Population Pharmacokinetic Analysis of Pexidartinib in Healthy Subjects and Patients With Tenosynovial Giant Cell Tumor (TGCT) or Other Solid Tumors

Ophelia Yin,¹ Jia Kang,² William Knebel,² Hamim Zahir,¹ Michiel van de Sande,³ William D. Tap,⁴ Hans Gelderblom,³ Silvia Stacchiotti,⁵ Jon Greenberg,⁶ Dale Shuster,⁶ Andrew J. Wagner⁷

¹Quantitative Clinical Pharmacology and Translational Sciences, Daiichi Sankyo, Inc., Basking Ridge, NJ, USA; ²Metrum Research Group, Tariffville, CT, USA; ³Leiden University Medical Center, Leiden, Netherlands; ⁴Memorial Sloan Kettering Cancer Center and Weill Cornell Medical College, New York, NY, USA; ⁵Fondazione IRCCS Istituto Nazionale dei Tumori, Milan, Italy; ⁶Global Oncology R&D, Daiichi Sankyo, Inc., Basking Ridge, NJ, USA; ⁷Dana-Farber Cancer Institute, Boston, MA, USA

INTRODUCTION

Background

- Pexidartinib is a novel oral small-molecule inhibitor that selectively targets colony-stimulating factor 1 receptor (CSF1R), KIT proto-oncogene receptor tyrosine kinase (KIT), and FMSlike tyrosine kinase 3 (FLT3) harboring an internal tandem duplication mutation^{1,2}
- Pexidartinib has demonstrated significant tumor response
- Observed concentration-time profiles of pexidartinib showed:
- A bi-exponential decay of pexidartinib concentrations, viewed on a semi-log scale plot (Figure 1a)
- Steady-state concentrations of pexidartinib were maintained over time upon multiple dosing (**Figure 1b**)

Figure 1. Observed Concentration-Time Profiles of Pexidartinib in (a) Healthy Subjects Following Single Oral Dose of 200 mg to 2400 mg and (b) Study PLX108-01 and ENLIVEN Patients

- Final full model included the following covariate-parameter relationships (**Table 2**):
 - Body weight (WT); sex; race (Asian [N = 8] vs. non-Asian [N = 367]); baseline values of CRCL, AST, and TBIL; and study population (healthy subjects vs. patients) on CL/F
 - WT on Q/F, V2/F, and V3/F
 - A formulation effect on F1 was fixed in the model to account for a 17% higher observed pexidartinib exposure with the phase 3 formulation compared with the phase 1 formulation

and improvements in function in patients with symptomatic tenosynovial giant cell tumor (TGCT) that is associated with severe morbidity or functional limitations not amenable to improvement with surgery³

Objective

- To develop a population pharmacokinetics (PK) model for pexidartinib in healthy subjects and patients with TGCT
- To identify and estimate effects of potential covariates, such as demographic and clinical factors, that affect pexidartinib PK

METHODS

Data Source and Study Design

- Data were from nine clinical studies with a total of 375 subjects who contributed a total of 8430 PK samples (Table 1)
 - Seven phase 1 clinical pharmacology studies in healthy subjects (N = 159), with serial PK samples collected up to 144 hours or 192 hours post-dose
 - Phase 1 Study PLX108-01 in patients with TGCT and other solid tumors (N = 132), with 5 to 6 PK samples per patients collected on Cycle 1 Day 1 and Cycle 1 Day 15, and predose PK samples on Days 1, 8, and 16 of Cycle 1
 - Phase 3 Study PLX108-10 (ENLIVEN) in patients with TGCT (N = 84), with 7 PK samples collected on Cycle 1 Day 15, and random samples on Cycle 3 Day 1 and Cycle 5 Day 1
- Serum concentrations of pexidartinib were determined by a validated liquid chromatography-tandem mass spectrometry method, with lower limit of quantification (LLOQ) of 2.5 ng/mL





RESULTS

Table 2. Parameter Estimate From the FinalPopulation PK Model for Pexidartinib

| Parameter CL/F(exp(θ1)) | Estimate* | 95%Cl** |
|--|-------------------------|-------------------|
| $CL/F(exp(\theta_1))$ | 5.83 L/hr | (5.43, 6.27) |
| · (<i>WT</i> /80) ^{0.75} | | |
| · (<i>CRCL</i> _{<90} /90) ^{<i>θ</i>⁹} | -0.0941 | (-0.402, 0.214) |
| · (Asian)·exp(θ_{10}) | 1.27 | (1.05, 1.54) |
| · (<i>AST</i> _{>80} /80) ^{θ11} | 0.0709 | (-0.180, 0.322) |
| · (<i>TBIL</i> _{>20.5} /20.5) ^{θ12} | 0.244 | (0.183, 0.306) |
| · (<i>StHT</i>)·exp(θ_{13}) | 1.26 | (1.16, 1.36) |
| · (<i>Female</i>)·exp(θ_{14}) | 0.869 | (0.808, 0.934) |
| $V_c/F(\exp(\theta_2))$ | 98.0 L | (90.0, 107) |
| · (WT /80) ¹ | | |
| $V_{p}/F(\exp(\theta_{3}))$ | 116 L | (106, 128) |
| · (<i>WT</i> /80) ¹ | | |
| $Q/F(\exp(\theta_4))$ | 20.7 L/hr | (17.9, 23.8) |
| · (<i>WT</i> /80) ^{0.75} | | |
| $KA(\exp(\theta_5))$ | 6.82 hr-1 | (5.09, 9.14) |
| $ALAG1(exp(\theta_6))$ | 0.387 hr | (0.385, 0.390) |
| $D1(\exp(\theta_7))$ | 1.22 hr | (1.20, 1.25) |
| $F1_{Phase1}(\exp(\theta_8))$ | 0.855 Fixed | |
| Ω _{1.1} CL/F | 0.0860 (%CV = 30) | (0.0633, 0.109) |
| $\Omega_{2.1}COV_{V_c/F-CL/F}$ | 0.0774 (corr = 0.504) | (0.0425, 0.112) |
| Ω _{2.2} V _c /F | 0.274 (%CV = 56.1) | (0.207, 0.341) |
| $\Omega_{3.1}COV_{V_p/F-CL/F}$ | 0.0149 (corr = 0.110) | (-0.0178, 0.0476) |
| $\Omega_{3.2}COV_{V_p/F-V_c/F}$ | -0.0467 (corr = -0.193) | (-0.105, 0.0111) |
| Ω _{3.3} V _p /F | 0.213 (%CV = 48.8) | (0.152, 0.275) |
| Ω _{4.4} Q/F | 0.406 (%CV = 70.8) | (0.271, 0.541) |
| Ω _{5.5} KA | 1.31 (%CV = 165) | (0.648, 1.98) |
| Ω _{6.6} Ph1Form | 0.101 (%CV = 32.6) | (0.0592, 0.143) |
| Ω _{7.7} ΙΟV <i>ΚΑ(η</i> ₇₋₁₁) | 1.83 (%CV = 229) | (1.26, 2.40) |
| Ω _{12.12} IOV <i>F1(η</i> ₁₂₋₂₁) | 0.0652 (%CV = 25.9) | (0.0560, 0.0743) |
| Σ _{1.1,prop,pat (ε1)} | 0.0883 (%CV = 29.7) | (0.0839, 0.0927) |
| $\Sigma_{2.2,\text{prop,ht}(\epsilon_2)}$ | 0.0384 (%CV = 19.6) | (0.0377, 0.0391) |

- Covariate effects on pexidartinib exposure are illustrated by the forest plots (Figure 2):
- Model estimated a 21% lower steady-state AUC from 0-24 hours (AUC_{0-24,ss}) of pexidartinib in healthy subjects compared with patients
- Effect of race (Asian vs. non-Asian) on AUC_{0-24,ss} fell partially outside the range of 80% to 125%, but with a wide 95% confidence interval
- All other covariates (sex, CRCL, AST, TBIL) showed a less than 20% effect on pexidartinib exposure, except WT, for which a low value of 53 kg resulted in an approximately 36% increase in AUC_{0-24,ss} compared with the median value of 80 kg
- Current analysis dataset included a relatively narrow range of CRCL values and a small number of renally impaired subjects, which have limited the evaluation of CRCL effect
- Covariate effects on steady-state C_{max} ($C_{max,ss}$) are similar to those on AUC_{0-24,ss}
- Visual predictive check suggested that the final model described the observed data and was suitable for simulation (Figure 3)

Figure 2. Forest Plot of Covariate Effects on Pexidartinib (a) AUC_{0-24,ss} and (b) C_{max,ss}

Table 1. Summary of Studies Included in Population PKAnalysis

| Study | Phase | N | Number of PK Samples | Description | Dose Regimen |
|-----------|-------|-----|-------------------------|---|---|
| U114 | 1 | 30 | 1728 | Relative BA study in HV | 400 mg single doses |
| U116 | 1 | 36 | 1824 | Relative BA study in HV | 600 mg single doses |
| U117 | 1 | 18 | 1119 | Dose proportionality study in HV | 200, 400, and 600 mg single doses |
| U118 | 1 | 16 | 334 | Drug-drug interaction with itraconazole in HV | 600 mg single doses |
| U119 | 1 | 16 | 333 | Drug-drug interaction with rifampin in HV | 600 mg single doses |
| U120 | 1 | 16 | 323 | Drug-drug interaction with esomeprazole in HV | 600 mg single doses |
| U121 | 1 | 27 | 589 | Food effect study in HV | 1200, 1800, 2400 mg |
| PLX108-01 | 1 | 132 | 1726 | Dose-ranging study in patients with TGCT or other solid tumor | 200 to 1200 mg/day |
| ENLIVEN | 3 | 84 | 454 | Phase 3 study in patients with TGCT | Part 1: 1000 mg/day for 2 weeks, followed by 800 mg/day Part 2: 800 mg/day |

D Reference Subjec Reference subject WT: 53 WT: 5 WT: 69 WT: WT: 8 WT: 89 WT: 108 WT: 108 CRCL: 54. CRCL: 54.7 CRCL: 66.4 CRCL: 66.4 0.997 CRCL: 86 CRCL: 86 CRCL: 89.3 CRCL: 89.2 0.999 AST: 8 AST: 81 0.994 AST: 8 AST: 88 0.979 AST: 11 AST: 111 0.957 AST: 15 AST: 159 -TBIL: 20.5 TBIL: 20.52 0.981 TBIL: 22.2 TBIL: 22.23 TBIL: 27.3 TBIL: 27.36 TBIL: 44.4 TBIL: 44.46 Race: Asia Race: Asian Sex: Female Sex: Female Phase 1 studies in healthy subject Phase 1 Studies in healthy subjects 0.6 0.8 1.0 1.2 1.4 0.6 0.8 1.0 1.2 1.4 Relative C_{max} s Relative AUC_{0-24 se}

AUC_{0-24,ss} and C_{max,ss} were derived for the dose of 800 mg/day. The reference is a male, non-Asian patient with median covariate values of WT 80 kg, CRCL \geq 90 mL/min, AST \leq 80 U/L, and TBIL \leq 20.5 µmol/L. The values shown next to WT, CRCL, AST, and TBIL were the 5th, 25th, 75th, and 95th percentiles of observed values of each covariate. Dots and solid horizontal lines are median and 95% CI of (a) relative AUC_{0-24,ss} and (b) relative C_{max,ss}. The gray-shaded region represents a range of 80% to 125%.

AST = aspartate aminotransferase (U/L), AUC_{0-24,ss} = steady-state area under the plasma concentration-time curve from 0 to 24 hours, $C_{max,ss}$ = steady-state maximum plasma concentration, CRCL = creatinine clearance (mL/min), TBIL = total bilirubin (µmol/L), WT = body weight (kg).

Phase 1 formulation was used in studies PLX108-01 and U114, whereas phase 3 formulation was used in all other studies.

BA = bioavailability, HV = healthy volunteers, N = number of subjects, PK = pharmacokinetics, TGCT = tenosynovial giant cell tumor.

Population PK Analysis

- The structural PK model was a two-compartment model with sequential zero- and first-order absorption and lag time, and linear elimination from the central compartment
- Inter-individual variability (IIV) was included on clearance from central compartment (CL/F), central volume of distribution (V2/F), peripheral volume of distribution (V3/F), inter-compartmental clearance (Q/F), and absorption rate constant (Ka)

Figure 3. Visual Predictive Check Plots for (a) Studies in Healthy Subjects, (b) Study PLX108-01, and (c) ENLIVEN Study



*Estimates of θ modeled in the log domain were exponentiated and are reported in the table. **95% CI was derived from standard errors obtained from the NONMEM \$COVARIANCE step. AST = aspartate aminotransferase (U/L), ALAG1 = lag time, CI = confidence interval, CL/F = apparent clearance, CV = coefficient of variation, corr = correlation, COV = covariance, CRCL = creatinine clearance (mL/min), D1 = duration of zero-order deposition, F1_{Phase1} = relative bioavailability of Phase 1 formulation to Phase 3 formulation, IOV = inter-occasion variability (variance), KA = first-order absorption rate constant, PK = pharmacokinetic, Q/F = apparent inter-compartmental clearance, StHT = Phase 1 studies with healthy subjects, TBIL = total bilirubin (µmol/L), Vc/F = apparent central compartment volume, Vp/F = apparent peripheral compartment volume, WT = body weight (kg), θ = fixed effect parameter, Ω = inter-individual covariance matrix, $\Omega_{6.6}$ Ph1Form = inter-individual variability of F1 for the Phase 1 formulation, $\Sigma_{1.1,prop,pat}$ = proportional residual variability for studies in patients, $\Sigma_{1.1,prop,ht}$ = proportional residual variability for Studies in patients, $\Sigma_{1.1,prop,ht}$ = proportional residual variability for Studies in patients, $\Sigma_{1.1,prop,ht}$ = proportional residual variability for Phase 1 studies in healthy subjects.

 For the dose regimen of 800 mg/day (400 mg BID) in the ENLIVEN Study, the mean of individual pexidartinib AUC₀₋₁₂ was 77465.1 hr*ng/mL at steady state and 21529.3 hr*ng/mL on Day 1, representing an accumulation ratio of 3.5 (Table 3)

Table 3. Post Hoc PK Parameters of Pexidartinib in the ENLIVEN Study

| PK variables | Mean (SD) | Median (P5-P95) |
|--|--|--|
| AUC _{0-12h} (hr*ng/mL) on Day 1 | 21529.3 (5231.1) | 20737.7 (14011.5-30731.2) |
| AUC _{0-12h} (hr*ng/mL) at steady state | 77465.1 (24974.6) | 72462.7 (47845.9-127464.3) |
| C _{max} (ng/mL) on Day 1 | 3523.6 (1093.1) | 3247.8 (2188.4-5384.6) |
| C _{max} (ng/mL) at steady state | 8625.3 (2745.7) | 7992.8 (5373.6-13834.1) |
| | PK variablesAUC_0-12h (hr*ng/mL) on Day 1AUC_0-12h (hr*ng/mL) at steady stateAUC_0-12h (hr*ng/mL) at steady stateCmax (ng/mL) on Day 1Cmax (ng/mL) at steady state | PK variables Mean (SD) AUC _{0-12h} (hr*ng/mL) on Day 1 21529.3 (5231.1) AUC _{0-12h} (hr*ng/mL) at steady state 77465.1 (24974.6) C _{max} (ng/mL) on Day 1 3523.6 (1093.1) C _{max} (ng/mL) at steady state 8625.3 (2745.7) |

- Inter-occasion variability (IOC) was added to Ka and relative bioavailability (F1)
- A proportional residual error model was used to describe the residual variability
- The following covariates were evaluated on the absorption and/or disposition parameters using a full-model approach:
 - Subject demographics (age, sex, body weight, and race)
 - Liver and renal function parameters (aspartate aminotransferase [AST], alanine aminotransferase [ALT], total bilirubin [TBIL], serum creatinine clearance [CRCL])
 - Drug formulations (phase 1 vs. phase 3 formulation)
 - Study populations (healthy volunteers vs. patients)
- The final full model was assessed by goodness-of-fit plot and visual predictive check

PK Simulations

 Based on the final population PK model and individual post hoc PK parameters, the area under the plasma concentration-time curve (AUC) and maximum plasma concentration (C_{max}) were generated for the dose regimen of 800 mg/day (400 mg BID) in the ENLIVEN Study



data. Blue-shaded area represents the 95% prediction band of the simulated median, and red-shaded areas represent the 95% prediction bands of the simulated 5th and 95th percentiles

| 800 mg/day | Accumulation ratio | 3.6 (0.8) | 3.5 (2.7-4.5) |
|------------|--------------------|-----------|---------------|
| 800 mg/day | CL/F (L/hr) | 5.6 (1.6) | 5.5 (3.1-8.4) |

 AUC_{0-12h} = area under the plasma concentration-time curve from zero to 12 hours, C_{max} = maximum plasma concentration, CL/F = apparent clearance, SD = standard deviation, PK = pharmacokinetic, P5 = 5th percentile, P95 = 95th percentile.

CONCLUSIONS

 A population PK model was successfully developed for pexidartinib. No clinically meaningful effects on pexidartinib exposure were identified for demographic characteristics, such as WT, sex, race, study population (healthy subjects vs. patients), and renal and hepatic functional parameters

The observed pexidartinib concentrations (ng/mL) are plotted (black dots) versus time after dose (hr) with the 5th (red dashed line), 50th (blue line), and 95th (red dashed line) percentiles of the observed

- Evaluation of renal function effect was limited by the analysis dataset, which included a relatively narrow range of CRCL values and a limited number of renally impaired subjects. Therefore, results from a dedicated renal impairment study should be considered in conjunction with current analysis for dose recommendations in renal impairment subjects
- The model was also used to generate individual exposure metrics in subsequent exposureresponse analyses

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