

A population pharmacokinetic model is beneficial in quantifying hair concentrations of ritonavir-boosted atazanavir: A study of HIV-infected Zimbabwean adolescents.

Bernard Ngara (✉ bernardngara4@gmail.com)

University of Zimbabwe College of Health Sciences <https://orcid.org/0000-0002-8355-5954>

Simbarashe Zvada

Stellenbosch University

Tariro Dianah Chawana

University of Zimbabwe College of Health Sciences

Babill Stray-Pedersen

Oslo University Hospital

Charles Fungai Brian Nhachi

University of Zimbabwe College of Health Sciences

Simbarashe Rusakaniko

University of Zimbabwe College of Health Sciences

Research article

Keywords: Pharmacokinetic modelling, HIV/AIDS, Adolescents, Adherence, Hair, NONMEM

Posted Date: March 4th, 2020

DOI: <https://doi.org/10.21203/rs.3.rs-15958/v1>

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Version of Record: A version of this preprint was published on August 3rd, 2020. See the published version at <https://doi.org/10.1186/s40360-020-00437-y>.

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3 **boosted atazanavir: A study of HIV-infected Zimbabwean adolescents.**

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6 ¹*Conflict of interest*

7 ²*Funding*

8 ³*Correspondence*

¹ The authors do not have a commercial or other association that might pose a conflict of interest

² The corresponding author obtained doctorate scholarship from Letten Foundation, Norway. This research was also commissioned by the National Institute for Health Research, using Official Development Assistance funding 16/136/33.

³ Bernard Ngara, University of Zimbabwe College of Health Sciences, Department of Community Medicine, Mazowe Street, Parirenyatwa Complex, P.O Box A178 Avondale, Harare, Zimbabwe, email: bernardngara4@gmail.com.

1 **Authors:**

Name	Title	Affiliation/Address	Authorship Position
Bernard Ngara bernardngara4@gmail.com +263 776 971 400 https://orcid.org/0000-0002-8355-5954 .	Mr.	University of Zimbabwe College of Health Sciences Department of Community Medicine Mazowe Street, Parirenyatwa Complex P.O Box A178 Avondale Harare, Zimbabwe	1
Simbarashe Zvada	Dr.	Stellenbosch University, Department of Clinical Pharmacology, Private Bag X1, Matieland, 7602, Stellenbosch, South Africa	2
Tariro Dianah Chawana	Dr.	University of Zimbabwe College of Health Sciences Department of Clinical Pharmacology Mazowe Street, Parirenyatwa Complex P.O Box A178 Avondale Harare, Zimbabwe	3
Babill Stray-Pedersen	Prof.	Oslo University Hospital Institute of Clinical Medicine Women's Clinic 0027 OSLO Norway	4

Charles Fungai Brian Nhachi	Prof.	University of Zimbabwe College of Health Sciences Department of Clinical Pharmacology Mazowe Street, Parirenyatwa Complex P.O Box A178 Avondale Harare, Zimbabwe	5
Simbarashe Rusakaniko	Prof.	University of Zimbabwe College of Health Sciences Department of Community Medicine Mazowe Street, Parirenyatwa Complex P.O Box A178 Avondale Harare, Zimbabwe	6

1

1 **Abstract**

2 **Background**

3 Adolescents experience higher levels of non-adherence to HIV treatment. Drug concentration in hair promises to be
4 reliable for assessing exposure to antiretroviral (ARV) drugs. Pharmacokinetic modelling can explore utility of drug
5 in hair. We aimed at developing and validating a pharmacokinetic model based on atazanavir/ritonavir (ATV/r) in
6 hair and identify factors associated with variabilities in hair accumulation.

7 **Methods**

8 We based the study on secondary data analysis whereby data from a previous study on Zimbabwean adolescents
9 which collected hair samples at enrolment and three months follow-up was used in model development. We
10 performed model development in NONMEM (version 7.3) ADVAN 13.

11 **Results**

12 There is 16% / 18% of the respective ATV/r in hair as a ratio of steady-state trough plasma concentrations. At
13 follow-up, we estimated an increase of 30% /42% of respective ATV/r in hair. We associated a unit increase in
14 adherence score with 2% increase in hair concentration both ATV/r. Thinner participants had 54% higher while
15 overweight had 21% lower atazanavir in hair compared to normal weight participants. Adolescents receiving care
16 from fellow siblings had atazanavir in hair at least 54% less compared to other forms of care.

17 **Conclusion**

18 The determinants of increased ATV/r concentrations in hair found in our analysis are monitoring at follow up event,
19 body mass index, and caregiver status. Measuring drug concentration in hair is feasibly accomplished and could be
20 more accurate for monitoring ARV drugs exposure.

21

22 **Keywords**

23 Pharmacokinetic modelling, HIV/AIDS, Adolescents, Adherence, Hair, NONMEM

24

25

1 **Background**

2 About 36.9 million people were living with Human Immunodeficiency Virus (HIV) worldwide in 2017. Of these,
3 approximately 3 million were children and adolescents under 20 years of age [1]. Zimbabwe has a prevalence of
4 13.3%, with 1.3 million people living with HIV including 77,000 children and adolescents [2]. Poor adherence to
5 treatment leads to sub-optimal drug exposure limiting treatment efficacy [3,4]. It has been estimated that between 20
6 to 50% of adolescents experience adherence-related antiretroviral (ARV) drug treatment failure [5–11].

7 It is desirable to have a routine assessment of adherence and exposure to ARV drugs available for use by healthcare
8 providers. The methods used for assessing adherence and exposure to ARV drugs include self-reported missed
9 doses, monitoring pharmacy refills and conducting pill counts, use of electronic monitoring devices, measuring
10 ARV concentration in plasma or hair [12–14]. Quantifying ARV drugs in hair provides information of both steady-
11 state pharmacokinetics and long term adherence and has shown to predict well the relationship between drug
12 exposure and treatment outcomes when compared to other approaches [15–21].

13 Some suggest that hair uptake most external substances or their metabolites from the systemic circulation through
14 the hair bulb blood supply by passive diffusion from blood into growing hair cells at the base of the follicle and then
15 bound in the hair shaft [22–25]. Once the drug accumulates into the growing hair, we can detect it long after
16 elimination from the systematic circulation, unlike in conventional biological samples such as blood and urine [26–
17 30]. The scalp hair fibre grows at an average rate of 0.5 to 1.5 centimetres per month [31]. Thus the amount of drug
18 in hair is constantly increasing until the next hair cut or when all the drug is removed from the systematic
19 circulation.

20 In Zimbabwe and other resources limited settings, pharmacokinetic (PK) modelling applied focused primarily on
21 systemic exposure to ARV drugs. It based the models used on data generated by quantifying drugs mostly from
22 single time-point plasma samples [32–35]. Hypothetically hair PK parameters can provide additional information
23 about the patient's drug exposure overtime, hence the need to determine and apply hair PK for prediction of drug
24 amount in the hair in relation to exposure in plasma.

25 Using single time-point plasma samples is considered unreliable when assessing drug exposure in populations at risk
26 of non-adherence. Measuring ARV concentrations in hair promises to be more accurate and feasibly accomplished.

1 However, there are very few studies which analysed drug concentration measured in hair using PK approaches. The
2 aim of the work reported in this paper was to develop and validate a population PK model for ATV/r concentrations
3 in hair and explore factors associated with increased or reduced concentrations assuming a direct relationship
4 between ratio in plasma and hair.

5 **Methods**

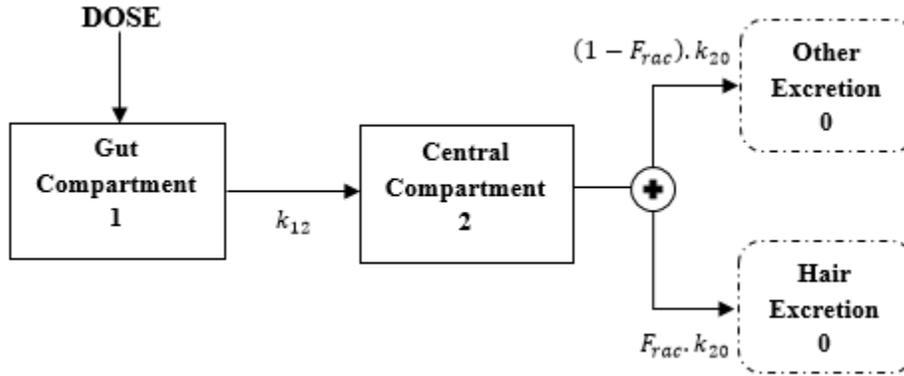
6 **Source of data**

7 The study used secondary data from a previously published study conducted in Zimbabwe [15]. They collected the
8 data between January 2015 and May 2016. It comprised 50 adolescents aged between 10 to 18 years and were on
9 ATV/r (300/100 mg) based 2nd line HIV treatment for at least 6 months. They enrolled these participants at a public
10 health hospital in Harare, Zimbabwe, and randomized to either adherence intervention or standard of care arms.
11 They excluded participants if they were on anti-TB treatment, did not prefer home be followed-ups, had viral load
12 <1 000 copies/ml within the previous 2 months, or were on ATV/r as 1st line treatment. The goal of the primary
13 study was to test the impact of a home-based modified directly administered adherence intervention on virologic
14 outcome. They collected questionnaire data, blood samples and hair samples cut closest to the scalp at baseline and
15 at 90 days follow-up using standard protocol described in the primary study [15]. They did the procedures taken in
16 preparation and analysis of hair samples at the University of California, San Francisco Hair Analytical Laboratory.
17 A standard protocol for preparation and analysis of ATV in hair was developed, validated, peer reviewed and
18 approved by the Division of AIDS Clinical Pharmacology and Quality Assurance program [36].

19

20 **Pharmacokinetic modelling**

21 We developed a population PK model to describe ATV/r concentrations in hair. We fixed parameters describing the
22 PK of atazanavir and ritonavir in plasma based on previously published estimates got from studies conducted in
23 almost similar settings [36,37] with inter-individual (IIV) and inter-occasion (IOV) variabilities estimated
24 sometimes. We did model development using the first-order conditional estimation method with interaction (FOCE-
25 I) in NONMEM (version 7.3) ADVAN 13 [38]. We schematically presented the structural population PK model
26 applied to both atazanavir and ritonavir concentrations using Figure 1.



1
2 **Figure 1**

3 The model describes the concentration of drug in hair at steady-state trough plasma concentrations in the body 24
4 hours after a dose. Given that there were no plasma concentration data, we used a simplified model which estimate
5 ratios of concentrations between hair and plasma. We describe the rate of change of amount of drug between
6 compartments in Figure 1 using the differential equations:

7
$$\frac{dA_1}{dt} = -k_{12} \cdot A_1 \quad (1)$$

8
$$\frac{dA_2}{dt} = k_{12} \cdot A_1 - k_{20} \cdot A_2 \quad (2)$$

9
$$\frac{dA_3}{dt} = F_{rac}(k_{20}) \cdot A_2 \quad (3)$$

10
$$C = \frac{A_3}{V_h} \quad (4)$$

11 Equation 1 describe drug absorption into the plasma circulation i.e. central compartment at a rate (k_{12}) proportional
12 to the dose amount (A_1); Equation 2 describe input from Equation 1 and total elimination of the drug at rate ($k_{20} =$
13 $\frac{CL}{V_c}$, where CL and V_c represents clearance and apparent volume of distribution of the bioavailable drug proportional
14 to the amount in the central compartment (A_2); Equation 3 describe a ratio (F_{rac}) of hair concentration relative to
15 steady-state plasma trough concentration. Equation 4 predicts the drug concentration C in hair using the ratio of
16 amount of drug in the hair (A_3) and apparent volume of distribution (V_h) of the bioavailable drug in the hair.

17 We tested all covariates in Table 1 during covariate analysis. We selected the optimal covariates relationships
18 through clinical and prior assessment of statistical significance testing using the Stepwise Covariate Model building
19 (SCM) method as implemented in Perl-speaks-NONMEM (PsN). We tested relations on the F_{rac} parameter only.

1 **Model evaluation**

2 We used the change in objective function value (ΔOFV) provided from NONMEM model output at 5% level of
3 significance (i.e. $\Delta OFV > 3.83$, Chi-square 1-degree of freedom) in forward selection process and then at 1% level
4 of significance (i.e. $\Delta OFV > 6.64$, Chi-square 1-degree of freedom) in the backward deletion process, to make
5 discriminations between hierarchical models. We performed bootstrap analysis and 90% confidence intervals on the
6 final covariate models by re-sampling 1000 times in PsN as part of model evaluation numerically. We used
7 graphical assessment of the standard goodness-of-fit plots as part of model evaluation graphically [39]. Both
8 proportional and additive or combined error models were tested and discriminated by means of change in objective
9 function value (ΔOFV).

10 **Results**

11 **Study participant details**

12 We used fifty participant data in the analysis. The mean (standard deviation) age in years of the study participants
13 was 15.8 (1.8). Fifty-four percent were female. The majority (89%) of these adolescents were attending secondary
14 school, while others were still primary school. Ten percent of the study participants were under the care of fellow
15 siblings below the age of 19 years while others were under the care of parents or relatives. Eighty-two percent of the
16 adolescents were on tenofovir, lamivudine and ATV/r. The median length of hair samples was 1cm (range 0.5cm to
17 1.5cm). The mean (standard deviation) weight of hair samples was 2.0g (0.15g). The mean (standard deviation) drug
18 concentration for atazanavir / ritonavir was 2.5ng/mg hair (2ng/mg hair)/0.5ng/mg hair (0.4ng/mg hair) respectively.
19 The mean (standard deviation) adherence level of a visual analogue scale of 0 to 100% was 84.1% (18.1%). Further
20 we present details on characteristics of the study population in Table 1.

1 Table 1: Summary statistics describing data variables of the original study.

Variable	Response
Length of hair (cm), median (range)	1 (0.5 – 1.5)
Hair weight (grams), mean (Standard Deviation; Range)	2.0 (0.15; 1.76-2.28)
Samples Below limit of quantification	
Atazanavir	18 (18)
Ritonavir	12 (12)
Drug regimen: recruitment + follow-up, n (%)	
Tenofovir/Lamivudine/Atazanavir-ritonavir	75 (82)
Abacavir/Didanosine /Atazanavir-ritonavir	6 (7)
Zidovudine/Lamivudine/Atazanavir-ritonavir	6 (7)
Abacavir/Lamivudine/Atazanavir-ritonavir	3 (3)
Tenofovir/ Emtricitabine /Atazanavir-ritonavir	2 (2)
Body mass index-for-age, n (%)	
Normal	25 (54)
Overweight	7 (15)
Thinness	14 (30)
Age (years), mean (Standard Deviation; Range)	15.8 (1.8; 11-18)
Gender, n (%)	
Female	27 (54)
Caregiver, n (%)	
Parent	10 (20)
Grandparent	20 (40)
Sibling	5 (10)
Aunt/uncle	15 (30)
Level of education, n (%)	
Secondary school	39 (89)
Primary school	8 (9)
Dropped	1 (2)
WHO disease progression stage, n (%)	
Early	16 (32)
Late	34 (68)
Adherence by visual inspection of analogue scale, mean (Standard Deviation; Range)	84.2 (18.1; 30-100)
Atazanavir concentration (ng/mg), mean (Standard Deviation; Range)	2.5 (2.0; 0.07-8.65)
Ritonavir concentration (ng/mg), mean (Standard Deviation; Range)	0.5 (0.4; 0.01-1.39)

1 **Population pharmacokinetic modelling**

2 **Atazanavir model:** We fixed the parameters describing the steady-state population pharmacokinetics of atazanavir
3 in the plasma to 0.44 litres per hour for k_{12} , 10 litres per hour for CL/F and 63.4 litres for V_c [36,37]. We included
4 body weight as a covariate on CL/F and V_c through allometric scaling, fixing the exponents to 0.75 and 1 for CL/F
5 and V_c respectively [40]. We initially estimated V_h for atazanavir in hair but later on fixed it to 1 because the model
6 estimates were close to 1 and also to stabilise the final model results. We estimated that ATV concentration
7 measured in hair is approximately 16% the amount of atazanavir plasma trough concentration after adjusting for
8 covariate effects. Covariate model results show that participants had ATV concentration 30% less at enrolment than
9 that at follow-up event (p-value < 0.0001). A unit increase in self-reported adherence score increased ATV
10 concentration by 2% (p-value = 0.0004). Thinner participants had 54% higher ATV concentration, while overweight
11 participants had 21% lower ATV concentration compared to participants with normal body mass index-for-age (p-
12 value = 0.0165). Participants receiving care from a parent and uncle or aunt had atazanavir in hair 53% and 12%
13 higher respectively, while those receiving care from fellow siblings had atazanavir in hair 54% lower compared to
14 participants receiving care from grandparents (p-value = 0.0406). We present the detailed results in Table 2.

1 **Table 2.** Final model parameters describing joint fixed plasma and hair pharmacokinetics of atazanavir

Parameter	Population mean (SE as %)	1000 samples bootstrap medians (90% CI)	Variability (SE as %)	1000 samples bootstrap medians (90% CI)
k_{12} (litres hour ⁻¹)	0.44 fixed	0.44 fixed	0.45 fixed	0.44 fixed
CL/F (litres hour ⁻¹)	10 fixed	10 fixed	1.04 (99)	0.97 (0.50 – 1.98)
V_c (litres)	63.4 fixed	63.4 fixed	0.50 fixed	0.50 fixed
F_{rac}	0.16 (16)	0.15 (0.06 to 0.26)		
V_h (litres)	1 fixed	1 fixed		
Occasion (Follow-up) $_F_{rac}$	*	*		
Occasion (Enrolment) $_F_{rac}$	-0.30 (23)	-0.27 (-0.50 to -0.07)		
Adherence score $_F_{rac}$	0.02 (18)	0.015 (0.004 to 0.017)		
Body Mass Index-for-age (Normal) $_F_{rac}$	*	*		
Body Mass Index-for-age (Thin) $_F_{rac}$	0.54 (22)	0.49 (0.06 to 0.74)		
Body Mass Index-for-age (Overweight) $_F_{rac}$	-0.21 (121)	-0.15 (-0.26 to -0.05)		
Guardian (Grandparent) $_F_{rac}$	*	*		
Guardian (Parent) $_F_{rac}$	0.53 (56)	0.55 (0.03 to 2.58)		
Guardian (Uncle/Aunt) $_F_{rac}$	0.12 (177)	0.17 (0.05 to 1.60)		
Guardian (Sibling) $_F_{rac}$	-0.54 (35)	-0.60 (-0.92 to -0.27)		
ε_{ADD}	0.30 (1)	0.29 (0.14 to 0.44)		
ε_{PROP}	0.50 (2)	0.50 (0.39 to 0.61)		
Ω	1	1		

2 k_{12} : Absorption rate constant; CL/F : apparent drug clearance; V_c and V_h : apparent volume of distribution in the central and hair compartments, respectively; F_{rac}
3 amount of drug cleared into the hair as a proportion of the amount of drug in plasma at steady-state troughs; $FACTOR_{F_{rac}}$: effect of covariate on F_{rac} ; ε_{ADD} and
4 ε_{PROP} : additive and proportional error terms, respectively; Ω : residual error; SE: standard error. *: reference group.

1 **Ritonavir model:** We fixed the parameters describing the steady-state population pharmacokinetics of ritonavir in
2 the plasma to 2.31 litres per hour for k_{12} , 12.8 litres per hour for CL/F and 105 litres for V_c [36,37]. We initially
3 estimated V_h for ritonavir in hair but later on fixed it to 1 because the model estimates were close to 1 and also to
4 stabilise the final model results. We estimated that ritonavir concentration measured in hair is approximately 18%
5 the amount of ritonavir plasma trough concentration after adjusting for covariate effects. Covariate model results
6 show that participants had ritonavir fraction 42% less at enrolment than that at follow-up event (p-value = 0.0003).
7 A unit increase in self-reported adherence score increased ritonavir concentrations by 2% (p-value = 0.0245). We
8 present the detailed results in Table 3.

1 **Table 3.** Final model parameters describing joint fixed plasma and hair pharmacokinetics of ritonavir

Parameter	Population mean (SE as %)	1000 samples bootstrap medians (90% CI)	Variability (SE as %)	1000 samples bootstrap medians (90% CI)
k_{12} (litres hour ⁻¹)	2.31 fixed	2.31 fixed	0.45 fixed	0.45 fixed
CL/F (litres hour ⁻¹)	12.8 fixed	12.8 fixed	0.28 (31)	0.28 (0.01 to 0.68)
V_c (litres)	105 fixed	105 fixed	0.50 fixed	0.50 fixed
F_{rac}	0.18 (16)	0.18 (0.14 to 0.21)		
V_h (litres)	1 fixed	1 fixed		
Occasion (Follow-up) $_F_{rac}$	*	*		
Occasion (Enrolment) $_F_{rac}$	-0.42 (22)	-0.39 (-0.56 to -0.21)		
Adherence score $_F_{rac}$	0.02 (47)	0.014 (0.008 to 0.017)		
ε_{ADD}	0.34 (95)	0.36 (0.04 to 0.63)		
ε_{PROP}	0.26 (26)	0.24 (0.13 to 0.31)		
Ω	1	1		

2 k_{12} : Absorption rate constant; CL/F : apparent drug clearance; V_c and V_h : apparent volume of distribution in the central and hair compartments, respectively; F_{rac} :

3 amount of drug cleared into the hair as a proportion of the amount of drug in plasma; $FACTOR_{F_{rac}}$: effect of covariate on F_{rac} ; ε_{ADD} and ε_{PROP} : additive and

4 proportional error terms, respectively; Ω : residual error; SE : standard error. *: reference category.

1 **Model diagnostics.**

2 There was no huge variation between all the estimated final model parameters and those got using 1000 samples
3 bootstrap. All the estimated final model were falling within the 90% confidence intervals. Figure 2 presents the basic
4 goodness-of-fit plots showing the population model predictions versus observations and the residual error plots for
5 both atazanavir and ritonavir final models. The results show low bias, and fairly good precision showing fairly
6 acceptable predictive performance.

7
8 **Discussion**

9 This is a breakthrough study to perform joint pharmacokinetic modelling of plasma and hair drug concentrations,
10 determine the relationship between exposure of the drug in hair and that to plasma. Several studies have used drug
11 concentration as a tool for measuring antiretroviral drug exposure in situations where non-adherence to treatment
12 maybe a challenge. However the choice of the multivariate statistical models involving hair concentration as the
13 outcome variable in these studies lacked the dose component which plays a critical role when optimising the
14 relationship between drug exposure and treatment outcomes [5,15,17–20]. A non-linear mixed effect PK model has
15 an advantage that includes the dose component. The main purpose of the current model is basically to inform future
16 study design that involve measuring drug concentrations in hair. Later in the discussion, we will present some
17 limitations and recommendations that can improve the power of this method.

18
19 The amount of atazanavir concentration determined in hair is approximately 16% of steady-state plasma trough. We
20 estimated an almost similar ratio of 18% for ritonavir as part of model testing. In our conceptual framework, we are
21 interested in finding covariates that affect drug exposure to improve the dosing strategies. We used the SCM in
22 identifying covariates associated with variation in hair drug exposure. The drug that accumulates in hair comes from
23 plasma, therefore one of the major assumption is that the covariates found to have an association with accumulation
24 of drug in hair in our results are a function of an altered plasma PK profile.

25
26 ATV/r concentrations increased on follow-up occasion irrespective of study arm which could result from design
27 biases the original study could not eliminate. By being involved in a study, participants are more aware that they are
28 under investigation, hence they adhere more to treatment, increasing hair drug concentrations in both arms. The

1 primary study randomized participants to study arms without blinding, so this could have introduced the biases.
2 High body mass index (BMI)-for-age decreased atazanavir concentrations in hair, while low body mass index-for-
3 age increased atazanavir accumulation in hair. These findings concur with earlier studies in adults[41,42]. They
4 associated low BMI with high plasma drug concentrations, often resulting in supra-therapeutic drug concentrations,
5 with subsequent drug toxicity, side effects and defaulting treatment. Furthermore, the same literature associates high
6 BMI with low plasma drug concentrations, often resulting in sub-therapeutic drug concentrations and subsequent
7 treatment failure and drug resistance. To the best of our knowledge, our study is the first to show this association in
8 adolescents and using drug concentrations in hair.

9
10 We associated receiving care from siblings with lower drug concentrations in hair. Based on current knowledge, this
11 is the 1st study to prove this association using hair samples. Results from a different previous study showed higher
12 self-reported adherence in children and youth who stayed with their parents and grandparents, than those who stayed
13 with siblings [43]. Siblings of HIV-infected children and youth are often immature themselves and still coming to
14 terms with burdens associated with child-headed families, orphanhood and poverty. The needs of HIV-infected
15 children come in as an extra burden that the siblings may not manage the pressure that comes with the burden,
16 leading to missed doses and hospital visits, and subsequent treatment failure in the HIV-infected adolescents.

17
18 A major limitation of the modelling approach applied in the article is we had a small sample size and that we got a
19 few times-points of drug concentration data from each participant. An additional number of participants coupled
20 with having several or segmental measurement of drug concentration from the hair and additionally measuring drug
21 concentration in plasma will improve the power of the modelling framework that used in this paper. Using prior
22 estimates on the plasma PK model could have led to an underestimation of steady state trough concentration because
23 of unavailability of adherence data, however including prior estimates in the form of both fixed and random effects
24 on the plasma PK model could have reduced the bias. Most of the study participants (82%) were on a uniform drug
25 combination, however the unavailability of data on non-HIV/AIDS linked co-therapies limited investigations of the
26 drug-drug interactions which is also very critical to test during PK analysis.

27
28

1 **Conclusion**

2 We have showed some work which can complement the efforts being taken by other scientists to establish the use of
3 measuring drug concentration in hair at HIV/AIDS points of care. Most important determinants of increased
4 concentrations in hair were monitoring at follow up event, BMI-for-age and caregiver. Measuring ARV
5 concentrations in hair promises to be more accurate and feasibly accomplished. It is crucial to perform follow-up
6 work which involves establishing the relationship between hair drug concentrations and a measure of treatment
7 response such as viral loads. Comparing the predictive accuracy for exposure-response models when exposure of
8 interest is plasma or hair drug concentrations is necessary to perform.

9

10 **Abbreviations**

11 ARV - Antiretroviral

12 ATV/r - Atazanavir/ritonavir

13 NONMEM – Nonlinear Mixed Effect Modelling

14 HIV – Human Immunodeficiency Virus

15 AIDS – Acquired Immunodeficiency Syndrome

16 BMI - **B**ody Mass Index

17 PK – Pharmacokinetic

18 IIV - Inter-individual variability

19 IOV - Inter-occasion variability

20 FOCE-I - First-order conditional estimation method with interaction

21 PsN - Perl-speaks-NONMEM

22 SCM - Stepwise Covariate Model building

23 Δ OFV - Change in objective function value

24 Ng/mg – nanogram per milligram

25 k_{12} - Absorption rate constant

26 CL/F - Apparent drug clearance

27 V_c - Apparent volume of distribution in the central compartments

28 V_h - Apparent volume of distribution in the hair compartments

1 F_{rac} - Amount of drug cleared into the hair as a proportion of the amount of drug in plasma at steady-state troughs;

2 FACTOR_ F_{rac} - effect of covariate on F_{rac}

3 ϵ_{ADD} - Additive error term

4 ϵ_{PROP} - Proportional error term

5 Ω - Residual error

6 SE - Standard error

7 P-value – Probability value

8 CI – Confidence Interval

9

10 **Declarations**

11 **Ethics approval and consent to participate**

12 The study got approval to use secondary data from the institutional and national ethical review committees [Joint
13 Research Ethics Committee (JREC/101/18) and Medical Research Council of Zimbabwe (MRCZ/A/2301)
14 respectively].

15

16 **Consent for publication**

17 N/A

18

19 **Availability of data and material**

20 The primary study did not publish the data to the public, however the data can be available upon request and
21 approval from the principal investigators of the primary study.

22

23 **Competing interests**

24 The authors declare that there is no conflict of interest.

25

26 **Funding**

27 The corresponding author obtained doctorate scholarship from Letten Foundation, Norway. This research is also
28 commissioned by the National Institute for Health Research, using Official Development Assistance (ODA) funding

1 16/136/33. The views expressed in this publication are those of the authors only and the funders did not play any
2 role in the design of the study; collection, analysis, interpretation of data, and in writing the manuscript.

3

4 **Authors' Contributions**

5 B.N, S.P.Z, and S.R model development and validation and statistical interpretation of model results. T.D.C and C.N
6 provided the data. T.D.C, B.S.P and C.N clinical and pharmacological interpretation and review of model results.

7 B.N, S.P.Z and T.D.C wrote the manuscript.

8 **Acknowledgements**

9 Authors acknowledge the entire research team of the original study which include researchers, research nurses, the
10 trained community workers, participants, and a team from the University of California San Francisco (United States
11 of America) which worked on the hair assays for their effort in producing the data which we used for secondary
12 analysis. We also acknowledge the University of Cape Town (South Africa) for providing a computing facility with
13 the NONMEM software.

14

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6 **Figure legends**

7 **Figure 1:** Schematic representation of the structural population PK model used to predict atazanavir and ritonavir
8 concentrations measured in hair.

9 **Figure 2:** Basic goodness-of-fit plots for the final model for atazanavir (panel A) and ritonavir (panel B). *Upper left*
10 *panel:* The observations are plotted versus the population predictions. *Upper right panel:* The observations are
11 plotted against the individual predictions. *Lower left panel:* The individually weighted residuals are plotted versus
12 time after dose. *Lower right panel:* The conditional weighted residuals are shown versus the individual predictions.
13 The open black circles represents observed data. The red line is a locally weighted scatter-plot smoother (LOESS),
14 while the solid line is identity or zero

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Figures

A

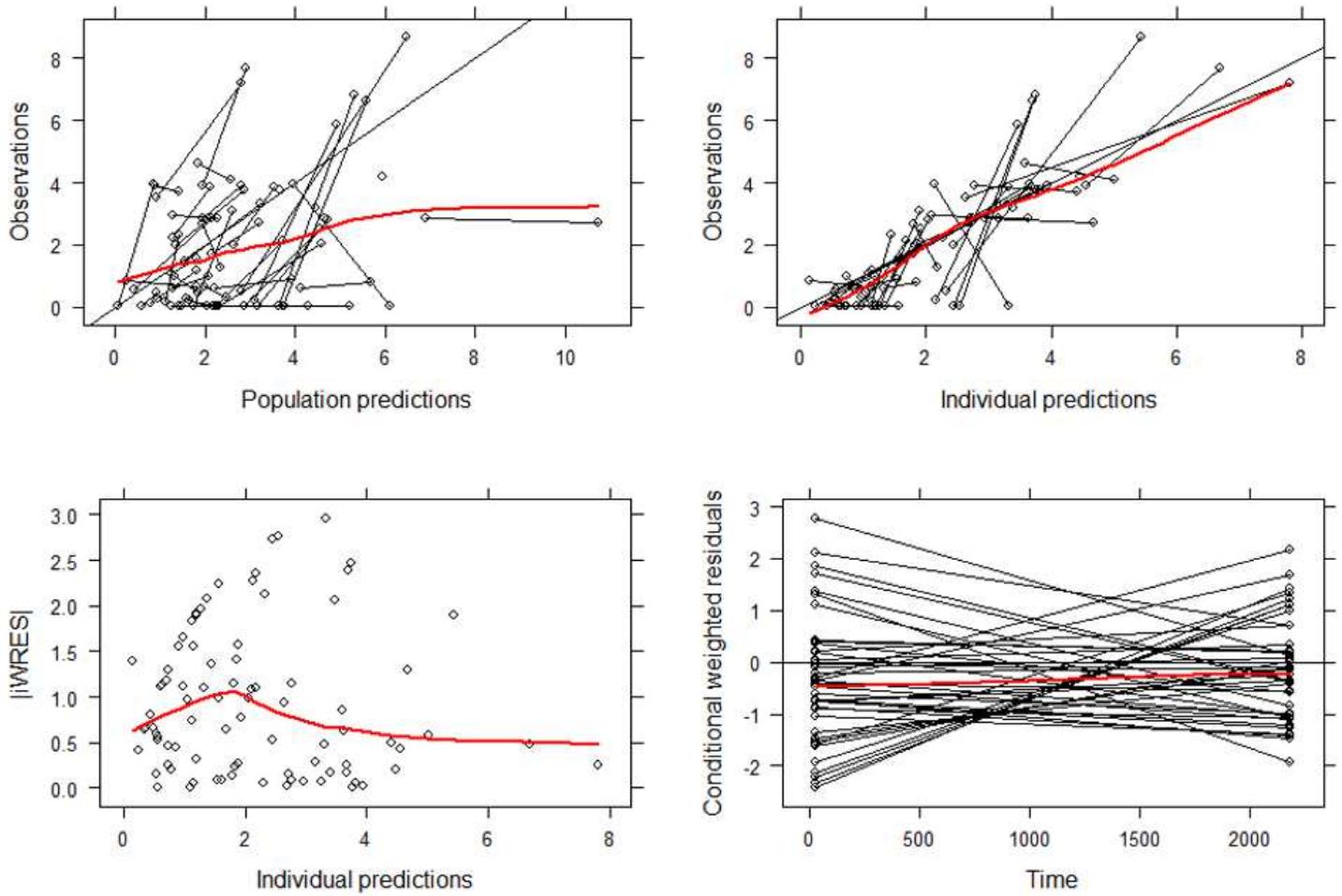
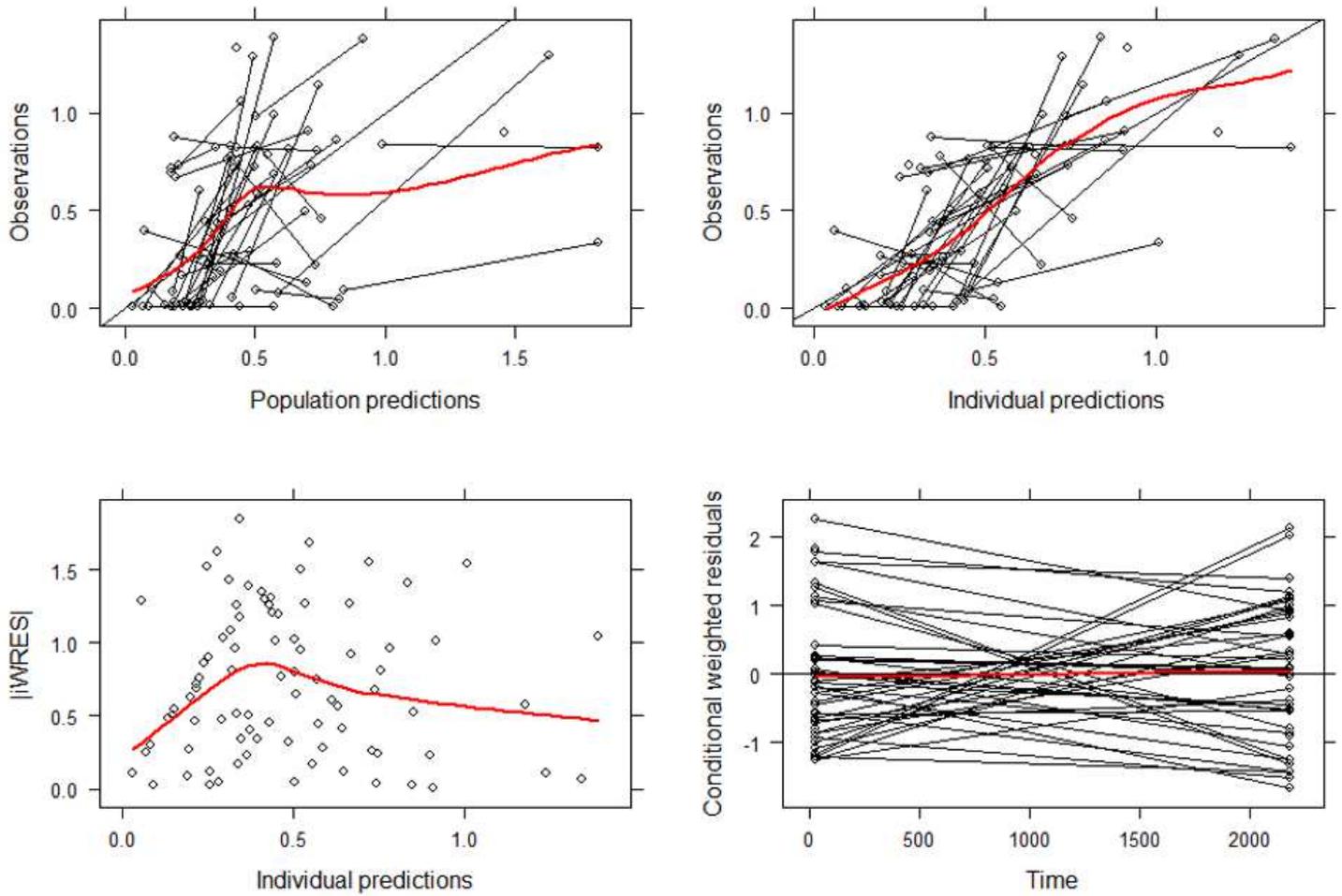


Figure 1

Schematic representation of the structural population PK model used to predict atazanavir and ritonavir concentrations measured in hair.

B**Figure 2**

Basic goodness-of-fit plots for the final model for atazanavir (panel A) and ritonavir (panel B). Upper left panel: The observations are plotted versus the population predictions. Upper right panel: The observations are plotted against the individual predictions. Lower left panel: The individually weighted residuals are plotted versus time after dose. Lower right panel: The conditional weighted residuals are shown versus the individual predictions. The open black circles represents observed data. The red line is a locally weighted scatter-plot smoother (LOESS), while the solid line is identity or zero.