Population Pharmacokinetics of Crenolanib, a Type I FLT3 Inhibitor, in Patients with **Relapsed/Refractory Acute Myeloid Leukemia**

ABSTRACT

Background: Crenolanib besylate is a selective and potent type I tyrosine kinase inhibitor (TKI) of wild-type and mutant FMS-like tyrosine kinase 3 (FLT3). Crenolanib is being studied in patients with relapsed/refractory FLT3+ AML with FLT3-ITD and/or FLT3-D835 mutations. This abstract summarizes the population pharmacokinetics (PK) of crenolanib in patients with relapsed or refractory FLT3+ AML.

Methods: Serum blood samples were collected on Day 1 at pre-dose and at 30 min, 60 min, 120 min, 4 h, 8 h Using a one-compartment model with first-order absorption using non-linear mixed-effects modeling, our day 1 and 24 h after single dose of crenolanib, and steady state levels were obtained on Day 15. All crenolanib serum Pharmacokinetic Evaluations pharmacokinetic data were extrapolated to predict the steady-state median, 10th percentile and 90th percentile concentrations were measured by a validated LC-MS/MS. The crenolanib population pharmacokinetics and individual concentrations, and overall range of crenolanib concentrations (Figure 5 and 6). post-hoc estimates were determined by non-linear mixed effects modeling. The inter-individual and inter-occasion With 100 mg TID dosing, steady state was expected to be reached on day 4 (Figure 5 and 7). Sample Collection Serial blood samples were obtained from 55 patients after crenolanib administration during (day 1 vs day 15 studies) variability of the parameters was assumed to be log normally distributed. The area under 4000 cycle 1 on day 1 and day 15 at the following time points: pre-dose, 30 min (±10), 60 min (±15), 120 min (±15), 100 mg TID the concentration-time curve from 0 to 24 hours (AUC; nmol hr/L), maximum concentration (C_{max}; nM), and time 300 mg QD Blue 10th-90th percentile 4 hours (±1), 8 hours (±2) and 24 hours (± 4). median to maximum concentration (T_{max}; hr) were determined from the concentration-time profile simulated using each - - - threshold individual's post-hoc estimated pharmacokinetic parameters. concentration

liquid chromatography-mass spectrometry (LC-MS/MS) method. **Results:** Crenolanib was found to be rapidly absorbed with a T_{max} of 2-3 hours after administration. Crenolanib serum concentration data in cancer patients were best described by a one-compartment PK model with first-order Pharmacokinetic Modeling The crenolanib population pharmacokinetics and individual post-hoc estimates were oral absorption. The model was parameterized in terms of ka (1/hr), the first-order absorption; T_{lag} (hr), lag time; determined by non-linear mixed effects modeling via Monolix (version 4.3.0, www.monolix.org) using the Stochastic CL (L/hr), apparent oral clearance; and V (L), apparent volume. The predictive accuracy of the model used to Approximation Expectation-Maximization (SAEM) approach. The inter-individual and inter-occasion (day 1 vs day forecast these values showed that most of the predicted values are close to the actual values with approximately 15 studies) variability of the parameters was assumed to be log normally distributed. A proportional residual error equal distribution of points on either side of the diagonal line. Hence, the model can be said to be unbiased and model was used with assumed normal distribution of the residuals. fairly accurate. In an attempt to maximize the trough levels and lower C_{max} of crenolanib to continually inhibit FLT3, Using a one-compartment model with first-order absorption using non-linear mixed-effects modeling, day 1 patients were administered crenolanib on a TID schedule. Comparable AUC was reached when crenolanib was pharmacokinetic data were extrapolated to predict the steady-state median, 10th percentile and 90th percentile administered three times daily as opposed to once daily. By day 4 of TID dosing, patients can be expected to reach concentrations, and overall range of crenolanib concentrations. an approximate steady state. During the fourth day of dosing, the predicted median C_{max} was 478 nM (10th percentile The pharmacokinetic model is parameterized in terms of ka (1/hr), the first-order absorption; T_{lag} (hr), lag time; CL of 169 nM and 90th percentile of 1478 nM) and the predicted median trough value was 290 nM (10th percentile of 89 (L/hr), apparent oral clearance; and V (L), apparent volume. The individual post-hoc parameter values were used nM and 90th percentile of 943). 100 mg TID dosing should maintain constant inhibition of FLT3, potentially increasing as the estimates of each patient's pharmacokinetic parameters. In addition, the area under the concentration-time their rate of response. curve (AUC) from 0 to 24 hours (AUC_{last}; hr*nM), maximum concentration (C_{max} ; nM), time to maximum concentration **Conclusions:** Crenolanib is rapidly absorbed with first order pharmacokinetics. Administering crenolanib on a TID T_{max}; hr) and half-life (t_{1/2}; hr) were determined from the concentration-time profile simulated using each individual's schedule allows for a total dose of 300 mg per day of crenolanib, while still maintaining adequate pharmacokinetic post-hoc estimated pharmacokinetic parameters.

properties to lower C_{max} and achieve higher trough levels. Pharmacokinetic profiling of crenolanib over a longer period of time showed that crenolanib does not accumulate with extended use.

Introduction

Features of an Ideal FLT3 Kinase Inhibitor

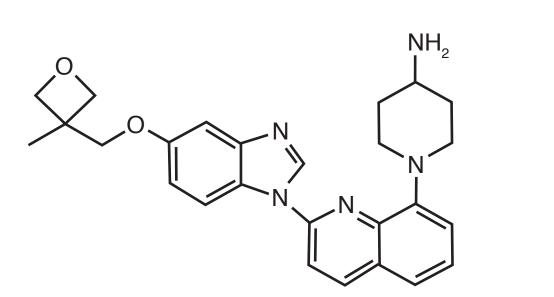
Good oral bioavailability

#3695

- Little interpatient variability
- Low protein binding
- Predictable metabolism and no active metabolite
- Clearance that does not depend on hepatic function Short half-life allowing for combination with cytotoxic chemotherapy
- Little drug accumulation
- No induction of metabolic enzymes that will reduce drug level

Crenolanib - a Potent FLT3 Inhibitor

- **Crenolanib** is a highly potent and specific inhibitor of FMS-like tyrosine kinase 3 (FLT3), Platelet- derived growth factor receptor alpha (PDGFR α) and PDGFR β (Figures 1 and 2) (1-3).
- This orally-administered, type I tyrosine kinase inhibitor (TKI) is able to intercept both inactive and active kinases (1) and is effective against FLT3 D835 mutants that are commonly resistant to other TKIs (4).



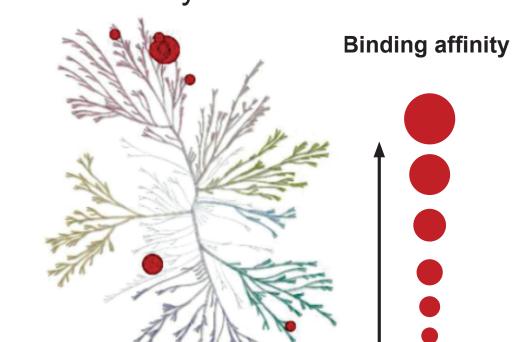


Figure 1. Crenolanib besylate molecular structure.

Figure 2. Kinome scan of crenolanib (5).

• About 27 - 37% of acute myeloid leukemia (AML) patients have mutation(s) in FLT3 (7). • Crenolanib is currently being evaluated in phase II clinical trials in patients with FLT3-mutant,

relapsed / refractory AML.

Prior Pharmacokinetic Studies of Crenolanib

- In vitro data indicate that crenolanib is metabolized via cytochrome P450 (CYP)-3A (AROG data on file).
- The hepatic extraction ratio was estimated to be moderate (0.35-0.53) in human.
- Previous dose finding studies have shown that 300 mg per day is safe for patients with glioma, and TID dosing effectively lowers the C_{max} to minimize toxicity while maintaining a trough level well above the minimum effective concentration (150 nM; 8).
- Food ingestion was found to have no significant effect on crenolanib absorption.
- Crenolanib clinical PK data have been collected in two studies investigating crenolanib in patients with FLT3 mutant AML.

Objectives

• To determine the pharmacokinetics of crenolanib in AML patients The actual serum concentration of crenolanib on day 1 is plotted against concentrations of crenolanib To compare the pharmacokinetics of crenolanib with body surface area (BSA)-based dosing versus that of predicted by a one-compartment model with first-order oral absorption (Figure 4). fixed dosing Most of the predicted values were close to the actual values, and there was approximately equal distribution

Patients and Trial Design

- Two phase II, single center, non-randomized, open-label, single agent studies (ARO-004 and ARO-005; Table 1) evaluated crenolanib in patients with relapsed or refractory AML and FLT3-ITD and/or activating kinase domain mutation.
- Patients received crenolanib orally daily until disease progression or unacceptable toxicity.
- Twenty-six (26) patients received BSA-based dosing of 200 mg/m² per day (divided into
- three doses), and twenty-nine (29) patients received 100 mg TID (three times per day).
- On the first day of treatment, a single dose of crenolanib (66.7 mg/m² or 100 mg) was given to each patient for the purpose of pharmacokinetic study, with the exception of four patients who received three doses.
- Data from a total of 54 patients are presented here.

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Study	# of Patients/ PK Samples	Sampling Intensity	Population	Crenolanib Regimen
ARO-004 NCT01522469	14/107	Serial samples on day 1 and 15	Relapsed / refractory AML with FLT3 mutation	N = 10: 100 mg TID N = 4: 200 mg/m ²
ARO-005 NCT01657682	41/352			N = 19: 100 mg TID N = 22: 200mg/m ²

Table 1. Dosing information across single agent crenolanib studies

Assessment of Crenolanib Concentrations All crenolanib serum concentrations were measured by a validated

The predictive accuracy of the model was verified by a plot of individual predicted concentrations versus actual concentrations for each measured time point.

Patient Characteristics

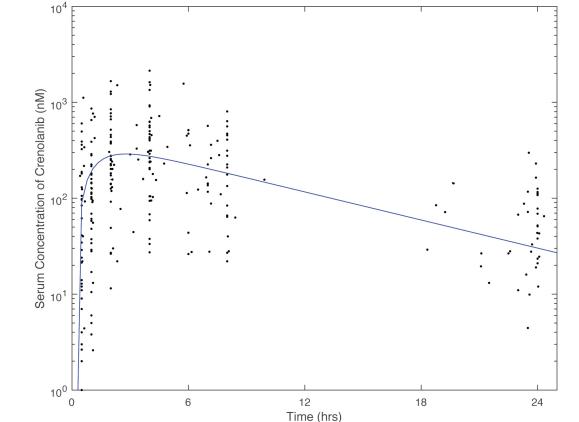
Baseline Characteristics	All Patients (N = 54*)	
Female, N (%)	30 (56)	
Age (years), median (range)	59.5 (21 – 87)	
≤ 60	28	
61-87	26	
Weight (kg), median (range)	69 (48 – 127)	
BSA (m ²), median (range)	1.74 (1.43 – 2.54)	
Creatinine (mg/dL), median (range)	0.67 (0.34 – 1.93)	
Bilirubin (mg/dL), median (range)	0.5 (0.2 – 1.1)	
Alanine aminotransferase (U/L), median (range)	35 (8 – 106)	
Aspartate aminotransferase (U/L), median (range)	29 (9 – 94)	

Table 2. Patient characteristic across two single agent crenolanib studies.
 *One patient was enrolled twice under separate indications and disease episodes. This patient contributed to two sets of samples for pharmacokinetic analysis.

Adequate Exposure Achieved

C_{max} Median is 311 nM and T_{max} is ~2 Hours

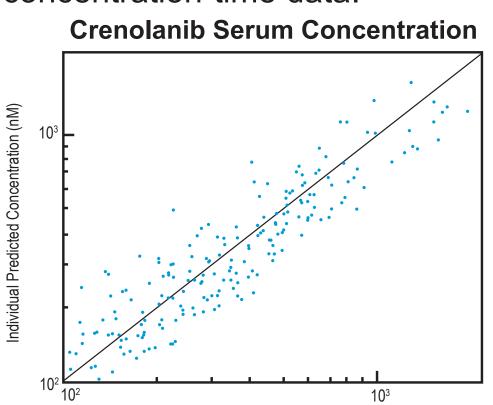
- On the first day of treatment, a single dose of crenolanib was given to all but four patients.
- The concentration of crenolanib in blood was monitored over time.



Parameters	Day 1, Median (Range) (N = 55)				
C _{max}	311 nM (44.94 – 1496.75 nM)				
T _{max}	1.96 hours (0.49 – 8.33 hours)				
t _{1/2}	5.97 hours (2.49 – 13.82 hours)				
AUC _{last}	3483 hr*nM (472.31 – 15362.64 hr*nM)				
C _{max} : highest observed concentration T _{max} : time of Cmax t _{1/2} : half-life AUC _{last} : AUC from 0 to 24 hours was estimated using the trapezoidal rule Table 3. Key PK parameters of a single dose of crenolanib Data are the median (range).					

PK Model Evaluation

of points on either side of the diagonal line. As such, the first-order oral absorption model appeared to accurately describe the serum concentration-time data.



Actual Concentration (nM

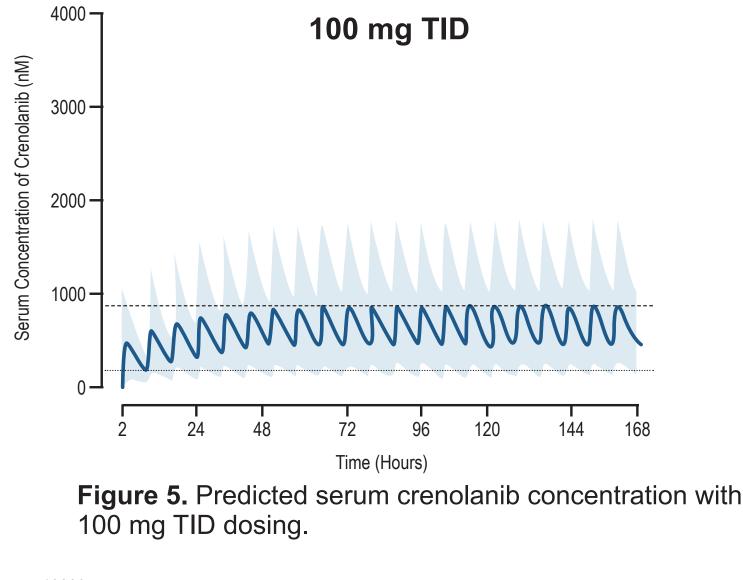
Figure 4. Plot of predicted crenolanib concentration versus actual concentration

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PK Modeling for TID Dosing

- **TID Dosing Lowered C**_{max} and Increased C_{trough}
- Simulations were performed to predict the anticipated peak and trough concentrations (Figure 5 and 6). • Based on experimental observations, the target concentration range was set to approximately 150 nM to 1000 nM



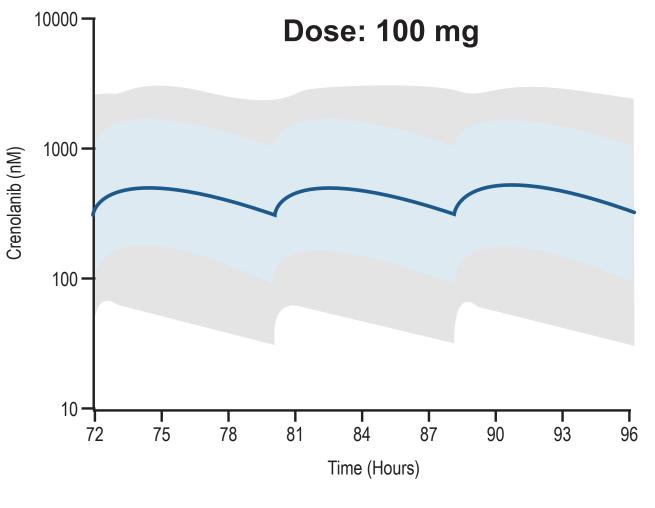
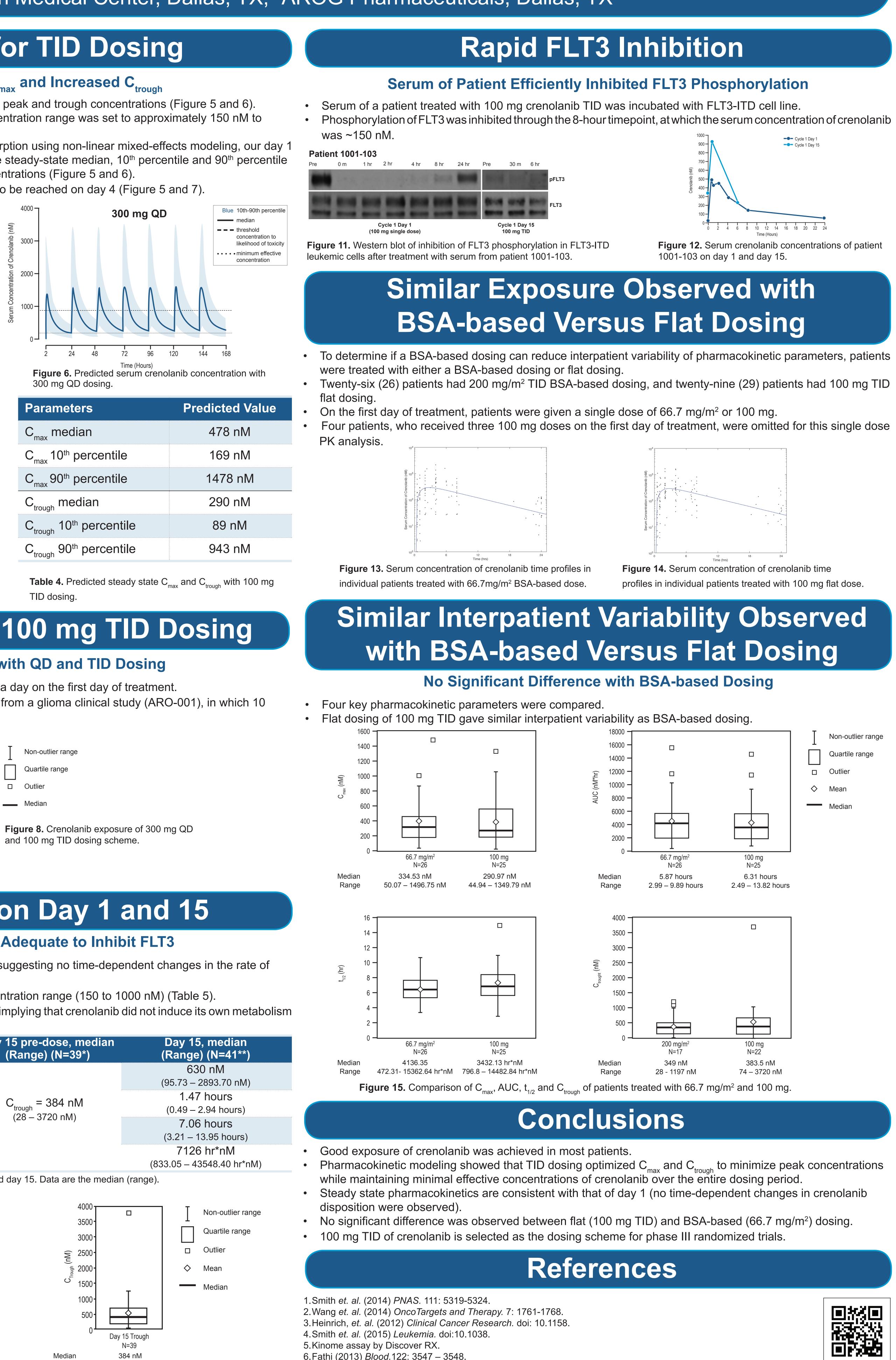


Figure 7. Predicted steady state serum concentration of crenolanib with 100 mg TID dosing.

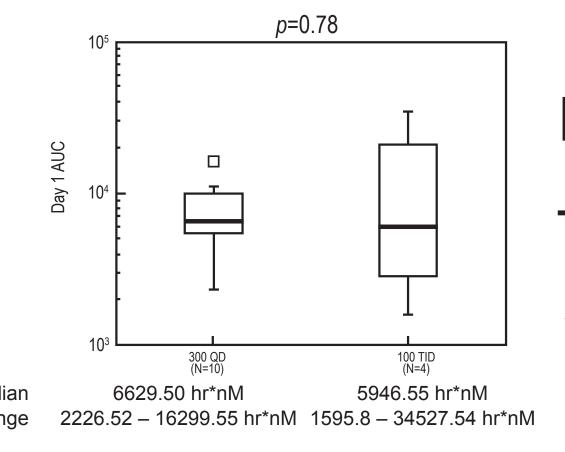


Parameters	Predicted Value
C _{max} median	478 nM
C _{max} 10 th percentile	169 nM
C _{max} 90 th percentile	1478 nM
C _{trough} median	290 nM
C _{trough} 10 th percentile	89 nM
C _{trough} 90 th percentile	943 nM

300 mg QD Versus 100 mg TID Dosing

Similar Drug Exposure with QD and TID Dosing

• Four patients received 100 mg crenolanib three times a day on the first day of treatment. • Their drug exposure (AUC) was compared to patients from a glioma clinical study (ARO-001), in which 10 patients have taken 300 mg crenolanib once daily.



and 100 mg TID dosing scheme.

Consistent PK on Day 1 and 15

Crenolanib Trough Level Adequate to Inhibit FLT3

- T_{max} was reasonably consistent on day 1 and day 15, suggesting no time-dependent changes in the rate of crenolanib absorption (Table 5).
- The median C_{trough} value is well within the target concentration range (150 to 1000 nM) (Table 5). • Crenolanib clearance was similar on day 1 and day 15, implying that crenolanib did not induce its own metabolism

(Figure 9). Day 1, median Day 15 pre-dose, median Parameters (Range) (N=55) (Range) (N=39*) 311 nM (44.94 – 1496.75 nM) 1.96 hours C_{trough} = 384 nM (0.49 – 8.33 hours) (28 – 3720 nM) 5.97 hours (2.49 – 13.82 hours) 3483 hr*nM AUC_{last} (472.31 – 15362.64 hr*nM) Table 5. Comparison of pharmacokinetic parameters between day 1 and day 15. Data are the median (range). Two patients, who had a dose-hold just before day 15, were censored. ** Only 41 out of 55 patients participated in the pharmacokinetic study on day 1

Day 15

61.04 L/hr

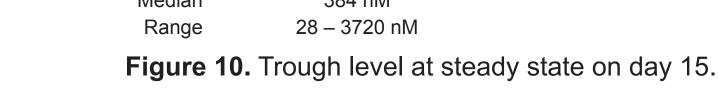
10.67-568.64 L/h

58.90 L/h

13.09 – 591.85 L/h

Figure 9. Comparison of clearance on day 1 and day 15

Range



Day 15 Trough

N=30

4.Smith et. al. (2015) Leukemia. doi:10.1038. 5. Kinome assay by Discover RX.

3. Heinrich, et. al. (2012) Clinical Cancer Research. doi: 10.1158.

6.Fathi (2013) Blood.122: 3547 - 3548.

7.Patel et. al. (2012) NEJM. 366: 1079-1089. 8.Galanis et. al. (2014) Blood 123: 94-100.

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