

# Population pharmacokinetic model for Ritonavir in HIV-infected patients treated with Lopinavir/Ritonavir (Kaletra®)



VALIDATION SET

65 (37/28)

156

108

48

48/60/22

INDEX SET

198 (110/88)

954

539

415

288/80/136

Demographic and clinical data of patients included in this study

A Martín Suárez<sup>1</sup>, D Santos Buelga<sup>1</sup>, E López<sup>2</sup>, S Cabrera<sup>3</sup>, R López<sup>4</sup>, E Ribera<sup>5</sup>, A Domínguez-Gil<sup>2</sup>, MJ García<sup>1</sup> <sup>1</sup>Department of Pharmacy and Pharmaceutical Technology, <sup>2</sup>Pharmacy Service. University Hospital; University of Salamanca, Salamanca, Spain. <sup>3</sup>Pharmacy Institute. Universidad Austral de Chile, Valdivia, Chile.

<sup>4</sup>Service of Biochemistry (section of pharmacology), <sup>5</sup>Clinic Hospital of Barcelona and Service of Infectious Diseases Vall'd Hebron, Barcelona, Spain.

#### INTRODUCTION

Kaletra® is a fixed dose coformulation of two HIV-1 protease inhibitors (lopinavir [LPV] 400mg/ritonavir [RTV] 100mg). The pharmacoenhancing effect of RTV on LPV resulted in a highly potent, clinically effective antiretroviral drug with a high genetic barrier to viral resistance.

# **OBJECTIVE**

Having the population model previously obtained for LPV with the same data-set (D Santos Buelga. PAGE 2009), the aim of this study was to develop and validate a population pharmacokinetic (PK) model for RTV used as a booster in HIV-infected patients treated with Kaletra®

**METHODS** PATIENTS: HIV-infected subjects, treated with Kaletra® twice daily

ANALITICAL ASSAY: HPLC with UV detection.

PHARMACOKINETIC ANALYSIS:

<u>PK Model</u>: one-compartment model with first-order absorption and elimination including the absorption lag-time (ALAG)

PK parameters estimated: Clearance(CL/F), distribution volume (V/F), Ka, ALAG

✓ Error model: Proportional (interindividual) and additive (residual)

Software: NONMEM V.I (FOCE, Interaction); Xpose (GAM)

<u>Covariates analysed</u>: age, sex, height, total body weight (TBW), body mass index (BMI), LPV trough concentration  $(CL_{LPV})$ , LPV clearance  $(CL_{LPV})$ , total bilirrubin, hepatitis C virus co-infection (HCV), and concomitant saquinavir (SQV), tenofovir (TFV) and atazanvir (ATV)

Nº SQV/TFV/ATV concentrations HCV (Y/N) 455/499 91/65 40.0 ± 8.18 42.2 ± 9.30 Age (years) Weight (kg) 69.2 ± 14.6 68.6 ± 12.4 BMI (kg/m<sup>2</sup>)  $23.9 \pm 4.21$  $23.4 \pm 3.40$ LPV concentration (mg/l) 7.92 ± 3.45 7.36± 3.61 RTV concentration (mg/l)  $0.86 \pm 0.57$  $0.79 \pm 0.53$ LPV clearance  $(\mathrm{CL}_{\mathrm{LPV}})$  (l/h)  $4.30 \pm 1.4$  $5.33 \pm 1.58$ 

Nº patients (male/female)

Nº RTV and LPV concentrations

Trough steady-state

- From full PK profiles

EXTERNAL VALIDATION: Comparison of model-predicted and observed concentrations obtained in the validation data. Mean prediction error (MPE) and standardised mean prediction errors (SMPE) were used.



## CONCLUSIONS

Inclusion of the absorption lag-time leads to the best fitting for the basic model.

•The inclusion of CL<sub>LPV</sub> in the model elicited a decrease in the objective function value (-658.66 to -801.18) and a reduction in the interindividual variability of RTV clearance (45% to 29%)

•RTV clearance was also significantly influenced by SQV concomitant treatment as a categorical covariate (0/1). This effect has been quantified (RTV clearance decreased by 25%).

•No covariates were found to explain the high variability of other PK parameters estimated.

•The validation results obtained confirm the adequacy of the proposed model.

## REFERENCES

• BS Kappelhoff et als. Br J Clin Pharmacol 59(2):174-182,2004. J Moltó et als. Clin Pharmacokinet 46(1):85-92,2007.

•E Ribera et als. Antimicrob Agents Chemother 48:4256-62,2004. •N von Henting. Drugs Today 43(4):221-47,2007.