

Physiologically-based Pharmacokinetic (PBPK) Modeling of the Combination of Ivacaftor and Lefamulin in Cystic Fibrosis Patients

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INTRODUCTION

- Lefamulin is a first-in-class pleuromutilin antibiotic approved as both IV and oral formulations for the treatment of community-acquired bacterial pneumonia in adults.¹
- The spectrum of activity for lefamulin includes activity against common causes of CAPB, including: *S. pneumoniae*, *Haemophilus influenzae*, *Moraxella catarrhalis*, *Mycoplasma pneumoniae*, *Chlamydia pneumoniae*, *Legionella pneumophila*, and *Staphylococcus aureus* including both methicillin susceptible and methicillin resistant strains.¹
- Given its potent activity against *S. aureus* and the propensity for patients with cystic fibrosis (CF) to be infected with this pathogen, lefamulin is currently being evaluated as a potential treatment for bacterial exacerbations in CF patients caused by *S. aureus*.²
- Lefamulin undergoes hepatic metabolism and is both a substrate and moderate inhibitor of CYP3A.¹
- Ivacaftor is a CF transmembrane conductance regulator (CFTR) modulator that is a sensitive substrate of CYP3A.³
- Given that CFTR modulators, including ivacaftor, have become a foundational therapy for many patients with CF, it is important to evaluate the potential for drug-drug interactions between lefamulin and ivacaftor to support lefamulin's clinical development as a therapeutic for exacerbations in CF patients.
- Through use of PBPK modeling, we characterized the drug-drug interaction potential for lefamulin as compared with itraconazole, fluconazole, and ketoconazole.

METHODS

PBPK Model Development and Validation

- PBPK models were derived and simulated using Version 20 of the Simcyp™ Simulator (www.simcyp.com) utilizing the minimal PBPK model with the non-physiological single adjustable compartment. Absorption was simulated using the Advanced Dissolution, Absorption, and Metabolism (ADAM) model. The chosen minimal PBPK model separates the liver and gastro-intestinal tract from the rest of the body to consider metabolism and inhibition/induction of the enzymes in these organs (Figure 1).
- Lefamulin:** A PBPK model for lefamulin was previously developed from clinical and in vitro data and validated against observed single and multiple dosing regimens (Data on File, Nabriva Therapeutics). Simulations for the final lefamulin PBPK model during validation experiments revealed mean AUC_{0-∞} and C_{max} values within 1.07-fold that of observed values in healthy volunteers.
- Ivacaftor:** A previously published PBPK model for ivacaftor⁴ was used as a basis for development of a minimal PBPK model herein. Once the ivacaftor model was developed, it was validated by simulating concentration-time profiles for repeat oral doses (5 to 14 days) of 50 mg QD and 150 mg BID, comparing exposures with published observed data.^{7,8,9}

PBPK Model Verification For Drug-Drug Interaction Studies

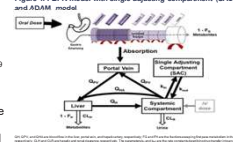
- Ivacaftor:** To characterize the contribution of CYP3A4 to the overall clearance for ivacaftor, simulations were run for combinations with itraconazole, fluconazole and ketoconazole (Simcyp Version 20 Standard Compound Library) and the ivacaftor exposures were compared with observed clinical data.^{7,8,9}
- Lefamulin:** The previously developed PBPK model for lefamulin was similarly validated in its original development. Namely, simulations were made and compared with observed clinical drug-drug interaction studies with ketoconazole, rifampin, and midazolam. Of note, the final enzyme competitive inhibition constant (K_i) varied between IV and oral formulations of lefamulin. (Table 1, Data on File, Nabriva Therapeutics)

Evaluation of Lefamulin and Ivacaftor Drug-Drug Interaction Potential

- Cystic Fibrosis Population:** For all drug-drug interaction simulations, patient demographic model inputs were derived from the literature and included a proportion of females (48.2%), average albumin concentration (41.9 g/L), a weighted distribution for age, and corrective coefficients for height and weight as a function of gender relative to the North European Caucasian population (preidentified in Simcyp Simulator).^{5,6}
- Given the availability of clinical pharmacokinetic data for ivacaftor as a single dose to cystic fibrosis patients, validations studies were conducted to evaluate predictability of the PBPK model in the cystic fibrosis population.⁸

- Simulations of 10 trials of 10 patients with CF (48.2% female) aged 18 to 35 years were generated with the following dosing regimens:

Figure 1. PBPK model with single adjusting compartment (SAC) and ADAM model



Experiment A: Simulation of a single Oral Ivacaftor Dose Administered within a Multi-dose Regimen of IV or Oral Lefamulin in Patients with CF	
Ivacaftor (single dose)	Lefamulin (steady state)
150 mg oral	150 mg IV
Combination therapy	150 mg oral on Day 7
	IV: 350 mg BID IV given as 1 hr infusion
	PO: 600 mg BID orally for 15 days in fed state

Experiment B: Simulation of Multiple Oral Doses of Ivacaftor Administered with a Single IV or PO Dose of Lefamulin in Patients with CF	
Ivacaftor dose (steady state)	Lefamulin (single dose)
150 mg BID x 8 days	150 mg IV
Combination therapy	150 mg BID x 8 days
	PO: 600 mg BID orally in fed state on day 7

Experiment C: Simulation of Multiple Oral Doses of Ivacaftor and Ivacaftor SD Administered with Multiple IV or Oral Lefamulin Doses in Patients with CF	
Ivacaftor dose (steady state)	Lefamulin (steady state)
150 mg BID orally x 8 days	150 mg IV given as 1 hr infusion on day 7
Combination therapy	150 mg BID x 8 days
	IV: 350 mg IV given BID as 1 hr infusion
	PO: 600 mg BID orally in fed state x 8 days

Statistical Analysis

- Statistical comparisons were performed using least square geometric mean ratios (GMR) for ivacaftor pharmacokinetic profiles with and without lefamulin. Specific comparisons were evaluated for maximum plasma concentration (C_{max}) and area under the plasma concentration-time curve (AUC). Classification of the strength of inhibition was as follows: changes in GMR by <1.25-fold was considered insignificant, 1.25-2.0-fold was considered weak inhibition, 2.0 – 5.0-fold was considered moderate inhibition, and >5.0-fold was considered strong inhibition.

RESULTS

Table 1. Input Parameters for ivacaftor and lefamulin PBPK Models

PARAMETER	Ivacaftor	Lefamulin
Physicochemical and Binding Parameters		
Molecular Weight (g/mol)	392.49	507.74
log P	5.98	2.8
Compound type	Diprotic acid	Monoprotic base
pKa	9.4, 11.6	9.41
ES ₁	0.55	0.71
f _u	0.001	0.121
Main binding protein – Albumin		
Absorption model – ADAM model		
f _{u, gut}	0.001	1
Caco-2 P _{app} (x10 ⁻⁶ cm/s)	11.9	-
P _{app} (obs) (x10 ⁻⁶ cm/s)	2.48	-
MechPeff model P _{app, eff} (x10 ⁻⁶ cm/s)	-	21400
Regional Permeability (x10⁻⁶ cm/s)		
Diadenum	-	0.994
Jejunum I/II	-	4/1.45
Ileum I/II/IV	-	0.62/0.62/0.61/0.587
Colon	-	0.01
Formulation type		
Solubility (mg/mL)	0.003	500 (pH 7.4)
Distribution model – minimal PBPK model (Ivacaftor = full PBPK model)		
V _{ss} (L/kg)	1.74	1.79
SAC Q (L/h)	3.752	-
V _z (L/kg)	0.87	-
K _{sc} scalar	-	1
Elimination parameters – Enzyme kinetics		
Intrinsic clearance	97%	-
CYP3A4 K _m (μM)	4	-
CYP3A4 V _{max} (pmol/min/mg protein)	1519.2	-
Biliary CL _{int} (μL/min/10 ⁶ cells)	30	-
CL _{int} (L/h)	-	1.6
CYP2A4 CL _{int} (μL/min/mg)	-	0.153
Additional HLM CL _{int} (μL/min/mg)	-	39
f _u	0.105	1
Transport parameters		
P _{app} (pmol/min)	-	403.8
P _{app} K _m (μM)	-	10
Interaction parameters		
CYP3A4 K _i (μM) (PO administration)	-	0.2
I _{max}	-	0.98
CYP3A4 K _i (μM) (IV administration)	-	0.86
f _u	-	0.98
CYP2C8 K _i (μM)	-	41
f _u	-	1

PBPK Model Development and Validation

- Table 1 depicts the final input parameters for both ivacaftor and lefamulin in their respective PBPK models.
- In validation studies, simulations for the final ivacaftor PBPK model revealed mean AUC_{0-∞} and C_{max} values within 1.28-fold that of the observed values in healthy volunteers.
- Simulations of cystic fibrosis patients given single doses of ivacaftor resulted in AUC_{0-∞} and C_{max} values within 1.4-fold of the observed values, with simulated exposures being greater than observed.

PBPK Model Verification For Drug-Drug Interaction Studies

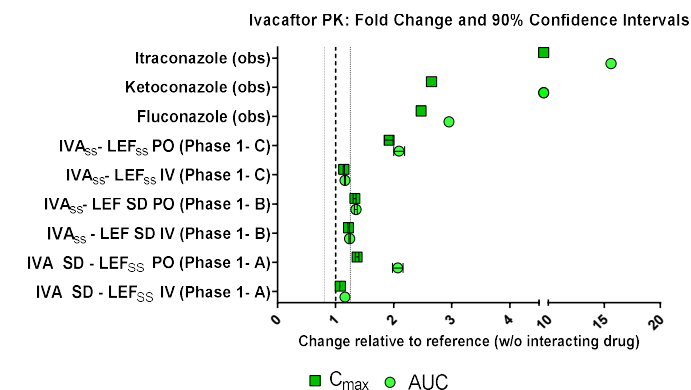
- Simulated geometric mean ratios for AUC and C_{max} were within 1.25, 1.25, and 1.5-fold of the observed values for itraconazole, fluconazole, and ketoconazole, respectively.

Evaluation of Lefamulin and Ivacaftor Drug-Drug Interaction Potential

- The results of the simulated drug-drug interaction between lefamulin and the sensitive CYP3A substrate, ivacaftor, are shown in Figure 1 as compared with observed clinical data for the moderate and strong CYP3A inhibitors fluconazole, ketoconazole, and itraconazole.
- Simulations combining oral ivacaftor with IV lefamulin resulted in non-significant impacts on ivacaftor pharmacokinetics with GMRs for AUC of ≤1.24 and C_{max} of ≤1.22.

- Simulated effects of oral lefamulin on the pharmacokinetics of ivacaftor were classified as mild to moderate inhibition with GMRs of ≤2.09 for AUC and ≤1.92 for C_{max}, with greater inhibition observed when lefamulin was given as multiple doses (i.e. Experiment A and C) vs. as a single dose (Experiment B).

Figure 1: Impact of Lefamulin (LEF) and Comparators on the Pharmacokinetics (PK) of Ivacaftor (IVA)



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CONCLUSIONS

- Simulations of PBPK models for lefamulin and ivacaftor were in reasonable agreement with observed clinical pharmacokinetic data with a ≤1.4-fold difference in AUC and C_{max} values.
- For drug-drug interaction simulation studies of ivacaftor combined with IV lefamulin in cystic fibrosis patients, lefamulin's effects on ivacaftor were non-significant.
- Simulated changes in ivacaftor AUC when administered with oral lefamulin in cystic fibrosis patients was classified as weak to moderate.
- Lefamulin simulations resulted in a lesser impact on ivacaftor AUC as compared to combinations with fluconazole (a known moderate CYP3A inhibitor).
- Taken together, these results suggest a low liability for a CYP3A-mediated drug-drug interaction between lefamulin and ivacaftor in cystic fibrosis patients.

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