (1.31, 5.98) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.19 (0.49, 2.88) | |
| | | | P-value | 0.6948 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 5 (3.8) | 4 (3.4) |
| | | | 95% CI ^a | (0.53, 7.04) | (0.12, 6.66) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.12 (0.31, 4.06) | |
| | | | P-value | 0.8662 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9170 | |
| | | | | | |
| Nervous system disorders | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 33 (14.3) | 18 (7.3) |
| | | | 95% CI ^a | (9.82, 18.88) | (4.05, 10.53) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.97 (1.14, 3.40) | |
| | | | P-value | 0.0149 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 22 (16.7) | 9 (7.6) |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.19 (1.05, 4.56) | |
| | | | P-value | 0.0370 | |
| | | | Test for interaction ^b | 0.6772 | |
| | | | rest for interaction | 0.0772 | |
| | Dizziness | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 6 (2.6) | 1 (0.4) |
| | | 95% CI ^a | (0.55, 4.67) | (0.00, 1.20) | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 6.44 (0.78, 53.11) | |
| | | | P-value | 0.0834 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 4 (3.0) | 3 (2.5) |
| | | | 95% CI ^a | (0.11, 5.95) | (0.00, 5.38) |
| | · | | | | |
| | | | Relative Risk (95% CI) ^a | 1.19 (0.27, 5.22) | |
| | | | P-value | 0.8157 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4751 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Headache | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 28 (12.2) | 17 (6.9) |
| | | | 95% CI ^a | (7.95, 16.40) | (3.73, 10.04) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.77 (1.00, 3.14) | |
| | | | P-value | 0.0520 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 19 (14.4) | 7 (5.9) |
| | | | 95% CI ^a | (8.41, 20.38) | (1.67, 10.19) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.43 (1.06, 5.57) | |
| | | | P-value | 0.0364 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4922 | |
| | | | | | |
| Skin and subcutaneous tissue disorders | Overall | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 47 (20.4) | 19 (7.7) |
| | | | 95% CI ^a | (15.22, 25.65) | (4.37, 11.02) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|-------------------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.66 (1.61, 4.39) | |
| | | | P-value | 0.0001 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 14 (10.6) | 4 (3.4) |
| | | 95% CI ^a | (5.35, 15.86) | (0.12, 6.66) | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.13 (1.06, 9.24) | |
| | | | P-value | 0.0390 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1683 | |
| | | | | | |
| | Acne | Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 33 (14.3) | 10 (4.0) |
| | | | 95% CI ^a | (9.82, 18.88) | (1.59, 6.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.54 (1.79, 7.03) | |
| | | | P-value | 0.0003 | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | | n (%) | 13 (9.8) | 0 |
| | | 95% CI ^a | (4.77, 14.93) | (0.00, 3.08) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|----------------------------|---|---|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 23.34 (1.40, 389.13) | |
| | | | P-value | 0.0282 | |
| | | | Test for interaction ^b | 0.7370 | 1 |
| | | | rest for interaction | 0.7370 | |
| | Dermatitis atopic | c Age (years) group: <40 | N | 230 | 247 |
| | | | n (%) | 16 (7.0) | 10 (4.0) |
| | | | 95% CI ^a | (3.67, 10.24) | (1.59, 6.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.72 (0.80, 3.71) | I |
| | | | P-value | 0.1679 | |
| | | | | | |
| | | | | | |
| | | Age (years) group: >=40 | N | 132 | 118 |
| | | Age (years) group: >=40 | N n (%) | 132 | 118 |
| | | Age (years) group: >=40 | | | |
| | | Age (years) group: >=40 | n (%) 95% CI ^a | 1 (0.8) | 4 (3.4) |
| | | Age (years) group: >=40 | n (%) | 1 (0.8) | 4 (3.4) |
| | | Age (years) group: >=40 | n (%) 95% CI ^a | 1 (0.8) (0.00, 2.24) | 4 (3.4) |
| | | Age (years) group: >=40 | n (%) 95% CI ^a Relative Risk (95% CI) ^a | 1 (0.8) (0.00, 2.24) 0.22 (0.03, 1.97) | 4 (3.4) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|----------------------------|-----------------------|----------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| Overall | Overall | Sex: Male | N | 193 | 204 |
| Overall | Overall | Sex. Male | n (%) | 100 (51.8) | 68 (33.3) |
| | | | 95% CI ^a | (44.76, 58.86) | (26.86, 39.80) |
| | | | , | (11.70, 30.00) | (20.00, 37.00) |
| | | T | Relative Risk (95% CI) ^a | 1.55 (1.23, 1.97) | |
| | | | P-value | 0.0003 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 97 (57.4) | 61 (37.9) |
| | | | 95% CI ^a | (49.94, 64.85) | (30.39, 45.38) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.51 (1.20, 1.92) | |
| | | | P-value | 0.0006 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7096 | |
| | | | | | |
| Gastrointestinal disorders | Overall | Sex: Male | N | 193 | 204 |
| | | | n (%) | 21 (10.9) | 4 (2.0) |
| | | | 95% CI ^a | (6.49, 15.27) | (0.06, 3.86) |
| | | | Relative Risk (95% CI) ^a | 5.55 (1.94, 15.87) | |
| | | | P-value | 0.0014 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 49 (29.0) | 7 (4.3) |
| | | | 95% CI ^a | (22.15, 35.83) | (1.20, 7.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 6.67 (3.11, 14.29) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0005 | |
| | | To contract | To a second | | |
| | Nausea | Sex: Male | N | 193 | 204 |
| | | | n (%) | 21 (10.9) | 2 (1.0) |
| | | | 95% CI ^a | (6.49, 15.27) | (0.00, 2.33) |
| | | | Relative Risk (95% CI) ^a | 11 10 (2 (4 46 70) | |
| | | | P-value | 11.10 (2.64, 46.70) | |
| | | | 1 -value | 0.0010 | |
| | | Sex: | N | 169 | 161 |
| | | Female | | | |
| | | | n (%) | 49 (29.0) | 6 (3.7) |
| | | | 95% CI ^a | (22.15, 35.83) | (0.80, 6.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 7.78 (3.43, 17.66) | |
| | | | P-value | <.0001 | |
| | | | | | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--|-----------------------|----------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Vomiting | Sex: Male | N | 193 | 204 |
| | | | n (%) | 2 (1.0) | 3 (1.5) |
| | | | 95% CI ^a | (0.00, 2.46) | (0.00, 3.12) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.70 (0.12, 4.17) | |
| | | | P-value | 0.6997 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 9 (5.3) | 3 (1.9) |
| | | | 95% CI ^a | (1.94, 8.71) | (0.00, 3.95) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.86 (0.79, 10.37) | |
| | | | P-value | 0.1102 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0936 | |
| | | | | | |
| General disorders and administration site conditions | Overall | Sex: Male | N | 193 | 204 |
| | | | n (%) | 3 (1.6) | 3 (1.5) |
| | | | 95% CI ^a | (0.00, 3.30) | (0.00, 3.12) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.22, 5.17) | |
| | | | P-value | 0.9455 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|---------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 7 (4.1) | 2 (1.2) |
| | | | 95% CI ^a | (1.14, 7.15) | (0.00, 2.95) |
| | | | Relative Risk (95% CI) ^a | 2.22 (0.70, 15.01) | |
| | | | | 3.33 (0.70, 15.81) | |
| | | | P-value | 0.1294 | |
| | | | Test for interaction ^b | 0.1904 | |
| | | | I | | |
| | Fatigue | Sex: Male | N | 193 | 204 |
| | | | n (%) | 3 (1.6) | 3 (1.5) |
| | | | 95% CI ^a | (0.00, 3.30) | (0.00, 3.12) |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.22, 5.17) | |
| | | | P-value | 0.9455 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 7 (4.1) | 2 (1.2) |
| | | | 95% CI ^a | (1.14, 7.15) | (0.00, 2.95) |
| | | | Relative Risk (95% CI) ^a | 3.33 (0.70, 15.81) | |
| | | | P-value | 0.1294 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1904 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------------|-----------------------|-----------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| Infections and infestations | Overall | ıll Sex: Male N | N | 193 | 204 |
| | | | n (%) | 40 (20.7) | 41 (20.1) |
| | | | 95% CI ^a | (15.01, 26.44) | (14.60, 25.60) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.03 (0.70, 1.52) | |
| | | | P-value | 0.8768 | |
| | | | | | |
| | | | N | 169 | 161 |
| | | | n (%) | 28 (16.6) | 35 (21.7) |
| | | | 95% CI ^a | (10.96, 22.17) | (15.37, 28.11) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.76 (0.49, 1.19) | |
| | | | P-value | 0.2343 | |
| | | | | | |
| | | | Test for interaction ^b | 0.3320 | |
| | | | | | |
| | COVID-19 | Sex: Male | N | 193 | 204 |
| | | | n (%) | 7 (3.6) | 5 (2.5) |
| | | | 95% CI ^a | (0.99, 6.26) | (0.33, 4.57) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.48 (0.48, 4.58) | |
| | | | P-value | 0.4969 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Sex: Female | N | 169 | 161 | |
| | | | n (%) | 8 (4.7) | 7 (4.3) |
| | | 95 | 95% CI ^a | (1.53, 7.94) | (1.20, 7.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.09 (0.40, 2.93) | |
| | | | P-value | 0.8665 | |
| | | | Test for interaction ^b | 0.7869 | |
| | | | | | |
| | Conjunctivitis | Sex: Male | N | 193 | 204 |
| | | | n (%) | 7 (3.6) | 20 (9.8) |
| | | | 95% CI ^a | (0.99, 6.26) | (5.72, 13.88) |
| | | | Relative Risk (95% CI) ^a | 0.37 (0.16, 0.86) | |
| | | | P-value | 0.0200 | |
| | | la. | l., | 1.50 | 1.51 |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 1 (0.6) | 15 (9.3) |
| | | | 95% CI ^a | (0.00, 1.75) | (4.83, 13.81) |
| | | | Relative Risk (95% CI) ^a | 0.06 (0.01, 0.48) | I |
| | | | P-value | 0.00 (0.01, 0.48) | |
| | | | 1 | 0.0073 | |
| | | | Test for interaction ^b | 0.4937 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--------------------|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Folliculitis | Sex: Male | N | 193 | 204 |
| | | | n (%) | 10 (5.2) | 2 (1.0) |
| | | | 95% CI ^a | (2.05, 8.31) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 5.28 (1.17, 23.81) | |
| | | | P-value | 0.0302 | |
| | - | - | | | |
| | | Sex: N Female n (% | N | 169 | 161 |
| | | | n (%) | 2 (1.2) | 1 (0.6) |
| | | | 95% CI ^a | (0.00, 2.81) | (0.00, 1.83) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.91 (0.17, 20.81) | |
| | | | P-value | 0.5972 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0729 | |
| | <u> </u> | | | | |
| | Herpes simplex | Sex: Male | N | 193 | 204 |
| | | | n (%) | 8 (4.1) | 2 (1.0) |
| | | | 95% CI ^a | (1.33, 6.96) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.23 (0.91, 19.66) | |
| | | | P-value | 0.0660 | Ť |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|---------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 4 (2.4) | 3 (1.9) |
| | | | 95% CI ^a | (0.07, 4.66) | (0.00, 3.95) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.27 (0.29, 5.59) | |
| | | | P-value | 0.7516 | |
| | | | I= | | |
| | | | Test for interaction ^b | 0.2368 | |
| | Nasopharyngitis | Sex: Male | N | 193 | 204 |
| | Nasopharyngius | Sex. Wate | n (%) | 7 (3.6) | 7 (3.4) |
| | | | 95% CI ^a | (0.99, 6.26) | (0.93, 5.93) |
| | | - | | (****, **=*) | (0,,0,0,0) |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.38, 2.96) | |
| | | | P-value | 0.9159 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 7 (4.1) | 5 (3.1) |
| | | | 95% CI ^a | (1.14, 7.15) | (0.43, 5.79) |
| | | | D 1 .: D: 1 (050) GD3 | | |
| | | | Relative Risk (95% CI) ^a | 1.33 (0.43, 4.12) | |
| | | | P-value | 0.6165 | |
| | | | Test for interaction ^b | 0.7611 | T |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------------------|-----------|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Oral herpes | Sex: Male | N | 193 | 204 |
| | | | n (%) | 3 (1.6) | 7 (3.4) |
| | | | 95% CI ^a | (0.00, 3.30) | (0.93, 5.93) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.45 (0.12, 1.73) | |
| | | | P-value | 0.2461 | |
| | - | | | | |
| | | Female n | N | 169 | 161 |
| | | | n (%) | 6 (3.6) | 8 (5.0) |
| | | | 95% CI ^a | (0.76, 6.34) | (1.61, 8.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.71 (0.25, 2.01) | |
| | | | P-value | 0.5249 | |
| | | | | | |
| | | | Test for interaction ^b | 0.8754 | |
| | | | | | |
| | Upper respiratory tract infection | Sex: Male | N | 193 | 204 |
| | | | n (%) | 6 (3.1) | 3 (1.5) |
| | | | 95% CI ^a | (0.66, 5.56) | (0.00, 3.12) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.11 (0.54, 8.33) | |
| | | | P-value | 0.2849 | 1 |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|----------------|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 4 (2.4) | 6 (3.7) |
| | | | 95% CI ^a | (0.07, 4.66) | (0.80, 6.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.64 (0.18, 2.21) | |
| | | | P-value | 0.4754 | |
| | | | | | _ |
| | | | Test for interaction ^b | 0.2174 | |
| | | To contract | 1 | | |
| Investigations | Overall | Sex: Male | N | 193 | 204 |
| | | | n (%) | 21 (10.9) | 12 (5.9) |
| | | | 95% CI ^a | (6.49, 15.27) | (2.65, 9.11) |
| | | | Relative Risk (95% CI) ^a | 1.85 (0.94, 3.66) | |
| | | | P-value | 0.0769 | |
| | | | | 010707 | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 17 (10.1) | 14 (8.7) |
| | | | 95% CI ^a | (5.52, 14.59) | (4.34, 13.05) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.16 (0.59, 2.27) | |
| | | | P-value | 0.6717 | |
| | | | | | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|--|----------------|-------------------------------------|------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Blood creatine phosphokinase increased | Sex: Male | N | 193 | 204 |
| | | | n (%) | 9 (4.7) | 7 (3.4) |
| | | | 95% CI ^a | (1.69, 7.64) | (0.93, 5.93) |
| | | | Relative Risk (95% CI) ^a | 1.36 (0.52, 3.58) | T |
| | | | P-value | 0.5345 | |
| | <u> </u> | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 5 (3.0) | 6 (3.7) |
| | | | 95% CI ^a | (0.40, 5.51) | (0.80, 6.65) |
| | | | Relative Risk (95% CI) ^a | 0.79 (0.25, 2.55) | |
| | | | P-value | 0.6983 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4755 | |
| | Natural killer cell count decreased | Sex: Male | N | 193 | 204 |
| | T WILLIAM TELLON GOLD GOULD GEOLOGIC | Ben mare | n (%) | 5 (2.6) | 0 |
| | | | 95% CI ^a | (0.35, 4.83) | (0.00, 1.79) |
| | | | Relative Risk (95% CI) ^a | 10.60 (0.58, 192.65) | 1 |
| | | | P-value | 0.1107 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------------|----------------|-------------------------------------|---------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 5 (3.0) | 0 |
| | | ç | 95% CI ^a | (0.40, 5.51) | (0.00, 2.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 9.56 (0.53, 173.52) | |
| | | | P-value | 0.1270 | |
| | | | I= | | |
| | | | Test for interaction ^b | 0.8670 | |
| | SARS-CoV-2 test positive | Sex: Male | N | 193 | 204 |
| | 5711tb 66 + 2 test positive | Sem mare | n (%) | 7 (3.6) | 5 (2.5) |
| | | | 95% CI ^a | (0.99, 6.26) | (0.33, 4.57) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.48 (0.48, 4.58) | |
| | | | P-value | 0.4969 | |
| | II. | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 8 (4.7) | 8 (5.0) |
| | | | 95% CI ^a | (1.53, 7.94) | (1.61, 8.33) |
| | | | D 1 (' D' 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 0.95 (0.37, 2.48) | |
| | | | P-value | 0.9208 | |
| | | | Test for interaction ^b | 0.6334 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------------|-----------------------|---------------------|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| Nervous system disorders | Overall | Sex: Male | N | 193 | 204 |
| | | | n (%) | 28 (14.5) | 12 (5.9) |
| | | | 95% CI ^a | (9.54, 19.48) | (2.65, 9.11) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.47 (1.29, 4.71) | |
| | | | P-value | 0.0062 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 27 (16.0) | 15 (9.3) |
| | | 95% CI ^a | 95% CI ^a | (10.45, 21.50) | (4.83, 13.81) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.71 (0.95, 3.10) | |
| | | | P-value | 0.0747 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7148 | |
| | | | | | |
| | Dizziness | Sex: Male | N | 193 | 204 |
| | | | n (%) | 4 (2.1) | 2 (1.0) |
| | | | 95% CI ^a | (0.06, 4.08) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.11 (0.39, 11.41) | |
| | | | P-value | 0.3842 | Î |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 6 (3.6) | 2 (1.2) |
| | | | 95% CI ^a | (0.76, 6.34) | (0.00, 2.95) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.86 (0.59, 13.95) | |
| | | | P-value | 0.1943 | |
| | | | I | | |
| | | | Test for interaction ^b | 0.5557 | |
| | Headache | Sex: Male | N | 193 | 204 |
| | Treadache | Sex. Mare | n (%) | 24 (12.4) | 10 (4.9) |
| | | | 95% CI ^a | (7.78, 17.09) | (1.94, 7.86) |
| | | | | , , , | |
| | | | Relative Risk (95% CI) ^a | 2.54 (1.25, 5.16) | |
| | | | P-value | 0.0103 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 23 (13.6) | 14 (8.7) |
| | | | 95% CI ^a | (8.44, 18.78) | (4.34, 13.05) |
| | | | D 1 .: D: 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 1.57 (0.83, 2.93) | |
| | | | P-value | 0.1623 | |
| | | | Test for interaction ^b | 0.5872 | T |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--|-----------------------|----------------|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| Skin and subcutaneous tissue disorders | Overall | Sex: Male | N | 193 | 204 |
| | | | n (%) | 34 (17.6) | 12 (5.9) |
| | | | 95% CI ^a | (12.24, 22.99) | (2.65, 9.11) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.99 (1.60, 5.61) | |
| | | | P-value | 0.0006 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 27 (16.0) | 11 (6.8) |
| | | 9 | 95% CI ^a | (10.45, 21.50) | (2.94, 10.73) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.34 (1.20, 4.56) | |
| | | | P-value | 0.0126 | |
| | | | | | |
| | | | Test for interaction ^b | 0.5853 | |
| | | | | | |
| | Acne | Sex: Male | N | 193 | 204 |
| | | | n (%) | 24 (12.4) | 4 (2.0) |
| | | | 95% CI ^a | (7.78, 17.09) | (0.06, 3.86) |
| | - | - | | | |
| | | | Relative Risk (95% CI) ^a | 6.34 (2.24, 17.94) | |
| | | | P-value | 0.0005 | Î |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|----------------|-------------------------------------|---------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 22 (13.0) | 6 (3.7) |
| | | | 95% CI ^a | (7.94, 18.09) | (0.80, 6.65) |
| | | | D 1 1 D11 (050) GD3 | | |
| | | | Relative Risk (95% CI) ^a | 3.49 (1.45, 8.39) | |
| | | | P-value | 0.0052 | |
| | | 1 | Test for interaction ^b | 0.5050 | T |
| | | | Test for interaction | 0.7950 | |
| | Dermatitis atopic | Sex: Male | N | 193 | 204 |
| | | | n (%) | 11 (5.7) | 8 (3.9) |
| | | | 95% CI ^a | (2.43, 8.97) | (1.26, 6.59) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.45 (0.60, 3.54) | |
| | | | P-value | 0.4099 | |
| | | | | | |
| | | Sex: Female | N | 169 | 161 |
| | | | n (%) | 6 (3.6) | 6 (3.7) |
| | | | 95% CI ^a | (0.76, 6.34) | (0.80, 6.65) |
| | | | D-1-4 Di-1- (050/ CD8 | | |
| | | | Relative Risk (95% CI) ^a | 0.95 (0.31, 2.89) | |
| | | | P-value | 0.9318 | |
| | | | Test for interaction ^b | 0.5105 | 1 |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used. The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

^{.5} was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | | | |
| Overall | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 92 (52.0) | 68 (34.9) |
| | | 95% CI ^a | (44.62, 59.34) | (28.18, 41.56) | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.49 (1.17, 1.89) | |
| | | | P-value | 0.0010 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 90 (60.0) | 55 (41.7) |
| | | | 95% CI ^a | (52.16, 67.84) | (33.26, 50.08) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.44 (1.13, 1.83) | |
| | | | P-value | 0.0030 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 9 (52.9) | 4 (21.1) |
| | | | 95% CI ^a | (29.21, 76.67) | (2.72, 39.38) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.51 (0.94, 6.70) | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|----------------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 6 (33.3) | 2 (10.5) |
| | | | 95% CI ^a | (11.56, 55.11) | (0.00, 24.33) |
| | | 1 | Relative Risk (95% CI) ^a | 3.17 (0.73, 13.70) | T |
| | | P-value | 0.1230 | | |
| | 7 | | | | |
| | | | Test for interaction ^b | 0.8773 | |
| | | | | | |
| Gastrointestinal disorders | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 39 (22.0) | 6 (3.1) |
| | | | 95% CI ^a | (15.93, 28.14) | (0.65, 5.50) |
| | | | Relative Risk (95% CI) ^a | 7.16 (3.11, 16.51) | T |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 30 (20.0) | 5 (3.8) |
| | | 95% CI ^a | (13.60, 26.40) | (0.53, 7.04) | |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 5.28 (2.11, 13.21) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 1 (5.9) | 0 |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.29 (0.08, 64.21) | |
| | | | P-value | 0.6252 | |
| | | | | | |
| | | America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.0039 | |
| | · | | | | |
| | Nausea | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 39 (22.0) | 4 (2.1) |
| | | | 95% CI ^a | (15.93, 28.14) | (0.06, 4.04) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 10.74 (3.92, 29.45) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | m (0/) | 30 (20.0) | 4 (3.0) |
| | | | n (%) | 30 (20.0) | + (5.0) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 6.60 (2.39, 18.24) | |
| | | | P-value | 0.0003 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 1 (5.9) | 0 |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 17.65) |
| | | | | | |
| | T T | | Relative Risk (95% CI) ^a | 2.29 (0.08, 64.21) | |
| | | P-value | 0.6252 | | |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.0019 | |
| | | | | | |
| | Vomiting | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 7 (4.0) | 3 (1.5) |
| | | | 95% CI ^a | (1.08, 6.83) | (0.00, 3.27) |
| | | | | | |
| | | | | | |
| | | I | Relative Risk (95% CI) ^a | 2.57 (0.68, 9.79) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|---|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 3 (2.3) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 4.82) |
| | | | Relative Risk (95% CI) ^a | 1.17 (0.27, 5.15) | T |
| | | | P-value | 0.8322 | |
| | | | | 0.0322 | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.8619 | |
| | | | | | |
| General disorders and administration site conditions | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 6 (3.4) | 2 (1.0) |
| | | | 95% CI ^a | (0.72, 6.06) | (0.00, 2.44) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.31 (0.68, 16.16) | |
| | | | P-value | 0.1399 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 3 (2.3) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 4.82) |
| | | | D 1 .: D: 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 1.17 (0.27, 5.15) | |
| | | | P-value | 0.8322 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | - | | | |
| | | | Test for interaction ^b | 0.8540 | |
| | | | | | |
| | Fatigue | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 6 (3.4) | 2 (1.0) |
| | | | 95% CI ^a | (0.72, 6.06) | (0.00, 2.44) |
| | - | | | | |
| | | | Relative Risk (95% CI) ^a | 3.31 (0.68, 16.16) | |
| | | | P-value | 0.1399 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------------|-----------------------|--|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 3 (2.3) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 4.82) |
| | | | D-1-4: Di-1- (050/ CD) | | |
| | | | Relative Risk (95% CI) ^a | 1.17 (0.27, 5.15) | |
| | | | P-value | 0.8322 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.8540 | |
| | | | | | |
| Infections and infestations | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 24 (13.6) | 38 (19.5) |
| | | | 95% CI ^a | (8.52, 18.60) | (13.93, 25.05) |
| | - | | | | |
| | | | Relative Risk (95% CI) ^a | 0.70 (0.44, 1.11) | |
| | | | P-value | 0.1294 | T . |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2' (N=365) |
|-------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 41 (27.3) | 36 (27.3) |
| | | | 95% CI ^a | (20.20, 34.47) | (19.68, 34.87) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.00 (0.68, 1.47) | |
| | | | P-value | 0.9909 | |
| | | | 1 | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 15.30) |
| | | | Relative Risk (95% CI) ^a | 2.24 (0.22, 22.51) | |
| | | | P-value | 0.4948 | |
| | - | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 15.30) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.07, 15.64) | |
| | | | P-value | 0.9686 | |
| | | | Test for interaction ^b | 0.5521 | I |
| | | | 103t 101 interaction | 0.5531 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | COVID-19 | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 4 (2.3) | 4 (2.1) |
| | | | 95% CI ^a | (0.07, 4.45) | (0.06, 4.04) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.10 (0.28, 4.34) | |
| | | | P-value | 0.8899 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 11 (7.3) | 7 (5.3) |
| | | | 95% CI ^a | (3.16, 11.51) | (1.48, 9.13) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.38 (0.55, 3.46) | |
| | | | P-value | 0.4890 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 18.53) | (0.00, 15.30) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | T T | Relative Risk (95% CI) ^a | 0.51 (0.02, 14.40) | |
| | | | P-value | 0.6952 | |
| | | | Test for interaction ^b | 0.9070 | |
| | | | | | |
| | Conjunctivitis | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 4 (2.3) | 18 (9.2) |
| | | | 95% CI ^a | (0.07, 4.45) | (5.17, 13.29) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.24 (0.08, 0.71) | |
| | | | P-value | 0.0095 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 3 (2.0) | 16 (12.1) |
| | | | 95% CI ^a | (0.00, 4.24) | (6.55, 17.69) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.17 (0.05, 0.55) | |
| | | | P-value | 0.0035 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 1 (5.9) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 15.30) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.12 (0.08, 16.52) | |
| | | | P-value | 0.9355 | |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.2628 | |
| | | | | | |
| | Folliculitis | US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 5 (2.8) | 3 (1.5) |
| | | | 95% CI ^a | (0.38, 5.27) | (0.00, 3.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.84 (0.45, 7.57) | |
| | | | P-value | 0.4006 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 5 (3.3) | 0 |
| | | | 95% CI ^a | (0.46, 6.21) | (0.00, 2.76) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 8.83 (0.49, 160.17) | |
| | | | P-value | 0.1406 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 0 |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.59 (0.22, 94.96) | |
| | | | P-value | 0.3244 | |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.7124 | |
| | ' | | | | |
| | Herpes simplex | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 3 (1.7) | 1 (0.5) |
| | | | 95% CI ^a | (0.00, 3.60) | (0.00, 1.52) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.31 (0.35, 31.48) | |
| | | | P-value | 0.2986 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 9 (6.0) | 4 (3.0) |
| | | | 11 (70) | > (0.0) | . (5.0) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.98 (0.62, 6.28) | |
| | | | P-value | 0.2461 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.9031 | |
| | | | | | |
| | Nasopharyngitis | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 2 (1.1) | 4 (2.1) |
| | | | 95% CI ^a | (0.00, 2.69) | (0.06, 4.04) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.55 (0.10, 2.97) | |
| | | | P-value | 0.4880 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 11 (7.3) | 8 (6.1) |
| | | | 95% CI ^a | (3.16, 11.51) | (1.99, 10.13) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2' (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.21 (0.50, 2.92) | |
| | | | P-value | 0.6712 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 0 |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.17 (0.08, 60.76) | |
| | | | P-value | 0.6494 | |
| | | | | | |
| | | | Test for interaction ^b | 0.8600 | |
| | | | | | |
| | Oral herpes | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 1 (0.6) | 5 (2.6) |
| | | | 95% CI ^a | (0.00, 1.67) | (0.35, 4.78) |
| | | | - | | |
| | | | Relative Risk (95% CI) ^a | 0.22 (0.03, 1.87) | |
| | | | P-value | 0.1654 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 8 (5.3) | 10 (7.6) |
| | | | 95% CI ^a | (1.74, 8.93) | (3.06, 12.09) |
| | | | D 1 .: D: 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 0.70 (0.29, 1.73) | |
| | | | P-value | 0.4446 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.9522 | |
| | | | | | |
| | Upper respiratory tract infection | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 7 (4.0) | 6 (3.1) |
| | | | 95% CI ^a | (1.08, 6.83) | (0.65, 5.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.29 (0.44, 3.75) | |
| | | | P-value | 0.6461 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 3 (2.0) | 3 (2.3) |
| | | 95% CI ^a | (0.00, 4.24) | (0.00, 4.82) | |
| | | | Relative Risk (95% CI) ^a | 0.00 (0.10, 1.20) | 1 |
| | | | | 0.88 (0.18, 4.29) | |
| | | | P-value | 0.8742 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.9776 | |
| | | | | | |
| Investigations | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 12 (6.8) | 13 (6.7) |
| | | | 95% CI ^a | (3.08, 10.48) | (3.17, 10.17) |
| | - | | | · | |
| | | | Relative Risk (95% CI) ^a | 1.02 (0.48, 2.17) | |
| | | | P-value | 0.9653 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|--|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 23 (15.3) | 11 (8.3) |
| | | | 95% CI ^a | (9.57, 21.10) | (3.62, 13.05) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.84 (0.93, 3.63) | |
| | | | P-value | 0.0785 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 3 (16.7) | 2 (10.5) |
| | | | 95% CI ^a | (0.00, 33.88) | (0.00, 24.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.58 (0.30, 8.40) | |
| | | | P-value | 0.5894 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4656 | |
| | | | | | |
| | Blood creatine phosphokinase increased | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 8 (4.5) | 8 (4.1) |
| | | | 95% CI ^a | (1.46, 7.58) | (1.32, 6.89) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|---------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Î | Relative Risk (95% CI) ^a | 1.10 (0.42, 2.87) | |
| | | | P-value | 0.8430 | |
| | | Di | N | 150 | 132 |
| | | Region of enrollment: Europe | | 150 | |
| | | | n (%) | 4 (2.7) | 4 (3.0) |
| | | 9 | 95% CI ^a | (0.09, 5.24) | (0.11, 5.95) |
| | | | Relative Risk (95% CI) ^a | 0.88 (0.22, 3.45) | |
| | | | P-value | 0.8545 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | 1 |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 2 (11.1) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 25.63) | (0.00, 15.30) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.11 (0.21, 21.32) | |
| | | | P-value | 0.5265 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9243 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-------------------------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Natural killer cell count decreased | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 10 (6.7) | 0 |
| | | | 95% CI ^a | (2.67, 10.66) | (0.00, 2.76) |
| | | | Relative Risk (95% CI) ^a | 17.67 (1.04, 299.56) | |
| | | | P-value | 0.0468 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | T | 1 | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | NE | |
| | CARC CoV 2 test mositive | Danian of annullments | N | 177 | 195 |
| | SARS-CoV-2 test positive | Region of enrollment: US/Canada/Australia | IN . | 1// | 193 |
| | | | n (%) | 4 (2.3) | 5 (2.6) |
| | | | 95% CI ^a | (0.07, 4.45) | (0.35, 4.78) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 0.88 (0.24, 3.23) | |
| | | | P-value | 0.8489 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 10 (6.7) | 7 (5.3) |
| | | | 95% CI ^a | (2.67, 10.66) | (1.48, 9.13) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.26 (0.49, 3.21) | |
| | | | P-value | 0.6322 | |
| | · | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 15.30) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.06 (0.07, 15.64) | |
| | | | P-value | 0.9686 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9659 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| Nervous system disorders | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 24 (13.6) | 13 (6.7) |
| | | | 95% CI ^a | (8.52, 18.60) | (3.17, 10.17) |
| | | | Relative Risk (95% CI) ^a | 2.02 (1.07. 2.97) | 1 |
| | | | P-value | 2.03 (1.07, 3.87) | |
| | | | r-value | 0.0306 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 27 (18.0) | 12 (9.1) |
| | | | 95% CI ^a | (11.85, 24.15) | (4.19, 14.00) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.98 (1.05, 3.75) | |
| | | | P-value | 0.0360 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 2 (10.5) |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 24.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.12 (0.18, 7.09) | |
| | | | P-value | 0.9061 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 2 (11.1) | 0 |
| | | | 95% CI ^a | (0.00, 25.63) | (0.00, 17.65) |
| | | 1 | Relative Risk (95% CI) ^a | 4 22 (0 21 80 87) | 1 |
| | | | P-value | 4.33 (0.21, 89.87) | + |
| | | | r-value | 0.3432 | |
| | | | Test for interaction ^b | 0.9048 | |
| | | | | | |
| | Dizziness | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 4 (2.3) | 1 (0.5) |
| | | | 95% CI ^a | (0.07, 4.45) | (0.00, 1.52) |
| | | T | Relative Risk (95% CI) ^a | 4.41 (0.50, 39.06) | T |
| | | | P-value | 0.1828 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 6 (4.0) | 3 (2.3) |
| | | | 95% CI ^a | (0.86, 7.14) | (0.00, 4.82) |
| | | | Relative Risk (95% CI) ^a | 1.76 (0.45, 6.90) | |
| | | | P-value | 0.4173 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.9848 | |
| | | | | | |
| | Headache | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 21 (11.9) | 12 (6.2) |
| | | | 95% CI ^a | (7.10, 16.63) | (2.78, 9.53) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.93 (0.98, 3.80) | |
| | | | P-value | 0.0583 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 22 (14.7) | 10 (7.6) |
| | | | 95% CI ^a | (9.01, 20.33) | (3.06, 12.09) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.94 (0.95, 3.94) | |
| | | | P-value | 0.0682 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 2 (10.5) |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 24.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.12 (0.18, 7.09) | |
| | | | P-value | 0.9061 | |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 2 (11.1) | 0 |
| | | | 95% CI ^a | (0.00, 25.63) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.33 (0.21, 89.87) | |
| | | | P-value | 0.3432 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9443 | |
| | | | | | |
| Skin and subcutaneous tissue disorders | Overall | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 34 (19.2) | 14 (7.2) |
| | | | 95% CI ^a | (13.41, 25.01) | (3.56, 10.80) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.68 (1.49, 4.82) | |
| | | | P-value | 0.0010 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|-------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 21 (14.0) | 7 (5.3) |
| | | | 95% CI ^a | (8.45, 19.55) | (1.48, 9.13) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.64 (1.16, 6.01) | |
| | | | P-value | 0.0208 | |
| | | | l.v. | | 1 10 |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 5 (29.4) | 2 (10.5) |
| | | | 95% CI ^a | (7.75, 51.07) | (0.00, 24.33) |
| | | | Relative Risk (95% CI) ^a | 2.79 (0.62, 12.57) | |
| | | | P-value | 0.1805 | |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 0 |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.17 (0.08, 60.76) | |
| | | | P-value | 0.6494 | |
| | | | Test for interaction ^b | 0.4840 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Acne | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 29 (16.4) | 7 (3.6) |
| | | 95% CI ^a | (10.93, 21.84) | (0.98, 6.20) | |
| | | 1 | Relative Risk (95% CI) ^a | 4.56 (2.05, 10.16) | 1 |
| | | | P-value | 4.56 (2.05, 10.16) 0.0002 | |
| | | | 1-value | 0.0002 | |
| | | D : 6 11 4 E | XI . | 150 | 122 |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 13 (8.7) | 1 (0.8) |
| | | | 95% CI ^a | (4.16, 13.17) | (0.00, 2.24) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 11.44 (1.52, 86.28) | |
| | | | P-value | 0.0181 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | n (%) | 4 (23.5) | 2 (10.5) |
| | | | 95% CI ^a | (3.37, 43.69) | (0.00, 24.33) |
| | | 1 | Relative Risk (95% CI) ^a | 2.24 (0.47, 10.70) | I |
| | | | P-value | 0.3141 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.1633 | |
| | | | | | |
| | Dermatitis atopic | Region of enrollment: US/Canada/Australia | N | 177 | 195 |
| | | | n (%) | 5 (2.8) | 8 (4.1) |
| | | | 95% CI ^a | (0.38, 5.27) | (1.32, 6.89) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.69 (0.23, 2.07) | |
| | | | P-value | 0.5056 | |
| | | | | | |
| | | Region of enrollment: Europe | N | 150 | 132 |
| | | | n (%) | 9 (6.0) | 6 (4.5) |
| | | | 95% CI ^a | (2.20, 9.80) | (0.99, 8.10) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.32 (0.48, 3.61) | |
| | | | P-value | 0.5886 | |
| | | | | | |
| | | Region of enrollment: Asia | N | 17 | 19 |
| | | | | | |
| | | | n (%) 95% CI ^a | 2 (11.8) | 0 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 4.59 (0.22, 94.96) | |
| | | | P-value | 0.3244 | Ì |
| | | | | | |
| | | Region of enrollment: Latin America | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 0 |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.17 (0.08, 60.76) | |
| | | | P-value | 0.6494 | |
| | | | | | |
| | | | Test for interaction ^b | | T . |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---------------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | | | |
| Overall | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 121 (56.0) | 76 (34.5) |
| | | | 95% CI ^a | (49.40, 62.64) | (28.26, 40.83) |
| | | | Relative Risk (95% CI) ^a | 1.62 (1.31, 2.01) | 1 |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 76 (52.1) | 53 (36.6) |
| | | | 95% CI ^a | (43.95, 60.16) | (28.71, 44.39) |
| | | | Relative Risk (95% CI) ^a | 1.42 (1.09, 1.86) | 1 |
| | | | P-value | 0.0089 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4029 | |
| | | | | | |
| astrointestinal disorders | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 45 (20.8) | 7 (3.2) |
| | | | 95% CI ^a | (15.42, 26.25) | (0.86, 5.50) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 6.55 (3.02, 14.20) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 25 (17.1) | 4 (2.8) |
| | | | 95% CI ^a | (11.01, 23.23) | (0.09, 5.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 6.21 (2.22, 17.39) | |
| | | | P-value | 0.0005 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4477 | |
| | | | | | |
| | Nausea | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 45 (20.8) | 7 (3.2) |
| | | | 95% CI ^a | (15.42, 26.25) | (0.86, 5.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 6.55 (3.02, 14.20) | |
| | | | P-value | <.0001 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 25 (17.1) | 1 (0.7) |
| | 95% CI ^a | (11.01, 23.23) | (0.00, 2.04) | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 24.83 (3.41, 180.82) | |
| | | | P-value | 0.0015 | |
| | | | Test for interaction ^b | 0.7016 | |
| | | | rest for interaction | 0.7010 | |
| | Vomiting | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 2 (0.9) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.00, 2.16) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.55 (0.50, 12.98) | |
| | | | P-value | 0.2608 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 6 (4.1) | 4 (2.8) |
| | | | 95% CI ^a | (0.89, 7.33) | (0.09, 5.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.49 (0.43, 5.17) | |
| | | | P-value | 0.5300 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9906 | 1 |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------------|--|-------------------------------------|------------------------------|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| General disorders and administration ite conditions | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 6 (2.8) | 3 (1.4) |
| | | | 95% CI ^a | (0.59, 4.97) | (0.00, 2.90) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.04 (0.52, 8.04) | |
| | | | P-value | 0.3098 | |
| | | | | | |
| | | Severe | N | 146 | 145 |
| | | | n (%) | 4 (2.7) | 2 (1.4) |
| | | | 95% CI ^a | (0.09, 5.39) | (0.00, 3.28) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.99 (0.37, 10.68) | |
| | | | P-value | 0.4238 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9801 | |
| | | | | | |
| | Fatigue | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 6 (2.8) | 3 (1.4) |
| | | | 95% CI ^a | (0.59, 4.97) | (0.00, 2.90) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-----------------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.04 (0.52, 8.04) | |
| | | | P-value | 0.3098 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 4 (2.7) | 2 (1.4) |
| | | | 95% CI ^a | (0.09, 5.39) | (0.00, 3.28) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.99 (0.37, 10.68) | |
| | | | P-value | 0.4238 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9801 | |
| | | | | | |
| infections and infestations | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 44 (20.4) | 45 (20.5) |
| | | | 95% CI ^a | (15.00, 25.74) | (15.12, 25.78) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.00 (0.69, 1.44) | |
| | | | P-value | 0.9826 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 24 (16.4) | 31 (21.4) |
| | | 95% CI ^a | | | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|--|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 0.77 (0.48, 1.24) | |
| | | | P-value | 0.2840 | |
| | | | Test for interaction ^b | 0.4225 | 1 |
| | | | Test for interaction | 0.4225 | |
| | COVID-19 Baseline disease severi Moderate | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 8 (3.7) | 7 (3.2) |
| | | | 95% CI ^a | (1.19, 6.22) | (0.86, 5.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.16 (0.43, 3.15) | |
| | | | P-value | 0.7652 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 7 (4.8) | 5 (3.4) |
| | | | 95% CI ^a | (1.33, 8.26) | (0.48, 6.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.39 (0.45, 4.28) | |
| | | | P-value | 0.5656 | |
| | | | Test for interaction ^b | 0.7757 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|--|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Conjunctivitis Baseline disease seve Moderate | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 3 (1.4) | 23 (10.5) |
| | | | 95% CI ^a | (0.00, 2.95) | (6.41, 14.50) |
| | | , | D 1 .: D: 1 (050) GD3 | | |
| | | | Relative Risk (95% CI) ^a | 0.13 (0.04, 0.44) | |
| | | | P-value | 0.0009 | |
| | | | | | |
| | | Baseline disease severity: N Severe | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 12 (8.3) |
| | | | 95% CI ^a | (0.47, 6.37) | (3.79, 12.76) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.41 (0.15, 1.14) | |
| | | | P-value | 0.0893 | |
| | | | I | | |
| | | | Test for interaction ^b | 0.2401 | |
| | Folliculitis | Baseline disease severity: | N | 216 | 220 |
| | Tonicultis | Moderate Moderate | 11 | 210 | 220 |
| | | | n (%) | 9 (4.2) | 1 (0.5) |
| | | | 95% CI ^a | (1.50, 6.83) | (0.00, 1.34) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 9.17 (1.17, 71.74) | |
| | | | P-value | 0.0348 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | n | n (%) | 3 (2.1) | 2 (1.4) |
| | | | 95% CI ^a | (0.00, 4.36) | (0.00, 3.28) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.49 (0.25, 8.78) | |
| | | | P-value | 0.6597 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1476 | |
| | | | | | |
| | Herpes simplex | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 7 (3.2) | 1 (0.5) |
| | | | 95% CI ^a | (0.88, 5.60) | (0.00, 1.34) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 7.13 (0.88, 57.46) | |
| | | | P-value | 0.0651 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 4 (2.8) |
| | | | 95% CI ^a | (0.47, 6.37) | (0.09, 5.42) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.24 (0.34, 4.53) | |
| | | | P-value | 0.7433 | |
| | | | Test for interaction ^b | 0.3854 | |
| | | | Test for interaction | 0.3034 | |
| | Nasopharyngitis | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 9 (4.2) | 10 (4.5) |
| | | | 95% CI ^a | (1.50, 6.83) | (1.79, 7.30) |
| | · | | | | |
| | | | Relative Risk (95% CI) ^a | 0.92 (0.38, 2.21) | |
| | | | P-value | 0.8465 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 2 (1.4) |
| | | | 95% CI ^a | (0.47, 6.37) | (0.00, 3.28) |
| | | | 30 70 01 | (0.77, 0.37) | |
| | | | 90 /0 61 | (0.47, 0.57) | (****,**23) |
| | | | Relative Risk (95% CI) ^a | 2.48 (0.49, 12.59) | (2.22) |
| | | | | | (3103, 1203) |
| | | | Relative Risk (95% CI) ^a | 2.48 (0.49, 12.59) | (0000) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------|-----------------------------------|--|-------------------------------------|------------------------------|--------------------------------|
| ystem Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Oral herpes | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 5 (2.3) | 7 (3.2) |
| | | | 95% CI ^a | (0.31, 4.32) | (0.86, 5.50) |
| | | | D 1 D' 1 (050) GD3 | | |
| | | | Relative Risk (95% CI) ^a | 0.73 (0.23, 2.26) | |
| | | | P-value | 0.5818 | |
| | | | | | |
| | | Baseline disease severity: Nevere | N | 146 | 145 |
| | | | n (%) | 4 (2.7) | 8 (5.5) |
| | | | 95% CI ^a | (0.09, 5.39) | (1.80, 9.23) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.50 (0.15, 1.61) | |
| | | | P-value | 0.2442 | |
| | | | m of the h | | |
| | | | Test for interaction ^b | 0.4927 | |
| | Upper respiratory tract infection | Rasalina disaasa savarity | N | 216 | 220 |
| | opper respiratory tract infection | Moderate | 14 | 210 | 220 |
| | | | n (%) | 9 (4.2) | 5 (2.3) |
| | | | 95% CI ^a | (1.50, 6.83) | (0.30, 4.24) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.83 (0.62, 5.38) | |
| | | | P-value | 0.2700 | |
| | | | | | |
| | | Baseline disease severity: N Severe | N | 146 | 145 |
| | | | n (%) | 1 (0.7) | 4 (2.8) |
| | | | 95% CI ^a | (0.00, 2.02) | (0.09, 5.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.25 (0.03, 2.19) | |
| | | | P-value | 0.2102 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0828 | |
| | | | | | |
| Investigations | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 24 (11.1) | 15 (6.8) |
| | | | 95% CI ^a | (6.92, 15.30) | (3.49, 10.15) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.63 (0.88, 3.02) | |
| | | | P-value | 0.1209 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 14 (9.6) | 11 (7.6) |
| | | | 95% CI ^a | (4.81, 14.37) | (3.28, 11.90) |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|--|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.26 (0.59, 2.69) | |
| | | | P-value | 0.5433 | |
| | | | Test for interaction ^b | 0.5006 | 1 |
| | | | Test for interaction | 0.5906 | |
| | Blood creatine phosphokinase increased | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 9 (4.2) | 7 (3.2) |
| | | | 95% CI ^a | (1.50, 6.83) | (0.86, 5.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.31 (0.50, 3.45) | |
| | | | P-value | 0.5857 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 6 (4.1) |
| | | | 95% CI ^a | (0.47, 6.37) | (0.90, 7.38) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.83 (0.26, 2.65) | |
| | | | P-value | 0.7501 | |
| | | | | | |
| | | | Test for interaction ^b | 0.5547 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|--|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Natural killer cell count decreased Baseline disease severity: Moderate | N | 216 | 220 | |
| | | | n (%) | 6 (2.8) | 0 |
| | | | 95% CI ^a | (0.59, 4.97) | (0.00, 1.66) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 12.25 (0.69, 217.98) | |
| | | | P-value | 0.0880 | |
| | | | | | |
| | | Baseline disease severity: N Severe | N | 146 | 145 |
| | | | n (%) | 4 (2.7) | 0 |
| | | | 95% CI ^a | (0.09, 5.39) | (0.00, 2.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 7.97 (0.43, 149.45) | |
| | | | P-value | 0.1651 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9343 | |
| | | | | | |
| | SARS-CoV-2 test positive | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 10 (4.6) | 8 (3.6) |
| | | | 95% CI ^a | (1.83, 7.43) | (1.16, 6.11) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 1.27 (0.51, 3.16) | |
| | | | P-value | 0.6032 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 5 (3.4) |
| | | | 95% CI ^a | (0.47, 6.37) | (0.48, 6.42) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.99 (0.29, 3.36) | |
| | | | P-value | 0.9912 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7215 | |
| | | | | | |
| Nervous system disorders | Overall | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 30 (13.9) | 18 (8.2) |
| | | | 95% CI ^a | (9.28, 18.50) | (4.56, 11.80) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.70 (0.98, 2.95) | |
| | | | P-value | 0.0609 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 25 (17.1) | 9 (6.2) |
| | | 95% CI ^a | (11.01, 23.23) | (2.28, 10.13) | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2' (N=365) |
|--------------------|-----------------------|--|---|--|-----------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.76 (1.33, 5.70) | |
| | | | P-value | 0.0062 | |
| | | 1 | Test for interaction ^b | 0.2787 | 1 |
| | | | Test for interaction | 0.2787 | |
| | Dizziness | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 4 (1.9) | 3 (1.4) |
| | | | 95% CI ^a | (0.05, 3.65) | (0.00, 2.90) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.36 (0.31, 6.00) | I |
| | | | P-value | 0.6863 | |
| | | | | | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | N n (%) | 6 (4.1) | 145 |
| | | | | | |
| | | | n (%) | 6 (4.1) | 1 (0.7) |
| | | | n (%) | 6 (4.1) | 1 (0.7) |
| | | | n (%) 95% CI ^a | 6 (4.1) (0.89, 7.33) | 1 (0.7) |
| | | | n (%) 95% CI ^a Relative Risk (95% CI) ^a | 6 (4.1) (0.89, 7.33) 5.96 (0.73, 48.88) | 1 (0.7) |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | Headache | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 27 (12.5) | 15 (6.8) |
| | | | 95% CI ^a | (8.09, 16.91) | (3.49, 10.15) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.83 (1.00, 3.35) | |
| | | | P-value | 0.0487 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 20 (13.7) | 9 (6.2) |
| | | | 95% CI ^a | (8.12, 19.28) | (2.28, 10.13) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.21 (1.04, 4.68) | |
| | | | P-value | 0.0392 | |
| | | | I | | |
| | | | Test for interaction ^b | 0.6865 | |
| Skin and subcutaneous tissue disorders | Overall | Baseline disease severity: | N | 216 | 220 |
| okin and subcutaneous tissue disorders | Overall | Moderate | IN . | 210 | 220 |
| | | | n (%) | 34 (15.7) | 9 (4.1) |
| | | | 95% CI ^a | (10.88, 20.60) | (1.47, 6.71) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 3.85 (1.89, 7.83) | |
| | | | P-value | 0.0002 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 27 (18.5) | 14 (9.7) |
| | | | 95% CI ^a | (12.20, 24.79) | (4.85, 14.46) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.92 (1.05, 3.50) | |
| | | | P-value | 0.0347 | |
| | | | | | |
| | | | Test for interaction ^b | 0.6464 | |
| | | | | | |
| | Acne | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 29 (13.4) | 2 (0.9) |
| | | | 95% CI ^a | (8.88, 17.97) | (0.00, 2.16) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 14.77 (3.57, 61.13) | |
| | | | P-value | 0.0002 | |
| | | | | | |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 17 (11.6) | 8 (5.5) |
| | | | 95% CI ^a | (6.44, 16.85) | (1.80, 9.23) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2V (N=365) |
|--------------------|-----------------------|--|-------------------------------------|------------------------------|--------------------------------|
| System Organ Class | MedDRA Preferred Term | Subgroup | | n (%) | n (%) |
| | | | Relative Risk (95% CI) ^a | 2.11 (0.94, 4.74) | |
| | | | P-value | 0.0701 | |
| | | | Test for interaction ^b | 0.1246 | 1 |
| | | | Test for interaction | 0.1346 | |
| | Dermatitis atopic | Baseline disease severity: Moderate | N | 216 | 220 |
| | | | n (%) | 7 (3.2) | 8 (3.6) |
| | | | 95% CI ^a | (0.88, 5.60) | (1.16, 6.11) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.89 (0.33, 2.41) | |
| | | | P-value | 0.8208 | |
| | | | | | 1 |
| | | Baseline disease severity: Severe | N | 146 | 145 |
| | | | n (%) | 10 (6.8) | 6 (4.1) |
| | | | 95% CI ^a | (2.75, 10.95) | (0.90, 7.38) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.66 (0.62, 4.44) | |
| | | | P-value | 0.3163 | |
| | | | | | |
| | | | Test for interaction ^b | 0.3289 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:18)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:29)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | . , | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 173 (75.2) | 161 (65.2) |
| | | | 95% CI ^a | (69.64, 80.80) | (59.24, 71.12) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.15 (1.03, 1.30) | |
| | | | P-value | 0.0169 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 94 (71.2) | 76 (64.4) |
| | | | 95% CI ^a | (63.49, 78.94) | (55.77, 73.05) |
| | | | Relative Risk (95% CI) ^a | 1 11 (0.02 1.21) | |
| | | | P-value | 1.11 (0.93, 1.31) | |
| | | | r-value | 0.2538 | |
| | | | Test for interaction ^b | 0.5853 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 141 (73.1) | 127 (62.3) |
| | | | 95% CI ^a | (66.80, 79.32) | (55.60, 68.91) |
| | | | Relative Risk (95% CI) ^a | 1.17 (1.02, 1.35) | |
| | | | P-value | 0.0220 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 126 (74.6) | 110 (68.3) |
| | | | 95% CI ^a | (67.99, 81.12) | (61.14, 75.51) |
| | | | Relative Risk (95% CI) ^a | 1.09 (0.95, 1.25) | I |
| | | | P-value | 0.2123 | |
| | | | | | |
| | | | Test for interaction ^b | 0.6078 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|---------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 133 (75.1) | 128 (65.6) |
| | | | 95% CI ^a | (68.77, 81.51) | (58.98, 72.31) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.14 (1.00, 1.31) | |
| | | | P-value | 0.0452 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 111 (74.0) | 90 (68.2) |
| | | | 95% CI ^a | (66.98, 81.02) | (60.24, 76.13) |
| | | | D 1 (' D' 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 1.09 (0.93, 1.26) | |
| | | | P-value | 0.2855 | |
| | _ | | | | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| Region of enrollment: Asia | Overall | Overall | n (%) | 11 (64.7) | 10 (52.6) |
| Region of enrollment: Asia | Overall | Overall | | · | · |
| Region of enrollment: Asia | Overall | Overall | n (%) | 11 (64.7) | 10 (52.6) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|---------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 12 (66.7) | 9 (47.4) |
| | | | 95% CI ^a | (44.89, 88.44) | (24.92, 69.82) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.41 (0.79, 2.50) | |
| | | | P-value | 0.2446 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9324 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|---|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | , , |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 162 (75.0) | 146 (66.4) |
| | | | 95% CI ^a | (69.23, 80.77) | (60.12, 72.61) |
| | | | *************************************** | | |
| | | | Relative Risk (95% CI) ^a | 1.13 (1.00, 1.28) | |
| | | | P-value | 0.0486 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 105 (71.9) | 91 (62.8) |
| | | | 95% CI ^a | (64.63, 79.21) | (54.89, 70.63) |
| | 0 | -70-1 | -7- | | |
| | | | Relative Risk (95% CI) ^a | 1.15 (0.98, 1.35) | |
| | | | P-value | 0.0977 | |
| | | | | | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| Subgroup | Ставв | 101111 | | (/0) | (/0) |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 19 (8.3) | 15 (6.1) |
| | | | 95% CI ^a | (4.70, 11.82) | (3.09, 9.05) |
| | | | Relative Risk (95% CI) ^a | 1.36 (0.71, 2.61) | |
| | | | P-value | 0.3555 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 12 (9.1) | 7 (5.9) |
| | | | 95% CI ^a | (4.19, 14.00) | (1.67, 10.19) |
| | | | Relative Risk (95% CI) ^a | 1.53 (0.62, 3.76) | |
| | | | P-value | 0.3517 | |
| | | | | | |
| | | | Test for interaction ^b | 0.8103 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|---|------------------------------|--------------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Sex: Male | Overell | Overall | N | 193 | 204 |
| Sex. Male | Overan | Overall | n (%) | 19 (9.8) | 12 (5.9) |
| | | | 95% CI ^a | (5.64, 14.05) | (2.65, 9.11) |
| | ľ | | Polotive Piels (050/ CI) | 1 57 (0.00 0.05) | |
| | | | Relative Risk (95% CI) ^a P-value | 1.67 (0.83, 3.35) 0.1467 | |
| | | | 1 | 0.1407 | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 12 (7.1) | 10 (6.2) |
| | | | 95% CI ^a | (3.23, 10.97) | (2.48, 9.94) |
| | | | Relative Risk (95% CI) ^a | 1.14 (0.51, 2.57) | |
| | | | P-value | 0.7464 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4216 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) | |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|-----------|
| | System Organ | MedDRA Preferred | | | | |
| Subgroup | Class | Term | | n (%) | n (%) | |
| | | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 | |
| | | | n (%) | 14 (7.9) | 11 (5.6) | |
| | | | 95% CI ^a | (3.93, 11.89) | (2.40, 8.88) | |
| | | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.40 (0.65, 3.01) | | |
| | | | P-value | 0.3853 | | |
| | | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 | |
| | | | | n (%) | 15 (10.0) | 11 (8.3) |
| | | | 95% CI ^a | (5.20, 14.80) | (3.62, 13.05) | |
| | | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.20 (0.57, 2.52) | | |
| | | | P-value | 0.6301 | | |
| | | | 1 | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 | |
| | | | n (%) | 1 (5.9) | 0 | |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 17.65) | |
| | | | Relative Risk (95% CI) ^a | 2.20 (0.09 (4.21) | T | |
| | | | P-value | 2.29 (0.08, 64.21) | | |
| | | | r-value | 0.6252 | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 1 (5.6) | 0 |
| | | | 95% CI ^a | (0.00, 16.14) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.17 (0.08, 60.76) | |
| | | | P-value | 0.6494 | |
| | | | | | |
| | | | Test for interaction ^b | 0.9965 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 18 (8.3) | 11 (5.0) |
| | | | 95% CI ^a | (4.65, 12.02) | (2.12, 7.88) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.67 (0.81, 3.45) | |
| | | | P-value | 0.1680 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 13 (8.9) | 11 (7.6) |
| | | | 95% CI ^a | (4.28, 13.52) | (3.28, 11.90) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.17 (0.54, 2.53) | |
| | | | P-value | 0.6832 | |
| | | 20 | | | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 4 (1.7) | 1 (0.4) |
| | | | 95% CI ^a | (0.05, 3.43) | (0.00, 1.20) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.30 (0.48, 38.15) | |
| | | | P-value | 0.1908 | |
| | | | | | |
| Age (years) group: >=40 | Overall Overall | | N | 132 | 118 |
| | | | n (%) | 5 (3.8) | 1 (0.8) |
| | | | 95% CI ^a | (0.53, 7.04) | (0.00, 2.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.47 (0.53, 37.71) | |
| | | | P-value | 0.1688 | |
| | | | | | |
| | | | Test for interaction ^b | 0.4405 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|----------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| Sex. Wate | Overan | Overan | n (%) | 5 (2.6) | 1 (0.5) |
| | | | 95% CI ^a | (0.35, 4.83) | (0.00, 1.45) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 5.28 (0.62, 44.83) | |
| | | | P-value | 0.1270 | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| Sex. I ciliale | Overan | Overall | n (%) | 4 (2.4) | 1 (0.6) |
| | | | 95% CI ^a | (0.07, 4.66) | (0.00, 1.83) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.81 (0.43, 33.73) | |
| | | | P-value | 0.2292 | |
| | | | | | |
| | | | Test for interaction ^b | 0.8457 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|---------|---------------------|-------------------------------------|---------------------------------|--------------------------------|
| | Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 5 (2.8) | 1 (0.5) |
| | | | 95% CI ^a | (0.38, 5.27) | (0.00, 1.52) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 5.51 (0.65, 46.70) | |
| | | | P-value | 0.1177 | |
| | | | l | 1-0 | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 1 (0.8) |
| | | | 95% CI ^a | (0.09, 5.24) | (0.00, 2.24) |
| | | | Relative Risk (95% CI) ^a | 3.52 (0.40, 31.10) | |
| | | | P-value | 0.2576 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 0 | 0 |
| D : C 11 / I / A : | 0 " | 0 11 | N | 10 | 10 |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | Test for interaction ^b | 0.0626 | |
| | | | 1 est for interaction | 0.9636 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Suogroup | Class | Term | | п (/0) | 11 (70) |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 4 (1.9) | 2 (0.9) |
| | | | 95% CI ^a | (0.05, 3.65) | (0.00, 2.16) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.04 (0.38, 11.01) | |
| | | | P-value | 0.4084 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 5 (3.4) | 0 |
| | | | 95% CI ^a | (0.47, 6.37) | (0.00, 2.51) |
| | | | Relative Risk (95% CI) ^a | 0.07 (0.55, 190.75) | |
| | | | | 9.97 (0.55, 180.76) | - |
| | | | P-value | 0.1200 | |
| | | | Test for interaction ^b | 0.2721 | |

Included data up to 28 days after last dose of study.

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| <u> </u> | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 34 (14.8) | 10 (4.0) |
| | | | 95% CI ^a | (10.20, 19.37) | (1.59, 6.51) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.65 (1.85, 7.22) | |
| | | | P-value | 0.0002 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 14 (10.6) | 1 (0.8) |
| | | | 95% CI ^a | (5.35, 15.86) | (0.00, 2.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 12.52 (1.67, 93.73) | |
| | | | P-value | 0.0139 | |
| | | | | | |
| | | | Test for interaction ^b | 0.7300 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:26)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| C M - 1 - | O11 | O11 | N | 102 | 204 |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 26 (13.5) | 4 (2.0) |
| | | | 95% CI ^a | (8.65, 18.29) | (0.06, 3.86) |
| | | | Relative Risk (95% CI) ^a | 6.87 (2.44, 19.32) | |
| | | | P-value | 0.0003 | |
| | _ | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 22 (13.0) | 7 (4.3) |
| | | | 95% CI ^a | (7.94, 18.09) | (1.20, 7.50) |
| | | | D 1 4' D' 1 (050/ CD3 | | |
| | | | Relative Risk (95% CI) ^a | 2.99 (1.32, 6.82) | |
| | | | P-value | 0.0090 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:26)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|---|-------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 30 (16.9) | 8 (4.1) |
| | | | 95% CI ^a | (11.42, 22.48) | (1.32, 6.89) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.13 (1.95, 8.77) | |
| | | | P-value | 0.0002 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 14 (9.3) | 1 (0.8) |
| | | | 95% CI ^a | (4.68, 13.99) | (0.00, 2.24) |
| | | | | | |
| | | | D 1 (' D' 1 (050) CD3 | | |
| | | | Relative Risk (95% CI) ^a | 12.32 (1.64, 92.43) | |
| | | | Relative Risk (95% CI) ^a P-value | 12.32 (1.64, 92.43) 0.0146 | |
| | | | P-value | 0.0146 | |
| Region of enrollment: Asia | Overall | Overall | P-value | 0.0146 17 | 19 |
| Region of enrollment: Asia | Overall | Overall | P-value N n (%) | 0.0146 17 4 (23.5) | 2 (10.5) |
| Region of enrollment: Asia | Overall | Overall | P-value | 0.0146 17 | · |
| Region of enrollment: Asia | Overall | Overall | P-value N n (%) | 0.0146 17 4 (23.5) | 2 (10.5) |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:26)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-----------------------------------|---------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.1721 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:26)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| ouog.oup | Class | 101111 | | (,,,) | (/0) |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 30 (13.9) | 3 (1.4) |
| | | | 95% CI ^a | (9.28, 18.50) | (0.00, 2.90) |
| | | | Relative Risk (95% CI) ^a | 10.19 (3.16, 32.88) | T |
| | | | P-value | 0.0001 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 18 (12.3) | 8 (5.5) |
| | | | 95% CI ^a | (7.00, 17.66) | (1.80, 9.23) |
| | | | D 1 1 D 1 1 (050) GD3 | | |
| | | | Relative Risk (95% CI) ^a | 2.23 (1.00, 4.98) | |
| | | | P-value | 0.0490 | |
| | | | Test for interaction ^b | 0.1911 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:26)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| C 1 | | | | | , |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 7 (3.0) | 2 (0.8) |
| | | | 95% CI ^a | (0.82, 5.26) | (0.00, 1.93) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.76 (0.79, 17.91) | |
| | | | P-value | 0.0965 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 4 (3.0) | 1 (0.8) |
| | | | 95% CI ^a | (0.11, 5.95) | (0.00, 2.50) |
| | | | Relative Risk (95% CI) ^a | 2.50 (0.44.24.54) | |
| | | | ` ' | 3.58 (0.41, 31.54) | |
| | | | P-value | 0.2514 | |
| | | | Test for interaction ^b | 0.9812 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 9 (4.7) | 2 (1.0) |
| | | | 95% CI ^a | (1.69, 7.64) | (0.00, 2.33) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.76 (1.04, 21.74) | |
| | | | P-value | 0.0443 | |
| Sex: Female | O11 | O11 | N | 169 | 161 |
| Sex: Female | Overall | Overall | | | |
| | | | n (%) | 2 (1.2) | 1 (0.6) |
| | | | 95% CI ^a | (0.00, 2.81) | (0.00, 1.83) |
| | | | Relative Risk (95% CI) ^a | 1.91 (0.17, 20.81) | |
| | | | P-value | 0.5972 | |
| | | | | | |
| | | | Test for interaction ^b | 0.1122 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 4 (2.3) | 3 (1.5) |
| | | | 95% CI ^a | (0.07, 4.45) | (0.00, 3.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.47 (0.33, 6.47) | |
| | | | P-value | 0.6113 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 5 (3.3) | 0 |
| | | | 95% CI ^a | (0.46, 6.21) | (0.00, 2.76) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 8.83 (0.49, 160.17) | |
| | | | P-value | 0.1406 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 0 |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 17.65) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 4.59 (0.22, 94.96) | |
| | | | P-value | 0.3244 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|---------|-----------------------------|-----------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.5893 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 8 (3.7) | 1 (0.5) |
| | | | 95% CI ^a | (1.19, 6.22) | (0.00, 1.34) |
| | | | 7 | | |
| | | | Relative Risk (95% CI) ^a | 8.15 (1.03, 64.60) | |
| | | | P-value | 0.0470 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 3 (2.1) | 2 (1.4) |
| | | | 95% CI ^a | (0.00, 4.36) | (0.00, 3.28) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.49 (0.25, 8.78) | |
| | | | P-value | 0.6597 | |
| | | | | | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| Subgroup | Class | Term | | 11 (70) | 11 (70) |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 4 (1.7) | 25 (10.1) |
| | | | 95% CI ^a | (0.05, 3.43) | (6.36, 13.88) |
| | | | Relative Risk (95% CI) ^a | 0.17 (0.06, 0.49) | |
| | | | P-value | 0.0009 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 6 (4.5) | 14 (11.9) |
| | | | 95% CI ^a | (0.99, 8.10) | (6.03, 17.70) |
| | | | Relative Risk (95% CI) ^a | 0.38 (0.15, 0.96) | |
| | | | P-value | 0.0417 | |
| | | | | | |
| | | | Test for interaction ^b | 0.8350 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 9 (4.7) | 22 (10.8) |
| | | | 95% CI ^a | (1.69, 7.64) | (6.53, 15.04) |
| | | | Relative Risk (95% CI) ^a | 0.43 (0.20, 0.92) | 1 |
| | | | P-value | 0.0285 | |
| | | | | | |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 1 (0.6) | 17 (10.6) |
| | | | 95% CI ^a | (0.00, 1.75) | (5.81, 15.31) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.06 (0.01, 0.42) | |
| | | | P-value | 0.0049 | |
| | | I | The Control of the | | |
| | | | Test for interaction ^b | 0.3281 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 4 (2.3) | 22 (11.3) |
| | | | 95% CI ^a | (0.07, 4.45) | (6.84, 15.72) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.20 (0.07, 0.57) | |
| | | | P-value | 0.0026 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 4 (2.7) | 16 (12.1) |
| | | | 95% CI ^a | (0.09, 5.24) | (6.55, 17.69) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.22 (0.08, 0.64) | |
| | | | P-value | 0.0056 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 2 (11.8) | 1 (5.3) |
| | | | 95% CI ^a | (0.00, 27.08) | (0.00, 15.30) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 2.24 (0.22, 22.51) | |
| | | | P-value | 0.4948 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-----------------------------------|---------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.1573 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|--------------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 3 (1.4) | 28 (12.7) |
| | | | 95% CI ^a | (0.00, 2.95) | (8.32, 17.13) |
| | | | Relative Risk (95% CI) ^a | 0.11 (0.03, 0.35) | I |
| | | | P-value | 0.0002 | |
| | | | | | |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 7 (4.8) | 11 (7.6) |
| | | | 95% CI ^a | (1.33, 8.26) | (3.28, 11.90) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.63 (0.25, 1.58) | |
| | | | P-value | 0.3280 | |
| | | | | | |
| | | | Test for interaction ^b | 0.0225 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Proportion of Subjects with Serious Superinfections of Infections (Treated with Antibiotics, Antiviral or Fungicidal Agents for a Duration > 2 Weeks) Treatment-Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set (Protocol B7451050)

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Proportion of Subjects with Severe Superinfections of Infections (Treated with Antibiotics, Antiviral or Fungicidal Agents for a Duration > 2 Weeks) Treatment - Emergent Adverse Events by Subgroup (All Causalities) - Safety Analysis Set (Protocol B7451050)

Table not created

No participant meets the reporting criteria

Included data up to 28 days after last dose of study.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

Age group (<40, >=40)

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Age (years) group: <40 | Overall | Overall | N | 230 | 247 |
| | | | n (%) | 5 (2.2) | 8 (3.2) |
| | | | 95% CI ^a | (0.29, 4.06) | (1.03, 5.45) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.67 (0.22, 2.02) | |
| | | | P-value | 0.4786 | |
| | | | | | |
| Age (years) group: >=40 | Overall | Overall | N | 132 | 118 |
| | | | n (%) | 4 (3.0) | 1 (0.8) |
| | | | 95% CI ^a | (0.11, 5.95) | (0.00, 2.50) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 3.58 (0.41, 31.54) | |
| | | | P-value | 0.2514 | |
| | .11 | | | | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Sex

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------|---------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | Organ | MedDRA Preferred Term | | n (%) | n (%) |
| Sex: Male | Overall | Overall | N | 193 | 204 |
| | | | n (%) | 2 (1.0) | 4 (2.0) |
| | | | 95% CI ^a | (0.00, 2.46) | (0.06, 3.86) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.53 (0.10, 2.85) | |
| | | | P-value | 0.4585 | |
| a 1 | 0 11 | 0 11 | | 4.50 | 1.51 |
| Sex: Female | Overall | Overall | N | 169 | 161 |
| | | | n (%) | 7 (4.1) | 5 (3.1) |
| | | | 95% CI ^a | (1.14, 7.15) | (0.43, 5.79) |
| | | | Relative Risk (95% CI) ^a | 1.33 (0.43, 4.12) | |
| | | | P-value | 0.6165 | |
| | | | · | | |
| | | | Test for interaction ^b | 0.4137 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|---|-----------------|---------------------|-------------------------------------|------------------------------|--------------------------------|
| | System Organ | MedDRA Preferred | | | |
| Subgroup | Class | Term | | n (%) | n (%) |
| | | | | | |
| Region of enrollment: US/Canada/Australia | Overall | Overall | N | 177 | 195 |
| | | | n (%) | 5 (2.8) | 3 (1.5) |
| | | | 95% CI ^a | (0.38, 5.27) | (0.00, 3.27) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.84 (0.45, 7.57) | |
| | | | P-value | 0.4006 | |
| | | | | | |
| Region of enrollment: Europe | Overall | Overall | N | 150 | 132 |
| | | | n (%) | 3 (2.0) | 6 (4.5) |
| | | | 95% CI ^a | (0.00, 4.24) | (0.99, 8.10) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 0.44 (0.11, 1.72) | |
| | | | P-value | 0.2388 | |
| | | | | | |
| Region of enrollment: Asia | Overall | Overall | N | 17 | 19 |
| | | | n (%) | 1 (5.9) | 0 |
| | | | 95% CI ^a | (0.00, 17.07) | (0.00, 17.65) |
| | | | D 1 (' D' 1 (050) CD' | | |
| | | | Relative Risk (95% CI) ^a | 2.29 (0.08, 64.21) | |
| | | | P-value | 0.6252 | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Region

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-----------------------------------|---------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| Region of enrollment: Latin America | Overall | Overall | N | 18 | 19 |
| | | | n (%) | 0 | 0 |
| | | | | | |
| | | | Test for interaction ^b | 0.5066 | |
| | | | | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Baseline disease severity

| | | | | Abrocitinib 200mg QD (N=362) | Dupilumab 300mg Q2W (N=365) |
|-------------------------------------|--------------------------|-----------------------------|-------------------------------------|------------------------------|-----------------------------|
| Subgroup | System Organ Class | MedDRA Preferred Term | | n (%) | n (%) |
| | | | | | |
| Baseline disease severity: Moderate | Overall | Overall | N | 216 | 220 |
| | | | n (%) | 7 (3.2) | 5 (2.3) |
| | | | 95% CI ^a | (0.88, 5.60) | (0.30, 4.24) |
| | | | | | |
| | | | Relative Risk (95% CI) ^a | 1.43 (0.46, 4.42) | |
| | | | P-value | 0.5391 | |
| | | | | | -0 |
| Baseline disease severity: Severe | Overall | Overall | N | 146 | 145 |
| | | | n (%) | 2 (1.4) | 4 (2.8) |
| | | | 95% CI ^a | (0.00, 3.26) | (0.09, 5.42) |
| | Con | | -0- | | |
| | | | Relative Risk (95% CI) ^a | 0.50 (0.09, 2.67) | |
| | | | P-value | 0.4146 | |
| | | | | | |
| | | | Test for interaction ^b | | |

Included data up to 28 days after last dose of study.

The two sided P-value was based on a type 3 wald test with the null hypothesis: all the parameters associated with the interaction effect (treatment by subgroup category) were zero. MedDRA v24.0 coding dictionary applied.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (02:02) Source Data: adae Table Generation: 26SEP2021 (07:28)

Subjects were counted only once per treatment per event. .5 was added to 0 count cells when RR can't be calculated.

a. Results calculated with normal approximation.

b. For test of treatment and subgroup interaction, a log regression model with covariates treatment, subgroup category and treatment by subgroup interaction was used.

Non-Severe Treatment Emergent Subset Flag Adverse Events, overall and by SOC/PT - Safety Population

| System Organ Class (SOC) | Abrocitinib 200mg QD | Dupilumab 300mg Q2W | Total Population |
|--|----------------------|---------------------|------------------|
| Preferred Term (PT) | N=362 | N=365 | N=727 |
| Overall | 195 (53.9%) | 129 (35.3%) | 324 (44.6%) |
| Infections and infestations | 67 (18.5%) | 76 (20.8%) | 143 (19.7%) |
| Conjunctivitis | 8 (2.2%) | 35 (9.6%) | 43 (5.9%) |
| COVID-19 | 14 (3.9%) | 12 (3.3%) | 26 (3.6%) |
| Nasopharyngitis | 14 (3.9%) | 12 (3.3%) | 26 (3.6%) |
| Oral herpes | 9 (2.5%) | 15 (4.1%) | 24 (3.3%) |
| Upper respiratory tract infection | 10 (2.8%) | 9 (2.5%) | 19 (2.6%) |
| Herpes simplex | 12 (3.3%) | 5 (1.4%) | 17 (2.3%) |
| Folliculitis | 12 (3.3%) | 3 (0.8%) | 15 (2.1%) |
| Skin and subcutaneous tissue disorders | 60 (16.6%) | 23 (6.3%) | 83 (11.4%) |
| Acne | 46 (12.7%) | 10 (2.7%) | 56 (7.7%) |
| Dermatitis atopic | 16 (4.4%) | 14 (3.8%) | 30 (4.1%) |
| Gastrointestinal disorders | 70 (19.3%) | 11 (3.0%) | 81 (11.1%) |
| Nausea | 70 (19.3%) | 8 (2.2%) | 78 (10.7%) |
| Vomiting | 11 (3.0%) | 6 (1.6%) | 17 (2.3%) |
| Nervous system disorders | 54 (14.9%) | 27 (7.4%) | 81 (11.1%) |
| Headache | 46 (12.7%) | 24 (6.6%) | 70 (9.6%) |
| Dizziness | 10 (2.8%) | 4 (1.1%) | 14 (1.9%) |
| Investigations | 38 (10.5%) | 26 (7.1%) | 64 (8.8%) |
| SARS-CoV-2 test positive | 15 (4.1%) | 13 (3.6%) | 28 (3.9%) |
| Blood creatine phosphokinase increased | 14 (3.9%) | 13 (3.6%) | 27 (3.7%) |
| Natural killer cell count decreased | 10 (2.8%) | 0 | 10 (1.4%) |
| General disorders and administration site conditions | 10 (2.8%) | 5 (1.4%) | 15 (2.1%) |
| Fatigue | 10 (2.8%) | 5 (1.4%) | 15 (2.1%) |

Note(s):
Safety Population
CTCAE: Common Terminology Criteria for Adverse Events; PT: (MedDRA) Preferred Term; SOC: (MedDRA) System Organ Class.
Analysis Cut Off date: 06AUG2021
[[root]\02 Programs\02.05 Safety Incidences SOCPT.sas] run Thursday, November 11, 2021 at 11:26:34

Responderanalysen JADE DARE (Woche 12)

Binary Outcome Analysis: SCORAD-75 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300n | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|---------------------------|-----------|------------------|------------|------------|--------------|-----------------|---------------------------|-------------------|---------------|
| | | | | | | | _ | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 362 | 133 (36.7%) | 365 | 93 (25.5%) | 1.70 | 1.44 | 11.3% | 0.0010+* | |
| Population | | (31.8%, | | (21.1%, | (1.24, 2.34) | (1.16, 1.80) | (4.6%, 18.0%) | | |
| 1 | | 41.9%) | | 30.3%) | , , | , , , | , , , | | |
| Baseline Disease Severity | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 216 | 80 (37.0%) | 220 | 58 (26.4%) | 1.64 | 1.40 | 10.7% | 0.0180* | 0.8589 |
| disease (IGA=3) | | (30.6%, | | (20.7%, | (1.09, 2.47) | (1.06, 1.86) | (2.0%, 19.4%) | | |
| | | 43.9%) | | 32.7%) | (, , | (,, | (,, | | |
| Severe baseline disease | 146 | 53 (36.3%) | 145 | 35 (24.1%) | 1.79 | 1.50 | 12.2% | 0.0298* | |
| (IGA=4) | | (28.5%, | | (17.4%, | (1.08, 2.98) | (1.05, 2.16) | (1.7%, 22.6%) | 0.0-7-0 | |
| (1011 .) | | 44.7%) | | 31.9%) | (1100, 2170) | (1.02, 2.10) | (11,70, 221070) | | |
| Gender | | ,0) | | 211770) | | | | | |
| Week 12 | | | | | | | | | |
| Male | 193 | 64 (33.2%) | 204 | 47 (23.0%) | 1.66 | 1.44 | 10.1% | 0.0258* | 0.6462 |
| 111110 | 173 | (26.6%, | 204 | (17.4%, | (1.06, 2.58) | (1.04, 1.98) | (1.3%, 18.9%) | 0.0230 | 0.0402 |
| | | 40.3%) | | 29.4%) | (1.00, 2.50) | (1.04, 1.70) | (1.570, 10.570) | | |
| Female | 169 | 69 (40.8%) | 161 | 46 (28.6%) | 1.73 | 1.43 | 12.3% | 0.0211* | |
| Temale | 109 | , , | 101 | , | (1.09, 2.73) | (1.05, 1.94) | (2.1%, 22.4%) | 0.0211 | |
| | | (33.3%, | | (21.7%, | (1.09, 2.75) | (1.03, 1.94) | (2.1%, 22.4%) | | |
| | | 48.6%) | | 36.2%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-75 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|-----------|-----------------|-------------|------------|---------------|--|-------------------|-----------------|---------------|--|
| | | - 1 | | - | | | | CMH or Logistic | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 77 (43.5%) | 195 | 47 (24.1%) | 2.42 | 1.80 | 19.4% | 0.0001* | 0.0045* | |
| | | (36.1%, | | (18.3%, | (1.56, 3.78) | (1.34, 2.44) | (9.9%, 28.9%) | | | |
| | | 51.1%) | | 30.7%) | , , , | , , , | , | | | |
| Europe | 150 | 36 (24.0%) | 132 | 32 (24.2%) | 0.99 | 0.99 | -0.2% | 1.0000 | | |
| | | (17.4%, | | (17.2%, | (0.57, 1.70) | (0.65, 1.50) | (-10.3%, 9.8%) | | | |
| | | 31.6%) | | 32.5%) | (3.3.3) | (, | (,, | | | |
| Asia | 17 | 5 (29.4%) | 19 | 5 (26.3%) | 1.17 | 1.12 | 3.1% | 1.0000 | | |
| 1.014 | | (10.3%, | | (9.1%, | (0.27, 5.02) | (0.39, 3.20) | (-26.2%, 32.4%) | | | |
| | | 56.0%) | | 51.2%) | (0.27, 0.02) | (0.05, 5.20) | (2012/0, 521.170) | | | |
| Latin America | 18 | 15 (83.3%) | 19 | 9 (47.4%) | 5.56 | 1.76 | 36.0% | 0.0382* | | |
| | 10 | (58.6%, | | (24.4%, | (1.20, 25.71) | (1.05, 2.95) | (7.7%, 64.3%) | ****** | | |
| | | 96.4%) | | 71.1%) | (1.20, 23.71) | (1.03, 2.93) | (7.770, 01.370) | | | |
| Age Subgroup | | 70.470) | | 71.170) | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 230 | 80 (34.8%) | 247 | 64 (25.9%) | 1.53 | 1.34 | 8.9% | 0.0366* | 0.3334 | |
| vio years | 250 | (28.6%, | 217 | (20.6%, | (1.03, 2.26) | (1.02, 1.77) | (0.6%, 17.1%) | 0.0500 | 0.5551 | |
| | | 41.3%) | | 31.8%) | (1.03, 2.20) | (1.02, 1.77) | (0.070, 17.170) | | | |
| >=40 years | 132 | 53 (40.2%) | 118 | 29 (24.6%) | 2.06 | 1.63 | 15.6% | 0.0104* | | |
| >=40 years | 132 | | 110 | | | (1.12, 2.39) | (4.2%, 27.0%) | 0.0104** | | |
| | | (31.7%, | | (17.1%, | (1.19, 3.55) | (1.12, 2.39) | (4.2%, 27.0%) | | | |
| | | 49.0%) | | 33.4%) | | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-90 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | | |
|--------------------------------------|-----------|-------------------|------------|---------------|--------------|-----------------|---------------------------|-------------------------------|--------------|--|
| | | | | | | 1 | | CMH or Logistic P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Overall | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety | 362 | 60 (16.6%) | 365 | 24 (6.6%) | 2.82 | 2.52 | 10.0% | <0.0001+* | | |
| Population | | (12.9%, 20.8%) | | (4.3%, 9.6%) | (1.71, 4.64) | (1.61, 3.96) | (5.4%, 14.6%) | | | |
| Baseline Disease Severity Week 12 | | | | | | | | | | |
| Moderate baseline | 216 | 33 (15.3%) | 220 | 16 (7.3%) | 2.30 | 2.10 | 8.0% | 0.0096* | 0.3015 | |
| disease (IGA=3) | | (10.8%, 20.8%) | | (4.2%, 11.5%) | (1.23, 4.31) | (1.19, 3.70) | (2.1%, 13.9%) | | | |
| Severe baseline disease | 146 | 27 (18.5%) | 145 | 8 (5.5%) | 3.89 | 3.35 | 13.0% | 0.0009* | | |
| (IGA=4) | | (12.6%, 25.8%) | | (2.4%, 10.6%) | (1.70, 8.88) | (1.58, 7.13) | (5.7%, 20.3%) | | | |
| Gender Week 12 | | | | | | | | | | |
| Male | 193 | 33 (17.1%) | 204 | 9 (4.4%) | 4.47 | 3.88 | 12.7% | <0.0001* | 0.2318 | |
| | | (12.1%, 23.2%) | | (2.0%, 8.2%) | (2.08, 9.61) | (1.90, 7.89) | (6.7%, 18.7%) | | | |
| Female | 169 | 27 (16.0%) | 161 | 15 (9.3%) | 1.85 | 1.71 | 6.7% | 0.0976 | | |
| | | (10.8%, | | (5.3%, | (0.95, 3.62) | (0.95, 3.10) | (-0.5%, 13.8%) | | | |
| | | 22.4%) | | 14.9%) | | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-90 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|-----------|-----------------|------------|--------------|---------------|--|-----------------|-------------------|--------------|--|
| | | | | | | | - | CMH or Logistic F | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 32 (18.1%) | 195 | 8 (4.1%) | 5.16 | 4.41 | 14.0% | < 0.0001* | 0.0573 | |
| | | (12.7%, | | (1.8%, 7.9%) | (2.31, 11.53) | (2.09, 9.31) | (7.7%, 20.3%) | | | |
| | | 24.6%) | | | | | | | | |
| Europe | 150 | 19 (12.7%) | 132 | 8 (6.1%) | 2.25 | 2.09 | 6.6% | 0.0693 | | |
| - | | (7.8%, | | (2.7%, | (0.95, 5.32) | (0.95, 4.62) | (-0.1%, 13.3%) | | | |
| | | 19.1%) | | 11.6%) | | | | | | |
| Asia | 17 | 0 (0.0%) | 19 | 2 (10.5%) | 0.20 | 0.22 | -9.7% | 0.4873 | | |
| | | | | (1.3%, | (0.01, 4.47) | (0.01, 4.33) | (-26.1%, 6.6%) | | | |
| | | | | 33.1%) | | | | | | |
| Latin America | 18 | 9 (50.0%) | 19 | 6 (31.6%) | 2.17 | 1.58 | 18.4% | 0.3245 | | |
| | | (26.0%, | | (12.6%, | (0.57, 8.26) | (0.71, 3.55) | (-12.7%, 49.6%) | | | |
| | | 74.0%) | | 56.6%) | | | | | | |
| Age Subgroup | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 230 | 35 (15.2%) | 247 | 16 (6.5%) | 2.59 | 2.35 | 8.7% | 0.0027* | 0.4752 | |
| • | | (10.8%, | | (3.7%, | (1.39, 4.82) | (1.34, 4.13) | (3.2%, 14.3%) | | | |
| | | 20.5%) | | 10.3%) | | | | | | |
| >=40 years | 132 | 25 (18.9%) | 118 | 8 (6.8%) | 3.21 | 2.79 | 12.2% | 0.0049* | | |
| • | | (12.6%, | | (3.0%, | (1.39, 7.44) | (1.31, 5.95) | (4.1%, 20.2%) | | | |
| | | 26.7%) | | 12.9%) | , | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-100 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitir | nib 200mg QD Dupi | lumab 300 | mg Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | | |
|----------------------------------|-----------|-------------------|-----------|------------------|---------------|-----------------|---|-------------------------------|--------------|--|
| | | | | | | | CMH or Logistic 1 | CMH or Logistic P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Overall | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety | 362 | 21 (5.8%) | 365 | 6 (1.6%) | 3.68 | 3.52 | 4.1% | 0.0031+* | | |
| Population | | (3.6%, 8.7%) | | (0.6%, 3.5%) | (1.47, 9.24) | (1.44, 8.61) | (1.4%, 6.9%) | | | |
| Baseline Disease Severity | | | | , , , | | | , | | | |
| Week 12 | | | | | | | | | | |
| Moderate baseline | 216 | 10 (4.6%) | 220 | 3 (1.4%) | 3.51 | 3.40 | 3.3% | 0.0519 | 0.4451 | |
| disease (IGA=3) | | (2.2%, 8.3%) | | (0.3%, 3.9%) | (0.95, 12.94) | (0.95, 12.17) | (0.1%, 6.5%) | | | |
| Severe baseline disease | 146 | 11 (7.5%) | 145 | 3 (2.1%) | 3.86 | 3.64 | 5.5% | 0.0516 | | |
| (IGA=4) | | (3.8%, | | (0.4%, 5.9%) | (1.05, 14.13) | (1.04, 12.78) | (0.6%, 10.3%) | | | |
| , | | 13.1%) | | (,, | (, , | (12 , 11 = 7 | (, | | | |
| Gender | | / | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Male | 193 | 11 (5.7%) | 204 | 3 (1.5%) | 4.05 | 3.88 | 4.2% | 0.0284* | 0.9547 | |
| | | (2.9%, | | (0.3%, 4.2%) | (1.11, 14.74) | (1.10, 13.68) | (0.6%, 7.9%) | 0.0_0 | 0.50 | |
| | | 10.0%) | | (******, **=***) | (,) | (====, =====) | (************************************** | | | |
| Female | 169 | 10 (5.9%) | 161 | 3 (1.9%) | 3.31 | 3.18 | 4.1% | 0.0872 | | |
| | 10) | (2.9%, | 101 | (0.4%, 5.3%) | (0.89, 12.26) | (0.89, 11.33) | (-0.1%, 8.2%) | 3.0072 | | |
| | | 10.6%) | | (0.170, 5.570) | (0.0), 12.20) | (0.05, 11.55) | (0.170, 0.270) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-100 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitir | nib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|-------------------------|-----------|-------------------------------|------------|------------------------------|-----------------------|--|-------------------------|----------------------------|--------------------------|
| | | • | | - | | - | _ | CMH or Logistic I | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Region | | (55,0 C1) | | (55,70 CI) | (5070 C1) | (33/0 C1) | (5570 C1) | , and [1] | meraetion |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 177 | 12 (6.8%) (3.6%, 11.5%) | 195 | 2 (1.0%) (0.1%, 3.7%) | 7.02 (1.55, 31.81) | 6.61 (1.50, 29.13) | 5.8% (1.8%, 9.7%) | 0.0047* | 0.5209 |
| Europe | 150 | 8 (5.3%) (2.3%, 10.2%) | 132 | 3 (2.3%) (0.5%, 6.5%) | 2.42 (0.63, 9.33) | 2.35 (0.64, 8.66) | 3.1% (-1.3%, 7.5%) | 0.2280 | |
| Asia | 17 | 0 (0.0%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 0.35 (0.01, 9.24) | 0.37 (0.02, 8.53) | -4.7% (-18.5%, 9.1%) | 1.0000 | |
| Latin America | 18 | 1 (5.6%) (0.1%, 27.3%) | 19 | 0 (0.0%) | 3.34 (0.13, 87.52) | 3.16 (0.14, 72.84) | 5.4% (-8.5%, 19.3%) | 0.4865 | |
| Age Subgroup Week 12 | | , | | | | | | | |
| <40 years | 230 | 12 (5.2%) (2.7%, 8.9%) | 247 | 5 (2.0%) (0.7%, 4.7%) | 2.66 (0.92, 7.68) | 2.58 (0.92, 7.20) | 3.2% (-0.2%, 6.6%) | 0.0823 | 0.3360 |
| >=40 years | 132 | 9 (6.8%) (3.2%, 12.5%) | 118 | 1 (0.8%) (0.0%, 4.6%) | 8.56 (1.07, 68.62) | 8.05 (1.03, 62.56) | 6.0% (1.4%, 10.6%) | 0.0209* | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: EASI-75 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--|-----------|----------------------------------|-------------|----------------------------------|----------------------|--|-------------------------|----------------------------|--------------------------|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | (>0,0 01) | | (5070 01) | (5570 C1) | (5070-02) | (5570 62) | , alde [1] | moraemon | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 274 (75.7%) (70.9%, 80.0%) | 365 | 223 (61.1%) (55.9%, 66.1%) | 1.98 (1.44, 2.72) | 1.24 (1.12, 1.37) | 14.6% (7.9%, 21.3%) | <0.0001+* | | |
| Baseline Disease Severity Week 12 | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 160 (74.1%) (67.7%, 79.8%) | 220 | 137 (62.3%) (55.5%, 68.7%) | 1.73 (1.15, 2.60) | 1.19 (1.04, 1.35) | 11.8% (3.1%, 20.5%) | 0.0101* | 0.2971 | |
| Severe baseline disease (IGA=4) | 146 | 114 (78.1%) (70.5%, 84.5%) | 145 | 86 (59.3%) (50.8%, 67.4%) | 2.44 (1.46, 4.08) | 1.32 (1.12, 1.54) | 18.8% (8.3%, 29.2%) | 0.0006* | | |
| Gender Week 12 | | , | | 211117 | | | | | | |
| Male | 193 | 148 (76.7%) (70.1%, 82.5%) | 204 | 111 (54.4%) (47.3%, 61.4%) | 2.76 (1.79, 4.25) | 1.41 (1.22, 1.63) | 22.3% (13.2%, 31.3%) | <0.0001* | 0.0347* | |
| Female | 169 | 126 (74.6%) (67.3%, 80.9%) | 161 | 112 (69.6%) (61.8%, 76.6%) | 1.28 (0.79, 2.08) | 1.07 (0.94, 1.23) | 5.0% (-4.7%, 14.7%) | 0.3280 | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: EASI-75 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------|-----------|------------------|------------|-------------|---|--|---|-------------------|---------------|--|--|
| | | • | | - | | - | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Region | | | | | | | | | | | |
| Week 12 | | | | | | | | | | | |
| US/Canada/Australia | 177 | 137 (77.4%) | 195 | 116 (59.5%) | 2.33 | 1.30 | 17.9% | 0.0002* | 0.6301 | | |
| | | (70.5%, | | (52.2%, | (1.48, 3.67) | (1.13, 1.50) | (8.7%, 27.2%) | | | | |
| | | 83.3%) | | 66.4%) | | | | | | | |
| Europe | 150 | 110 (73.3%) | 132 | 78 (59.1%) | 1.90 | 1.24 | 14.2% | 0.0159* | | | |
| • | | (65.5%, | | (50.2%, | (1.15, 3.14) | (1.05, 1.47) | (3.3%, 25.2%) | | | | |
| | | 80.2%) | | 67.6%) | | | | | | | |
| Asia | 17 | 11 (64.7%) | 19 | 12 (63.2%) | 1.07 | 1.02 | 1.5% | 1.0000 | | | |
| | | (38.3%, | | (38.4%, | (0.27, 4.18) | (0.63, 1.67) | (-29.9%, 33.0%) | | | | |
| | | 85.8%) | | 83.7%) | , , , | | , | | | | |
| Latin America | 18 | 16 (88.9%) | 19 | 17 (89.5%) | 0.94 | 0.99 | -0.6% | 1.0000 | | | |
| | | (65.3%, | | (66.9%, | (0.12, 7.50) | (0.79, 1.24) | (-20.6%, 19.4%) | | | | |
| | | 98.6%) | | 98.7%) | | | | | | | |
| Age Subgroup | | , | | , | | | | | | | |
| Week 12 | | | | | | | | | | | |
| <40 years | 230 | 171 (74.3%) | 247 | 157 (63.6%) | 1.66 | 1.17 | 10.8% | 0.0134* | 0.1457 | | |
| • | | (68.2%, | | (57.2%, | (1.12, 2.46) | (1.04, 1.32) | (2.5%, 19.0%) | | | | |
| | | 79.9%) | | 69.6%) | | | | | | | |
| >=40 years | 132 | 103 (78.0%) | 118 | 66 (55.9%) | 2.80 | 1.40 | 22.1% | 0.0002* | | | |
| • | | (70.0%, | | (46.5%, | (1.62, 4.85) | (1.16, 1.68) | (10.7%, 33.5%) | | | | |
| | | 84.8%) | | 65.1%) | , | , , , , , , , | , ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EASI-90 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--|-----------|----------------------------------|-------------|----------------------------------|----------------------|--|------------------------|----------------------------|--------------------------|--|--|
| | | | | | | CMH or | | | | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Overall | | (5570 C1) | | (5070 01) | (5570 C1) | (5070 02) | (50,0 01) | , and [1] | III CIUCUSII | | |
| Week 12 | | | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 171 (47.2%) (42.0%, 52.5%) | 365 | 122 (33.4%) (28.6%, 38.5%) | 1.79 (1.32, 2.41) | 1.41 (1.18, 1.69) | 13.8% (6.8%, 20.9%) | 0.0001+* | | | |
| Baseline Disease Severity Week 12 | | , | | , | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 104 (48.1%) (41.3%, 55.0%) | 220 | 78 (35.5%) (29.1%, 42.2%) | 1.69 (1.15, 2.48) | 1.36 (1.08, 1.70) | 12.7% (3.5%, 21.9%) | 0.0087* | 0.7880 | | |
| Severe baseline disease (IGA=4) | 146 | 67 (45.9%) (37.6%, 54.3%) | 145 | 44 (30.3%) (23.0%, 38.5%) | 1.95 (1.20, 3.15) | 1.51 (1.12, 2.05) | 15.5% (4.5%, 26.6%) | 0.0079* | | | |
| Gender Week 12 | | 2// | | 201270) | | | | | | | |
| Male | 193 | 89 (46.1%) (38.9%, 53.4%) | 204 | 55 (27.0%) (21.0%, 33.6%) | 2.32 (1.52, 3.53) | 1.71 (1.30, 2.25) | 19.2% (9.9%, 28.5%) | 0.0001* | 0.1618 | | |
| Female | 169 | 82 (48.5%) (40.8%, 56.3%) | 161 | 67 (41.6%) (33.9%, 49.6%) | 1.32 (0.86, 2.04) | 1.17 (0.92, 1.48) | 6.9% (-3.8%, 17.6%) | 0.2246 | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: EASI-90 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------|-----------|------------------|-------------|------------|-----------------|--|-------------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Region | | | | | | | | | | | |
| Week 12 | | | | | | | | | | | |
| US/Canada/Australia | 177 | 93 (52.5%) | 195 | 62 (31.8%) | 2.38 | 1.65 | 20.7% | < 0.0001* | 0.1067 | | |
| | | (44.9%, | | (25.3%, | (1.56, 3.62) | (1.29, 2.12) | (10.9%, 30.6%) | | | | |
| | | 60.1%) | | 38.8%) | | , , , | , | | | | |
| Europe | 150 | 58 (38.7%) | 132 | 43 (32.6%) | 1.30 | 1.19 | 6.1% | 0.3202 | | | |
| 1 | | (30.8%, | | (24.7%, | (0.80, 2.13) | (0.86, 1.63) | (-5.1%, 17.3%) | | | | |
| | | 47.0%) | | 41.3%) | , , , | | , | | | | |
| Asia | 17 | 5 (29.4%) | 19 | 5 (26.3%) | 1.17 | 1.12 | 3.1% | 1.0000 | | | |
| | | (10.3%, | | (9.1%, | (0.27, 5.02) | (0.39, 3.20) | (-26.2%, 32.4%) | | | | |
| | | 56.0%) | | 51.2%) | , , , | | , , , | | | | |
| Latin America | 18 | 15 (83.3%) | 19 | 12 (63.2%) | 2.92 | 1.32 | 20.2% | 0.2691 | | | |
| | | (58.6%, | | (38.4%, | (0.62, 13.76) | (0.88, 1.97) | (-7.5%, 47.9%) | | | | |
| | | 96.4%) | | 83.7%) | , | | , | | | | |
| Age Subgroup | | , | | , | | | | | | | |
| Week 12 | | | | | | | | | | | |
| <40 years | 230 | 102 (44.3%) | 247 | 81 (32.8%) | 1.63 | 1.35 | 11.6% | 0.0110* | 0.3564 | | |
| , | | (37.8%, | | (27.0%, | (1.13, 2.37) | (1.07, 1.70) | (2.9%, 20.2%) | | | | |
| | | 51.0%) | | 39.0%) | | | | | | | |
| >=40 years | 132 | 69 (52.3%) | 118 | 41 (34.7%) | 2.06 | 1.50 | 17.5% | 0.0072* | | | |
| | | (43.4%, | | (26.2%, | (1.24, 3.43) | (1.12, 2.02) | (5.4%, 29.6%) | | | | |
| | | 61.0%) | | 44.1%) | , , , , , , , , | , , , _ , _ , | (= 1.7, = 2.0070) | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EASI-100 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--|-----------|---------------------------------|------------|---|-----------------------|--|------------------------|----------------------------|--------------------------|--|--|
| | | | | _ | | | _ | CMH or Logistic I | P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Overall | | , , | | , | ` | , , | , | | | | |
| Week 12 | | | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 56 (15.5%) (11.9%, 19.6%) | 365 | 20 (5.5%) (3.4%, 8.3%) | 3.16 (1.85, 5.39) | 2.83 (1.73, 4.61) | 10.0% (5.6%, 14.4%) | <0.0001+* | | | |
| Baseline Disease Severity Week 12 | | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 34 (15.7%) (11.2%, 21.3%) | 220 | 13 (5.9%) (3.2%, 9.9%) | 2.97 (1.52, 5.81) | 2.66 (1.45, 4.91) | 9.8% (4.1%, 15.6%) | 0.0011* | 0.9467 | | |
| Severe baseline disease (IGA=4) | 146 | 22 (15.1%) (9.7%, 21.9%) | 145 | 7 (4.8%) (2.0%, 9.7%) | 3.50 (1.44, 8.47) | 3.12 (1.38, 7.08) | 10.2% (3.5%, 17.0%) | 0.0054* | | | |
| Gender Week 12 | | , | | | | | | | | | |
| Male | 193 | 30 (15.5%) (10.7%, 21.4%) | 204 | 8 (3.9%) (1.7%, 7.6%) | 4.51 (2.01, 10.11) | 3.96 (1.86, 8.43) | 11.6% (5.9%, 17.4%) | 0.0001* | 0.4518 | | |
| Female | 169 | 26 (15.4%) (10.3%, 21.7%) | 161 | 12 (7.5%) (3.9%, 12.7%) | 2.26 (1.10, 4.64) | 2.06 (1.08, 3.95) | 7.9% (1.1%, 14.7%) | 0.0257* | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: EASI-100 response at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitini | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|------------|-----------------|------------|--------------|---------------|--|---|-------------------|---------------------|--|
| | | | | | | _ | _ | CMH or Logistic F | istic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 31 (17.5%) | 195 | 8 (4.1%) | 4.96 | 4.27 | 13.4% | < 0.0001* | 0.1169 | |
| | | (12.2%, | | (1.8%, 7.9%) | (2.21, 11.12) | (2.02, 9.04) | (7.2%, 19.7%) | | | |
| | | 23.9%) | | | | | | | | |
| Europe | 150 | 17 (11.3%) | 132 | 7 (5.3%) | 2.28 | 2.14 | 6.0% | 0.0874 | | |
| • | | (6.7%, | | (2.2%, | (0.92, 5.69) | (0.91, 4.99) | (-0.3%, 12.4%) | | | |
| | | 17.5%) | | 10.6%) | | | | | | |
| Asia | 17 | 1 (5.9%) | 19 | 2 (10.5%) | 0.53 | 0.56 | -4.6% | 1.0000 | | |
| | | (0.1%, | | (1.3%, | (0.04, 6.44) | (0.06, 5.63) | (-22.4%, 13.1%) | | | |
| | | 28.7%) | | 33.1%) | | . , , | , , , | | | |
| Latin America | 18 | 7 (38.9%) | 19 | 3 (15.8%) | 3.39 | 2.46 | 23.1% | 0.1510 | | |
| | | (17.3%, | | (3.4%, | (0.72, 16.07) | (0.75, 8.09) | (-4.8%, 51.0%) | | | |
| | | 64.3%) | | 39.6%) | | , , , | , | | | |
| Age Subgroup | | , | | , | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 230 | 34 (14.8%) | 247 | 13 (5.3%) | 3.12 | 2.81 | 9.5% | 0.0006* | 0.7809 | |
| • | | (10.5%, | | (2.8%, 8.8%) | (1.60, 6.08) | (1.52, 5.19) | (4.2%, 14.9%) | | | |
| | | 20.0%) | | , , , | | , , | | | | |
| >=40 years | 132 | 22 (16.7%) | 118 | 7 (5.9%) | 3.17 | 2.81 | 10.7% | 0.0096* | | |
| - , | | (10.7%, | | (2.4%, | (1.30, 7.73) | (1.25, 6.34) | (3.1%, 18.4%) | | | |
| | | 24.1%) | | 11.8%) | (,) | (,, | (====================================== | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-2 in POEM total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------------|-----------|-----------------|-------------|------------|--------------|--|---------------|---------------|--------------|--|
| | | | | | | CMH or Logistic F | | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Overall | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety | 358 | 108 (30.2%) | 363 | 61 (16.8%) | 2.14 | 1.79 | 13.4% | <0.0001+* | | |
| Population | | (25.5%, | | (13.1%, | (1.50, 3.05) | (1.36, 2.37) | (7.2%, 19.5%) | | | |
| • | | 35.2%) | | 21.1%) | | , , , | , | | | |
| Baseline Disease Severity | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Moderate baseline | 213 | 65 (30.5%) | 219 | 34 (15.5%) | 2.39 | 1.97 | 15.0% | 0.0002* | 0.5525 | |
| disease (IGA=3) | | (24.4%, | | (11.0%, | (1.50, 3.81) | (1.36, 2.84) | (7.2%, 22.8%) | | | |
| | | 37.2%) | | 21.0%) | | | | | | |
| Severe baseline disease | 145 | 43 (29.7%) | 144 | 27 (18.8%) | 1.83 | 1.58 | 10.9% | 0.0391* | | |
| (IGA=4) | | (22.4%, | | (12.7%, | (1.05, 3.17) | (1.04, 2.41) | (1.1%, 20.7%) | | | |
| | | 37.8%) | | 26.1%) | | | | | | |
| Gender | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Male | 190 | 56 (29.5%) | 202 | 28 (13.9%) | 2.60 | 2.13 | 15.6% | 0.0002* | 0.4891 | |
| | | (23.1%, | | (9.4%, | (1.56, 4.31) | (1.41, 3.20) | (7.6%, 23.7%) | | | |
| | | 36.5%) | | 19.4%) | | | | | | |
| Female | 168 | 52 (31.0%) | 161 | 33 (20.5%) | 1.74 | 1.51 | 10.5% | 0.0328* | | |
| | | (24.1%, | | (14.5%, | (1.05, 2.88) | (1.03, 2.21) | (1.1%, 19.8%) | | | |
| | | 38.5%) | | 27.6%) | | . , , | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-2 in POEM total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|-----------|-----------------|-------------|------------|---------------|--|---|-------------------|-----------------------------|--|
| | | | | _ | | _ | _ | CMH or Logistic F | H or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 51 (28.8%) | 193 | 29 (15.0%) | 2.29 | 1.92 | 13.8% | 0.0015* | 0.8576 | |
| | | (22.3%, | | (10.3%, | (1.37, 3.82) | (1.28, 2.88) | (5.4%, 22.2%) | | | |
| | | 36.1%) | | 20.9%) | | | | | | |
| Europe | 150 | 43 (28.7%) | 132 | 21 (15.9%) | 2.12 | 1.80 | 12.8% | 0.0150* | | |
| 1 | | (21.6%, | | (10.1%, | (1.18, 3.81) | (1.13, 2.87) | (3.2%, 22.3%) | | | |
| | | 36.6%) | | 23.3%) | | | | | | |
| Asia | 14 | 7 (50.0%) | 19 | 8 (42.1%) | 1.38 | 1.19 | 7.9% | 0.7325 | | |
| | | (23.0%, | | (20.3%, | (0.34, 5.51) | (0.56, 2.50) | (-26.4%, 42.2%) | | | |
| | | 77.0%) | | 66.5%) | (332 , 232 , | (112.1) | (,, | | | |
| Latin America | 17 | 7 (41.2%) | 19 | 3 (15.8%) | 3.73 | 2.61 | 25.4% | 0.1394 | | |
| | | (18.4%, | | (3.4%, | (0.78, 17.88) | (0.80, 8.52) | (-3.2%, 54.0%) | | | |
| | | 67.1%) | | 39.6%) | (3.1.2) | (, , , , , , , , , , , , , , , , , , , | (= = = = = = = = = = = = = = = = = = = | | | |
| Age Subgroup | | , | | , | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 226 | 63 (27.9%) | 246 | 42 (17.1%) | 1.88 | 1.63 | 10.8% | 0.0056* | 0.2639 | |
| , | | (22.1%, | | (12.6%, | (1.21, 2.92) | (1.16, 2.31) | (3.3%, 18.3%) | | | |
| | | 34.2%) | | 22.4%) | , , , , | () -, , | (====, | | | |
| >=40 years | 132 | 45 (34.1%) | 117 | 19 (16.2%) | 2.67 | 2.10 | 17.9% | 0.0014* | | |
| J | 102 | (26.1%, | | (10.1%, | (1.45, 4.91) | (1.31, 3.38) | (7.4%, 28.3%) | 3.001. | | |
| | | 42.8%) | | 24.2%) | (1, 1.51) | (1.51, 5.50) | (,,3, 20.370) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0 in POEM total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--|-----------|---------------------------------|------------|---------------------------|-----------------------|--|------------------------|----------------------------|--------------------------|--|
| | | | | - | | | _ | CMH or Logistic I | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | (2570 C1) | | (2370 CI) | (95% CI) | (2370 CI) | (9370 CI) | value [1] | meraction | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety Population | 359 | 49 (13.6%) (10.3%, 17.6%) | 365 | 11 (3.0%) (1.5%, 5.3%) | 5.13 (2.62, 10.05) | 4.54 (2.40, 8.59) | 10.7% (6.7%, 14.6%) | <0.0001+* | | |
| Baseline Disease Severity Week 12 | | , | | | | | | | | |
| Moderate baseline disease (IGA=3) | 214 | 33 (15.4%) (10.9%, 21.0%) | 220 | 8 (3.6%) (1.6%, 7.0%) | 4.83 (2.18, 10.73) | 4.24 (2.00, 8.97) | 11.8% (6.3%, 17.2%) | <0.0001* | 0.4476 | |
| Severe baseline disease (IGA=4) | 145 | 16 (11.0%) (6.4%, 17.3%) | 145 | 3 (2.1%) (0.4%, 5.9%) | 5.87 (1.67, 20.61) | 5.33 (1.59, 17.91) | 9.0% (3.4%, 14.6%) | 0.0033* | | |
| Gender Week 12 | | , | | | | | | | | |
| Male | 191 | 23 (12.0%) (7.8%, 17.5%) | 204 | 3 (1.5%) (0.3%, 4.2%) | 9.17 (2.71, 31.08) | 8.19 (2.50, 26.83) | 10.6% (5.7%, 15.5%) | <0.0001* | 0.9376 | |
| Female | 168 | 26 (15.5%) (10.4%, 21.8%) | 161 | 8 (5.0%) (2.2%, 9.6%) | 3.50 (1.54, 7.99) | 3.11 (1.45, 6.68) | 10.5% (4.1%, 16.9%) | 0.0019* | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0 in POEM total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|-----------|-----------------|------------|-----------------------------------|----------------|--|-----------------------|-------------------|---------------|--|
| | | | | | | | • | CMH or Logistic F | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 21 (11.9%) | 195 | 3 (1.5%) | 8.62 | 7.71 | 10.3% | < 0.0001* | 0.8118 | |
| | | (7.5%, | | (0.3%, 4.4%) | (2.52, 29.42) | (2.34, 25.41) | (5.3%, 15.4%) | | | |
| | | 17.6%) | | , , , | , , , | , , , | ` ' ' | | | |
| Europe | 150 | 23 (15.3%) | 132 | 7 (5.3%) | 3.23 | 2.89 | 10.0% | 0.0067* | | |
| • | | (10.0%, | | (2.2%, | (1.34, 7.81) | (1.28, 6.52) | (3.1%, 16.9%) | | | |
| | | 22.1%) | | 10.6%) | , , , | , , , | , | | | |
| Asia | 15 | 3 (20.0%) | 19 | 0 (0.0%) | 10.92 | 8.75 | 19.4% | 0.0760 | | |
| | | (4.3%, | | , , | (0.52, 229.91) | (0.49, 157.34) | (-2.0%, 40.8%) | | | |
| | | 48.1%) | | | | , | , | | | |
| Latin America | 17 | 2 (11.8%) | 19 | 1 (5.3%) | 2.40 | 2.24 | 6.5% | 0.5929 | | |
| | | (1.5%, | | (0.1%, | (0.20, 29.13) | (0.22, 22.51) | (-11.8%, 24.8%) | | | |
| | | 36.4%) | | 26.0%) | , | , , , | , , , | | | |
| Age Subgroup | | , | | , | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 227 | 28 (12.3%) | 247 | 6 (2.4%) | 5.65 | 5.08 | 9.9% | < 0.0001* | 0.6465 | |
| • | | (8.4%, | | (0.9%, 5.2%) | (2.29, 13.92) | (2.14, 12.04) | (5.2%, 14.6%) | | | |
| | | 17.3%) | | , , , , | , | , , , | | | | |
| >=40 years | 132 | 21 (15.9%) | 118 | 5 (4.2%) | 4.28 | 3.75 | 11.7% | 0.0031* | | |
| | | (10.1%, | | (1.4%, 9.6%) | (1.56, 11.74) | (1.46, 9.64) | (4.5%, 18.9%) | | | |
| | | 23.3%) | | (, , , , , , , , , , , , , , , , | (,, | 3,2101) | (= = = , = = = , = , | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in CDLQI total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------------------------|-----------|------------------------|-------------|------------------------|----------------|--|----------------|----------------------------|--------------------------|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | , | | , | , | · / | , | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety | 358 | 126 (35.2%) | 361 | 110 (30.5%) | 1.24 | 1.16 | 4.7% | 0.1758+ | | |
| Population | | (30.2%, 40.4%) | | (25.8%, 35.5%) | (0.91, 1.69) | (0.94, 1.43) | (-2.1%, 11.6%) | | | |
| Baseline Disease Severity Week 12 | | , | | , | | | | | | |
| Moderate baseline | 212 | 77 (36.3%) | 217 | 68 (31.3%) | 1.25 | 1.16 | 5.0% | 0.3076 | 0.9146 | |
| disease (IGA=3) | | (29.8%, 43.2%) | | (25.2%, 38.0%) | (0.84, 1.87) | (0.89, 1.51) | (-4.0%, 13.9%) | | | |
| Severe baseline disease | 146 | 49 (33.6%) | 144 | 42 (29.2%) | 1.23 | 1.15 | 4.4% | 0.4492 | | |
| (IGA=4) | | (26.0%, 41.8%) | | (21.9%, 37.3%) | (0.75, 2.02) | (0.82, 1.62) | (-6.3%, 15.1%) | | | |
| Gender | | , | | , | | | | | | |
| Week 12 | | | | | | | | | | |
| Male | 191 | 73 (38.2%) | 201 | 61 (30.3%) | 1.42 | 1.26 | 7.9% | 0.1106 | 0.3205 | |
| | | (31.3%, | | (24.1%, | (0.93, 2.16) | (0.96, 1.66) | (-1.5%, 17.2%) | | | |
| | | 45.5%) | | 37.2%) | | | | | | |
| Female | 167 | 53 (31.7%) | 160 | 49 (30.6%) | 1.05 | 1.04 | 1.1% | 0.9050 | | |
| | | (24.8%, | | (23.6%, | (0.66, 1.68) | (0.75, 1.43) | (-8.9%, 11.2%) | | | |
| | | 39.4%) | | 38.4%) | | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in CDLQI total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|-----------|-----------------|-------------|------------|--------------|--|------------------|-------------------|---------------|--|
| | | | | | | _ | _ | CMH or Logistic F | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 174 | 69 (39.7%) | 191 | 66 (34.6%) | 1.24 | 1.15 | 5.1% | 0.3301 | 0.7702 | |
| | | (32.3%, | | (27.8%, | (0.81, 1.90) | (0.88, 1.50) | (-4.8%, 15.0%) | | | |
| | | 47.3%) | | 41.8%) | | | , | | | |
| Europe | 150 | 46 (30.7%) | 132 | 35 (26.5%) | 1.23 | 1.16 | 4.2% | 0.5099 | | |
| 1 | | (23.4%, | | (19.2%, | (0.73, 2.06) | (0.80, 1.68) | (-6.4%, 14.7%) | | | |
| | | 38.7%) | | 34.9%) | | . , , | , , , | | | |
| Asia | 16 | 3 (18.8%) | 19 | 4 (21.1%) | 0.87 | 0.89 | -2.3% | 1.0000 | | |
| | | (4.0%, | | (6.1%, | (0.16, 4.60) | (0.23, 3.41) | (-28.8%, 24.2%) | | | |
| | | 45.6%) | | 45.6%) | (** **, ***, | (| (,, | | | |
| Latin America | 18 | 8 (44.4%) | 19 | 5 (26.3%) | 2.24 | 1.69 | 18.1% | 0.3133 | | |
| | | (21.5%, | | (9.1%, | (0.56, 8.91) | (0.68, 4.21) | (-12.2%, 48.4%) | | | |
| | | 69.2%) | | 51.2%) | (0.00, 0.00) | (3.33, 1.22) | (====,=, ====,, | | | |
| Age Subgroup | | 0,12,0) | | 21.270) | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 227 | 67 (29.5%) | 243 | 67 (27.6%) | 1.10 | 1.07 | 1.9% | 0.6830 | 0.3267 | |
| | | (23.7%, | | (22.1%, | (0.74, 1.64) | (0.80, 1.43) | (-6.2%, 10.1%) | | | |
| | | 35.9%) | | 33.6%) | () | (, | (11 17, | | | |
| >=40 years | 131 | 59 (45.0%) | 118 | 43 (36.4%) | 1.43 | 1.24 | 8.6% | 0.1972 | | |
| - 10 30415 | 131 | (36.3%, | 110 | (27.8%, | (0.86, 2.38) | (0.91, 1.68) | (-3.6%, 20.8%) | 0.1772 | | |
| | | 54.0%) | | 45.8%) | (0.00, 2.30) | (0.51, 1.00) | (3.070, 20.070) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: PP-NRS total score 4-point improvement at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------------|-----------|------------------|-------------|-------------|--------------|--|----------------|-------------------|---------------|--|
| | | | | | | | - | CMH or Logistic 1 | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Overall | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety | 357 | 235 (65.8%) | 364 | 224 (61.5%) | 1.20 | 1.07 | 4.3% | 0.2346+ | | |
| Population | | (60.7%, | | (56.3%, | (0.89, 1.63) | (0.96, 1.19) | (-2.7%, 11.3%) | | | |
| | | 70.7%) | | 66.6%) | | | | | | |
| Baseline Disease Severity | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Moderate baseline | 213 | 132 (62.0%) | 219 | 131 (59.8%) | 1.09 | 1.04 | 2.2% | 0.6935 | 0.4025 | |
| disease (IGA=3) | | (55.1%, | | (53.0%, | (0.74, 1.61) | (0.89, 1.21) | (-7.0%, 11.4%) | | | |
| | | 68.5%) | | 66.4%) | | | | | | |
| Severe baseline disease | 144 | 103 (71.5%) | 145 | 93 (64.1%) | 1.40 | 1.12 | 7.4% | 0.2082 | | |
| (IGA=4) | | (63.4%, | | (55.8%, | (0.86, 2.31) | (0.95, 1.31) | (-3.3%, 18.1%) | | | |
| | | 78.7%) | | 71.9%) | | | | | | |
| Gender | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| Male | 191 | 124 (64.9%) | 204 | 120 (58.8%) | 1.30 | 1.10 | 6.1% | 0.2161 | 0.5999 | |
| | | (57.7%, | | (51.7%, | (0.86, 1.95) | (0.95, 1.29) | (-3.5%, 15.7%) | | | |
| | | 71.7%) | | 65.6%) | | | | | | |
| Female | 166 | 111 (66.9%) | 160 | 104 (65.0%) | 1.09 | 1.03 | 1.9% | 0.7275 | | |
| | | (59.2%, | | (57.1%, | (0.69, 1.72) | (0.88, 1.20) | (-8.4%, 12.2%) | | | |
| | | 74.0%) | | 72.4%) | , | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline PP-NRS score <4.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: PP-NRS total score 4-point improvement at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300ı | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-------------------------|-----------|------------------|-------------|-------------|---------------|--|----------------|-------------------------------|--------------|--|
| | | | | | | | | CMH or Logistic P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 174 | 116 (66.7%) | 195 | 118 (60.5%) | 1.31 | 1.10 | 6.2% | 0.2350 | 0.1069 | |
| | | (59.1%, | | (53.3%, | (0.85, 2.00) | (0.94, 1.29) | (-3.7%, 16.0%) | | | |
| | | 73.6%) | | 67.4%) | | | | | | |
| Europe | 148 | 96 (64.9%) | 131 | 81 (61.8%) | 1.14 | 1.05 | 3.0% | 0.6201 | | |
| • | | (56.6%, | | (52.9%, | (0.70, 1.86) | (0.88, 1.26) | (-8.3%, 14.4%) | | | |
| | | 72.5%) | | 70.2%) | | | | | | |
| Asia | 17 | 8 (47.1%) | 19 | 14 (73.7%) | 0.32 | 0.64 | -26.6% | 0.1711 | | |
| | | (23.0%, | | (48.8%, | (0.08, 1.28) | (0.36, 1.13) | (-57.5%, 4.3%) | | | |
| | | 72.2%) | | 90.9%) | | | | | | |
| Latin America | 18 | 15 (83.3%) | 19 | 11 (57.9%) | 3.64 | 1.44 | 25.4% | 0.1510 | | |
| | | (58.6%, | | (33.5%, | (0.78, 16.93) | (0.93, 2.23) | (-2.7%, 53.5%) | | | |
| | | 96.4%) | | 79.7%) | | | | | | |
| Age Subgroup Week 12 | | | | | | | | | | |
| <40 years | 227 | 135 (59.5%) | 246 | 148 (60.2%) | 0.97 | 0.99 | -0.7% | 0.9253 | 0.0495* | |
| • | | (52.8%, | | (53.7%, | (0.67, 1.40) | (0.85, 1.15) | (-9.5%, 8.2%) | | | |
| | | 65.9%) | | 66.3%) | | | | | | |
| >=40 years | 130 | 100 (76.9%) | 118 | 76 (64.4%) | 1.84 | 1.19 | 12.5% | 0.0358* | | |
| • | | (68.7%, | | (55.1%, | (1.06, 3.21) | (1.01, 1.41) | (1.2%, 23.8%) | | | |
| | | 83.9%) | | 73.0%) | , , , | | , , , | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline PP-NRS score <4.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--|-----------|----------------------------------|-------------|---------------------------------|----------------------|--|------------------------|-------------------------------|--------------------------|--|--|
| | | | | | | | _ | CMH or Logistic P-val [2] for | | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Overall | | , | | , , | ` | • | | | | | |
| Week 12 | | | | | | | | | | | |
| Full Analysis Set Safety Population | 362 | 135 (37.3%) (32.3%, 42.5%) | 365 | 91 (24.9%) (20.6%, 29.7%) | 1.79 (1.30, 2.46) | 1.50 (1.20, 1.87) | 12.4% (5.7%, 19.0%) | 0.0003+* | | | |
| Baseline Disease Severity Week 12 | | , | | , | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 80 (37.0%) (30.6%, 43.9%) | 220 | 59 (26.8%) (21.1%, 33.2%) | 1.61 (1.07, 2.41) | 1.38 (1.04, 1.83) | 10.2% (1.5%, 18.9%) | 0.0240* | 0.4747 | | |
| Severe baseline disease (IGA=4) | 146 | 55 (37.7%) (29.8%, 46.1%) | 145 | 32 (22.1%) (15.6%, 29.7%) | 2.13 (1.27, 3.58) | 1.71 (1.18, 2.47) | 15.6% (5.2%, 26.0%) | 0.0047* | | | |
| Gender Week 12 | | , | | _,,,, | | | | | | | |
| Male | 193 | 71 (36.8%) (30.0%, 44.0%) | 204 | 46 (22.5%) (17.0%, 28.9%) | 2.00 (1.29, 3.10) | 1.63 (1.19, 2.23) | 14.2% (5.3%, 23.1%) | 0.0021* | 0.5906 | | |
| Female | 169 | 64 (37.9%) (30.5%, 45.6%) | 161 | 45 (28.0%) (21.2%, 35.6%) | 1.57 (0.99, 2.50) | 1.35 (0.99, 1.86) | 9.9% (-0.2%, 20.0%) | 0.0615 | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| - | Abrocitini | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---------------------|------------|-----------------|-------------|------------|---------------|--|----------------|-------------------|---------------|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 177 | 69 (39.0%) | 195 | 50 (25.6%) | 1.85 | 1.52 | 13.3% | 0.0075* | 0.8815 | |
| | | (31.8%, | | (19.7%, | (1.19, 2.88) | (1.12, 2.06) | (3.9%, 22.8%) | | | |
| | | 46.6%) | | 32.4%) | | | | | | |
| Europe | 150 | 53 (35.3%) | 132 | 34 (25.8%) | 1.57 | 1.37 | 9.6% | 0.0936 | | |
| - | | (27.7%, | | (18.5%, | (0.94, 2.63) | (0.96, 1.97) | (-1.1%, 20.3%) | | | |
| | | 43.5%) | | 34.1%) | | | | | | |
| Asia | 17 | 7 (41.2%) | 19 | 4 (21.1%) | 2.63 | 1.96 | 20.1% | 0.2814 | | |
| | | (18.4%, | | (6.1%, | (0.61, 11.37) | (0.69, 5.53) | (-9.6%, 49.8%) | | | |
| | | 67.1%) | | 45.6%) | | | | | | |
| Latin America | 18 | 6 (33.3%) | 19 | 3 (15.8%) | 2.67 | 2.11 | 17.5% | 0.2691 | | |
| | | (13.3%, | | (3.4%, | (0.55, 12.88) | (0.62, 7.20) | (-9.7%, 44.8%) | | | |
| | | 59.0%) | | 39.6%) | | | | | | |
| Age Subgroup | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 230 | 73 (31.7%) | 247 | 59 (23.9%) | 1.48 | 1.33 | 7.9% | 0.0652 | 0.0658 | |
| • | | (25.8%, | | (18.7%, | (0.99, 2.22) | (0.99, 1.78) | (-0.2%, 15.9%) | | | |
| | | 38.2%) | | 29.7%) | | | | | | |
| >=40 years | 132 | 62 (47.0%) | 118 | 32 (27.1%) | 2.38 | 1.73 | 19.9% | 0.0016* | | |
| • | | (38.2%, | | (19.3%, | (1.40, 4.05) | (1.22, 2.45) | (8.2%, 31.5%) | | | |
| | | 55.8%) | | 36.1%) | , | | , , , | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--|-----------|---|-------------|---|----------------------|--|-------------------------|----------------------------|--------------------------|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | (************************************** | | (************************************** | (*******/ | (************************************** | (| | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety Population | 302 | 120 (39.7%) (34.2%, 45.5%) | 309 | 123 (39.8%) (34.3%, 45.5%) | 0.99 (0.71, 1.38) | 0.99 (0.82, 1.21) | -0.2% (-7.9%, 7.5%) | 0.9591+ | | |
| Baseline Disease Severity Week 12 | | , | | , | | | | | | |
| Moderate baseline disease (IGA=3) | 177 | 57 (32.2%) (25.4%, 39.6%) | 184 | 66 (35.9%) (28.9%, 43.3%) | 0.85 (0.55, 1.31) | 0.90 (0.67, 1.20) | -3.7% (-13.4%, 6.1%) | 0.5056 | 0.3014 | |
| Severe baseline disease (IGA=4) | 125 | 63 (50.4%) (41.3%, 59.5%) | 125 | 57 (45.6%) (36.7%, 54.7%) | 1.21 (0.74, 1.99) | 1.11 (0.85, 1.43) | 4.8% (-7.6%, 17.2%) | 0.5268 | | |
| Gender Week 12 | | , | | , | | | | | | |
| Male | 161 | 59 (36.6%) (29.2%, 44.6%) | 172 | 59 (34.3%) (27.2%, 41.9%) | 1.11 (0.71, 1.74) | 1.07 (0.80, 1.43) | 2.3% (-7.9%, 12.6%) | 0.7310 | 0.4648 | |
| Female | 141 | 61 (43.3%) (35.0%, 51.9%) | 137 | 64 (46.7%) (38.1%, 55.4%) | 0.87 (0.54, 1.40) | 0.93 (0.71, 1.20) | -3.5% (-15.1%, 8.2%) | 0.6298 | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 12 - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|---|-----------|-----------------|-------------|------------|---|--|---|-------------------|---------------|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Region | | | | | | | | | | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 138 | 41 (29.7%) | 155 | 53 (34.2%) | 0.81 | 0.87 | -4.5% | 0.4528 | 0.2849 | |
| | | (22.2%, | | (26.8%, | (0.50, 1.33) | (0.62, 1.22) | (-15.2%, 6.2%) | | | |
| | | 38.1%) | | 42.2%) | , , , | ` ' ' | , | | | |
| Europe | 131 | 65 (49.6%) | 119 | 54 (45.4%) | 1.19 | 1.09 | 4.2% | 0.5279 | | |
| 1 | | (40.8%, | | (36.2%, | (0.72, 1.95) | (0.84, 1.42) | (-8.1%, 16.6%) | | | |
| | | 58.5%) | | 54.8%) | , , , | ` ' ' | , | | | |
| Asia | 16 | 3 (18.8%) | 19 | 8 (42.1%) | 0.32 | 0.45 | -23.4% | 0.1667 | | |
| | | (4.0%, | | (20.3%, | (0.07, 1.50) | (0.14, 1.40) | (-52.7%, 5.9%) | | | |
| | | 45.6%) | | 66.5%) | (0101, 1100) | (******) | (====================================== | | | |
| Latin America | 17 | 11 (64.7%) | 16 | 8 (50.0%) | 1.83 | 1.29 | 14.7% | 0.4905 | | |
| | | (38.3%, | | (24.7%, | (0.45, 7.41) | (0.71, 2.36) | (-18.7%, 48.1%) | | | |
| | | 85.8%) | | 75.3%) | (0112, 1112) | (*** = , = = = , | (,, | | | |
| Age Subgroup | | 00.070) | | , | | | | | | |
| Week 12 | | | | | | | | | | |
| <40 years | 198 | 79 (39.9%) | 209 | 88 (42.1%) | 0.91 | 0.95 | -2.2% | 0.6872 | 0.4273 | |
| , | | (33.0%, | | (35.3%, | (0.61, 1.36) | (0.75, 1.20) | (-11.8%, 7.4%) | | | |
| | | 47.1%) | | 49.1%) | (, , , , , , , , , , , , , , , , , , , | (************************************** | (| | | |
| >=40 years | 104 | 41 (39.4%) | 100 | 35 (35.0%) | 1.21 | 1.13 | 4.4% | 0.5635 | | |
| | 201 | (30.0%, | -00 | (25.7%, | (0.68, 2.13) | (0.79, 1.61) | (-8.8%, 17.7%) | 0.000 | | |
| | | 49.5%) | | 45.2%) | (0.00, 2.15) | (0), 1.01) | (0.073, 17.770) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Total mortality at 12 weeks (TEAE leading to death) - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|-----------|-------------------|-------------|------------|------------------|--|----------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 2 (0.6%) | 365 | 0 (0.0%) | >99.99 | >99.99 | 0.6% | 0.9162+ | | | |
| | | (0.1%, 2.0%) | | | (>99.99, >99.99) | (>99.99, >99.99) | (-9.7%, 10.8%) | | | | |
| Baseline Disease Severity | | | | | | | | | | | |
| Moderate baseline | 216 | 1 (0.5%) | 220 | 0(0.0%) | 3.07 | 3.06 | 0.5% | 0.4954 | 0.8525 | | |
| disease (IGA=3) | | (0.0%, 2.6%) | | | (0.12, 75.77) | (0.13, 74.59) | (-0.8%, 1.7%) | | | | |
| Severe baseline disease | 146 | 1 (0.7%) | 145 | 0(0.0%) | 3.00 | 2.98 | 0.7% | 1.0000 | | | |
| (IGA=4) | | (0.0%, 3.8%) | | | (0.12, 74.25) | (0.12, 72.54) | (-1.2%, 2.6%) | | | | |
| Gender | | , , , | | | , | , | , | | | | |
| Male | 193 | 1 (0.5%) | 204 | 0 (0.0%) | 3.19 | 3.17 | 0.5% | 0.4861 | 0.9677 | | |
| | | (0.0%, 2.9%) | | | (0.13, 78.71) | (0.13, 77.35) | (-0.9%, 1.9%) | | | | |
| Female | 169 | 1 (0.6%) | 161 | 0 (0.0%) | 2.88 | 2.86 | 0.6% | 1.0000 | | | |
| | | (0.0%, 3.3%) | | | (0.12, 71.10) | (0.12, 69.67) | (-1.1%, 2.2%) | | | | |
| Region | | , , , | | | , | , | , | | | | |
| US/Canada/Australia | 177 | 1 (0.6%) | 195 | 0 (0.0%) | 3.32 | 3.30 | 0.6% | 0.4758 | 0.9997 | | |
| | | (0.0%, 3.1%) | | | (0.13, 82.10) | (0.14, 80.57) | (-0.9%, 2.1%) | | | | |
| Europe | 150 | 1 (0.7%) | 132 | 0 (0.0%) | 2.66 | 2.64 | 0.6% | 1.0000 | | | |
| | | (0.0%, 3.7%) | | | (0.11, 65.83) | (0.11, 64.31) | (-1.3%, 2.5%) | | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 | 1.11 | 0.3% | NE | | | |
| | - / | 0 (0.070) | | 0 (0.070) | (0.02, 59.20) | (0.02, 53.16) | (-9.9%, 10.5%) | 1,2 | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 | 1.05 | 0.1% | NE | | | |
| Laun America | 10 | 0 (0.0%) | 19 | 0 (0.070) | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | NE | | | |
| | | | | • | (0.02, 33.92) | (0.02, 30.43) | (-3.6%, 10.1%) | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Total mortality at 12 weeks (TEAE leading to death) - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|-----------------|-------------|------------|---------------|--|---------------|---------------|--------------|--|
| | | | | | | CMH or Logistic P-val [2] for | | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 0 (0.0%) | 247 | 0 (0.0%) | 1.07 | 1.07 | 0.0% | NE | 0.2880 | |
| • | | | | | (0.02, 54.34) | (0.02, 53.88) | (-0.8%, 0.8%) | | | |
| >=40 years | 132 | 2 (1.5%) | 118 | 0 (0.0%) | 4.54 | 4.47 | 1.5% | 0.4996 | | |
| • | | (0.2%, 5.4%) | | | (0.22, 95.54) | (0.22, 92.25) | (-1.1%, 4.0%) | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

| | Abrocitin | ib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg | | | |
|---------------------------|-----------|----------------------------------|-------------|----------------------------------|-----------------------|----------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 228 (63.0%) (57.8%, 68.0%) | 365 | 182 (49.9%) (44.6%, 55.1%) | 1.71 (1.27, 2.30) | 1.26 (1.11, 1.44) | 13.1% (6.0%, 20.3%) | 0.000 | |
| Baseline Disease Severity | | , | | , | | | | | |
| Moderate baseline | 216 | 144 (66.7%) | 220 | 110 (50.0%) | 2.00 | 1.33 | 16.7% | 0.0005* | 0.1736 |
| disease (IGA=3) | | (60.0%, 72.9%) | | (43.2%, 56.8%) | (1.36, 2.95) | (1.13, 1.57) | (7.5%, 25.8%) | | |
| Severe baseline disease | 146 | 84 (57.5%) | 145 | 72 (49.7%) | 1.37 | 1.16 | 7.9% | 0.1968 | |
| (IGA=4) | | (49.1%, 65.7%) | | (41.3%, 58.1%) | (0.87, 2.18) | (0.93, 1.44) | (-3.5%, 19.3%) | | |
| Gender | | , | | , | | | | | |
| Male | 193 | 117 (60.6%) (53.3%, 67.6%) | 204 | 94 (46.1%) (39.1%, 53.2%) | 1.80 (1.21, 2.68) | 1.32 (1.09, 1.59) | 14.5% (4.8%, 24.3%) | 0.0048* | 0.8388 |
| Female | 169 | 111 (65.7%) (58.0%, 72.8%) | 161 | 88 (54.7%) (46.6%, 62.5%) | 1.59 (1.02, 2.47) | 1.20 (1.01, 1.44) | 11.0% (0.5%, 21.5%) | 0.0435* | |
| Region | | , | | ŕ | | | | | |
| US/Canada/Australia | 177 | 118 (66.7%) (59.2%, 73.6%) | 195 | 100 (51.3%) (44.0%, 58.5%) | 1.90 (1.25, 2.89) | 1.30 (1.09, 1.54) | 15.4% (5.5%, 25.3%) | 0.0031* | 0.5431 |
| Europe | 150 | 92 (61.3%) (53.0%, 69.2%) | 132 | 66 (50.0%) (41.2%, 58.8%) | 1.59 (0.99, 2.55) | 1.23 (0.99, 1.52) | 11.3% (-0.2%, 22.9%) | 0.0712 | |
| Asia | 17 | 8 (47.1%) (23.0%, 72.2%) | 19 | 10 (52.6%) (28.9%, 75.6%) | 0.80 (0.22, 2.97) | 0.89 (0.46, 1.73) | -5.6% (-38.2%, 27.1%) | 1.0000 | |
| Latin America | 18 | 10 (55.6%) (30.8%, 78.5%) | 19 | 6 (31.6%) (12.6%, 56.6%) | 2.71 (0.71, 10.36) | 1.76 (0.81, 3.84) | 24.0% (-7.1%, 55.0%) | 0.1914 | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| P- | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|---------------------|-------------|------------------------|----------------|--|----------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 145 (63.0%) | 247 | 120 (48.6%) | 1.81 | 1.30 | 14.5% | 0.0017* | 0.6440 | |
| • | | (56.5%, | | (42.2%, | (1.25, 2.60) | (1.10, 1.53) | (5.6%, 23.3%) | | | |
| | | 69.3%) | | 55.0%) | | | | | | |
| >=40 years | 132 | 83 (62.9%) | 118 | 62 (52.5%) | 1.53 | 1.20 | 10.3% | 0.1234 | | |
| • | | (54.0%, | | (43.1%, | (0.92, 2.54) | (0.96, 1.49) | (-1.9%, 22.5%) | | | |
| | | 71.1%) | | 61.8%) | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300 | mg Q2W | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|-----------------------------------|-----------|--------------------------|------------|--------------------------|--|-----------------------|------------------------|----------------------------|--------------------------|--|
| | | | | | | | _ | CMH or Logistic 1 | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | | , , | | , | | | |
| Safety Set | 362 | 7 (1.9%) (0.8%, 3.9%) | 365 | 5 (1.4%) (0.4%, 3.2%) | 1.42 (0.45, 4.52) | 1.41 (0.45, 4.41) | 0.6% (-1.3%, 2.4%) | 0.5510+ | | |
| Baseline Disease Severity | | (,, | | (,, | (3. 3, 3.) | () | (12 11, 11 11, | | | |
| Moderate baseline disease (IGA=3) | 216 | 3 (1.4%) (0.3%, 4.0%) | 220 | 3 (1.4%) (0.3%, 3.9%) | 1.02 (0.20, 5.10) | 1.02 (0.21, 4.99) | 0.0% (-2.2%, 2.2%) | 1.0000 | 0.5017 | |
| Severe baseline disease | 146 | 4 (2.7%) | 145 | 2 (1.4%) | 2.01 | 1.99 | 1.4% | 0.6842 | | |
| (IGA=4) | | (0.8%, 6.9%) | | (0.2%, 4.9%) | (0.36, 11.17) | (0.37, 10.68) | (-1.9%, 4.6%) | | | |
| Gender | | (, | | (, | , , , | , , , | , , , | | | |
| Male | 193 | 4 (2.1%) (0.6%, 5.2%) | 204 | 2 (1.0%) (0.1%, 3.5%) | 2.14 (0.39, 11.81) | 2.11 (0.39, 11.41) | 1.1% (-1.3%, 3.5%) | 0.4379 | 0.5386 | |
| Female | 169 | 3 (1.8%) (0.4%, 5.1%) | 161 | 3 (1.9%) (0.4%, 5.3%) | 0.95 (0.19, 4.79) | 0.95 (0.20, 4.65) | -0.1% (-3.0%, 2.8%) | 1.0000 | | |
| Region | | (******,*******) | | (311,74, 212,74) | (****, ****) | (3.23, 3.35) | (===,=, ===, | | | |
| US/Canada/Australia | 177 | 4 (2.3%) (0.6%, 5.7%) | 195 | 3 (1.5%) (0.3%, 4.4%) | 1.48 (0.33, 6.71) | 1.47 (0.33, 6.47) | 0.7% (-2.1%, 3.5%) | 0.7128 | 0.9990 | |
| Europe | 150 | 3 (2.0%) (0.4%, 5.7%) | 132 | 2 (1.5%) (0.2%, 5.4%) | 1.33 (0.22, 8.06) | 1.32 (0.22, 7.78) | 0.5% (-2.6%, 3.5%) | 1.0000 | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 (0.02, 59.20) | 1.11 (0.02, 53.16) | 0.3% (-9.9%, 10.5%) | NE | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------|------------|--------------|---------------|--|---------------|---------------|--------------|
| | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 4 (1.7%) | 247 | 4 (1.6%) | 1.08 | 1.07 | 0.1% | 1.0000 | 0.5005 |
| • | | (0.5%, 4.4%) | | (0.4%, 4.1%) | (0.27, 4.35) | (0.27, 4.24) | (-2.2%, 2.4%) | | |
| >=40 years | 132 | 3 (2.3%) | 118 | 1 (0.8%) | 2.72 | 2.68 | 1.4% | 0.6244 | |
| • | | (0.5%, 6.5%) | | (0.0%, 4.6%) | (0.28, 26.52) | (0.28, 25.43) | (-1.6%, 4.5%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|------------------------------------|-----------|--------------------------|------------|--------------------------|------------------------|--|--|---|--|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction | | |
| Overall | | (2370 CI) | | ()570 CI) | (3370 CI) | (55% C1) | (3370 CI) | varae [1] | meraction | | |
| Safety Set | 362 | 5 (1.4%) (0.4%, 3.2%) | 365 | 2 (0.5%) (0.1%, 2.0%) | 2.54 (0.49, 13.19) | 2.52 (0.49, 12.91) | 0.8% (-0.6%, 2.3%) | 0.2504+ | | | |
| Baseline Disease Severity | | (| | (, | (===, ===, | (, , , , , , , , , , , , , , , , , , , | (,, | | | | |
| Moderate baseline disease (IGA=3) | 216 | 4 (1.9%) (0.5%, 4.7%) | 220 | 2 (0.9%) (0.1%, 3.2%) | 2.06 (0.37, 11.35) | 2.04 (0.38, 11.01) | 0.9% (-1.2%, 3.1%) | 0.4462 | 0.8551 | | |
| Severe baseline disease (IGA=4) | 146 | 1 (0.7%) (0.0%, 3.8%) | 145 | 0 (0.0%) | 3.00 (0.12, 74.25) | 2.98 (0.12, 72.54) | 0.7% (-1.2%, 2.6%) | 1.0000 | | | |
| Gender | | (0.070, 5.070) | | | (0.12, 725) | (0112, 72101) | (1.270, 2.070) | | | | |
| Male | 193 | 3 (1.6%) (0.3%, 4.5%) | 204 | 1 (0.5%) (0.0%, 2.7%) | 3.21 (0.33, 31.08) | 3.17 (0.33, 30.22) | 1.1% (-0.9%, 3.1%) | 0.3597 | 0.7294 | | |
| Female | 169 | 2 (1.2%) (0.1%, 4.2%) | 161 | 1 (0.6%) (0.0%, 3.4%) | 1.92 (0.17, 21.34) | 1.91 (0.17, 20.81) | 0.6% (-1.5%, 2.6%) | 1.0000 | | | |
| Region | | (*****, *****) | | (*****,*****) | (5151, 25151) | (*****, =*****) | (-10 / 10 / 10 / 10 / 10 / 10 / 10 / 10 | | | | |
| US/Canada/Australia | 177 | 2 (1.1%) (0.1%, 4.0%) | 195 | 2 (1.0%) (0.1%, 3.7%) | 1.10 (0.15, 7.91) | 1.10 (0.16, 7.74) | 0.1% (-2.0%, 2.2%) | 1.0000 | 0.7499 | | |
| Europe | 150 | 3 (2.0%) (0.4%, 5.7%) | 132 | 0 (0.0%) | 6.29 (0.32, 122.87) | 6.17 (0.32, 118.27) | 1.9% (-0.7%, 4.6%) | 0.2504 | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 (0.02, 59.20) | 1.11 (0.02, 53.16) | 0.3% (-9.9%, 10.5%) | NE | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | (-9.8%, 10.1%) (-9.8%, 10.1%) | NE | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitir | ib 200mg QD Dupi | lumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------|------------|--------------|---------------|--|---------------|---------------|--------------|
| | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 2 (0.9%) | 247 | 1 (0.4%) | 2.16 | 2.15 | 0.5% | 0.6113 | 0.5703 |
| • | | (0.1%, 3.1%) | | (0.0%, 2.2%) | (0.19, 23.96) | (0.20, 23.53) | (-1.0%, 1.9%) | | |
| >=40 years | 132 | 3 (2.3%) | 118 | 1 (0.8%) | 2.72 | 2.68 | 1.4% | 0.6244 | |
| , | | (0.5%, 6.5%) | | (0.0%, 4.6%) | (0.28, 26.52) | (0.28, 25.43) | (-1.6%, 4.5%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| Adiocidi | nib 200mg QD Dupi | iumad 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|----------|---------------------------------|---|--|---|--|---|---|---|--|--|
| | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| | | | | | | | | | | |
| 362 | 10 (2.8%) | 365 | 4 (1.1%) | 2.56 | 2.52 | 1.7% | 0.1021+ | | | |
| | (1.3%, 5.0%) | | (0.3%, 2.8%) | (0.80, 8.25) | (0.80, 7.96) | (-0.3%, 3.7%) | | | | |
| | | | | | | | | | | |
| 216 | 8 (3.7%) | 220 | 3 (1.4%) | 2.78 | 2.72 | 2.3% | 0.1378 | 0.3852 | | |
| | (1.6%, 7.2%) | | (0.3%, 3.9%) | (0.73, 10.63) | (0.73, 10.10) | (-0.6%, 5.3%) | | | | |
| 146 | 2 (1.4%) | 145 | 1 (0.7%) | 2.00 | 1.99 | 0.7% | 1.0000 | | | |
| | (0.2%, 4.9%) | | (0.0%, 3.8%) | (0.18, 22.30) | (0.18, 21.67) | (-1.6%, 3.0%) | | | | |
| | (, | | (, | , | , , , | , , , | | | | |
| 193 | 2(1.0%) | 204 | 1 (0.5%) | 2.13 | 2.11 | 0.5% | 0.6139 | 0.2710 | | |
| | (0.1%, 3.7%) | | | (0.19, 23.63) | (0.19, 23.13) | (-1.2%, 2.3%) | | | | |
| 169 | 8 (4.7%) | 161 | 3 (1.9%) | 2.62 | 2.54 | 2.9% | 0.2202 | | | |
| | (2.1%, 9.1%) | | (0.4%, 5.3%) | (0.68, 10.04) | (0.69, 9.41) | (-1.0%, 6.7%) | | | | |
| | (, , | | (,, | (, , , , , , , , , , , , , , , , , , , | (,, | (,, | | | | |
| 177 | 6 (3.4%) | 195 | 2 (1.0%) | 3.39 | 3.31 | 2.4% | 0.1579 | 0.9284 | | |
| | ` , | | ` , | (0.67, 17.00) | (0.68, 16.16) | (-0.7%, 5.4%) | | | | |
| 150 | ` ' ' | 132 | , , | ` ' ' | , , , | | 0.6880 | | | |
| | ` / | | ` / | | | | ******* | | | |
| 17 | ` ' ' | 19 | , , | | | | NE | | | |
| | = (5.070) | | = (=.070) | | | | 112 | | | |
| 18 | 0 (0 0%) | 19 | 0 (0 0%) | | | | NF | | | |
| 10 | 0 (0.070) | 1) | 0 (0.070) | | | | ILL | | | |
| • | 362 216 146 193 169 | n (95% CI) 362 10 (2.8%) (1.3%, 5.0%) 216 8 (3.7%) (1.6%, 7.2%) 146 2 (1.4%) (0.2%, 4.9%) 193 2 (1.0%) (0.1%, 3.7%) 169 8 (4.7%) (2.1%, 9.1%) 177 6 (3.4%) (1.3%, 7.2%) 150 4 (2.7%) (0.7%, 6.7%) 17 0 (0.0%) | n (95% CI) n 362 10 (2.8%) 365 (1.3%, 5.0%) 216 8 (3.7%) 220 (1.6%, 7.2%) 146 2 (1.4%) 145 (0.2%, 4.9%) 193 2 (1.0%) 204 (0.1%, 3.7%) 169 8 (4.7%) 161 (2.1%, 9.1%) 177 6 (3.4%) 195 (1.3%, 7.2%) 150 4 (2.7%) 132 (0.7%, 6.7%) 17 0 (0.0%) 19 | n (95% CI) n (95% CI) 362 10 (2.8%) (1.3%, 5.0%) 365 4 (1.1%) (0.3%, 2.8%) 216 8 (3.7%) (1.6%, 7.2%) (0.3%, 3.9%) 146 2 (1.4%) (0.2%, 4.9%) 145 1 (0.7%) (0.0%, 3.8%) 193 2 (1.0%) (0.1%, 3.7%) (0.0%, 2.7%) 169 8 (4.7%) (0.1%, 3.7%) (0.4%, 5.3%) 169 8 (4.7%) (2.1%, 9.1%) 161 3 (1.9%) (0.4%, 5.3%) 177 6 (3.4%) (1.3%, 7.2%) (0.1%, 3.7%) (0.1%, 3.7%) 150 4 (2.7%) (0.1%, 3.7%) 132 2 (1.5%) (0.2%, 5.4%) 17 0 (0.0%) 19 0 (0.0%) 19 0 (0.0%) | n (95% CI) n (95% CI) (95% CI) 362 10 (2.8%) (1.3%, 5.0%) 365 (0.3%, 2.8%) 4 (1.1%) (0.3%, 2.8%) 2.56 (0.80, 8.25) 216 8 (3.7%) (1.6%, 7.2%) 220 (0.3%, 3.9%) (0.73, 10.63) (0.73, 10.63) 146 2 (1.4%) (0.2%, 4.9%) 145 (0.0%, 3.8%) 10.7%) (0.0%, 3.8%) 2.00 (0.18, 22.30) 193 2 (1.0%) (0.1%, 3.7%) 204 (0.0%, 2.7%) (0.19, 23.63) (0.19, 23.63) 169 8 (4.7%) (2.1%, 9.1%) 161 (0.4%, 5.3%) 3.39 (0.68, 10.04) 177 6 (3.4%) (1.3%, 7.2%) 195 (0.1%, 3.7%) 2 (1.0%) (0.1%, 3.7%) 3.39 (0.1%, 3.7%) 150 4 (2.7%) (0.7%, 6.7%) 132 (1.5%) (0.2%, 5.4%) 1.78 (0.2%, 5.4%) (0.32, 9.88) (0.32, 9.88) 17 0 (0.0%) 19 (0.00%, 5.4%) (0.32, 9.88) (0.02, 5.4%) | n (95% CI) n (95% CI) (95% CI) (95% CI) 362 10 (2.8%) 365 4 (1.1%) 2.56 2.52 (1.3%, 5.0%) 220 3 (1.4%) 2.78 2.72 (1.6%, 7.2%) (0.3%, 3.9%) (0.73, 10.63) (0.73, 10.10) 146 2 (1.4%) 145 1 (0.7%) 2.00 1.99 (0.2%, 4.9%) (0.0%, 3.8%) (0.18, 22.30) (0.18, 21.67) 193 2 (1.0%) 204 1 (0.5%) 2.13 2.11 (0.1%, 3.7%) (0.0%, 2.7%) (0.19, 23.63) (0.19, 23.13) 169 8 (4.7%) 161 3 (1.9%) 2.62 2.54 (2.1%, 9.1%) (0.4%, 5.3%) (0.68, 10.04) (0.69, 9.41) 177 6 (3.4%) 195 2 (1.0%) 3.39 3.31 (1.3%, 7.2%) (0.1%, 3.7%) (0.67, 17.00) (0.68, 16.16) 150 4 (2.7%) 132 2 (1.5%) 1.78 1.76 (0.7%, 6.7%) (0.2%, 5.4%) (0.32, 9.88) <td>n (95% CI) n (95% CI) (95% CI) (95% CI) (95% CI) 362 10 (2.8%) 365 4 (1.1%) 2.56 2.52 1.7% (1.3%, 5.0%) 20 3 (1.4%) 2.78 2.72 2.3% (1.6%, 7.2%) (0.3%, 3.9%) (0.73, 10.63) (0.73, 10.10) (-0.6%, 5.3%) 146 2 (1.4%) 145 1 (0.7%) 2.00 1.99 0.7% (0.2%, 4.9%) (0.0%, 3.8%) (0.18, 22.30) (0.18, 21.67) (-1.6%, 3.0%) 193 2 (1.0%) 204 1 (0.5%) 2.13 2.11 0.5% (0.1%, 3.7%) (0.0%, 2.7%) (0.19, 23.63) (0.19, 23.13) (-1.2%, 2.3%) 169 8 (4.7%) 161 3 (1.9%) 2.62 2.54 2.9% (2.1%, 9.1%) (0.4%, 5.3%) (0.68, 10.04) (0.69, 9.41) (-1.0%, 6.7%) 177 6 (3.4%) 195 2 (1.0%) 3.39 3.31 2.4% (1.3%, 7.2%) (0.1%, 3.7%) (0.67, 17.00) (</td> <td>$\begin{array}{c ccccccccccccccccccccccccccccccccccc$</td> | n (95% CI) n (95% CI) (95% CI) (95% CI) (95% CI) 362 10 (2.8%) 365 4 (1.1%) 2.56 2.52 1.7% (1.3%, 5.0%) 20 3 (1.4%) 2.78 2.72 2.3% (1.6%, 7.2%) (0.3%, 3.9%) (0.73, 10.63) (0.73, 10.10) (-0.6%, 5.3%) 146 2 (1.4%) 145 1 (0.7%) 2.00 1.99 0.7% (0.2%, 4.9%) (0.0%, 3.8%) (0.18, 22.30) (0.18, 21.67) (-1.6%, 3.0%) 193 2 (1.0%) 204 1 (0.5%) 2.13 2.11 0.5% (0.1%, 3.7%) (0.0%, 2.7%) (0.19, 23.63) (0.19, 23.13) (-1.2%, 2.3%) 169 8 (4.7%) 161 3 (1.9%) 2.62 2.54 2.9% (2.1%, 9.1%) (0.4%, 5.3%) (0.68, 10.04) (0.69, 9.41) (-1.0%, 6.7%) 177 6 (3.4%) 195 2 (1.0%) 3.39 3.31 2.4% (1.3%, 7.2%) (0.1%, 3.7%) (0.67, 17.00) (| $\begin{array}{c ccccccccccccccccccccccccccccccccccc$ | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300i | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------|------------|--------------|---------------|--|---------------|---------------|--------------|
| | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 5 (2.2%) | 247 | 3 (1.2%) | 1.81 | 1.79 | 1.0% | 0.4907 | 0.3617 |
| • | | (0.7%, 5.0%) | | (0.3%, 3.5%) | (0.43, 7.65) | (0.43, 7.41) | (-1.4%, 3.3%) | | |
| >=40 years | 132 | 5 (3.8%) | 118 | 1 (0.8%) | 4.61 | 4.47 | 2.9% | 0.2175 | |
| • | | (1.2%, 8.6%) | | (0.0%, 4.6%) | (0.53, 40.01) | (0.53, 37.71) | (-0.7%, 6.6%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | nib 200mg QD Dup | ilumab 3001 | ng Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | | |
|-----------------------------------|-----------|----------------------------------|-------------|----------------------------------|-----------------------|----------------------|---------------------------|-------------------------------|--------------------------|--|
| | | | | | | _ | _ | CMH or Logistic P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 362 | 228 (63.0%) (57.8%, 68.0%) | 365 | 181 (49.6%) (44.3%, 54.8%) | 1.73 (1.29, 2.33) | 1.27 (1.12, 1.45) | 13.4% (6.2%, 20.5%) | 0.000 | | |
| Baseline Disease Severity | | ŕ | | , | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 144 (66.7%) (60.0%, 72.9%) | 220 | 110 (50.0%) (43.2%, 56.8%) | 2.00 (1.36, 2.95) | 1.33 (1.13, 1.57) | 16.7% (7.5%, 25.8%) | 0.0005* | 0.1984 | |
| Severe baseline disease (IGA=4) | 146 | 84 (57.5%) (49.1%, 65.7%) | 145 | 71 (49.0%) (40.6%, 57.4%) | 1.41 (0.89, 2.24) | 1.17 (0.95, 1.46) | 8.6% (-2.9%, 20.0%) | 0.1593 | | |
| Gender | | 001,707 | | 271.70) | | | | | | |
| Male | 193 | 117 (60.6%) (53.3%, 67.6%) | 204 | 93 (45.6%) (38.6%, 52.7%) | 1.84 (1.23, 2.74) | 1.33 (1.10, 1.61) | 15.0% (5.3%, 24.7%) | 0.0035* | 0.7986 | |
| Female | 169 | 111 (65.7%) (58.0%, 72.8%) | 161 | 88 (54.7%) (46.6%, 62.5%) | 1.59 (1.02, 2.47) | 1.20 (1.01, 1.44) | 11.0% (0.5%, 21.5%) | 0.0435* | | |
| Region | | .=, | | J=10 /11/ | | | | | | |
| US/Canada/Australia | 177 | 118 (66.7%) (59.2%, 73.6%) | 195 | 100 (51.3%) (44.0%, 58.5%) | 1.90 (1.25, 2.89) | 1.30 (1.09, 1.54) | 15.4% (5.5%, 25.3%) | 0.0031* | 0.6674 | |
| Europe | 150 | 92 (61.3%) (53.0%, 69.2%) | 132 | 66 (50.0%) (41.2%, 58.8%) | 1.59 (0.99, 2.55) | 1.23 (0.99, 1.52) | 11.3% (-0.2%, 22.9%) | 0.0712 | | |
| Asia | 17 | 8 (47.1%) (23.0%, 72.2%) | 19 | 9 (47.4%) (24.4%, 71.1%) | 0.99 (0.27, 3.66) | 0.99 (0.50, 1.98) | -0.3% (-33.0%, 32.4%) | 1.0000 | | |
| Latin America | 18 | 10 (55.6%) (30.8%, 78.5%) | 19 | 6 (31.6%) (12.6%, 56.6%) | 2.71 (0.71, 10.36) | 1.76 (0.81, 3.84) | 24.0% (-7.1%, 55.0%) | 0.1914 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|-----------|------------------|-------------|-------------|--------------|--|----------------|---------------|--------------|--|--|
| | | | | | | CMH or Logistic P-val [2] for | | | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 145 (63.0%) | 247 | 119 (48.2%) | 1.83 | 1.31 | 14.9% | 0.0013* | 0.6133 | | |
| | | (56.5%, | | (41.8%, | (1.27, 2.65) | (1.11, 1.54) | (6.0%, 23.7%) | | | | |
| | | 69.3%) | | 54.6%) | | | | | | | |
| >=40 years | 132 | 83 (62.9%) | 118 | 62 (52.5%) | 1.53 | 1.20 | 10.3% | 0.1234 | | | |
| • | | (54.0%, | | (43.1%, | (0.92, 2.54) | (0.96, 1.49) | (-1.9%, 22.5%) | | | | |
| | | 71.1%) | | 61.8%) | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrociti | nib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|----------|------------------|------------|--------------|---------------|--|----------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 7 (1.9%) | 365 | 5 (1.4%) | 1.42 | 1.41 | 0.6% | 0.5510+ | | | |
| • | | (0.8%, 3.9%) | | (0.4%, 3.2%) | (0.45, 4.52) | (0.45, 4.41) | (-1.3%, 2.4%) | | | | |
| Baseline Disease Severity | | | | | | | | | | | |
| Moderate baseline | 216 | 3 (1.4%) | 220 | 3 (1.4%) | 1.02 | 1.02 | 0.0% | 1.0000 | 0.5017 | | |
| disease (IGA=3) | | (0.3%, 4.0%) | | (0.3%, 3.9%) | (0.20, 5.10) | (0.21, 4.99) | (-2.2%, 2.2%) | | | | |
| Severe baseline disease | 146 | 4 (2.7%) | 145 | 2 (1.4%) | 2.01 | 1.99 | 1.4% | 0.6842 | | | |
| (IGA=4) | | (0.8%, 6.9%) | | (0.2%, 4.9%) | (0.36, 11.17) | (0.37, 10.68) | (-1.9%, 4.6%) | | | | |
| Gender | | (, | | (,, | , , , | , | . , , | | | | |
| Male | 193 | 4 (2.1%) | 204 | 2 (1.0%) | 2.14 | 2.11 | 1.1% | 0.4379 | 0.5386 | | |
| | | (0.6%, 5.2%) | | (0.1%, 3.5%) | (0.39, 11.81) | (0.39, 11.41) | (-1.3%, 3.5%) | | | | |
| Female | 169 | 3 (1.8%) | 161 | 3 (1.9%) | 0.95 | 0.95 | -0.1% | 1.0000 | | | |
| | | (0.4%, 5.1%) | | (0.4%, 5.3%) | (0.19, 4.79) | (0.20, 4.65) | (-3.0%, 2.8%) | | | | |
| Region | | (, , | | (,, | (===, ===, | (===, ===, | (, | | | | |
| US/Canada/Australia | 177 | 4 (2.3%) | 195 | 3 (1.5%) | 1.48 | 1.47 | 0.7% | 0.7128 | 0.9990 | | |
| | | (0.6%, 5.7%) | | (0.3%, 4.4%) | (0.33, 6.71) | (0.33, 6.47) | (-2.1%, 3.5%) | | | | |
| Europe | 150 | 3 (2.0%) | 132 | 2 (1.5%) | 1.33 | 1.32 | 0.5% | 1.0000 | | | |
| F- | | (0.4%, 5.7%) | | (0.2%, 5.4%) | (0.22, 8.06) | (0.22, 7.78) | (-2.6%, 3.5%) | -1000 | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 | 1.11 | 0.3% | NE | | | |
| | | = (=.070) | | = (51070) | (0.02, 59.20) | (0.02, 53.16) | (-9.9%, 10.5%) | 1,2 | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 | 1.05 | 0.1% | NE | | | |
| Laum / micrica | 10 | 0 (0.070) | 1) | 0 (0.070) | | | | IVL | | | |
| Laun America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | NE . | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------|------------|--------------|---------------|--|---------------|---------------|--------------|
| | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 4 (1.7%) | 247 | 4 (1.6%) | 1.08 | 1.07 | 0.1% | 1.0000 | 0.5005 |
| • | | (0.5%, 4.4%) | | (0.4%, 4.1%) | (0.27, 4.35) | (0.27, 4.24) | (-2.2%, 2.4%) | | |
| >=40 years | 132 | 3 (2.3%) | 118 | 1 (0.8%) | 2.72 | 2.68 | 1.4% | 0.6244 | |
| , | | (0.5%, 6.5%) | | (0.0%, 4.6%) | (0.28, 26.52) | (0.28, 25.43) | (-1.6%, 4.5%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300 | mg Q2W | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|-----------|-------------------|------------|--------------|--|----------------|----------------|-------------------|---------------|--|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 362 | 5 (1.4%) | 365 | 2 (0.5%) | 2.54 | 2.52 | 0.8% | 0.2504+ | | |
| • | | (0.4%, 3.2%) | | (0.1%, 2.0%) | (0.49, 13.19) | (0.49, 12.91) | (-0.6%, 2.3%) | | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline | 216 | 4 (1.9%) | 220 | 2 (0.9%) | 2.06 | 2.04 | 0.9% | 0.4462 | 0.8551 | |
| disease (IGA=3) | | (0.5%, 4.7%) | | (0.1%, 3.2%) | (0.37, 11.35) | (0.38, 11.01) | (-1.2%, 3.1%) | | | |
| Severe baseline disease | 146 | 1 (0.7%) | 145 | 0 (0.0%) | 3.00 | 2.98 | 0.7% | 1.0000 | | |
| (IGA=4) | | (0.0%, 3.8%) | | | (0.12, 74.25) | (0.12, 72.54) | (-1.2%, 2.6%) | | | |
| Gender | | , , , | | | | | | | | |
| Male | 193 | 3 (1.6%) | 204 | 1 (0.5%) | 3.21 | 3.17 | 1.1% | 0.3597 | 0.7294 | |
| | | (0.3%, 4.5%) | | (0.0%, 2.7%) | (0.33, 31.08) | (0.33, 30.22) | (-0.9%, 3.1%) | | | |
| Female | 169 | 2 (1.2%) | 161 | 1 (0.6%) | 1.92 | 1.91 | 0.6% | 1.0000 | | |
| | | (0.1%, 4.2%) | | (0.0%, 3.4%) | (0.17, 21.34) | (0.17, 20.81) | (-1.5%, 2.6%) | | | |
| Region | | , , , | | , , , | , , | | , , , | | | |
| US/Canada/Australia | 177 | 2 (1.1%) | 195 | 2 (1.0%) | 1.10 | 1.10 | 0.1% | 1.0000 | 0.7499 | |
| | | (0.1%, 4.0%) | | (0.1%, 3.7%) | (0.15, 7.91) | (0.16, 7.74) | (-2.0%, 2.2%) | | | |
| Europe | 150 | 3 (2.0%) | 132 | 0 (0.0%) | 6.29 | 6.17 | 1.9% | 0.2504 | | |
| | | (0.4%, 5.7%) | | . () | (0.32, 122.87) | (0.32, 118.27) | (-0.7%, 4.6%) | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 | 1.11 | 0.3% | NE | | |
| | -, | = (=.070) | | - (5.070) | (0.02, 59.20) | (0.02, 53.16) | (-9.9%, 10.5%) | 1.2 | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 | 1.05 | 0.1% | NE | | |
| Laur / micrica | 10 | 0 (0.070) | 1) | 0 (0.070) | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | T\L | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events at 12 weeks excluding progression events - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitir | nib 200mg QD Dupi | ilumab 300ı | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|--------------------------|-------------|--------------------------|-----------------------|--|-----------------------|--------------------------------|-------------------------------|--|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic Regression p- | P-val [2] for trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 2 (0.9%) (0.1%, 3.1%) | 247 | 1 (0.4%) (0.0%, 2.2%) | 2.16 (0.19, 23.96) | 2.15 (0.20, 23.53) | 0.5% (-1.0%, 1.9%) | 0.6113 | 0.5703 | |
| >=40 years | 132 | 3 (2.3%) (0.5%, 6.5%) | 118 | 1 (0.8%) (0.0%, 4.6%) | 2.72 (0.28, 26.52) | 2.68 (0.28, 25.43) | 1.4% (-1.6%, 4.5%) | 0.6244 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Superinfections at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | lumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|-----------|------------------------------|-----------|---------------------------|-----------------------|--|------------------------|---|--|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 14 (3.9%) (2.1%, 6.4%) | 365 | 10 (2.7%) (1.3%, 5.0%) | 1.43 (0.63, 3.26) | 1.41 (0.64, 3.14) | 1.1% (-1.5%, 3.7%) | 0.3949+ | | | |
| Baseline Disease Severity | | , , , | | , , , | , , | , , | , , , | | | | |
| Moderate baseline | 216 | 6 (2.8%) | 220 | 5 (2.3%) | 1.23 | 1.22 | 0.5% | 0.7697 | 0.5869 | | |
| disease (IGA=3) | | (1.0%, 5.9%) | | (0.7%, 5.2%) | (0.37, 4.09) | (0.38, 3.95) | (-2.4%, 3.5%) | | | | |
| Severe baseline disease | 146 | 8 (5.5%) | 145 | 5 (3.4%) | 1.62 | 1.59 | 2.0% | 0.5722 | | | |
| (IGA=4) | | (2.4%, 10.5%) | | (1.1%, 7.9%) | (0.52, 5.08) | (0.53, 4.74) | (-2.7%, 6.8%) | | | | |
| Gender | | , | | | | | | | | | |
| Male | 193 | 8 (4.1%) (1.8%, 8.0%) | 204 | 6 (2.9%) (1.1%, 6.3%) | 1.43 (0.49, 4.19) | 1.41 (0.50, 3.99) | 1.2% (-2.4%, 4.8%) | 0.5924 | 0.9565 | | |
| Female | 169 | 6 (3.6%) (1.3%, 7.6%) | 161 | 4 (2.5%) (0.7%, 6.2%) | 1.44 (0.40, 5.22) | 1.43 (0.41, 4.97) | 1.1% (-2.6%, 4.7%) | 0.7508 | | | |
| Region | | , , , | | , , , | | | | | | | |
| US/Canada/Australia | 177 | 8 (4.5%) (2.0%, 8.7%) | 195 | 4 (2.1%) (0.6%, 5.2%) | 2.26 (0.67, 7.64) | 2.20 (0.68, 7.19) | 2.5% (-1.2%, 6.1%) | 0.2422 | 0.5669 | | |
| Europe | 150 | 5 (3.3%) (1.1%, 7.6%) | 132 | 6 (4.5%) (1.7%, 9.6%) | 0.72 (0.22, 2.43) | 0.73 (0.23, 2.35) | -1.2% (-5.8%, 3.4%) | 0.7602 | | | |
| Asia | 17 | 1 (5.9%) (0.1%, 28.7%) | 19 | 0 (0.0%) | 3.55 (0.14, 93.01) | 3.33 (0.14, 76.75) | 5.8% (-8.7%, 20.3%) | 0.4722 | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Superinfections at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|-------------------|------------|--------------|--------------|--|---------------|-----------------|---------------|
| | | | | | | | | CMH or Logistic | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 6 (2.6%) | 247 | 6 (2.4%) | 1.08 | 1.07 | 0.2% | 1.0000 | 0.4081 |
| • | | (1.0%, 5.6%) | | (0.9%, 5.2%) | (0.34, 3.38) | (0.35, 3.28) | (-2.6%, 3.0%) | | |
| >=40 years | 132 | 8 (6.1%) | 118 | 4 (3.4%) | 1.84 | 1.79 | 2.7% | 0.3853 | |
| , | | (2.7%, | | (0.9%, 8.5%) | (0.54, 6.27) | (0.55, 5.79) | (-2.5%, 7.9%) | | |
| | | 11.6%) | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Herpes Zoster at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|-----------|-------------------|-------------|------------|------------------|--|----------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic P | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 2 (0.6%) | 365 | 0(0.0%) | >99.99 | >99.99 | 0.6% | 0.9162+ | | | |
| • | | (0.1%, 2.0%) | | | (>99.99, >99.99) | (>99.99, >99.99) | (-9.7%, 10.8%) | | | | |
| Baseline Disease Severity | | | | | | | | | | | |
| Moderate baseline | 216 | 1 (0.5%) | 220 | 0(0.0%) | 3.07 | 3.06 | 0.5% | 0.4954 | 0.8525 | | |
| disease (IGA=3) | | (0.0%, 2.6%) | | | (0.12, 75.77) | (0.13, 74.59) | (-0.8%, 1.7%) | | | | |
| Severe baseline disease | 146 | 1 (0.7%) | 145 | 0(0.0%) | 3.00 | 2.98 | 0.7% | 1.0000 | | | |
| (IGA=4) | | (0.0%, 3.8%) | | | (0.12, 74.25) | (0.12, 72.54) | (-1.2%, 2.6%) | | | | |
| Gender | | , , , | | | | | | | | | |
| Male | 193 | 1 (0.5%) | 204 | 0 (0.0%) | 3.19 | 3.17 | 0.5% | 0.4861 | 0.9677 | | |
| | | (0.0%, 2.9%) | | | (0.13, 78.71) | (0.13, 77.35) | (-0.9%, 1.9%) | | | | |
| Female | 169 | 1 (0.6%) | 161 | 0(0.0%) | 2.88 | 2.86 | 0.6% | 1.0000 | | | |
| | | (0.0%, 3.3%) | | | (0.12, 71.10) | (0.12, 69.67) | (-1.1%, 2.2%) | | | | |
| Region | | , , , | | | , | , | | | | | |
| US/Canada/Australia | 177 | 2 (1.1%) | 195 | 0 (0.0%) | 5.57 | 5.51 | 1.1% | 0.2257 | 0.7853 | | |
| | | (0.1%, 4.0%) | | | (0.27, 116.81) | (0.27, 113.90) | (-0.7%, 3.0%) | | | | |
| Europe | 150 | 0 (0.0%) | 132 | 0 (0.0%) | 0.88 | 0.88 | -0.0% | NE | | | |
| 1 | | ` / | | | (0.02, 44.68) | (0.02, 44.08) | (-1.4%, 1.3%) | | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 | 1.11 | 0.3% | NE | | | |
| | | ,, | | | (0.02, 59.20) | (0.02, 53.16) | (-9.9%, 10.5%) | | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 | 1.05 | 0.1% | NE | | | |
| | 10 | 2 (2.070) | | . (2.070) | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | 1,2 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Herpes Zoster at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|-----------------|-------------|------------|---------------|--|---------------|---------------|--------------|
| | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 0 (0.0%) | 247 | 0 (0.0%) | 1.07 | 1.07 | 0.0% | NE | 0.2880 |
| • | | | | | (0.02, 54.34) | (0.02, 53.88) | (-0.8%, 0.8%) | | |
| >=40 years | 132 | 2 (1.5%) | 118 | 0 (0.0%) | 4.54 | 4.47 | 1.5% | 0.4996 | |
| • | | (0.2%, 5.4%) | | | (0.22, 95.54) | (0.22, 92.25) | (-1.1%, 4.0%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

| | Abrocitii | nib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-----------------------------------|-----------|-------------------------------|-------------|-------------------------------|-----------------------|--|--------------------------|----------------------------|-----------------------------|--|
| | | | | | | | | CMH or Logistic | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 362 | 9 (2.5%) (1.1%, 4.7%) | 365 | 28 (7.7%) (5.2%, 10.9%) | 0.31 (0.14, 0.66) | 0.32 (0.16, 0.68) | -5.2% (-8.4%, -2.0%) | 0.001 | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 3 (1.4%) (0.3%, 4.0%) | 220 | 17 (7.7%) (4.6%, 12.1%) | 0.17 (0.05, 0.58) | 0.18 (0.05, 0.60) | -6.3% (-10.2%, -2.5%) | 0.0021* | 0.4179 | |
| Severe baseline disease (IGA=4) | 146 | 6 (4.1%) (1.5%, 8.7%) | 145 | 11 (7.6%) (3.8%, 13.2%) | 0.52 (0.19, 1.45) | 0.54 (0.21, 1.43) | -3.5% (-8.9%, 1.9%) | 0.2230 | | |
| Gender | | | | , | | | | | | |
| Male | 193 | 8 (4.1%) (1.8%, 8.0%) | 204 | 14 (6.9%) (3.8%, 11.2%) | 0.59 (0.24, 1.43) | 0.60 (0.26, 1.41) | -2.7% (-7.2%, 1.7%) | 0.2768 | 0.0986 | |
| Female | 169 | 1 (0.6%) (0.0%, 3.3%) | 161 | 14 (8.7%) (4.8%, 14.2%) | 0.06 (0.01, 0.48) | 0.07 (0.01, 0.51) | -8.1% (-12.6%, -3.6%) | 0.0003* | | |
| Region | | | | , | | | | | | |
| US/Canada/Australia | 177 | 4 (2.3%) (0.6%, 5.7%) | 195 | 15 (7.7%) (4.4%, 12.4%) | 0.28 (0.09, 0.85) | 0.29 (0.10, 0.87) | -5.4% (-9.8%, -1.1%) | 0.0188* | 0.4118 | |
| Europe | 150 | 3 (2.0%) (0.4%, 5.7%) | 132 | 12 (9.1%) (4.8%, 15.3%) | 0.20 (0.06, 0.74) | 0.22 (0.06, 0.76) | -7.1% (-12.5%, -1.7%) | 0.0140* | | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 2.40 (0.20, 29.13) | 2.24 (0.22, 22.51) | 6.5% (-11.8%, 24.8%) | 0.5929 | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Conjunctivitis at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| 9- | Abrocitin | nib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|------------------|-------------|------------|--------------|--|----------------|-------------------|--------------|--|
| | | | | | | | | CMH or Logistic I | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 4 (1.7%) | 247 | 18 (7.3%) | 0.23 | 0.24 | -5.5% | 0.0040* | 0.8323 | |
| | | (0.5%, 4.4%) | | (4.4%, | (0.08, 0.68) | (0.08, 0.69) | (-9.2%, -1.9%) | | | |
| | | | | 11.3%) | | | | | | |
| >=40 years | 132 | 5 (3.8%) | 118 | 10 (8.5%) | 0.43 | 0.45 | -4.7% | 0.1811 | | |
| | | (1.2%, 8.6%) | | (4.1%, | (0.14, 1.28) | (0.16, 1.27) | (-10.7%, 1.3%) | | | |
| | | | | 15.0%) | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | Abrocitin | ib 200mg QD Dupi | ilumab 300 | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------------------------|-----------|--------------------------------|------------|------------------------------|------------------------|--|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 36 (9.9%) (7.1%, 13.5%) | 365 | 6 (1.6%) (0.6%, 3.5%) | 6.61 (2.75, 15.88) | 6.05 (2.58, 14.18) | 8.3% (5.0%, 11.6%) | <0.0001+* | |
| Baseline Disease Severity | | ŕ | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 23 (10.6%) (6.9%, 15.5%) | 220 | 2 (0.9%) (0.1%, 3.2%) | 12.99 (3.02, 55.81) | 11.71 (2.80, 49.08) | 9.7% (5.4%, 14.0%) | <0.0001* | 0.3234 |
| Severe baseline disease (IGA=4) | 146 | 13 (8.9%) (4.8%, 14.7%) | 145 | 4 (2.8%) (0.8%, 6.9%) | 3.45 (1.10, 10.83) | 3.23 (1.08, 9.67) | 6.1% (0.8%, 11.5%) | 0.0427* | |
| Gender | | , | | | | | | | |
| Male | 193 | 19 (9.8%) (6.0%, 14.9%) | 204 | 3 (1.5%) (0.3%, 4.2%) | 7.32 (2.13, 25.14) | 6.69 (2.01, 22.26) | 8.4% (3.9%, 12.9%) | 0.0003* | 0.9659 |
| Female | 169 | 17 (10.1%) (6.0%, 15.6%) | 161 | 3 (1.9%) (0.4%, 5.3%) | 5.89 (1.69, 20.51) | 5.40 (1.61, 18.07) | 8.2% (3.2%, 13.2%) | 0.0020* | |
| Region | | , | | | | | | | |
| US/Canada/Australia | 177 | 22 (12.4%) (8.0%, 18.2%) | 195 | 4 (2.1%) (0.6%, 5.2%) | 6.78 (2.29, 20.08) | 6.06 (2.13, 17.24) | 10.4% (5.1%, 15.6%) | <0.0001* | 0.3836 |
| Europe | 150 | 11 (7.3%) (3.7%, 12.7%) | 132 | 1 (0.8%) (0.0%, 4.1%) | 10.37 (1.32, 81.42) | 9.68 (1.27, 73.98) | 6.6% (2.1%, 11.0%) | 0.0065* | |
| Asia | 17 | 3 (17.6%) (3.8%, 43.4%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 3.86 (0.36, 41.20) | 3.35 (0.38, 29.26) | 12.4% (-8.3%, 33.1%) | 0.3255 | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Acne (PT) at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| · | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|-----------------|------------|--------------|----------------|--|---------------|-------------------|---------------|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 25 (10.9%) | 247 | 6 (2.4%) | 4.90 | 4.47 | 8.4% | 0.0003* | 0.9090 | |
| • | | (7.2%, | | (0.9%, 5.2%) | (1.97, 12.17) | (1.87, 10.71) | (4.0%, 12.9%) | | | |
| | | 15.6%) | | | | | | | | |
| >=40 years | 132 | 11 (8.3%) | 118 | 0 (0.0%) | 22.43 | 20.58 | 8.2% | 0.0009* | | |
| • | | (4.2%, | | | (1.31, 384.98) | (1.23, 345.46) | (3.3%, 13.1%) | | | |
| | | 14.4%) | | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Folliculitis (PT) at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitii | nib 200mg QD Dupi | lumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|-----------|-------------------|-----------|--------------|---------------------------------------|--|----------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 7 (1.9%) | 365 | 3 (0.8%) | 2.38 | 2.35 | 1.1% | 0.1984+ | | | |
| - | | (0.8%, 3.9%) | | (0.2%, 2.4%) | (0.61, 9.27) | (0.61, 9.03) | (-0.6%, 2.8%) | | | | |
| Baseline Disease Severity | | | | | | | | | | | |
| Moderate baseline | 216 | 4 (1.9%) | 220 | 1 (0.5%) | 4.13 | 4.07 | 1.4% | 0.2124 | 0.6984 | | |
| disease (IGA=3) | | (0.5%, 4.7%) | | (0.0%, 2.5%) | (0.46, 37.27) | (0.46, 36.16) | (-0.6%, 3.4%) | | | | |
| Severe baseline disease | 146 | 3 (2.1%) | 145 | 2 (1.4%) | 1.50 | 1.49 | 0.7% | 1.0000 | | | |
| (IGA=4) | | (0.4%, 5.9%) | | (0.2%, 4.9%) | (0.25, 9.11) | (0.25, 8.78) | (-2.3%, 3.7%) | | | | |
| Gender | | | | | | | | | | | |
| Male | 193 | 7 (3.6%) | 204 | 2 (1.0%) | 3.80 | 3.70 | 2.6% | 0.0972 | 0.0511 | | |
| | | (1.5%, 7.3%) | | (0.1%, 3.5%) | (0.78, 18.53) | (0.78, 17.59) | (-0.3%, 5.6%) | | | | |
| Female | 169 | 0 (0.0%) | 161 | 1 (0.6%) | 0.32 | 0.32 | -0.6% | 0.4879 | | | |
| | | | | (0.0%, 3.4%) | (0.01, 7.80) | (0.01, 7.74) | (-2.3%, 1.1%) | | | | |
| Region | | | | | | | | | | | |
| US/Canada/Australia | 177 | 4 (2.3%) | 195 | 3 (1.5%) | 1.48 | 1.47 | 0.7% | 0.7128 | 0.9020 | | |
| | | (0.6%, 5.7%) | | (0.3%, 4.4%) | (0.33, 6.71) | (0.33, 6.47) | (-2.1%, 3.5%) | | | | |
| Europe | 150 | 2 (1.3%) | 132 | 0 (0.0%) | 4.46 | 4.40 | 1.3% | 0.5003 | | | |
| _ | | (0.2%, 4.7%) | | | (0.21, 93.77) | (0.21, 90.91) | (-1.0%, 3.6%) | | | | |
| Asia | 17 | 1 (5.9%) | 19 | 0 (0.0%) | 3.55 | 3.33 | 5.8% | 0.4722 | | | |
| | | (0.1%, | | | (0.14, 93.01) | (0.14, 76.75) | (-8.7%, 20.3%) | | | | |
| | | 28.7%) | | | · · · · · · · · · · · · · · · · · · · | , | | | | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 | 1.05 | 0.1% | NE | | | |
| | | . / | | . , | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Folliculitis (PT) at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitir | nib 200mg QD Dupi | lumab 300i | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|-------------------|------------|--------------|---------------|--|---------------|-------------------|---------------|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 2 (0.9%) | 247 | 2 (0.8%) | 1.07 | 1.07 | 0.1% | 1.0000 | 0.1512 |
| • | | (0.1%, 3.1%) | | (0.1%, 2.9%) | (0.15, 7.69) | (0.15, 7.56) | (-1.6%, 1.7%) | | |
| >=40 years | 132 | 5 (3.8%) | 118 | 1 (0.8%) | 4.61 | 4.47 | 2.9% | 0.2175 | |
| • | | (1.2%, 8.6%) | | (0.0%, 4.6%) | (0.53, 40.01) | (0.53, 37.71) | (-0.7%, 6.6%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Infections and infestations [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300r | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|---------------------------|-----------|---------------------------------|-------------|---------------------------------|----------------------|--|------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 74 (20.4%) (16.4%, 25.0%) | 365 | 71 (19.5%) (15.5%, 23.9%) | 1.06 (0.74, 1.53) | 1.05 (0.79, 1.41) | 1.0% (-4.8%, 6.8%) | 0.7384+ | |
| Baseline Disease Severity | | , | | , | | | | | |
| Moderate baseline | 216 | 49 (22.7%) | 220 | 39 (17.7%) | 1.36 | 1.28 | 5.0% | 0.2328 | 0.1004 |
| disease (IGA=3) | | (17.3%, 28.9%) | | (12.9%, 23.4%) | (0.85, 2.18) | (0.88, 1.86) | (-2.6%, 12.5%) | | |
| Severe baseline disease | 146 | 25 (17.1%) | 145 | 32 (22.1%) | 0.73 | 0.78 | -4.9% | 0.3045 | |
| (IGA=4) | | (11.4%, 24.2%) | | (15.6%, 29.7%) | (0.41, 1.31) | (0.48, 1.24) | (-14.1%, 4.2%) | | |
| Gender | | , | | , | | | | | |
| Male | 193 | 43 (22.3%) | 204 | 36 (17.6%) | 1.34 | 1.26 | 4.6% | 0.2600 | 0.1778 |
| | | (16.6%, 28.8%) | | (12.7%, 23.6%) | (0.82, 2.19) | (0.85, 1.88) | (-3.2%, 12.5%) | | |
| Female | 169 | 31 (18.3%) | 161 | 35 (21.7%) | 0.81 | 0.84 | -3.4% | 0.4921 | |
| | | (12.8%, 25.0%) | | (15.6%, 28.9%) | (0.47, 1.39) | (0.55, 1.30) | (-12.0%, 5.2%) | | |
| Region | | 20.070) | | 20.5 70) | | | | | |
| US/Canada/Australia | 177 | 36 (20.3%) (14.7%, | 195 | 34 (17.4%) (12.4%, | 1.21 (0.72, 2.03) | 1.17 (0.76, 1.78) | 2.9% (-5.1%, 10.9%) | 0.5081 | 0.1630 |
| | | 27.0%) | | 23.5%) | | | | | |
| Europe | 150 | 32 (21.3%) | 132 | 34 (25.8%) | 0.78 | 0.83 | -4.4% | 0.4006 | |
| | | (15.1%, 28.8%) | | (18.5%, 34.1%) | (0.45, 1.36) | (0.54, 1.26) | (-14.4%, 5.5%) | | |
| Asia | 17 | 1 (5.9%) | 19 | 2 (10.5%) | 0.53 | 0.56 | -4.6% | 1.0000 | |
| | | (0.1%, 28.7%) | | (1.3%, 33.1%) | (0.04, 6.44) | (0.06, 5.63) | (-22.4%, 13.1%) | | |
| Latin America | 18 | 5 (27.8%) | 19 | 1 (5.3%) | 6.92 | 5.28 | 22.5% | 0.0897 | |
| | | (9.7%, 53.5%) | | (0.1%, 26.0%) | (0.72, 66.51) | (0.68, 40.91) | (-0.5%, 45.5%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Infections and infestations [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------------|------------|------------------------|----------------|--|----------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 43 (18.7%) | 247 | 46 (18.6%) | 1.00 | 1.00 | 0.1% | 1.0000 | 0.7226 |
| • | | (13.9%, | | (14.0%, | (0.63, 1.59) | (0.69, 1.46) | (-6.9%, 7.1%) | | |
| | | 24.3%) | | 24.0%) | | | | | |
| >=40 years | 132 | 31 (23.5%) | 118 | 25 (21.2%) | 1.14 | 1.11 | 2.3% | 0.7615 | |
| • | | (16.5%, | | (14.2%, | (0.63, 2.08) | (0.70, 1.76) | (-8.0%, 12.6%) | | |
| | | 31.6%) | | 29.7%) | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Gastrointestinal disorders [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-----------------------------------|-----------|---------------------------------|------------|-------------------------------|-----------------------|--|---|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | (*******/ | | (*******/ | (| , | (************************************** | | | |
| Safety Set | 362 | 88 (24.3%) (20.0%, 29.1%) | 365 | 23 (6.3%) (4.0%, 9.3%) | 4.78 (2.94, 7.76) | 3.86 (2.50, 5.96) | 18.0% (12.9%, 23.1%) | <0.0001+* | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 55 (25.5%) (19.8%, 31.8%) | 220 | 18 (8.2%) (4.9%, 12.6%) | 3.83 (2.17, 6.79) | 3.11 (1.89, 5.12) | 17.3% (10.4%, 24.1%) | <0.0001* | 0.8460 | |
| Severe baseline disease (IGA=4) | 146 | 33 (22.6%) (16.1%, 30.3%) | 145 | 5 (3.4%) (1.1%, 7.9%) | 8.18 (3.09, 21.63) | 6.55 (2.63, 16.32) | 19.2% (11.7%, 26.6%) | <0.0001* | | |
| Gender | | , | | | | | | | | |
| Male | 193 | 31 (16.1%) (11.2%, 22.0%) | 204 | 7 (3.4%) (1.4%, 6.9%) | 5.39 (2.31, 12.55) | 4.68 (2.11, 10.38) | 12.6% (6.9%, 18.4%) | <0.0001* | 0.0150* | |
| Female | 169 | 57 (33.7%) (26.6%, 41.4%) | 161 | 16 (9.9%) (5.8%, 15.6%) | 4.61 (2.51, 8.46) | 3.39 (2.04, 5.66) | 23.8% (15.3%, 32.3%) | <0.0001* | | |
| Region | | , | | , | | | | | | |
| US/Canada/Australia | 177 | 50 (28.2%) (21.7%, 35.5%) | 195 | 10 (5.1%) (2.5%, 9.2%) | 7.28 (3.56, 14.90) | 5.51 (2.88, 10.53) | 23.1% (15.8%, 30.4%) | <0.0001* | 0.0142* | |
| Europe | 150 | 36 (24.0%) (17.4%, 31.6%) | 132 | 10 (7.6%) (3.7%, 13.5%) | 3.85 (1.83, 8.12) | 3.17 (1.64, 6.13) | 16.4% (8.2%, 24.6%) | 0.0002* | | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 3 (15.8%) (3.4%, 39.6%) | 0.71 (0.10, 4.86) | 0.75 (0.14, 3.94) | -4.0% (-26.5%, 18.4%) | 1.0000 | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Gastrointestinal disorders [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD CMH or Logistic P-val [2] for | | | | | |
|--------------------|-----------|---------------------------------|-------------|---------------------------------------|----------------------|---|-------------------------|----------------------------|--------------|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup | | |
| Age Subgroup | | | | · · · · · · · · · · · · · · · · · · · | | · | | | | | |
| <40 years | 230 | 62 (27.0%) (21.3%, 33.2%) | 247 | 16 (6.5%) (3.7%, 10.3%) | 5.33 (2.97, 9.56) | 4.16 (2.48, 7.00) | 20.5% (14.0%, 27.0%) | <0.0001* | 0.1790 | | |
| >=40 years | 132 | 26 (19.7%) (13.3%, 27.5%) | 118 | 7 (5.9%) (2.4%, 11.8%) | 3.89 (1.62, 9.34) | 3.32 (1.50, 7.37) | 13.8% (5.8%, 21.8%) | 0.0013* | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Skin and subcutaneous tissue disorders[SOC] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|---------------------------|-----------|---------------------------------|-------------|--------------------------------|----------------------|--|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic l Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 59 (16.3%) (12.6%, 20.5%) | 365 | 36 (9.9%) (7.0%, 13.4%) | 1.78 (1.14, 2.77) | 1.65 (1.12, 2.44) | 6.4% (1.6%, 11.3%) | 0.009 | |
| Baseline Disease Severity | | , | | , | | | | | |
| Moderate baseline | 216 | 35 (16.2%) | 220 | 20 (9.1%) | 1.93 | 1.78 | 7.1% | 0.0302* | 0.7526 |
| disease (IGA=3) | | (11.6%, 21.8%) | | (5.6%, 13.7%) | (1.08, 3.47) | (1.06, 2.99) | (0.9%, 13.3%) | | |
| Severe baseline disease | 146 | 24 (16.4%) | 145 | 16 (11.0%) | 1.59 | 1.49 | 5.4% | 0.2332 | |
| (IGA=4) | | (10.8%, 23.5%) | | (6.4%, 17.3%) | (0.80, 3.13) | (0.83, 2.69) | (-2.5%, 13.3%) | | |
| Gender | | , | | , | | | | | |
| Male | 193 | 29 (15.0%) (10.3%, 20.9%) | 204 | 17 (8.3%) (4.9%, 13.0%) | 1.95 (1.03, 3.67) | 1.80 (1.02, 3.17) | 6.7% (0.4%, 13.0%) | 0.0419* | 0.9193 |
| Female | 169 | 30 (17.8%) (12.3%, 24.4%) | 161 | 19 (11.8%) (7.3%, 17.8%) | 1.61 (0.87, 3.00) | 1.50 (0.88, 2.56) | 6.0% (-1.7%, 13.6%) | 0.1631 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 177 | 32 (18.1%) (12.7%, 24.6%) | 195 | 19 (9.7%) (6.0%, 14.8%) | 2.04 (1.11, 3.76) | 1.86 (1.09, 3.15) | 8.3% (1.3%, 15.4%) | 0.0234* | 0.4334 |
| Europe | 150 | 24 (16.0%) (10.5%, 22.9%) | 132 | 13 (9.8%) (5.3%, 16.3%) | 1.74 (0.85, 3.58) | 1.62 (0.86, 3.06) | 6.2% (-1.6%, 13.9%) | 0.1576 | |
| Asia | 17 | 3 (17.6%) (3.8%, 43.4%) | 19 | 3 (15.8%) (3.4%, 39.6%) | 1.14 (0.20, 6.60) | 1.12 (0.26, 4.81) | 1.9% (-22.6%, 26.3%) | 1.0000 | |
| Latin America | 18 | 0 (0.0%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 0.33 (0.01, 8.73) | 0.35 (0.02, 8.09) | -4.9% (-18.5%, 8.7%) | 1.0000 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

TABLE 2.16: Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Skin and subcutaneous tissue disorders SOC] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|------------------------|-------------|---------------------------------------|----------------|--|----------------|---|-----------------------------|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | trt*subgroup interaction | |
| Age Subgroup | | | | · · · · · · · · · · · · · · · · · · · | | | | | | |
| <40 years | 230 | 39 (17.0%) | 247 | 23 (9.3%) | 1.99 | 1.82 | 7.6% | 0.0143* | 0.5064 | |
| • | | (12.3%, | | (6.0%, | (1.15, 3.45) | (1.12, 2.95) | (1.6%, 13.7%) | | | |
| | | 22.4%) | | 13.6%) | | | | | | |
| >=40 years | 132 | 20 (15.2%) | 118 | 13 (11.0%) | 1.44 | 1.38 | 4.1% | 0.3560 | | |
| • | | (9.5%, | | (6.0%, | (0.68, 3.05) | (0.72, 2.64) | (-4.2%, 12.5%) | | | |
| | | 22.4%) | | 18.1%) | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Investigations [SOC] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|---------------------------|-----------|---------------------------------|-------------|--------------------------------|------------------------|--|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 52 (14.4%) (10.9%, 18.4%) | 365 | 33 (9.0%) (6.3%, 12.5%) | 1.69 (1.06, 2.68) | 1.59 (1.05, 2.40) | 5.3% (0.7%, 10.0%) | 0.025 | |
| Baseline Disease Severity | | , | | , | | | | | |
| Moderate baseline | 216 | 35 (16.2%) | 220 | 21 (9.5%) | 1.83 | 1.70 | 6.7% | 0.0449* | 0.4691 |
| disease (IGA=3) | | (11.6%, 21.8%) | | (6.0%, 14.2%) | (1.03, 3.26) | (1.02, 2.82) | (0.4%, 12.9%) | | |
| Severe baseline disease | 146 | 17 (11.6%) | 145 | 12 (8.3%) | 1.46 | 1.41 | 3.4% | 0.4343 | |
| (IGA=4) | | (6.9%, 18.0%) | | (4.3%, 14.0%) | (0.67, 3.18) | (0.70, 2.84) | (-3.5%, 10.2%) | | |
| Gender | | , | | , | | | | | |
| Male | 193 | 25 (13.0%) (8.6%, 18.5%) | 204 | 19 (9.3%) (5.7%, 14.2%) | 1.45 (0.77, 2.73) | 1.39 (0.79, 2.44) | 3.6% (-2.6%, 9.8%) | 0.2663 | 0.4406 |
| Female | 169 | 27 (16.0%) (10.8%, 22.4%) | 161 | 14 (8.7%) (4.8%, 14.2%) | 2.00 (1.01, 3.96) | 1.84 (1.00, 3.38) | 7.3% (0.2%, 14.3%) | 0.0471* | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 177 | 19 (10.7%) (6.6%, 16.3%) | 195 | 18 (9.2%) (5.6%, 14.2%) | 1.18 (0.60, 2.33) | 1.16 (0.63, 2.14) | 1.5% (-4.6%, 7.6%) | 0.7292 | 0.3324 |
| Europe | 150 | 27 (18.0%) (12.2%, 25.1%) | 132 | 14 (10.6%) (5.9%, 17.2%) | 1.85 (0.93, 3.70) | 1.70 (0.93, 3.10) | 7.4% (-0.7%, 15.5%) | 0.0914 | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 0 (0.0%) | 6.29 (0.28, 140.86) | 5.56 (0.29, 108.16) | 11.4% (-6.0%, 28.8%) | 0.2159 | |
| Latin America | 18 | 4 (22.2%) (6.4%, 47.6%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 5.14 (0.52, 51.29) | 4.22 (0.52, 34.28) | 17.0% (-4.7%, 38.6%) | 0.1797 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Investigations [SOC] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitini | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|------------|------------------|-------------|------------|--------------|--|---------------|-------------------------------|--------------|--|--|
| | | | | | | | | CMH or Logistic P-val [2] for | | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 30 (13.0%) | 247 | 24 (9.7%) | 1.39 | 1.34 | 3.3% | 0.3115 | 0.2524 | | |
| - | | (9.0%, | | (6.3%, | (0.79, 2.46) | (0.81, 2.23) | (-2.4%, 9.0%) | | | | |
| | | 18.1%) | | 14.1%) | | | | | | | |
| >=40 years | 132 | 22 (16.7%) | 118 | 9 (7.6%) | 2.42 | 2.19 | 9.0% | 0.0349* | | | |
| - | | (10.7%, | | (3.5%, | (1.07, 5.50) | (1.05, 4.56) | (1.1%, 17.0%) | | | | |
| | | 24.1%) | | 14.0%) | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nervous system disorders [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|---------------------------|-----------|------------------------------|------------|------------------------|-----------------------|--|------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall Safety Set | 362 | 60 (16.6%) | 365 | 25 (6.8%) | 2.70 | 2.42 | 9.7% | <0.0001+* | |
| Surety Set | 302 | (12.9%, 20.8%) | 303 | (4.5%, 9.9%) | (1.65, 4.42) | (1.55, 3.77) | (5.1%, 14.4%) | V0.0001 | |
| Baseline Disease Severity | | 20.070) | | | | | | | |
| Moderate baseline | 216 | 36 (16.7%) | 220 | 18 (8.2%) | 2.24 | 2.04 | 8.5% | 0.0086* | 0.5451 |
| disease (IGA=3) | | (12.0%, 22.3%) | | (4.9%, 12.6%) | (1.23, 4.09) | (1.19, 3.47) | (2.3%, 14.6%) | | |
| Severe baseline disease | 146 | 24 (16.4%) | 145 | 7 (4.8%) | 3.88 | 3.41 | 11.6% | 0.0019* | |
| (IGA=4) | | (10.8%, 23.5%) | | (2.0%, 9.7%) | (1.61, 9.32) | (1.51, 7.65) | (4.7%, 18.6%) | | |
| Gender | | | | | | | | | |
| Male | 193 | 30 (15.5%) | 204 | 10 (4.9%) | 3.57 | 3.17 | 10.6% | 0.0004* | 0.7065 |
| | | (10.7%, 21.4%) | | (2.4%, 8.8%) | (1.69, 7.52) | (1.59, 6.31) | (4.7%, 16.6%) | | |
| Female | 169 | 30 (17.8%) | 161 | 15 (9.3%) | 2.10 | 1.91 | 8.4% | 0.0361* | |
| | | (12.3%, 24.4%) | | (5.3%, 14.9%) | (1.08, 4.07) | (1.07, 3.41) | (1.1%, 15.7%) | | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 177 | 32 (18.1%) | 195 | 12 (6.2%) | 3.37 | 2.94 | 11.9% | 0.0004* | 0.6765 |
| | | (12.7%, 24.6%) | | (3.2%, 10.5%) | (1.67, 6.77) | (1.56, 5.52) | (5.3%, 18.5%) | | |
| Europe | 150 | 25 (16.7%) | 132 | 11 (8.3%) | 2.20 | 2.00 | 8.3% | 0.0483* | |
| • | | (11.1%, 23.6%) | | (4.2%, 14.4%) | (1.04, 4.67) | (1.02, 3.91) | (0.7%, 15.9%) | | |
| Asia | 17 | 2(11.8%) | 19 | | 1.13 | 1.12 | 1.2% | 1.0000 | |
| Asia | 1/ | (1.5%, | 19 | 2 (10.5%) (1.3%, | (0.14, 9.07) | (0.18, 7.09) | (-19.4%, 21.9%) | 1.0000 | |
| | | 36.4%) | | 33.1%) | | | | | |
| Latin America | 18 | 1 (5.6%) (0.1%, 27.3%) | 19 | 0 (0.0%) | 3.34 (0.13, 87.52) | 3.16 (0.14, 72.84) | 5.4% (-8.5%, 19.3%) | 0.4865 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nervous system disorders [SOC] – Safety Set JADE DARE (PF-04965842) - 2023 datacut

| - | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|-----------------|-------------|------------|--------------|--|-------------------|---------------|--------------|--|
| | | | | _ | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 38 (16.5%) | 247 | 17 (6.9%) | 2.68 | 2.40 | 9.6% | 0.0014* | 0.9605 | |
| • | | (12.0%, | | (4.1%, | (1.46, 4.89) | (1.39, 4.13) | (3.9%, 15.4%) | | | |
| | | 22.0%) | | 10.8%) | | | | | | |
| >=40 years | 132 | 22 (16.7%) | 118 | 8 (6.8%) | 2.75 | 2.46 | 9.9% | 0.0191* | | |
| • | | (10.7%, | | (3.0%, | (1.17, 6.44) | (1.14, 5.31) | (2.1%, 17.7%) | | | |
| | | 24.1%) | | 12.9%) | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nausea [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | QD | |
|------------------------------------|-----------|---------------------------------|------------|--------------------------|-------------------------|-------------------------|---------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic 1 Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | (5070 01) | | (>0 /0 01) | (5070 61) | (5070-02) | (507,0 02) | , and [1] | - Intertaction |
| Safety Set | 362 | 69 (19.1%) (15.1%, 23.5%) | 365 | 7 (1.9%) (0.8%, 3.9%) | 12.04 (5.45, 26.61) | 9.94 (4.63, 21.33) | 17.1% (12.9%, 21.4%) | <0.0001+* | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 45 (20.8%) (15.6%, 26.9%) | 220 | 6 (2.7%) (1.0%, 5.8%) | 9.39 (3.91, 22.52) | 7.64 (3.33, 17.53) | 18.1% (12.3%, 23.9%) | <0.0001* | 0.5278 |
| Severe baseline disease (IGA=4) | 146 | 24 (16.4%) (10.8%, 23.5%) | 145 | 1 (0.7%) (0.0%, 3.8%) | 28.33 (3.78, 212.45) | 23.84 (3.27, 173.87) | 15.7% (9.6%, 21.9%) | <0.0001* | |
| Gender | | , | | | | | | | |
| Male | 193 | 21 (10.9%) (6.9%, 16.2%) | 204 | 2 (1.0%) (0.1%, 3.5%) | 12.33 (2.85, 53.34) | 11.10 (2.64, 46.70) | 9.9% (5.3%, 14.5%) | <0.0001* | 0.0003* |
| Female | 169 | 48 (28.4%) (21.7%, 35.8%) | 161 | 5 (3.1%) (1.0%, 7.1%) | 12.38 (4.78, 32.04) | 9.15 (3.74, 22.39) | 25.3% (18.0%, 32.6%) | <0.0001* | |
| Region | | , | | | | | | | |
| US/Canada/Australia | 177 | 39 (22.0%) (16.2%, 28.9%) | 195 | 3 (1.5%) (0.3%, 4.4%) | 18.09 (5.48, 59.72) | 14.32 (4.51, 45.53) | 20.5% (14.1%, 26.8%) | <0.0001* | 0.0543 |
| Europe | 150 | 29 (19.3%) (13.3%, 26.6%) | 132 | 4 (3.0%) (0.8%, 7.6%) | 7.67 (2.62, 22.46) | 6.38 (2.30, 17.67) | 16.3% (9.3%, 23.3%) | <0.0001* | |
| Asia | 17 | 1 (5.9%) (0.1%, 28.7%) | 19 | 0 (0.0%) | 3.55 (0.14, 93.01) | 3.33 (0.14, 76.75) | 5.8% (-8.7%, 20.3%) | 0.4722 | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Nausea [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| 8 | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|-----------------|------------|--------------|---------------|--|----------------|------------------------------------|--------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic F Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 52 (22.6%) | 247 | 4 (1.6%) | 17.75 | 13.96 | 21.0% | < 0.0001* | 0.0166* |
| • | | (17.4%, | | (0.4%, 4.1%) | (6.30, 49.97) | (5.13, 37.99) | (15.4%, 26.6%) | | |
| | | 28.6%) | | | | | | | |
| >=40 years | 132 | 17 (12.9%) | 118 | 3 (2.5%) | 5.67 | 5.07 | 10.3% | 0.0040* | |
| • | | (7.7%, | | (0.5%, 7.3%) | (1.62, 19.86) | (1.52, 16.85) | (4.0%, 16.7%) | | |
| | | 19.8%) | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Headache [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-----------------------------------|-----------|--------------------------------|------------|-------------------------------|-----------------------|--|-------------------------|----------------------------|-----------------------------|--|
| | | | | _ | | _ | _ | CMH or Logistic 1 | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | | · · · · · · | | | | | |
| Safety Set | 362 | 43 (11.9%) (8.7%, 15.7%) | 365 | 18 (4.9%) (2.9%, 7.7%) | 2.60 (1.47, 4.60) | 2.41 (1.42, 4.09) | 6.9% (2.9%, 11.0%) | 0.000 | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 26 (12.0%) (8.0%, 17.1%) | 220 | 13 (5.9%) (3.2%, 9.9%) | 2.18 (1.09, 4.36) | 2.04 (1.08, 3.86) | 6.1% (0.8%, 11.5%) | 0.0290* | 0.6372 | |
| Severe baseline disease (IGA=4) | 146 | 17 (11.6%) (6.9%, 18.0%) | 145 | 5 (3.4%) (1.1%, 7.9%) | 3.69 (1.32, 10.29) | 3.38 (1.28, 8.91) | 8.2% (2.2%, 14.2%) | 0.0131* | | |
| Gender | | , | | | | | | | | |
| Male | 193 | 23 (11.9%) (7.7%, 17.3%) | 204 | 7 (3.4%) (1.4%, 6.9%) | 3.81 (1.59, 9.09) | 3.47 (1.53, 7.91) | 8.5% (3.3%, 13.7%) | 0.0019* | 0.4249 | |
| Female | 169 | 20 (11.8%) (7.4%, 17.7%) | 161 | 11 (6.8%) (3.5%, 11.9%) | 1.83 (0.85, 3.95) | 1.73 (0.86, 3.50) | 5.0% (-1.2%, 11.2%) | 0.1339 | | |
| Region | | | | , | | | | | | |
| US/Canada/Australia | 177 | 21 (11.9%) (7.5%, 17.6%) | 195 | 9 (4.6%) (2.1%, 8.6%) | 2.78 (1.24, 6.25) | 2.57 (1.21, 5.46) | 7.2% (1.6%, 12.8%) | 0.0127* | 0.9516 | |
| Europe | 150 | 19 (12.7%) (7.8%, 19.1%) | 132 | 7 (5.3%) (2.2%, 10.6%) | 2.59 (1.05, 6.37) | 2.39 (1.04, 5.50) | 7.4% (0.8%, 13.9%) | 0.0391* | | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 2 (10.5%) (1.3%, 33.1%) | 1.13 (0.14, 9.07) | 1.12 (0.18, 7.09) | 1.2% (-19.4%, 21.9%) | 1.0000 | | |
| Latin America | 18 | 1 (5.6%) (0.1%, 27.3%) | 19 | 0 (0.0%) | 3.34 (0.13, 87.52) | 3.16 (0.14, 72.84) | 5.4% (-8.5%, 19.3%) | 0.4865 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Headache [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitini | ib 200mg QD Dupi | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|------------|------------------|------------|--------------|---------------|--|---------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 26 (11.3%) | 247 | 14 (5.7%) | 2.12 | 1.99 | 5.6% | 0.0314* | 0.3699 | | |
| | | (7.5%, | | (3.1%, 9.3%) | (1.08, 4.17) | (1.07, 3.72) | (0.6%, 10.6%) | | | | |
| | | 16.1%) | | | | | | | | | |
| >=40 years | 132 | 17 (12.9%) | 118 | 4 (3.4%) | 4.21 | 3.80 | 9.5% | 0.0105* | | | |
| • | | (7.7%, | | (0.9%, 8.5%) | (1.38, 12.91) | (1.32, 10.97) | (2.9%, 16.1%) | | | | |
| | | 19.8%) | | | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Acne [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | ilumab 300ı | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-----------------------------------|-----------|--------------------------------|-------------|------------------------------|------------------------|--|-------------------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | | | , | , | , , | , , | | | |
| Safety Set | 362 | 36 (9.9%) (7.1%, 13.5%) | 365 | 6 (1.6%) (0.6%, 3.5%) | 6.61 (2.75, 15.88) | 6.05 (2.58, 14.18) | 8.3% (5.0%, 11.6%) | <0.0001+* | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 23 (10.6%) (6.9%, 15.5%) | 220 | 2 (0.9%) (0.1%, 3.2%) | 12.99 (3.02, 55.81) | 11.71 (2.80, 49.08) | 9.7% (5.4%, 14.0%) | <0.0001* | 0.3234 | |
| Severe baseline disease (IGA=4) | 146 | 13 (8.9%) (4.8%, 14.7%) | 145 | 4 (2.8%) (0.8%, 6.9%) | 3.45 (1.10, 10.83) | 3.23 (1.08, 9.67) | 6.1% (0.8%, 11.5%) | 0.0427* | | |
| Gender | | , | | | | | | | | |
| Male | 193 | 19 (9.8%) (6.0%, 14.9%) | 204 | 3 (1.5%) (0.3%, 4.2%) | 7.32 (2.13, 25.14) | 6.69 (2.01, 22.26) | 8.4% (3.9%, 12.9%) | 0.0003* | 0.9659 | |
| Female | 169 | 17 (10.1%) (6.0%, 15.6%) | 161 | 3 (1.9%) (0.4%, 5.3%) | 5.89 (1.69, 20.51) | 5.40 (1.61, 18.07) | 8.2% (3.2%, 13.2%) | 0.0020* | | |
| Region | | , | | | | | | | | |
| US/Canada/Australia | 177 | 22 (12.4%) (8.0%, 18.2%) | 195 | 4 (2.1%) (0.6%, 5.2%) | 6.78 (2.29, 20.08) | 6.06 (2.13, 17.24) | 10.4% (5.1%, 15.6%) | <0.0001* | 0.3836 | |
| Europe | 150 | 11 (7.3%) (3.7%, 12.7%) | 132 | 1 (0.8%) (0.0%, 4.1%) | 10.37 (1.32, 81.42) | 9.68 (1.27, 73.98) | 6.6% (2.1%, 11.0%) | 0.0065* | | |
| Asia | 17 | 3 (17.6%) (3.8%, 43.4%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 3.86 (0.36, 41.20) | 3.35 (0.38, 29.26) | 12.4% (-8.3%, 33.1%) | 0.3255 | | |
| Latin America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | 1.05 (0.02, 55.92) | 1.05 (0.02, 50.43) | 0.1% (-9.8%, 10.1%) | NE | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value <0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events at 12 weeks - Acne [PT] - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| · | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|-----------|-----------------|------------|--------------|----------------|--|---------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 25 (10.9%) | 247 | 6 (2.4%) | 4.90 | 4.47 | 8.4% | 0.0003* | 0.9090 | | |
| | | (7.2%, | | (0.9%, 5.2%) | (1.97, 12.17) | (1.87, 10.71) | (4.0%, 12.9%) | | | | |
| | | 15.6%) | | | | | | | | | |
| >=40 years | 132 | 11 (8.3%) | 118 | 0(0.0%) | 22.43 | 20.58 | 8.2% | 0.0009* | | | |
| - | | (4.2%, | | | (1.31, 384.98) | (1.23, 345.46) | (3.3%, 13.1%) | | | | |
| | | 14.4%) | | | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (PT)-PT Natural killer cell count decreased at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrociti | nib 200mg QD Dupi | lumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|---------------------------|----------|-------------------|------------|------------|------------------|--|----------------|-------------------|--------------|--|--|
| | | T (0) | | - (0/) | 0.70 | | n n | CMH or Logistic I | | | |
| **** | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Overall | | | | | | | | | | | |
| Safety Set | 362 | 7 (1.9%) | 365 | 0 (0.0%) | >99.99 | >99.99 | 1.9% | 0.7144+ | | | |
| | | (0.8%, 3.9%) | | | (>99.99, >99.99) | (>99.99, >99.99) | (-8.4%, 12.3%) | | | | |
| Baseline Disease Severity | | | | | | | | | | | |
| Moderate baseline | 216 | 4 (1.9%) | 220 | 0(0.0%) | 9.34 | 9.17 | 1.8% | 0.0594 | 0.9084 | | |
| disease (IGA=3) | | (0.5%, 4.7%) | | | (0.50, 174.51) | (0.50, 169.22) | (-0.1%, 3.8%) | | | | |
| Severe baseline disease | 146 | 3 (2.1%) | 145 | 0 (0.0%) | 7.10 | 6.95 | 2.0% | 0.2474 | | | |
| (IGA=4) | | (0.4%, 5.9%) | | | (0.36, 138.64) | (0.36, 133.41) | (-0.6%, 4.7%) | | | | |
| Gender | | , , , | | | , , , | • | , , , | | | | |
| Male | 193 | 2 (1.0%) | 204 | 0 (0.0%) | 5.34 | 5.28 | 1.0% | 0.2357 | 0.2508 | | |
| | | (0.1%, 3.7%) | | | (0.25, 111.93) | (0.26, 109.36) | (-0.7%, 2.8%) | | | | |
| Female | 169 | 5 (3.0%) | 161 | 0 (0.0%) | 10.80 | 10.48 | 2.9% | 0.0610 | | | |
| | | (1.0%, 6.8%) | | | (0.59, 196.90) | (0.58, 188.06) | (0.1%, 5.7%) | | | | |
| Region | | (=10,0,000,0) | | | (0.25, 0.50) | (0.00, 0.0000) | (31272, 21172) | | | | |
| US/Canada/Australia | 177 | 0 (0.0%) | 195 | 0 (0.0%) | 1.10 | 1.10 | 0.0% | NE | 0.1069 | | |
| | | (() () | | | (0.02, 55.80) | (0.02, 55.20) | (-1.0%, 1.1%) | | | | |
| Europe | 150 | 7 (4.7%) | 132 | 0 (0.0%) | 13.85 | 13.21 | 4.6% | 0.0158* | | | |
| 201010 | 130 | (1.9%, 9.4%) | 132 | 0 (0.070) | (0.78, 244.87) | (0.76, 229.14) | (1.0%, 8.2%) | 0.0150 | | | |
| Asia | 17 | 0 (0.0%) | 19 | 0 (0.0%) | 1.11 | 1.11 | 0.3% | NE | | | |
| 2 151U | 17 | 0 (0.070) | 17 | 0 (0.070) | (0.02, 59.20) | (0.02, 53.16) | (-9.9%, 10.5%) | NE | | | |
| Latin America | 18 | 0 (0 00/) | 10 | 0 (0 00/) | 1.05 | 1.05 | 0.1% | NE | | | |
| Laun America | 18 | 0 (0.0%) | 19 | 0 (0.0%) | | | | NE | | | |
| | | | | • | (0.02, 55.92) | (0.02, 50.43) | (-9.8%, 10.1%) | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (PT)-PT Natural killer cell count decreased at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|--------------------|-----------|------------------|------------|------------|----------------|--|---------------|-----------------|---------------|
| | | | | | | | | CMH or Logistic | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Age Subgroup | | | | | | | | | |
| <40 years | 230 | 4 (1.7%) | 247 | 0 (0.0%) | 9.83 | 9.66 | 1.7% | 0.0533 | 0.7901 |
| • | | (0.5%, 4.4%) | | | (0.53, 183.68) | (0.52, 178.48) | (-0.1%, 3.6%) | | |
| >=40 years | 132 | 3 (2.3%) | 118 | 0 (0.0%) | 6.41 | 6.26 | 2.2% | 0.2493 | |
| • | | (0.5%, 6.5%) | | | (0.33, 125.31) | (0.33, 120.00) | (-0.7%, 5.2%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | |
|---------------------------|-----------|--------------------------------|------------|-------------------------------|-----------------------|--|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 43 (11.9%) (8.7%, 15.7%) | 365 | 18 (4.9%) (2.9%, 7.7%) | 2.60 (1.47, 4.60) | 2.41 (1.42, 4.09) | 6.9% (2.9%, 11.0%) | 0.000 | |
| Baseline Disease Severity | | , | | | | | | | |
| Moderate baseline | 216 | 26 (12.0%) | 220 | 13 (5.9%) | 2.18 | 2.04 | 6.1% | 0.0290* | 0.6372 |
| disease (IGA=3) | | (8.0%, 17.1%) | | (3.2%, 9.9%) | (1.09, 4.36) | (1.08, 3.86) | (0.8%, 11.5%) | | |
| Severe baseline disease | 146 | 17 (11.6%) | 145 | 5 (3.4%) | 3.69 | 3.38 | 8.2% | 0.0131* | |
| (IGA=4) | | (6.9%, 18.0%) | | (1.1%, 7.9%) | (1.32, 10.29) | (1.28, 8.91) | (2.2%, 14.2%) | | |
| Gender | | , | | | | | | | |
| Male | 193 | 23 (11.9%) (7.7%, 17.3%) | 204 | 7 (3.4%) (1.4%, 6.9%) | 3.81 (1.59, 9.09) | 3.47 (1.53, 7.91) | 8.5% (3.3%, 13.7%) | 0.0019* | 0.4249 |
| Female | 169 | 20 (11.8%) (7.4%, 17.7%) | 161 | 11 (6.8%) (3.5%, 11.9%) | 1.83 (0.85, 3.95) | 1.73 (0.86, 3.50) | 5.0% (-1.2%, 11.2%) | 0.1339 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 177 | 21 (11.9%) (7.5%, 17.6%) | 195 | 9 (4.6%) (2.1%, 8.6%) | 2.78 (1.24, 6.25) | 2.57 (1.21, 5.46) | 7.2% (1.6%, 12.8%) | 0.0127* | 0.9516 |
| Europe | 150 | 19 (12.7%) (7.8%, 19.1%) | 132 | 7 (5.3%) (2.2%, 10.6%) | 2.59 (1.05, 6.37) | 2.39 (1.04, 5.50) | 7.4% (0.8%, 13.9%) | 0.0391* | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 2 (10.5%) (1.3%, 33.1%) | 1.13 (0.14, 9.07) | 1.12 (0.18, 7.09) | 1.2% (-19.4%, 21.9%) | 1.0000 | |
| Latin America | 18 | 1 (5.6%) (0.1%, 27.3%) | 19 | 0 (0.0%) | 3.34 (0.13, 87.52) | 3.16 (0.14, 72.84) | 5.4% (-8.5%, 19.3%) | 0.4865 | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point. The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Selected Adverse Events (PT)-PT Headache at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300 | mg Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|--------------------|-----------|------------------|-----------|--------------|---------------|--|---------------|-------------------------------|--------------|--|
| | | | | | | | | CMH or Logistic P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Age Subgroup | | | | | | | | | | |
| <40 years | 230 | 26 (11.3%) | 247 | 14 (5.7%) | 2.12 | 1.99 | 5.6% | 0.0314* | 0.3699 | |
| • | | (7.5%, | | (3.1%, 9.3%) | (1.08, 4.17) | (1.07, 3.72) | (0.6%, 10.6%) | | | |
| | | 16.1%) | | | | | | | | |
| >=40 years | 132 | 17 (12.9%) | 118 | 4 (3.4%) | 4.21 | 3.80 | 9.5% | 0.0105* | | |
| - | | (7.7%, | | (0.9%, 8.5%) | (1.38, 12.91) | (1.32, 10.97) | (2.9%, 16.1%) | | | |
| | | 19.8%) | | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (PT)-SOC Nervous system disorders at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300 | mg Q2W | | Dupilumab 300mg | Q2W vs. Abrocitinib 200mg | • | |
|---------------------------|-----------|---------------------------------|------------|-------------------------------|-----------------------|-----------------------|---------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 362 | 60 (16.6%) (12.9%, 20.8%) | 365 | 25 (6.8%) (4.5%, 9.9%) | 2.70 (1.65, 4.42) | 2.42 (1.55, 3.77) | 9.7% (5.1%, 14.4%) | <0.0001+* | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline | 216 | 36 (16.7%) | 220 | 18 (8.2%) | 2.24 | 2.04 | 8.5% | 0.0086* | 0.5451 |
| disease (IGA=3) | | (12.0%, 22.3%) | | (4.9%, 12.6%) | (1.23, 4.09) | (1.19, 3.47) | (2.3%, 14.6%) | | |
| Severe baseline disease | 146 | 24 (16.4%) | 145 | 7 (4.8%) | 3.88 | 3.41 | 11.6% | 0.0019* | |
| (IGA=4) | | (10.8%, 23.5%) | | (2.0%, 9.7%) | (1.61, 9.32) | (1.51, 7.65) | (4.7%, 18.6%) | | |
| Gender | | , | | | | | | | |
| Male | 193 | 30 (15.5%) (10.7%, 21.4%) | 204 | 10 (4.9%) (2.4%, 8.8%) | 3.57 (1.69, 7.52) | 3.17 (1.59, 6.31) | 10.6% (4.7%, 16.6%) | 0.0004* | 0.7065 |
| Female | 169 | 30 (17.8%) (12.3%, 24.4%) | 161 | 15 (9.3%) (5.3%, 14.9%) | 2.10 (1.08, 4.07) | 1.91 (1.07, 3.41) | 8.4% (1.1%, 15.7%) | 0.0361* | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 177 | 32 (18.1%) (12.7%, 24.6%) | 195 | 12 (6.2%) (3.2%, 10.5%) | 3.37 (1.67, 6.77) | 2.94 (1.56, 5.52) | 11.9% (5.3%, 18.5%) | 0.0004* | 0.6765 |
| Europe | 150 | 25 (16.7%) (11.1%, 23.6%) | 132 | 11 (8.3%) (4.2%, 14.4%) | 2.20 (1.04, 4.67) | 2.00 (1.02, 3.91) | 8.3% (0.7%, 15.9%) | 0.0483* | |
| Asia | 17 | 2 (11.8%) (1.5%, 36.4%) | 19 | 2 (10.5%) (1.3%, 33.1%) | 1.13 (0.14, 9.07) | 1.12 (0.18, 7.09) | 1.2% (-19.4%, 21.9%) | 1.0000 | |
| Latin America | 18 | 1 (5.6%) (0.1%, 27.3%) | 19 | 0 (0.0%) | 3.34 (0.13, 87.52) | 3.16 (0.14, 72.84) | 5.4% (-8.5%, 19.3%) | 0.4865 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Selected Adverse Events (PT)-SOC Nervous system disorders at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dupi | lumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|-----------|------------------|------------|------------|--------------|--|---------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 38 (16.5%) | 247 | 17 (6.9%) | 2.68 | 2.40 | 9.6% | 0.0014* | 0.9605 | | |
| • | | (12.0%, | | (4.1%, | (1.46, 4.89) | (1.39, 4.13) | (3.9%, 15.4%) | | | | |
| | | 22.0%) | | 10.8%) | | | | | | | |
| >=40 years | 132 | 22 (16.7%) | 118 | 8 (6.8%) | 2.75 | 2.46 | 9.9% | 0.0191* | | | |
| • | | (10.7%, | | (3.0%, | (1.17, 6.44) | (1.14, 5.31) | (2.1%, 17.7%) | | | | |
| | | 24.1%) | | 12.9%) | , | , | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (SOC)-SOC Skin and subcutaneous tissue disorders at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitin | ib 200mg QD Dup | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | |
|-----------------------------------|-----------|---------------------------------|-------------|--------------------------------|----------------------|--|-------------------------|----------------------------|-----------------------------|--|
| | | • | | - | | - | | CMH or Logistic 1 | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | · · · | | | | | | |
| Safety Set | 362 | 59 (16.3%) (12.6%, 20.5%) | 365 | 36 (9.9%) (7.0%, 13.4%) | 1.78 (1.14, 2.77) | 1.65 (1.12, 2.44) | 6.4% (1.6%, 11.3%) | 0.009 | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 216 | 35 (16.2%) (11.6%, 21.8%) | 220 | 20 (9.1%) (5.6%, 13.7%) | 1.93 (1.08, 3.47) | 1.78 (1.06, 2.99) | 7.1% (0.9%, 13.3%) | 0.0302* | 0.7526 | |
| Severe baseline disease (IGA=4) | 146 | 24 (16.4%) (10.8%, 23.5%) | 145 | 16 (11.0%) (6.4%, 17.3%) | 1.59 (0.80, 3.13) | 1.49 (0.83, 2.69) | 5.4% (-2.5%, 13.3%) | 0.2332 | | |
| Gender | | | | | | | | | | |
| Male | 193 | 29 (15.0%) (10.3%, 20.9%) | 204 | 17 (8.3%) (4.9%, 13.0%) | 1.95 (1.03, 3.67) | 1.80 (1.02, 3.17) | 6.7% (0.4%, 13.0%) | 0.0419* | 0.9193 | |
| Female | 169 | 30 (17.8%) (12.3%, 24.4%) | 161 | 19 (11.8%) (7.3%, 17.8%) | 1.61 (0.87, 3.00) | 1.50 (0.88, 2.56) | 6.0% (-1.7%, 13.6%) | 0.1631 | | |
| Region | | , | | 2.10.0) | | | | | | |
| US/Canada/Australia | 177 | 32 (18.1%) (12.7%, 24.6%) | 195 | 19 (9.7%) (6.0%, 14.8%) | 2.04 (1.11, 3.76) | 1.86 (1.09, 3.15) | 8.3% (1.3%, 15.4%) | 0.0234* | 0.4334 | |
| Europe | 150 | 24 (16.0%) (10.5%, 22.9%) | 132 | 13 (9.8%) (5.3%, 16.3%) | 1.74 (0.85, 3.58) | 1.62 (0.86, 3.06) | 6.2% (-1.6%, 13.9%) | 0.1576 | | |
| Asia | 17 | 3 (17.6%) (3.8%, 43.4%) | 19 | 3 (15.8%) (3.4%, 39.6%) | 1.14 (0.20, 6.60) | 1.12 (0.26, 4.81) | 1.9% (-22.6%, 26.3%) | 1.0000 | | |
| Latin America | 18 | 0 (0.0%) | 19 | 1 (5.3%) (0.1%, 26.0%) | 0.33 (0.01, 8.73) | 0.35 (0.02, 8.09) | -4.9% (-18.5%, 8.7%) | 1.0000 | | |

Notes

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Selected Adverse Events (SOC)-SOC Skin and subcutaneous tissue disorders at 12 weeks - Safety Set JADE DARE (PF-04965842) - 2023 datacut

| | Abrocitini | ib 200mg QD Dupi | ilumab 300n | ng Q2W | | Dupilumab 300mg Q2W vs. Abrocitinib 200mg QD | | | | | |
|--------------------|------------|------------------|-------------|------------|--------------|--|----------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Age Subgroup | | | | | | | | | | | |
| <40 years | 230 | 39 (17.0%) | 247 | 23 (9.3%) | 1.99 | 1.82 | 7.6% | 0.0143* | 0.5064 | | |
| - | | (12.3%, | | (6.0%, | (1.15, 3.45) | (1.12, 2.95) | (1.6%, 13.7%) | | | | |
| | | 22.4%) | | 13.6%) | | | | | | | |
| >=40 years | 132 | 20 (15.2%) | 118 | 13 (11.0%) | 1.44 | 1.38 | 4.1% | 0.3560 | | | |
| • | | (9.5%, | | (6.0%, | (0.68, 3.05) | (0.72, 2.64) | (-4.2%, 12.5%) | | | | |
| | | 22.4%) | | 18.1%) | | | | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

MMRM Analysen JADE DARE

Descriptive Summary of Total EASI Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | Dupilumab 300mg Q2W (N=365) | | | | | | | | |
|------------------------------|----------|---------------------------------|-------|--------|--------------------------------|-------|-------|-------------|-------|--------|------|-------|-------|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| 01 15 | D 11 | 262 (100.0) | 20.1 | 24.5 | 11.5 | 10.4 | 22.6 | 265 (100.0) | 20.1 | 24.5 | 11.0 | 10.0 | 22.5 |
| Observed Data | Baseline | 362 (100.0) | 28.1 | | 11.5 | 19.4 | 33.6 | 365 (100.0) | 28.1 | | 11.9 | 19.2 | 33.5 |
| | Week 2 | 350 (96.7) | 12.7 | 10.0 | 10.6 | 4.8 | 17.6 | 350 (95.9) | | | 11.3 | 7.8 | 19.7 |
| | Week 4 | 342 (94.5) | 7.6 | 5.2 | 8.8 | 1.8 | 10.0 | 351 (96.2) | | | 9.9 | 3.8 | 15.4 |
| | Week 8 | 336 (92.8) | 5.1 | 2.8 | 6.6 | 0.8 | 6.6 | 348 (95.3) | | | 8.2 | 2.4 | 10.6 |
| | Week 12 | 329 (90.9) | 4.1 | 2.4 | 5.8 | 0.7 | 5.6 | 341 (93.4) | 6.3 | 4.0 | 7.3 | 1.6 | 8.2 |
| | Week 16 | 324 (89.5) | 3.8 | 1.6 | 6.1 | 0.4 | 4.8 | 336 (92.1) | 5.3 | 3.4 | 6.7 | 1.2 | 6.7 |
| | Week 20 | 320 (88.4) | 3.5 | 1.4 | 5.9 | 0.3 | 4.3 | 333 (91.2) | 4.7 | 2.4 | 6.6 | 0.8 | 6.0 |
| | Week 26 | 301 (83.1) | 3.3 | 1.5 | 5.2 | 0.0 | 4.3 | 324 (88.8) | 4.2 | 2.4 | 5.8 | 0.6 | 5.6 |
| | | | | | | | | | | | | | |
| Change from Baseline | Week 2 | 350 (96.7) | -15.5 | -15.2 | 9.4 | -20.8 | -8.7 | 350 (95.9) | -12.8 | -11.0 | 10.3 | -18.5 | -5.5 |
| | Week 4 | 342 (94.5) | -20.7 | -19.3 | 10.2 | -25.9 | -15.1 | 351 (96.2) | -17.2 | -15.5 | 10.7 | -22.4 | -10.9 |
| | Week 8 | 336 (92.8) | -23.1 | -21.0 | 10.8 | -29.2 | -16.8 | 348 (95.3) | -20.2 | -18.4 | 10.7 | -24.8 | -13.6 |
| | Week 12 | 329 (90.9) | -24.0 | -21.4 | 11.0 | -29.3 | -17.3 | 341 (93.4) | -21.9 | -19.2 | 11.1 | -26.9 | -15.1 |
| | Week 16 | 324 (89.5) | -24.4 | -21.7 | 10.7 | -29.6 | -17.7 | 336 (92.1) | -22.9 | -20.8 | 11.5 | -27.9 | -15.7 |
| | Week 20 | 320 (88.4) | -24.8 | -21.8 | 10.9 | -30.0 | -17.8 | 333 (91.2) | -23.3 | -20.8 | 11.2 | -28.0 | -16.4 |
| | Week 26 | 301 (83.1) | -24.7 | -22.5 | 10.6 | -29.2 | -17.8 | 324 (88.8) | -23.8 | -21.8 | 11.3 | -28.5 | -16.8 |
| | | | | | | | | | | | | | |
| Percent Change from Baseline | Week 2 | 350 (96.7) | -56.3 | -60.1 | 28.0 | -80.6 | -33.9 | 350 (95.9) | -45.7 | -44.2 | 29.4 | -71.1 | -22.8 |
| | Week 4 | 342 (94.5) | -74.0 | -80.3 | 24.3 | -92.4 | -64.2 | 351 (96.2) | -61.9 | -65.3 | 26.6 | -84.2 | -44.8 |
| | Week 8 | 336 (92.8) | -81.9 | -89.5 | 22.6 | -96.8 | -74.9 | 348 (95.3) | -72.6 | -79.2 | 23.7 | -90.8 | -61.1 |
| | Week 12 | 329 (90.9) | -85.2 | -91.2 | 17.9 | -97.6 | -79.3 | 341 (93.4) | -78.0 | -83.9 | 20.3 | -94.1 | -67.5 |
| | Week 16 | 324 (89.5) | -87.1 | -93.2 | 17.6 | -98.3 | -82.8 | 336 (92.1) | -81.1 | -88.4 | 19.9 | -95.5 | -72.9 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

EASI = eczema area and severity index; N = N = number of subjects in the analysis set; N = N = number of subjects in the analysis set with observed data at the specified visit; N = N = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 15SEP2021 (01:53)

Descriptive Summary of Total EASI Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | Dupilumab 300mg Q2W (N=365) | | | | V | | | | |
|------------------------------|---------|---------------------------------|-------|-------|--------------------------------|--------|--------|------------|-------|-------|------|-------|-------|
| | | n (%) Mean Median SD Q1 Q3 | | | n (%) | Mean | Median | SD | Q1 | Q3 | | | |
| Percent Change from Baseline | Week 20 | 320 (88.4) | -88.3 | -94.6 | 16.8 | -99.0 | -84.0 | 333 (91.2) | -83.7 | -89.9 | 18.6 | -96.8 | -76.6 |
| | Week 26 | 301 (83.1) | -88.4 | -94.2 | 17.0 | -100.0 | -83.5 | 324 (88.8) | -85.2 | -91.1 | 17.8 | -97.3 | -80.1 |
| | | | | | | | | | | | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

EASI = eczema area and severity index; N = N = number of subjects in the analysis set; N = N = number of subjects in the analysis set with observed data at the specified visit; N = N = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 15SEP2021 (01:53)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|-------------------------|------------------------|
| | | | |
| | N | 362 | 365 |
| | | | |
| Week 2 | N1 | 351 | 353 |
| | N2 (%) | 350 (99.7) | 350 (99.2) |
| | LSM (SE) | -56.18 (1.53) | -45.72 (1.53) |
| | 95% CI | (-59.19, -53.18) | (-48.72, -42.72) |
| | · | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -10.46 | |
| | 95% CI | (-14.71, -6.22) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.37 | |
| | 95% CI of Hedges'g | (-0.52, -0.22) | |
| | | | |
| Week 4 | N1 | 350 | 352 |
| | N2 (%) | 342 (97.7) | 351 (99.7) |
| | LSM (SE) | -73.76 (1.36) | -61.76 (1.36) |
| | ESM (SE) | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; EASI = eczema area and severity index; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | -11.99 | |
| | 95% CI | (-15.77, -8.22) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.47 | |
| | 95% CI of Hedges'g | (-0.62, -0.32) | |
| | | | |
| Week 8 | N1 | 343 | 351 |
| | N2 (%) | 336 (98.0) | 348 (99.1) |
| | LSM (SE) | -81.70 (1.25) | -72.46 (1.24) |
| | 95% CI | (-84.16, -79.25) | (-74.89, -70.02) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -9.25 | |
| | 95% CI | (-12.70, -5.79) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.40 | |
| | 95% CI of Hedges'g | (-0.55, -0.25) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; EASI = eczema area and severity index; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|---------------------------|-------------------------|------------------------|
| Week 12 | N1 | 332 | 343 |
| | N2 (%) | 329 (99.1) | 341 (99.4) |
| | LSM (SE) | -84.40 (1.05) | -78.05 (1.04) |
| | 95% CI | (-86.46, -82.34) | (-80.09, -76.01) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -6.35 | |
| | 95% CI | (-9.25, -3.44) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.33 | |
| | 95% CI of Hedges'g | (-0.48, -0.18) | |
| | | | |
| Week 16 | N1 | 329 | 341 |
| | N2 (%) | 324 (98.5) | 336 (98.5) |
| | LSM (SE) | -86.39 (1.03) | -81.14 (1.02) |
| | 95% CI | (-88.41, -84.37) | (-83.14, -79.15) |
| | Aharasidin ib Danilhansah | | |
| | Abrocitinib - Dupilumab | 5.24 | |
| | LSM | -5.24 | |
| | 95% CI | (-8.09, -2.40) | |
| | Two-sided P-value | 0.0003 | |
| | Hedges'g | -0.28 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; EASI = eczema area and severity index; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W | |
|---------|-------------------------|-------------------------|------------------------|--|
| | 95% CI of Hedges'g | (-0.44, -0.13) | | |
| | | | | |
| Week 20 | N1 | 323 | 334 | |
| | N2 (%) | 320 (99.1) | 333 (99.7) | |
| | LSM (SE) | -87.40 (0.98) | -83.42 (0.97) | |
| | 95% CI | (-89.33, -85.47) | (-85.32, -81.52) | |
| | | | | |
| | Abrocitinib - Dupilumab | | | |
| | LSM | -3.98 | | |
| | 95% CI | (-6.69, -1.27) | | |
| | Two-sided P-value | 0.0040 | | |
| | Hedges'g | -0.23 | | |
| | 95% CI of Hedges'g | (-0.38, -0.07) | | |
| | | | | |
| Week 26 | N1 | 301 | 324 | |
| | N2 (%) | 301 (100.0) | 324 (100.0) | |
| | LSM (SE) | -87.03 (1.00) | -84.96 (0.98) | |
| | 95% CI | (-89.00, -85.05) | (-86.89, -83.02) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; EASI = eczema area and severity index; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 06SEP2021 (02:56)

| | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|-------------------------|-------------------------|------------------------|
| Abrocitinib - Dupilumab | | |
| LSM | -2.07 | |
| 95% CI | (-4.83, 0.69) | |
| Two-sided P-value | 0.1408 | |
| Hedges'g | -0.12 | |
| 95% CI of Hedges'g | (-0.28, 0.04) | |
| | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

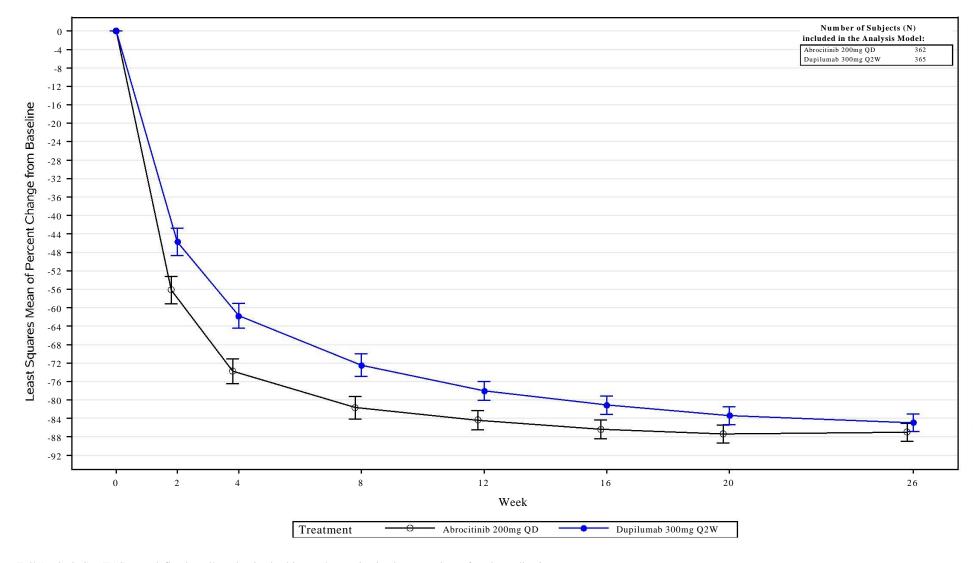
Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; EASI = eczema area and severity index; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 06SEP2021 (02:56)



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge.

Vertical line represented 95% confidence interval.

EASI = eczema area and severity index; OD = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adea Table Generation: 05SEP2021 (22:29)

Descriptive Summary of Peak Pruritus Numerical Rating Scale (PP-NRS), Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD)

(Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | |] | Dupiluı | mab 300m (N=365) | ng Q2V | V | | | |
|------------------------------|----------|---------------------------------|-------|--------|------|-------|---------|---------------------|--------|--------|------|-------|-------|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| Observed Data | Baseline | 362 (100.0) | 7.4 | 8.0 | 1.6 | 7.0 | 8.0 | 365 (100.0) | 7.4 | 7.0 | 1.6 | 6.0 | 9.0 |
| | Week 2 | 354 (97.8) | 3.8 | 4.0 | 2.3 | 2.0 | 6.0 | 355 (97.3) | | 5.0 | 2.2 | 3.0 | 7.0 |
| | Week 4 | 349 (96.4) | 3.2 | 3.0 | 2.3 | 1.0 | 5.0 | 351 (96.2) | 4.2 | 4.0 | 2.2 | 2.0 | 6.0 |
| | Week 8 | 345 (95.3) | 2.8 | 2.0 | 2.3 | 1.0 | 4.0 | 351 (96.2) | 3.6 | 3.0 | 2.4 | 2.0 | 5.0 |
| | Week 12 | 333 (92.0) | 2.6 | 2.0 | 2.3 | 1.0 | 4.0 | 347 (95.1) | 3.1 | 3.0 | 2.2 | 1.0 | 5.0 |
| | Week 16 | 331 (91.4) | 2.6 | 2.0 | 2.3 | 1.0 | 4.0 | 342 (93.7) | 3.0 | 3.0 | 2.2 | 1.0 | 4.0 |
| | Week 20 | 326 (90.1) | 2.5 | 2.0 | 2.2 | 1.0 | 4.0 | 338 (92.6) | 2.8 | 3.0 | 2.1 | 1.0 | 4.0 |
| | Week 26 | 313 (86.5) | 2.3 | 2.0 | 2.2 | 1.0 | 4.0 | 328 (89.9) | 2.7 | 2.0 | 2.1 | 1.0 | 4.0 |
| | | | | | | | | | | | | | |
| Change from Baseline | Week 2 | 354 (97.8) | -3.6 | -3.0 | 2.3 | -5.0 | -2.0 | 355 (97.3) | -2.3 | -2.0 | 2.1 | -4.0 | -1.0 |
| | Week 4 | 349 (96.4) | -4.3 | -4.0 | 2.5 | -6.0 | -2.0 | 351 (96.2) | -3.2 | -3.0 | 2.3 | -5.0 | -2.0 |
| | Week 8 | 345 (95.3) | -4.7 | -5.0 | 2.5 | -6.0 | -3.0 | 351 (96.2) | -3.8 | -4.0 | 2.5 | -5.0 | -2.0 |
| | Week 12 | 333 (92.0) | -4.8 | -5.0 | 2.5 | -7.0 | -3.0 | 347 (95.1) | -4.2 | -4.0 | 2.3 | -6.0 | -3.0 |
| | Week 16 | 331 (91.4) | -4.8 | -5.0 | 2.6 | -7.0 | -3.0 | 342 (93.7) | -4.3 | -4.0 | 2.3 | -6.0 | -3.0 |
| | Week 20 | 326 (90.1) | -4.9 | -5.0 | 2.5 | -7.0 | -3.0 | 338 (92.6) | -4.5 | -5.0 | 2.3 | -6.0 | -3.0 |
| | Week 26 | 313 (86.5) | -5.1 | -5.0 | 2.4 | -7.0 | -4.0 | 328 (89.9) | -4.6 | -5.0 | 2.4 | -6.0 | -3.0 |
| | | | | | | | | | | | | | |
| Percent Change from Baseline | Week 2 | 354 (97.8) | -48.2 | -50.0 | 30.0 | -71.4 | -25.0 | 355 (97.3) | -31.5 | -28.6 | 27.6 | -50.0 | -12.5 |
| | Week 4 | 349 (96.4) | -56.2 | -60.0 | 32.0 | -83.3 | -33.3 | 351 (96.2) | -42.9 | -42.9 | 29.0 | -66.7 | -22.2 |
| | Week 8 | 345 (95.3) | -62.0 | -66.7 | 31.8 | -87.5 | -37.5 | 351 (96.2) | -50.9 | -55.6 | 31.2 | -75.0 | -30.0 |
| | Week 12 | 333 (92.0) | -63.9 | -71.4 | 33.0 | -87.5 | -44.4 | 347 (95.1) | -57.1 | -60.0 | 29.0 | -80.0 | -37.5 |
| | Week 16 | 331 (91.4) | -64.0 | -71.4 | 33.9 | -87.5 | -50.0 | 342 (93.7) | -58.7 | -62.5 | 29.2 | -83.3 | -40.0 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

OD = observed data

N = number of subjects in the analysis set n = number of subjects in the analysis set with observed data at the specified visit

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 30AUG2021 (06:20)

Descriptive Summary of Peak Pruritus Numerical Rating Scale (PP-NRS), Change from Baseline and Percent Change from Baseline up to Week 26 (FAS,

(Protocol B7451050)

| | | Abrocitinib 200mg QD Dupilumab 300mg Q2W (N=362) (N=365) | | | | V | | | | | | | |
|------------------------------|---------|--|----------------------------|-------|------|-------|-------|------------|-------|-------|------|-------|-------|
| | | n (%) | n (%) Mean Median SD Q1 Q3 | | | n (%) | Mean | Median | SD | Q1 | Q3 | | |
| Percent Change from Baseline | Week 20 | 326 (90.1) | -65.6 | -71.4 | 31.2 | -88.9 | -44.4 | 338 (92.6) | -60.8 | -64.6 | 28.7 | -83.3 | -44.4 |
| | Week 26 | 313 (86.5) | -68.2 | -75.0 | 31.1 | -88.9 | -50.0 | 328 (89.9) | -62.8 | -71.4 | 29.0 | -85.7 | -44.4 |
| | | | | | | | | | | | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

OD = observed data

N = number of subjects in the analysis set n = number of subjects in the analysis set with observed data at the specified visit

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 30AUG2021 (06:20)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|----------------------------|----------------------------|
| | | | |
| | N | 362 | 365 |
| | | | |
| Week 2 | N1 | 354 | 355 |
| | N2 (%) | 354 (100.0) | 355 (100.0) |
| | LSM (SE) | -3.60 (0.11) | -2.37 (0.11) |
| | 95% CI | (-3.82, -3.38) | (-2.59, -2.15) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.23 | |
| | 95% CI | (-1.54, -0.92) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.58 | |
| | 95% CI of Hedges'g | (-0.73, -0.43) | |
| | | | |
| Week 4 | N1 | 350 | 352 |
| Week 4 | | | |
| Week 4 | N2 (%) | 349 (99.7) | 351 (99.7) |
| Week 4 | N2 (%) LSM (SE) | 349 (99.7) -4.24 (0.12) | 351 (99.7) -3.23 (0.12) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.01 | |
| | 95% CI | (-1.33, -0.68) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.46 | |
| | 95% CI of Hedges'g | (-0.61, -0.31) | |
| | | | |
| Week 8 | N1 | 345 | 351 |
| | N2 (%) | 345 (100.0) | 351 (100.0) |
| | LSM (SE) | -4.63 (0.12) | -3.81 (0.12) |
| | 95% CI | (-4.87, -4.39) | (-4.05, -3.58 |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.81 | |
| | 95% CI | (-1.15, -0.48) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.36 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| Week 12 | N1 | 334 | 347 |
| | N2 (%) | 333 (99.7) | 347 (100.0) |
| | LSM (SE) | -4.74 (0.12) | -4.24 (0.12) |
| | 95% CI | (-4.97, -4.51) | (-4.47, -4.01) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.50 | |
| | 95% CI | (-0.82, -0.17) | |
| | Two-sided P-value | 0.0029 | |
| | Hedges'g | -0.23 | |
| | 95% CI of Hedges'g | (-0.38, -0.08) | |
| | | | |
| Week 16 | N1 | 331 | 343 |
| | N2 (%) | 331 (100.0) | 342 (99.7) |
| | LSM (SE) | -4.76 (0.12) | -4.33 (0.12) |
| | 95% CI | (-5.00, -4.53) | (-4.56, -4.10) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.43 | |
| | 95% CI | (-0.76, -0.10) | |
| | Two-sided P-value | 0.0109 | |
| | Hedges'g | -0.20 | |
| | 95% CI of Hedges'g | (-0.35, -0.05) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | | | |
| Week 20 | N1 | 326 | 338 |
| | N2 (%) | 326 (100.0) | 338 (100.0) |
| | LSM (SE) | -4.83 (0.12) | -4.50 (0.12) |
| | 95% CI | (-5.05, -4.60) | (-4.73, -4.27) |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.32 | |
| | 95% CI | (-0.65, 0.00) | |
| | Two-sided P-value | 0.0497 | |
| | Hedges'g | -0.15 | |
| | 95% CI of Hedges'g | (-0.30, 0.00) | |
| | | | |
| Week 26 | N1 | 313 | 328 |
| | N2 (%) | 313 (100.0) | 328 (100.0) |
| | LSM (SE) | -4.92 (0.12) | -4.65 (0.12) |
| | 95% CI | (-5.15, -4.68) | (-4.88, -4.41) |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.27 | |
| | 95% CI | (-0.60, 0.06) | |
| | Two-sided P-value | 0.1068 | |
| | Hedges'g | -0.13 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 06SEP2021 (02:56)

| | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------------------|-------------------------|------------------------|
| 95% CI of Hedges'g | (-0.28, 0.03) | |
| | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

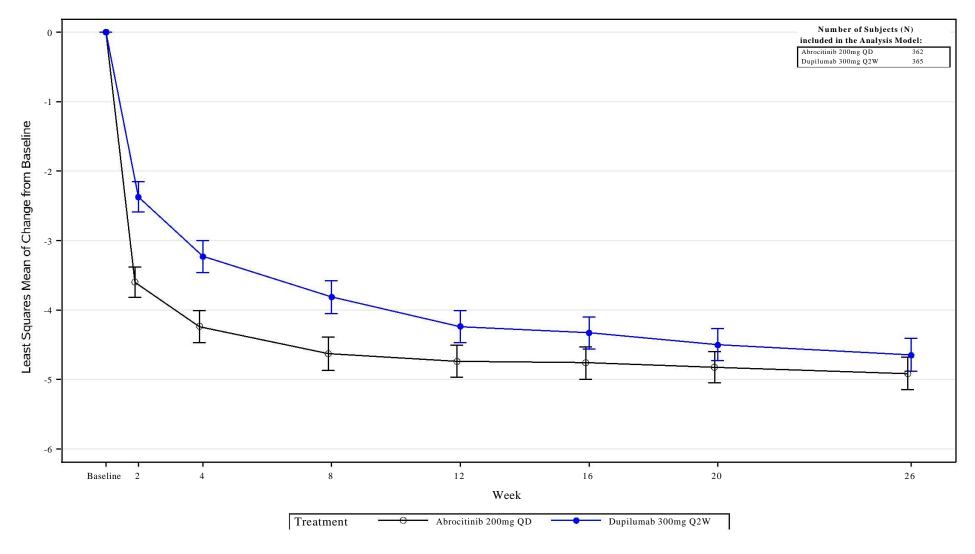
Data after dropout or use of rescue therapy was censored.

CI = confidence interval; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 06SEP2021 (02:56)



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge.

Vertical line represented 95% confidence interval.

OD = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adnr Table Generation: 05SEP2021 (22:29)

Descriptive Summary of DLQI, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | | Dupilumab 300mg Q2W (N=365) | | | | | | | |
|------------------------------|----------|---------------------------------|-------|--------|------|-----------------------------|-------|------------|-------|--------|------|-------|-------|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| Observed Data | Baseline | 361 (99.7) | 14.0 | 14.0 | 6.8 | 9.0 | 19.0 | 363 (99.5) | 14.2 | 14.0 | 6.3 | 9.0 | 19.0 |
| | Week 2 | 347 (95.9) | 5.4 | 4.0 | 5.1 | 2.0 | 8.0 | 351 (96.2) | | | 5.1 | 3.0 | 11.0 |
| | Week 12 | 329 (90.9) | 3.5 | 2.0 | 4.0 | 1.0 | 4.0 | 341 (93.4) | 4.3 | 2.0 | 4.7 | 1.0 | 6.0 |
| | Week 16 | 325 (89.8) | 3.2 | 2.0 | 3.9 | 1.0 | 4.0 | 338 (92.6) | 4.0 | 2.5 | 4.3 | 1.0 | 6.0 |
| | Week 20 | 320 (88.4) | 3.3 | 2.0 | 4.0 | 1.0 | 5.0 | 333 (91.2) | 3.8 | 2.0 | 4.1 | 1.0 | 5.0 |
| | Week 26 | 303 (83.7) | 3.6 | 2.0 | 4.7 | 1.0 | 5.0 | 325 (89.0) | 4.0 | 2.0 | 4.5 | 1.0 | 6.0 |
| | | | | | | | | | | | | | |
| Change from Baseline | Week 2 | 346 (95.6) | -8.6 | -8.0 | 6.1 | -12.0 | -4.0 | 349 (95.6) | -6.6 | -6.0 | 5.9 | -10.0 | -3.0 |
| | Week 12 | 328 (90.6) | -10.7 | -10.0 | 6.7 | -15.0 | -6.0 | 339 (92.9) | -9.6 | -9.0 | 6.1 | -13.0 | -6.0 |
| | Week 16 | 324 (89.5) | -10.8 | -10.0 | 6.7 | -15.0 | -6.0 | 336 (92.1) | -9.9 | -9.0 | 6.0 | -13.0 | -6.0 |
| | Week 20 | 319 (88.1) | -10.9 | -10.0 | 6.7 | -15.0 | -6.0 | 331 (90.7) | -10.1 | -9.0 | 6.1 | -14.0 | -6.0 |
| | Week 26 | 302 (83.4) | -10.5 | -11.0 | 6.6 | -15.0 | -5.0 | 323 (88.5) | -10.0 | -10.0 | 6.2 | -14.0 | -6.0 |
| | | | | | | | | | | | | | |
| Percent Change from Baseline | Week 2 | 345 (95.3) | -58.3 | -66.7 | 40.4 | -84.2 | -42.1 | 349 (95.6) | -42.6 | -50.0 | 38.6 | -70.6 | -25.0 |
| | Week 12 | 327 (90.3) | -70.6 | -83.3 | 63.5 | -93.8 | -62.5 | 339 (92.9) | -66.6 | -77.8 | 34.7 | -90.0 | -50.0 |
| | Week 16 | 323 (89.2) | -73.1 | -85.0 | 40.1 | -95.0 | -61.9 | 336 (92.1) | -68.6 | -77.4 | 31.9 | -90.0 | -56.4 |
| | Week 20 | 319 (88.1) | -73.2 | -85.0 | 38.7 | -95.2 | -66.7 | 331 (90.7) | -69.0 | -80.0 | 38.3 | -90.9 | -60.0 |
| | Week 26 | 301 (83.1) | -72.7 | -84.6 | 44.1 | -95.8 | -66.7 | 323 (88.5) | -68.5 | -80.0 | 37.6 | -90.9 | -56.5 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

DLQI = dermatology life quality index; OD = observed data. Data after dropout or use of rescue therapy was censored.

N = number of subjects in the analysis set.

n = number of subjects in the analysis set:

n = number of subjects in the analysis set with observed data at the specified visit.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 12SEP2021 (23:49)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | | | |
| | N | 361 | 363 |
| | | | |
| Week 2 | N1 | 349 | 351 |
| | N2 (%) | 346 (99.1) | 349 (99.4) |
| | LSM (SE) | -8.63 (0.24) | -6.67 (0.24) |
| | 95% CI | (-9.11, -8.16) | (-7.15, -6.20) |
| | <u> </u> | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.96 | |
| | 95% CI | (-2.64, -1.29) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.43 | |
| | 95% CI of Hedges'g | (-0.58, -0.28) | |
| | | | |
| Week 12 | N1 | 331 | 341 |
| | N2 (%) | 328 (99.1) | 339 (99.4) |
| | LSM (SE) | -10.69 (0.22) | -9.69 (0.22) |
| | 95% CI | (-11.13, -10.25) | (-10.12, -9.25 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; DLQI = Dermatology Life Quality Index; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 12SEP2021 (23:19)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.00 | |
| | 95% CI | (-1.62, -0.38) | |
| | Two-sided P-value | 0.0016 | |
| | Hedges'g | -0.25 | |
| | 95% CI of Hedges'g | (-0.40, -0.09) | |
| | <u> </u> | | |
| Week 16 | N1 | 328 | 339 |
| | N2 (%) | 324 (98.8) | 336 (99.1) |
| | LSM (SE) | -10.80 (0.22) | -10.04 (0.21) |
| | 95% CI | (-11.22, -10.38) | (-10.46, -9.62) |
| | <u> </u> | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.76 | |
| | 95% CI | (-1.35, -0.16) | |
| | Two-sided P-value | 0.0126 | |
| | Hedges'g | -0.19 | |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; DLQI = Dermatology Life Quality Index; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 12SEP2021 (23:19)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| Week 20 | N1 | 322 | 333 |
| | N2 (%) | 319 (99.1) | 331 (99.4) |
| | LSM (SE) | -10.77 (0.22) | -10.13 (0.21) |
| | 95% CI | (-11.19, -10.34) | (-10.55, -9.71) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.64 | |
| | 95% CI | (-1.23, -0.04) | |
| | Two-sided P-value | 0.0376 | |
| | Hedges'g | -0.16 | |
| | 95% CI of Hedges'g | (-0.32, -0.01) | |
| | | | |
| Week 26 | N1 | 302 | 323 |
| | N2 (%) | 302 (100.0) | 323 (100.0) |
| | LSM (SE) | -10.34 (0.24) | -10.04 (0.24) |
| | 95% CI | (-10.82, -9.86) | (-10.51, -9.57) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.30 | |
| | 95% CI | (-0.97, 0.37) | |
| | Two-sided P-value | 0.3814 | |
| | Hedges'g | -0.07 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; DLQI = Dermatology Life Quality Index; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 12SEP2021 (23:19)

| | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------------------|-------------------------|------------------------|
| 95% CI of Hedges'g | (-0.23, 0.09) | |
| | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; DLQI = Dermatology Life Quality Index; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

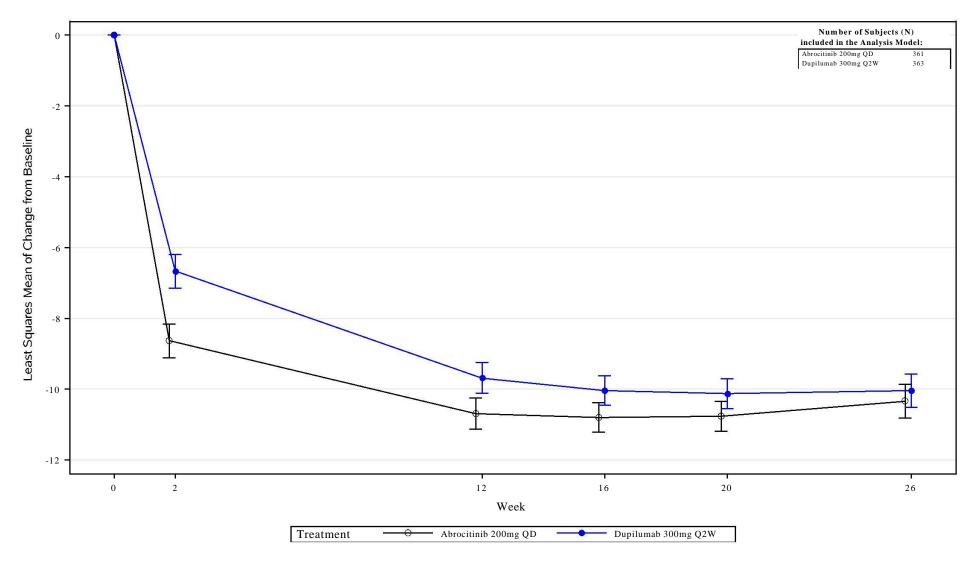
OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: adli Table Generation: 12SEP2021 (23:19)

Parameter: DLQI Score



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

Vertical line represented 95% confidence interval.

DLQI = dermatology life quality index; OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

PFIZER CONFIDENTIAL SDTM Creation: 22JUN2021 (01:49) Source Data: adli Table Generation: 14SEP2021 (23:40)

Descriptive Summary of EQ-5D VAS Score, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | | |] | Dupilumab 300mg Q2W (N=365) | | | | | |
|------------------------------|----------|---------------------------------|------|--------|------|------|------|--------------------------------|------|--------|------|------|------|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| Observed Data | Baseline | 362 (100.0) | 68.4 | 72.0 | 19.5 | 59.0 | 80.0 | 364 (99.7) | 67.7 | 70.0 | 18.3 | 58.5 | 80.0 |
| | Week 12 | 329 (90.9) | _ | | 15.3 | 75.0 | 90.0 | 341 (93.4) | | | 13.4 | 70.0 | |
| | Week 16 | 323 (89.2) | 80.8 | 83.0 | 14.9 | 75.0 | 90.0 | 337 (92.3) | 78.5 | 81.0 | 15.9 | 70.0 | 90.0 |
| | Week 26 | 303 (83.7) | 81.8 | 85.0 | 15.3 | 75.0 | 92.0 | 324 (88.8) | 82.5 | 85.0 | 13.5 | 76.0 | 91.0 |
| | | | | | | | | | | | | | |
| Change from Baseline | Week 12 | 329 (90.9) | 12.1 | 10.0 | 20.3 | 0.0 | 20.0 | 340 (93.2) | 11.7 | 10.0 | 18.5 | 0.0 | 20.0 |
| | Week 16 | 323 (89.2) | 12.2 | 9.0 | 20.2 | 0.0 | 22.0 | 336 (92.1) | 10.6 | 10.0 | 18.2 | 0.0 | 20.0 |
| | Week 26 | 303 (83.7) | 13.4 | 10.0 | 20.9 | 1.0 | 24.0 | 323 (88.5) | 14.6 | 12.0 | 18.0 | 3.0 | 25.0 |
| | | | | | | | | | | | | | |
| Percent Change from Baseline | Week 12 | 328 (90.6) | 32.0 | 12.6 | 73.1 | 0.0 | 34.4 | 340 (93.2) | 32.1 | 13.0 | 82.8 | 0.0 | 33.3 |
| | Week 16 | 322 (89.0) | 31.4 | 11.1 | 68.8 | 0.0 | 36.4 | 336 (92.1) | 26.7 | 12.5 | 64.7 | 0.0 | 33.3 |
| | Week 26 | 302 (83.4) | 33.8 | 13.9 | 73.1 | 1.3 | 40.0 | 323 (88.5) | 34.7 | 15.4 | 69.5 | 3.7 | 41.7 |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

EQ-5D VAS Score = EuroQuol Quality of Life 5-Dimension Visual Analog Scale Score; OD = observed data. Data after dropout or use of rescue therapy was censored. N = number of subjects in the analysis set.

n = number of subjects in the analysis set with observed data at the specified visit.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: ade5 Table Generation: 12SEP2021 (23:49)

Least Squares Mean of Change from Baseline in EQ-5D VAS Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | | | |
| | N | 362 | 364 |
| | | | |
| Week 12 | N1 | 331 | 342 |
| | N2 (%) | 329 (99.4) | 340 (99.4) |
| | LSM (SE) | 12.37 (0.74) | 11.55 (0.73) |
| | 95% CI | (10.92, 13.82) | (10.12, 12.98) |
| | · | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | 0.82 | |
| | 95% CI | (-1.22, 2.86) | |
| | Two-sided P-value | 0.4310 | |
| | Hedges'g | 0.06 | |
| | 95% CI of Hedges'g | (-0.09, 0.21) | |
| | · | | |
| Week 16 | N1 | 328 | 338 |
| | N2 (%) | 323 (98.5) | 336 (99.4) |
| | LSM (SE) | 12.57 (0.79) | 10.47 (0.78) |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; EQ-5D VAS Score = EuroQuol Quality of Life 5-Dimension Visual Analog Scale Score; SE = Standard Error; LSM = least squares mean;

N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: ade5 Table Generation: 12SEP2021 (23:19)

Least Squares Mean of Change from Baseline in EQ-5D VAS Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | 2.09 | |
| | 95% CI | (-0.08, 4.27) | |
| | Two-sided P-value | 0.0592 | |
| | Hedges'g | 0.15 | |
| | 95% CI of Hedges'g | (-0.01, 0.30) | |
| | | | |
| Week 26 | N1 | 303 | 323 |
| | N2 (%) | 303 (100.0) | 323 (100.0) |
| | LSM (SE) | 13.48 (0.76) | 14.30 (0.75) |
| | 95% CI | (11.98, 14.99) | (12.84, 15.76 |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.82 | |
| | 95% CI | (-2.91, 1.28) | |
| | Two-sided P-value | 0.4450 | |
| | Hedges'g | -0.06 | |
| | 95% CI of Hedges'g | (-0.22, 0.10) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; EQ-5D VAS Score = EuroQuol Quality of Life 5-Dimension Visual Analog Scale Score; SE = Standard Error; LSM = least squares mean;

N = number of subjects included in the analysis model;

N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

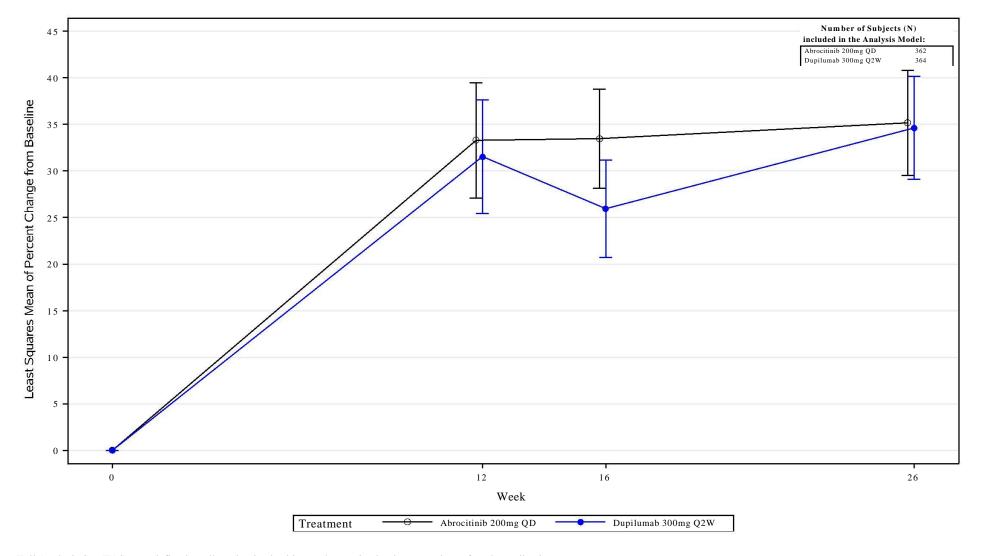
OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: ade5 Table Generation: 12SEP2021 (23:19)

Parameter: EQ-5D VAS Score



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Vertical line represented 95% confidence interval.

EQ-5D VAS Score = EuroQuol Quality of Life 5-Dimension Visual Analog Scale Score; OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction,

baseline disease severity, baseline value and an unstructured covariance matrix.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (01:50) Source Data: ade5 Table Generation: 13SEP2021 (01:30)

Descriptive Summary of POEM, Absolute Values, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

Parameter: Total Score

| | | | (N=362) | | | | | Dupilumab 300mg Q2W (N=365) | | | | |
|--------|--|---|--|---|---|---|---|---|---|---|---|---|
| | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| seline | 362 (100.0) | 20.4 | 21.0 | 5.8 | 17.0 | 25.0 | 365 (100.0) | 20.9 | 21.0 | 5 3 | 18.0 | 25.0 |
| eek 12 | ` ` | | | 6.1 | 2.0 | 9.0 | 341 (93.4) | | | 6.2 | 3.0 | 11.0 |
| eek 16 | 323 (89.2) | 6.2 | 4.0 | 5.8 | 2.0 | 9.0 | 337 (92.3) | 7.8 | 7.0 | 5.7 | 4.0 | 11.0 |
| eek 26 | 303 (83.7) | 6.5 | 5.0 | 6.4 | 1.0 | 10.0 | 322 (88.2) | 7.2 | 6.0 | 5.8 | 3.0 | 10.0 |
| | | | | | | | | | | | | |
| eek 12 | 327 (90.3) | -14.0 | -15.0 | 7.0 | -19.0 | -9.0 | 341 (93.4) | -12.7 | -13.0 | 6.9 | -18.0 | -8.0 |
| eek 16 | 323 (89.2) | -14.1 | -15.0 | 6.8 | -19.0 | -9.0 | 337 (92.3) | -12.9 | -13.0 | 6.8 | -18.0 | -8.0 |
| eek 26 | 303 (83.7) | -13.7 | -14.0 | 7.3 | -19.0 | -9.0 | 322 (88.2) | -13.5 | -14.0 | 6.6 | -18.0 | -9.0 |
| | | | | | | | | | | | | |
| eek 12 | 324 (89.5) | -66.5 | -77.8 | 70.5 | -91.7 | -54.5 | 341 (93.4) | -59.9 | -68.4 | 33.5 | -82.4 | -44.4 |
| eek 16 | 320 (88.4) | -67.0 | -76.9 | 60.8 | -91.1 | -53.6 | 337 (92.3) | -61.1 | -65.2 | 27.9 | -81.0 | -46.4 |
| eek 26 | 300 (82.9) | -65.7 | -76.5 | 58.4 | -91.7 | -52.6 | 322 (88.2) | -64.5 | -71.4 | 27.0 | -84.0 | -47.8 |
| ' (| eek 16 eek 26 eek 12 eek 16 eek 26 eek 12 | eek 12 327 (90.3) eek 16 323 (89.2) eek 26 303 (83.7) eek 12 327 (90.3) eek 16 323 (89.2) eek 26 303 (83.7) eek 12 324 (89.5) eek 12 320 (88.4) | eek 12 327 (90.3) 6.3 eek 16 323 (89.2) 6.2 eek 26 303 (83.7) 6.5 eek 12 327 (90.3) -14.0 eek 16 323 (89.2) -14.1 eek 26 303 (83.7) -13.7 eek 12 324 (89.5) -66.5 eek 16 320 (88.4) -67.0 | eek 12 327 (90.3) 6.3 4.0 eek 16 323 (89.2) 6.2 4.0 eek 26 303 (83.7) 6.5 5.0 eek 12 327 (90.3) -14.0 -15.0 eek 16 323 (89.2) -14.1 -15.0 eek 26 303 (83.7) -13.7 -14.0 eek 12 324 (89.5) -66.5 -77.8 eek 16 320 (88.4) -67.0 -76.9 | eek 12 327 (90.3) 6.3 4.0 6.1 eek 16 323 (89.2) 6.2 4.0 5.8 eek 26 303 (83.7) 6.5 5.0 6.4 eek 12 327 (90.3) -14.0 -15.0 7.0 eek 16 323 (89.2) -14.1 -15.0 6.8 eek 26 303 (83.7) -13.7 -14.0 7.3 eek 12 324 (89.5) -66.5 -77.8 70.5 eek 16 320 (88.4) -67.0 -76.9 60.8 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 -53.6 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 341 (93.4) eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 337 (92.3) eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 322 (88.2) eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 341 (93.4) eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 337 (92.3) eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 322 (88.2) eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 341 (93.4) eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 -53.6 337 (92.3) | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 341 (93.4) 8.0 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 337 (92.3) 7.8 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 322 (88.2) 7.2 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 341 (93.4) -12.7 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 337 (92.3) -12.9 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 322 (88.2) -13.5 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 341 (93.4) -59.9 eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 -53.6 337 (92.3) -61.1 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 341 (93.4) 8.0 6.0 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 337 (92.3) 7.8 7.0 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 322 (88.2) 7.2 6.0 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 341 (93.4) -12.7 -13.0 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 337 (92.3) -12.9 -13.0 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 322 (88.2) -13.5 -14.0 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 341 (93.4) -59.9 -68.4 eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 -53.6 337 (92.3) -61.1 -65.2 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 341 (93.4) 8.0 6.0 6.2 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 337 (92.3) 7.8 7.0 5.7 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 322 (88.2) 7.2 6.0 5.8 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 341 (93.4) -12.7 -13.0 6.9 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 337 (92.3) -12.9 -13.0 6.8 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 322 (88.2) -13.5 -14.0 6.6 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 341 (93.4) -59.9 -68.4 33.5 eek 16 320 (88.4) -67.0 -76.9 60.8 -91.1 -53.6 337 (92.3) -61.1 -65.2 27.9 | eek 12 327 (90.3) 6.3 4.0 6.1 2.0 9.0 341 (93.4) 8.0 6.0 6.2 3.0 eek 16 323 (89.2) 6.2 4.0 5.8 2.0 9.0 337 (92.3) 7.8 7.0 5.7 4.0 eek 26 303 (83.7) 6.5 5.0 6.4 1.0 10.0 322 (88.2) 7.2 6.0 5.8 3.0 eek 12 327 (90.3) -14.0 -15.0 7.0 -19.0 -9.0 341 (93.4) -12.7 -13.0 6.9 -18.0 eek 16 323 (89.2) -14.1 -15.0 6.8 -19.0 -9.0 337 (92.3) -12.9 -13.0 6.8 -18.0 eek 26 303 (83.7) -13.7 -14.0 7.3 -19.0 -9.0 322 (88.2) -13.5 -14.0 6.6 -18.0 eek 12 324 (89.5) -66.5 -77.8 70.5 -91.7 -54.5 341 (93.4) -59.9 -68.4 33.5 -82.4 eek 16 320 (88.4) -67.0 -76.9 60.8 - |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

POEM = patient-oriented eczema measure; OD = observed data. Data after dropout or use of rescue therapy was censored.

N = number of subjects in the analysis set.

n = number of subjects in the analysis set with observed data at the specified visit.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 05SEP2021 (10:05)

Least Squares Mean of Change from Baseline in POEM Total Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | | | |
| | N | 362 | 365 |
| | | | |
| Week 12 | N1 | 331 | 343 |
| | N2 (%) | 327 (98.8) | 341 (99.4) |
| | LSM (SE) | -14.23 (0.33) | -12.63 (0.32) |
| | 95% CI | (-14.87, -13.59) | (-13.26, -12.01) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.60 | |
| | 95% CI | (-2.49, -0.70) | |
| | Two-sided P-value | 0.0005 | |
| | Hedges'g | -0.27 | |
| | 95% CI of Hedges'g | (-0.42, -0.12) | |
| | | | |
| Week 16 | N1 | 326 | 338 |
| | N2 (%) | 323 (99.1) | 337 (99.7) |
| | 112 (70) | | |
| | LSM (SE) | -14.22 (0.31) | -12.83 (0.30) |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; POEM = patient-oriented eczema measure; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 02SEP2021 (23:49)

Least Squares Mean of Change from Baseline in POEM Total Score at Weeks 12, 16 and 26 - MMRM (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | -1.39 | |
| | 95% CI | (-2.24, -0.54) | |
| | Two-sided P-value | 0.0013 | |
| | Hedges'g | -0.25 | |
| | 95% CI of Hedges'g | (-0.40, -0.10) | |
| | · | | |
| Week 26 | N1 | 303 | 322 |
| | N2 (%) | 303 (100.0) | 322 (100.0) |
| | LSM (SE) | -13.81 (0.33) | -13.39 (0.33) |
| | 95% CI | (-14.47, -13.15) | (-14.03, -12.75 |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -0.42 | |
| | 95% CI | (-1.34, 0.50) | |
| | Two-sided P-value | 0.3684 | |
| | Hedges'g | -0.07 | |
| | 95% CI of Hedges'g | (-0.23, 0.09) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

CI = confidence interval; POEM = patient-oriented eczema measure; LSM = least squares mean; SE = Standard Error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint.

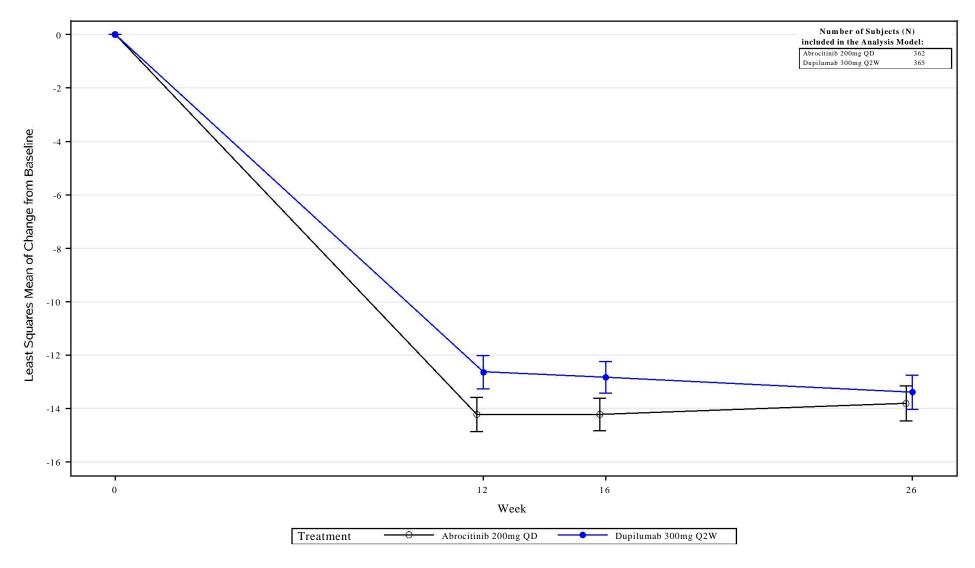
OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

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Parameter: Total Score



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

Vertical line represented 95% confidence interval.

POEM = patient-oriented eczema measure; OD = observed data. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix.

PFIZER CONFIDENTIAL SDTM Creation: 06AUG2021 (03:25) Source Data: adpm Table Generation: 05SEP2021 (09:35)

Output File: ./nda1_cdisc/B7451050_GBA/adpm_f201

Descriptive Summary of SCORAD Total Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | | Abrocitinib 200mg QD (N=362) | | | | |] | Dupiluı | nab 300n (N=365) | ng Q2W | | | | |
|------------------------------|----------|-------------|---------------------------------|--------|------|-------|-------|-------------|---------|---------------------|--------|-------|-------|--|--|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 | | |
| Observed Data | Baseline | 362 (100.0) | 67.8 | 66.4 | 12.8 | 58.9 | 76.8 | 365 (100.0) | 66.8 | 65.2 | 12.7 | 58.0 | 75.1 | | |
| | Week 2 | 349 (96.4) | 38.0 | 37.8 | 17.5 | 25.2 | 49.4 | 350 (95.9) | 44.4 | 44.9 | 18.0 | 30.4 | 56.6 | | |
| | Week 4 | 340 (93.9) | 27.5 | 25.7 | 15.8 | 17.5 | 35.8 | 351 (96.2) | 35.7 | 33.8 | 17.1 | 23.0 | 47.4 | | |
| | Week 8 | 335 (92.5) | 22.9 | 21.6 | 15.2 | 12.1 | 30.9 | 348 (95.3) | 29.4 | 27.3 | 15.6 | 18.6 | 39.2 | | |
| | Week 12 | 328 (90.6) | 20.9 | 19.3 | 14.7 | 10.7 | 29.3 | 341 (93.4) | 26.1 | 24.6 | 14.8 | 16.1 | 33.7 | | |
| | Week 16 | 324 (89.5) | 19.4 | 18.0 | 14.9 | 8.0 | 26.9 | 336 (92.1) | 23.5 | 21.2 | 14.7 | 14.3 | 31.2 | | |
| | Week 20 | 319 (88.1) | 18.3 | 16.4 | 14.8 | 7.2 | 25.8 | 333 (91.2) | 21.6 | 19.4 | 14.7 | 11.9 | 30.0 | | |
| | Week 26 | 300 (82.9) | 18.0 | 16.4 | 14.8 | 5.6 | 25.7 | 323 (88.5) | 20.9 | 18.9 | 14.6 | 11.5 | 28.4 | | |
| | | | | | | | | | | | | | | | |
| Change from Baseline | Week 2 | 349 (96.4) | -30.0 | -29.5 | 15.2 | -40.8 | -18.8 | 350 (95.9) | -22.4 | -20.2 | 16.5 | -34.1 | -10.5 | | |
| | Week 4 | 340 (93.9) | -40.6 | -40.5 | 15.7 | -51.0 | -30.7 | 351 (96.2) | -31.0 | -29.9 | 15.9 | -41.1 | -19.4 | | |
| | Week 8 | 335 (92.5) | -45.0 | -45.0 | 16.8 | -57.3 | -34.8 | 348 (95.3) | -37.2 | -36.7 | 16.8 | -49.1 | -25.5 | | |
| | Week 12 | 328 (90.6) | -46.9 | -47.5 | 17.6 | -57.8 | -35.3 | 341 (93.4) | -40.7 | -39.5 | 16.5 | -51.9 | -30.3 | | |
| | Week 16 | 324 (89.5) | -48.6 | -48.9 | 16.9 | -58.8 | -38.0 | 336 (92.1) | -43.3 | -43.2 | 17.5 | -53.9 | -32.0 | | |
| | Week 20 | 319 (88.1) | -49.7 | -50.6 | 17.5 | -60.6 | -38.4 | 333 (91.2) | -45.1 | -43.9 | 17.4 | -55.9 | -34.2 | | |
| | Week 26 | 300 (82.9) | -50.0 | -50.4 | 17.3 | -60.6 | -39.1 | 323 (88.5) | -45.8 | -46.4 | 17.4 | -57.0 | -33.8 | | |
| | | | | | | | | | | | | | | | |
| Percent Change from Baseline | Week 2 | 349 (96.4) | -44.6 | -45.3 | 22.4 | -59.7 | -27.0 | 350 (95.9) | -33.5 | -31.1 | 24.0 | -51.5 | -15.3 | | |
| | Week 4 | 340 (93.9) | -59.8 | -61.7 | 21.3 | -73.9 | -47.9 | 351 (96.2) | -46.9 | -46.5 | 22.5 | -63.5 | -30.1 | | |
| | Week 8 | 335 (92.5) | -66.2 | -68.6 | 22.4 | -81.2 | -54.8 | 348 (95.3) | -55.7 | -56.8 | 22.5 | -72.5 | -41.6 | | |
| | Week 12 | 328 (90.6) | -68.7 | -70.0 | 21.8 | -84.7 | -56.4 | 341 (93.4) | -60.7 | -62.2 | 21.0 | -76.1 | -49.3 | | |
| | Week 16 | 324 (89.5) | -71.4 | -73.0 | 21.0 | -88.1 | -59.6 | 336 (92.1) | -64.4 | -67.3 | 21.2 | -78.9 | -52.9 | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set; n = number of subjects in the analysis set with observed data at the specified visit; OD = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 15SEP2021 (02:28)

Descriptive Summary of SCORAD Total Score, Change from Baseline and Percent Change from Baseline up to Week 26 (FAS, OD) (Protocol B7451050)

| | | Abrocitinib 200mg QD (N=362) | | | | Dupilumab 300mg Q2W (N=365) | | | | | | | |
|---------------------------------|---------|---------------------------------|-------|--------|------|--------------------------------|-------|------------|-------|--------|------|-------|-------|
| | | n (%) | Mean | Median | SD | Q1 | Q3 | n (%) | Mean | Median | SD | Q1 | Q3 |
| Percent Change from Baseline | Week 20 | 319 (88.1) | -72.9 | -75.6 | 21.4 | -89.7 | -60.9 | 333 (91.2) | -67.3 | -70.0 | 21.2 | -81.6 | -54.7 |
| | Week 26 | 300 (82.9) | -73.4 | -75.9 | 21.3 | -91.6 | -62.4 | 323 (88.5) | -68.3 | -70.3 | 21.3 | -83.2 | -57.5 |
| | | | | | | | | | | | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

SCORAD = scoring atopic dermatitis; N = number of subjects in the analysis set; n = number of subjects in the analysis set with observed data at the specified visit; OD = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 15SEP2021 (02:28)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|-------------------------|------------------------|
| | | | |
| | N | 362 | 365 |
| | | | |
| Week 2 | N1 | 350 | 353 |
| | N2 (%) | 349 (99.7) | 350 (99.2) |
| | LSM (SE) | -44.52 (1.24) | -33.47 (1.24) |
| | 95% CI | (-46.96, -42.09) | (-35.90, -31.05) |
| | | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -11.05 | |
| | 95% CI | (-14.49, -7.61) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.48 | |
| | 95% CI of Hedges'g | (-0.63, -0.33) | |
| | | | |
| Week 4 | N1 | 349 | 352 |
| | N2 (%) | 340 (97.4) | 351 (99.7) |
| | LSM (SE) | -59.58 (1.18) | -46.83 (1.17) |
| | | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; SCORAD = scoring atopic dermatitis; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data. Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------|-------------------------|-------------------------|------------------------|
| | Abrocitinib - Dupilumab | | |
| | LSM | -12.75 | |
| | 95% CI | (-16.02, -9.49) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.58 | |
| | 95% CI of Hedges'g | (-0.74, -0.43) | |
| | <u> </u> | | |
| Week 8 | N1 | 342 | 351 |
| | N2 (%) | 335 (98.0) | 348 (99.1) |
| | LSM (SE) | -65.84 (1.22) | -55.64 (1.20) |
| | 95% CI | (-68.23, -63.45) | (-58.00, -53.27 |
| | <u> </u> | | |
| | Abrocitinib - Dupilumab | | |
| | LSM | -10.20 | |
| | 95% CI | (-13.57, -6.84) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.46 | |
| | 95% CI of Hedges'g | (-0.61, -0.30) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; SCORAD = scoring atopic dermatitis; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data. Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|----------------------------|-------------------------|------------------------|
| Week 12 | N1 | 331 | 343 |
| | N2 (%) | 328 (99.1) | 341 (99.4) |
| | LSM (SE) | -67.90 (1.17) | -60.59 (1.16) |
| | 95% CI | (-70.20, -65.60) | (-62.86, -58.32) |
| | A han sidin ih Danilham ah | | |
| | Abrocitinib - Dupilumab | 7.01 | |
| | LSM | -7.31 | |
| | 95% CI | (-10.54, -4.08) | |
| | Two-sided P-value | <.0001 | |
| | Hedges'g | -0.34 | |
| | 95% CI of Hedges'g | (-0.50, -0.19) | |
| Week 16 | N1 | 328 | 341 |
| WCCK 10 | N2 (%) | 324 (98.8) | 336 (98.5) |
| | | | |
| | LSM (SE) | -70.58 (1.16) | -64.44 (1.14) |
| | 95% CI | (-72.85, -68.31) | (-66.67, -62.20) |
| | Abrocitinib - Dupilumab | | |
| | • | -6.14 | |
| | LSM | | |
| | 95% CI | (-9.33, -2.96) | |
| | Two-sided P-value | 0.0002 | |
| | Hedges'g | -0.29 | |
| | 95% CI of Hedges'g | (-0.45, -0.14) | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; SCORAD = scoring atopic dermatitis; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data. Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (02:56)

| | | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|---------|-------------------------|-------------------------|------------------------|
| | | | |
| Week 20 | N1 | 322 | 334 |
| | N2 (%) | 319 (99.1) | 333 (99.7) |
| | LSM (SE) | -71.78 (1.19) | -66.84 (1.17) |
| | 95% CI | (-74.11, -69.45) | (-69.13, -64.54) |
| | Abrocitinib - Dupilumab | | |
| | LSM | -4.94 | |
| | 95% CI | (-8.22, -1.67) | |
| | Two-sided P-value | 0.0031 | |
| | Hedges'g | -0.23 | |
| | 95% CI of Hedges'g | (-0.39, -0.08) | |
| | | | |
| Week 26 | N1 | 300 | 323 |
| | N2 (%) | 300 (100.0) | 323 (100.0) |
| | LSM (SE) | -71.47 (1.23) | -68.21 (1.20) |
| | 95% CI | (-73.88, -69.06) | (-70.57, -65.86) |
| | Abrocitinib - Dupilumab | 1 | |
| | LSM | -3.26 | |
| | 95% CI | (-6.63, 0.11) | |
| | Two-sided P-value | 0.0578 | |
| | Hedges'g | -0.15 | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication. Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; SCORAD = scoring atopic dermatitis; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data. Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge. Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (02:56)

| | Abrocitinib 200mg QD | Dupilumab 300mg Q2W |
|--------------------|-------------------------|------------------------|
| 95% CI of Hedges'g | (-0.31, 0.01) | |
| | | |

Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

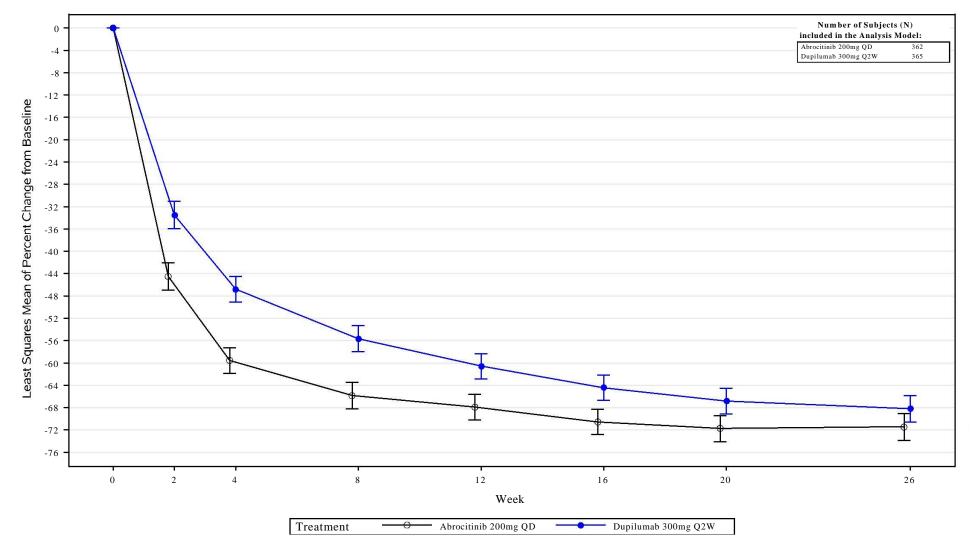
Baseline was defined as the measurement collected on or prior to Day 1.

Data after dropout or use of rescue therapy was censored.

CI = confidence interval; SCORAD = scoring atopic dermatitis; LSM = least squares mean; SE = standard error; N = number of subjects included in the analysis model; N1 = number of subjects at risk/eligible at the timepoint; N2 = number of subjects included in the analysis and with valid score at the timepoint; OD = observed data. Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge.

Hedges'g was defined as the LSM difference divided by pooled standard deviation.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (02:56)



Full Analysis Set (FAS) was defined as all randomized subjects who received at least one dose of study medication.

Baseline was defined as the measurement collected on or prior to Day 1. Data after dropout or use of rescue therapy was censored.

Mixed Model Repeated Measure (MMRM) contained fixed factors of treatment, visit, treatment by visit interaction, baseline disease severity, baseline value and an unstructured covariance matrix. Compound symmetry covariance matrix was used if the model with unstructured covariance did not converge.

Vertical line represented 95% confidence interval.

SCORAD = scoring atopic dermatitis; OD = observed data.

PFIZER CONFIDENTIAL SDTM Creation: 09AUG2021 (01:16) Source Data: adda Table Generation: 06SEP2021 (05:51)

Responderanalysen JADE TEEN (Woche 12)

| | PF-04965842 200mg QD | | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|---------------------------|----------------------|------------|----|------------|---------------|---------------|---|-------------------------------|--------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 93 | 32 (34.4%) | 96 | 12 (12.5%) | 3.58 | 2.71 | 21.8% | 0.0004+* | |
| Population | | (24.9%, | | (6.6%, | (1.71, 7.48) | (1.50, 4.90) | (10.0%, 33.6%) | | |
| • | | 45.0%) | | 20.8%) | | | | | |
| Baseline Disease Severity | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 20 (32.8%) | 57 | 9 (15.8%) | 2.60 | 2.08 | 17.0% | 0.0352* | 0.3459 |
| disease (IGA=3) | | (21.3%, | | (7.5%, | (1.07, 6.34) | (1.03, 4.18) | (1.9%, 32.1%) | | |
| | | 46.0%) | | 27.9%) | | | | | |
| Severe baseline disease | 32 | 12 (37.5%) | 39 | 3 (7.7%) | 7.20 | 4.88 | 29.8% | 0.0031* | |
| (IGA=4) | | (21.1%, | | (1.6%, | (1.81, 28.57) | (1.50, 15.80) | (11.1%, 48.6%) | | |
| | | 56.3%) | | 20.9%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 22 (39.3%) | 44 | 5 (11.4%) | 5.05 | 3.46 | 27.9% | 0.0028* | 0.2167 |
| | | (26.5%, | | (3.8%, | (1.72, 14.78) | (1.42, 8.39) | (12.1%, 43.8%) | | |
| | | 53.2%) | | 24.6%) | | | | | |
| Female | 37 | 10 (27.0%) | 52 | 7 (13.5%) | 2.38 | 2.01 | 13.6% | 0.1702 | |
| | | (13.8%, | | (5.6%, | (0.81, 6.99) | (0.84, 4.79) | (-3.5%, 30.6%) | | |
| | | 44.1%) | | 25.8%) | , , , , , , | , , , , , , | , | | |

Notes

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965842 200mg QD | | Placebo | | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------------------|------------|---------|------------|----------------|----------------|-------------------------|-------------------------------|--------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 7 (25.0%) | 33 | 9 (27.3%) | 0.89 | 0.92 | -2.3% | 1.0000 | 0.0464* |
| | | (10.7%, | | (13.3%, | (0.28, 2.80) | (0.39, 2.14) | (-24.4%, 19.8%) | | |
| | | 44.9%) | | 45.5%) | | | | | |
| Europe | 28 | 10 (35.7%) | 27 | 0 (0.0%) | 31.22 | 20.28 | 34.4% | 0.0007* | |
| | | (18.6%, | | | (1.72, 565.97) | (1.25, 329.82) | (16.3%, 52.6%) | | |
| | | 55.9%) | | | | | | | |
| Asia | 28 | 13 (46.4%) | 29 | 2 (6.9%) | 11.70 | 6.73 | 39.5% | 0.0008* | |
| | | (27.5%, | | (0.8%, | (2.32, 58.94) | (1.67, 27.17) | (18.9%, 60.2%) | | |
| | | 66.1%) | | 22.8%) | | | | | |
| Latin America | 9 | 2 (22.2%) | 7 | 1 (14.3%) | 1.71 | 1.56 | 7.9% | 1.0000 | |
| | | (2.8%, | | (0.4%, | (0.12, 23.94) | (0.17, 13.87) | (-29.6%, 45.5%) | | |
| | | 60.0%) | | 57.9%) | | | | | |
| Weight Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 17 (37.0%) | 53 | 7 (13.2%) | 3.85 | 2.80 | 23.7% | 0.0091* | 0.7273 |
| | | (23.2%, | | (5.5%, | (1.42, 10.42) | (1.27, 6.14) | (7.1%, 40.4%) | | |
| | | 52.5%) | | 25.3%) | | | | | |
| > Median Value | 47 | 15 (31.9%) | 43 | 5 (11.6%) | 3.56 | 2.74 | 20.3% | 0.0242* | |
| | | (19.1%, | | (3.9%, | (1.17, 10.88) | (1.09, 6.91) | (3.9%, 36.7%) | | |
| | | 47.1%) | | 25.1%) | | | | | |
| Age Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 56 | 19 (33.9%) | 60 | 8 (13.3%) | 3.34 | 2.54 | 20.6% | 0.0147* | 0.7994 |
| | | (21.8%, | | (5.9%, | (1.32, 8.44) | (1.21, 5.34) | (5.5%, 35.7%) | | |
| | | 47.8%) | | 24.6%) | | | | | |
| > Median Value | 37 | 13 (35.1%) | 36 | 4 (11.1%) | 4.33 | 3.16 | 24.0% | 0.0252* | |
| | | (20.2%, | | (3.1%, | (1.25, 14.96) | (1.14, 8.79) | (5.5%, 42.5%) | | |
| | | 52.5%) | | 26.1%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965842 200mg QD Placebo | | | | | Placebo vs. PF-04965842 200mg QD | | | |
|---------------------------|------------------------------|------------------------|----|---|---------------|--|----------------|----------------------------|-----------------------------|
| | | - | | F | OR | RR | RD | CMH or Logistic I | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | (95% CI) | (95% CI) | (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | (* * * * * *) | | (************************************** | , | (, , , , , , , , , , , , , , , , , , , | (2.2.2.) | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 93 | 11 (11.8%) | 96 | 4 (4.2%) | 3.00 | 2.75 | 7.5% | 0.0591+ | |
| Population | | (6.1%, | | (1.1%, | (0.92, 9.80) | (0.91, 8.29) | (-0.2%, 15.2%) | | |
| <u>r</u> | | 20.2%) | | 10.3%) | (===,===, | (,, | (, , | | |
| Baseline Disease Severity | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 8 (13.1%) | 57 | 3 (5.3%) | 2.72 | 2.49 | 7.9% | 0.2071 | 0.8706 |
| disease (IGA=3) | | (5.8%, | | (1.1%, | (0.68, 10.80) | (0.70, 8.93) | (-2.4%, 18.1%) | | |
| | | 24.2%) | | 14.6%) | | | | | |
| Severe baseline disease | 32 | 3 (9.4%) | 39 | 1 (2.6%) | 3.93 | 3.66 | 6.8% | 0.3207 | |
| (IGA=4) | | (2.0%, | | (0.1%, | (0.39, 39.77) | (0.40, 33.48) | (-4.4%, 18.1%) | | |
| | | 25.0%) | | 13.5%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 9 (16.1%) | 44 | 2 (4.5%) | 4.02 | 3.54 | 11.5% | 0.1061 | 0.1836 |
| | | (7.6%, | | (0.6%, | (0.82, 19.67) | (0.80, 15.54) | (0.1%, 22.9%) | | |
| | | 28.3%) | | 15.5%) | | | | | |
| Female | 37 | 2 (5.4%) | 52 | 2 (3.8%) | 1.43 | 1.41 | 1.6% | 1.0000 | |
| | | (0.7%, | | (0.5%, | (0.19, 10.63) | (0.21, 9.53) | (-7.4%, 10.5%) | | |
| | | 18.2%) | | 13.2%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965842 200mg QD | | Placebo | | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------------------|------------|---------|------------|----------------|----------------|---|-------------------------------|--------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 2 (7.1%) | 33 | 3 (9.1%) | 0.77 | 0.79 | -1.9% | 1.0000 | 0.1135 |
| | | (0.9%, | | (1.9%, | (0.12, 4.96) | (0.14, 4.37) | (-15.6%, 11.7%) | | |
| | | 23.5%) | | 24.3%) | | | | | |
| Europe | 28 | 4 (14.3%) | 27 | 0 (0.0%) | 10.10 | 8.69 | 13.7% | 0.1115 | |
| _ | | (4.0%, | | | (0.52, 197.31) | (0.49, 154.06) | (-0.3%, 27.8%) | | |
| | | 32.7%) | | | | | | | |
| Asia | 28 | 5 (17.9%) | 29 | 0 (0.0%) | 13.81 | 11.38 | 17.3% | 0.0235* | |
| | | (6.1%, | | | (0.73, 262.60) | (0.66, 196.67) | (2.3%, 32.3%) | | |
| | | 36.9%) | | | | | | | |
| Latin America | 9 | 0 (0.0%) | 7 | 1 (14.3%) | 0.23 | 0.27 | -13.8% | 0.4375 | |
| | | | | (0.4%, | (0.01, 6.52) | (0.01, 5.70) | (-44.0%, 16.5%) | | |
| | | | | 57.9%) | , , , | | | | |
| Weight Subgroup | | | | ŕ | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 4 (8.7%) | 53 | 1 (1.9%) | 4.95 | 4.61 | 6.8% | 0.1802 | 0.8482 |
| | | (2.4%, | | (0.0%, | (0.53, 46.00) | (0.53, 39.78) | (-2.1%, 15.7%) | | |
| | | 20.8%) | | 10.1%) | , | , , , | , , , | | |
| > Median Value | 47 | 7 (14.9%) | 43 | 3 (7.0%) | 2.33 | 2.13 | 7.9% | 0.3202 | |
| | | (6.2%, | | (1.5%, | (0.56, 9.67) | (0.59, 7.74) | (-4.8%, 20.6%) | | |
| | | 28.3%) | | 19.1%) | . , , | | , , , , , , | | |
| Age Subgroup | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 56 | 6 (10.7%) | 60 | 2 (3.3%) | 3.48 | 3.21 | 7.4% | 0.1527 | 0.9259 |
| | | (4.0%, | | (0.4%, | (0.67, 18.02) | (0.68, 15.27) | (-1.9%, 16.7%) | | |
| | | 21.9%) | | 11.5%) | | | , | | |
| > Median Value | 37 | 5 (13.5%) | 36 | 2 (5.6%) | 2.66 | 2.43 | 8.0% | 0.4297 | |
| | | (4.5%, | | (0.7%, | (0.48, 14.68) | (0.50, 11.74) | (-5.4%, 21.3%) | | |
| | | 28.8%) | | 18.7%) | (== =, = ===) | (====, =====, | (= = = = = = = = = = = = = = = = = = = | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: SCORAD-100 response at week 12 - Full Analysis Set Safety Population JADE TEEN $\,$ (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo vs | . PF-04965842 200mg QD | | |
|---------------------------|---------|---------------|----|--------------|----------------|----------------|---|-------------------|---------------|
| | | | | | | | | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 93 | 3 (3.2%) | 96 | 1 (1.0%) | 2.90 | 2.80 | 2.0% | 0.3447+ | |
| Population | | (0.7%, 9.1%) | | (0.0%, 5.7%) | (0.29, 28.68) | (0.30, 26.18) | (-2.1%, 6.1%) | | |
| Baseline Disease Severity | | , , , , | | , , , | , | , , | , | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 3 (4.9%) | 57 | 1 (1.8%) | 2.90 | 2.80 | 3.2% | 0.6194 | 0.5059 |
| disease (IGA=3) | | (1.0%, | | (0.0%, 9.4%) | (0.29, 28.68) | (0.30, 26.18) | (-3.2%, 9.6%) | | |
| , , | | 13.7%) | | , , , | , , | , , | , , , | | |
| Severe baseline disease | 32 | 0 (0.0%) | 39 | 0 (0.0%) | 1.22 | 1.21 | 0.3% | NE | |
| (IGA=4) | | , , | | , , | (0.02, 62.95) | (0.02, 59.45) | (-5.1%, 5.7%) | | |
| Gender | | | | | , , | , , | , , , | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 3 (5.4%) | 44 | 0 (0.0%) | 5.82 | 5.53 | 5.0% | 0.2533 | 0.1699 |
| | | (1.1%, | | , , | (0.29, 115.75) | (0.29, 104.25) | (-1.9%, 12.0%) | | |
| | | 14.9%) | | | , ,, | , , , , , , , | , | | |
| Female | 37 | 0 (0.0%) | 52 | 1 (1.9%) | 0.46 | 0.46 | -1.5% | 1.0000 | |
| | - | - (-/-/-/ | | (0.0%, | (0.02, 11.55) | (0.02, 11.11) | (-7.3%, 4.2%) | | |
| | | | | 10.3%) | (***=, -1100) | (5.52, 1111) | (1.270, 1.270) | | |

Notes

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD |) | |
|----------------------------|----------|------------------------------|----|------------------------------|------------------------|------------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Region | | , | | , , | , | , , | , | . , | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) (0.1%, 18.3%) | 33 | 1 (3.0%) (0.1%, 15.8%) | 1.19 (0.07, 19.86) | 1.18 (0.08, 17.99) | 0.5% (-8.5%, 9.6%) | 1.0000 | 0.9486 |
| Europe | 28 | 1 (3.6%) (0.1%, 18.3%) | 27 | 0 (0.0%) | 3.00 (0.12, 76.91) | 2.90 (0.12, 68.15) | 3.4% (-6.0%, 12.8%) | 1.0000 | |
| Asia | 28 | 1 (3.6%) (0.1%, 18.3%) | 29 | 0 (0.0%) | 3.22 (0.13, 82.38) | 3.10 (0.13, 73.12) | 3.5% (-5.8%, 12.8%) | 0.4912 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |
| Weight Subgroup Week 12 | | | | | , | , , | | | |
| <= Median Value | 46 | 2 (4.3%) (0.5%, 14.8%) | 53 | 0 (0.0%) | 6.01 (0.28, 128.50) | 5.74 (0.28, 116.67) | 4.4% (-2.5%, 11.3%) | 0.2134 | 0.3183 |
| > Median Value | 47 | 1 (2.1%) (0.1%, 11.3%) | 43 | 1 (2.3%) (0.1%, 12.3%) | 0.91 (0.06, 15.06) | 0.91 (0.06, 14.18) | -0.2% (-6.3%, 5.9%) | 1.0000 | |
| Age Subgroup Week 12 | | , | | , | | | | | |
| <= Median Value | 56 | 2 (3.6%) (0.4%, 12.3%) | 60 | 1 (1.7%) (0.0%, 8.9%) | 2.19 (0.19, 24.79) | 2.14 (0.20, 22.98) | 1.9% (-3.9%, 7.7%) | 0.6089 | 0.8841 |
| > Median Value | 37 | 1 (2.7%) (0.1%, 14.2%) | 36 | 0 (0.0%) | 3.00 (0.12, 76.09) | 2.92 (0.12, 69.43) | 2.6% (-4.6%, 9.8%) | 1.0000 | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|---------------------------|----------|--------------|----|------------|---------------|--------------|-------------------------|-------------------|--------------|
| | | _ | | | | | _ | CMH or Logistic P | -val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 67 (71.3%) | 96 | 39 (40.6%) | 3.46 | 1.71 | 29.7% | <0.0001+* | |
| Population | | (61.0%, | | (30.7%, | (1.89, 6.31) | (1.31, 2.24) | (16.2%, 43.3%) | | |
| • | | 80.1%) | | 51.1%) | | | | | |
| Baseline Disease Severity | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 43 (70.5%) | 57 | 31 (54.4%) | 2.00 | 1.30 | 16.1% | 0.0874 | 0.0873 |
| disease (IGA=3) | | (57.4%, | | (40.7%, | (0.94, 4.28) | (0.97, 1.73) | (-1.2%, 33.4%) | | |
| | | 81.5%) | | 67.6%) | | | | | |
| Severe baseline disease | 33 | 24 (72.7%) | 39 | 8 (20.5%) | 10.33 | 3.55 | 52.2% | < 0.0001* | |
| (IGA=4) | | (54.5%, | | (9.3%, | (3.47, 30.77) | (1.85, 6.81) | (32.4%, 72.0%) | | |
| | | 86.7%) | | 36.5%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 40 (71.4%) | 44 | 14 (31.8%) | 5.36 | 2.24 | 39.6% | 0.0001* | 0.4468 |
| | | (57.8%, | | (18.6%, | (2.27, 12.65) | (1.41, 3.57) | (21.5%, 57.8%) | | |
| | | 82.7%) | | 47.6%) | | | | | |
| Female | 38 | 27 (71.1%) | 52 | 25 (48.1%) | 2.65 | 1.48 | 23.0% | 0.0332* | |
| | | (54.1%, | | (34.0%, | (1.09, 6.44) | (1.04, 2.09) | (3.2%, 42.8%) | | |
| | | 84.6%) | | 62.4%) | | | | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

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The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------|---------------------|----|------------------------|----------------|----------------|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Region | | (93% CI) | 11 | (93% CI) | (93% CI) | (9370 C1) | (93% CI) | value [1] | Interaction |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 20 (71.4%) | 33 | 20 (60.6%) | 1.63 | 1.18 | 10.8% | 0.4269 | 0.0303* |
| | | (51.3%, | | (42.1%, | (0.55, 4.77) | (0.82, 1.69) | (-12.8%, 34.4%) | | |
| | | 86.8%) | | 77.1%) | , , , | | , | | |
| Europe | 29 | 22 (75.9%) | 27 | 5 (18.5%) | 13.83 | 4.10 | 57.3% | <0.0001* | |
| • | | (56.5%, | | (6.3%, | (3.80, 50.28) | (1.81, 9.28) | (36.0%, 78.7%) | | |
| | | 89.7%) | | 38.1%) | | | | | |
| Asia | 28 | 22 (78.6%) | 29 | 11 (37.9%) | 6.00 | 2.07 | 40.6% | 0.0030* | |
| | | (59.0%, | | (20.7%, | (1.86, 19.40) | (1.25, 3.43) | (17.3%, 63.9%) | | |
| | | 91.7%) | | 57.7%) | | | | | |
| Latin America | 9 | 3 (33.3%) | 7 | 3 (42.9%) | 0.67 | 0.78 | -9.5% | 1.0000 | |
| | | (7.5%, | | (9.9%, | (0.09, 5.13) | (0.22, 2.74) | (-57.4%, 38.4%) | | |
| | | 70.1%) | | 81.6%) | | | | | |
| Weight Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 35 (76.1%) | 53 | 22 (41.5%) | 4.48 | 1.83 | 34.6% | 0.0006* | 0.4191 |
| | | (61.2%, | | (28.1%, | (1.88, 10.71) | (1.28, 2.62) | (16.5%, 52.7%) | | |
| | | 87.4%) | | 55.9%) | | | | | |
| > Median Value | 48 | 32 (66.7%) | 43 | 17 (39.5%) | 3.06 | 1.69 | 27.1% | 0.0120* | |
| | | (51.6%, | | (25.0%, | (1.30, 7.20) | (1.11, 2.57) | (7.3%, 46.9%) | | |
| | | 79.6%) | | 55.6%) | | | | | |
| Age Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | . = |
| <= Median Value | 57 | 40 (70.2%) | 60 | 25 (41.7%) | 3.29 | 1.68 | 28.5% | 0.0028* | 0.7006 |
| | | (56.6%, | | (29.1%, | (1.53, 7.08) | (1.19, 2.38) | (11.3%, 45.7%) | | |
| | | 81.6%) | | 55.1%) | | | | | |
| > Median Value | 37 | 27 (73.0%) | 36 | 14 (38.9%) | 4.24 | 1.88 | 34.1% | 0.0047* | |
| | | (55.9%, | | (23.1%, | (1.58, 11.39) | (1.19, 2.95) | (12.7%, 55.5%) | | |
| 2 | | 86.2%) | | 56.5%) | | | | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|---------------------------|----------|--------------|----|------------|---------------|--------------|-------------------------|-------------------|---------------|
| | | - | | | | | _ | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 46 (48.9%) | 96 | 17 (17.7%) | 4.40 | 2.72 | 30.8% | <0.0001+* | |
| Population | | (38.5%, | | (10.7%, | (2.26, 8.54) | (1.69, 4.37) | (18.2%, 43.5%) | | |
| 1 | | 59.5%) | | 26.8%) | , , | ` ' ' | | | |
| Baseline Disease Severity | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 31 (50.8%) | 57 | 12 (21.1%) | 3.88 | 2.41 | 29.8% | 0.0011* | 0.9846 |
| disease (IGA=3) | | (37.7%, | | (11.4%, | (1.72, 8.72) | (1.38, 4.23) | (13.4%, 46.2%) | | |
| | | 63.9%) | | 33.9%) | | | | | |
| Severe baseline disease | 33 | 15 (45.5%) | 39 | 5 (12.8%) | 5.67 | 3.55 | 32.6% | 0.0033* | |
| (IGA=4) | | (28.1%, | | (4.3%, | (1.77, 18.12) | (1.44, 8.72) | (12.7%, 52.6%) | | |
| | | 63.6%) | | 27.4%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 29 (51.8%) | 44 | 7 (15.9%) | 5.68 | 3.26 | 35.9% | 0.0003* | 0.4307 |
| | | (38.0%, | | (6.6%, | (2.17, 14.87) | (1.58, 6.72) | (18.9%, 52.8%) | | |
| | | 65.3%) | | 30.1%) | | | | | |
| Female | 38 | 17 (44.7%) | 52 | 10 (19.2%) | 3.40 | 2.33 | 25.5% | 0.0113* | |
| | | (28.6%, | | (9.6%, | (1.33, 8.71) | (1.20, 4.50) | (6.4%, 44.6%) | | |
| | | 61.7%) | | 32.5%) | , , , , , | , , , , , , | , , , , , , , | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------|---------------|----|------------|---|---------------|-------------------------|------------------------------------|--------------|
| Tris (D. 1) | | Events (%) | | Events (%) | OR (050) GD | RR | RD | CMH or Logistic 1 Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 12 (42.9%) | 33 | 10 (30.3%) | 1.73 | 1.41 | 12.6% | 0.4231 | 0.2462 |
| | | (24.5%, | | (15.6%, | (0.60, 4.95) | (0.72, 2.77) | (-11.6%, 36.7%) | | |
| | | 62.8%) | | 48.7%) | | | | | |
| Europe | 29 | 15 (51.7%) | 27 | 2 (7.4%) | 13.39 | 6.98 | 44.3% | 0.0004* | |
| | | (32.5%, | | (0.9%, | (2.67, 67.27) | (1.76, 27.73) | (23.6%, 65.0%) | | |
| | | 70.6%) | | 24.3%) | | | | | |
| Asia | 28 | 16 (57.1%) | 29 | 4 (13.8%) | 8.33 | 4.14 | 43.3% | 0.0008* | |
| | | (37.2%, | | (3.9%, | (2.28, 30.39) | (1.58, 10.87) | (21.1%, 65.6%) | | |
| | | 75.5%) | | 31.7%) | | | | | |
| Latin America | 9 | 3 (33.3%) | 7 | 1 (14.3%) | 3.00 | 2.33 | 19.0% | 0.5846 | |
| | | (7.5%, | | (0.4%, | (0.24, 37.67) | (0.30, 17.88) | (-21.2%, 59.3%) | | |
| | | 70.1%) | | 57.9%) | (, , , , , , , , , , , , , , , , , , , | (, | , , , , , , , , , , , , | | |
| Weight Subgroup | | , | | 2.13,10, | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 23 (50.0%) | 53 | 8 (15.1%) | 5.63 | 3.31 | 34.9% | 0.0002* | 0.6171 |
| Tilodian varao | | (34.9%, | | (6.7%, | (2.18, 14.52) | (1.64, 6.68) | (17.5%, 52.3%) | 0.0002 | 0.0171 |
| | | 65.1%) | | 27.6%) | (2.10, 11.32) | (1.01, 0.00) | (17.570, 52.570) | | |
| > Median Value | 48 | 23 (47.9%) | 43 | 9 (20.9%) | 3.48 | 2.29 | 27.0% | 0.0087* | |
| / Wiedian Value | 40 | (33.3%, | 43 | (10.0%, | (1.37, 8.79) | (1.19, 4.39) | (8.3%, 45.6%) | 0.0007 | |
| | | 62.8%) | | 36.0%) | (1.57, 0.77) | (1.15, 4.55) | (0.570, 45.070) | | |
| Age Subgroup | | 02.070) | | 30.070) | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 57 | 27 (47.4%) | 60 | 12 (20.0%) | 3.60 | 2.37 | 27.4% | 0.0030* | 0.5035 |
| - Median value | 31 | (34.0%, | 00 | (10.8%, | (1.59, 8.16) | (1.33, 4.21) | (10.9%, 43.8%) | 0.0030 | 0.3033 |
| | | , | | 32.3%) | (1.39, 6.10) | (1.33, 4.21) | (10.9%, 43.0%) | | |
| . 34 1' 37 1 | 27 | 61.0%) | 26 | , | 6.54 | 2.70 | 27.50 | 0.0010* | |
| > Median Value | 37 | 19 (51.4%) | 36 | 5 (13.9%) | 6.54 | 3.70 | 37.5% | 0.0010* | |
| | | (34.4%, | | (4.7%, | (2.09, 20.54) | (1.55, 8.84) | (17.8%, 57.1%) | | |
| | | 68.1%) | | 29.5%) | | | | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | . PF-04965842 200mg QD | | |
|--------------------------------------|----------|---------------|----|--------------|----------------|----------------|------------------------|------------------------------------|-------------------------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic I Regression p- | P-val [2] for trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | · | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 8 (8.5%) | 96 | 2 (2.1%) | 4.16 | 3.79 | 6.1% | 0.0590+ | |
| Population | | (3.7%, 16.1%) | | (0.3%, 7.3%) | (0.85, 20.30) | (0.85, 17.01) | (-0.2%, 12.4%) | | |
| Baseline Disease Severity Week 12 | | , | | | | | | | |
| Moderate baseline | 61 | 7 (11.5%) | 57 | 2 (3.5%) | 3.56 | 3.27 | 8.0% | 0.1650 | 0.4321 |
| disease (IGA=3) | | (4.7%, | | (0.4%, | (0.71, 17.94) | (0.71, 15.10) | (-1.3%, 17.3%) | | |
| | | 22.2%) | | 12.1%) | | | | | |
| Severe baseline disease | 33 | 1 (3.0%) | 39 | 0 (0.0%) | 3.65 | 3.53 | 3.2% | 0.4583 | |
| (IGA=4) | | (0.1%, 15.8%) | | | (0.14, 92.55) | (0.15, 83.84) | (-4.6%, 10.9%) | | |
| Gender | | , | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 6 (10.7%) | 44 | 0 (0.0%) | 11.46 | 10.26 | 10.3% | 0.0332* | 0.1811 |
| | | (4.0%, | | | (0.63, 209.14) | (0.59, 177.39) | (1.5%, 19.1%) | | |
| | | 21.9%) | | | | | | | |
| Female | 38 | 2 (5.3%) | 52 | 2 (3.8%) | 1.39 | 1.37 | 1.4% | 1.0000 | |
| | | (0.6%, | | (0.5%, | (0.19, 10.33) | (0.20, 9.29) | (-7.4%, 10.2%) | | |
| | | 17.7%) | | 13.2%) | | | | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | . PF-04965842 200mg QD | | |
|---------------------|----------|------------------------|----|------------------------|----------------|----------------|------------------------|----------------------------|-----------------------------|
| | | | | | | | • | CMH or Logistic 1 | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Region | | | | | , , | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) | 33 | 2 (6.1%) | 0.57 | 0.59 | -2.5% | 1.0000 | 0.2550 |
| | | (0.1%, | | (0.7%, | (0.05, 6.69) | (0.06, 6.16) | (-13.1%, 8.2%) | | |
| | | 18.3%) | | 20.2%) | | | | | |
| Europe | 29 | 4 (13.8%) | 27 | 0 (0.0%) | 9.71 | 8.40 | 13.2% | 0.1124 | |
| | | (3.9%, | | | (0.50, 189.38) | (0.47, 149.04) | (-0.5%, 26.9%) | | |
| | | 31.7%) | | | | | | | |
| Asia | 28 | 3 (10.7%) | 29 | 0 (0.0%) | 8.10 | 7.24 | 10.4% | 0.1120 | |
| | | (2.3%, | | | (0.40, 164.32) | (0.39, 134.12) | (-2.3%, 23.1%) | | |
| | | 28.2%) | | | | | | | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| | | | | | (0.01, 44.65) | (0.02, 36.05) | (-22.8%, 20.3%) | | |
| Weight Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 5 (10.9%) | 53 | 1 (1.9%) | 6.34 | 5.76 | 9.0% | 0.0937 | 0.4285 |
| | | (3.6%, | | (0.0%, | (0.71, 56.42) | (0.70, 47.54) | (-0.7%, 18.7%) | | |
| | | 23.6%) | | 10.1%) | | | | | |
| > Median Value | 48 | 3 (6.3%) | 43 | 1 (2.3%) | 2.80 | 2.69 | 3.9% | 0.6189 | |
| | | (1.3%, | | (0.1%, | (0.28, 27.98) | (0.29, 24.88) | (-4.3%, 12.1%) | | |
| | | 17.2%) | | 12.3%) | | | | | |
| Age Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 57 | 5 (8.8%) | 60 | 2 (3.3%) | 2.79 | 2.63 | 5.4% | 0.2640 | 0.7276 |
| | | (2.9%, | | (0.4%, | (0.52, 14.99) | (0.53, 13.02) | (-3.2%, 14.1%) | | |
| | | 19.3%) | | 11.5%) | | | | | |
| > Median Value | 37 | 3 (8.1%) | 36 | 0 (0.0%) | 7.41 | 6.82 | 7.9% | 0.2397 | |
| | | (1.7%, | | | (0.37, 148.68) | (0.36, 127.44) | (-2.1%, 17.8%) | | |
| | | 21.9%) | | | | | | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-2 in POEM total score at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|---------------------------|----------|--------------|----|------------|---------------|---------------|-------------------------|-------------------|---------------|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 18 (19.1%) | 95 | 6 (6.3%) | 3.59 | 3.08 | 13.0% | 0.0077 + * | |
| Population | | (11.8%, | | (2.4%, | (1.35, 9.54) | (1.27, 7.45) | (3.7%, 22.3%) | | |
| | | 28.6%) | | 13.2%) | | | | | |
| Baseline Disease Severity | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 11 (18.0%) | 56 | 3 (5.4%) | 3.89 | 3.37 | 12.7% | 0.0460* | 0.9046 |
| disease (IGA=3) | | (9.4%, | | (1.1%, | (1.02, 14.75) | (0.99, 11.45) | (1.4%, 24.0%) | | |
| | | 30.0%) | | 14.9%) | | | | | |
| Severe baseline disease | 33 | 7 (21.2%) | 39 | 3 (7.7%) | 3.23 | 2.76 | 13.5% | 0.1698 | |
| IGA=4) | | (9.0%, | | (1.6%, | (0.76, 13.68) | (0.77, 9.82) | (-2.7%, 29.8%) | | |
| | | 38.9%) | | 20.9%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 11 (19.6%) | 44 | 4 (9.1%) | 2.44 | 2.16 | 10.6% | 0.1683 | 0.7260 |
| | | (10.2%, | | (2.5%, | (0.72, 8.29) | (0.74, 6.33) | (-2.9%, 24.0%) | | |
| | | 32.4%) | | 21.7%) | | | | | |
| Female | 38 | 7 (18.4%) | 51 | 2 (3.9%) | 5.53 | 4.70 | 14.5% | 0.0343* | |
| | | (7.7%, | | (0.5%, | (1.08, 28.37) | (1.03, 21.36) | (1.1%, 27.9%) | | |
| | | 34.3%) | | 13.5%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | | |
|---------------------|----------|------------------------|---------|------------------------|----------------|-------------------------|------------------|----------------------------|-----------------------------|
| | | • | | | | | | CMH or Logistic 1 | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 5 (17.9%) | 32 | 6 (18.8%) | 0.94 | 0.95 | -0.9% | 1.0000 | 0.1239 |
| | | (6.1%, | | (7.2%, | (0.25, 3.50) | (0.33, 2.78) | (-20.5%, 18.7%) | | |
| | | 36.9%) | | 36.4%) | | | | | |
| Europe | 29 | 8 (27.6%) | 27 | 0 (0.0%) | 21.74 | 15.87 | 26.5% | 0.0046* | |
| | | (12.7%, | | | (1.19, 398.13) | (0.96, 262.30) | (9.7%, 43.4%) | | |
| | | 47.2%) | | | , , , | | , , , | | |
| Asia | 28 | 5 (17.9%) | 29 | 0 (0.0%) | 13.81 | 11.38 | 17.3% | 0.0235* | |
| | | (6.1%, | | ` / | (0.73, 262.60) | (0.66, 196.67) | (2.3%, 32.3%) | | |
| | | 36.9%) | | | (, | (====, | (12 11, 12 11, | | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| | | (/ | | (, , , , , | (0.01, 44.65) | (0.02, 36.05) | (-22.8%, 20.3%) | | |
| Weight Subgroup | | | | | (6.61, 1.162) | (0.02, 20.02) | (22.070, 20.070) | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 8 (17.4%) | 52 | 0 (0.0%) | 23.18 | 19.17 | 17.1% | 0.0017* | 0.3671 |
| | | (7.8%, | | * (***,*) | (1.30, 413.93) | (1.14, 323.23) | (5.8%, 28.4%) | | |
| | | 31.4%) | | | (====, =====, | (===, =====, | (===,=,===,=, | | |
| > Median Value | 48 | 10 (20.8%) | 43 | 6 (14.0%) | 1.62 | 1.49 | 6.9% | 0.4230 | |
| Tribular variation | | (10.5%, | | (5.3%, | (0.54, 4.92) | (0.59, 3.76) | (-8.6%, 22.3%) | 0250 | |
| | | 35.0%) | | 27.9%) | (0.31, 1.52) | (0.35, 3.70) | (0.070, 22.370) | | |
| Age Subgroup | | 22.070) | | 2.1370) | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 57 | 13 (22.8%) | 59 | 5 (8.5%) | 3.19 | 2.69 | 14.3% | 0.0413* | 0.6253 |
| | | (12.7%, | | (2.8%, | (1.06, 9.64) | (1.03, 7.06) | (1.3%, 27.3%) | 2.0112 | 0.0200 |
| | | 35.8%) | | 18.7%) | (1.00, 7.0.) | (1.05, 7.00) | (1.570, 27.570) | | |
| > Median Value | 37 | 5 (13.5%) | 36 | 1 (2.8%) | 5.47 | 4.86 | 10.7% | 0.1992 | |
| - 1.1eGiun vuiuc | 51 | (4.5%, | 30 | (0.1%, | (0.61, 49.35) | (0.60, 39.63) | (-1.5%, 23.0%) | 0.1772 | |
| | | 28.8%) | | 14.5%) | (0.01, 49.33) | (0.00, 39.03) | (-1.5/0, 25.0/0) | | |

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0 in POEM total score at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-049658 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | | |
|--|-----------|------------------------------|----|------------------------------|-----------------------|-----------------------|-------------------------|--|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | , | | , | , | | | | | |
| Week 12 | | | | | | | | | | |
| Full Analysis Set Safety Population | 94 | 6 (6.4%) (2.4%, 13.4%) | 95 | 4 (4.2%) (1.2%, 10.4%) | 1.52 (0.41, 5.54) | 1.48 (0.44, 5.04) | 2.1% (-4.3%, 8.5%) | 0.5272+ | | |
| Baseline Disease Severity Week 12 | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 4 (6.6%) (1.8%, 15.9%) | 56 | 3 (5.4%) (1.1%, 14.9%) | 1.24 (0.26, 5.80) | 1.22 (0.29, 5.23) | 1.2% (-7.4%, 9.8%) | 1.0000 | 0.7282 | |
| Severe baseline disease (IGA=4) | 33 | 2 (6.1%) (0.7%, 20.2%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 2.45 (0.21, 28.32) | 2.36 (0.22, 24.91) | 3.5% (-6.0%, 13.0%) | 0.5900 | | |
| Gender Week 12 | | / | | | | | | | | |
| Male | 56 | 3 (5.4%) (1.1%, 14.9%) | 44 | 2 (4.5%) (0.6%, 15.5%) | 1.19 (0.19, 7.44) | 1.18 (0.21, 6.75) | 0.8% (-7.7%, 9.3%) | 1.0000 | 0.6328 | |
| Female | 38 | 3 (7.9%) (1.7%, 21.4%) | 51 | 2 (3.9%) (0.5%, 13.5%) | 2.10 (0.33, 13.24) | 2.01 (0.35, 11.46) | 4.0% (-6.1%, 14.1%) | 0.6473 | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | 842 200mg QD | | | | Placebo vs | | | |
|-------------|------------------------|----|---------------------|----------------|------------------|---|-------------------------------|--|
| Events (0/) | | | | | | | CMH or Logistic P-val [2] for | |
| n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| | , | | | , , | | | | |
| | | | | | | | | |
| 28 | 2 (7.1%) | 32 | 4 (12.5%) | 0.54 | 0.57 | -5.4% | 0.6754 | 0.4377 |
| | (0.9%, | | (3.5%, | (0.09, 3.19) | (0.11, 2.89) | (-20.3%, 9.6%) | | |
| | 23.5%) | | 29.0%) | , , | ` ' ' | , | | |
| 29 | 1 (3.4%) | 27 | 0 (0.0%) | 2.89 | 2.80 | 3.2% | 1.0000 | |
| | (0.1%, | | , , , | (0.11, 74.15) | (0.12, 65.93) | (-6.0%, 12.4%) | | |
| | , , | | | , | | | | |
| 28 | | 29 | 0 (0.0%) | 8.10 | 7.24 | 10.4% | 0.1120 | |
| | , | | (, | | (0.39, 134.12) | (-2.3%, 23.1%) | | |
| | | | | (0.10, 1010_) | (3.62), 20.1122) | (====================================== | | |
| 9 | , | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| | (0.0,0) | | * (***,*) | | | | | |
| | | | | (0.01, 1.1.02) | (0.02, 20.02) | (22.070, 20.070) | | |
| | | | | | | | | |
| 46 | 2 (4.3%) | 52 | 0 (0.0%) | 5.90 | 5.64 | 4.4% | 0.2178 | 0.4530 |
| | | 02 | 0 (0.070) | | | | 0.2170 | 0 |
| | ` ' | | | (0.20, 120.12) | (0.20, 11) | (2.570, 11.570) | | |
| 48 | , | 43 | 4 (9 3%) | 0.89 | 0.90 | -1.0% | 1 0000 | |
| .0 | | | | | | | 1.0000 | |
| | | | | (0.21, 3.70) | (0.21, 3.30) | (12.770, 10.770) | | |
| | 20.070) | | 22.170) | | | | | |
| | | | | | | | | |
| 57 | 2 (3.5%) | 59 | 3 (5.1%) | 0.68 | 0.69 | -1.6% | 1.0000 | 0.1579 |
| ٥, | ` ' | | ` ' | | | | 2,0000 | 0.1577 |
| | | | | (0.11,22) | (0.12, 0.50) | (0.570, 5.070) | | |
| 37 | , | 36 | , | 4 24 | 3 89 | 8.0% | 0.3575 | |
| 57 | | 30 | | | | | 0.5515 | |
| | | | | (0.43, 39.94) | (0.40, 33.17) | (-3.370, 19.470) | | |
| | 28 29 28 | 28 | 28 | 28 | 28 | 28 | 28 | 28 2 (7.1%) 32 4 (12.5%) 0.54 0.57 -5.4% 0.6754 (0.9%, 3.5%, 0.09, 3.19) (0.11, 2.89) (-20.3%, 9.6%) 23.5%) 29.0%) 2.89 2.80 3.2% 1.0000 (0.1%, 17.8%) 28 3 (10.7%) 29 0 (0.0%) 8.10 7.24 10.4% 0.1120 (2.3%, 2.82%) 9 0 (0.0%) 7 0 (0.0%) 0.79 0.80 -1.3% NE (0.00, 44.65) (0.01, 44.65) (0.02, 36.05) (-22.8%, 20.3%) 14.8%) 14.8% 14.8% 14.8% 14.8% 14.8% 14.8% 14.8% 14.8% 14.8% 12.1% 12.1% 12.1% 14.1% 15.1% 10.1% 12.1% 12.1% 14.1% 13.1% 13.1% 12.1% 12.1% 14.1% 13.1% |

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in CDLQI total score at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------------|----------|--------------|----|------------|--------------|--------------|-------------------------|-------------------|---------------|
| | | • | | | | | • | CMH or Logistic F | P-val [2] for |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 21 (22.3%) | 96 | 13 (13.5%) | 1.78 | 1.59 | 8.2% | 0.1382+ | |
| Population | | (14.4%, | | (7.4%, | (0.83, 3.83) | (0.85, 2.98) | (-2.6%, 19.0%) | | |
| • | | 32.1%) | | 22.0%) | | | | | |
| Baseline Disease Severity | | , | | • | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 61 | 16 (26.2%) | 57 | 10 (17.5%) | 1.67 | 1.50 | 8.7% | 0.2757 | 0.8366 |
| disease (IGA=3) | | (15.8%, | | (8.7%, | (0.69, 4.07) | (0.74, 3.02) | (-6.1%, 23.5%) | | |
| | | 39.1%) | | 29.9%) | | | | | |
| Severe baseline disease | 33 | 5 (15.2%) | 39 | 3 (7.7%) | 2.14 | 1.97 | 7.5% | 0.4563 | |
| (IGA=4) | | (5.1%, | | (1.6%, | (0.47, 9.74) | (0.51, 7.63) | (-7.4%, 22.3%) | | |
| | | 31.9%) | | 20.9%) | | | | | |
| Gender | | ŕ | | , | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 11 (19.6%) | 44 | 4 (9.1%) | 2.44 | 2.16 | 10.6% | 0.1683 | 0.9548 |
| | | (10.2%, | | (2.5%, | (0.72, 8.29) | (0.74, 6.33) | (-2.9%, 24.0%) | | |
| | | 32.4%) | | 21.7%) | | | | | |
| Female | 38 | 10 (26.3%) | 52 | 9 (17.3%) | 1.71 | 1.52 | 9.0% | 0.3113 | |
| | | (13.4%, | | (8.2%, | (0.62, 4.73) | (0.68, 3.38) | (-8.4%, 26.4%) | | |
| | | 43.1%) | | 30.3%) | | ` ' ' | ` ' | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | PF-04965842 200mg QD | | Placebo | | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------------------|------------------------|---------|---------------------|----------------|---|-------------------------|----------------------------|-----------------------------|
| | | • | | | | | • | CMH or Logistic I | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 6 (21.4%) | 33 | 6 (18.2%) | 1.23 | 1.18 | 3.2% | 0.7590 | 0.0577 |
| | | (8.3%, | | (7.0%, | (0.35, 4.34) | (0.43, 3.25) | (-16.9%, 23.4%) | | |
| | | 41.0%) | | 35.5%) | | | | | |
| Europe | 29 | 10 (34.5%) | 27 | 1 (3.7%) | 13.68 | 9.31 | 30.8% | 0.0056* | |
| • | | (17.9%, | | (0.1%, | (1.61, 116.20) | (1.28, 67.95) | (12.1%, 49.5%) | | |
| | | 54.3%) | | 19.0%) | | | | | |
| Asia | 28 | 5 (17.9%) | 29 | 6 (20.7%) | 0.83 | 0.86 | -2.8% | 1.0000 | |
| | | (6.1%, | | (8.0%, | (0.22, 3.12) | (0.30, 2.51) | (-23.3%, 17.6%) | | |
| | | 36.9%) | | 39.7%) | (, , | (, , , , , , , , , , , , , , , , , , , | (,, | | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| | | (/ | | ` / | (0.01, 44.65) | (0.02, 36.05) | (-22.8%, 20.3%) | | |
| Weight Subgroup | | | | | (0.000) | (333_, 23332) | (==:::, =:::, | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 9 (19.6%) | 53 | 7 (13.2%) | 1.60 | 1.48 | 6.4% | 0.4241 | 0.6557 |
| | | (9.4%, | | (5.5%, | (0.54, 4.70) | (0.60, 3.66) | (-8.3%, 21.0%) | | |
| | | 33.9%) | | 25.3%) | (0.0.1, 11.0) | (3.33, 2.33) | (0.070, = 0.070) | | |
| > Median Value | 48 | 12 (25.0%) | 43 | 6 (14.0%) | 2.06 | 1.79 | 11.0% | 0.2917 | |
| Tributari varao | .0 | (13.6%, | | (5.3%, | (0.70, 6.07) | (0.74, 4.36) | (-5.0%, 27.1%) | 0.2717 | |
| | | 39.6%) | | 27.9%) | (0.70, 0.07) | (6.7.1, 1.156) | (5.070, 27.170) | | |
| Age Subgroup | | 57.070) | | 21.570) | | | | | |
| Week 12 | | | | | | | | | |
| = Median Value | 57 | 12 (21.1%) | 60 | 9 (15.0%) | 1.51 | 1.40 | 6.1% | 0.4729 | 0.5333 |
| | | (11.4%, | 00 | (7.1%, | (0.58, 3.92) | (0.64, 3.08) | (-7.9%, 20.0%) | 2> | 0.0000 |
| | | 33.9%) | | 26.6%) | (0.00, 0.02) | (0.0., 2.00) | (1.570, 20.073) | | |
| > Median Value | 37 | 9 (24.3%) | 36 | 4 (11.1%) | 2.57 | 2.19 | 13.2% | 0.2210 | |
| 1,100/aii vaiuo | 51 | (11.8%, | 50 | (3.1%, | (0.71, 9.27) | (0.74, 6.48) | (-4.0%, 30.4%) | 0.2210 | |
| | | 41.2%) | | 26.1%) | (0.71, 9.27) | (0.74, 0.48) | (-4.070, 30.470) | | |

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: PP-NRS total score 4-point improvement at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|--|----------|---------------------------------|----|---------------------------------|----------------------|---|-------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | (* * * * * / | | (| , | (************************************** | (| | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety Population | 90 | 41 (45.6%) (35.0%, 56.4%) | 96 | 25 (26.0%) (17.6%, 36.0%) | 2.34 (1.26, 4.33) | 1.73 (1.15, 2.61) | 19.1% (5.5%, 32.6%) | 0.0067+* | |
| Baseline Disease Severity Week 12 | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 58 | 30 (51.7%) (38.2%, 65.0%) | 57 | 15 (26.3%) (15.5%, 39.7%) | 3.00 (1.37, 6.56) | 1.97 (1.19, 3.24) | 25.4% (8.2%, 42.6%) | 0.0073* | 0.1868 |
| Severe baseline disease (IGA=4) | 32 | 11 (34.4%) (18.6%, 53.2%) | 39 | 10 (25.6%) (13.0%, 42.1%) | 1.52 (0.55, 4.23) | 1.34 (0.65, 2.75) | 8.7% (-12.7%, 30.1%) | 0.4459 | |
| Gender Week 12 | | , | | , | | | | | |
| Male | 53 | 26 (49.1%) (35.1%, 63.2%) | 44 | 13 (29.5%) (16.8%, 45.2%) | 2.30 (0.99, 5.33) | 1.66 (0.97, 2.83) | 19.5% (0.5%, 38.6%) | 0.0627 | 0.7702 |
| Female | 37 | 15 (40.5%) (24.8%, 57.9%) | 52 | 12 (23.1%) (12.5%, 36.8%) | 2.27 (0.91, 5.70) | 1.76 (0.93, 3.30) | 17.5% (-2.1%, 37.0%) | 0.1024 | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline PP-NRS score <4.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | PF-04965 | 5842 200mg QD | Placebo | | | Placebo v | s. PF-04965842 200mg QD | | | |
|---------------------|------------|------------------------|-----------|------------------------|----------------|----------------|-------------------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic F Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Region | | (23/0 CI) | 11 | (7370 C1) | (7370 C1) | (7370 CI) | (7570 C1) | value [1] | meraction | |
| Week 12 | | | | | | | | | | |
| US/Canada/Australia | 26 | 14 (53.8%) | 33 | 16 (48.5%) | 1.24 | 1.11 | 5.4% | 0.7948 | 0.5058 | |
| | | (33.4%, | | (30.8%, | (0.44, 3.47) | (0.67, 1.83) | (-20.3%, 31.0%) | | | |
| | | 73.4%) | | 66.5%) | | | | | | |
| Europe | 28 | 13 (46.4%) | 27 | 2 (7.4%) | 10.83 | 6.27 | 39.0% | 0.0019* | | |
| | | (27.5%, | | (0.9%, | (2.14, 54.77) | (1.56, 25.21) | (18.1%, 60.0%) | | | |
| | | 66.1%) | | 24.3%) | | | | | | |
| Asia | 28 | 12 (42.9%) | 29 | 7 (24.1%) | 2.36 | 1.78 | 18.7% | 0.1665 | | |
| | | (24.5%, | | (10.3%, | (0.76, 7.32) | (0.82, 3.85) | (-5.3%, 42.8%) | | | |
| | 0 | 62.8%) | 7 | 43.5%) | 5.55 | 4.44 | 21.50/ | 0.4667 | | |
| Latin America | 8 | 2 (25.0%) | 7 | 0 (0.0%) | 5.77 | 4.44 | 21.5% | 0.4667 | | |
| | | (3.2%, | | | (0.23, 143.37) | (0.25, 79.42) | (-12.2%, 55.3%) | | | |
| Weight Subgroup | | 65.1%) | | | | | | | | |
| Week 12 | | | | | | | | | | |
| <= Median Value | 44 | 21 (47.7%) | 53 | 11 (20.8%) | 3.49 | 2.30 | 27.0% | 0.0087* | 0.2980 | |
| | | (32.5%, | | (10.8%, | (1.43, 8.48) | (1.25, 4.23) | (8.6%, 45.3%) | | | |
| | | 63.3%) | | 34.1%) | | | | | | |
| > Median Value | 46 | 20 (43.5%) | 43 | 14 (32.6%) | 1.59 | 1.34 | 10.9% | 0.3830 | | |
| | | (28.9%, | | (19.1%, | (0.67, 3.78) | (0.78, 2.30) | (-9.1%, 31.0%) | | | |
| | | 58.9%) | | 48.5%) | | | | | | |
| Age Subgroup | | | | | | | | | | |
| Week 12 | <i>~</i> . | 20 (51 00() | 60 | 10 (20 00() | 2.51 | 1.72 | 21.00/ | 0.0221* | 0.5000 | |
| <= Median Value | 54 | 28 (51.9%) | 60 | 18 (30.0%) | 2.51 | 1.73 | 21.9% | 0.0221* | 0.5223 | |
| | | (37.8%, | | (18.8%, | (1.17, 5.42) | (1.09, 2.75) | (4.2%, 39.5%) | | | |
| . M. di W.L | 26 | 65.7%) | 26 | 43.2%) | 2.24 | 1.00 | 1670/ | 0.1077 | | |
| > Median Value | 36 | 13 (36.1%) | 36 | 7 (19.4%) | 2.34 | 1.86 | 16.7% | 0.1877 | | |
| | | (20.8%, 53.8%) | | (8.2%, 36.0%) | (0.80, 6.82) | (0.84, 4.11) | (-3.7%, 37.0%) | | | |

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline PP-NRS score <4.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|--------------------------------------|----------|-------------------|----|------------------|--------------|--------------|-------------------------|------------------------------------|-------------------------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic F Regression p- | P-val [2] for trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | , | | , | , | , | , | . , | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 94 | 18 (19.1%) | 96 | 10 (10.4%) | 2.03 | 1.83 | 8.7% | 0.0935+ | |
| Population | | (11.8%, 28.6%) | | (5.1%, 18.3%) | (0.88, 4.67) | (0.89, 3.77) | (-1.4%, 18.7%) | | |
| Baseline Disease Severity Week 12 | | • | | , | | | | | |
| Moderate baseline | 61 | 12 (19.7%) | 57 | 6 (10.5%) | 2.08 | 1.87 | 9.1% | 0.2051 | 0.9027 |
| disease (IGA=3) | | (10.6%, | | (4.0%, | (0.72, 5.98) | (0.75, 4.65) | (-3.6%, 21.9%) | | |
| | | 31.8%) | | 21.5%) | | | | | |
| Severe baseline disease | 33 | 6 (18.2%) | 39 | 4 (10.3%) | 1.94 | 1.77 | 7.9% | 0.4962 | |
| (IGA=4) | | (7.0%, | | (2.9%, | (0.50, 7.58) | (0.55, 5.75) | (-8.3%, 24.2%) | | |
| | | 35.5%) | | 24.2%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 56 | 12 (21.4%) | 44 | 5 (11.4%) | 2.13 | 1.89 | 10.1% | 0.2834 | 0.6841 |
| | | (11.6%, | | (3.8%, | (0.69, 6.58) | (0.72, 4.95) | (-4.2%, 24.3%) | | |
| | | 34.4%) | | 24.6%) | | | | | |
| Female | 38 | 6 (15.8%) | 52 | 5 (9.6%) | 1.76 | 1.64 | 6.2% | 0.5170 | |
| | | (6.0%, | | (3.2%, | (0.50, 6.27) | (0.54, 4.99) | (-7.9%, 20.3%) | | |
| | | 31.3%) | | 21.0%) | | | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|----------------------|----------|---------------------|----|------------------------|-----------------|-----------------|-------------------------|-------------------------------|-----------------------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgrouj interaction |
| Region | | (3570 61) | | (5570 61) | (5570 61) | (50,0 01) | (50,0 01) | , and [1] | menan |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 28 | 8 (28.6%) | 33 | 9 (27.3%) | 1.07 | 1.05 | 1.3% | 1.0000 | 0.8159 |
| | | (13.2%, | | (13.3%, | (0.35, 3.28) | (0.47, 2.35) | (-21.3%, 23.9%) | | |
| | | 48.7%) | | 45.5%) | (3.22, 2.23) | (****, =:e=/ | (==== ,=, === ,= , | | |
| Europe | 29 | 5 (17.2%) | 27 | 0 (0.0%) | 12.35 | 10.27 | 16.5% | 0.0522 | |
| | _, | (5.8%, | _, | * (***,*) | (0.65, 234.92) | (0.59, 177.29) | (1.9%, 31.2%) | | |
| | | 35.8%) | | | (5152, 22 13 2) | (0.05, 17, 125) | (===,=,==,=, | | |
| Asia | 28 | 4 (14.3%) | 29 | 1 (3.4%) | 4.67 | 4.14 | 10.8% | 0.1936 | |
| 7 1514 | | (4.0%, | | (0.1%, | (0.49, 44.64) | (0.49, 34.82) | (-3.7%, 25.4%) | 0.1750 | |
| | | 32.7%) | | 17.8%) | (0.13, 1.110.) | (61.5, 51.62) | (51.70, 251.70) | | |
| Latin America | 9 | 1 (11.1%) | 7 | 0 (0.0%) | 2.65 | 2.40 | 8.8% | 1.0000 | |
| Eutin / Interior | Ź | (0.3%, | , | 0 (0.070) | (0.09, 75.29) | (0.11, 51.32) | (-19.0%, 36.5%) | 1.0000 | |
| | | 48.2%) | | | (0.0), 73.2) | (0.11, 31.32) | (17.070, 30.370) | | |
| Weight Subgroup | | 10.270) | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 46 | 8 (17.4%) | 53 | 4 (7.5%) | 2.58 | 2.30 | 9.8% | 0.2162 | 0.8132 |
| Tirodian varao | .0 | (7.8%, | 00 | (2.1%, | (0.72, 9.21) | (0.74, 7.16) | (-3.2%, 22.9%) | 0.2102 | 0.0152 |
| | | 31.4%) | | 18.2%) | (0.72, 3.21) | (01, 1, 7,110) | (5.270, 22.570) | | |
| > Median Value | 48 | 10 (20.8%) | 43 | 6 (14.0%) | 1.62 | 1.49 | 6.9% | 0.4230 | |
| y integratif variate | .0 | (10.5%, | | (5.3%, | (0.54, 4.92) | (0.59, 3.76) | (-8.6%, 22.3%) | 0250 | |
| | | 35.0%) | | 27.9%) | (3.2.1, 1.2.2) | (3.53, 21.3) | (5.575, ==1575) | | |
| Age Subgroup | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 57 | 12 (21.1%) | 60 | 9 (15.0%) | 1.51 | 1.40 | 6.1% | 0.4729 | 0.5179 |
| | | (11.4%, | | (7.1%, | (0.58, 3.92) | (0.64, 3.08) | (-7.9%, 20.0%) | | |
| | | 33.9%) | | 26.6%) | . , , | | | | |
| > Median Value | 37 | 6 (16.2%) | 36 | 1 (2.8%) | 6.77 | 5.84 | 13.4% | 0.1070 | |
| | | (6.2%, | | (0.1%, | (0.77, 59.42) | (0.74, 46.11) | (0.4%, 26.5%) | | |
| | | 32.0%) | | 14.5%) | , , , , , | , , , , , | , ,, | | |

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: EQ-5D VAS 15-point improvement at week 12 - Full Analysis Set Safety Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------------|----------|--------------|----|------------|---------------|--------------|-------------------------|-------------------|---------------|
| | | - | | | | | _ | CMH or Logistic I | P-val [2] for |
| *** ** / 5 | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Full Analysis Set Safety | 75 | 41 (54.7%) | 73 | 32 (43.8%) | 1.53 | 1.24 | 10.7% | 0.1990+ | |
| Population | | (42.7%, | | (32.2%, | (0.80, 2.93) | (0.89, 1.73) | (-5.4%, 26.8%) | | |
| · F | | 66.2%) | | 55.9%) | (, | (*****) | (, , | | |
| Baseline Disease Severity | | , | | , | | | | | |
| Week 12 | | | | | | | | | |
| Moderate baseline | 47 | 25 (53.2%) | 40 | 19 (47.5%) | 1.26 | 1.12 | 5.7% | 0.6694 | 0.4882 |
| disease (IGA=3) | • • | (38.1%, | .0 | (31.5%, | (0.54, 2.92) | (0.73, 1.71) | (-15.4%, 26.7%) | 0.000 | 0.1002 |
| discuse (IGH=3) | | 67.9%) | | 63.9%) | (0.5 1, 2.52) | (0.73, 1.71) | (13.170, 20.770) | | |
| Severe baseline disease | 28 | 16 (57.1%) | 33 | 13 (39.4%) | 2.05 | 1.45 | 17.7% | 0.2037 | |
| (IGA=4) | 20 | (37.2%, | 33 | (22.9%, | | (0.85, 2.47) | (-7.0%, 42.5%) | 0.2037 | |
| (IGA=4) | | , , | | | (0.74, 5.71) | (0.83, 2.47) | (-7.0%, 42.3%) | | |
| G 1 | | 75.5%) | | 57.9%) | | | | | |
| Gender | | | | | | | | | |
| Week 12 | | | | | | | | | |
| Male | 46 | 26 (56.5%) | 33 | 16 (48.5%) | 1.38 | 1.17 | 8.0% | 0.5022 | 0.8854 |
| | | (41.1%, | | (30.8%, | (0.56, 3.39) | (0.76, 1.80) | (-14.2%, 30.3%) | | |
| | | 71.1%) | | 66.5%) | | | | | |
| Female | 29 | 15 (51.7%) | 40 | 16 (40.0%) | 1.61 | 1.29 | 11.7% | 0.4624 | |
| | | (32.5%, | | (24.9%, | (0.61, 4.22) | (0.77, 2.17) | (-12.0%, 35.4%) | | |
| | | 70.6%) | | 56.7%) | , | | , | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------|----------|--------------|----|------------|---------------|--------------|-------------------------|-------------------------------|--------------|
| | | _ | | | | | _ | CMH or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Region | | | | | | | | | |
| Week 12 | | | | | | | | | |
| US/Canada/Australia | 18 | 8 (44.4%) | 21 | 12 (57.1%) | 0.60 | 0.78 | -12.7% | 0.5273 | 0.1577 |
| | | (21.5%, | | (34.0%, | (0.17, 2.14) | (0.41, 1.47) | (-43.9%, 18.5%) | | |
| | | 69.2%) | | 78.2%) | | | | | |
| Europe | 24 | 12 (50.0%) | 24 | 11 (45.8%) | 1.18 | 1.09 | 4.2% | 1.0000 | |
| | | (29.1%, | | (25.6%, | (0.38, 3.67) | (0.60, 1.97) | (-24.1%, 32.4%) | | |
| | | 70.9%) | | 67.2%) | | | | | |
| Asia | 26 | 17 (65.4%) | 23 | 7 (30.4%) | 4.32 | 2.15 | 34.9% | 0.0222* | |
| | | (44.3%, | | (13.2%, | (1.30, 14.34) | (1.09, 4.23) | (8.7%, 61.2%) | | |
| | | 82.8%) | | 52.9%) | | | | | |
| Latin America | 7 | 4 (57.1%) | 5 | 2 (40.0%) | 2.00 | 1.43 | 17.1% | 1.0000 | |
| | | (18.4%, | | (5.3%, | (0.19, 20.61) | (0.41, 4.99) | (-39.3%, 73.6%) | | |
| | | 90.1%) | | 85.3%) | | | | | |
| Weight Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 38 | 21 (55.3%) | 41 | 17 (41.5%) | 1.74 | 1.33 | 13.8% | 0.2635 | 0.7077 |
| | | (38.3%, | | (26.3%, | (0.72, 4.25) | (0.84, 2.12) | (-8.0%, 35.6%) | | |
| | | 71.4%) | | 57.9%) | | | | | |
| > Median Value | 37 | 20 (54.1%) | 32 | 15 (46.9%) | 1.33 | 1.15 | 7.2% | 0.6324 | |
| | | (36.9%, | | (29.1%, | (0.52, 3.44) | (0.72, 1.85) | (-16.4%, 30.8%) | | |
| | | 70.5%) | | 65.3%) | | | | | |
| Age Subgroup | | | | | | | | | |
| Week 12 | | | | | | | | | |
| <= Median Value | 43 | 21 (48.8%) | 42 | 17 (40.5%) | 1.40 | 1.21 | 8.4% | 0.5151 | 0.6272 |
| | | (33.3%, | | (25.6%, | (0.59, 3.31) | (0.75, 1.95) | (-12.7%, 29.4%) | | |
| | | 64.5%) | | 56.7%) | | | | | |
| > Median Value | 32 | 20 (62.5%) | 31 | 15 (48.4%) | 1.78 | 1.29 | 14.1% | 0.3152 | |
| | | (43.7%, | | (30.2%, | (0.65, 4.85) | (0.82, 2.03) | (-10.2%, 38.4%) | | |
| | | 78.9%) | | 66.9%) | | • • • | • | | |

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline EQ5D-VAS >85.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Stratified Cochran-Mantel-Haenszel p-value presented for overall analysis, stratified by by disease activity (moderate, severe) at enrollment.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Total mortality (TEAE leading to death) - Safety Set JADE TEEN $\,$ (PF-04965842) - 2023 datacut

No adverse events of this type occurred

| | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|-----------------------------------|----------|---------------------------------|----|---------------------------------|----------------------------------|----------------------|--------------------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 94 | 59 (62.8%) (52.2%, 72.5%) | 96 | 50 (52.1%) (41.6%, 62.4%) | 1.55 (0.87, 2.77) | 1.21 (0.94, 1.54) | 10.7% (-3.3%, 24.7%) | 0.1341+ | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 37 (60.7%) (47.3%, 72.9%) | 57 | 30 (52.6%) (39.0%, 66.0%) | 1.39 (0.67, 2.88) | 1.15 (0.84, 1.58) | 8.0% (-9.8%, 25.9%) | 0.4578 | 0.5912 | |
| Severe baseline disease (IGA=4) | 33 | 22 (66.7%) (48.2%, 82.0%) | 39 | 20 (51.3%) (34.8%, 67.6%) | 1.90 (0.73, 4.95) | 1.30 (0.88, 1.92) | 15.4% (-7.1%, 37.9%) | 0.2336 | | |
| Gender | | , | | , | | | | | | |
| Male | 56 | 32 (57.1%) (43.2%, 70.3%) | 44 | 22 (50.0%) (34.6%, 65.4%) | 1.33 (0.60, 2.95) | 1.14 (0.79, 1.66) | 7.1% (-12.5%, 26.8%) | 0.5464 | 0.3850 | |
| Female | 38 | 27 (71.1%) (54.1%, 84.6%) | 52 | 28 (53.8%) (39.5%, 67.8%) | 2.10 (0.87, 5.11) | 1.32 (0.96, 1.82) | 17.2% (-2.6%, 37.0%) | 0.1267 | | |
| Region | | , | | , | | | | | | |
| US/Canada/Australia | 28 | 12 (42.9%) (24.5%, 62.8%) | 33 | 12 (36.4%) (20.4%, 54.9%) | 1.31 (0.47, 3.68) | 1.18 (0.63, 2.19) | 6.5% (-18.1%, 31.1%) | 0.7929 | 0.8874 | |
| Europe | 29 | 20 (69.0%) (49.2%, 84.7%) | 27 | 16 (59.3%) (38.8%, 77.6%) | 1.53 (0.51, 4.59) | 1.16 (0.78, 1.73) | 9.7% (-15.3%, 34.7%) | 0.5785 | | |
| Asia | 28 | 22 (78.6%) (59.0%, 91.7%) | 29 | 19 (65.5%) (45.7%, 82.1%) | 1.93 (0.59, 6.30) | 1.20 (0.86, 1.66) | 13.1% (-10.0%, 36.1%) | 0.3786 | | |
| Latin America | 9 | 5 (55.6%) (21.2%, 86.3%) | 7 | 3 (42.9%) (9.9%, 81.6%) | 1.67 (0.23, 12.22) | 1.30 (0.46, 3.65) | 12.7% (-36.3%, 61.7%) | 1.0000 | | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|--------------------|----------|---------------------------------|----|---------------------------------|----------------------|----------------------|-------------------------|----------------------------|--------------------------|
| | | | | | | | <u> </u> | CMH or Logistic I | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Weight Subgroup | | (>=/* ==/ | | (22,22) | (2272 23) | (2272 22) | (3273 23) | [-] | |
| <= Median Value | 46 | 28 (60.9%) (45.4%, 74.9%) | 53 | 31 (58.5%) (44.1%, 71.9%) | 1.10 (0.49, 2.47) | 1.04 (0.75, 1.44) | 2.4% (-17.0%, 21.7%) | 0.8398 | 0.2466 |
| > Median Value | 48 | 31 (64.6%) (49.5%, 77.8%) | 43 | 19 (44.2%) (29.1%, 60.1%) | 2.30 (0.99, 5.36) | 1.46 (0.98, 2.17) | 20.4% (0.3%, 40.5%) | 0.0601 | |
| Age Subgroup | | , | | | | | | | |
| = Median Value | 57 | 35 (61.4%) (47.6%, 74.0%) | 60 | 33 (55.0%) (41.6%, 67.9%) | 1.30 (0.62, 2.72) | 1.12 (0.82, 1.52) | 6.4% (-11.4%, 24.2%) | 0.5746 | 0.4677 |
| > Median Value | 37 | 24 (64.9%) (47.5%, 79.8%) | 36 | 17 (47.2%) (30.4%, 64.5%) | 2.06 (0.81, 5.28) | 1.37 (0.90, 2.09) | 17.6% (-4.8%, 40.1%) | 0.1601 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-0496 | 5842 200mg QD | Placebo | | | Placebo v | s. PF-04965842 200mg QD | | |
|------------------------------------|---------|------------------------------|---------|------------------------------|-----------------------|-----------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 2 (2.1%) (0.3%, 7.5%) | 96 | 2 (2.1%) (0.3%, 7.3%) | 1.02 (0.14, 7.41) | 1.02 (0.15, 7.10) | 0.0% (-4.0%, 4.1%) | 0.9830+ | |
| Baseline Disease Severity | | , , , | | , , , | , , , | , , , , | | | |
| Moderate baseline disease (IGA=3) | 61 | 2 (3.3%) (0.4%, 11.3%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 1.90 (0.17, 21.52) | 1.87 (0.17, 20.05) | 1.5% (-4.1%, 7.1%) | 1.0000 | 0.4155 |
| Severe baseline disease (IGA=4) | 33 | 0 (0.0%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 0.38 (0.02, 9.72) | 0.39 (0.02, 9.32) | -2.3% (-9.4%, 4.9%) | 1.0000 | |
| Gender | | | | , | | | | | |
| Male | 56 | 0 (0.0%) | 44 | 0 (0.0%) | 0.79 (0.02, 40.48) | 0.79 (0.02, 39.02) | -0.2% (-4.1%, 3.7%) | NE | 0.7330 |
| Female | 38 | 2 (5.3%) (0.6%, 17.7%) | 52 | 2 (3.8%) (0.5%, 13.2%) | 1.39 (0.19, 10.33) | 1.37 (0.20, 9.29) | 1.4% (-7.4%, 10.2%) | 1.0000 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) (0.1%, 18.3%) | 33 | 1 (3.0%) (0.1%, 15.8%) | 1.19 (0.07, 19.86) | 1.18 (0.08, 17.99) | 0.5% (-8.5%, 9.6%) | 1.0000 | 0.7489 |
| Europe | 29 | 0 (0.0%) | 27 | 1 (3.7%) (0.1%, 19.0%) | 0.30 (0.01, 7.67) | 0.31 (0.01, 7.33) | -3.7% (-13.2%, 5.8%) | 0.4821 | |
| Asia | 28 | 1 (3.6%) (0.1%, 18.3%) | 29 | 0 (0.0%) | 3.22 (0.13, 82.38) | 3.10 (0.13, 73.12) | 3.5% (-5.8%, 12.8%) | 0.4912 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|--------------------|----------|------------------------------|----|------------------------------|------------------------|-----------------------|-------------------------|-------------------------------|--------------------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Weight Subgroup | | , | | | , | • • • | | | |
| <= Median Value | 46 | 0 (0.0%) | 53 | 2 (3.8%) (0.5%, 13.0%) | 0.22 (0.01, 4.73) | 0.23 (0.01, 4.67) | -3.6% (-9.9%, 2.8%) | 0.4974 | 0.1132 |
| > Median Value | 48 | 2 (4.2%) (0.5%, 14.3%) | 43 | 0 (0.0%) | 4.68 (0.22, 100.20) | 4.49 (0.22, 90.99) | 4.0% (-2.9%, 10.9%) | 0.4960 | |
| Age Subgroup | | | | | | | | | |
| <= Median Value | 57 | 1 (1.8%) (0.0%, 9.4%) | 60 | 2 (3.3%) (0.4%, 11.5%) | 0.52 (0.05, 5.87) | 0.53 (0.05, 5.65) | -1.6% (-7.3%, 4.1%) | 1.0000 | 0.3688 |
| > Median Value | 37 | 1 (2.7%) (0.1%, 14.2%) | 36 | 0 (0.0%) | 3.00 (0.12, 76.09) | 2.92 (0.12, 69.43) | 2.6% (-4.6%, 9.8%) | 1.0000 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|--------------------------------------|---------|------------------------------|----|------------------------------|---------------------------------------|-----------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic 1 Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | · · · · · · · · · · · · · · · · · · · | | | | |
| Safety Set | 94 | 1 (1.1%) (0.0%, 5.8%) | 96 | 2 (2.1%) (0.3%, 7.3%) | 0.51 (0.05, 5.67) | 0.51 (0.05, 5.54) | -1.0% (-4.5%, 2.5%) | 0.5714+ | |
| Baseline Disease Severity | | , , , | | , , , | ` ' ' | , , , | ` ' ' | | |
| Moderate baseline disease (IGA=3) | 61 | 1 (1.6%) (0.0%, 8.8%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 0.93 (0.06, 15.28) | 0.93 (0.06, 14.59) | -0.1% (-4.8%, 4.6%) | 1.0000 | 0.6186 |
| Severe baseline disease (IGA=4) | 33 | 0 (0.0%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 0.38 (0.02, 9.72) | 0.39 (0.02, 9.32) | -2.3% (-9.4%, 4.9%) | 1.0000 | |
| Gender | | | | , | | | | | |
| Male | 56 | 1 (1.8%) (0.0%, 9.6%) | 44 | 1 (2.3%) (0.1%, 12.0%) | 0.78 (0.05, 12.86) | 0.79 (0.05, 12.21) | -0.5% (-6.1%, 5.1%) | 1.0000 | 0.7967 |
| Female | 38 | 0 (0.0%) | 52 | 1 (1.9%) (0.0%, 10.3%) | 0.45 (0.02, 11.25) | 0.45 (0.02, 10.83) | -1.5% (-7.2%, 4.1%) | 1.0000 | |
| Region | | | | , | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) (0.1%, 18.3%) | 33 | 1 (3.0%) (0.1%, 15.8%) | 1.19 (0.07, 19.86) | 1.18 (0.08, 17.99) | 0.5% (-8.5%, 9.6%) | 1.0000 | 0.9113 |
| Europe | 29 | 0 (0.0%) | 27 | 1 (3.7%) (0.1%, 19.0%) | 0.30 (0.01, 7.67) | 0.31 (0.01, 7.33) | -3.7% (-13.2%, 5.8%) | 0.4821 | |
| Asia | 28 | 0 (0.0%) | 29 | 0 (0.0%) | 1.04 (0.02, 53.95) | 1.03 (0.02, 50.42) | 0.1% (-6.5%, 6.6%) | NE | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|---------|------------------------------|----|------------------------------|----------------------------------|-----------------------|-------------------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 0 (0.0%) | 53 | 1 (1.9%) (0.0%, 10.1%) | 0.38 (0.01, 9.47) | 0.38 (0.02, 9.18) | -1.7% (-7.0%, 3.6%) | 1.0000 | 0.7219 | | |
| > Median Value | 48 | 1 (2.1%) (0.1%, 11.1%) | 43 | 1 (2.3%) (0.1%, 12.3%) | 0.89 (0.05, 14.74) | 0.90 (0.06, 13.89) | -0.2% (-6.3%, 5.8%) | 1.0000 | | | |
| Age Subgroup | | | | | | | | | | | |
| <= Median Value | 57 | 1 (1.8%) (0.0%, 9.4%) | 60 | 1 (1.7%) (0.0%, 8.9%) | 1.05 (0.06, 17.25) | 1.05 (0.07, 16.43) | 0.1% (-4.6%, 4.8%) | 1.0000 | 0.5154 | | |
| > Median Value | 37 | 0 (0.0%) | 36 | 1 (2.8%) (0.1%, 14.5%) | 0.32 (0.01, 8.00) | 0.32 (0.01, 7.71) | -2.7% (-10.1%, 4.6%) | 0.4932 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|--------------------------------------|---------|-------------------------------|------|------------------------------|------------------------|-----------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic 1 Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | - 11 | (2570 CI) | - 11 | (5570 C1) | (7370 C1) | (33% C1) | (5570 C1) | value [1] | meraction |
| Safety Set | 94 | 2 (2.1%) (0.3%, 7.5%) | 96 | 2 (2.1%) (0.3%, 7.3%) | 1.02 (0.14, 7.41) | 1.02 (0.15, 7.10) | 0.0% (-4.0%, 4.1%) | 0.9830+ | |
| Baseline Disease Severity | | , , , | | , , , | | , , , | , , , | | |
| Moderate baseline disease (IGA=3) | 61 | 2 (3.3%) (0.4%, 11.3%) | 57 | 0 (0.0%) | 4.83 (0.23, 102.84) | 4.68 (0.23, 95.39) | 3.2% (-2.3%, 8.6%) | 0.4963 | 0.1295 |
| Severe baseline disease (IGA=4) | 33 | 0 (0.0%) | 39 | 2 (5.1%) (0.6%, 17.3%) | 0.22 (0.01, 4.83) | 0.24 (0.01, 4.73) | -4.8% (-13.3%, 3.7%) | 0.4965 | |
| Gender | | | | , | | | | | |
| Male | 56 | 2 (3.6%) (0.4%, 12.3%) | 44 | 1 (2.3%) (0.1%, 12.0%) | 1.59 (0.14, 18.16) | 1.57 (0.15, 16.77) | 1.3% (-5.3%, 7.9%) | 1.0000 | 0.5289 |
| Female | 38 | 0 (0.0%) | 52 | 1 (1.9%) (0.0%, 10.3%) | 0.45 (0.02, 11.25) | 0.45 (0.02, 10.83) | -1.5% (-7.2%, 4.1%) | 1.0000 | |
| Region | | | | , | | | | | |
| US/Canada/Australia | 28 | 0 (0.0%) | 33 | 0 (0.0%) | 1.18 (0.02, 61.15) | 1.17 (0.02, 57.26) | 0.3% (-6.0%, 6.5%) | NE | 0.4550 |
| Europe | 29 | 0 (0.0%) | 27 | 0 (0.0%) | 0.93 (0.02, 48.62) | 0.93 (0.02, 45.47) | -0.1% (-6.8%, 6.6%) | NE | |
| Asia | 28 | 0 (0.0%) | 29 | 2 (6.9%) (0.8%, 22.8%) | 0.19 (0.01, 4.20) | 0.21 (0.01, 4.13) | -6.6% (-17.6%, 4.4%) | 0.4912 | |
| Latin America | 9 | 2 (22.2%) (2.8%, 60.0%) | 7 | 0 (0.0%) | 5.00 (0.20, 122.74) | 4.00 (0.22, 72.01) | 18.8% (-12.9%, 50.4%) | 0.4750 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events-Treatment Emergent Adverse Events leading to therapy discontinuation - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | Placebo | | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|----------|------------------------------|---------|------------------------------|----------------------------------|-----------------------|------------------------|----------------------------|--------------------------|--|--|
| | | | | | | | - | CMH or Logistic I | P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 1 (2.2%) (0.1%, 11.5%) | 53 | 2 (3.8%) (0.5%, 13.0%) | 0.57 (0.05, 6.46) | 0.58 (0.05, 6.15) | -1.6% (-8.2%, 5.0%) | 1.0000 | 0.4345 | | |
| > Median Value | 48 | 1 (2.1%) (0.1%, 11.1%) | 43 | 0 (0.0%) | 2.75 (0.11, 69.24) | 2.69 (0.11, 64.43) | 1.9% (-3.8%, 7.7%) | 1.0000 | | | |
| Age Subgroup | | | | | | | | | | | |
| <= Median Value | 57 | 1 (1.8%) (0.0%, 9.4%) | 60 | 2 (3.3%) (0.4%, 11.5%) | 0.52 (0.05, 5.87) | 0.53 (0.05, 5.65) | -1.6% (-7.3%, 4.1%) | 1.0000 | 0.3688 | | |
| > Median Value | 37 | 1 (2.7%) (0.1%, 14.2%) | 36 | 0 (0.0%) | 3.00 (0.12, 76.09) | 2.92 (0.12, 69.43) | 2.6% (-4.6%, 9.8%) | 1.0000 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|-----------------------------------|----------|---------------------------------|----|---------------------------------|-----------------------|----------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 59 (62.8%) (52.2%, 72.5%) | 96 | 50 (52.1%) (41.6%, 62.4%) | 1.55 (0.87, 2.77) | 1.21 (0.94, 1.54) | 10.7% (-3.3%, 24.7%) | 0.1341+ | |
| Baseline Disease Severity | | , | | , | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 37 (60.7%) (47.3%, 72.9%) | 57 | 30 (52.6%) (39.0%, 66.0%) | 1.39 (0.67, 2.88) | 1.15 (0.84, 1.58) | 8.0% (-9.8%, 25.9%) | 0.4578 | 0.5912 |
| Severe baseline disease (IGA=4) | 33 | 22 (66.7%) (48.2%, 82.0%) | 39 | 20 (51.3%) (34.8%, 67.6%) | 1.90 (0.73, 4.95) | 1.30 (0.88, 1.92) | 15.4% (-7.1%, 37.9%) | 0.2336 | |
| Gender | | 0=10,10, | | , | | | | | |
| Male | 56 | 32 (57.1%) (43.2%, 70.3%) | 44 | 22 (50.0%) (34.6%, 65.4%) | 1.33 (0.60, 2.95) | 1.14 (0.79, 1.66) | 7.1% (-12.5%, 26.8%) | 0.5464 | 0.3850 |
| Female | 38 | 27 (71.1%) (54.1%, 84.6%) | 52 | 28 (53.8%) (39.5%, 67.8%) | 2.10 (0.87, 5.11) | 1.32 (0.96, 1.82) | 17.2% (-2.6%, 37.0%) | 0.1267 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 12 (42.9%) (24.5%, 62.8%) | 33 | 12 (36.4%) (20.4%, 54.9%) | 1.31 (0.47, 3.68) | 1.18 (0.63, 2.19) | 6.5% (-18.1%, 31.1%) | 0.7929 | 0.8874 |
| Europe | 29 | 20 (69.0%) (49.2%, 84.7%) | 27 | 16 (59.3%) (38.8%, 77.6%) | 1.53 (0.51, 4.59) | 1.16 (0.78, 1.73) | 9.7% (-15.3%, 34.7%) | 0.5785 | |
| Asia | 28 | 22 (78.6%) (59.0%, 91.7%) | 29 | 19 (65.5%) (45.7%, 82.1%) | 1.93 (0.59, 6.30) | 1.20 (0.86, 1.66) | 13.1% (-10.0%, 36.1%) | 0.3786 | |
| Latin America | 9 | 5 (55.6%) (21.2%, 86.3%) | 7 | 3 (42.9%) (9.9%, 81.6%) | 1.67 (0.23, 12.22) | 1.30 (0.46, 3.65) | 12.7% (-36.3%, 61.7%) | 1.0000 | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Any Treatment Emergent Adverse Events excluding progression events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|----------|---------------------------------|----|---------------------------------|----------------------------------|----------------------|-------------------------|----------------------------|--------------------------|--|--|
| | | | | | | | • | CMH or Logistic F | P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 28 (60.9%) (45.4%, 74.9%) | 53 | 31 (58.5%) (44.1%, 71.9%) | 1.10 (0.49, 2.47) | 1.04 (0.75, 1.44) | 2.4% (-17.0%, 21.7%) | 0.8398 | 0.2466 | | |
| > Median Value | 48 | 31 (64.6%) (49.5%, 77.8%) | 43 | 19 (44.2%) (29.1%, 60.1%) | 2.30 (0.99, 5.36) | 1.46 (0.98, 2.17) | 20.4% (0.3%, 40.5%) | 0.0601 | | | |
| Age Subgroup | | , | | , | | | | | | | |
| <= Median Value | 57 | 35 (61.4%) (47.6%, 74.0%) | 60 | 33 (55.0%) (41.6%, 67.9%) | 1.30 (0.62, 2.72) | 1.12 (0.82, 1.52) | 6.4% (-11.4%, 24.2%) | 0.5746 | 0.4677 | | |
| > Median Value | 37 | 24 (64.9%) (47.5%, 79.8%) | 36 | 17 (47.2%) (30.4%, 64.5%) | 2.06 (0.81, 5.28) | 1.37 (0.90, 2.09) | 17.6% (-4.8%, 40.1%) | 0.1601 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events excluding progression events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|-----------------------------------|---------|------------------------------|----|------------------------------|-----------------------|-----------------------|--------------------------|------------------------------------|-------------------------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic l Regression p- | P-val [2] for trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | (* * * * * *) | | (*********) | (| , | () | | |
| Safety Set | 94 | 2 (2.1%) (0.3%, 7.5%) | 96 | 2 (2.1%) (0.3%, 7.3%) | 1.02 (0.14, 7.41) | 1.02 (0.15, 7.10) | 0.0% (-4.0%, 4.1%) | 0.9830+ | |
| Baseline Disease Severity | | (0.00,00,000,00) | | (0.070, 7.070) | (*****, *****) | (3122, 1123) | (,, | | |
| Moderate baseline disease (IGA=3) | 61 | 2 (3.3%) (0.4%, 11.3%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 1.90 (0.17, 21.52) | 1.87 (0.17, 20.05) | 1.5% (-4.1%, 7.1%) | 1.0000 | 0.4155 |
| Severe baseline disease (IGA=4) | 33 | 0 (0.0%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 0.38 (0.02, 9.72) | 0.39 (0.02, 9.32) | -2.3% (-9.4%, 4.9%) | 1.0000 | |
| Gender | | | | ŕ | | | | | |
| Male | 56 | 0 (0.0%) | 44 | 0 (0.0%) | 0.79 (0.02, 40.48) | 0.79 (0.02, 39.02) | -0.2% (-4.1%, 3.7%) | NE | 0.7330 |
| Female | 38 | 2 (5.3%) (0.6%, 17.7%) | 52 | 2 (3.8%) (0.5%, 13.2%) | 1.39 (0.19, 10.33) | 1.37 (0.20, 9.29) | 1.4% (-7.4%, 10.2%) | 1.0000 | |
| Region | | | | | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) (0.1%, 18.3%) | 33 | 1 (3.0%) (0.1%, 15.8%) | 1.19 (0.07, 19.86) | 1.18 (0.08, 17.99) | 0.5% (-8.5%, 9.6%) | 1.0000 | 0.7489 |
| Europe | 29 | 0 (0.0%) | 27 | 1 (3.7%) (0.1%, 19.0%) | 0.30 (0.01, 7.67) | 0.31 (0.01, 7.33) | -3.7% (-13.2%, 5.8%) | 0.4821 | |
| Asia | 28 | 1 (3.6%) (0.1%, 18.3%) | 29 | 0 (0.0%) | 3.22 (0.13, 82.38) | 3.10 (0.13, 73.12) | 3.5% (-5.8%, 12.8%) | 0.4912 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events-Severe Treatment Emergent Adverse Events excluding progression events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| - | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|----------|------------------------------|----|------------------------------|----------------------------------|-----------------------|------------------------|----------------------------|--------------------------|--|--|
| | | | | | | | | CMH or Logistic I | P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 0 (0.0%) | 53 | 2 (3.8%) (0.5%, 13.0%) | 0.22 (0.01, 4.73) | 0.23 (0.01, 4.67) | -3.6% (-9.9%, 2.8%) | 0.4974 | 0.1132 | | |
| > Median Value | 48 | 2 (4.2%) (0.5%, 14.3%) | 43 | 0 (0.0%) | 4.68 (0.22, 100.20) | 4.49 (0.22, 90.99) | 4.0% (-2.9%, 10.9%) | 0.4960 | | | |
| Age Subgroup | | | | | | | | | | | |
| = Median Value | 57 | 1 (1.8%) (0.0%, 9.4%) | 60 | 2 (3.3%) (0.4%, 11.5%) | 0.52 (0.05, 5.87) | 0.53 (0.05, 5.65) | -1.6% (-7.3%, 4.1%) | 1.0000 | 0.3688 | | |
| > Median Value | 37 | 1 (2.7%) (0.1%, 14.2%) | 36 | 0 (0.0%) | 3.00 (0.12, 76.09) | 2.92 (0.12, 69.43) | 2.6% (-4.6%, 9.8%) | 1.0000 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events excluding progression events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------------|---------|---------------|----|--------------|---------------|---------------|-------------------------|----------------------------------|--------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 1 (1.1%) | 96 | 2 (2.1%) | 0.51 | 0.51 | -1.0% | 0.5714+ | |
| | | (0.0%, 5.8%) | | (0.3%, 7.3%) | (0.05, 5.67) | (0.05, 5.54) | (-4.5%, 2.5%) | | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline | 61 | 1 (1.6%) | 57 | 1 (1.8%) | 0.93 | 0.93 | -0.1% | 1.0000 | 0.6186 |
| disease (IGA=3) | | (0.0%, 8.8%) | | (0.0%, 9.4%) | (0.06, 15.28) | (0.06, 14.59) | (-4.8%, 4.6%) | | |
| Severe baseline disease | 33 | 0 (0.0%) | 39 | 1 (2.6%) | 0.38 | 0.39 | -2.3% | 1.0000 | |
| (IGA=4) | | | | (0.1%, | (0.02, 9.72) | (0.02, 9.32) | (-9.4%, 4.9%) | | |
| | | | | 13.5%) | | | | | |
| Gender | | | | , | | | | | |
| Male | 56 | 1 (1.8%) | 44 | 1 (2.3%) | 0.78 | 0.79 | -0.5% | 1.0000 | 0.7967 |
| | | (0.0%, 9.6%) | | (0.1%, | (0.05, 12.86) | (0.05, 12.21) | (-6.1%, 5.1%) | | |
| | | | | 12.0%) | | | | | |
| Female | 38 | 0 (0.0%) | 52 | 1 (1.9%) | 0.45 | 0.45 | -1.5% | 1.0000 | |
| | | , , | | (0.0%, | (0.02, 11.25) | (0.02, 10.83) | (-7.2%, 4.1%) | | |
| | | | | 10.3%) | | , , , | , | | |
| Region | | | | , | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) | 33 | 1 (3.0%) | 1.19 | 1.18 | 0.5% | 1.0000 | 0.9113 |
| | | (0.1%, | | (0.1%, | (0.07, 19.86) | (0.08, 17.99) | (-8.5%, 9.6%) | | |
| | | 18.3%) | | 15.8%) | . , , | , , , | ` ' ' | | |
| Europe | 29 | 0 (0.0%) | 27 | 1 (3.7%) | 0.30 | 0.31 | -3.7% | 0.4821 | |
| ı | | - (/ | | (0.1%, | (0.01, 7.67) | (0.01, 7.33) | (-13.2%, 5.8%) | | |
| | | | | 19.0%) | (,, | (, | (,,, | | |
| Asia | 28 | 0 (0.0%) | 29 | 0 (0.0%) | 1.04 | 1.03 | 0.1% | NE | |
| . 2020 | 20 | 0 (0.070) | | 0 (0.070) | (0.02, 53.95) | (0.02, 50.42) | (-6.5%, 6.6%) | 112 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| Julii / iiilci ica | 9 | 0 (0.070) | , | 0 (0.070) | (0.01, 44.65) | (0.02, 36.05) | (-22.8%, 20.3%) | 1417 | |
| | | | | | (0.01, 44.03) | (0.02, 30.03) | (-22.0%, 20.3%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events-Serious Treatment Emergent Adverse Events excluding progression events - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-0496 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|---------|------------------------------|----|------------------------------|----------------------------------|-----------------------|-------------------------|-------------------|---------------|--|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 0 (0.0%) | 53 | 1 (1.9%) (0.0%, 10.1%) | 0.38 (0.01, 9.47) | 0.38 (0.02, 9.18) | -1.7% (-7.0%, 3.6%) | 1.0000 | 0.7219 | | |
| > Median Value | 48 | 1 (2.1%) (0.1%, 11.1%) | 43 | 1 (2.3%) (0.1%, 12.3%) | 0.89 (0.05, 14.74) | 0.90 (0.06, 13.89) | -0.2% (-6.3%, 5.8%) | 1.0000 | | | |
| Age Subgroup | | | | | | | | | | | |
| <= Median Value | 57 | 1 (1.8%) (0.0%, 9.4%) | 60 | 1 (1.7%) (0.0%, 8.9%) | 1.05 (0.06, 17.25) | 1.05 (0.07, 16.43) | 0.1% (-4.6%, 4.8%) | 1.0000 | 0.5154 | | |
| > Median Value | 37 | 0 (0.0%) | 36 | 1 (2.8%) (0.1%, 14.5%) | 0.32 (0.01, 8.00) | 0.32 (0.01, 7.71) | -2.7% (-10.1%, 4.6%) | 0.4932 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Superinfections - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

No adverse events of this type occurred

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Herpes Zoster - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

No adverse events of this type occurred

| | PF-04965842 200mg QD | | Placebo | | | Placebo vs. PF-04965842 200mg QD | | | |
|--------------------------------------|----------------------|------------------------|---------|------------------------------|-----------------------|----------------------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 0 (0.0%) | 96 | 2 (2.1%) (0.3%, 7.3%) | <0.01 (<0.01, NE) | <0.01 (<0.01, NE) | -4.4% (-25.4%, 16.7%) | 0.6846+ | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 0 (0.0%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 0.31 (0.01, 7.67) | 0.31 (0.01, 7.50) | -1.8% (-6.4%, 2.9%) | 0.4831 | 0.9055 |
| Severe baseline disease (IGA=4) | 33 | 0 (0.0%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 0.38 (0.02, 9.72) | 0.39 (0.02, 9.32) | -2.3% (-9.4%, 4.9%) | 1.0000 | |
| Gender | | | | , | | | | | |
| Male | 56 | 0 (0.0%) | 44 | 1 (2.3%) (0.1%, 12.0%) | 0.26 (0.01, 6.45) | 0.26 (0.01, 6.31) | -2.5% (-8.2%, 3.3%) | 0.4400 | 0.8254 |
| Female | 38 | 0 (0.0%) | 52 | 1 (1.9%) (0.0%, 10.3%) | 0.45 (0.02, 11.25) | 0.45 (0.02, 10.83) | -1.5% (-7.2%, 4.1%) | 1.0000 | |
| Region | | | | , | | | | | |
| US/Canada/Australia | 28 | 0 (0.0%) | 33 | 0 (0.0%) | 1.18 (0.02, 61.15) | 1.17 (0.02, 57.26) | 0.3% (-6.0%, 6.5%) | NE | 0.7289 |
| Europe | 29 | 0 (0.0%) | 27 | 0 (0.0%) | 0.93 (0.02, 48.62) | 0.93 (0.02, 45.47) | -0.1% (-6.8%, 6.6%) | NE | |
| Asia | 28 | 0 (0.0%) | 29 | 2 (6.9%) (0.8%, 22.8%) | 0.19 (0.01, 4.20) | 0.21 (0.01, 4.13) | -6.6% (-17.6%, 4.4%) | 0.4912 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Conjunctivitis - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-049658 | 342 200mg QD |] | Placebo | | Placebo vs | . PF-04965842 200mg QD | | |
|-------------------|-----------|--------------|----|------------|---|---------------|------------------------|-------------------------------|--------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| | | Events (%) | | Events (%) | OR | RR | RD | Regression p- | trt*subgroup |
| isit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Veight Subgroup | | | | | | | | | |
| = Median Value | 46 | 0 (0.0%) | 53 | 2 (3.8%) | 0.22 | 0.23 | -3.6% | 0.4974 | 0.3760 |
| | | | | (0.5%, | (0.01, 4.73) | (0.01, 4.67) | (-9.9%, 2.8%) | | |
| | | | | 13.0%) | | , , , | , , , | | |
| Median Value | 48 | 0 (0.0%) | 43 | 0 (0.0%) | 0.90 | 0.90 | -0.1% | NE | |
| | | | | , , | (0.02, 46.17) | (0.02, 44.31) | (-4.3%, 4.1%) | | |
| ge Subgroup | | | | | | | | | |
| = Median Value | 57 | 0 (0.0%) | 60 | 2 (3.3%) | 0.20 | 0.21 | -3.2% | 0.4960 | 0.4160 |
| | | | | (0.4%, | (0.01, 4.33) | (0.01, 4.29) | (-8.8%, 2.3%) | | |
| | | | | 11.5%) | , | · , , , , , , | , , , , , , | | |
| Median Value | 37 | 0 (0.0%) | 36 | 0 (0.0%) | 0.97 | 0.97 | -0.0% | NE | |
| | | | | - (/-/ | (0.02, 50.37) | (0.02, 47.80) | (-5.2%, 5.2%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

| | PF-04965842 200mg QD | | Placebo | | Placebo vs. PF-04965842 200mg QD | | | | | |
|-----------------------------------|----------------------|-------------------------------|---------|-------------------------------|----------------------------------|------------------------|---------------------------|----------------------------|-----------------------------|--|
| | | - | | | | | _ | CMH or Logistic | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 94 | 5 (5.3%) (1.7%, 12.0%) | 96 | 1 (1.0%) (0.0%, 5.7%) | 5.34 (0.61, 46.58) | 5.11 (0.61, 42.89) | 4.3% (-0.7%, 9.2%) | 0.0917+ | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 3 (4.9%) (1.0%, 13.7%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 2.90 (0.29, 28.68) | 2.80 (0.30, 26.18) | 3.2% (-3.2%, 9.6%) | 0.6194 | 0.6039 | |
| Severe baseline disease (IGA=4) | 33 | 2 (6.1%) (0.7%, 20.2%) | 39 | 0 (0.0%) | 6.27 (0.29, 135.37) | 5.88 (0.29, 118.36) | 6.1% (-3.3%, 15.5%) | 0.2066 | | |
| Gender | | , | | | | | | | | |
| Male | 56 | 3 (5.4%) (1.1%, 14.9%) | 44 | 0 (0.0%) | 5.82 (0.29, 115.75) | 5.53 (0.29, 104.25) | 5.0% (-1.9%, 12.0%) | 0.2533 | 0.7608 | |
| Female | 38 | 2 (5.3%) (0.6%, 17.7%) | 52 | 1 (1.9%) (0.0%, 10.3%) | 2.83 (0.25, 32.44) | 2.74 (0.26, 29.09) | 3.3% (-4.7%, 11.4%) | 0.5711 | | |
| Region | | | | | | | | | | |
| US/Canada/Australia | 28 | 0 (0.0%) | 33 | 0 (0.0%) | 1.18 (0.02, 61.15) | 1.17 (0.02, 57.26) | 0.3% (-6.0%, 6.5%) | NE | 0.2121 | |
| Europe | 29 | 1 (3.4%) (0.1%, 17.8%) | 27 | 0 (0.0%) | 2.89 (0.11, 74.15) | 2.80 (0.12, 65.93) | 3.2% (-6.0%, 12.4%) | 1.0000 | | |
| Asia | 28 | 4 (14.3%) (4.0%, 32.7%) | 29 | 0 (0.0%) | 10.84 (0.56, 211.32) | 9.31 (0.52, 165.33) | 13.9% (-0.1%, 27.8%) | 0.0518 | | |
| Latin America | 9 | 0 (0.0%) | 7 | 1 (14.3%) (0.4%, 57.9%) | 0.23 (0.01, 6.52) | 0.27 (0.01, 5.70) | -13.8% (-44.0%, 16.5%) | 0.4375 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value <0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Acne (PT) - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------|----------|------------------------|----|------------------------|---|---|----------------|----------------------------|--------------------------|--|
| | | | | | | | _ | CMH or Logistic F | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Weight Subgroup | | (2370 CI) | | (7570 CI) | (5570 CI) | (2370 C1) | (55% C1) | varae [1] | meraction | |
| | 10 | 2 (6 50() | 52 | 1 (1 00/) | 2.62 | 2.46 | 1.60/ | 0.2240 | 0.0056 | |
| <= Median Value | 46 | 3 (6.5%) | 53 | 1 (1.9%) | 3.63 | 3.46 | 4.6% | 0.3349 | 0.8956 | |
| | | (1.4%, | | (0.0%, | (0.36, 36.15) | (0.37, 32.10) | (-3.4%, 12.7%) | | | |
| | | 17.9%) | | 10.1%) | | | | | | |
| > Median Value | 48 | 2 (4.2%) | 43 | 0(0.0%) | 4.68 | 4.49 | 4.0% | 0.4960 | | |
| | | (0.5%, | | | (0.22, 100.20) | (0.22, 90.99) | (-2.9%, 10.9%) | | | |
| | | 14.3%) | | | (, , , , , , , , , , , , , , , , , , , | (, , , , , , , , , , , , , , , , , , , | (,, | | | |
| Age Subgroup | | 1 110/0) | | | | | | | | |
| <= Median Value | 57 | 4 (7.0%) | 60 | 0 (0.0%) | 10.18 | 9.47 | 6.9% | 0.0533 | 0.1962 | |
| ~= iviculan value | 31 | ` , | 00 | 0 (0.070) | (0.54, 193.44) | (0.52, 171.95) | (-0.3%, 14.2%) | 0.0555 | 0.1702 | |
| | | (1.9%, | | | (0.54, 195.44) | (0.52, 171.95) | (-0.3%, 14.2%) | | | |
| | | 17.0%) | | | | | | | | |
| > Median Value | 37 | 1 (2.7%) | 36 | 1 (2.8%) | 0.97 | 0.97 | -0.1% | 1.0000 | | |
| | | (0.1%, | | (0.1%, | (0.06, 16.16) | (0.06, 14.97) | (-7.6%, 7.4%) | | | |
| | | 14.2%) | | 14.5%) | , , | , , , | | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-0496 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|---------------------------|---------|---------------|----|--------------|-------------------|-----------------|-------------------------|----------------------------------|--------------|
| W. 4/D. 14 | | Events (%) | | Events (%) | OR (OS) | RR (050), GD | RD | CMH or Logistic Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 2 (2.1%) | 96 | 1 (1.0%) | 2.07 | 2.04 | 1.1% | 0.5493+ | |
| | | (0.3%, 7.5%) | | (0.0%, 5.7%) | (0.18, 23.17) | (0.19, 22.15) | (-2.5%, 4.6%) | | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline | 61 | 1 (1.6%) | 57 | 0 (0.0%) | 2.85 | 2.81 | 1.6% | 1.0000 | 0.8146 |
| disease (IGA=3) | | (0.0%, 8.8%) | | | (0.11, 71.43) | (0.12, 67.52) | (-2.9%, 6.1%) | | |
| Severe baseline disease | 33 | 1 (3.0%) | 39 | 1 (2.6%) | 1.19 | 1.18 | 0.5% | 1.0000 | |
| (IGA=4) | | (0.1%, | | (0.1%, | (0.07, 19.75) | (0.08, 18.17) | (-7.2%, 8.1%) | | |
| | | 15.8%) | | 13.5%) | , | , | | | |
| Gender | | , | | , | | | | | |
| Male | 56 | 1 (1.8%) | 44 | 1 (2.3%) | 0.78 | 0.79 | -0.5% | 1.0000 | 0.4201 |
| | | (0.0%, 9.6%) | | (0.1%, | (0.05, 12.86) | (0.05, 12.21) | (-6.1%, 5.1%) | | |
| | | (,, | | 12.0%) | (, | , | (== ==, == ==, | | |
| Female | 38 | 1 (2.6%) | 52 | 0 (0.0%) | 4.20 | 4.08 | 2.9% | 0.4222 | |
| | | (0.1%, | | * (***/*) | (0.17, 105.95) | (0.17, 97.43) | (-3.7%, 9.5%) | | |
| | | 13.8%) | | | (0.0.1, 0.0.15.0) | (3121, 31112) | (211,72,732,72) | | |
| Region | | / | | | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) | 33 | 0 (0.0%) | 3.65 | 3.52 | 3.7% | 0.4590 | 0.9049 |
| | 20 | (0.1%, | | = (=.070) | (0.14, 93.32) | (0.15, 83.07) | (-5.3%, 12.7%) | 01.000 | 0.50.5 |
| | | 18.3%) | | | (0.1., /2.2) | (0.12, 02.07) | (5.570, 12.770) | | |
| Europe | 29 | 0 (0.0%) | 27 | 0 (0.0%) | 0.93 | 0.93 | -0.1% | NE | |
| Lurope | 2) | 0 (0.070) | 21 | 0 (0.070) | (0.02, 48.62) | (0.02, 45.47) | (-6.8%, 6.6%) | T.L | |
| Asia | 28 | 1 (3.6%) | 29 | 1 (3.4%) | 1.04 | 1.04 | 0.1% | 1.0000 | |
| rssia | 46 | (0.1%, | 29 | (0.1%, | (0.06, 17.43) | (0.07, 15.77) | (-9.4%, 9.7%) | 1.0000 | |
| | | 18.3%) | | 17.8%) | (0.00, 17.43) | (0.07, 13.77) | (-9.4%, 9.1%) | | |
| r A | | , | 7 | , | 0.70 | 0.00 | 1.20/ | NIE | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 | 0.80 | -1.3% | NE | |
| | | | | | (0.01, 44.65) | (0.02, 36.05) | (-22.8%, 20.3%) | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events of Special Interest (AESI)-Folliculitis (PT) - Safety Set JADE TEEN $\,$ (PF-04965842) - 2023 datacut

| | PF-049658 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | | |
|--------------------|-----------|--------------------|----|------------------------------|---|---|-------------------------|-------------------|---------------|--|
| | | • | | | | | | CMH or Logistic F | P-val [2] for | |
| V:-:4 / D1-4: | _ | Events (%) | _ | Events (%) | OR | RR | RD | Regression p- | trt*subgroup | |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction | |
| Weight Subgroup | | | | | | | | | | |
| <= Median Value | 46 | 0 (0.0%) | 53 | 1 (1.9%) (0.0%, 10.1%) | 0.38 (0.01, 9.47) | 0.38 (0.02, 9.18) | -1.7% (-7.0%, 3.6%) | 1.0000 | 0.1986 | |
| > Median Value | 48 | 2 (4.2%) (0.5%, | 43 | 0 (0.0%) | 4.68 (0.22, 100.20) | 4.49 (0.22, 90.99) | 4.0% (-2.9%, 10.9%) | 0.4960 | | |
| | | 14.3%) | | | (0.22, 100.20) | (0.22, 90.99) | (-2.9%, 10.9%) | | | |
| Age Subgroup | | | | | | | | | | |
| <= Median Value | 57 | 0 (0.0%) | 60 | 1 (1.7%) (0.0%, 8.9%) | 0.34 (0.01, 8.64) | 0.35 (0.01, 8.43) | -1.6% (-6.2%, 3.0%) | 1.0000 | 0.1649 | |
| > Median Value | 37 | 2 (5.4%) (0.7%, | 36 | 0 (0.0%) | 5.14 (0.24, 110.89) | 4.87 (0.24, 98.02) | 5.2% (-3.5%, 13.9%) | 0.4932 | | |
| | | 18.2%) | | | (, , , , , , , , , , , , , , , , , , , | (, , , , , , , , , , , , , , , , , , , | (, , | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Infections and infestations [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | |
|--------------------------------------|----------|---------------------------------|----|---------------------------------|----------------------------------|----------------------|---------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic 1 Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall Safety Set | 94 | 34 (36.2%) (26.5%, 46.7%) | 96 | 30 (31.3%) (22.2%, 41.5%) | 1.25 (0.68, 2.28) | 1.16 (0.78, 1.73) | 4.9% (-8.5%, 18.3%) | 0.4727+ | |
| Baseline Disease Severity | | 40.770) | | 41.570) | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 19 (31.1%) (19.9%, 44.3%) | 57 | 17 (29.8%) (18.4%, 43.4%) | 1.06 (0.49, 2.33) | 1.04 (0.61, 1.80) | 1.3% (-15.3%, 17.9%) | 1.0000 | 0.4245 |
| Severe baseline disease (IGA=4) | 33 | 15 (45.5%) (28.1%, 63.6%) | 39 | 13 (33.3%) (19.1%, 50.2%) | 1.67 (0.64, 4.33) | 1.36 (0.76, 2.44) | 12.1% (-10.4%, 34.6%) | 0.3380 | |
| Gender | | 0010,0) | | , | | | | | |
| Male | 56 | 18 (32.1%) (20.3%, 46.0%) | 44 | 12 (27.3%) (15.0%, 42.8%) | 1.26 (0.53, 3.01) | 1.18 (0.64, 2.18) | 4.9% (-13.1%, 22.8%) | 0.6638 | 0.8071 |
| Female | 38 | 16 (42.1%) (26.3%, 59.2%) | 52 | 18 (34.6%) (22.0%, 49.1%) | 1.37 (0.58, 3.25) | 1.22 (0.72, 2.06) | 7.5% (-12.8%, 27.8%) | 0.5141 | |
| Region | | | | .,, | | | | | |
| US/Canada/Australia | 28 | 7 (25.0%) (10.7%, 44.9%) | 33 | 6 (18.2%) (7.0%, 35.5%) | 1.50 (0.44, 5.13) | 1.38 (0.52, 3.62) | 6.8% (-13.9%, 27.6%) | 0.5471 | 0.2252 |
| Europe | 29 | 17 (58.6%) (38.9%, 76.5%) | 27 | 10 (37.0%) (19.4%, 57.6%) | 2.41 (0.82, 7.06) | 1.58 (0.89, 2.82) | 21.6% (-4.0%, 47.1%) | 0.1194 | |
| Asia | 28 | 10 (35.7%) (18.6%, 55.9%) | 29 | 13 (44.8%) (26.4%, 64.3%) | 0.68 (0.24, 1.98) | 0.80 (0.42, 1.51) | -9.1% (-34.5%, 16.2%) | 0.5919 | |
| Latin America | 9 | 0 (0.0%) | 7 | 1 (14.3%) (0.4%, 57.9%) | 0.23 (0.01, 6.52) | 0.27 (0.01, 5.70) | -13.8% (-44.0%, 16.5%) | 0.4375 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Infections and infestations [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | | |
|--------------------|----------|---------------------------------|----|---------------------------------|----------------------------------|----------------------|--------------------------|----------------------------|--------------------------|--|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | | |
| Weight Subgroup | | | | | | | | | | | |
| <= Median Value | 46 | 13 (28.3%) (16.0%, 43.5%) | 53 | 20 (37.7%) (24.8%, 52.1%) | 0.65 (0.28, 1.52) | 0.75 (0.42, 1.33) | -9.5% (-27.9%, 9.0%) | 0.3941 | 0.0284* | | |
| > Median Value | 48 | 21 (43.8%) (29.5%, 58.8%) | 43 | 10 (23.3%) (11.8%, 38.6%) | 2.57 (1.03, 6.37) | 1.88 (1.00, 3.53) | 20.5% (1.6%, 39.4%) | 0.0480* | | | |
| Age Subgroup | | , | | | | | | | | | |
| <= Median Value | 57 | 18 (31.6%) (19.9%, 45.2%) | 60 | 21 (35.0%) (23.1%, 48.4%) | 0.86 (0.40, 1.85) | 0.90 (0.54, 1.51) | -3.4% (-20.5%, 13.6%) | 0.8446 | 0.1229 | | |
| > Median Value | 37 | 16 (43.2%) (27.1%, 60.5%) | 36 | 9 (25.0%) (12.1%, 42.2%) | 2.29 (0.84, 6.19) | 1.73 (0.88, 3.40) | 18.2% (-3.1%, 39.6%) | 0.1396 | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Gastrointestinal disorders [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|-----------------------------------|----------|---------------------------------|----|-------------------------------|-----------------------|-----------------------|-------------------------|-------------------------------|--------------|
| | | Events (%) | | Events (%) | OR | RR | RD | CMH or Logistic Regression p- | trt*subgroup |
| Visit / Population | n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | value [1] | interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 22 (23.4%) (15.3%, 33.3%) | 96 | 8 (8.3%) (3.7%, 15.8%) | 3.36 (1.41, 8.00) | 2.81 (1.32, 5.99) | 15.1% (4.9%, 25.3%) | 0.003 | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 15 (24.6%) (14.5%, 37.3%) | 57 | 5 (8.8%) (2.9%, 19.3%) | 3.39 (1.14, 10.06) | 2.80 (1.09, 7.22) | 15.8% (2.8%, 28.9%) | 0.0274* | 0.8087 |
| Severe baseline disease (IGA=4) | 33 | 7 (21.2%) (9.0%, 38.9%) | 39 | 3 (7.7%) (1.6%, 20.9%) | 3.23 (0.76, 13.68) | 2.76 (0.77, 9.82) | 13.5% (-2.7%, 29.8%) | 0.1698 | |
| Gender | | , | | , | | | | | |
| Male | 56 | 11 (19.6%) (10.2%, 32.4%) | 44 | 3 (6.8%) (1.4%, 18.7%) | 3.34 (0.87, 12.82) | 2.88 (0.86, 9.70) | 12.8% (0.0%, 25.6%) | 0.0848 | 0.4915 |
| Female | 38 | 11 (28.9%) (15.4%, 45.9%) | 52 | 5 (9.6%) (3.2%, 21.0%) | 3.83 (1.20, 12.19) | 3.01 (1.14, 7.95) | 19.3% (2.8%, 35.8%) | 0.0252* | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 6 (21.4%) (8.3%, 41.0%) | 33 | 2 (6.1%) (0.7%, 20.2%) | 4.23 (0.78, 22.93) | 3.54 (0.77, 16.15) | 15.4% (-1.9%, 32.6%) | 0.1272 | 0.2652 |
| Europe | 29 | 4 (13.8%) (3.9%, 31.7%) | 27 | 3 (11.1%) (2.4%, 29.2%) | 1.28 (0.26, 6.33) | 1.24 (0.31, 5.05) | 2.7% (-14.6%, 19.9%) | 1.0000 | |
| Asia | 28 | 10 (35.7%) (18.6%, 55.9%) | 29 | 2 (6.9%) (0.8%, 22.8%) | 7.50 (1.47, 38.32) | 5.18 (1.24, 21.57) | 28.8% (8.8%, 48.8%) | 0.0099* | |
| Latin America | 9 | 2 (22.2%) (2.8%, 60.0%) | 7 | 1 (14.3%) (0.4%, 57.9%) | 1.71 (0.12, 23.94) | 1.56 (0.17, 13.87) | 7.9% (-29.6%, 45.5%) | 1.0000 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Gastrointestinal disorders [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo vs | . PF-04965842 200mg QD | | |
|--------------------|----------|---------------------------------|----|-------------------------------|-----------------------|-----------------------|-------------------------|-------------------------------|--------------------------|
| | | | | | | | | CMH or Logistic P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Weight Subgroup | | | | | | | | | |
| <= Median Value | 46 | 12 (26.1%) (14.3%, 41.1%) | 53 | 4 (7.5%) (2.1%, 18.2%) | 4.32 (1.29, 14.55) | 3.46 (1.20, 9.98) | 18.5% (4.0%, 33.1%) | 0.0150* | 0.4991 |
| > Median Value | 48 | 10 (20.8%) (10.5%, 35.0%) | 43 | 4 (9.3%) (2.6%, 22.1%) | 2.57 (0.74, 8.89) | 2.24 (0.76, 6.62) | 11.5% (-2.9%, 25.9%) | 0.1544 | |
| Age Subgroup | | | | | | | | | |
| <= Median Value | 57 | 14 (24.6%) (14.1%, 37.8%) | 60 | 6 (10.0%) (3.8%, 20.5%) | 2.93 (1.04, 8.26) | 2.46 (1.01, 5.95) | 14.6% (1.1%, 28.1%) | 0.0492* | 0.9383 |
| > Median Value | 37 | 8 (21.6%) (9.8%, 38.2%) | 36 | 2 (5.6%) (0.7%, 18.7%) | 4.69 (0.92, 23.86) | 3.89 (0.89, 17.10) | 16.1% (0.8%, 31.3%) | 0.0854 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Nervous system disorders [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo vs. PF-04965842 200mg QD | | | | |
|--------------------------------------|----------|--------------------------------|----|-------------------------------|------------------------|----------------------------------|--------------------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic 1 Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 94 | 14 (14.9%) (8.4%, 23.7%) | 96 | 9 (9.4%) (4.4%, 17.1%) | 1.69 (0.69, 4.12) | 1.59 (0.72, 3.49) | 5.5% (-3.7%, 14.8%) | 0.2429+ | | |
| Baseline Disease Severity | | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 9 (14.8%) (7.0%, | 57 | 6 (10.5%) (4.0%, | 1.47 (0.49, 4.43) | 1.40 (0.53, 3.69) | 4.2% (-7.7%, 16.2%) | 0.5855 | 0.7462 | |
| Severe baseline disease (IGA=4) | 33 | 26.2%) 5 (15.2%) (5.1%, | 39 | 21.5%) 3 (7.7%) (1.6%, | 2.14 (0.47, 9.74) | 1.97 (0.51, 7.63) | 7.5% (-7.4%, 22.3%) | 0.4563 | | |
| Gender | | 31.9%) | | 20.9%) | | | | | | |
| Male | 56 | 8 (14.3%) (6.4%, 26.2%) | 44 | 3 (6.8%) (1.4%, 18.7%) | 2.28 (0.57, 9.15) | 2.10 (0.59, 7.44) | 7.5% (-4.3%, 19.3%) | 0.3384 | 0.7567 | |
| Female | 38 | 6 (15.8%) (6.0%, 31.3%) | 52 | 6 (11.5%) (4.4%, 23.4%) | 1.44 (0.43, 4.86) | 1.37 (0.48, 3.92) | 4.3% (-10.2%, 18.7%) | 0.7548 | | |
| Region | | , | | | | | | | | |
| US/Canada/Australia | 28 | 2 (7.1%) (0.9%, 23.5%) | 33 | 3 (9.1%) (1.9%, 24.3%) | 0.77 (0.12, 4.96) | 0.79 (0.14, 4.37) | -1.9% (-15.6%, 11.7%) | 1.0000 | 0.0313* | |
| Europe | 29 | 1 (3.4%) (0.1%, 17.8%) | 27 | 3 (11.1%) (2.4%, 29.2%) | 0.29 (0.03, 2.93) | 0.31 (0.03, 2.81) | -7.7% (-21.3%, 5.9%) | 0.3434 | | |
| Asia | 28 | 8 (28.6%) (13.2%, 48.7%) | 29 | 1 (3.4%) (0.1%, 17.8%) | 11.20 (1.30, 96.79) | 8.29 (1.11, 62.02) | 25.1% (7.1%, 43.1%) | 0.0119* | | |
| Latin America | 9 | 3 (33.3%) (7.5%, 70.1%) | 7 | 2 (28.6%) (3.7%, 71.0%) | 1.25 (0.15, 10.70) | 1.17 (0.26, 5.19) | 4.8% (-40.7%, 50.2%) | 1.0000 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Nervous system disorders [SOC] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------|----------|-------------------------------|----|-------------------------------|----------------------------------|-----------------------|--------------------------|----------------------------|--------------------------|--|
| | | | | | | | • | CMH or Logistic I | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Weight Subgroup | | | | , , , | , , | , | , , | | | |
| <= Median Value | 46 | 7 (15.2%) (6.3%, 28.9%) | 53 | 6 (11.3%) (4.3%, 23.0%) | 1.41 (0.44, 4.53) | 1.34 (0.49, 3.71) | 3.9% (-9.5%, 17.3%) | 0.7666 | 0.7085 | |
| > Median Value | 48 | 7 (14.6%) (6.1%, 27.8%) | 43 | 3 (7.0%) (1.5%, 19.1%) | 2.28 (0.55, 9.43) | 2.09 (0.58, 7.58) | 7.6% (-5.0%, 20.2%) | 0.3233 | | |
| Age Subgroup | | | | | | | | | | |
| <= Median Value | 57 | 6 (10.5%) (4.0%, 21.5%) | 60 | 7 (11.7%) (4.8%, 22.6%) | 0.89 (0.28, 2.83) | 0.90 (0.32, 2.52) | -1.1% (-12.5%, 10.2%) | 1.0000 | 0.0748 | |
| > Median Value | 37 | 8 (21.6%) (9.8%, 38.2%) | 36 | 2 (5.6%) (0.7%, 18.7%) | 4.69 (0.92, 23.86) | 3.89 (0.89, 17.10) | 16.1% (0.8%, 31.3%) | 0.0854 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo vs | s. PF-04965842 200mg QD | | |
|-----------------------------------|----------|---------------------------------|----|-------------------------------|-------------------------|-------------------------|-------------------------|----------------------------|--------------------------|
| | | | | | | | | CMH or Logistic 1 | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | (2070-0-5) | | (2272-02) | (2272-22) | (2011-00) | (>= /* = -/ | [-] | |
| Safety Set | 94 | 17 (18.1%) (10.9%, 27.4%) | 96 | 1 (1.0%) (0.0%, 5.7%) | 20.97 (2.73, >99.99) | 17.36 (2.36, >99.99) | 17.0% (9.0%, 25.1%) | <0.0001+* | |
| Baseline Disease Severity | | , | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 11 (18.0%) (9.4%, 30.0%) | 57 | 1 (1.8%) (0.0%, 9.4%) | 12.32 (1.54, 98.85) | 10.28 (1.37, 77.10) | 16.3% (6.0%, 26.5%) | 0.0044* | 0.8605 |
| Severe baseline disease (IGA=4) | 33 | 6 (18.2%) (7.0%, 35.5%) | 39 | 0 (0.0%) | 18.67 (1.01, 345.30) | 15.29 (0.89, 261.74) | 17.9% (4.2%, 31.5%) | 0.0071* | |
| Gender | | | | | | | | | |
| Male | 56 | 8 (14.3%) (6.4%, 26.2%) | 44 | 0 (0.0%) | 15.60 (0.87, 278.15) | 13.42 (0.80, 226.35) | 13.8% (4.1%, 23.5%) | 0.0086* | 0.3386 |
| Female | 38 | 9 (23.7%) (11.4%, 40.2%) | 52 | 1 (1.9%) (0.0%, 10.3%) | 15.83 (1.91, 131.29) | 12.32 (1.63, 93.14) | 21.8% (7.7%, 35.8%) | 0.0016* | |
| Region | | , | | | | | | | |
| US/Canada/Australia | 28 | 3 (10.7%) (2.3%, 28.2%) | 33 | 0 (0.0%) | 9.20 (0.45, 186.12) | 8.21 (0.44, 152.39) | 10.6% (-1.9%, 23.1%) | 0.0910 | 0.0582 |
| Europe | 29 | 2 (6.9%) (0.8%, 22.8%) | 27 | 0 (0.0%) | 5.00 (0.23, 109.01) | 4.67 (0.23, 93.02) | 6.5% (-4.5%, 17.6%) | 0.4916 | |
| Asia | 28 | 10 (35.7%) (18.6%, 55.9%) | 29 | 0 (0.0%) | 33.49 (1.85, 606.14) | 21.72 (1.33, 353.95) | 34.5% (16.5%, 52.6%) | 0.0003* | |
| Latin America | 9 | 2 (22.2%) (2.8%, 60.0%) | 7 | 1 (14.3%) (0.4%, 57.9%) | 1.71 (0.12, 23.94) | 1.56 (0.17, 13.87) | 7.9% (-29.6%, 45.5%) | 1.0000 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Nausea [PT] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------|----------|---------------------------------|----|------------------------------|----------------------------------|-------------------------|------------------------|----------------------------|--------------------------|--|
| | | | | | | | • | CMH or Logistic I | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Weight Subgroup | | | | | | | | | | |
| <= Median Value | 46 | 10 (21.7%) (10.9%, 36.4%) | 53 | 0 (0.0%) | 30.78 (1.75, 541.83) | 24.13 (1.45, 400.74) | 21.4% (9.2%, 33.6%) | 0.0003* | 0.2778 | |
| > Median Value | 48 | 7 (14.6%) (6.1%, 27.8%) | 43 | 1 (2.3%) (0.1%, 12.3%) | 7.17 (0.84, 60.89) | 6.27 (0.80, 48.93) | 12.3% (1.3%, 23.2%) | 0.0616 | | |
| Age Subgroup | | , | | | | | | | | |
| <= Median Value | 57 | 11 (19.3%) (10.0%, 31.9%) | 60 | 1 (1.7%) (0.0%, 8.9%) | 14.11 (1.76, 113.28) | 11.58 (1.54, 86.83) | 17.6% (6.9%, 28.4%) | 0.0017* | 0.8169 | |
| > Median Value | 37 | 6 (16.2%) (6.2%, 32.0%) | 36 | 0 (0.0%) | 15.06 (0.82, 278.09) | 12.66 (0.74, 216.78) | 15.8% (3.2%, 28.3%) | 0.0251* | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Upper respiratory tract infection [PT] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 5842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|------------------------------------|----------|--------------------------------|----|--------------------------------|------------------------|------------------------|--------------------------|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | (2272 23) | | (2272 22) | (*********) | (20,000) | (22,12,22) | [-] | |
| Safety Set | 94 | 10 (10.6%) (5.2%, 18.7%) | 96 | 10 (10.4%) (5.1%, 18.3%) | 1.02 (0.41, 2.59) | 1.02 (0.45, 2.34) | 0.2% (-8.5%, 9.0%) | 0.9603+ | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 5 (8.2%) (2.7%, 18.1%) | 57 | 4 (7.0%) (1.9%, 17.0%) | 1.18 (0.30, 4.64) | 1.17 (0.33, 4.14) | 1.2% (-8.4%, 10.7%) | 1.0000 | 0.8914 |
| Severe baseline disease (IGA=4) | 33 | 5 (15.2%) (5.1%, 31.9%) | 39 | 6 (15.4%) (5.9%, 30.5%) | 0.98 (0.27, 3.57) | 0.98 (0.33, 2.94) | -0.2% (-16.9%, 16.4%) | 1.0000 | |
| Gender | | , | | , | | | | | |
| Male | 56 | 4 (7.1%) (2.0%, 17.3%) | 44 | 2 (4.5%) (0.6%, 15.5%) | 1.62 (0.28, 9.25) | 1.57 (0.30, 8.19) | 2.6% (-6.5%, 11.7%) | 0.6920 | 0.8279 |
| Female | 38 | 6 (15.8%) (6.0%, 31.3%) | 52 | 8 (15.4%) (6.9%, 28.1%) | 1.03 (0.33, 3.26) | 1.03 (0.39, 2.71) | 0.4% (-14.8%, 15.6%) | 1.0000 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 2 (7.1%) (0.9%, 23.5%) | 33 | 0 (0.0%) | 6.32 (0.29, 137.37) | 5.86 (0.29, 117.23) | 7.2% (-3.8%, 18.1%) | 0.2066 | 0.6428 |
| Europe | 29 | 4 (13.8%) (3.9%, 31.7%) | 27 | 5 (18.5%) (6.3%, 38.1%) | 0.70 (0.17, 2.95) | 0.74 (0.22, 2.49) | -4.7% (-24.0%, 14.6%) | 0.7249 | |
| Asia | 28 | 4 (14.3%) (4.0%, 32.7%) | 29 | 5 (17.2%) (5.8%, 35.8%) | 0.80 (0.19, 3.35) | 0.83 (0.25, 2.77) | -3.0% (-21.9%, 15.9%) | 1.0000 | |
| Latin America | 9 | 0 (0.0%) | 7 | 0 (0.0%) | 0.79 (0.01, 44.65) | 0.80 (0.02, 36.05) | -1.3% (-22.8%, 20.3%) | NE | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value <0.05

Binary Outcome Analysis: Adverse Events by SOC/PT-Any Treatment Emergent Adverse Events - Upper respiratory tract infection [PT] - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------|----------|-------------------------------|----|-------------------------------|----------------------------------|-----------------------|--------------------------|----------------------------|--------------------------|--|
| | | | | | | | • | CMH or Logistic 1 | P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| Weight Subgroup | | , | | , | ` | | | | | |
| <= Median Value | 46 | 5 (10.9%) (3.6%, 23.6%) | 53 | 7 (13.2%) (5.5%, 25.3%) | 0.80 (0.24, 2.72) | 0.82 (0.28, 2.42) | -2.3% (-15.1%, 10.5%) | 0.7674 | 0.5153 | |
| > Median Value | 48 | 5 (10.4%) (3.5%, 22.7%) | 43 | 3 (7.0%) (1.5%, 19.1%) | 1.55 (0.35, 6.91) | 1.49 (0.38, 5.88) | 3.4% (-8.1%, 15.0%) | 0.7174 | | |
| Age Subgroup | | | | | | | | | | |
| <= Median Value | 57 | 5 (8.8%) (2.9%, 19.3%) | 60 | 8 (13.3%) (5.9%, 24.6%) | 0.63 (0.19, 2.04) | 0.66 (0.23, 1.89) | -4.6% (-15.9%, 6.7%) | 0.5598 | 0.1605 | |
| > Median Value | 37 | 5 (13.5%) (4.5%, 28.8%) | 36 | 2 (5.6%) (0.7%, 18.7%) | 2.66 (0.48, 14.68) | 2.43 (0.50, 11.74) | 8.0% (-5.4%, 21.3%) | 0.4297 | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (PT)-PT Natural killer cell count decreased - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

No adverse events of this type occurred

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | D | |
|---------------------------|----------|-------------------------------|----|-------------------------------|------------------------|------------------------|--------------------------|---|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction |
| Overall | | | | | | | | | |
| Safety Set | 94 | 8 (8.5%) (3.7%, 16.1%) | 96 | 7 (7.3%) (3.0%, 14.4%) | 1.18 (0.41, 3.40) | 1.17 (0.44, 3.09) | 1.2% (-6.5%, 8.9%) | 0.7555+ | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline | 61 | 6 (9.8%) | 57 | 5 (8.8%) | 1.13 | 1.12 | 1.1% | 1.0000 | 0.9821 |
| disease (IGA=3) | | (3.7%, 20.2%) | | (2.9%, 19.3%) | (0.33, 3.94) | (0.36, 3.47) | (-9.4%, 11.5%) | | |
| Severe baseline disease | 33 | 2 (6.1%) | 39 | 2 (5.1%) | 1.19 | 1.18 | 0.9% | 1.0000 | |
| (IGA=4) | | (0.7%, 20.2%) | | (0.6%, 17.3%) | (0.16, 8.97) | (0.18, 7.94) | (-9.8%, 11.6%) | | |
| Gender | | · · | | ŕ | | | | | |
| Male | 56 | 4 (7.1%) (2.0%, 17.3%) | 44 | 1 (2.3%) (0.1%, 12.0%) | 3.31 (0.36, 30.71) | 3.14 (0.36, 27.13) | 4.9% (-3.2%, 12.9%) | 0.3808 | 0.4782 |
| Female | 38 | 4 (10.5%) (2.9%, 24.8%) | 52 | 6 (11.5%) (4.4%, 23.4%) | 0.90 (0.24, 3.45) | 0.91 (0.28, 3.01) | -1.0% (-14.1%, 12.0%) | 1.0000 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 2 (7.1%) (0.9%, 23.5%) | 33 | 3 (9.1%) (1.9%, 24.3%) | 0.77 (0.12, 4.96) | 0.79 (0.14, 4.37) | -1.9% (-15.6%, 11.7%) | 1.0000 | 0.5578 |
| Europe | 29 | 1 (3.4%) (0.1%, 17.8%) | 27 | 2 (7.4%) (0.9%, 24.3%) | 0.45 (0.04, 5.23) | 0.47 (0.04, 4.84) | -4.0% (-15.9%, 7.9%) | 0.6045 | |
| Asia | 28 | 2 (7.1%) (0.9%, 23.5%) | 29 | 0 (0.0%) | 5.57 (0.26, 121.27) | 5.17 (0.26, 103.18) | 7.0% (-4.2%, 18.1%) | 0.2368 | |
| Latin America | 9 | 3 (33.3%) (7.5%, 70.1%) | 7 | 2 (28.6%) (3.7%, 71.0%) | 1.25 (0.15, 10.70) | 1.17 (0.26, 5.19) | 4.8% (-40.7%, 50.2%) | 1.0000 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Selected Adverse Events (PT)-PT Headache - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------|----------|------------------------|----|------------------------|----------------------------------|----------------|-----------------|----------------------------|--------------------------|--|
| | | | | | | | | CMH or Logistic F | P-val [2] for | |
| Visit / Population | | Events (%) (95% CI) | | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction | |
| | n | (93% CI) | n | (93% CI) | (93% CI) | (93% CI) | (93% CI) | value [1] | meracuon | |
| Weight Subgroup | | | | | | | | | | |
| <= Median Value | 46 | 3 (6.5%) | 53 | 4 (7.5%) | 0.85 | 0.86 | -1.0% | 1.0000 | 0.5678 | |
| | | (1.4%, | | (2.1%, | (0.18, 4.03) | (0.20, 3.66) | (-11.1%, 9.0%) | | | |
| | | 17.9%) | | 18.2%) | | | | | | |
| > Median Value | 48 | 5 (10.4%) | 43 | 3 (7.0%) | 1.55 | 1.49 | 3.4% | 0.7174 | | |
| | | (3.5%, | | (1.5%, | (0.35, 6.91) | (0.38, 5.88) | (-8.1%, 15.0%) | | | |
| | | 22.7%) | | 19.1%) | (0.55, 0.51) | (0.50, 5.00) | (0.170, 10.070) | | | |
| Age Subgroup | | 22.770) | | 17.170) | | | | | | |
| <= Median Value | 57 | 5 (8.8%) | 60 | 5 (8.3%) | 1.06 | 1.05 | 0.4% | 1.0000 | 0.7901 | |
| = Median value | 37 | ` / | 60 | ` / | | | | 1.0000 | 0.7901 | |
| | | (2.9%, | | (2.8%, | (0.29, 3.87) | (0.32, 3.44) | (-9.7%, 10.6%) | | | |
| | | 19.3%) | | 18.4%) | | | | | | |
| Median Value | 37 | 3 (8.1%) | 36 | 2 (5.6%) | 1.50 | 1.46 | 2.6% | 1.0000 | | |
| | | (1.7%, | | (0.7%, | (0.24, 9.55) | (0.26, 8.23) | (-9.0%, 14.1%) | | | |
| | | 21.9%) | | 18.7%) | , , , , , | , , , , , , | , , , , , , , , | | | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-04965 | 5842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | | |
|--------------------------------------|----------|--------------------------------|----|-------------------------------|----------------------------------|-----------------------|--------------------------|---|--|--|
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | CMH or Logistic I Regression p- value [1] | P-val [2] for trt*subgroup interaction | |
| Overall | | | | | | | | | | |
| Safety Set | 94 | 14 (14.9%) (8.4%, 23.7%) | 96 | 9 (9.4%) (4.4%, 17.1%) | 1.69 (0.69, 4.12) | 1.59 (0.72, 3.49) | 5.5% (-3.7%, 14.8%) | 0.2429+ | | |
| Baseline Disease Severity | | , | | , | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 9 (14.8%) (7.0%, 26.2%) | 57 | 6 (10.5%) (4.0%, 21.5%) | 1.47 (0.49, 4.43) | 1.40 (0.53, 3.69) | 4.2% (-7.7%, 16.2%) | 0.5855 | 0.7462 | |
| Severe baseline disease (IGA=4) | 33 | 5 (15.2%) (5.1%, 31.9%) | 39 | 3 (7.7%) (1.6%, 20.9%) | 2.14 (0.47, 9.74) | 1.97 (0.51, 7.63) | 7.5% (-7.4%, 22.3%) | 0.4563 | | |
| Gender | | 31.570) | | 20.570) | | | | | | |
| Male | 56 | 8 (14.3%) (6.4%, 26.2%) | 44 | 3 (6.8%) (1.4%, 18.7%) | 2.28 (0.57, 9.15) | 2.10 (0.59, 7.44) | 7.5% (-4.3%, 19.3%) | 0.3384 | 0.7567 | |
| Female | 38 | 6 (15.8%) (6.0%, 31.3%) | 52 | 6 (11.5%) (4.4%, 23.4%) | 1.44 (0.43, 4.86) | 1.37 (0.48, 3.92) | 4.3% (-10.2%, 18.7%) | 0.7548 | | |
| Region | | , | | | | | | | | |
| US/Canada/Australia | 28 | 2 (7.1%) (0.9%, 23.5%) | 33 | 3 (9.1%) (1.9%, 24.3%) | 0.77 (0.12, 4.96) | 0.79 (0.14, 4.37) | -1.9% (-15.6%, 11.7%) | 1.0000 | 0.0313* | |
| Europe | 29 | 1 (3.4%) (0.1%, 17.8%) | 27 | 3 (11.1%) (2.4%, 29.2%) | 0.29 (0.03, 2.93) | 0.31 (0.03, 2.81) | -7.7% (-21.3%, 5.9%) | 0.3434 | | |
| Asia | 28 | 8 (28.6%) (13.2%, 48.7%) | 29 | 1 (3.4%) (0.1%, 17.8%) | 11.20 (1.30, 96.79) | 8.29 (1.11, 62.02) | 25.1% (7.1%, 43.1%) | 0.0119* | | |
| Latin America | 9 | 3 (33.3%) (7.5%, 70.1%) | 7 | 2 (28.6%) (3.7%, 71.0%) | 1.25 (0.15, 10.70) | 1.17 (0.26, 5.19) | 4.8% (-40.7%, 50.2%) | 1.0000 | | |

Notes

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[*] p-value < 0.05

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

Binary Outcome Analysis: Selected Adverse Events (PT)-SOC Nervous system disorders - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | Placebo vs. PF-04965842 200mg QD | | | | |
|--------------------|----------|-------------------------------|----|-------------------------------|----------------------------------|-----------------------|--------------------------|----------------------------|--------------------------|
| | | | | | | | | CMH or Logistic F | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Weight Subgroup | | | | | · · · · · · | | | | |
| <= Median Value | 46 | 7 (15.2%) (6.3%, 28.9%) | 53 | 6 (11.3%) (4.3%, 23.0%) | 1.41 (0.44, 4.53) | 1.34 (0.49, 3.71) | 3.9% (-9.5%, 17.3%) | 0.7666 | 0.7085 |
| > Median Value | 48 | 7 (14.6%) (6.1%, 27.8%) | 43 | 3 (7.0%) (1.5%, 19.1%) | 2.28 (0.55, 9.43) | 2.09 (0.58, 7.58) | 7.6% (-5.0%, 20.2%) | 0.3233 | |
| Age Subgroup | | · · | | | | | | | |
| <= Median Value | 57 | 6 (10.5%) (4.0%, 21.5%) | 60 | 7 (11.7%) (4.8%, 22.6%) | 0.89 (0.28, 2.83) | 0.90 (0.32, 2.52) | -1.1% (-12.5%, 10.2%) | 1.0000 | 0.0748 |
| > Median Value | 37 | 8 (21.6%) (9.8%, 38.2%) | 36 | 2 (5.6%) (0.7%, 18.7%) | 4.69 (0.92, 23.86) | 3.89 (0.89, 17.10) | 16.1% (0.8%, 31.3%) | 0.0854 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|-----------------------------------|----------|-------------------------------|----|---|-----------------------|-----------------------|---------------------------|-------------------------------|-----------------------------|
| | | • | | | | | | CMH or Logistic P-val [2] for | |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Overall | | | | , | , , | , | | | |
| Safety Set | 94 | 9 (9.6%) (4.5%, 17.4%) | 96 | 7 (7.3%) (3.0%, 14.4%) | 1.35 (0.48, 3.78) | 1.31 (0.51, 3.38) | 2.3% (-5.6%, 10.2%) | 0.5712+ | |
| Baseline Disease Severity | | | | | | | | | |
| Moderate baseline disease (IGA=3) | 61 | 6 (9.8%) (3.7%, 20.2%) | 57 | 6 (10.5%) (4.0%, 21.5%) | 0.93 (0.28, 3.06) | 0.93 (0.32, 2.73) | -0.7% (-11.6%, 10.2%) | 1.0000 | 0.3635 |
| Severe baseline disease (IGA=4) | 33 | 3 (9.1%) (1.9%, 24.3%) | 39 | 1 (2.6%) (0.1%, 13.5%) | 3.80 (0.38, 38.41) | 3.55 (0.39, 32.49) | 6.5% (-4.5%, 17.5%) | 0.3266 | |
| Gender | | 2.1.570) | | 15.570) | | | | | |
| Male | 56 | 5 (8.9%) (3.0%, 19.6%) | 44 | 3 (6.8%) (1.4%, 18.7%) | 1.34 (0.30, 5.94) | 1.31 (0.33, 5.18) | 2.1% (-8.4%, 12.7%) | 1.0000 | 0.9266 |
| Female | 38 | 4 (10.5%) (2.9%, 24.8%) | 52 | 4 (7.7%) (2.1%, 18.5%) | 1.41 (0.33, 6.04) | 1.37 (0.37, 5.13) | 2.8% (-9.3%, 15.0%) | 0.7172 | |
| Region | | , | | , | | | | | |
| US/Canada/Australia | 28 | 1 (3.6%) (0.1%, 18.3%) | 33 | 2 (6.1%) (0.7%, 20.2%) | 0.57 (0.05, 6.69) | 0.59 (0.06, 6.16) | -2.5% (-13.1%, 8.2%) | 1.0000 | 0.4491 |
| Europe | 29 | 2 (6.9%) (0.8%, 22.8%) | 27 | 1 (3.7%) (0.1%, 19.0%) | 1.93 (0.16, 22.55) | 1.86 (0.18, 19.38) | 3.2% (-8.5%, 14.8%) | 1.0000 | |
| Asia | 28 | 6 (21.4%) (8.3%, 41.0%) | 29 | 3 (10.3%) (2.2%, 27.4%) | 2.36 (0.53, 10.57) | 2.07 (0.57, 7.49) | 11.1% (-7.7%, 29.9%) | 0.2973 | |
| Latin America | 9 | 0 (0.0%) | 7 | 1 (14.3%) (0.4%, 57.9%) | 0.23 (0.01, 6.52) | 0.27 (0.01, 5.70) | -13.8% (-44.0%, 16.5%) | 0.4375 | |

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point. The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

^[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

^[*] p-value < 0.05

Binary Outcome Analysis: Selected Adverse Events (SOC)-SOC Skin and subcutaneous tissue disorders - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965 | 842 200mg QD | | Placebo | | Placebo v | s. PF-04965842 200mg QD | | |
|--------------------|----------|-------------------------------|----|------------------------------|-----------------------|-----------------------|--------------------------|----------------------------|--------------------------|
| | | | | | | | • | CMH or Logistic F | P-val [2] for |
| Visit / Population | n | Events (%) (95% CI) | n | Events (%) (95% CI) | OR (95% CI) | RR (95% CI) | RD (95% CI) | Regression p- value [1] | trt*subgroup interaction |
| Weight Subgroup | | | | | | | | | |
| <= Median Value | 46 | 4 (8.7%) (2.4%, 20.8%) | 53 | 5 (9.4%) (3.1%, 20.7%) | 0.91 (0.23, 3.63) | 0.92 (0.26, 3.23) | -0.7% (-12.1%, 10.6%) | 1.0000 | 0.4199 |
| > Median Value | 48 | 5 (10.4%) (3.5%, 22.7%) | 43 | 2 (4.7%) (0.6%, 15.8%) | 2.38 (0.44, 12.98) | 2.24 (0.46, 10.95) | 5.8% (-4.9%, 16.5%) | 0.4396 | |
| Age Subgroup | | · · | | | | | | | |
| <= Median Value | 57 | 8 (14.0%) (6.3%, 25.8%) | 60 | 4 (6.7%) (1.8%, 16.2%) | 2.29 (0.65, 8.06) | 2.11 (0.67, 6.61) | 7.4% (-3.6%, 18.4%) | 0.2317 | 0.0885 |
| > Median Value | 37 | 1 (2.7%) (0.1%, 14.2%) | 36 | 3 (8.3%) (1.8%, 22.5%) | 0.31 (0.03, 3.08) | 0.32 (0.04, 2.97) | -5.6% (-16.1%, 4.8%) | 0.3575 | |

Notes:

Number of subjects: Safety Set

Analysis on overall population is calculated based on unstratified models, for OR, RR, and RD, using generalized linear models with logit, log, and identity link, respectively.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

[+] Unstratified Wald p-value for the risk difference, from a generalized linear model with identity link.

[*] p-value < 0.05

Safety incidence analysis: Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| System Organ Class (SOC) | PF-04965842 200mg QD | Placebo | Total Population |
|-----------------------------------|----------------------|-----------|------------------|
| Preferred Term (PT) | N=94 | N=96 | N=190 |
| Overall | 2 (2.1%) | 2 (2.1%) | 4 (2.1%) |
| Infections and infestations | 0 | 2 (2.1%) | 2 (1.1%) |
| Upper respiratory tract infection | 0 | 1 (1.0%) | 1 (0.5%) |
| Wound abscess | 0 | 1 (1.0%) | 1 (0.5%) |
| Gastrointestinal disorders | 1 (1.1%) | 0 | 1 (0.5%) |
| Gastrooesophageal reflux disease | 1 (1.1%) | 0 | 1 (0.5%) |
| Nausea | 1 (1.1%) | 0 | 1 (0.5%) |
| Vomiting | 1 (1.1%) | 0 | 1 (0.5%) |
| Nervous system disorders | 1 (1.1%) | 0 | 1 (0.5%) |
| Headache | 1 (1.1%) | 0 | 1 (0.5%) |

Safety incidence analysis: Severe Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| System Organ Class (SOC) | PF-04965842 200mg QD | Placebo | Total Population |
|--------------------------|----------------------|-----------|------------------|
| Preferred Term (PT) | N=94 | N=96 | N=190 |
| Overall | 0 (0.0%) | 0 (0.0%) | 0 (0.0%) |

Safety incidence analysis: Serious Treatment Emergent Adverse Events leading to study drug discontinuation, overall and by SOC/PT - Safety Set JADE TEEN (PF-04965842) - 2023 datacut

| System Organ Class (SOC) | PF-04965842 200mg QD | Placebo | Total Population |
|--------------------------|----------------------|-----------|------------------|
| Preferred Term (PT) | N=94 | N=96 | N=190 |
| Overall | 0 (0.0%) | 0 (0.0%) | 0 (0.0%) |

MMRM Analysen JADE TEEN

| | PF-04965842 200mg QD | Placebo |
|--|---------------------------|--------------------------|
| /ariable / analysis | N=94 | N=96 |
| Baseline | | |
| Return Rate (% subjects with missing data) | | |
| Results at Baseline | 0 (0.0%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 94 | 96 |
| Mean (SD) | 29.52 (12.20) | 29.17 (12.68) |
| p25, median, p75 Min,Max | 19.8, 25.4, 37.7 16, 68 | 19.5, 24.5, 34.9 9, 64 |

| | PF-04965842 200mg QD | Placebo |
|--|-------------------------------|-------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 2 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 2 | 2 (2.1%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 92 | 91 |
| Mean (SD) | 13.93 (12.13) | 20.78 (12.31) |
| p25, median, p75 Min,Max | 6.2, 11.0, 16.8 0, 59 | 12.9, 17.6, 25.0 3, 64 |
| Descriptive Results of Change from baseline | | |
| N with data | 92 | 91 |
| Mean (SD) | -54.69 (28.44) | -29.38 (27.22) |
| p25, median, p75 Min,Max | -75.7, -56.6, -41.8 -98, 29 | -49.6, -28.5, -10.7 -84, 39 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -54.55 (3.55) | -27.56 (3.51) |
| | (-61.58,-47.53) | (-34.51,-20.62) |
| Treatment arm difference (SE), (95% CI), p-value | | -26.99 (4.99) |
| | | (-36.87,-17.10) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.62 (-0.92, -0.32) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|-------------------------------|--------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 3 (3.2%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 91 | 96 |
| Mean (SD) | 7.58 (9.34) | 16.83 (12.91) |
| p25, median, p75 Min,Max | 2.1, 4.3, 10.4 0, 54 | 8.2, 13.2, 20.3 1, 68 |
| Descriptive Results of Change from baseline | | |
| N with data | 91 | 96 |
| Mean (SD) | -75.18 (23.88) | -41.48 (45.08) |
| p25, median, p75 Min,Max | -92.0, -82.7, -66.7 -100, 0 | -68.8, -50.2, -24.6 -93, 300 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -74.26 (3.82) | -41.71 (3.74) |
| | (-81.80,-66.71) | (-49.08,-34.33) |
| Treatment arm difference (SE), (95% CI), p-value | | -32.55 (5.35) |
| · · · · · · · · · · · · · · · · · · · | | (-43.10,-22.00) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.61 (-0.91, -0.32) |

| | PF-04965842 200mg QD | Placebo |
|--|--------------------------------|-------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 5 (5.3%) | 6 (6.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 89 | 90 |
| Mean (SD) | 6.78 (9.26) | 12.98 (13.58) |
| p25, median, p75 Min,Max | 1.0, 2.9, 8.0 0, 52 | 4.2, 9.2, 16.5 0, 70 |
| Descriptive Results of Change from baseline | | |
| N with data | 89 | 90 |
| Mean (SD) | -78.32 (24.92) | -56.85 (33.17) |
| p25, median, p75 Min,Max | -95.5, -88.1, -71.7 -100, -2 | -82.1, -65.7, -39.6 -99, 48 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -77.87 (3.03) | -57.56 (2.99) |
| | (-83.86,-71.89) | (-63.46,-51.66) |
| Treatment arm difference (SE), (95% CI), p-value | | -20.31 (4.26) |
| - | | (-28.73,-11.90) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.63 (-0.93, -0.33) |

| | PF-04965842 200mg QD | Placebo |
|--|--------------------------------|--------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 4 (4.3%) | 6 (6.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 90 |
| Mean (SD) | 5.97 (8.69) | 11.81 (12.64) |
| p25, median, p75 Min,Max | 0.8, 2.6, 6.6 0, 46 | 2.8, 7.0, 16.0 0, 60 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 90 |
| Mean (SD) | -80.98 (24.13) | -62.41 (31.16) |
| p25, median, p75 Min,Max | -96.7, -90.2, -74.1 -100, 50 | -85.8, -70.8, -44.5 -100, 24 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -80.68 (2.86) | -63.65 (2.84) |
| | (-86.33,-75.02) | (-69.25,-58.05) |
| Treatment arm difference (SE), (95% CI), p-value | | -17.02 (4.04) |
| | | (-24.99,-9.06) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.63 (-0.93, -0.33) |

| | PF-04965842 200mg QD | Placebo |
|---|----------------------|-------------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | -71.84 (2.79) | -47.62 (2.74) |
| | (-77.34,-66.34) | (-53.02,-42.22) |
| Treatment arm difference (SE), (95% CI), p-value | | -24.22 (3.91) |
| • | | (-31.94,-16.50) p<.0001 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | -0.90 (-1.20, -0.60) |
| | | |

Full Analysis Set Population

| PF-04965842 200mg QD | Placebo |
|---------------------------|----------------------------------|
| N=94 | N=96 |
| | |
| | |
| 1 (1.1%) | 0 (0.0%) |
| | |
| 93 | 96 |
| 66.24 (13.33) | 68.51 (13.39) |
| 56.4, 66.1, 76.4 32, 94 | 57.9, 68.3, 78.7 40, 101 |
| | N=94 1 (1.1%) 93 66.24 (13.33) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|------------------------------|------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 2 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 2 | 2 (2.1%) | 3 (3.1%) |
| Descriptive Results of Actual Values | | |
| N with data | 92 | 93 |
| Mean (SD) | 40.81 (19.26) | 56.11 (16.97) |
| p25, median, p75 Min,Max | 26.5, 38.7, 55.0 7, 95 | 47.0, 57.5, 66.1 18, 94 |
| Descriptive Results of Change from baseline | | |
| N with data | 92 | 93 |
| Mean (SD) | -38.76 (23.78) | -18.50 (19.88) |
| p25, median, p75 Min,Max | -54.1, -38.7, -22.2 -90, 7 | -30.1, -16.7, -4.5 -67, 47 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -38.63 (2.30) | -18.86 (2.26) |
| | (-43.16,-34.10) | (-23.33,-14.39) |
| Treatment arm difference (SE), (95% CI), p-value | | -19.77 (3.23) |
| | | (-26.15,-13.40) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.74 (-1.04, -0.44) |

MMRM models: SCA01-Total Score percent change from baseline, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965842 200mg QD | Placebo |
|--|-------------------------------|-------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 5 (5.3%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 89 | 96 |
| Mean (SD) | 28.22 (17.88) | 47.95 (17.16) |
| p25, median, p75 Min,Max | 15.5, 25.0, 35.1 0, 82 | 34.8, 47.5, 59.0 17, 95 |
| Descriptive Results of Change from baseline | | |
| N with data | 89 | 96 |
| Mean (SD) | -57.56 (23.56) | -29.99 (20.95) |
| p25, median, p75 Min,Max | -73.2, -61.3, -42.1 -100, 5 | -48.3, -28.9, -16.3 -73, 24 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -56.86 (2.34) | -30.14 (2.27) |
| | (-61.47,-52.25) | (-34.63,-25.66) |
| Treatment arm difference (SE), (95% CI), p-value | | -26.72 (3.26) |
| | | (-33.16,-20.28) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.74 (-1.04, -0.44) |

MMRM models: SCA01-Total Score percent change from baseline, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965842 200mg QD | Placebo |
|--|--------------------------------|-------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 5 (5.3%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 89 | 91 |
| Mean (SD) | 24.87 (17.27) | 41.24 (20.26) |
| p25, median, p75 Min,Max | 13.3, 22.1, 31.6 0, 75 | 25.1, 40.1, 51.7 6, 99 |
| Descriptive Results of Change from baseline | | |
| N with data | 89 | 91 |
| Mean (SD) | -62.40 (24.49) | -39.91 (24.23) |
| p25, median, p75 Min,Max | -78.7, -65.2, -53.3 -100, 29 | -56.0, -41.9, -19.6 -92, 13 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -61.98 (2.54) | -40.02 (2.49) |
| | (-66.98, -56.97) | (-44.93,-35.11) |
| Treatment arm difference (SE), (95% CI), p-value | | -21.95 (3.56) |
| | | (-28.98,-14.93) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.75 (-1.05, -0.45) |

MMRM models: SCA01-Total Score percent change from baseline, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965842 200mg QD | Placebo |
|--|--------------------------------|--------------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 5 (5.3%) | 7 (7.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 89 | 89 |
| Mean (SD) | 23.55 (17.35) | 38.55 (21.17) |
| p25, median, p75 Min,Max | 11.6, 18.6, 31.6 0, 90 | 23.0, 35.9, 51.9 0, 88 |
| Descriptive Results of Change from baseline | | |
| N with data | 89 | 89 |
| Mean (SD) | -64.58 (24.88) | -43.19 (28.25) |
| p25, median, p75 Min,Max | -82.1, -67.3, -52.2 -100, 45 | -65.3, -42.0, -21.7 -100, 22 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -64.27 (2.78) | -44.52 (2.75) |
| | (-69.76,-58.78) | (-49.94,-39.09) |
| Treatment arm difference (SE), (95% CI), p-value | | -19.76 (3.91) |
| | | (-27.48,-12.03) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.75 (-1.06, -0.45) |

MMRM models: SCA01-Total Score percent change from baseline, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965842 200mg QD | Placebo |
|---|----------------------|-------------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | -55.44 (2.08) | -33.39 (2.04) |
| | (-59.55,-51.32) | (-37.41,-29.36) |
| Treatment arm difference (SE), (95% CI), p-value | | -22.05 (2.92) |
| • | | (-27.82,-16.29) p<.0001 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | -1.10 (-1.41, -0.79) |
| | | |

Full Analysis Set Population

MMRM models: POEM01-Total Score, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| PF-04965842 200mg QD | Placebo |
|--------------------------|-------------------------------|
| N=94 | N=96 |
| | |
| | |
| 0 (0.0%) | 1 (1.0%) |
| | |
| 94 | 95 |
| 19.20 (6.22) | 19.84 (5.87) |
| 15.0, 20.0, 24.0 5, 28 | 16.0, 21.0, 24.0 6, 28 |
| | N=94 0 (0.0%) 94 19.20 (6.22) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 2 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 2 | 2 (2.1%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 92 | 91 |
| Mean (SD) | 11.24 (7.32) | 16.33 (7.49) |
| p25, median, p75 Min,Max | 6.0, 10.0, 16.0 0, 28 | 10.0, 17.0, 22.0 3, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 92 | 91 |
| Mean (SD) | -8.16 (7.24) | -3.59 (6.29) |
| p25, median, p75 Min,Max | -14.0, -7.0, -3.0 -25, 4 | -6.0, -2.0, 0.0 -24, 10 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -8.20 (0.67) | -3.42 (0.67) |
| | (-9.53,-6.88) | (-4.75,-2.10) |
| Treatment arm difference (SE), (95% CI), p-value | | -4.78 (0.95) |
| | | (-6.65,-2.90) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.56 (-0.86, -0.27) |

| | PF-04965842 200mg QD | Placebo |
|--|-----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 3 (3.2%) | 1 (1.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 91 | 95 |
| Mean (SD) | 8.76 (6.53) | 14.84 (6.70) |
| p25, median, p75 Min,Max | 4.0, 7.0, 12.0 0, 28 | 10.0, 15.0, 20.0 0, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 91 | 95 |
| Mean (SD) | -10.62 (6.44) | -5.00 (6.07) |
| p25, median, p75 Min,Max | -16.0, -10.0, -6.0 -26, 2 | -8.0, -5.0, -1.0 -20, 8 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -10.62 (0.59) | -4.87 (0.58) |
| | (-11.79,-9.44) | (-6.02,-3.72) |
| Treatment arm difference (SE), (95% CI), p-value | | -5.75 (0.83) |
| | | (-7.39,-4.10) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.56 (-0.85, -0.27) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 4 (4.3%) | 6 (6.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 90 |
| Mean (SD) | 8.77 (6.76) | 14.51 (7.21) |
| p25, median, p75 Min,Max | 4.0, 7.0, 13.0 0, 28 | 9.0, 14.0, 20.0 0, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 90 |
| Mean (SD) | -10.64 (7.51) | -5.51 (7.06) |
| p25, median, p75 Min,Max | -17.0, -9.5, -5.0 -26, 5 | -10.0, -6.0, 0.0 -28, 9 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -10.60 (0.68) | -5.40 (0.67) |
| | (-11.94,-9.26) | (-6.73,-4.07) |
| Treatment arm difference (SE), (95% CI), p-value | | -5.20 (0.96) |
| | | (-7.10,-3.31) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.57 (-0.87, -0.27) |

| | PF-04965842 200mg QD | Placebo |
|--|-----------------------------|----------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 4 (4.3%) | 7 (7.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 89 |
| Mean (SD) | 8.53 (6.77) | 12.93 (7.34) |
| p25, median, p75 Min,Max | 3.0, 7.5, 12.0 0, 28 | 7.0, 12.0, 18.0 0, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 89 |
| Mean (SD) | -10.86 (7.14) | -7.10 (8.22) |
| p25, median, p75 Min,Max | -17.0, -10.0, -6.0 -27, 7 | -12.0, -6.0, -1.0 -28, 8 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -10.85 (0.71) | -6.99 (0.71) |
| | (-12.25,-9.44) | (-8.39,-5.59) |
| Treatment arm difference (SE), (95% CI), p-value | | -3.86 (1.01) |
| | | (-5.84,-1.87) p=0.0002 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.57 (-0.87, -0.27) |

| | PF-04965842 200mg QD | Placebo |
|---|----------------------|-----------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | -10.07 (0.56) | -5.17 (0.55) |
| | (-11.17,-8.97) | (-6.26,-4.08) |
| Treatment arm difference (SE), (95% CI), p-value | | -4.90 (0.78) |
| • | | (-6.44,-3.35) p<.0001 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | -0.91 (-1.21, -0.61) |
| ` , | | , , , |

Full Analysis Set Population

| NI O4 | |
|-----------------------|-----------------------|
| N=94 | N=96 |
| | |
| | |
| 0 (0.0%) | 0 (0.0%) |
| | |
| 94 | 96 |
| 6.81 (1.97) | 7.16 (1.69) |
| 6.0, 7.0, 8.0 1, 10 | 6.0, 7.0, 8.0 4, 10 |
| | 94 6.81 (1.97) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo | |
|--|---------------------------|-------------------------|--|
| Variable / analysis | N=94 | N=96 | |
| Week 2 | | | |
| Return Rate (% subjects with missing data) | | | |
| Results at Week 2 | 3 (3.2%) | 1 (1.0%) | |
| Descriptive Results of Actual Values | | | |
| N with data | 91 | 95 | |
| Mean (SD) | 4.16 (2.17) | 5.96 (2.22) | |
| p25, median, p75 Min,Max | 3.0, 4.0, 5.0 0, 10 | 5.0, 6.0, 8.0 0, 10 | |
| Descriptive Results of Change from baseline | | | |
| N with data | 91 | 95 | |
| Mean (SD) | -2.69 (2.59) | -1.20 (1.99) | |
| p25, median, p75 Min,Max | -4.0, -2.0, -1.0 -10, 3 | -2.0, -1.0, 0.0 -9, 3 | |
| Least Square Means estimates of change from baseline | | | |
| Treatment arm estimate (SE), (95% CI) | -2.74 (0.21) | -1.10 (0.21) | |
| | (-3.16,-2.31) | (-1.52,-0.69) | |
| Treatment arm difference (SE), (95% CI), p-value | | -1.64 (0.30) | |
| | | (-2.23,-1.04) p<.0001 | |
| Hedge G analysis of change from baseline | | | |
| Treatment arm difference estimate (95% CI) | | -0.55 (-0.85, -0.26) | |

| | PF-04965842 200mg QD | Placebo |
|--|---------------------------|-------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 9 (9.6%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 85 | 91 |
| Mean (SD) | 3.34 (2.12) | 5.25 (2.26) |
| p25, median, p75 Min,Max | 2.0, 3.0, 5.0 0, 9 | 4.0, 5.0, 7.0 0, 10 |
| Descriptive Results of Change from baseline | | |
| N with data | 85 | 91 |
| Mean (SD) | -3.55 (2.61) | -1.96 (2.16) |
| p25, median, p75 Min,Max | -5.0, -3.0, -2.0 -10, 3 | -3.0, -2.0, 0.0 -9, 1 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -3.54 (0.22) | -1.86 (0.22) |
| | (-3.98,-3.10) | (-2.29,-1.43) |
| Treatment arm difference (SE), (95% CI), p-value | | -1.69 (0.31) |
| | | (-2.30,-1.07) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.57 (-0.87, -0.26) |

| | PF-04965842 200mg QD | Placebo |
|--|--------------------------|--------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 8 (8.5%) | 7 (7.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 86 | 89 |
| Mean (SD) | 3.03 (2.18) | 4.87 (2.60) |
| p25, median, p75 Min,Max | 1.0, 3.0, 5.0 0, 10 | 3.0, 5.0, 7.0 0, 10 |
| Descriptive Results of Change from baseline | | |
| N with data | 86 | 89 |
| Mean (SD) | -3.81 (2.62) | -2.31 (2.44) |
| p25, median, p75 Min,Max | -6.0, -4.0, -2.0 -9, 3 | -4.0, -2.0, 0.0 -10, 2 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -3.94 (0.24) | -2.29 (0.24) |
| | (-4.42,-3.46) | (-2.77,-1.82) |
| Treatment arm difference (SE), (95% CI), p-value | | -1.65 (0.34) |
| - - | | (-2.32,-0.97) p<.0001 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.57 (-0.87, -0.27) |

| | PF-04965842 200mg QD | Placebo |
|--|---------------------------|--------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 21 (22.3%) | 16 (16.7%) |
| Descriptive Results of Actual Values | | |
| N with data | 73 | 80 |
| Mean (SD) | 3.00 (2.20) | 4.40 (2.49) |
| p25, median, p75 Min,Max | 2.0, 3.0, 4.0 0, 10 | 2.5, 4.0, 6.0 0, 10 |
| Descriptive Results of Change from baseline | | |
| N with data | 73 | 80 |
| Mean (SD) | -3.97 (2.55) | -2.64 (2.68) |
| p25, median, p75 Min,Max | -6.0, -4.0, -2.0 -10, 1 | -4.0, -2.0, 0.0 -10, 2 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -3.93 (0.25) | -2.60 (0.24) |
| | (-4.42,-3.43) | (-3.08,-2.12) |
| Treatment arm difference (SE), (95% CI), p-value | | -1.32 (0.35) |
| | | (-2.02,-0.63) p=0.0002 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.61 (-0.93, -0.28) |

| | PF-04965842 200mg QD | Placebo |
|---|----------------------|-----------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | -3.54 (0.19) | -1.96 (0.19) |
| | (-3.92,-3.15) | (-2.34,-1.59) |
| Treatment arm difference (SE), (95% CI), p-value | | -1.57 (0.27) |
| • | | (-2.11,-1.04) p<.0001 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | -0.84 (-1.14, -0.54) |
| | | |

Full Analysis Set Population

MMRM models: CDLQ1-Total Score, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| PF-04965842 200mg QD | Placebo |
|-------------------------|-------------------------------|
| N=94 | N=96 |
| | |
| | |
| 0 (0.0%) | 0 (0.0%) |
| | |
| 94 | 96 |
| 13.62 (7.02) | 14.04 (6.69) |
| 8.0, 13.0, 19.0 3, 30 | 9.0, 14.0, 19.0 2, 30 |
| | N=94 0 (0.0%) 94 13.62 (7.02) |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 2 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 2 | 2 (2.1%) | 4 (4.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 92 | 92 |
| Mean (SD) | 7.57 (5.78) | 9.84 (6.29) |
| p25, median, p75 Min,Max | 3.0, 6.0, 11.0 0, 26 | 5.0, 9.0, 14.0 1, 25 |
| Descriptive Results of Change from baseline | | |
| N with data | 92 | 92 |
| Mean (SD) | -6.16 (6.51) | -4.24 (4.28) |
| p25, median, p75 Min,Max | -9.0, -5.0, -2.0 -26, 11 | -7.0, -4.0, -1.0 -17, 8 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -6.20 (0.49) | -4.15 (0.49) |
| | (-7.17,-5.23) | (-5.11,-3.19) |
| Treatment arm difference (SE), (95% CI), p-value | | -2.05 (0.69) |
| | | (-3.42,-0.69) p=0.0034 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.48 (-0.77, -0.18) |

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 3 (3.2%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 91 | 96 |
| Mean (SD) | 6.13 (5.05) | 8.58 (5.89) |
| p25, median, p75 Min,Max | 2.0, 5.0, 9.0 0, 21 | 4.0, 8.0, 12.0 0, 27 |
| Descriptive Results of Change from baseline | | |
| N with data | 91 | 96 |
| Mean (SD) | -7.53 (6.23) | -5.46 (5.42) |
| p25, median, p75 Min,Max | -11.0, -7.0, -3.0 -27, 3 | -8.0, -5.0, -2.0 -23, 7 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -7.57 (0.47) | -5.37 (0.46) |
| | (-8.50,-6.63) | (-6.28,-4.46) |
| Treatment arm difference (SE), (95% CI), p-value | | -2.20 (0.66) |
| | | (-3.50,-0.90) p=0.0010 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.47 (-0.76, -0.18) |

| | PF-04965842 200mg QD | Placebo |
|--|-----------------------------|-----------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 4 (4.3%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 91 |
| Mean (SD) | 5.61 (5.33) | 7.96 (6.38) |
| p25, median, p75 Min,Max | 2.0, 4.0, 7.0 0, 26 | 3.0, 7.0, 11.0 0, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 91 |
| Mean (SD) | -8.16 (6.96) | -6.10 (5.53) |
| p25, median, p75 Min,Max | -12.0, -8.0, -3.0 -27, 12 | -10.0, -6.0, -2.0 -20, 11 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -8.11 (0.52) | -5.99 (0.52) |
| | (-9.14,-7.07) | (-7.01,-4.97) |
| Treatment arm difference (SE), (95% CI), p-value | | -2.11 (0.74) |
| | | (-3.57,-0.66) p=0.0046 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.48 (-0.78, -0.18) |

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|----------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 4 (4.3%) | 6 (6.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 90 |
| Mean (SD) | 5.14 (4.74) | 7.82 (6.24) |
| p25, median, p75 Min,Max | 2.0, 4.0, 8.0 0, 29 | 3.0, 6.5, 12.0 0, 28 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 90 |
| Mean (SD) | -8.57 (6.57) | -6.44 (6.37) |
| p25, median, p75 Min,Max | -13.0, -8.0, -4.0 -23, 9 | -10.0, -6.0, -2.0 -22, 9 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | -8.58 (0.51) | -6.26 (0.50) |
| | (-9.59,-7.58) | (-7.26,-5.27) |
| Treatment arm difference (SE), (95% CI), p-value | | -2.32 (0.72) |
| · · · · · · · · · · · · · | | (-3.74,-0.91) p=0.0014 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | -0.48 (-0.78, -0.18) |

| | PF-04965842 200mg QD | Placebo |
|---|----------------------|------------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | -7.61 (0.42) | -5.44 (0.41) |
| | (-8.44,-6.78) | (-6.26,-4.63) |
| Treatment arm difference (SE), (95% CI), p-value | | -2.17 (0.59) |
| · · · · · · · · · · · · · · · · · · · | | (-3.34,-1.01) p=0.0003 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | -0.54 (-0.83, -0.24) |
| | | |

Full Analysis Set Population

MMRM models: EQ5D03-EQ VAS Score, Total population - Full Analysis Set Population JADE TEEN (PF-04965842) - 2023 datacut

| | PF-04965842 200mg QD | Placebo |
|--|---------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Baseline | | |
| Return Rate (% subjects with missing data) | | |
| Results at Baseline | 0 (0.0%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 94 | 96 |
| Mean (SD) | 64.94 (21.61) | 63.51 (24.83) |
| p25, median, p75 Min,Max | 50.0, 65.0, 82.0 9, 100 | 48.0, 65.5, 85.0 5, 100 |

Full Analysis Set Population

| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 2 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 2 | 2 (2.1%) | 4 (4.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 92 | 92 |
| Mean (SD) | 76.36 (17.10) | 70.70 (21.78) |
| p25, median, p75 Min,Max | 64.0, 80.0, 90.0 22, 100 | 60.0, 75.0, 88.5 0, 100 |
| Descriptive Results of Change from baseline | | |
| N with data | 92 | 92 |
| Mean (SD) | 11.80 (21.67) | 7.50 (21.65) |
| p25, median, p75 Min,Max | -0.5, 9.5, 25.0 -45, 75 | -4.0, 3.0, 19.5 -45, 78 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | 11.92 (1.76) | 7.00 (1.75) |
| | (8.45,15.39) | (3.55,10.44) |
| Treatment arm difference (SE), (95% CI), p-value | | 4.93 (2.48) |
| | | (0.03,9.82) p=0.0484 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | 0.33 (0.04, 0.62) |

Full Analysis Set Population

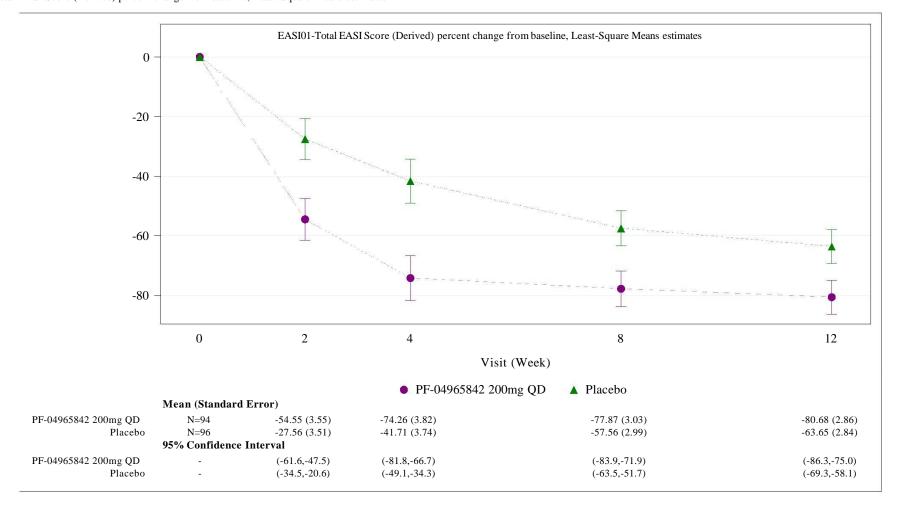
| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|----------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 4 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 4 | 3 (3.2%) | 0 (0.0%) |
| Descriptive Results of Actual Values | | |
| N with data | 91 | 96 |
| Mean (SD) | 79.03 (15.91) | 72.46 (20.58) |
| p25, median, p75 Min,Max | 73.0, 80.0, 90.0 20, 100 | 62.5, 75.0, 89.5 12, 100 |
| Descriptive Results of Change from baseline | | |
| N with data | 91 | 96 |
| Mean (SD) | 14.47 (19.45) | 8.95 (22.06) |
| p25, median, p75 Min,Max | 0.0, 11.0, 26.0 -27, 77 | -4.5, 5.0, 21.0 -40, 76 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | 14.47 (1.64) | 8.64 (1.60) |
| | (11.23,17.71) | (5.48,11.80) |
| Treatment arm difference (SE), (95% CI), p-value | | 5.83 (2.30) |
| | | (1.30,10.36) p=0.0119 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | 0.32 (0.04, 0.61) |

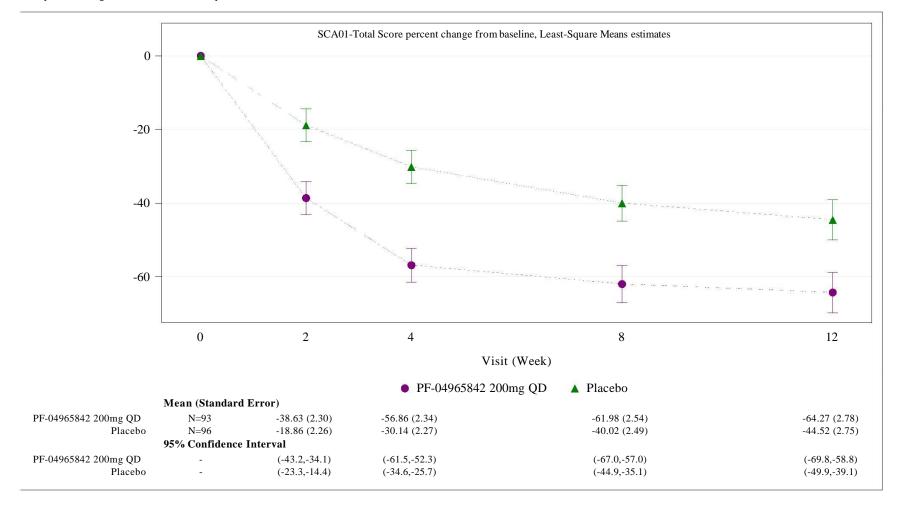
| | PF-04965842 200mg QD | Placebo |
|--|----------------------------|----------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 8 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 8 | 4 (4.3%) | 5 (5.2%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 91 |
| Mean (SD) | 78.97 (19.73) | 71.92 (20.54) |
| p25, median, p75 Min,Max | 74.0, 85.0, 93.0 18, 100 | 60.0, 74.0, 90.0 17, 100 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 91 |
| Mean (SD) | 14.48 (22.87) | 8.82 (21.50) |
| p25, median, p75 Min,Max | -3.0, 11.5, 31.0 -51, 74 | -2.0, 5.0, 21.0 -42, 76 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | 14.45 (1.84) | 8.28 (1.82) |
| | (10.82,18.08) | (4.70,11.86) |
| Treatment arm difference (SE), (95% CI), p-value | | 6.16 (2.59) |
| | | (1.06,11.26) p=0.0181 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | 0.33 (0.04, 0.62) |

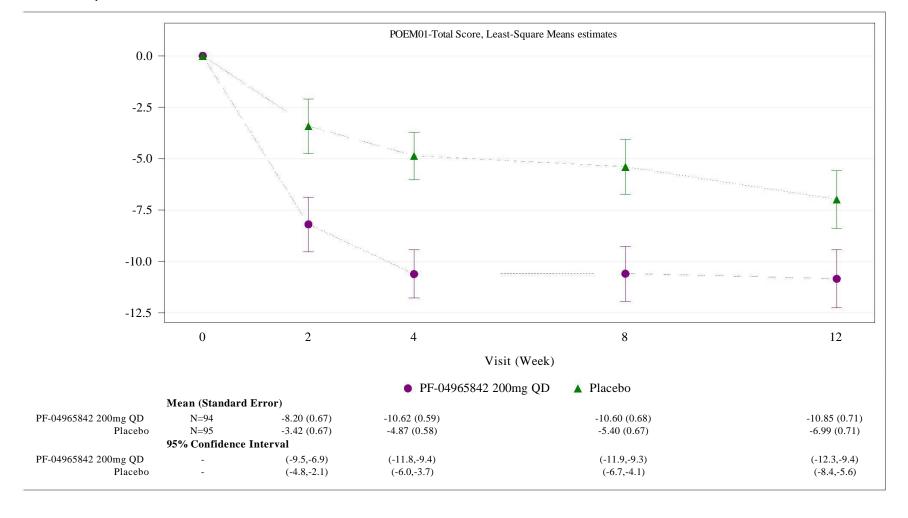
| | PF-04965842 200mg QD | Placebo |
|--|---------------------------|---------------------------|
| Variable / analysis | N=94 | N=96 |
| Week 12 | | |
| Return Rate (% subjects with missing data) | | |
| Results at Week 12 | 4 (4.3%) | 6 (6.3%) |
| Descriptive Results of Actual Values | | |
| N with data | 90 | 90 |
| Mean (SD) | 80.12 (17.68) | 72.94 (21.97) |
| p25, median, p75 Min,Max | 71.0, 85.0, 92.0 0, 100 | 62.0, 75.0, 91.0 0, 100 |
| Descriptive Results of Change from baseline | | |
| N with data | 90 | 90 |
| Mean (SD) | 15.71 (21.19) | 10.32 (23.92) |
| p25, median, p75 Min,Max | 1.0, 13.5, 30.0 -39, 61 | -3.0, 6.5, 24.0 -41, 90 |
| Least Square Means estimates of change from baseline | | |
| Treatment arm estimate (SE), (95% CI) | 15.54 (1.83) | 9.79 (1.82) |
| | (11.93,19.15) | (6.21,13.38) |
| Treatment arm difference (SE), (95% CI), p-value | | 5.75 (2.58) |
| | | (0.66,10.84) p=0.0270 |
| Hedge G analysis of change from baseline | | |
| Treatment arm difference estimate (95% CI) | | 0.33 (0.04, 0.63) |

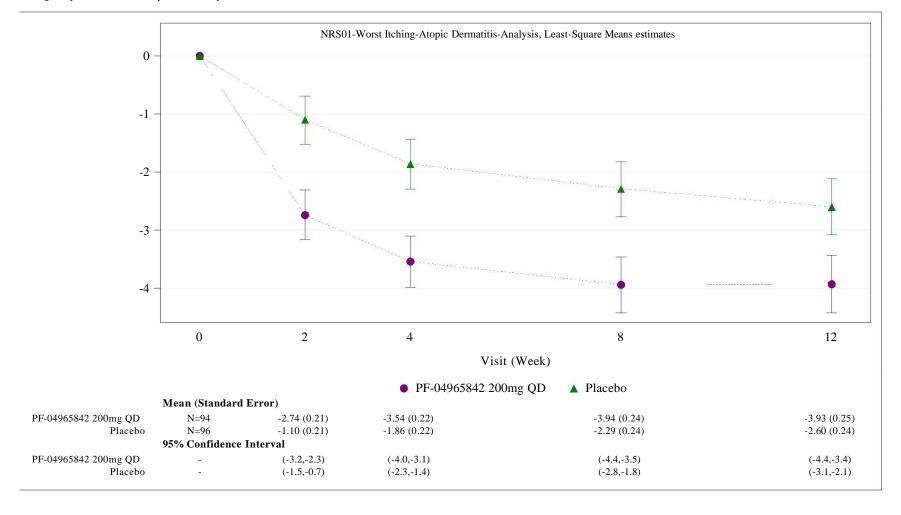
| | PF-04965842 200mg QD | Placebo |
|---|----------------------|----------------------|
| Variable / analysis | N=94 | N=96 |
| Overall post-baseline | | |
| Least Square Means estimates of change from baseline | | |
| Treatment arm average post-baseline estimate (SE), (95% CI) | 14.10 (1.42) | 8.43 (1.39) |
| | (11.30,16.89) | (5.68,11.17) |
| Treatment arm difference (SE), (95% CI), p-value | | 5.67 (1.99) |
| · //· | | (1.75,9.59) p=0.0048 |
| Hedge G analysis of average post-baseline change | | |
| Treatment arm difference estimate (95% CI) | | 0.41 (0.13, 0.70) |
| | | (,,, |

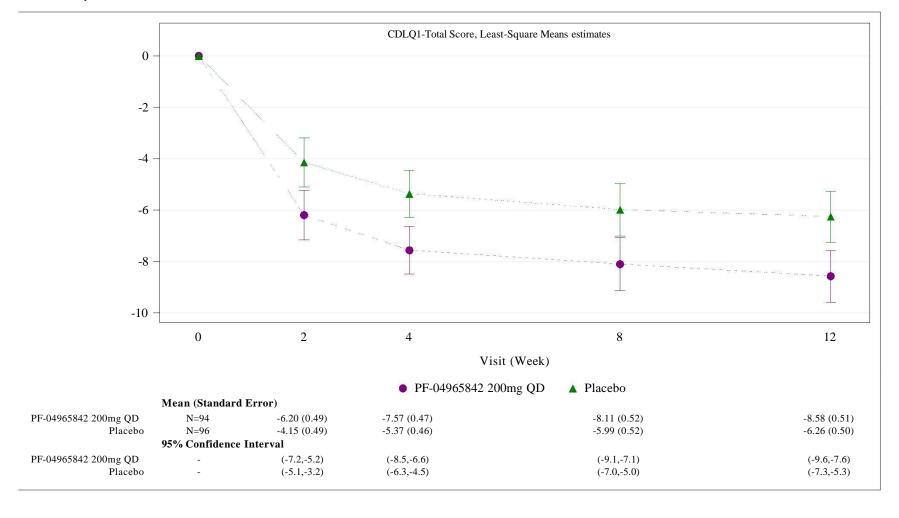
Full Analysis Set Population

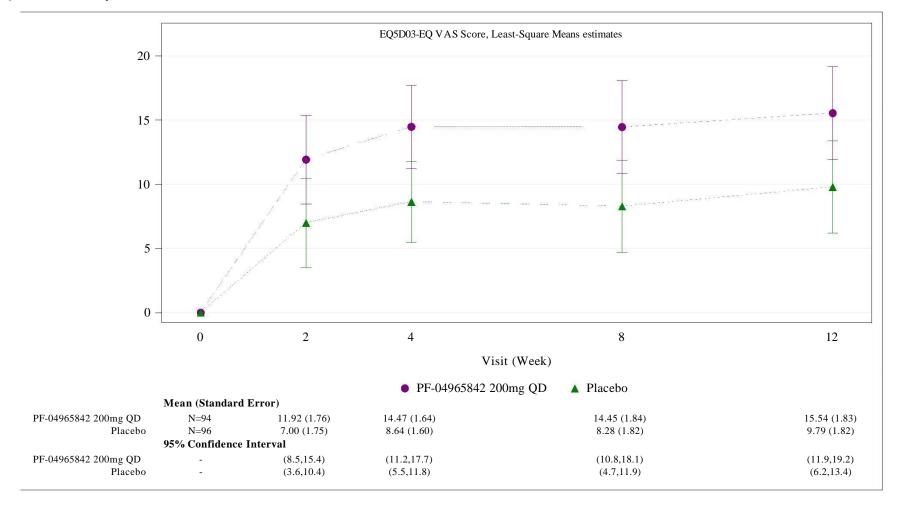












Zusatzanalysen JADE DARE und JADE TEEN (Woche 2 bis Woche 26)

Binary Outcome Analysis: SCORAD-90 response - Full Analysis Set Safety Population JADE TEEN and DARE (PF-04965842) - 2023 datacuts

| Abrocitinib 200mg QD [DARE] | | Dupiluma | Dupilumab 300mg Q2W [DARE] | | Abrocitinib 200mg QD [TEEN] | |
|-----------------------------|----------------|----------|----------------------------|----|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| - Visit Week 2 | | | | | | |
| 362 | 7 (1.9%) | 365 | 2 (0.5%) | 93 | 1 (1.1%) | |
| | (0.8%, 3.9%) | | (0.1%, 2.0%) | | (0.0%, 5.8%) | |
| - Visit Week 4 | | | | | | |
| 362 | 28 (7.7%) | 365 | 7 (1.9%) | 93 | 6 (6.5%) | |
| | (5.2%, 11.0%) | | (0.8%, 3.9%) | | (2.4%, 13.5% | |
| - Visit Week 8 | | | | | | |
| 362 | 48 (13.3%) | 365 | 15 (4.1%) | 93 | 9 (9.7%) | |
| | (9.9%, 17.2%) | | (2.3%, 6.7%) | | (4.5%, 17.6% | |
| - Visit Week 12 | | | | | | |
| 362 | 60 (16.6%) | 365 | 24 (6.6%) | 93 | 11 (11.8%) | |
| | (12.9%, 20.8%) | | (4.3%, 9.6%) | | (6.1%, 20.2% | |
| - Visit Week 16 | | | | | | |
| 362 | 71 (19.6%) | 365 | 31 (8.5%) | 0 | 0 | |
| | (15.6%, 24.1%) | | (5.8%, 11.8%) | | | |
| - Visit Week 20 | | | | | | |
| 362 | 79 (21.8%) | 365 | 44 (12.1%) | 0 | 0 | |
| | (17.7%, 26.4%) | | (8.9%, 15.8%) | | | |
| - Visit Week 26 | | | | | | |
| 362 | 80 (22.1%) | 365 | 52 (14.2%) | 0 | 0 | |
| | (17.9%, 26.7%) | | (10.8%, 18.3%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

| Abrocitinil | 200mg QD [DARE] | Dupilumat | 300mg Q2W [DARE] | Abrocitinib 200 | Abrocitinib 200mg QD [TEEN] | |
|-----------------|-----------------|-----------|------------------|-----------------|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| Visit Week 2 | | | | | | |
| 362 | 0 (0.0%) | 365 | 1 (0.3%) | 93 | 0 (0.0%) | |
| | | | (0.0%, 1.5%) | | | |
| Visit Week 4 | | | | | | |
| 362 | 11 (3.0%) | 365 | 3 (0.8%) | 93 | 1 (1.1%) | |
| | (1.5%, 5.4%) | | (0.2%, 2.4%) | | (0.0%, 5.8%) | |
| · Visit Week 8 | | | | | | |
| 362 | 15 (4.1%) | 365 | 4 (1.1%) | 93 | 3 (3.2%) | |
| | (2.3%, 6.7%) | | (0.3%, 2.8%) | | (0.7%, 9.1%) | |
| · Visit Week 12 | | | | | | |
| 362 | 21 (5.8%) | 365 | 6 (1.6%) | 93 | 3 (3.2%) | |
| | (3.6%, 8.7%) | | (0.6%, 3.5%) | | (0.7%, 9.1%) | |
| Visit Week 16 | | | | | | |
| 362 | 29 (8.0%) | 365 | 9 (2.5%) | 0 | 0 | |
| | (5.4%, 11.3%) | | (1.1%, 4.6%) | | | |
| · Visit Week 20 | | | | | | |
| 362 | 36 (9.9%) | 365 | 21 (5.8%) | 0 | 0 | |
| | (7.1%, 13.5%) | | (3.6%, 8.7%) | | | |
| Visit Week 26 | | | | | | |
| 362 | 37 (10.2%) | 365 | 22 (6.0%) | 0 | 0 | |
| | (7.3%, 13.8%) | | (3.8%, 9.0%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

| Abrocitini | b 200mg QD [DARE] | Dupiluma | Dupilumab 300mg Q2W [DARE] | | Abrocitinib 200mg QD [TEEN] | |
|-----------------|-------------------|----------|----------------------------|----|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| - Visit Week 2 | | | | | | |
| 362 | 7 (1.9%) | 365 | 3 (0.8%) | 94 | 0 (0.0%) | |
| | (0.8%, 3.9%) | | (0.2%, 2.4%) | | | |
| - Visit Week 4 | | | | | | |
| 362 | 27 (7.5%) | 365 | 9 (2.5%) | 94 | 5 (5.3%) | |
| | (5.0%, 10.7%) | | (1.1%, 4.6%) | | (1.7%, 12.0%) | |
| · Visit Week 8 | | | | | | |
| 362 | 45 (12.4%) | 365 | 13 (3.6%) | 94 | 9 (9.6%) | |
| | (9.2%, 16.3%) | | (1.9%, 6.0%) | | (4.5%, 17.4%) | |
| · Visit Week 12 | | | | | | |
| 362 | 56 (15.5%) | 365 | 20 (5.5%) | 94 | 8 (8.5%) | |
| | (11.9%, 19.6%) | | (3.4%, 8.3%) | | (3.7%, 16.1%) | |
| · Visit Week 16 | | | | | | |
| 362 | 70 (19.3%) | 365 | 33 (9.0%) | 0 | 0 | |
| | (15.4%, 23.8%) | | (6.3%, 12.5%) | | | |
| Visit Week 20 | | | | | | |
| 362 | 74 (20.4%) | 365 | 45 (12.3%) | 0 | 0 | |
| | (16.4%, 25.0%) | | (9.1%, 16.1%) | | | |
| · Visit Week 26 | , | | , | | | |
| 362 | 79 (21.8%) | 365 | 50 (13.7%) | 0 | 0 | |
| | (17.7%, 26.4%) | | (10.3%, 17.7%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: Achieving 0-2 in POEM total score - Full Analysis Set Safety Population JADE TEEN and DARE (PF-04965842) - 2023 datacuts

| Abrocitini | b 200mg QD [DARE] | Dupiluma | b 300mg Q2W [DARE] | Abrocitinib 200 | Abrocitinib 200mg QD [TEEN] | |
|-----------------|-------------------|----------|--------------------|-----------------|-----------------------------|--|
| | Events (%) | _ | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| · Visit Week 2 | | | | | | |
| 358 | 0 (0.0%) | 363 | 0 (0.0%) | 94 | 7 (7.4%) | |
| | | | | | (3.0%, 14.7%) | |
| · Visit Week 4 | | | | | | |
| 358 | 1 (0.3%) | 363 | 0 (0.0%) | 94 | 14 (14.9%) | |
| | (0.0%, 1.5%) | | | | (8.4%, 23.7%) | |
| - Visit Week 8 | | | | | | |
| 358 | 0 (0.0%) | 363 | 0 (0.0%) | 94 | 15 (16.0%) | |
| | | | | | (9.2%, 25.0%) | |
| · Visit Week 12 | | | | | | |
| 358 | 108 (30.2%) | 363 | 61 (16.8%) | 94 | 18 (19.1%) | |
| | (25.5%, 35.2%) | | (13.1%, 21.1%) | | (11.8%, 28.6%) | |
| · Visit Week 16 | | | | | | |
| 358 | 104 (29.1%) | 363 | 55 (15.2%) | 0 | 0 | |
| | (24.4%, 34.1%) | | (11.6%, 19.3%) | | | |
| · Visit Week 20 | | | | | | |
| 358 | 0 (0.0%) | 363 | 2 (0.6%) | 0 | 0 | |
| | | | (0.1%, 2.0%) | | | |
| · Visit Week 26 | | | | | | |
| 358 | 106 (29.6%) | 363 | 69 (19.0%) | 0 | 0 | |
| | (24.9%, 34.6%) | | (15.1%, 23.4%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point. The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

Binary Outcome Analysis: Achieving 0 in POEM total score - Full Analysis Set Safety Population JADE TEEN and DARE (PF-04965842) - 2023 datacuts

| Abrocitin | ib 200mg QD [DARE] | Dupiluma | Dupilumab 300mg Q2W [DARE] | | Abrocitinib 200mg QD [TEEN] | |
|---------------|------------------------------|----------|----------------------------|----|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| Visit Week 2 | | | | | | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 94 | 4 (4.3%) (1.2%, 10.5%) | |
| Visit Week 4 | | | | | | |
| 359 | 1 (0.3%) (0.0%, 1.5%) | 365 | 0 (0.0%) | 94 | 5 (5.3%) (1.7%, 12.0%) | |
| Visit Week 8 | , , | | | | , | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 94 | 3 (3.2%) (0.7%, 9.0%) | |
| Visit Week 12 | | | | | (, | |
| 359 | 49 (13.6%) (10.3%, 17.6%) | 365 | 11 (3.0%) (1.5%, 5.3%) | 94 | 6 (6.4%) (2.4%, 13.4%) | |
| Visit Week 16 | | | | | | |
| 359 | 45 (12.5%) (9.3%, 16.4%) | 365 | 16 (4.4%) (2.5%, 7.0%) | 0 | 0 | |
| Visit Week 20 | | | | | | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 0 | 0 | |
| Visit Week 26 | | | | | | |
| 359 | 49 (13.6%) (10.3%, 17.6%) | 365 | 26 (7.1%) (4.7%, 10.3%) | 0 | 0 | |

Notes:

Number of subjects: Full Analysis Set Safety Population Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: Achieving 0-1 in DLQI total score - Full Analysis Set Safety Population JADE TEEN and DARE (PF-04965842) - 2023 datacuts

| Abrocitini | Abrocitinib 200mg QD [DARE] | | b 300mg Q2W [DARE] | Abrocitinib 200 | Abrocitinib 200mg QD [TEEN] | |
|-----------------|-----------------------------|-----|--------------------|-----------------|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| - Visit Week 2 | | | | | | |
| 358 | 81 (22.6%) | 361 | 23 (6.4%) | 94 | 8 (8.5%) | |
| | (18.4%, 27.3%) | | (4.1%, 9.4%) | | (3.7%, 16.1%) | |
| · Visit Week 4 | | | | | | |
| 358 | 3 (0.8%) | 361 | 1 (0.3%) | 94 | 17 (18.1%) | |
| | (0.2%, 2.4%) | | (0.0%, 1.5%) | | (10.9%, 27.4%) | |
| - Visit Week 8 | | | | | | |
| 358 | 0 (0.0%) | 361 | 0 (0.0%) | 94 | 15 (16.0%) | |
| | | | | | (9.2%, 25.0%) | |
| - Visit Week 12 | | | | | | |
| 358 | 126 (35.2%) | 361 | 110 (30.5%) | 94 | 21 (22.3%) | |
| | (30.2%, 40.4%) | | (25.8%, 35.5%) | | (14.4%, 32.1%) | |
| - Visit Week 16 | | | | | | |
| 358 | 147 (41.1%) | 361 | 107 (29.6%) | 0 | 0 | |
| | (35.9%, 46.4%) | | (25.0%, 34.6%) | | | |
| · Visit Week 20 | | | | | | |
| 358 | 154 (43.0%) | 361 | 119 (33.0%) | 0 | 0 | |
| | (37.8%, 48.3%) | | (28.1%, 38.1%) | | | |
| Visit Week 26 | | | | | | |
| 358 | 137 (38.3%) | 361 | 114 (31.6%) | 0 | 0 | |
| | (33.2%, 43.5%) | | (26.8%, 36.6%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point. The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score - Full Analysis Set Safety Population JADE TEEN and DARE (PF-04965842) - 2023 datacuts

| Abrocitini | b 200mg QD [DARE] | Dupiluma | Dupilumab 300mg Q2W [DARE] | | Abrocitinib 200mg QD [TEEN] | |
|----------------|-------------------|----------|----------------------------|----|-----------------------------|--|
| | Events (%) | • | Events (%) | | Events (%) | |
| n | (95% CI) | n | (95% CI) | n | (95% CI) | |
| Overall | | | | | | |
| Visit Week 2 | | | | | | |
| 362 | 62 (17.1%) | 365 | 17 (4.7%) | 94 | 10 (10.6%) | |
| | (13.4%, 21.4%) | | (2.7%, 7.4%) | | (5.2%, 18.7%) | |
| · Visit Week 4 | | | | | | |
| 362 | 101 (27.9%) | 365 | 41 (11.2%) | 94 | 16 (17.0%) | |
| | (23.3%, 32.8%) | | (8.2%, 14.9%) | | (10.1%, 26.2%) | |
| · Visit Week 8 | | | | | | |
| 362 | 125 (34.5%) | 365 | 76 (20.8%) | 94 | 25 (26.6%) | |
| | (29.6%, 39.7%) | | (16.8%, 25.4%) | | (18.0%, 36.7%) | |
| Visit Week 12 | | | | | | |
| 362 | 135 (37.3%) | 365 | 91 (24.9%) | 94 | 18 (19.1%) | |
| | (32.3%, 42.5%) | | (20.6%, 29.7%) | | (11.8%, 28.6%) | |
| Visit Week 16 | | | | | | |
| 362 | 130 (35.9%) | 365 | 99 (27.1%) | 0 | 0 | |
| | (31.0%, 41.1%) | | (22.6%, 32.0%) | | | |
| Visit Week 20 | , , , | | , | | | |
| 362 | 139 (38.4%) | 365 | 98 (26.8%) | 0 | 0 | |
| | (33.4%, 43.6%) | | (22.4%, 31.7%) | | | |
| Visit Week 26 | | | | | | |
| 362 | 139 (38.4%) | 365 | 114 (31.2%) | 0 | 0 | |
| | (33.4%, 43.6%) | | (26.5%, 36.3%) | | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Zusatzanalysen JADE DARE (Woche 2 bis Woche 26)

Binary Outcome Analysis: SCORAD-90 response - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 200 | Omg QD [DARE] Dupil | lumab 300mg Q | 2W [DARE] | Abr | ocitinib 200mg QD [DARE] vs. Dupilu | ımab 300mg Q2W [DARE] | |
|-----------------|---------------------|---------------|----------------|---------------|-------------------------------------|-----------------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| - Visit Week 2 | | | | | | | |
| 362 | 7 (1.9%) | 365 | 2 (0.5%) | 3.58 | 3.53 | 1.4% | 0.0913+ |
| | (0.8%, 3.9%) | | (0.1%, 2.0%) | (0.74, 17.35) | (0.74, 16.87) | (-0.2%, 3.0%) | |
| - Visit Week 4 | | | | | | | |
| 362 | 28 (7.7%) | 365 | 7 (1.9%) | 4.29 | 4.03 | 5.8% | 0.0003+* |
| | (5.2%, 11.0%) | | (0.8%, 3.9%) | (1.85, 9.95) | (1.78, 9.12) | (2.7%, 8.9%) | |
| - Visit Week 8 | | | | | | | |
| 362 | 48 (13.3%) | 365 | 15 (4.1%) | 3.57 | 3.23 | 9.2% | <0.0001+* |
| | (9.9%, 17.2%) | | (2.3%, 6.7%) | (1.96, 6.50) | (1.84, 5.66) | (5.1%, 13.2%) | |
| - Visit Week 12 | 2 | | | | | | |
| 362 | 60 (16.6%) | 365 | 24 (6.6%) | 2.82 | 2.52 | 10.0% | <0.0001+* |
| | (12.9%, 20.8%) | | (4.3%, 9.6%) | (1.72, 4.65) | (1.61, 3.96) | (5.4%, 14.6%) | |
| - Visit Week 16 | • | | | | | | |
| 362 | 71 (19.6%) | 365 | 31 (8.5%) | 2.63 | 2.31 | 11.1% | < 0.0001+* |
| | (15.6%, 24.1%) | | (5.8%, 11.8%) | (1.68, 4.12) | (1.55, 3.43) | (6.1%, 16.1%) | |
| - Visit Week 20 |) | | | | | | |
| 362 | 79 (21.8%) | 365 | 44 (12.1%) | 2.04 | 1.81 | 9.8% | 0.0004+* |
| | (17.7%, 26.4%) | | (8.9%, 15.8%) | (1.36, 3.04) | (1.29, 2.54) | (4.4%, 15.2%) | |
| - Visit Week 26 | • | | | | | | |
| 362 | 80 (22.1%) | 365 | 52 (14.2%) | 1.71 | 1.55 | 7.9% | 0.0061+* |
| | (17.9%, 26.7%) | | (10.8%, 18.3%) | (1.16, 2.51) | (1.13, 2.13) | (2.3%, 13.4%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: SCORAD-100 response - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 200 | Abrocitinib 200mg QD [DARE] Dupilumab 300mg Q2W [DARE] | | Abr | ocitinib 200mg QD [DARE] vs. Dupilu | mab 300mg Q2W [DARE] | | |
|-----------------|--|-----|--------------|-------------------------------------|----------------------|---------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| - Visit Week 2 | | | | | | | |
| 362 | 0 (0.0%) | 365 | 1 (0.3%) | < 0.01 | NE | -0.3% | 0.3193+ |
| | | | (0.0%, 1.5%) | (NE, NE) | (NE, NE) | (-0.8%, 0.3%) | |
| - Visit Week 4 | | | | | | | |
| 362 | 11 (3.0%) | 365 | 3 (0.8%) | 3.78 | 3.70 | 2.2% | 0.0298+* |
| | (1.5%, 5.4%) | | (0.2%, 2.4%) | (1.05, 13.67) | (1.04, 13.14) | (0.2%, 4.2%) | |
| - Visit Week 8 | | | | | | | |
| 362 | 15 (4.1%) | 365 | 4 (1.1%) | 3.90 | 3.78 | 3.0% | 0.0101+* |
| | (2.3%, 6.7%) | | (0.3%, 2.8%) | (1.28, 11.87) | (1.27, 11.28) | (0.7%, 5.4%) | |
| - Visit Week 12 | | | | | | | |
| 362 | 21 (5.8%) | 365 | 6 (1.6%) | 3.68 | 3.53 | 4.2% | 0.0031+* |
| | (3.6%, 8.7%) | | (0.6%, 3.5%) | (1.47, 9.24) | (1.44, 8.64) | (1.4%, 6.9%) | |
| - Visit Week 16 | | | | | | | |
| 362 | 29 (8.0%) | 365 | 9 (2.5%) | 3.44 | 3.25 | 5.5% | 0.0008+* |
| | (5.4%, 11.3%) | | (1.1%, 4.6%) | (1.61, 7.39) | (1.56, 6.77) | (2.3%, 8.8%) | |
| - Visit Week 20 | | | | | | | |
| 362 | 36 (9.9%) | 365 | 21 (5.8%) | 1.81 | 1.73 | 4.2% | 0.0357+* |
| | (7.1%, 13.5%) | | (3.6%, 8.7%) | (1.03, 3.16) | (1.03, 2.90) | (0.3%, 8.1%) | |
| - Visit Week 26 | | | | | | | |
| 362 | 37 (10.2%) | 365 | 22 (6.0%) | 1.77 | 1.70 | 4.2% | 0.0386+* |
| | (7.3%, 13.8%) | | (3.8%, 9.0%) | (1.03, 3.07) | (1.02, 2.82) | (0.2%, 8.2%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

| Abrocitinib 200 | brocitinib 200mg QD [DARE] Dupilumab 300mg Q2W [DARE] | | Abro | ocitinib 200mg QD [DARE] vs. Dupilu | ımab 300mg Q2W [DARE] | | |
|-----------------|---|-----|----------------|-------------------------------------|-----------------------|---------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| - Visit Week 2 | | | | | | | |
| 362 | 7 (1.9%) | 365 | 3 (0.8%) | 2.38 | 2.35 | 1.1% | 0.1985+ |
| | (0.8%, 3.9%) | | (0.2%, 2.4%) | (0.61, 9.27) | (0.61, 9.03) | (-0.6%, 2.8%) | |
| - Visit Week 4 | | | | | | | |
| 362 | 27 (7.5%) | 365 | 9 (2.5%) | 3.19 | 3.02 | 5.0% | 0.0019+* |
| | (5.0%, 10.7%) | | (1.1%, 4.6%) | (1.48, 6.88) | (1.44, 6.34) | (1.9%, 8.1%) | |
| - Visit Week 8 | | | | | | | |
| 362 | 45 (12.4%) | 365 | 13 (3.6%) | 3.84 | 3.49 | 8.9% | <0.0001+* |
| | (9.2%, 16.3%) | | (1.9%, 6.0%) | (2.04, 7.26) | (1.92, 6.36) | (5.0%, 12.8%) | |
| - Visit Week 12 | 2 | | | | | | |
| 362 | 56 (15.5%) | 365 | 20 (5.5%) | 3.16 | 2.82 | 10.0% | <0.0001+* |
| | (11.9%, 19.6%) | | (3.4%, 8.3%) | (1.85, 5.38) | (1.73, 4.61) | (5.6%, 14.4%) | |
| - Visit Week 16 | ó | | | | | | |
| 362 | 70 (19.3%) | 365 | 33 (9.0%) | 2.41 | 2.14 | 10.3% | <0.0001+* |
| | (15.4%, 23.8%) | | (6.3%, 12.5%) | (1.55, 3.76) | (1.45, 3.15) | (5.3%, 15.3%) | |
| - Visit Week 20 |) | | | | | | |
| 362 | 74 (20.4%) | 365 | 45 (12.3%) | 1.83 | 1.66 | 8.1% | 0.0031+* |
| | (16.4%, 25.0%) | | (9.1%, 16.1%) | (1.22, 2.73) | (1.18, 2.33) | (2.8%, 13.5%) | |
| - Visit Week 26 | Ó | | | | | | |
| 362 | 79 (21.8%) | 365 | 50 (13.7%) | 1.76 | 1.59 | 8.1% | 0.0042 + * |
| | (17.7%, 26.4%) | | (10.3%, 17.7%) | (1.19, 2.59) | (1.15, 2.20) | (2.6%, 13.7%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: Achieving 0-2 in POEM total score - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 20 | 0mg QD [DARE] Dupil | umab 300mg Q | 2W [DARE] | Abr | ocitinib 200mg QD [DARE] vs. Dupilu | mab 300mg Q2W [DARE] | |
|-----------------------------------|---------------------|--------------|----------------|---------------|-------------------------------------|----------------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| - Visit Week 2 | | | | | | | |
| 358 | 0 (0.0%) | 363 | 0 (0.0%) | 1.01 | 1.01 | 0.0% | NE |
| | | | | (0.02, 51.24) | (0.02, 50.96) | (-0.5%, 0.5%) | |
| - Visit Week 4 | | | | | | | |
| 358 | 1 (0.3%) | 363 | 0 (0.0%) | NE | NE | 0.3% | 0.3140+ |
| | (0.0%, 1.5%) | | | (NE, NE) | (NE, NE) | (-0.3%, 0.8%) | |
| - Visit Week 8 | | | | | | | |
| 358 | 0 (0.0%) | 363 | 0 (0.0%) | 1.01 | 1.01 | 0.0% | NE |
| | | | | (0.02, 51.24) | (0.02, 50.96) | (-0.5%, 0.5%) | |
| Visit Week 12 | 2 | | | | | | |
| 358 | 108 (30.2%) | 363 | 61 (16.8%) | 2.14 | 1.80 | 13.4% | <0.0001+* |
| | (25.5%, 35.2%) | | (13.1%, 21.1%) | (1.50, 3.05) | (1.36, 2.37) | (7.2%, 19.5%) | |
| Visit Week 16 | 6 | | | | | | |
| 358 | 104 (29.1%) | 363 | 55 (15.2%) | 2.29 | 1.92 | 13.9% | <0.0001+* |
| | (24.4%, 34.1%) | | (11.6%, 19.3%) | (1.59, 3.31) | (1.43, 2.57) | (7.9%, 19.9%) | |
| Visit Week 20 |) | | | | | | |
| 358 | 0 (0.0%) | 363 | 2 (0.6%) | < 0.01 | NE | -0.6% | 0.1599+ |
| | | | (0.1%, 2.0%) | (NE, NE) | (NE, NE) | (-1.3%, 0.2%) | |
| Visit Week 20 | 6 | | | | | | |
| 358 | 106 (29.6%) | 363 | 69 (19.0%) | 1.79 | 1.56 | 10.6% | 0.0009+* |
| | (24.9%, 34.6%) | | (15.1%, 23.4%) | (1.27, 2.54) | (1.19, 2.03) | (4.4%, 16.8%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0-2.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: Achieving 0 in POEM total score - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 200 | Abrocitinib 200mg QD [DARE] Dupilumab 300mg Q2W [DARE] | | | Abr | Abrocitinib 200mg QD [DARE] vs. Dupilumab 300mg Q2W [DARE] | | | |
|-----------------|--|-----|---------------|---------------|--|---------------|-----------------|--|
| | Events (%) | | Events (%) | OR | RR | RD | | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] | |
| Overall | | | | | | | | |
| - Visit Week 2 | | | | | | | | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 1.02 | 1.02 | 0.0% | NE | |
| | | | | (0.02, 51.38) | (0.02, 51.10) | (-0.5%, 0.5%) | | |
| Visit Week 4 | | | | | | | | |
| 359 | 1 (0.3%) | 365 | 0 (0.0%) | NE | NE | 0.3% | 0.3133+ | |
| | (0.0%, 1.5%) | | | (NE, NE) | (NE, NE) | (-0.3%, 0.8%) | | |
| - Visit Week 8 | | | | | | | | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 1.02 | 1.02 | 0.0% | NE | |
| | | | | (0.02, 51.38) | (0.02, 51.10) | (-0.5%, 0.5%) | | |
| Visit Week 12 | | | | | | | | |
| 359 | 49 (13.6%) | 365 | 11 (3.0%) | 5.09 | 4.53 | 10.6% | <0.0001+* | |
| | (10.3%, 17.6%) | | (1.5%, 5.3%) | (2.60, 9.96) | (2.39, 8.57) | (6.7%, 14.6%) | | |
| Visit Week 16 | | | | | | | | |
| 359 | 45 (12.5%) | 365 | 16 (4.4%) | 3.13 | 2.86 | 8.2% | <0.0001+* | |
| | (9.3%, 16.4%) | | (2.5%, 7.0%) | (1.73, 5.64) | (1.65, 4.96) | (4.1%, 12.2%) | | |
| Visit Week 20 | | | | | | | | |
| 359 | 0 (0.0%) | 365 | 0 (0.0%) | 1.02 | 1.02 | 0.0% | NE | |
| | | | | (0.02, 51.38) | (0.02, 51.10) | (-0.5%, 0.5%) | | |
| Visit Week 26 | | | | | | | | |
| 359 | 49 (13.6%) | 365 | 26 (7.1%) | 2.06 | 1.92 | 6.5% | 0.0040+* | |
| | (10.3%, 17.6%) | | (4.7%, 10.3%) | (1.25, 3.40) | (1.22, 3.01) | (2.1%, 11.0%) | | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline POEM 0.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point. The 95% confidence intervals for the event rates are exact Clopper-Pearson intervals.

Binary Outcome Analysis: Achieving 0-1 in DLQI total score - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 20 | brocitinib 200mg QD [DARE] Dupilumab 300mg Q2W [DARE] | | | Abr | ocitinib 200mg QD [DARE] vs. Dupilo | ımab 300mg Q2W [DARE] | |
|----------------|---|-----|----------------|---------------|-------------------------------------|-----------------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| Visit Week 2 | | | | | | | |
| 358 | 81 (22.6%) | 361 | 23 (6.4%) | 4.30 | 3.55 | 16.3% | <0.0001+* |
| | (18.4%, 27.3%) | | (4.1%, 9.4%) | (2.63, 7.01) | (2.29, 5.51) | (11.2%, 21.3%) | |
| Visit Week 4 | ļ | | | | | | |
| 358 | 3 (0.8%) | 361 | 1 (0.3%) | 3.04 | 3.03 | 0.6% | 0.3123+ |
| | (0.2%, 2.4%) | | (0.0%, 1.5%) | (0.31, 29.39) | (0.32, 28.95) | (-0.5%, 1.6%) | |
| Visit Week 8 | ; | | | | | | |
| 358 | 0 (0.0%) | 361 | 0 (0.0%) | 1.01 | 1.01 | 0.0% | NE |
| | | | | (0.02, 50.96) | (0.02, 50.68) | (-0.5%, 0.5%) | |
| Visit Week 1 | 2 | | | | | | |
| 358 | 126 (35.2%) | 361 | 110 (30.5%) | 1.24 | 1.16 | 4.7% | 0.1777+ |
| | (30.2%, 40.4%) | | (25.8%, 35.5%) | (0.91, 1.69) | (0.94, 1.42) | (-2.1%, 11.6%) | |
| Visit Week 1 | .6 | | | | | | |
| 358 | 147 (41.1%) | 361 | 107 (29.6%) | 1.65 | 1.39 | 11.4% | 0.0014+* |
| | (35.9%, 46.4%) | | (25.0%, 34.6%) | (1.21, 2.25) | (1.13, 1.69) | (4.5%, 18.4%) | |
| Visit Week 2 | 0 | | | | | | |
| 358 | 154 (43.0%) | 361 | 119 (33.0%) | 1.54 | 1.30 | 10.1% | 0.0055+* |
| | (37.8%, 48.3%) | | (28.1%, 38.1%) | (1.13, 2.08) | (1.08, 1.58) | (3.0%, 17.1%) | |
| Visit Week 2 | 6 | | | | | | |
| 358 | 137 (38.3%) | 361 | 114 (31.6%) | 1.34 | 1.21 | 6.7% | 0.0601+ |
| | (33.2%, 43.5%) | | (26.8%, 36.6%) | (0.99, 1.83) | (0.99, 1.48) | (-0.3%, 13.6%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Number of subjects: Full Analysis Set Safety Population, excluding subjects with baseline DLQI 0-1.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.

Binary Outcome Analysis: Achieving 0-1 in PP-NRS total score - Full Analysis Set Safety Population JADE DARE (PF-04965842) - 2023 datacut

| Abrocitinib 200mg QD [DARE] | | Dupilumab 300mg Q2W [DARE] | | Abrocitinib 200mg QD [DARE] vs. Dupilumab 300mg Q2W [DARE] | | | |
|-----------------------------|----------------|----------------------------|----------------|--|--------------|----------------|-----------------|
| | Events (%) | | Events (%) | OR | RR | RD | |
| n | (95% CI) | n | (95% CI) | (95% CI) | (95% CI) | (95% CI) | CMH p-value [+] |
| Overall | | | | | | | |
| - Visit Week 2 | | | | | | | |
| 362 | 62 (17.1%) | 365 | 17 (4.7%) | 4.23 | 3.68 | 12.5% | <0.0001+* |
| | (13.4%, 21.4%) | | (2.7%, 7.4%) | (2.42, 7.39) | (2.19, 6.16) | (8.0%, 16.9%) | |
| - Visit Week 4 | | | | | | | |
| 362 | 101 (27.9%) | 365 | 41 (11.2%) | 3.06 | 2.48 | 16.7% | <0.0001+* |
| | (23.3%, 32.8%) | | (8.2%, 14.9%) | (2.05, 4.55) | (1.78, 3.46) | (11.0%, 22.3%) | |
| - Visit Week 8 | | | | | | | |
| 362 | 125 (34.5%) | 365 | 76 (20.8%) | 2.01 | 1.66 | 13.7% | <0.0001+* |
| | (29.6%, 39.7%) | | (16.8%, 25.4%) | (1.44, 2.80) | (1.30, 2.12) | (7.3%, 20.1%) | |
| - Visit Week 1 | 2 | | | | | | |
| 362 | 135 (37.3%) | 365 | 91 (24.9%) | 1.79 | 1.50 | 12.4% | 0.0003+* |
| | (32.3%, 42.5%) | | (20.6%, 29.7%) | (1.30, 2.46) | (1.20, 1.87) | (5.7%, 19.0%) | |
| - Visit Week 1 | 6 | | | | | | |
| 362 | 130 (35.9%) | 365 | 99 (27.1%) | 1.51 | 1.32 | 8.8% | 0.0108+* |
| | (31.0%, 41.1%) | | (22.6%, 32.0%) | (1.10, 2.06) | (1.07, 1.65) | (2.1%, 15.5%) | |
| - Visit Week 2 | 0 | | | | | | |
| 362 | 139 (38.4%) | 365 | 98 (26.8%) | 1.70 | 1.43 | 11.5% | 0.0009 + * |
| | (33.4%, 43.6%) | | (22.4%, 31.7%) | (1.24, 2.32) | (1.15, 1.77) | (4.8%, 18.3%) | |
| - Visit Week 2 | 6 | | | | | | |
| 362 | 139 (38.4%) | 365 | 114 (31.2%) | 1.37 | 1.23 | 7.2% | 0.0427+* |
| | (33.4%, 43.6%) | | (26.5%, 36.3%) | (1.01, 1.86) | (1.01, 1.50) | (0.3%, 14.1%) | |

Notes:

Number of subjects: Full Analysis Set Safety Population

Analysis on overall population is calculated based on stratified CMH (Cochran-Mantel-Haenszel) models, for OR, RR, and RD, stratified by by disease activity (moderate, severe) at enrollment.

Non-responder imputation is applied: subjects without a value at the corresponding visit is considered a non-responder. If a subject withdrew from the study or used rescue therapy, then this subject was counted as non-responder after that point.