#### Model-Informed Drug Development

#### **2021 Virtual Conference**

Efavirenz Physiologically Based Pharmacokinetic Model Development and Validation as a Moderate CYP3A4 Inducer for Drug-Drug Interaction Predictions

#### Inger M. Darling, Joel Owen, and Viera Lukacova Presented at AAPS PharmSci 360 in 2019

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### **Background/Purpose**

- Efavirenz is an antiretroviral medication used to treat and prevent HIV/AIDS.
- Efavirenz is eliminated primarily by CYP2B6 and CYP3A4 oxidative metabolism with further glucuronidation to inactive metabolites. Due to substantial induction of both CYP3A4 and CYP2B6 enzyme systems with daily dosing of efavirenz, dose adjustments for many coadministered medications are necessary [1].
- Strong, moderate, and weak inducers are drugs that decrease the AUC of sensitive index substrates of a given metabolic pathway by ≥ 80%, ≥ 50% to < 80%, and ≥ 20% to < 50%, respectively [2].</li>
- Efavirenz (dosed at 600 mg QD in adults) is a moderate CYP3A4 inducer. As such, a fully validated physiologically based pharmacokinetic (PBPK) model for efavirenz including validated induction potentials could be a useful standard compound for PBPK drug-drug interaction (DDI) simulations.





### Objectives

- Develop and validate a PBPK model for efavirenz in humans using PK data obtained from literature.
- Validate the efavirenz PBPK model as a moderate CYP3A4 inducer for DDI predictions using available induction parameters and observed PK data from DDI studies with CYP3A4 substrate(s) reported in literature.





## Methods

- GastroPlus<sup>®</sup> (Version 9.5) [3] was used to develop and validate the PBPK model and run DDI simulations.
- ADMET Predictor<sup>®</sup> (Version 8.1) [4] was applied to obtain in silico predicted estimates of key physicochemical and biopharmaceutical properties from the chemical structure of efavirenz where experimental values were not found in literature.
- Plasma concentration versus time (Cp-time) profiles from literature sources were acquired through a process of digitization.
- The parameters for the efavirenz PBPK model were derived separately for absorption, distribution, and elimination.
- The Advanced Compartmental Absorption and Transit (ACAT<sup>™</sup>) model was used to describe the intestinal dissolution, absorption, and metabolism of efavirenz [5].
- The PBPK physiologies, including organ weights, volumes, and blood flows, were generated by the Population Estimates for Age-Related Physiology (PEAR Physiology™) module.
- Efavirenz tissue distribution was modeled using a perfusion-limited model for all tissues, and tissue/plasma partition coefficients (Kps) were predicted by the default Lukacova method [6].





## Methods (cont.)

#### Mechanistic Assumptions Used for PBPK Model

- Efavirenz was absorbed from the gut via a passive diffusion process only.
- Efavirenz is metabolized primarily by CYP2B6 and CYP3A4 isozymes. For modeling purposes, metabolism was described using CYP2B6 and CYP3A4 enzymes only. It was assumed that renal elimination was negligible and was not incorporated in the current model [1].
- The induction of CYP2B6 and CYP3A4 by efavirenz was included in the model to predict steady-state efavirenz PK and for DDI simulations.





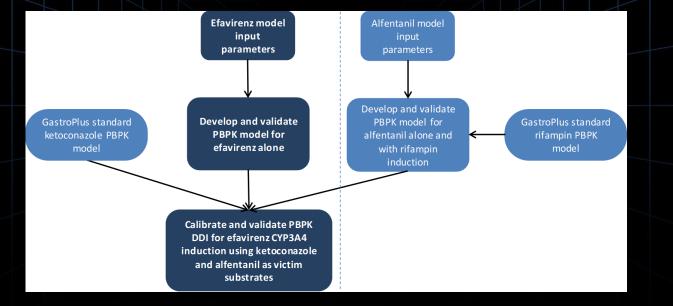
## Methods (cont.)

- Mean particle radius of 5  $\mu m$  was fitted to match the absorption phase and Tmax for the evaluated profiles.
- Literature values for CYP3A4 and CYP2B6 in vitro metabolic Km and Vmax in human liver microsomes [7] were used as starting points for efavirenz metabolism.
- The Vmax values were subsequently adjusted to match the observed efavirenz Cptime profiles for both single- and multiple-dose administrations of efavirenz.
- The initial values for EC50 and Emax were obtained from literature [8].
- The Emax for CYP3A4 was further optimized and validated against data from DDI studies with ketoconazole and alfentanil, respectively.
- The PBPK model for ketoconazole has been previously developed as a standard CYP3A4 substrate and competitive inhibitor for DDI assessments in GastroPlus.



## Methods (cont.)

• The PBPK model for alfentanil as a CYP3A4 substrate was also developed and validated as part of this study. The flow diagram provides an overview of the modeling steps.





# Results





#### Physicochemical and Biopharmaceutical Properties for Efavirenz Model Development

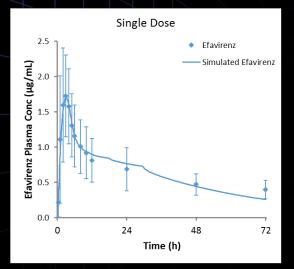
Efavirenz Property	Final Value <sup>a</sup>	
logP (at pH = 7.4)	4.6	
Aqueous Solubility at pH 6.99	9.0 mg/L	
pKa (Acid)	10.2	
Permeability: Caco-2 Papp	$0.25 \text{ cm/s} \times 10^5$	
Permeability: Human Peff	$1.07 \text{ cm/s} \times 10^{4b}$	
Fraction Unbound in Plasma (Human)	0.22%	
Blood/Plasma Concentration Ratio (Human)	0.74	
Mean Particle Radius	5 µm	
CYP3A4 Km	3.137 μg/mL	
CYP3A4 Vmax,PBPK	0.000562 mg/s/mg-enzyme	
CYP3A4 Vmax,GUT	0.2 mg/s	
CYP2B6 Km	3.137 μg/mL	
CYP2B6 Vmax,PBPK	0.050 mg/s/mg-enzyme	
CYP3A4 Emax	9.9	
CYP3A4 EC <sub>50</sub>	1.0 µM	
CYP2B6 Emax	5.76	
CYP2B6 EC50	0.82 µM	
<sup>a</sup> Depresents a mix of experimental predicted and fitted values		

<sup>a</sup> Represents a mix of experimental, predicted, and fitted values.

<sup>b</sup> Caco-2 Papp was converted to human Peff using the built-in ABSCa conversion.



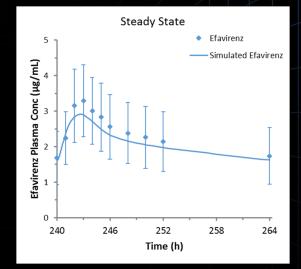
Representative Model-Predicted Mean Single-Dose (600 mg) and Steady-State Efavirenz (600 mg QD) Plasma Concentrations (lines) Versus Observed Mean (SD, points)



CYP2B6 PBPK V<sub>max</sub>: 0.118 mg/s/mg-enzyme Literature plasma concentration data from [9] (Cho, et al., Drug Metab Pharmacokinet. 2016).

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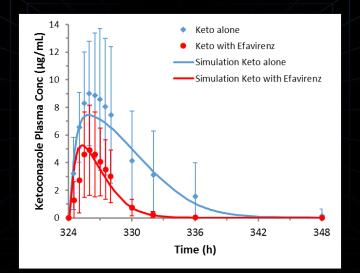
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Final DDI model CYP2B6 V<sub>max</sub>: 0.5 mg/s/mg-enzyme Literature plasma concentration data from [10] (Garg, et al. Br J Clin Pharmacol. 2013).



Efavirenz CYP3A4 Induction Calibration: Mean (SD) Observed (points) and Model-Predicted (lines) Ketoconazole Plasma Concentrations Following a Single 400-mg Dose With (red) and Without (blue) Efavirenz Induction (600 mg QD for 14 days)



Literature data from reference [11] (Sriwiriyajan, et al. Eur J Clin Pharmacol. 2007).



Efavirenz CYP3A4 Induction Validation: Mean (SD) Observed (points) and Model-Predicted (lines) Alfentanil Plasma Concentrations Following Single Doses of Either 15 μg/kg IV Alfentanil (A) or 43 μg/kg PO Alfentanil (B) With (red) and Without (blue) Efavirenz Induction (600 mg QD for 14 Days)

Alfentanil PO alone

Alfentanil with Efavirenz

Simulation Alfentanil PO alone

Simulation Alfentanil with Efavirenz

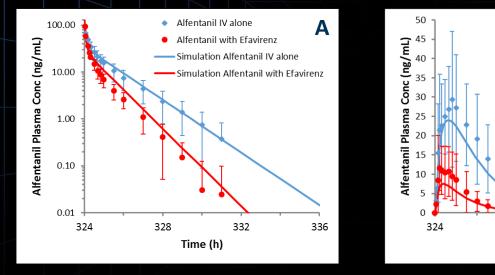
332

328

Time (h)

B

336



Literature data from reference [12] (Kharasch, et al. Clin Pharmacol Ther. 2012).





Efavirenz Induction Model Validation: Summary of Predicted and Observed DDI of Ketoconazole and Alfentanil Single-Dose Administration With and Without Efavirenz Induction (600 mg QD)

		% Reduction With Efavirenz Induction	
Dosage [Reference]	Parameter	Predicted	Observed
Ketoconazole 400 mg PO [11]	Cmax	29	44
	AU Cinf	65	72
Alfentanil 43 µg/kg PO [12]	C <sub>max</sub>	68	57
	AU Cinf	78	78
Alfentanil 15 µg/kg IV [12]	AU Cinf	37	46

[11]: Sriwiriyajan, et al. Eur J Clin Pharmacol. 2007 May;63(5):479-483.

[12]: Kharasch, et al. Clin Pharmacol Ther. 2012 Apr;91(4):673-684.





#### Physicochemical and Biopharmaceutical Properties for Alfentanil Model Development

Alfentanil Property	Final Value <sup>a</sup>
logP (at pH = 7.4)	1.5
Aqueous Solubility	0.87 mg/L
pKa (Base)	6.5
Permeability: Caco-2 Papp	$1.2 \text{ cm/s} \times 10^5$
Permeability: Human Peff	$2.49 \text{ cm/s} \times 10^{4b}$
Fraction Unbound in Plasma (Human)	8.6%
Blood/Plasma Concentration Ratio (Human)	0.66
CYP3A4 Km	6.419 µg/mL
CYP3A4 Vmax, PBPK	0.00129 mg/s/mg-enzyme
CYP3A4 Vmax,GUT	0.452 mg/s

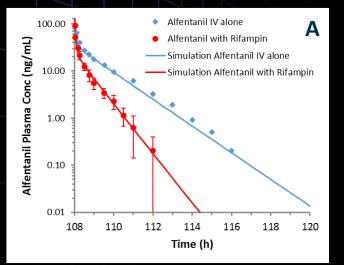
\* Represents a mix of experimental, predicted, and fitted values.

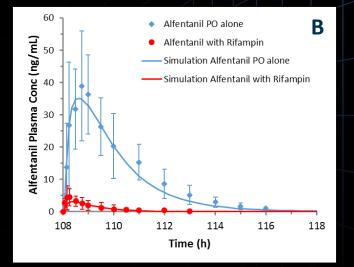
<sup>b</sup> Caco-2 Papp was converted to human Peff using the built-in ABSCa conversion.





Alfentanil Model and DDI Validation: Mean (SD) Observed (points) and Model-Predicted (lines) Alfentanil Plasma Concentrations Following Single Doses of Either 15 μg/mL IV Alfentanil (A) or 60 μg/kg PO Alfentanil (B) With (red) and Without (blue) Rifampin Induction (600 mg QD for 5 days)





Literature data from reference [13] (Kharasch, et al. Clin Pharmacol Ther. 2004).





#### Conclusions

- Developed PBPK model for efavirenz accurately reproduces the efavirenz exposures from published single-dose and steady-state studies.
- DDI simulations with efavirenz as the perpetrator with either ketoconazole or alfentanil as the victim accurately predict the observed exposures due to the induction of CYP3A4 by efavirenz.
- Developed and validated PBPK model for efavirenz is suitable to be used as a standard perpetrator to evaluate the impact of moderate CYP3A4 induction on CYP3A4-metabolized compounds.
- Developed and validated PBPK model for alfentanil is suitable to be used as a standard victim for DDI interaction predictions with CYP3A4 inducers.



#### References

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**Questions & Answers** 

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