

# Population PK and Viral Dynamic Modeling of S/GSK1349572 in Patients with HIV Infection

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# Objectives

**Develop a population PK-viral dynamic model to characterize the relationship between plasma 572 concentrations and HIV viral decline in HIV-infected patients**



- Explore the underlying mechanism
- Optimize dose selection for future studies
- Address clinical relevant questions  
e.g., Resistance, DDI, ...

# 572 exhibited potent antiviral activity following 10-day monotherapy in INI-naïve HIV patients



\*\* Approved dose for etravirine is 200mg BID

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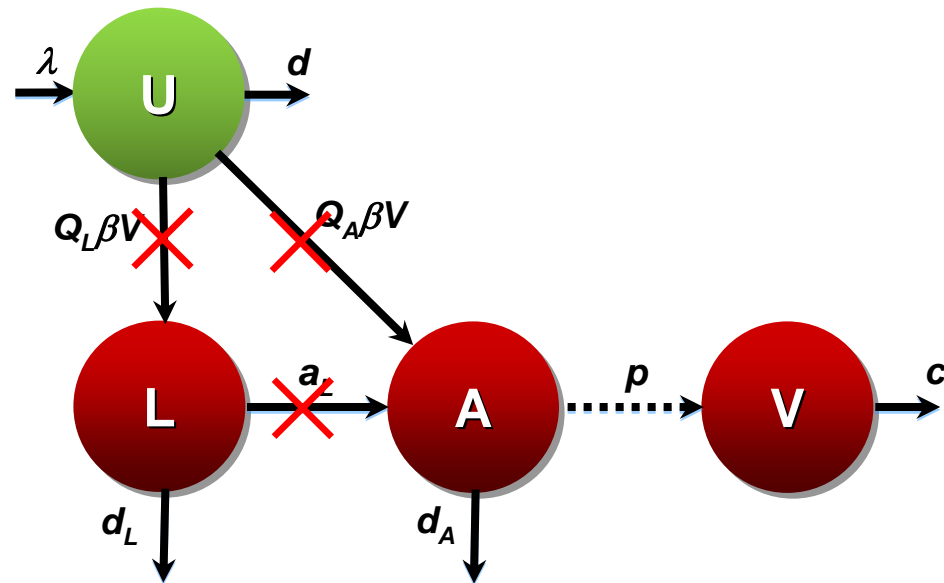
# Proposed Hypotheses for Potent Antiviral Activity of 572

**U: Uninfected CD4+ cells**

**A: Actively infected cells**

**L: Latently infected cells**

**V: Infectious virus**



Funk, et al., JAIDS 2001; 26:397-404.

\*Murray, et al. AIDS 2007, 21:2315-2321.

**Hypothesis 1:** INI acts like traditional anti-HIV drugs with increased potency in suppressing the viral infectivity

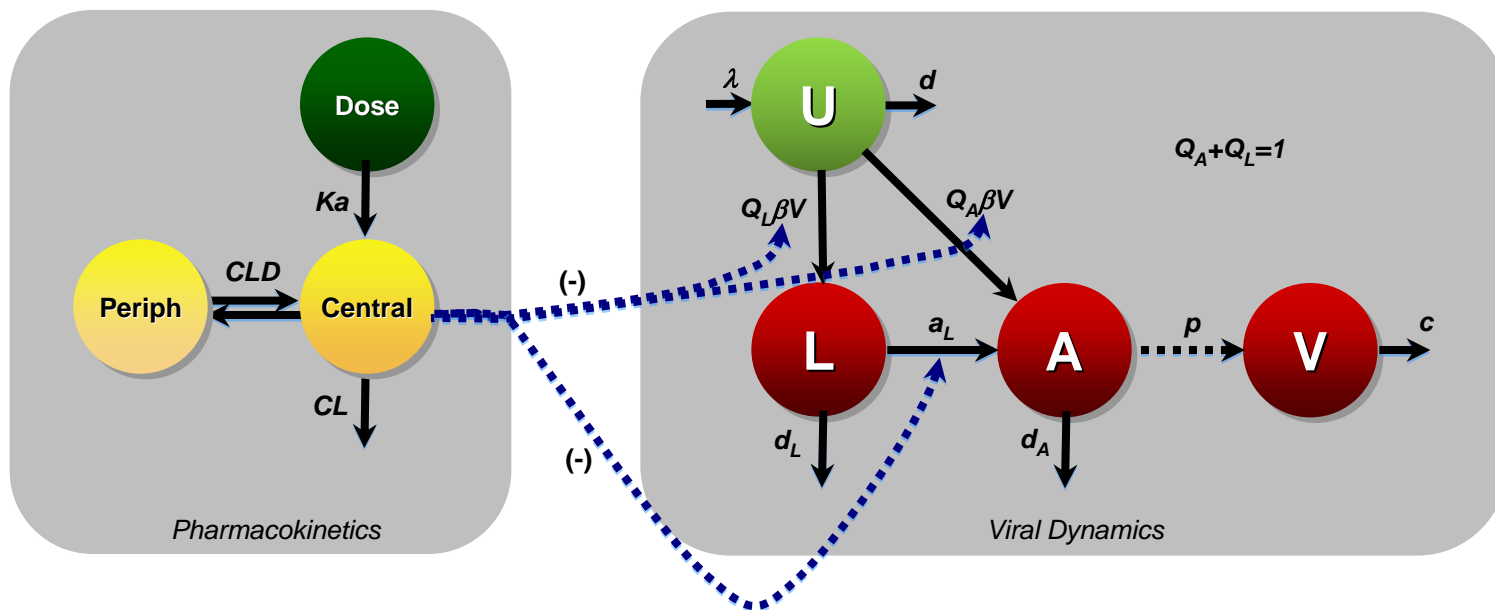
**Hypothesis 2\*:** In addition to suppression of viral infectivity, INI also blocks viral DNA integration in latently infected cells

# Phase 2a Study Design of 572

- Randomized, parallel, double-blind, dose ranging, placebo-controlled;
- INI naïve HIV infected adults;
- Daily dose of 2mg, 10mg, 50mg or placebo over 10 days;
- Active treatment (n=8) vs matching placebo (n=2) in each dose cohort;
- Rich PK samples on Days 1 and 10 with additional pre-dose samples in between;
- Daily viral load up to Day 11 plus additional viral load on Days 14 & 20 were obtained



# PK/PD Model Structure



Model 1: 
$$I_{\beta}(C) = 1 - \frac{C}{C + IC50}$$

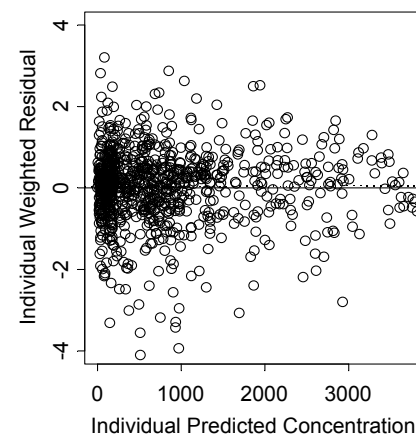
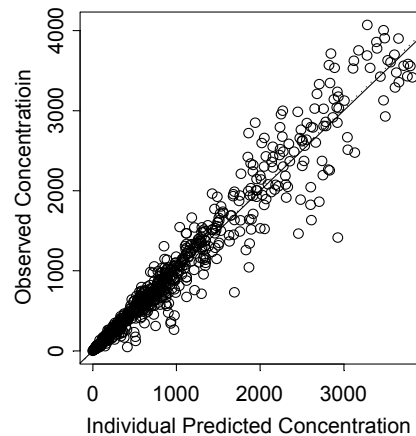
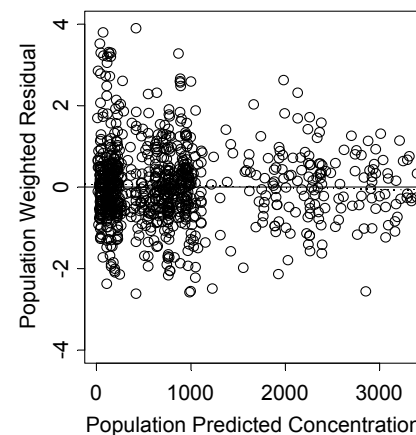
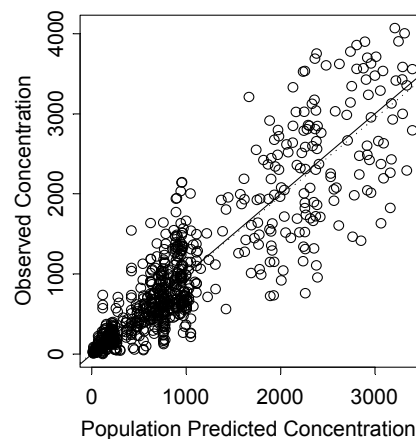
Model 2: 
$$I_{\beta}(C) = 1 - \frac{C}{C + IC50} \quad \& \quad I_{a_L}(C) = 1 - \alpha$$

# Population PK Results

Parameter	Estimate	%CV
$k_a$ (1/hr)	1.47	60.8
$CL/F$ (L/hr)	1.19	33.2
$V_c/F$ (L)	17.9	34.6
$CLD/F$ (L/hr)	0.26	-
$V_p/F$ (L)	1.73	-
$Alag$ (hr)	0.18	60.8
$Age \sim CL/F$	0.34	-
$\sigma^2$	0.034	-

$CL/F$  increases with age by  $\left(\frac{age}{38}\right)^{0.34}$

## Goodness of Fit



# Viral Dynamic Model Results

<u>Input Variable</u>	<u>Symbol</u>	<u>Units</u>	<u>Typical Value</u>
Death (clearance) rate of uninfected CD4+ cells	d	1/day	0.006 <sup>a</sup>
Death (clearance) rate of actively infected CD4+ cells	d <sub>A</sub>	1/day	0.72 <sup>a</sup>
Death (clearance) rate of latently infected CD4+ cells	d <sub>L</sub>	1/day	0.03 <sup>a</sup>
Reactivation rate of latently infected CD4+ cells	a <sub>L</sub>	1/day	0.036 <sup>b</sup>
Fraction of infected CD4+ cells that become actively infected	Q <sub>A</sub>	—	0.97 <sup>a</sup>
Fraction of infected CD4+ cells that become latently infected	Q <sub>L</sub>	—	0.029 <sup>a</sup>
p=viral production rate from actively infected CD4+ cells; c=viral death (clearance) rate	p/c	copies/cell	35.4 <sup>a</sup>

<sup>a</sup>Funk, et al., JAIDS 2001; 26:397-404.

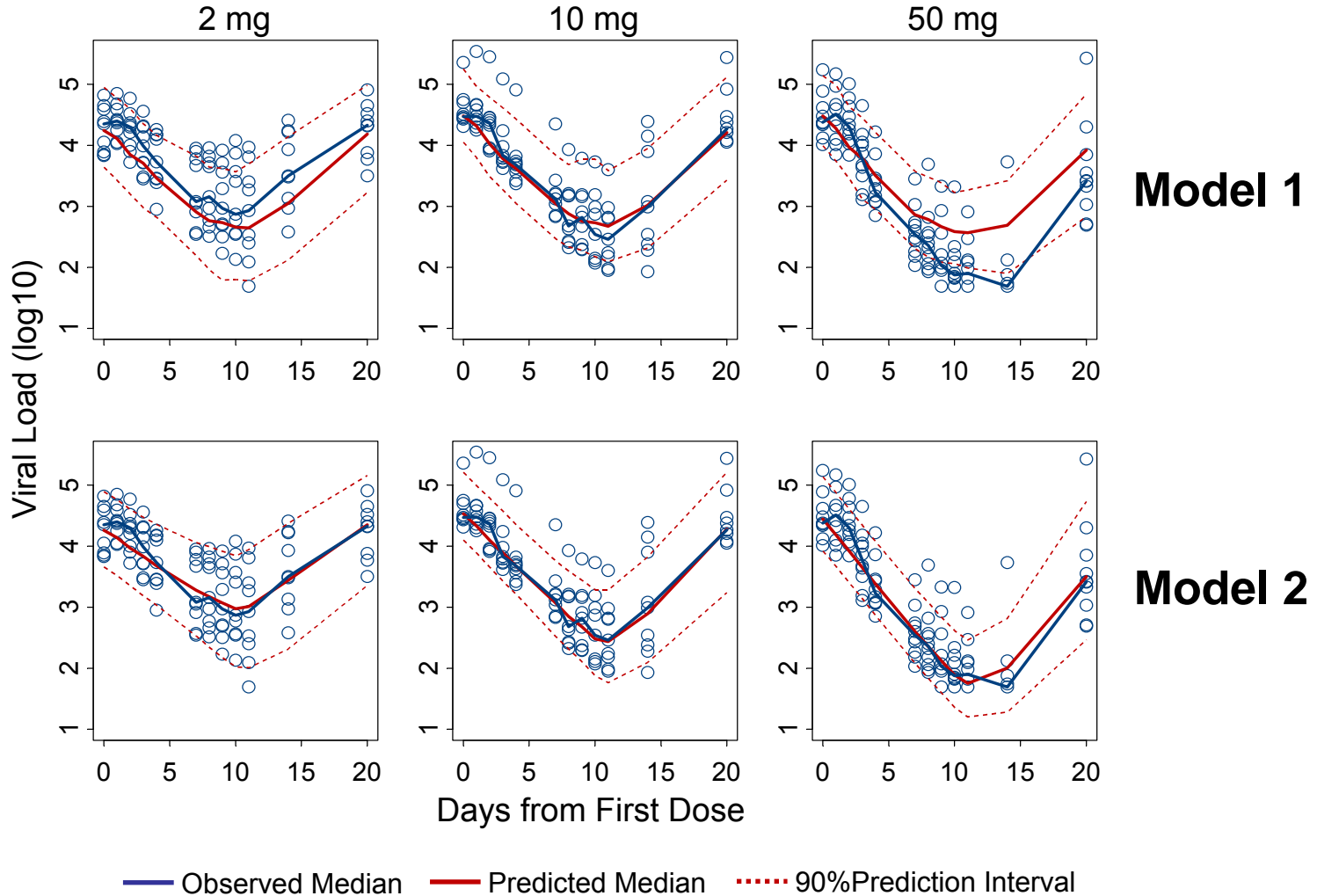
<sup>b</sup>Phillips, Science 1996; 271:497-499.

$$\text{Model 1: } I_{\beta}(C) = 1 - \frac{C}{C + IC50}$$

$$\text{Model 2: } I_{\beta}(C) = 1 - \frac{C}{C + IC50} \quad \& \quad I_{a_L}(C) = 1 - \alpha$$

Parameter	Estimate (%CV)	
	Model 1	Model 2
$R_0$	9.4 (41)	10.3 (39)
$IC50$ (ng/mL)	7.56 (231)	72.7 (70)
$\alpha$	-	0.96
$d_A$ (1/day)	0.63	0.65
$\sigma^2$	0.076	0.048
Parameter	Derived (%CV)	
	Model 1	Model 2
$\beta$ (1/(copies/ $\mu$ L)/day)	0.0020 (85)	0.0018 (86)
$\lambda$ (cells/ $\mu$ L/day)	0.66 (78)	0.64 (77)
<b>Objective Function</b>	-501.6	-667.4

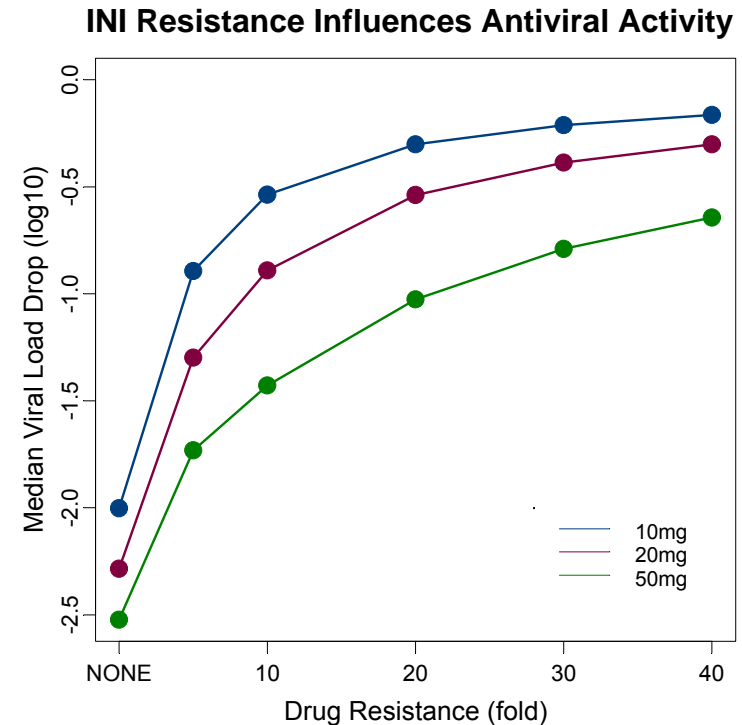
# Model Evaluation and Comparison



# Simulation: Dose-Response in Patients with INI Resistance

## Simulation Conditions:

- Baseline characteristics of INI resistant patients were assumed similar to INI naïve patients
- Drug resistance to INI was introduced by
  - Up to 40-fold increase in IC50
  - Drug effect on latently infected cells reduced by 50%
- Dosing: 10, 20, 50mg QD
- End point: Viral load drop at Day 10
- Number of study replicates: 200



Viral decline by 572 is influenced by the degree of resistance to INI

# Conclusions

- PK of 572 is best characterized by a 2-compartment model with moderate inter-subject variability and relatively small residual errors.
- PK-viral dynamic model adequately described the effect of 572 on viral load in HIV-infected patients.
- The best fit model indicates 572 may act on additional mechanism in HIV life cycle, supporting the potent antiviral activity with 572.
- This model was used to guide dose selection in Phase 2b study (Ivy Song Abstract 50).