



Characterization of rifampicin pharmacokinetics in Tanzanian and South African populations



Yuan J. Pétermann¹, Bibie Said^{2,3}, Margaretha L. Sariko⁴, Yann Thoma⁵, Stellah G. Mpagama², Chantal Csajka^{1,6,7}, Monia Guidi^{1,6,8}

1 Centre for Research and Innovation in Clinical Pharmaceutical Sciences, Lausanne University Hospital and University of Lausanne, Lausanne, Switzerland. 2 Kibong'oto Infectious Diseases Hospital, Sanya Juu, United Republic of Tanzania. 3 The Nelson Mandela African Institution of Science and Technology, Arusha, United Republic of Tanzania. 4 Kilimanjaro Clinical Research Institute Kilimanjaro, Moshi, United Republic of Tanzania. 5 School of Engineering and Management Vaud, HES-SO University of Applied Sciences and Arts Western Switzerland, 1401 Yverdon-les-Bains, Switzerland.. 6 Institute of Pharmaceutical Sciences of Western Switzerland, University of Geneva, University of Lausanne, Geneva & Lausanne, Switzerland. 7 School of Pharmaceutical Sciences, University of Geneva, University of Lausanne, Geneva & Lausanne, Switzerland. 8 Service of Clinical Pharmacology, Lausanne University Hospital and University of Lausanne, Lausanne, Switzerland.

Context

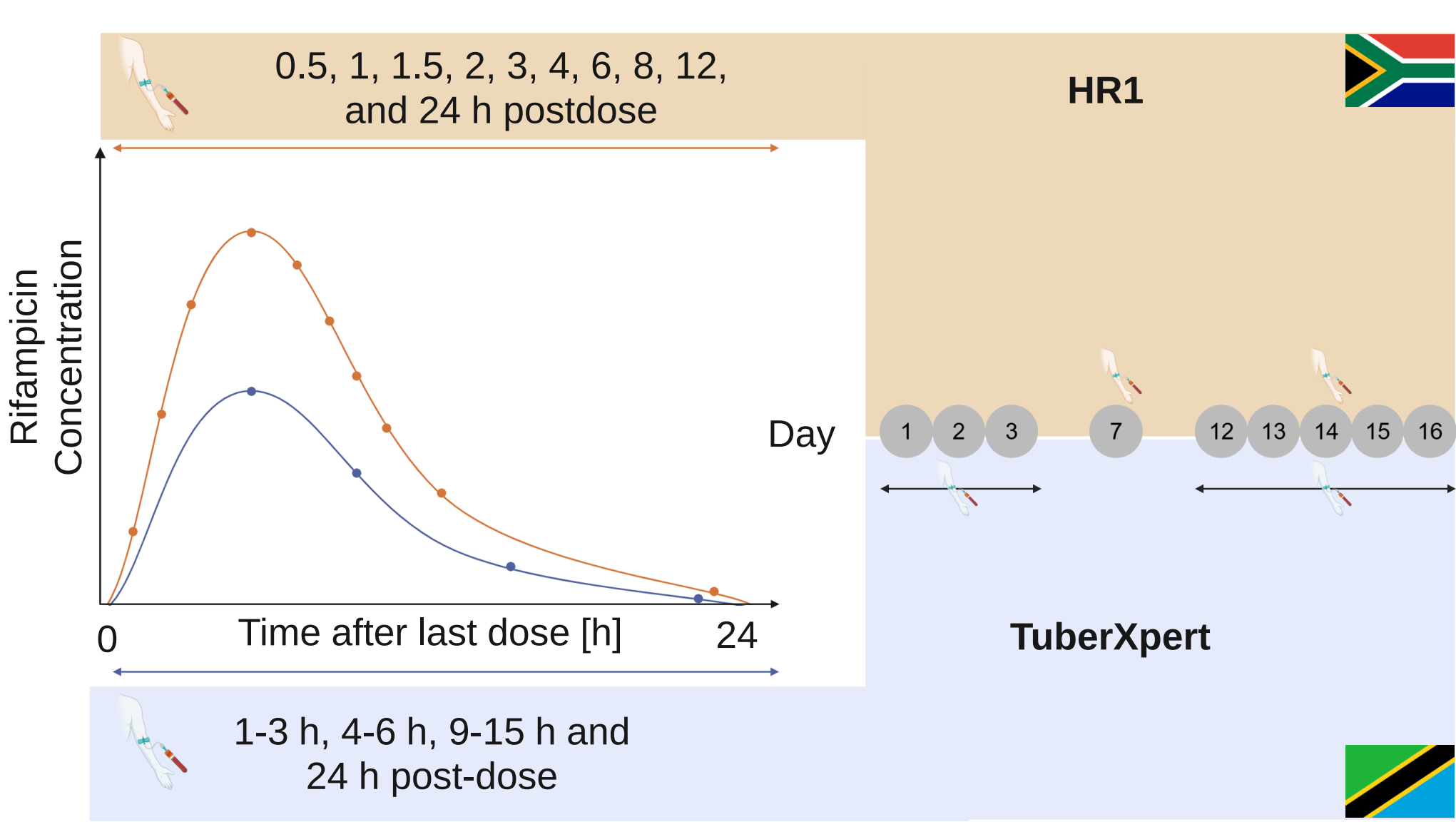
- Rifampicin is a cornerstone first-line anti-tuberculosis (TB) drug, but its **non-linear, highly variable and autoinductive pharmacokinetics (PK)** can result in subtherapeutic exposures.
- Genetic polymorphisms of the **organic anion-transporting polypeptide 1B1 (OATP1B1)** - which differ across ethnic groups - can substantially alter rifampicin PK.
- These complexities call for **optimized, individualized dosing strategies to ensure efficacy**.

Objectives

- Describe and characterize autoinductive and saturable PK** of rifampicin using the population approach.
- Identify and quantify individual and population variability** susceptible to alter rifampicin PK.

Methods

- A stepwise analysis comparing various compartmental models with linear and non-linear absorption and elimination were conducted with MonolixSuite™ (v2024R1). Inter-individual variability (IIV), and inter-occasion variability (IOV) on PK parameters were sequentially tested. Likewise, study effect was assessed as a covariate effect.
- Different residual error models were evaluated to characterize the unexplained variability.
- Visual Predictive Checks (VPC, n=1000) and bootstrap evaluations (n=2000) were performed to assess performance of the models.



- Data were collected through prospective rifampicin plasma concentrations sampled from TB patients at Kibong'oto Infectious Diseases Hospital, Tanzania, within the frame of the TuberXpert project¹ and completed with retrospective data from the South African HIGHRIF1 (HR1) study.

Results

Table 1. Characteristics of the TuberXpert (Tanzania, n_{patients}=42, n_{samples} = 278) and HR1 (South Africa, n_{patients}=92, n_{samples} = 1945) population

Population Characteristics	Median (range) or n (%)		n (%)			n (%)		
	TuberXpert	HR1	TuberXpert		HR1	TuberXpert		HR1
Age [years]	44 (25-65)	NA	Severely Underweight	14 (26)	NA	10 [mg/kg]	38 (90)	8 (9)
Female	11 (74)	NA	Moderately Underweight	3 (5)	NA	20 [mg/kg]	0 (0)	15 (16)
Weight [kg]	49 (31-65)	NA	Mildly Underweight	9 (17)	NA	25 [mg/kg]	1 (2)	15 (16)
Body Mass Index [kg/m²]	17.35 (12.10-24.70)	NA	No Malnutrition	16 (30)	NA	30 [mg/kg]	3 (7)	15 (16)
FFM [kg]	42.51 (19.94-50.44)	44.09 (28.50-57.86)	Overweight	9 (17)	NA	35 [mg/kg]	0 (0)	14 (15)
HIV positive patients	20 (48)	NA	Obese	3 (5)	NA	40 [mg/kg]	0 (0)	15 (16)
DM patients	6 (14)	NA				50 [mg/kg]	0 (0)	10 (11)

HIV, Human Immunodeficiency Virus; **DM**, diabetes mellitus; **FFM**, fat-free mass according to ²; malnutrition status according to ³

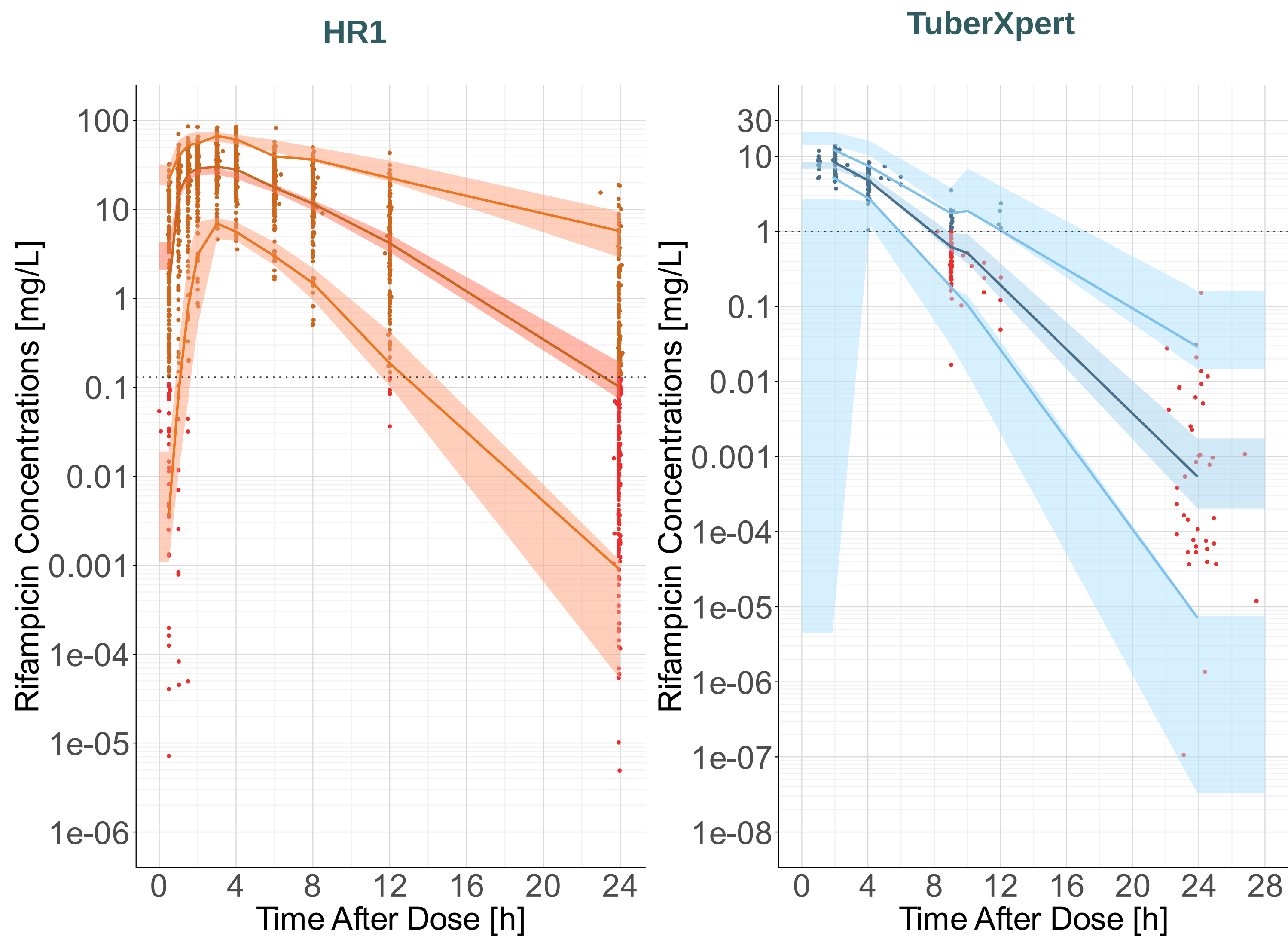


Figure 2: HR1 (Left) and TuberXpert (Right) VPC of the current model in log-transformed y axis. The points indicate the observed concentrations and the red dots their counterparts below the quantification limits (dotted lines, 0.13 mg/L and 1 mg/L for respectively TuberXpert and HR1). The solid lines represent the corresponding observed median and percentiles 5th and 95th. Shaded area depicts the prediction of the 90% confidence interval of the median and the 5th and 95th percentiles.

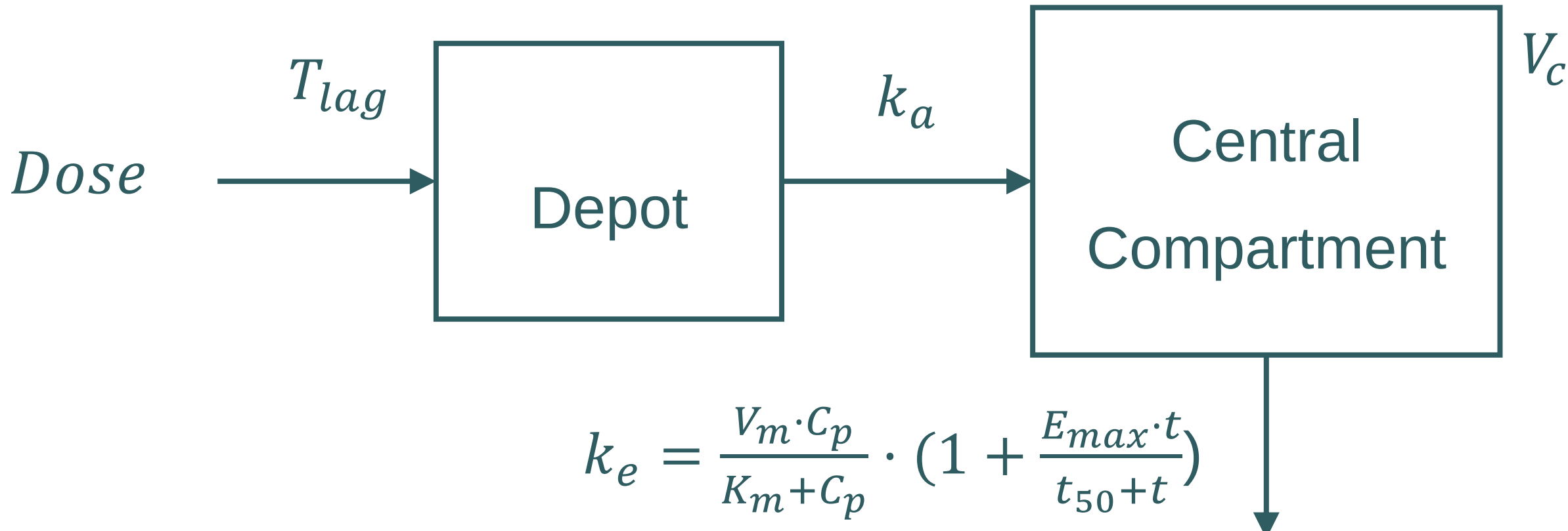


Figure 1. Rifampicin structural popPK model. The saturable component of the elimination rate (**ke**) was characterized by a Michaelis-Menten kinetics, with a maximum elimination rate **V_m** and a concentration **K_m** corresponding to half the maximal elimination rate. Autoinduction of **ke** is described by an **E_{max}** function depending on time (**t**), with half-life **t₅₀**, at which the autoinduction maximum effect (**E_{max}**) is reached. **K_a**, absorption rate constant; **V**, volume of distribution; **T_{lag}**, lag time.

Table 2. Final population PK parameters estimates and bootstrap results

Parameter	Final model		Bootstrap (n = 2000)	
	Estimate (RSE [%])	Median (CI _{95%})	Estimate (RSE [%])	Median (CI _{95%})
T_{lag} [h]	0.32 (12)	0.31 (0.24-0.38)		
ω_{Tlag} [CV%]	75 (22)	95 (47-161)		
	96 (17)	95 (45-112)		
ka [h⁻¹]	2.64 (6)	2.58 (2.33-2.89)		
ω_{ka} [CV%]	38 (12)	36 (27-45)		
	41 (3)	40 (38-42)		
V [L]	41 (3)	40 (38-42)		
ω_{Vc} [CV%]	18 (12)	19 (15-24)		
	124 (13)	125 (95-161)		
V_{m-HR1} [mg/h]	420 (10)	459 (354-601)		
ω_{Vm} [CV%]	38 (24)	33 (18-50)		
	27.5 (12)	30.1 (24.3-38.5)		
K_m [mg/L]	46 (23)	46 (25-73)		
ω_{Km} [CV%]	3.31 (15)	3.94 (2.87-5.27)		
	0.46 (16)	0.53 (0.23-0.88)		
E_{max-HR1}	24 (30)	21 (10-36)		
ω_{Emax} [CV%]	157 (20)	196 (134-287)		
	65 (23)	57 (30-92)		
t₅₀ [h]	83 (4)	105 (67-161)		
ω_{t50} [CV%]	25 (10)	24 (20-30)		
	23 (5)	24 (21-26)		
Correlation_{Vm-Km} [%]				
σ_{prop-TuberXpert} [%]				
σ_{prop-HR1} [%]				

ω, inter-individual variability; γ, inter-occasion variability; CI_{95%}, 95% confidence interval; σ_{prop}, proportional error; RSE, Relative standard error.

- Substantial study effect was observed**, underlining marked population differences in the elimination process of rifampicin.
- HR1 and TuberXpert populations increase their elimination by 3.76-fold and 1.38-fold**, respectively, once **autoinduction steady-state is reached**, after roughly a month of continuous treatment of rifampicin.

Conclusion

- We successfully characterized **rifampicin's non-linear, saturable kinetics and time-dependent autoinduction**.
- Significant inter-study differences**—likely due to ethnic variability in OATP1B1 polymorphism—were observed, possibly because Tanzanian patients are more prone to carry genotypes linked to reduced rifampicin exposure and thus faster eliminations⁴.
- Substantial interindividual variability in rifampicin PK** was observed, emphasizing the need to individualized and monitoring solutions to improve TB patients' care.
- Further investigations relying on these findings will contribute to the **development of a model-informed precision dosing software tailored to the Tanzanian routine clinical care**.