

Building and evaluation of a PBPK model for Fluconazole in healthy adults

Version	2.0-OSP12.3
based on <i>Model Snapshot and Evaluation Plan</i>	https://github.com/Open-Systems-Pharmacology/Fluconazole-Model/releases/tag/v2.0
OSP Version	12.3
Qualification Framework Version	3.6

This evaluation report and the corresponding PK-Sim project file are filed at:

<https://github.com/Open-Systems-Pharmacology/OSP-PBPK-Model-Library/>

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1 Introduction

Fluconazole is an antifungal medication for the treatment of a number of different fungal infections. It is metabolized by UGT2B7 and renally excreted by glomerular filtration. About 70% of fluconazole is excreted unchanged in urine during 120 hours. Fluconazole is a moderate inhibitor of CYP3A4 and can therefore serve as a perpetrator for CYP3A4 in DDI studies.

The herein presented PBPK model of fluconazole has been developed using published pharmacokinetic clinical data by Palkama et al. ([Palkama 1998](#)), Ripa et al. ([Ripa 1993](#)), Shiba et al. ([Shiba 1990](#)), Ahonen et al. ([Ahonen 1997](#)), Brammer et al. ([Brammer 1990](#)), Brammer et al. ([Brammer 1991](#)), Thorpe et al. ([Thorpe 1990](#) and [Varhe et al. \(Varhe 1996\)](#)). The model has then been evaluated by comparing observed data to simulations of both intravenously and orally administered fluconazole covering a dose range of 25 mg to 400 mg. Both single dose and multiple dose administration are represented in the development and evaluation data sets.

The presented model includes the following features:

- metabolism by UGT2B7,
- inhibition of CYP3A4,
- renal clearance by glomerular filtration,
- oral absorption with dissolution rate assigned to the Weibull function for low, medium and high doses.

2 Methods

2.1 Modeling strategy

The general concept of building a PBPK model has previously been described by Kuepfer et al. ([Kuepfer 2016](#)). Relevant information on anthropometric (height, weight) and physiological parameters (e.g. blood flows, organ volumes, binding protein concentrations, hematocrit, cardiac output) in adults was gathered from the literature and has been previously published ([Willmann 2007](#)). The information was incorporated into PK-Sim® and was used as default values for the simulations in adults.

The applied activity and variability of plasma proteins and active processes that are integrated into PK-Sim® are described in the publicly available PK-Sim® Ontogeny Database Version 7.3 ([PK-Sim Ontogeny Database Version 7.3](#)) or otherwise referenced for the specific process.

A mean model was built based on clinical data from studies with intravenous administration of fluconazole by Palkama et al. 1998 ([Palkama 1998](#)), Ripa et al. 1993 ([Ripa 1993](#)) and Shiba et al. 1990 ([Shiba 1990](#)). The studies conducted by Palkama et al. and Ripa et al. reported plasma concentrations of fluconazole and the study by Shiba reported fraction of unchanged fluconazole excreted to urine in addition to plasma concentrations. The mean PBPK model was developed using a typical male European individual. The relative tissue-specific expressions of enzymes predominantly being involved in the metabolism of fluconazole (UGT2B7) were considered ([Meyer 2012](#)).

A specific set of parameters (see below) was optimized to describe the distribution of fluconazole after intravenous administration using the Parameter Identification module provided in PK-Sim®. Structural model selection was mainly guided by visual inspection and total error of the resulting description of data, 95% confidence interval of the identified parameter values and biological plausibility.

Once the appropriate structural model was identified for the intravenous administration, the model was verified by simulating other clinical studies reporting pharmacokinetic concentration-time profiles after intravenous administration of fluconazole.

Thereafter additional parameters for oral administration of non-dissolved formulations (i.e. capsules) were identified. Administered doses of fluconazole were ranging from single dose of 25 mg to multiple dosing of 400 mg, why the applied Weibull function to describe the dissolution of fluconazole were divided into low, medium and high dose. The dissolution shape were kept between the doses while the dissolution time were separated.

The model was then verified by simulating further clinical studies reporting pharmacokinetic concentration-time profiles after oral administration of fluconazole.

Details about input data (physicochemical, *in vitro* and clinical) can be found in [Section 2.2](#).

Details about the structural model and its parameters can be found in [Section 2.3](#).

2.2 Data used

2.2.1 In vitro and physicochemical data

A literature search was performed to collect available information on physicochemical properties of fluconazole. The obtained information from literature is summarized in the table below, and is used for model building.

Parameter	Unit	Value	Source	Description
MW	g/mol	306.28	Debruyne 1993	Molecular weight
pK _a		2.03	Charoo 2014	acid dissociation constant of conjugate acid; compound type: weak base
Solubility (pH)	mg/mL	6.9 (7.4)	Charoo 2014	Aqueous Solubility
logP		0.5	Christofolletti 2016	Partition coefficient between octanol and water
		0.5	Charoo 2014	Partition coefficient between octanol and water
fu	%	89	Pfizer	Fraction unbound in plasma
	%	88	Christofolletti 2016	Fraction unbound in plasma
CL _{int} UGT2B7	1/min	0.005	Watt 2018	First order intrinsic clearance UGT2B7
P _{eff}	cm/s	0.000213	GastroPlus	Specific intestinal permeability
Ki CYP3A4	μmol/L	13.1	Sakaeda 2005	Non-competitive inhibition of CYP3A4
GFR fraction		0.17	Watt 2018	glomerular filtration fraction

2.2.2 Clinical data

A literature search was performed to collect available clinical data on fluconazole in adults.

The following publications were found in adults for model building:

Publication	Arm / Treatment / Information used for model building
Ahonen 1997	Plasma PK profiles in healthy subjects after single dose oral administration of 400 mg fluconazole
Brammer 1990	Plasma PK profiles in healthy subjects after multiple oral administration of 200, 300 and 400 mg fluconazole
Brammer 1991	Plasma PK profiles in healthy subjects after single dose oral administration of 50 mg fluconazole
Shiba 1990	Plasma PK profiles and urine data in healthy subjects after single iv and oral administration of 25 and 50 mg fluconazole and single oral dose of 100 mg fluconazole
Palkama 1998	Plasma PK profiles in healthy subjects after single dose iv and oral administration of 400 mg fluconazole
Ripa 1993	Plasma PK profiles in healthy subjects after single dose iv administration of 100 mg fluconazole
Thorpe 1990	Plasma PK profiles in healthy subjects after single dose oral administration of 100 mg fluconazole
Varhe 1996	Plasma PK profiles in healthy subjects after multiple oral administration of 100 and 200 mg fluconazole

The following table shows the data from the excretion studies ([Shiba 1990](#)) used for model building:

Observer	Value
Fraction excreted to urine of unchanged fluconazole after iv administration 25 mg	74%
Fraction excreted to urine of unchanged fluconazole after iv administration 25 mg	72%
Fraction excreted to urine of unchanged fluconazole after oral administration 25 mg	69%
Fraction excreted to urine of unchanged fluconazole after oral administration 50 mg	67%
Fraction excreted to urine of unchanged fluconazole after oral administration 100 mg	75%

The following dosing scenarios were simulated and compared to respective data for model verification:

Scenario	Data reference
iv 400 mg (60 min)	Ahonen 1997
iv 50 mg (2 min)	Brammer 1990
iv 25 mg (1 min)	Shiba 1990
iv 800 mg (240 min) once daily for 14 days	Sobue 2004
iv 100 mg (20 min)	Yeates 1994
po 400 mg	Palkama 1998
po 100 mg	Shiba 1990
po 50 mg once daily for 4 days	Varhe 1996

2.3 Model parameters and assumptions

2.3.1 Absorption

The parameter value for `Specific intestinal permeability` was used as in GastroPlus. Although, the permeability didn't seem to be the rate-limiting step in oral absorption as this parameter could not be identified when attempting to estimate a value using Parameter Identification. The default solubility was assumed to be the measured value in phosphate buffer (see [Section 2.2.1](#))

The dissolution of capsules/tablets (not always specified in the literature which formulation for oral administration was used) were implemented via empirical Weibull dissolution. A different Weibull function was used for different dose categories, low, medium high and high dose to distinguish the dissolution time across the dose range; see results of optimization in [Section 2.3.4](#).

2.3.2 Distribution

Fluconazole has a low protein bound (approx. 11 %) fraction in plasma (see [Section 2.2.1](#)). A value of 89% was used in this PBPK model for `Fraction unbound (plasma, reference value)`. The major binding partner was set to alpha1- acid glycoprotein(see [Section 2.2.1](#)).

An important parameter influencing the resulting volume of distribution is lipophilicity. The reported experimental logP values are in the range of 0.5-1.0 (see [Section 2.2.1](#)) which served as a starting value. Finally, the model parameters `Lipophilicity` was optimized to match best clinical data (see also [Section 2.3.4](#)).

After testing the available organ-plasma partition coefficient and cell permeability calculation methods built in PK-Sim, observed clinical data was best described by choosing the partition coefficient calculation by `Rodgers and Rowland` and cellular permeability calculation by `PK-Sim Standard`.

2.3.3 Metabolism and Elimination

One metabolic pathway was implement into the model via first order kinetics

- UGT2B7

The UGT2B7 expression profiles is based on high-sensitive real-time RT-PCR (Nishimura 2003). Absolute tissue-specific expressions were obtained by considering the respective absolute concentration in the liver. The `Specific clearance` of fluconazole accounted by UGT2B7 was identified to best describe observed clinical data after intravenous administration (see Section 2.3.4).

Additionally, renal clearance (which is the main elimination process for fluconazole) assumed to be mainly driven by glomerular filtration was implemented. The `GFR fraction` was identified to best describe the observed fraction of unchanged fluconazole that was renally excreted (see Section 2.3.4).

2.3.4 Automated Parameter Identification

This is the result of the final parameter identification for the intravenous model:

Model Parameter	Optimized Value	Unit
<code>Lipophilicity</code>	0.83	Log Units
<code>GFR fraction</code>	0.14	
<code>Specific clearance</code> (UGT2B7)	1.85E-3	1/min

This is the result of the final parameter identification for the dissolution parameters of a oral administered fluconazole:

Model Parameter	Optimized Value	Unit
<code>Dissolution time (50% dissolved) high dose</code>	105.81	min
<code>Dissolution time (50% dissolved) medium high dose</code>	75.14	min
<code>Dissolution time (50% dissolved) low dose</code>	33.40	min
<code>Dissolution shape</code>	2.14	

3 Results and Discussion

The PBPK model for fluconazole was developed and verified with clinical pharmacokinetic data.

The model was built and evaluated covering data from studies including in particular

- intravenous infusions and oral administrations (capsules).
- a dose range of 25 to 800 mg.

The model quantifies excretion via urine (glomerular filtration) and metabolism via UGT2B7.

The next sections show:

1. the final model input parameters for the building blocks: [Section 3.1](#).
2. the overall goodness of fit: [Section 3.2](#).
3. simulated vs. observed concentration-time profiles for the clinical studies used for model building and for model verification: [Section 3.3](#).

3.1 Fluconazole final input parameters

The compound parameter values of the final PBPK model are illustrated below.

Compound: Fluconazole

Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	6.9 mg/ml		Measurement_Solubility_Charoo2014	True
Reference pH	7		Measurement_Solubility_Charoo2014	True
Lipophilicity	0.8302390226 Log Units	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 1' on 2021-07-13 15:29	Optimized_Lipophilicity	True
Fraction unbound (plasma, reference value)	89 %		Measurement_Fraction_Unbound_Pfizer	True
Specific intestinal permeability (transcellular)	0.000213 cm/s		Measurement_Specific_Intestinal_Permeability_GastroPlus	True
F	2			
Is small molecule	Yes			
Molecular weight	306.328 g/mol			
Plasma protein binding partner	α 1-acid glycoprotein			

Calculation methods

Name	Value
Partition coefficients	Rodgers and Rowland
Cellular permeabilities	PK-Sim Standard

Processes

Metabolizing Enzyme: UGT2B7-Watt2018

Species: Human

Molecule: UGT2B7

Parameters

Name	Value	Value Origin
Intrinsic clearance	0.008 l/min	
Specific clearance	0.0018460294654 1/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 1' on 2021-07-13 15:29

Systemic Process: Glomerular Filtration-Watt2018

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	0.1433240785	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 1' on 2021-07-13 15:29

Inhibition: CYP3A4-Sakaeda2005

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
Ki	13.1 µmol/l	

Formulation: Fluconazole_Weibull_high dose

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	105.8071997888 min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Lag time	0 min	
Dissolution shape	2.1410681544	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Use as suspension	Yes	

Formulation: Fluconazole_Weibull_low dose

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	33.3965005792 min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Lag time	0 min	
Dissolution shape	2.1410681544	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Use as suspension	Yes	

Formulation: Fluconazole_Weibull_medium high dose

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	75.1395611322 min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Lag time	0 min	
Dissolution shape	2.1410681544	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification 2' on 2021-07-13 18:01
Use as suspension	Yes	

3.2 Fluconazole Diagnostics Plots

Below you find the goodness-of-fit visual diagnostic plots for the PBPK model performance of all data used presented in [Section 2.2.2](#).

The first plot shows observed versus simulated plasma concentration, the second weighted residuals versus time.

Table 3-1: GMFE for Goodness of fit plot for concentration in plasma.

Group	GMFE
Fluconazole iv (model building)	1.09
Fluconazole iv (model verification)	1.14
Fluconazole oral administration (model building)	1.20
Fluconazole oral administration (model verification)	1.26
All	1.18

- Fluconazole iv (model building)
- Fluconazole iv (model verification)
- ◆ Fluconazole oral administration (model building)
- ▲ Fluconazole oral administration (model verification)

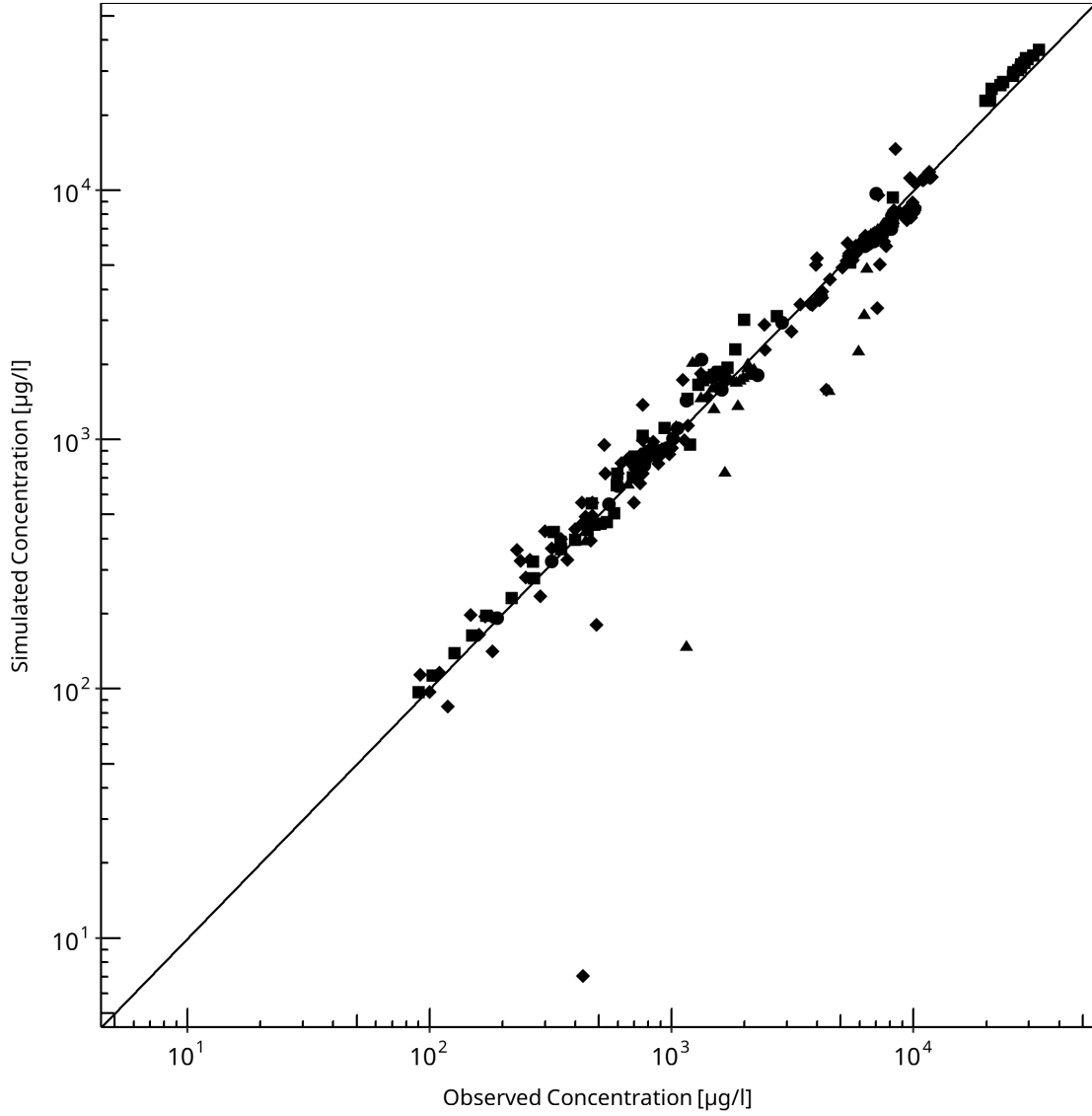


Figure 3-1: Goodness of fit plot for concentration in plasma.

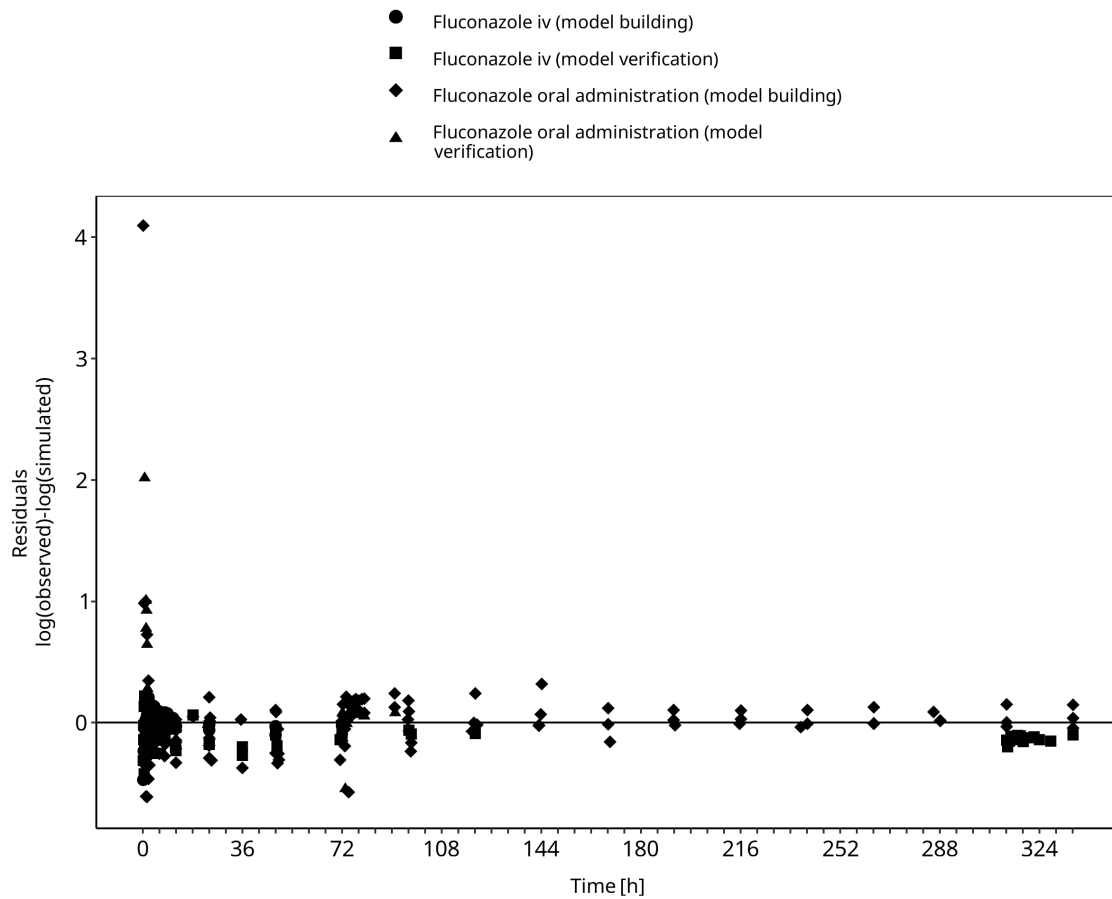


Figure 3-2: Goodness of fit plot for concentration in plasma.

3.3 Concentration-Time Profiles

Simulated versus observed concentration-time profiles of all data listed in [Section 2.2.2](#) are presented below.

3.3.1 Model Building

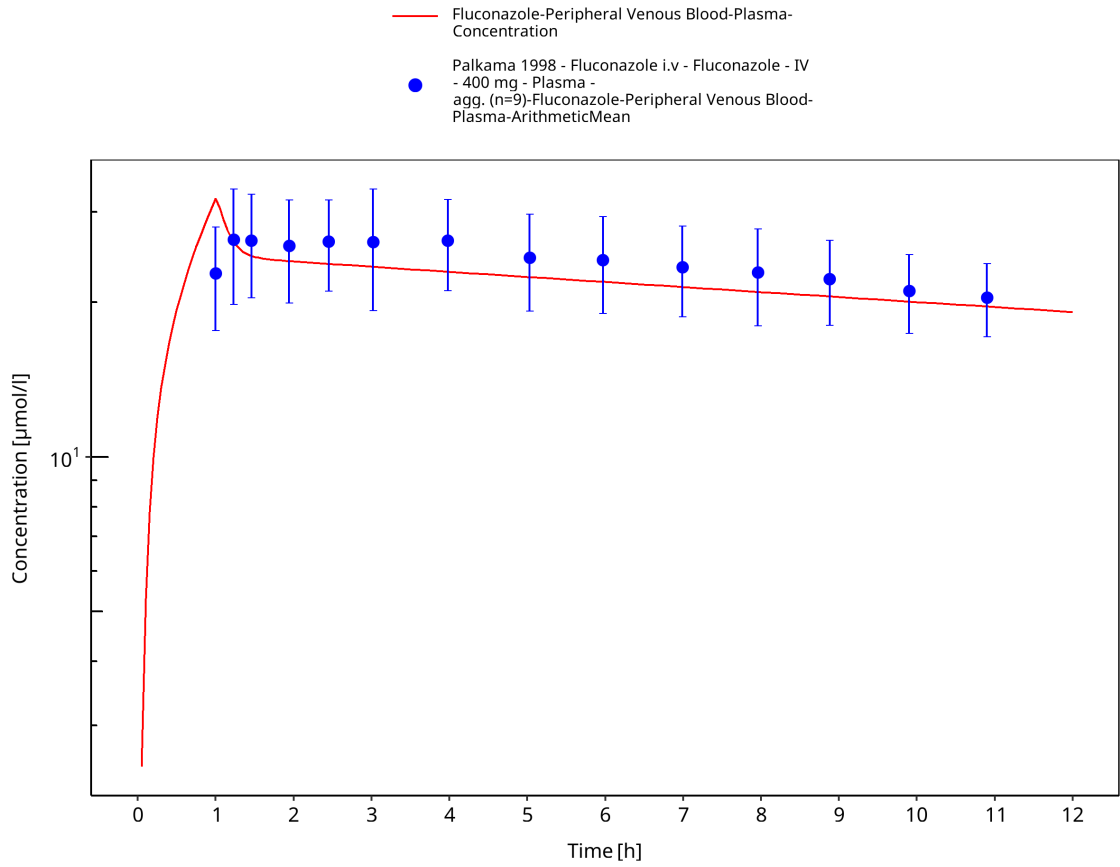


Figure 3-3: Time Profile Analysis

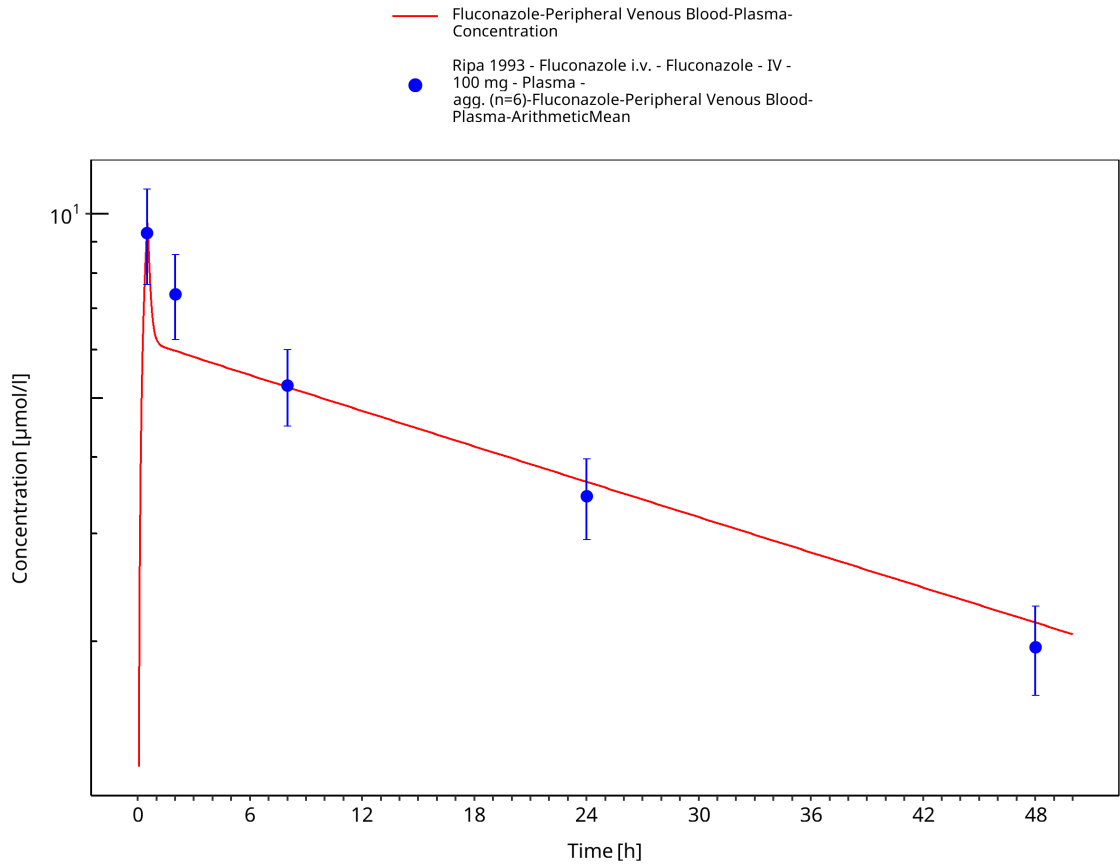


Figure 3-4: Time Profile Analysis

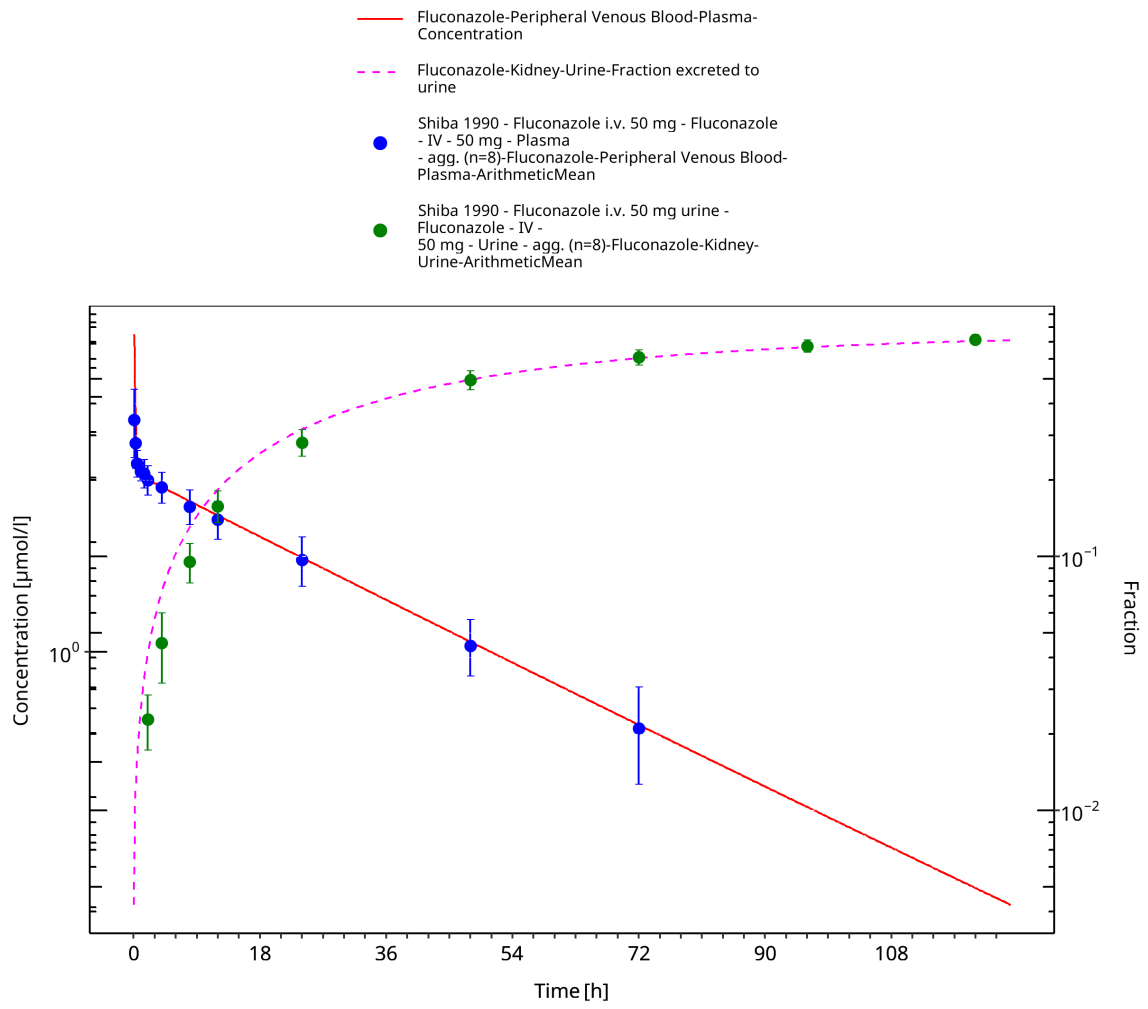


Figure 3-5: Time Profile Analysis

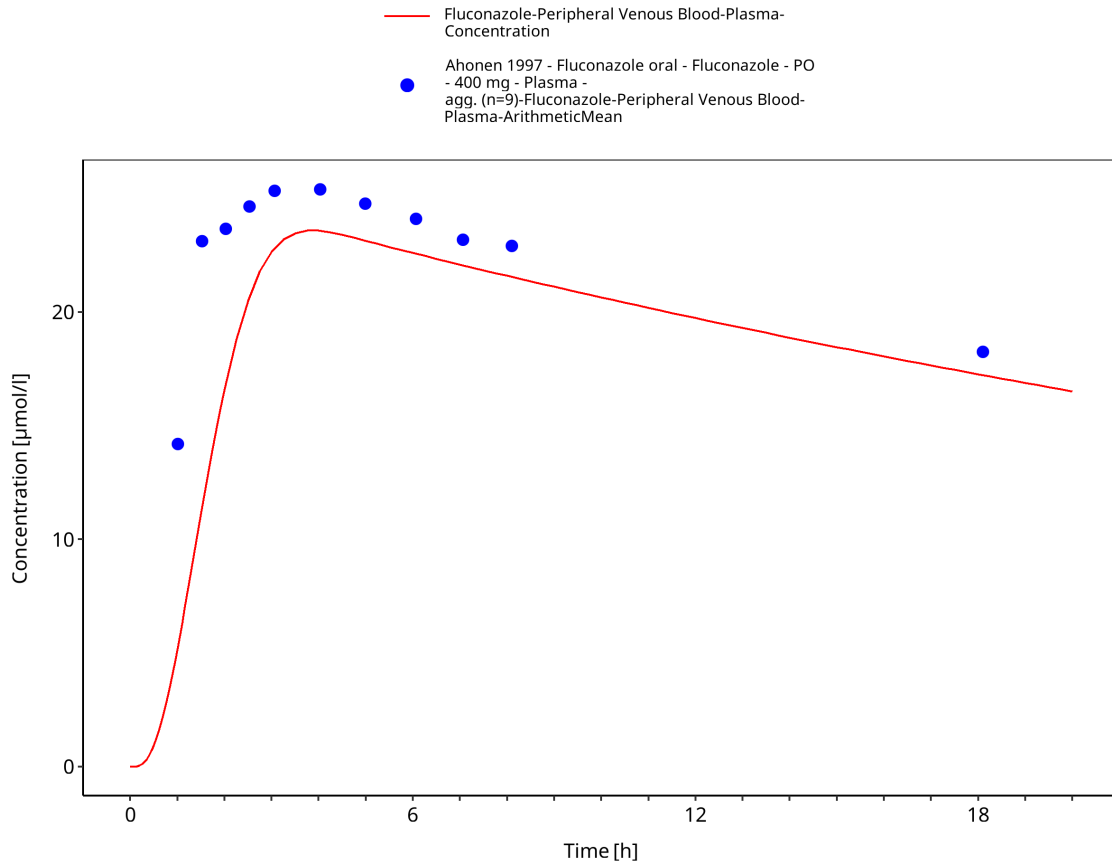


Figure 3-6: Time Profile Analysis

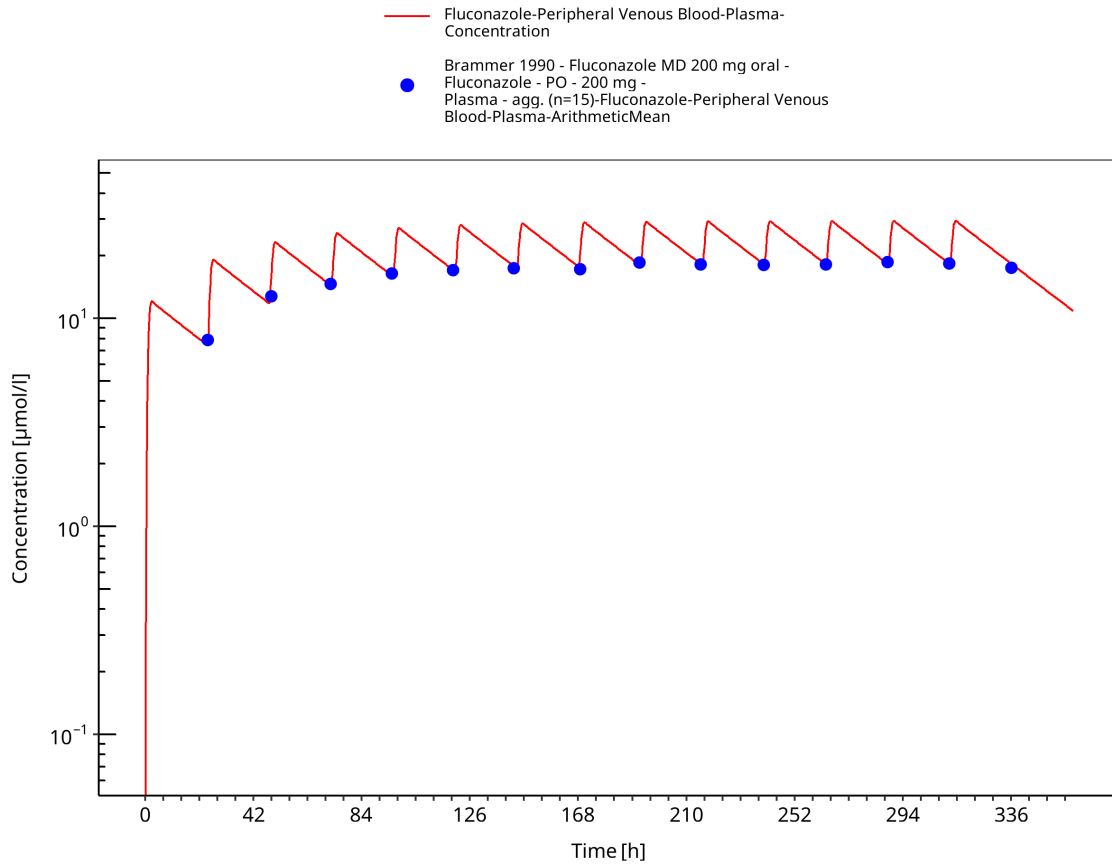


Figure 3-7: Time Profile Analysis

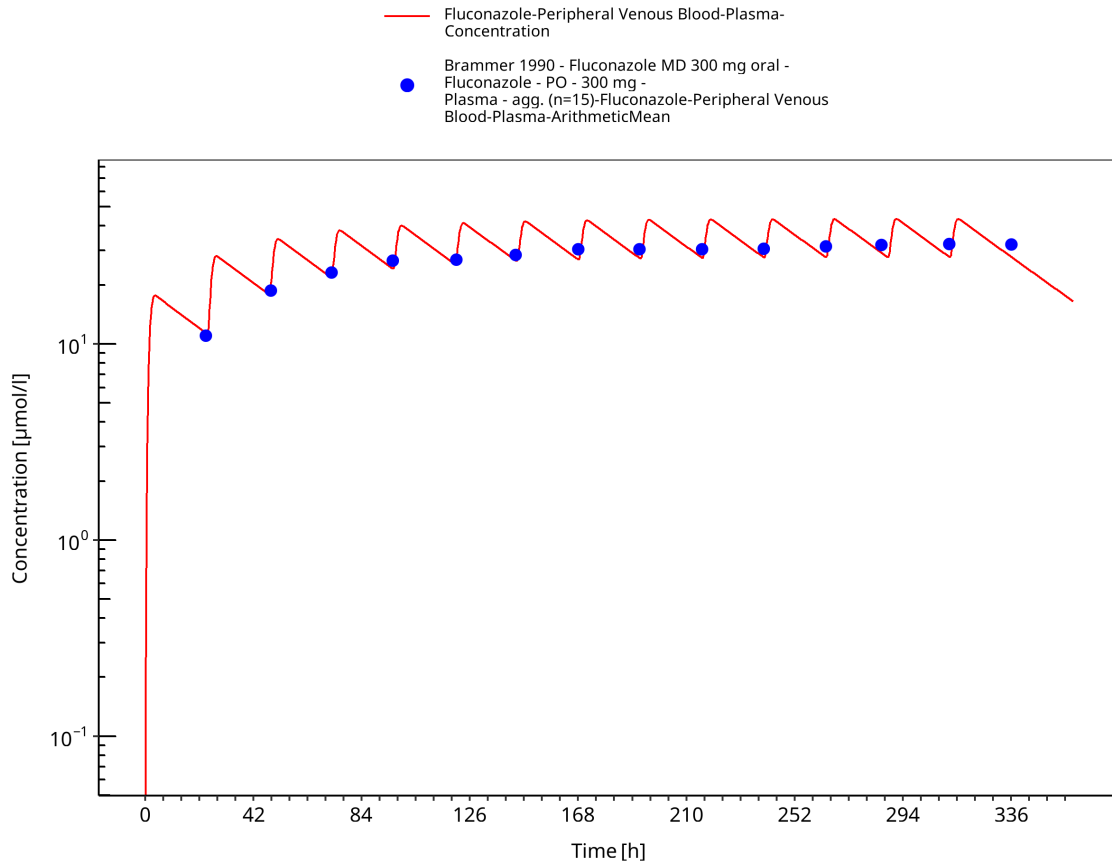


Figure 3-8: Time Profile Analysis

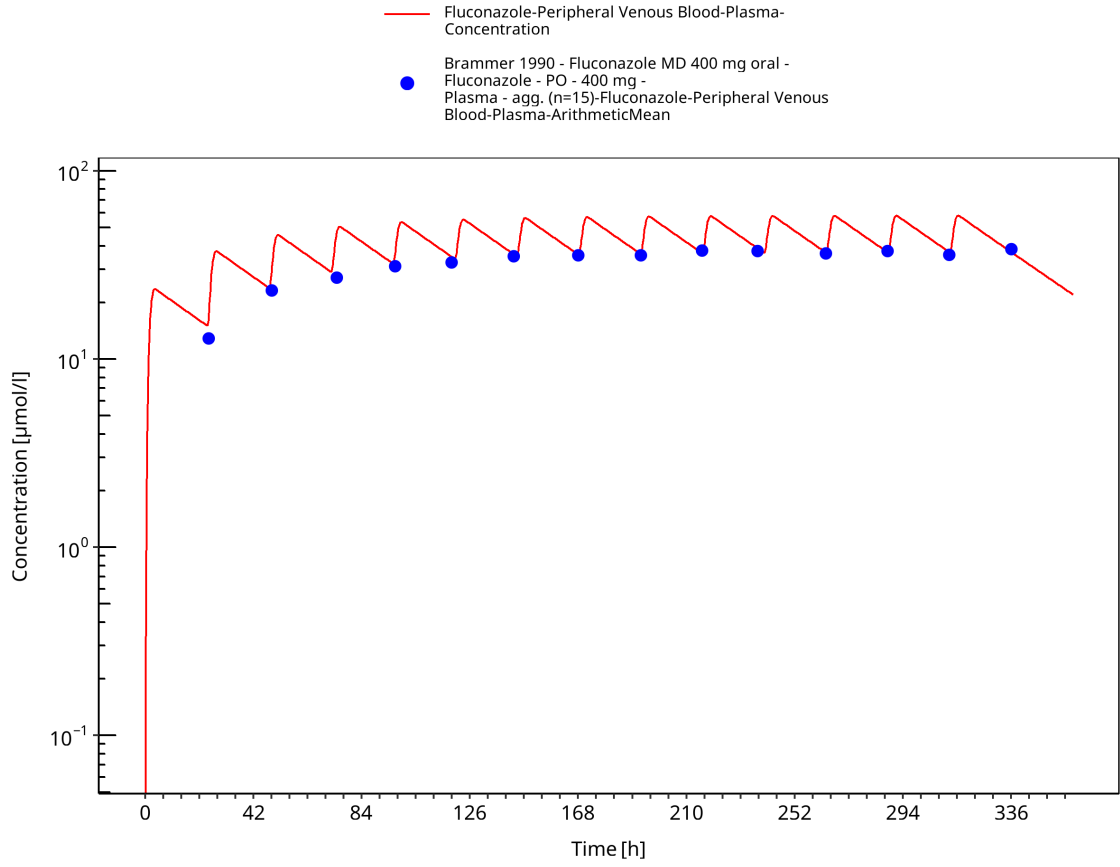


Figure 3-9: Time Profile Analysis

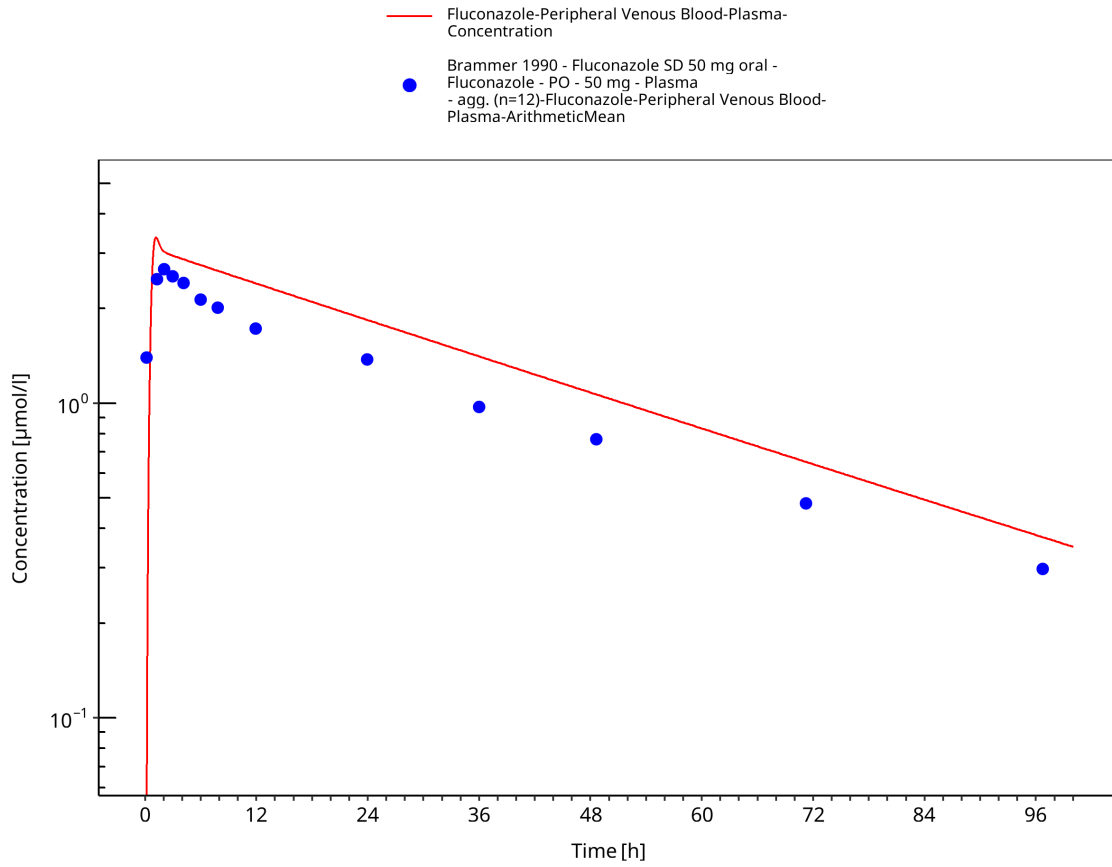


Figure 3-10: Time Profile Analysis

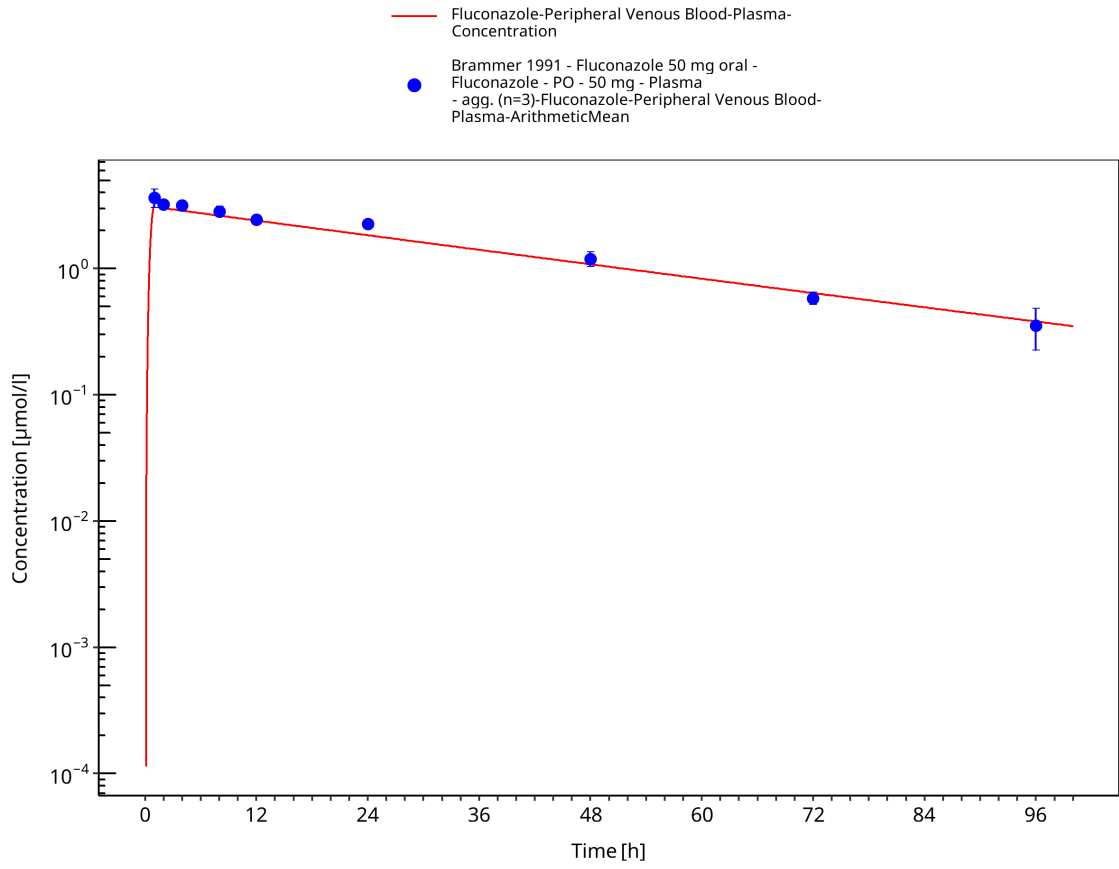


Figure 3-11: Time Profile Analysis

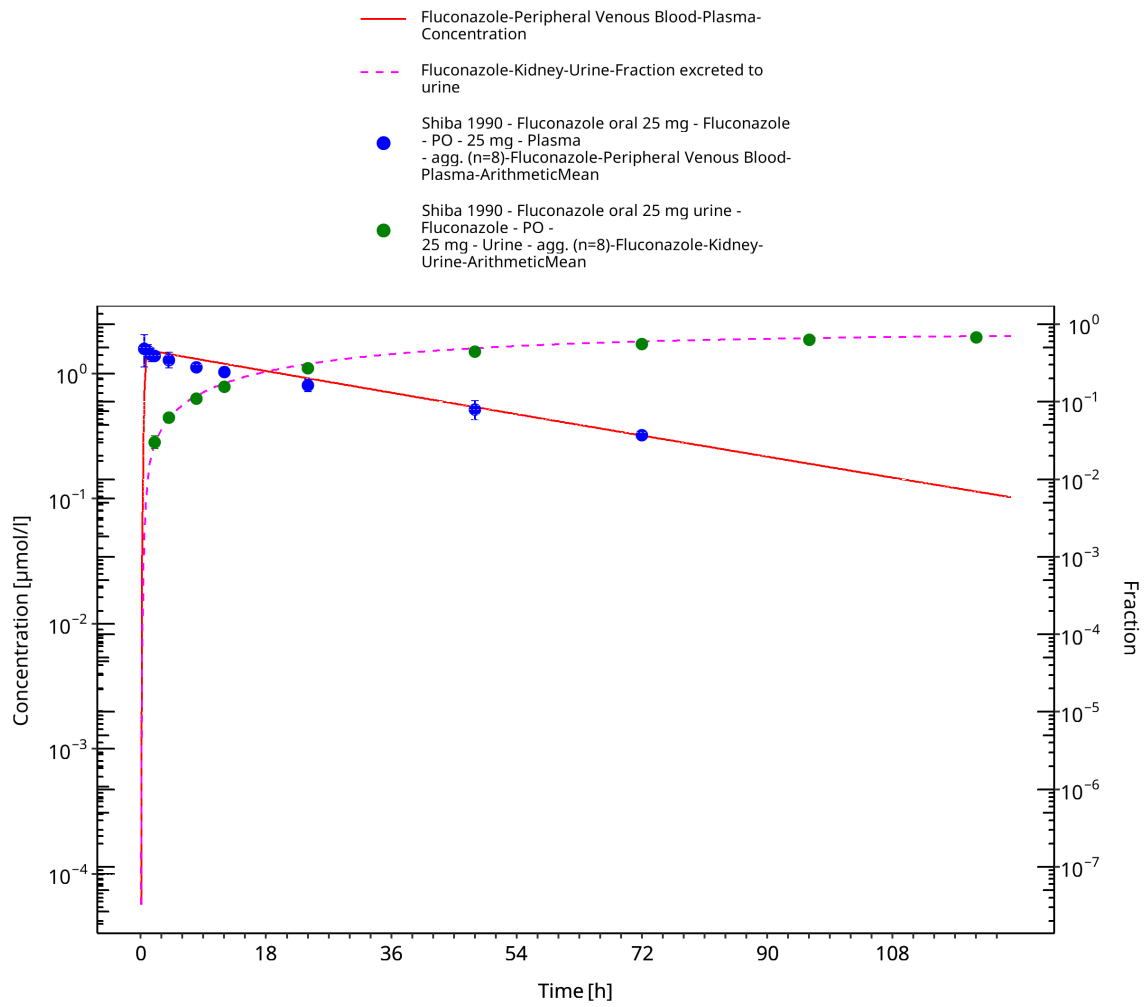


Figure 3-12: Time Profile Analysis

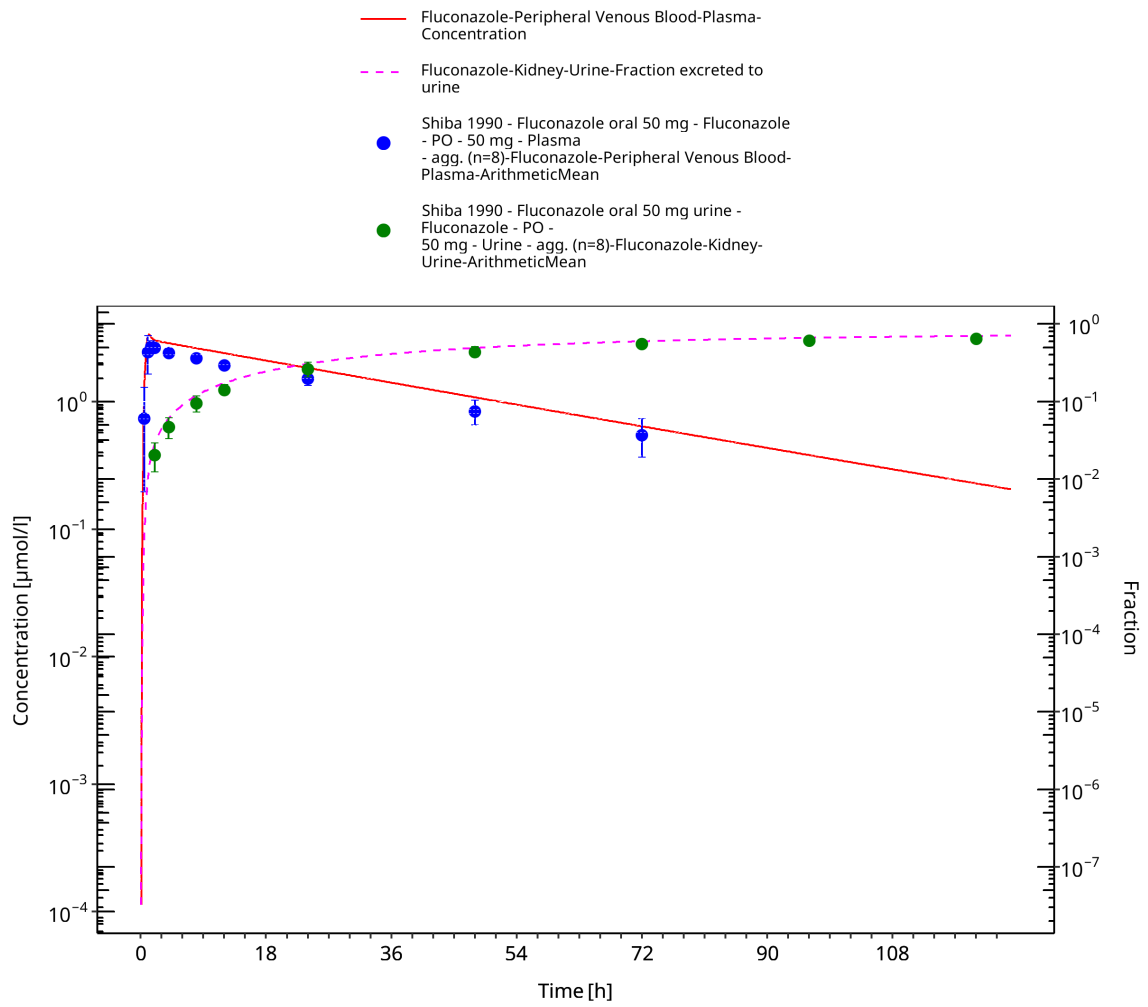


Figure 3-13: Time Profile Analysis

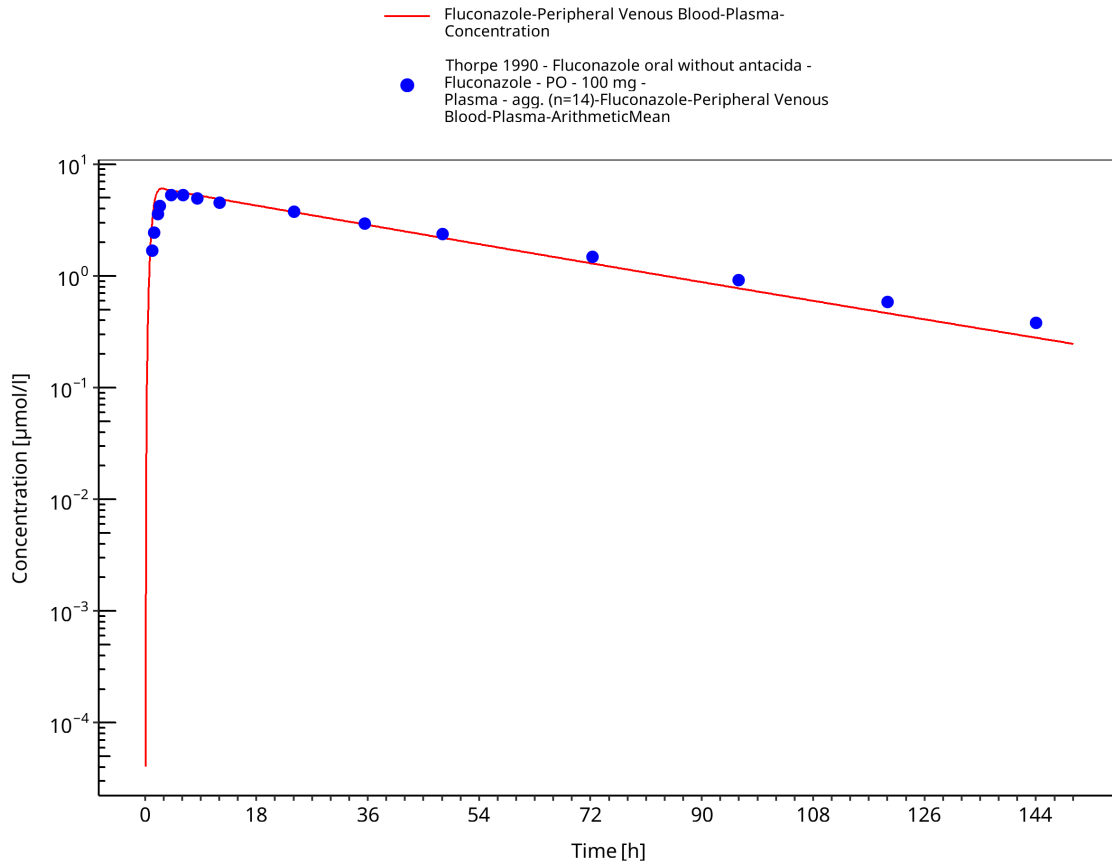


Figure 3-14: Time Profile Analysis

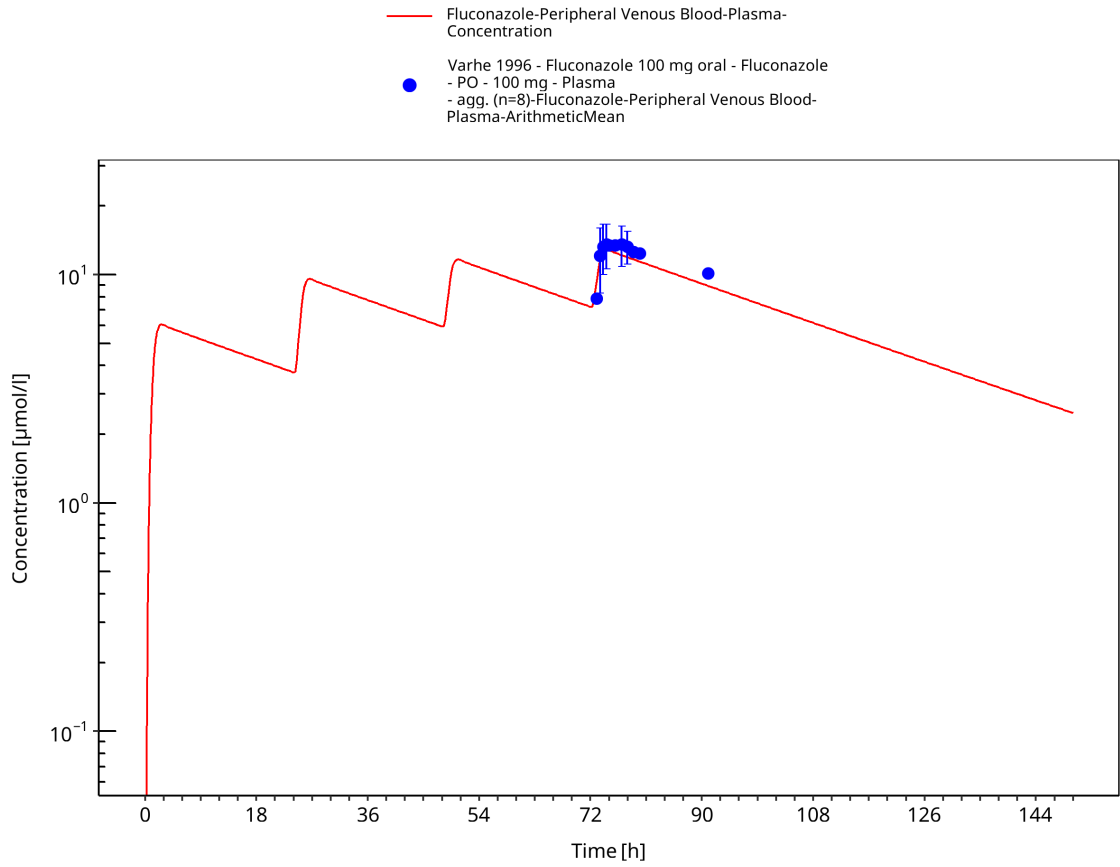


Figure 3-15: Time Profile Analysis

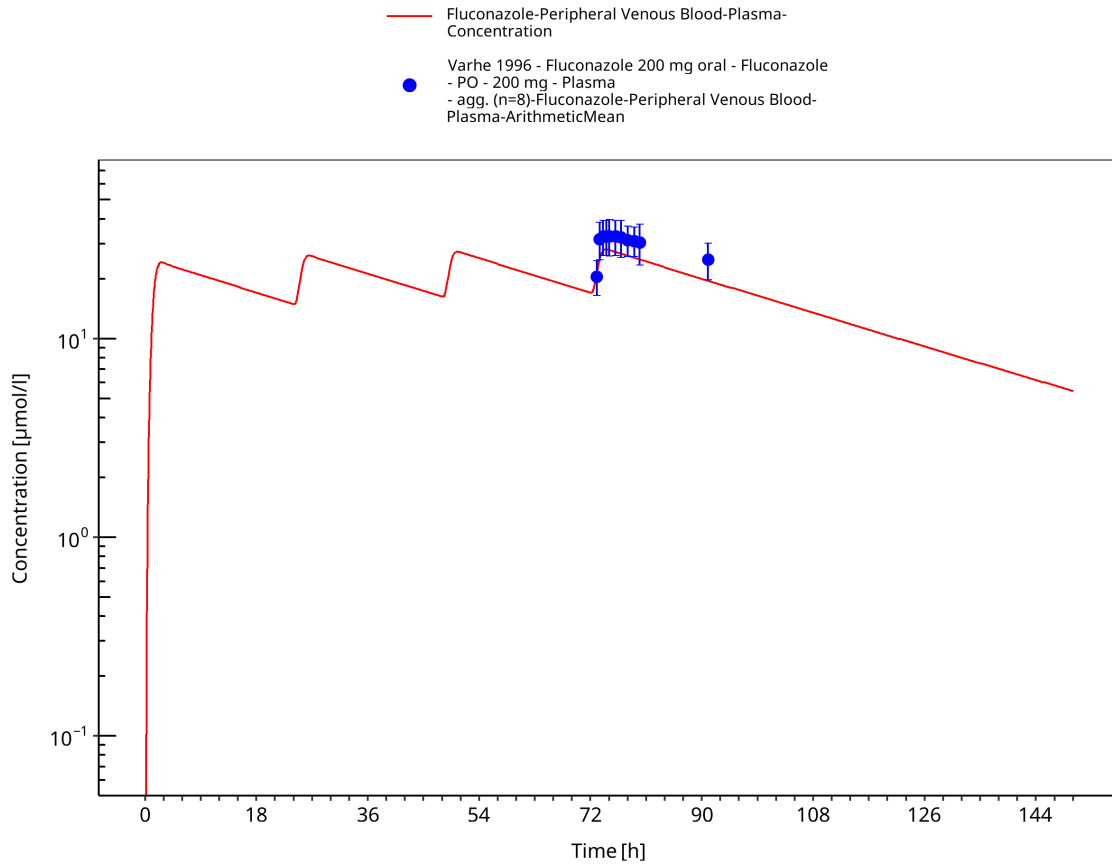


Figure 3-16: Time Profile Analysis

3.3.2 Model Verification

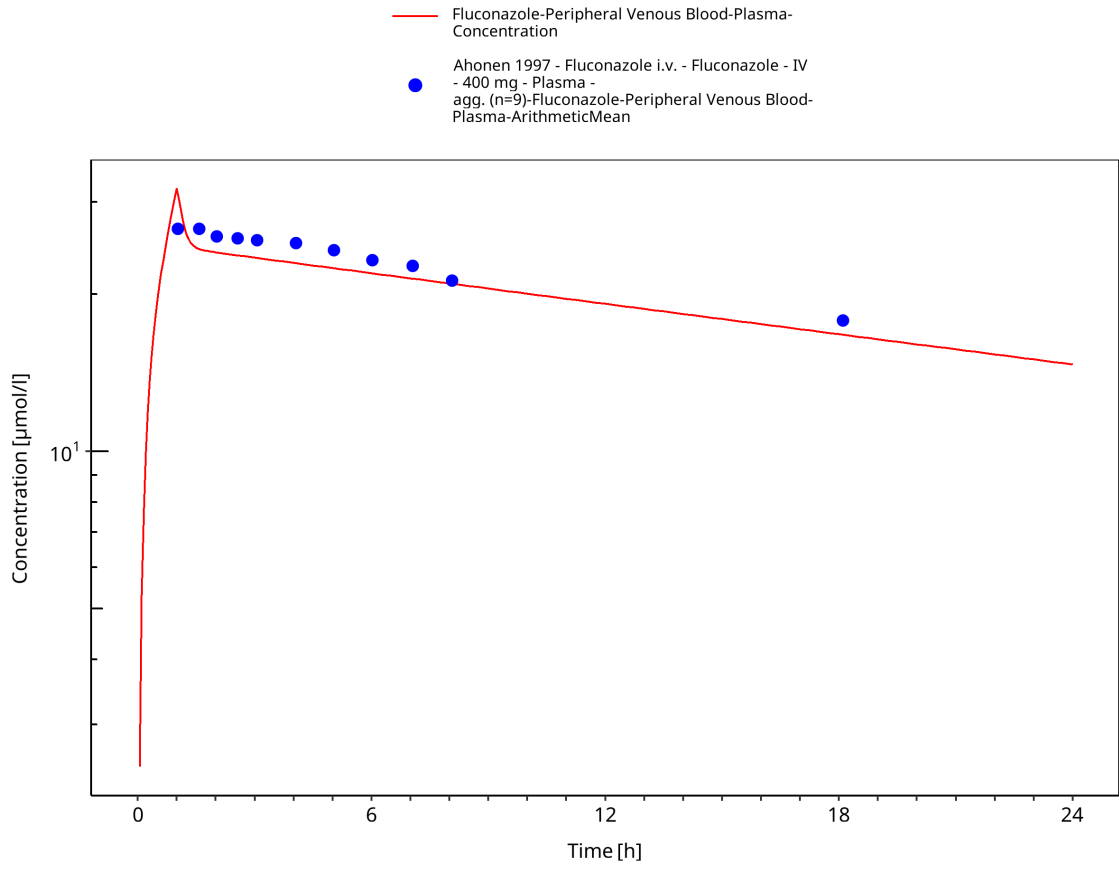


Figure 3-17: Time Profile Analysis

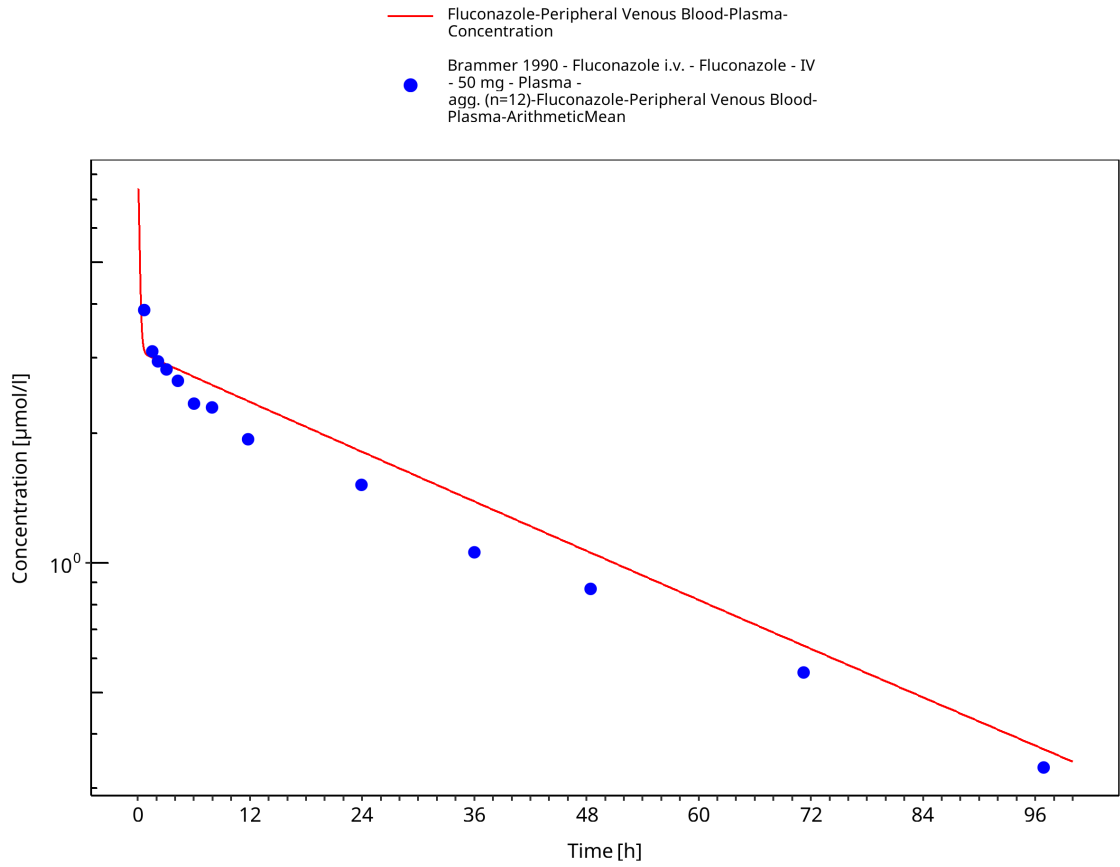


Figure 3-18: Time Profile Analysis

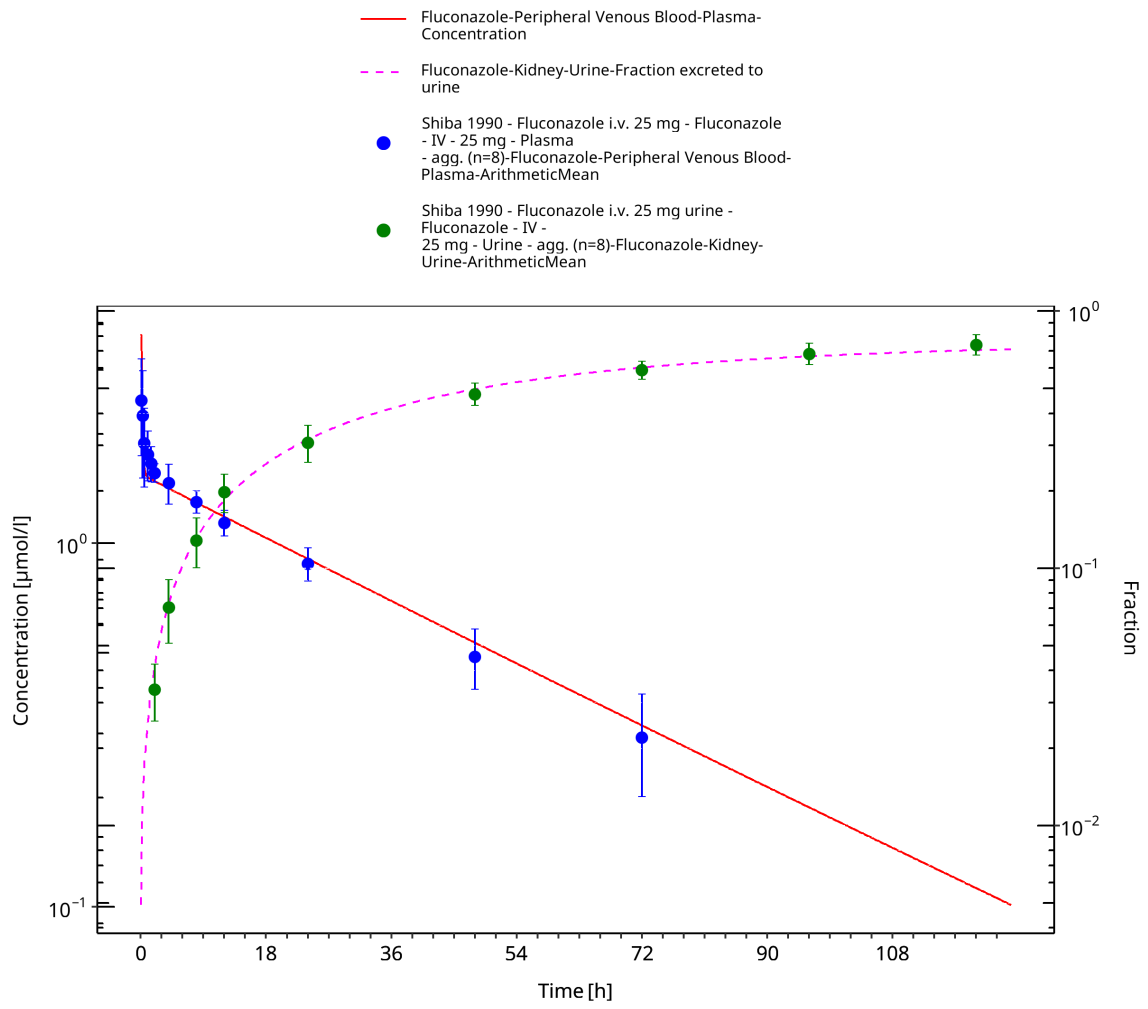


Figure 3-19: Time Profile Analysis

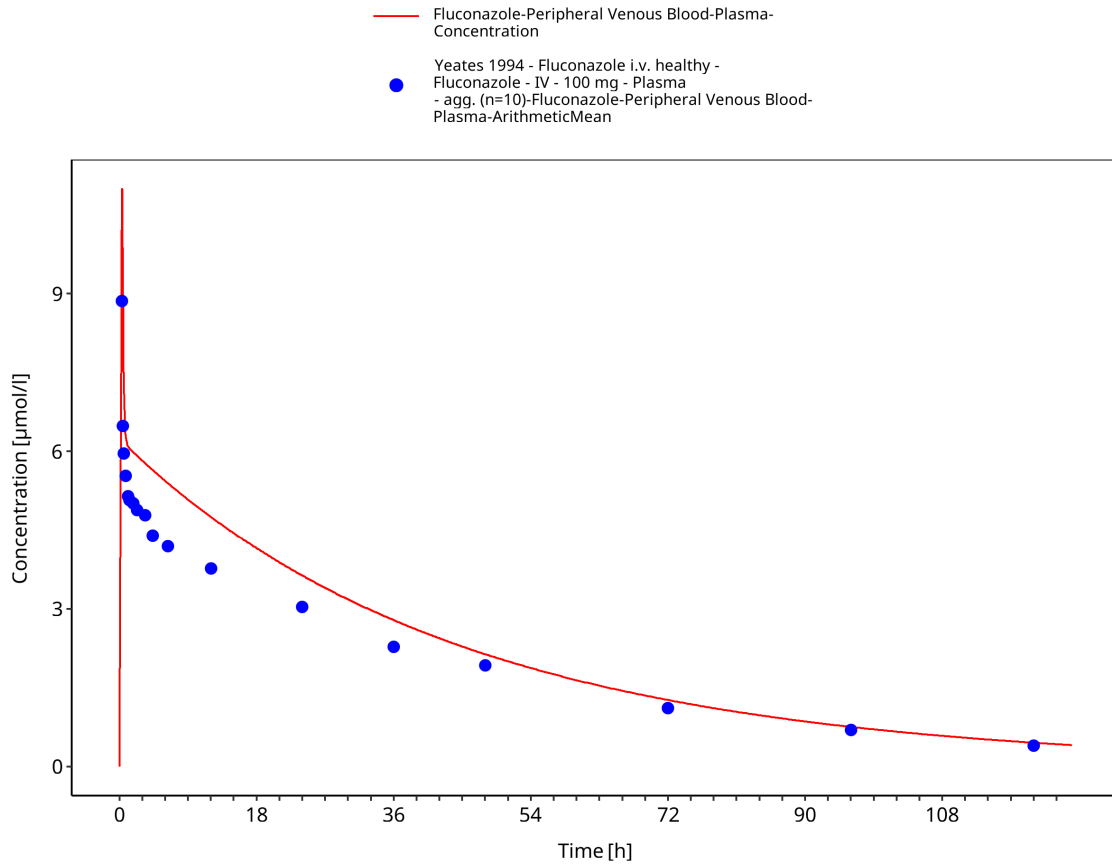


Figure 3-21: Time Profile Analysis

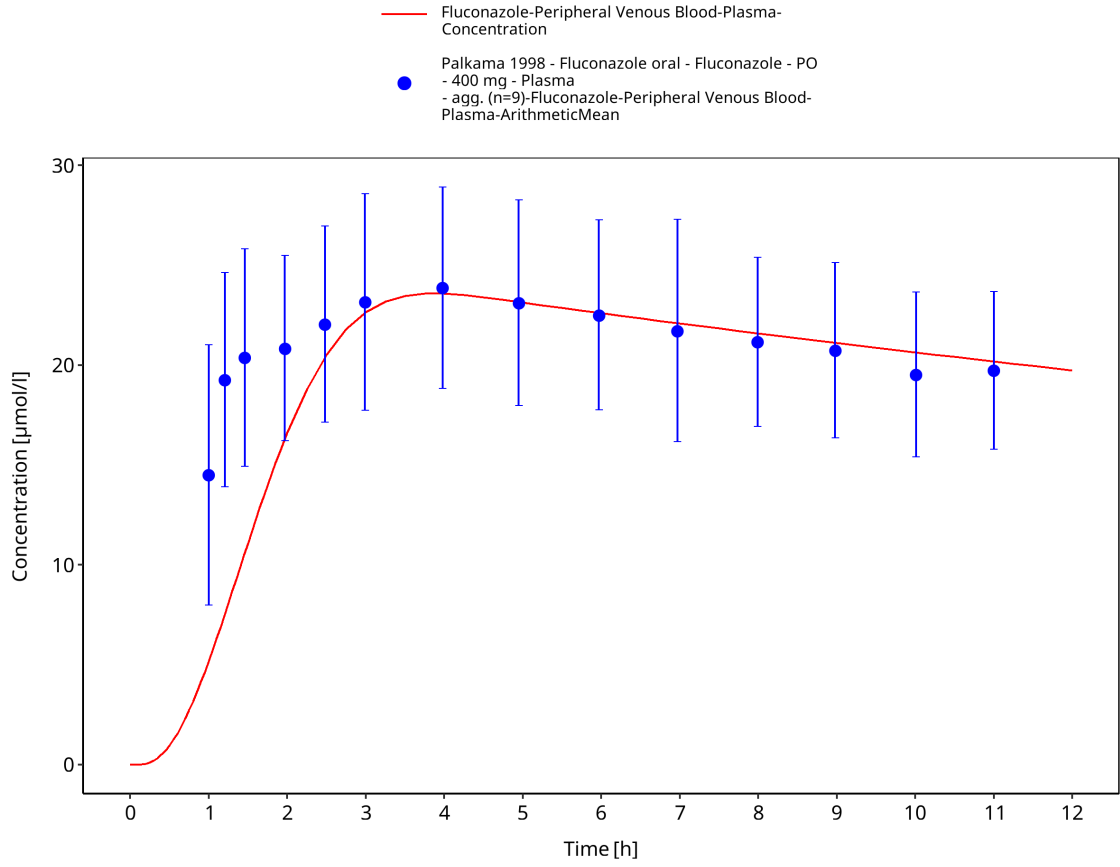


Figure 3-22: Time Profile Analysis

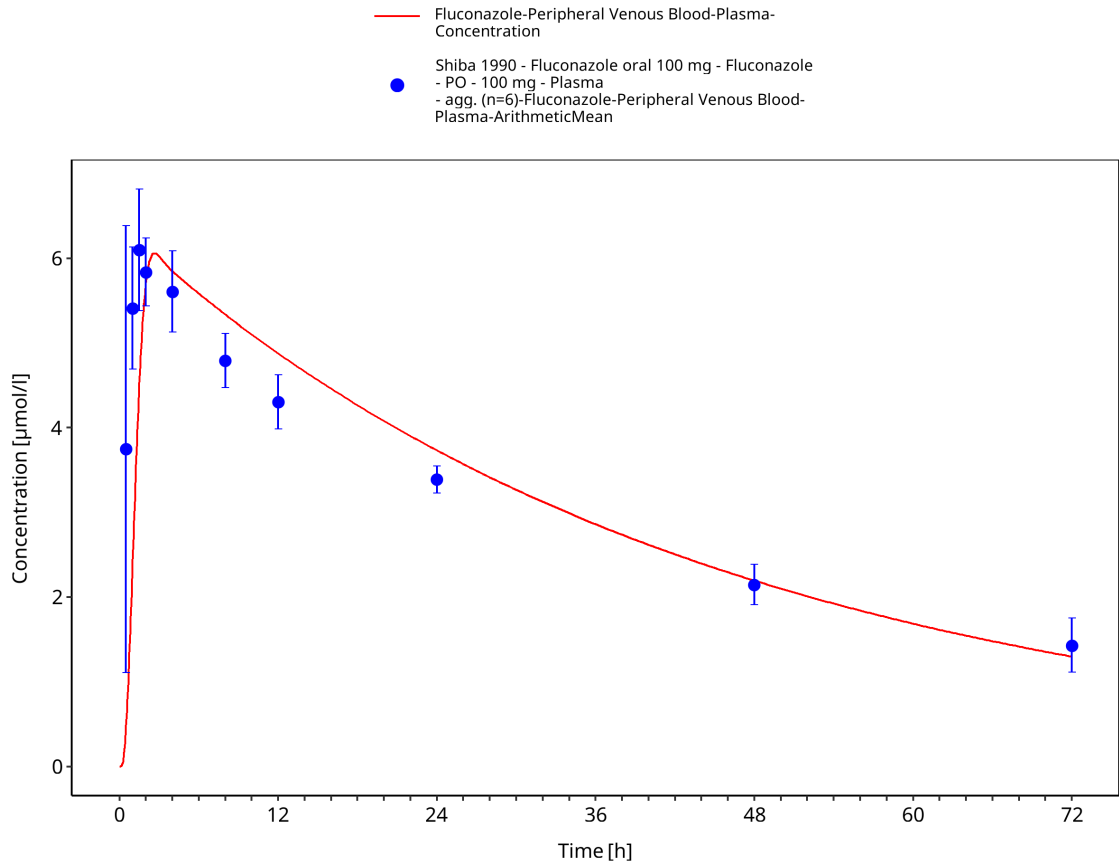


Figure 3-23: Time Profile Analysis

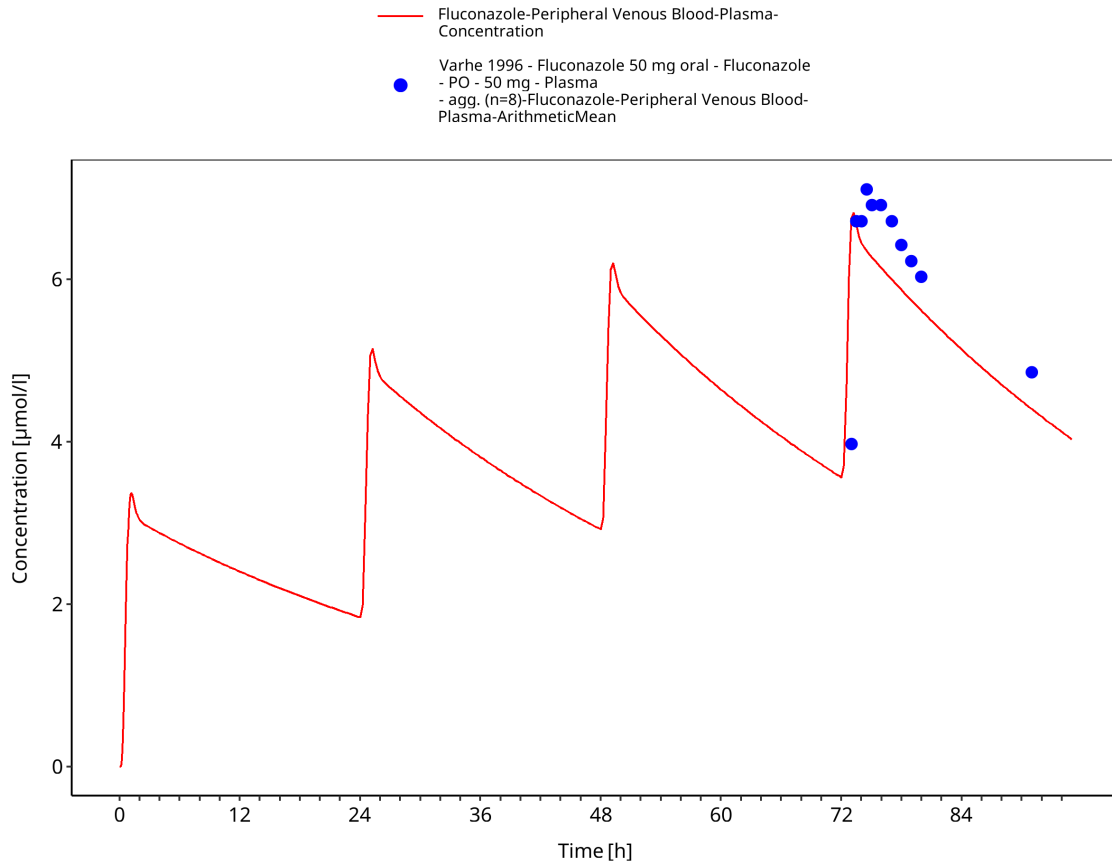


Figure 3-24: Time Profile Analysis

4 Conclusion

The presented PBPK model adequately describes the intravenous and oral pharmacokinetics of fluconazole in adults.

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