POPULATION PHARMACOKINETICS REPORT

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Study Drug:	Drug A
Indication(s):	Nothing
Study Number(s):	12345
Sponsor:	
Prepared By:	MMJ
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Approved By:	MMJ
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Figures in-text

Population PK Report	
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Tables in-text

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1.

2. EXECUTIVE SUMMARY

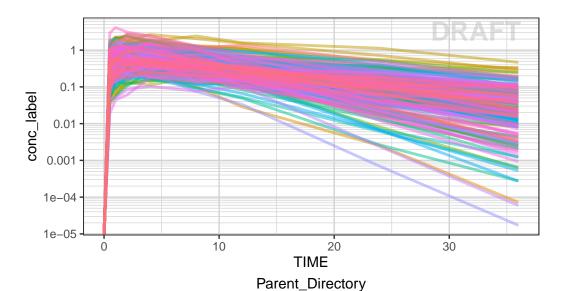
Drug X, an oral tablet which is being developed by company A and is used to treat Disease Y, has undergone N number of completed studies. A population PK analysis was performed to characterize the PK and identify sources of variability in the PK based on rich and sparse samples collected in Phase 1 and Phase 3 studies. As part of the population PK analysis, data from NN subjects was utilized where the doses were ranging from 5 mg- 100 mg Q3W. The data was adequately described using a 1-compartment model with a first-order absorption rate constant (Ka) with lag time (Tlag). A bootstrap method resulted in model reduction compared to reducing the full model with the additional of all the covariates like (age, sex, baseline body weight, race, baseline GFR, drug product) by removing covariates for which the 95Final Model-based simulations were performed to evaluate drug X CL under various conditions, estimate effective half-life, predict exposure metrics for 60 mg Q4W vs 30 mg Q2W dose regimens, and assess the clinical relevance of covariates of interest such as sex, hepatic function, renal function, race, manufacture process, and shorter infusion time in the final PPK model. Results suggest that exposures were higher in female subjects than males for subjects who received 60 mg Q4W. The predicted geometric means of drug X exposure (Cmin1, Cmax1, Cavg1, Cavd28, Cminss, Cmaxss, and Cavgss) at 60 mg Q4W and 30 mg Q2W are summarized in Table 5. As expected, Cavgss was similar across the two different regimens (difference < 5

SYNOBSIS

3. INTRODUCTION

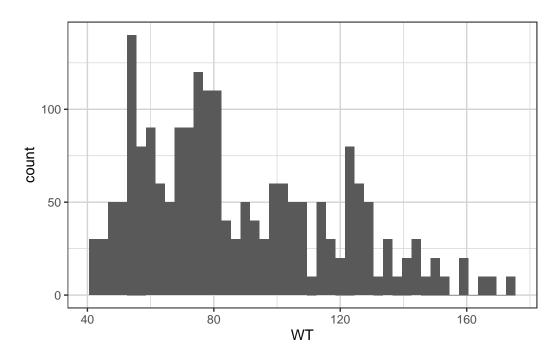
3.1. Exploratory Data Analysis

3.2. PK Plots



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3.3. Covariate Summary



4. METHODS

5. RESULTS

For each analysis (e.g. PopPK, PK/PD analysis, exposure-response analysis and simulations), an own subsection should be included.

5.1. Exploratory Data Analysis

Concentration-time profiles of drug X following oral administration of the first dose on Day 1 in healthy subjects are presented in Figure 2. Additional concentration-time profiles of drug X are presented in Appendix 2. A list of samples excluded from the analysis is presented in Appendix 2. Drug X was rapidly absorbed following oral administration and declined in a multi-exponential manner. Doses ranged from 5 mg to 100 mg. Drug X exposure was dose proportional, and the accumulation ratio based on 80 mg once daily dosing was 1.3 in Study 123.

5.2. Model Development

5.2.1. Base Model

A population PK analysis was performed based on rich and sparse samples collected in Phase 1 and Phase 3 studies in order to identify the structural model. Highlights of the base population PK analysis are presented below and in Appendix 2. 1- and 2-compartment models with linear elimination were tested. The 1-compartment model resulted in the lowest OFV. A first-order absorption rate constant (Ka) with lag time (Tlag) was used to characterize the rapid absorption of drug X. A mixed error model (additive and proportional) resulted in a substantially lower OFV relative to proportional or additive error models. Additional model refinements are presented below. An allometric function accounting for body weight effect on clearance (CL/F) and volume of distribution (V/F) was included in the model (run008). In addition, the effect of creatinine clearance was added on CL/F since the drug was previously demonstrated to undergo important renal excretion (run019).

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Reading ./run1.phi

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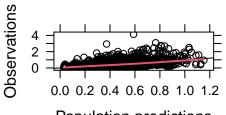
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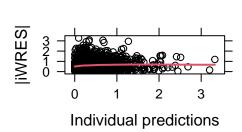
Estimates for \$prob	no.1,	subprob n	0.1,	method	foce
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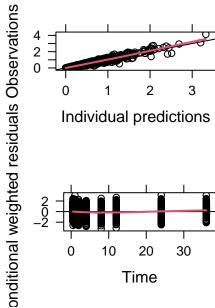
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THETA1	CL	9.996		0.03791
THETA2	V	109.1		0.04173
THETA3	KA	0.9377		0.04867
THETA4	PROP ERR	0.2021		0.02284
OMEGA(1,1)	PPV_CL	0.5285		0.04845
OMEGA(2,1)		0.5713		0.07815
OMEGA(2,2)	PPV_V	0.5701		0.04943
OMEGA(3,3)	PPV_KA	0.642		0.07221
SIGMA(1,1)		1	fix	_

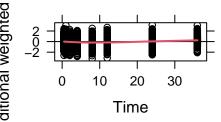
Basic goodness-of-fit plots (Run 1)



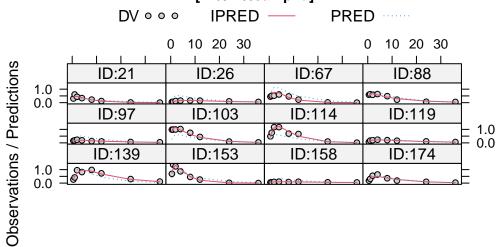
Population predictions





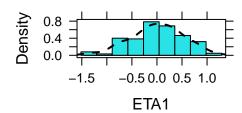


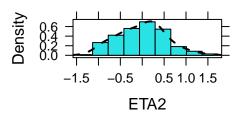
Individual plots (Run 1) [ID%in%samp.id]

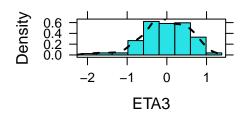


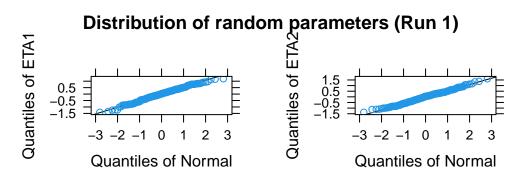
Time

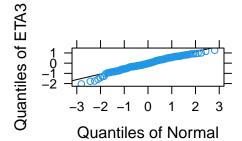
Distribution of ETAs (Run 1)



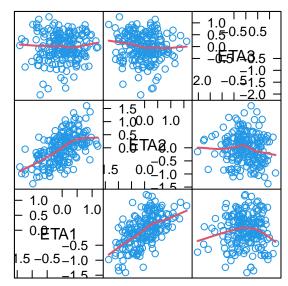




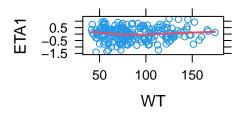


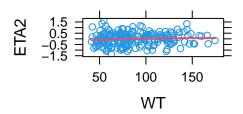


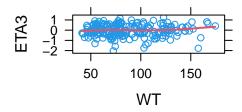
Scatterplot matrix of random parameters (Run 1)



Parameters vs. covariates (Run 1)







5.2.2. Covariate Model

Potential relationships between PK parameters (random effects) of drug X and categorical and continuous covariates from the base PK model (run019) are presented in Appendix 2. A stepwise covariate analysis was performed to identify sources of variability in PK parameters of drug X. Results for covariate analysis are presented in Appendix 2. A summary of covariates resulting in the maximum reduction of the OFV and included in each step of the analysis is presented in Table 6.

Covariates were evaluated using a forward inclusion approach with p<0.01 (Δ OFV>6.6349). The effect of fasted status on Ka resulted in the most important decrease in OFV as part of the first step of the analysis $(\Delta OVF = -322.051)$. In the second step, the effect of dose on CL/F resulted in the most important decrease in OFV (\triangle OVF = -161.224). In the 3rd and 4th steps, the effect of gender on V/F and CL/F resulted in the most important decrease in OFV (\triangle OVF = -50.805 and -72.726, respectively). In the 5th step, the effect of ESRD on CL/F resulted in the most important decrease in OFV (Δ OVF = -16.624). In the 6th step, the effect of dose on Ka resulted in the most important decrease in OFV (Δ OVF = -71.311). In the 7th step, the effect of formulation Ka resulted in the most important decrease in OFV ($\triangle OVF = -21.636$). In the 8th step, the effect of disease status (healthy subjects vs. narcolepsy/OSA patients) resulted in the most important decrease in OFV (Δ OVF = -17.684). Additional information is available in Appendix 2 (Section 12.36). During the backward testing, none of the covariate were removed. Additional information is available in Appendix 2. The bootstrap method was used to reduce the full model by removing covariates for which the 95% PIs included the null value relative to the reference population. Based on the estimates of the population PK model, concentration-time profiles of drug X were simulated (1000 replicates). Statistically significant covariates were retained in the reduced final model if the nonparametric 95% PIs excluded the null value relative to the reference population. All the covariates tested in the full model resulted in a statistically significant effect, and were retained in the final model.

Table 5.1.: Covariate Model Results

Step	Covariates	Base_OFV	New_OFV	ΔOFV
1	Fasted status on Ka	93576	93254	-322
2	Dose on CL/F	93254	93092	-161
3	Gender on V/F	93092	93042	-50
4	Dose on Ka	93042	92969	-72
5	Formulation on Ka	92881	92859	-21
6	Disease Status on CL/F	92859	92842	-17

5.2.3. Final Model

Typical population PK parameters of drug X derived with the final model (run005) are presented in Table 7. The continuous covariates (CRCL and weight) were centered to a reference value in the population PK analysis (116.5 mL/min and 92.5 kg). The reference value is <1% different than the median value in the Phase 3 studies.

The population estimates of CL/F and V/F for drug X were 19.49 L/h and 198.72 L, respectively, and are for a male patient who is 92.5 kg, has a CRCL of 116.5 mL/min, and is taking a dose of 150 mg. Based on the population PK model, the half-life of drug X was 7.07 h.

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For the OMEGA and SIGMA matrices, values are reported as standard deviations for the diagonal

Estimates for \$prob no.1, subprob no.1, method foce

Parameter	Label	Value	RSE
THETA1	CL	11.35	0.04252
THETA2	V	129.2	0.04959
THETA3	KA	0.9373	0.0487
THETA4	PROP ERR	0.2021	0.02282
OMEGA(1,1)	PPV_CL	0.5954	0.05392
OMEGA(2,1)		0.6779	0.05134
OMEGA(2,2)	PPV V	0.6847	0.04548

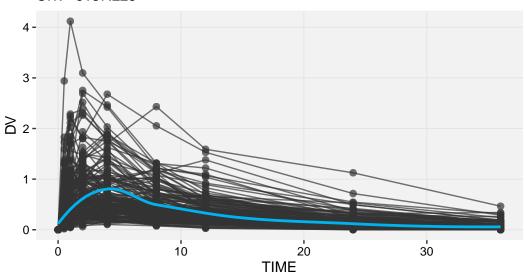
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OMEGA(3,3) PPV_KA 0.6431 0.07359 SIGMA(1,1) 1 fix -

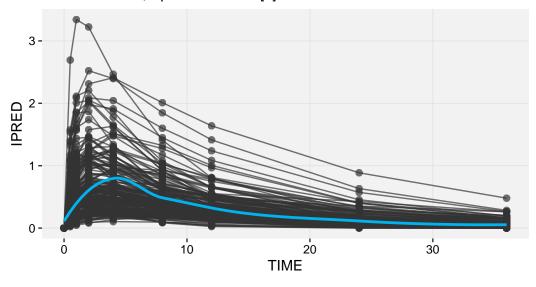
DV vs. TIME | run2

Ofv: -6187.225



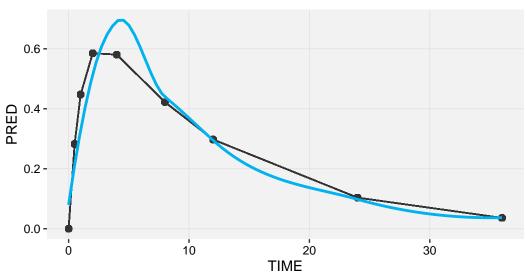
IPRED vs. TIME | run2

Ofv: -6187.225, Eps shrink: 17.1 [1]



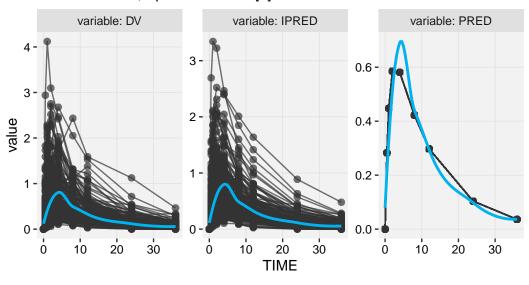
PRED vs. TIME | run2

Ofv: -6187.225



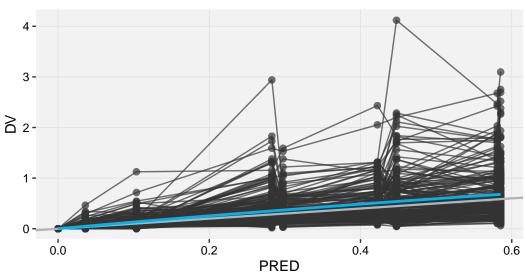
Observations, Individual and Population Predictions vs. TIM

Ofv: -6187.225, Eps shrink: 17.1 [1]



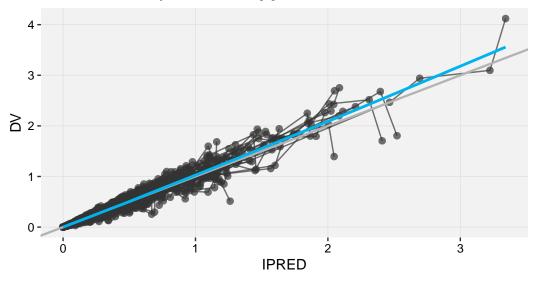
DV vs. PRED | run2

Ofv: -6187.225



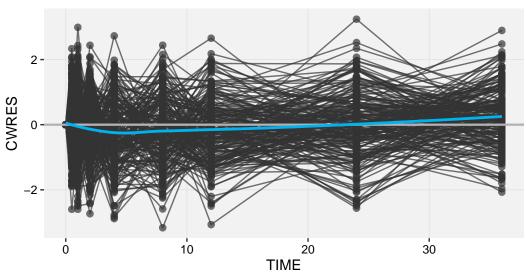
DV vs. IPRED | run2

Ofv: -6187.225, Eps shrink: 17.1 [1]



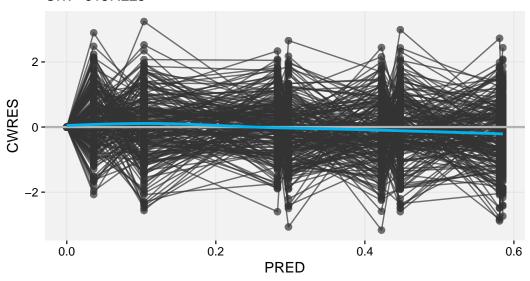
CWRES vs. TIME | run2

Ofv: -6187.225



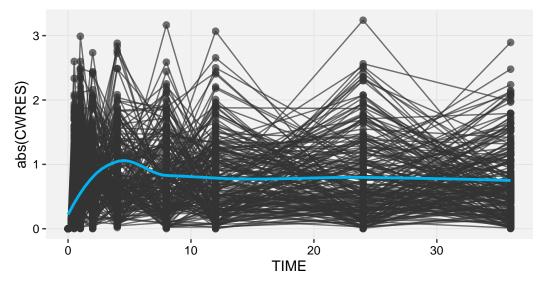
CWRES vs. PRED | run2

Ofv: -6187.225



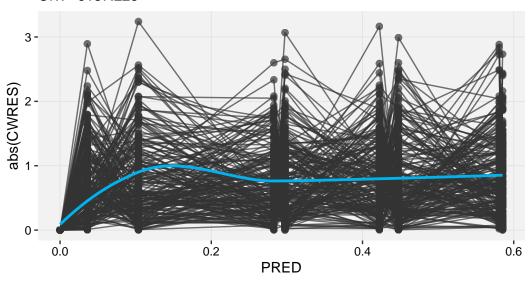
abs(CWRES) vs. TIME | run2

Ofv: -6187.225



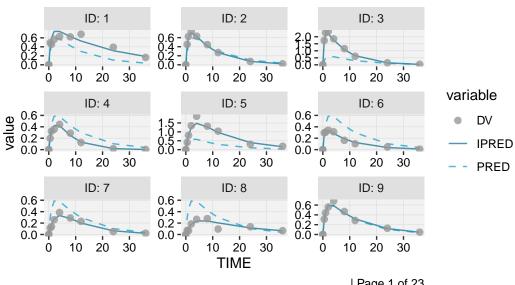
abs(CWRES) vs. PRED | run2

Ofv: -6187.225



Individual plots | run2

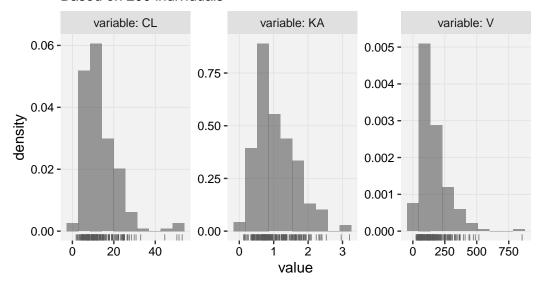
Ofv: -6187.225, Eps shrink: 17.1 [1]



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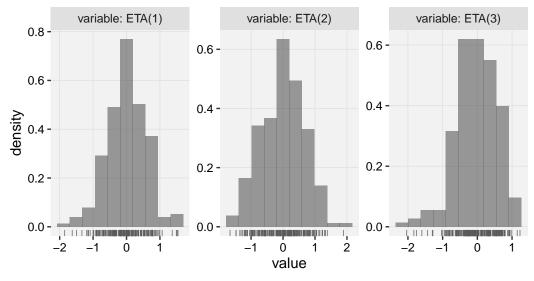
Parameter distribution | run2

Based on 200 individuals



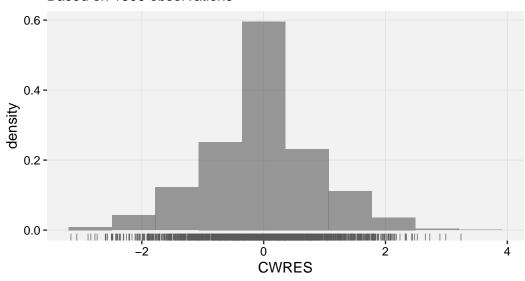
Eta distribution | run2

Based on 200 individuals, Eta shrink: 0.9 [1], 1.9 [2], 10.3 [3]



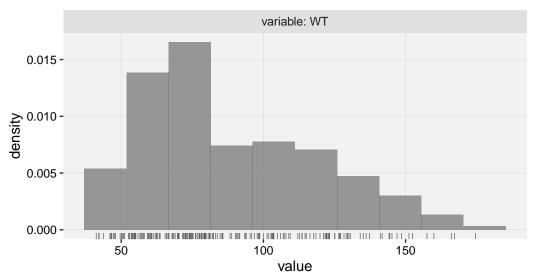
CWRES distribution | run2

Based on 1600 observations



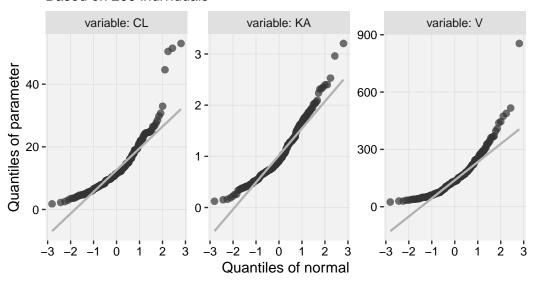
Continuous covariates distribution | run2

Based on 200 individuals



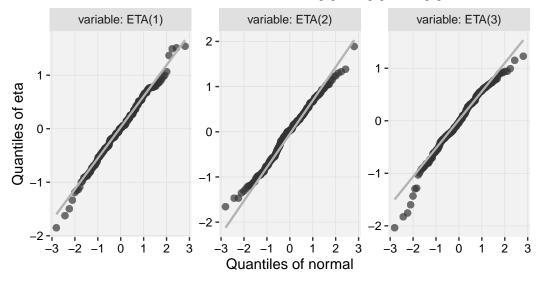
QQ plot of parameters | run2

Based on 200 individuals



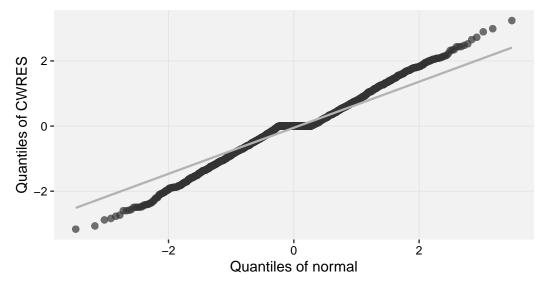
QQ plot of etas | run2

Based on 200 individuals, Eta shrink: 0.9 [1], 1.9 [2], 10.3 [3]



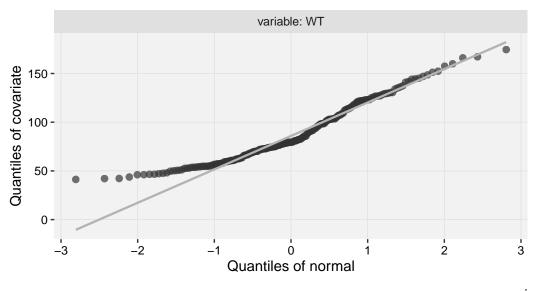
QQ plot of CWRES | run2

Based on 1600 observations



QQ plot of continuous covariates | run2

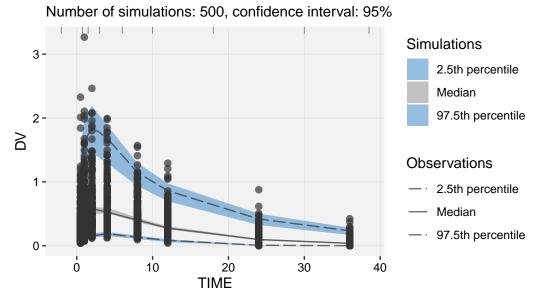
Based on 200 individuals



5.3. Model Evaluation

Visual predictive checks were performed for the full PPK model stratified by patient population and dosing regimens, and the results are in Figure 5. Overall, the pcVPC results indicate that the model adequately characterized the data and predicted concentrations to be used for E-R efficacy and safety analyses. The observed 5th, 50th (median), and 95th percentiles generally fall within the 90% PI (the shaded band) up to the first 60 days after the previous dose and the first 600 days after the first dose (trough concentrations). A pcVPC stratified by body weight is provided in Appendix 3.

Visual predictive checks | run2



_Template/isop-template-ms-main/example_ppk_nonmem_1/vpc_dir5

5.4. Model Application

Model-based simulations were performed to evaluate drug X CL under various conditions, estimate effective half-life, predict exposure metrics for 60 mg Q4W vs 30 mg Q2W dose regimens, and assess the clinical relevance of covariates of interest such as sex, hepatic function, renal function, race, manufacture process, and shorter infusion time in the final PPK model. The final model was used in these simulations.

5.4.1. Covariate Model Evaluation of Effect of Sex

Sex was a significant covariate on CL and VC (Figure 6), with male subjects having a higher CL and higher Vc than female subjects. Comparisons of model predicted drug exposure at 60 mg Q4W by sex are presented in Figure 6. In general, exposures were higher in female subjects than males for subjects who received 60 mg Q4W.

Sex was a significant covariate on CL and VC (Figure 6), with male subjects having a higher CL and higher Vc than female subjects. Comparisons of model predicted drug exposure at 60 mg Q4W by sex are presented in Figure 6. In general, exposures were higher in female subjects than males for subjects who received 60 mg Q4W.

5.4.2. Covariate Model Comparison of 30 mg Q2W and 60 Q4W Regimens

The predicted geometric means of drug X exposure (Cmin1, Cmax1, Cavg1, Cavd28, Cminss, Cmaxss, and Cavgss) at 60 mg Q4W and 30 mg Q2W are summarized in Table 5. As expected, Cavgss was similar across the two different regimens (difference < 5%). The exposures were higher with drug X 30 mg Q2W relative to 60 mg Q4W by approximately 51% for Cmind28 and 42% for Cminss. The exposures were lower with

drug X 30 mg Q2W relative to 60 mg Q4W by approximately 50% for Cmax1 and 31% for Cmaxss, which were also expected.

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6. DISCUSSION

7. CONCLUSION

Summary of major findings. Contextualize the PK model based simulations with regard to therapeutic window ?

8. REFERENCES

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9.

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