

# Building and evaluation of a PBPK model for raltegravir in adults

Version	2.0-OSP12.2
based on <i>Model Snapshot</i> and <i>Evaluation Plan</i>	<a href="https://github.com/Open-Systems-Pharmacology/Raltegravir-Model/releases/tag/v2.0">https://github.com/Open-Systems-Pharmacology/Raltegravir-Model/releases/tag/v2.0</a>
OSP Version	12.2
Qualification Framework Version	3.5

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

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The presented model building and evaluation report evaluates the performance of a PBPK model for raltegravir in adults.

Raltegravir, sold under the brand name Isentress, is an antiretroviral medication used to treat HIV/AIDS by blocking the establishment of post-integration HIV latency. It is also used as part of post exposure prophylaxis to prevent HIV infection following potential exposure. Raltegravir is only taken orally and is mainly metabolized by UGT1A1 (~70%) ([Kassahun 2007](#)). The final raltegravir model features metabolism by UGT1A1 and to a minor extent by UGT1A9. Additionally, there is excretion via glomerular filtration. The model adequately describes the pharmacokinetics of raltegravir in adults.

The raltegravir model is a whole-body PBPK model, allowing for dynamic translation between individuals with organs expressing UGT1A1. The raltegravir report demonstrates the level of confidence in the raltegravir PBPK model build with the OSP suite with regard to reliable predictions of raltegravir PK adults during model-informed drug development.

## 2 Methods

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### 2.1 Modeling strategy

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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 ([PK-Sim Ontogeny Database Version 7.3](#)). 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 ([Schlender 2016](#)) or otherwise referenced for the specific process.

First, a base mean model was built using data from the single dose escalation study to find an appropriate structure describing the PK of Raltegravir. The mean PK model was developed using a typical European individual. Unknown parameters were identified using the Parameter Identification module provided in PK-Sim®. Structural model selection was mainly guided by visual inspection of the resulting description of data and biological plausibility.

Once the appropriate structural model was identified, additional parameters for different formulations were identified.

A final PBPK model was established and simulations were compared to the reported data to evaluate model appropriateness and to assess model qualification, by means of diagnostics plots and predicted versus observed concentration-time profiles, of which the results support an adequate prediction of the PK in adults.

During model building, uncertainties in data quality, as well as study differences may cause not being able to adequately describe the PK of all reported clinical studies.

### 2.2 Data used

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#### 2.2.1 In vitro / physico-chemical data

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

Parameter	Unit	Raltegravir literature	Description
MW	g/mol	444.4163 ( <a href="#">drugbank.ca</a> )	Molecular weight
pKa		6.67 ( <a href="#">Moss 2012</a> )	Acid dissociation constant
Solubility (pH)	mg/L	Reference pH-dependent table ( <a href="#">Moss 2013</a> )	Solubility
logD (pH 7)		0.58 ( <a href="#">Moss 2012</a> )	Partition coefficient between octanol and water
fu		0.17 ( <a href="#">Laufer 2009</a> )	Fraction unbound
Km UGT1A1	μM	99 ( <a href="#">Kassahun 2007</a> )	Michaelis-Menten constant
Vmax UGT1A1	nmol/min/mg	0.89 ( <a href="#">Kassahun 2007</a> )	Maximum rate of reaction
Km UGT1A9	μM	296 ( <a href="#">Kassahun 2007</a> )	Michaelis-Menten constant
Vmax UGT1A9	nmol/min/mg	0.53 ( <a href="#">Kassahun 2007</a> )	Maximum rate of reaction

## 2.2.2 Clinical data

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

The following publications were found in adults for model building and evaluation:

Publication	Study description
<a href="#">Iwamoto 2008</a>	Single- and multiple-dose escalation study in healthy subjects
<a href="#">Iwamoto 2009</a>	Effects of ritonavir and efavirenz on safety, tolerability and pharmacokinetics of raltegravir
<a href="#">Markowitz 2006</a>	Monotherapy, followed by a longer term combination therapy of raltegravir versus efavirenz
<a href="#">Kassahun 2007</a>	Pharmacokinetics study in healthy adults
<a href="#">Rhee 2014</a>	Pediatric formulation study in healthy adults
<a href="#">Wenning 2009</a>	Effect of rifampin on the pharmacokinetics of raltegravir

## 2.3 Model parameters and assumptions

### 2.3.1 Absorption

As no intravenous data is currently available to study systemic clearance of raltegravir *in vivo*, only oral data was used for model building. For oral administration the following parameters play a role with regards to the absorption kinetics of a compound, which can be estimated with PBPK: solubility, lipophilicity and intestinal permeability. Moss et al. ([Moss 2013](#)) published values for raltegravir solubility in population groups with very low-, low-, medium-, high-, and very high intestinal luminal pH, after a single 400 mg dose of raltegravir. For the raltegravir PBPK model we have applied the medium pH group for creating a pH dependent solubility profile throughout the intestinal tract. The lipophilicity as well as pKa of raltegravir was also published by Moss et al ([Moss 2012](#), [Moss 2013](#)) to be 0.58 (as log partition coefficient between

octanol and water (pH 7) and 6.67 (acid)), respectively. These values were applied and fixed in the raltegravir PBPK model, without further optimization. Regarding intestinal transcellular permeability (Pint), Moss et al ([Moss 2012](#)) reported a range of apical to basolateral apparent permeability in Caco-2 monolayer at different pH values. Using published functions Pint can be calculated from Caco-2 cell membrane permeability measurements (Parrot et al. ([Parrot 2002](#))), Thelen et al. ([Thelen 2010](#), Sun et al. ([Sun 2002](#)) and Sjögren et al. ([Sjögren 2013](#))). However as no reference/calibrator compound was available to correct for inter-study variability, these functions could not be applied, and it was decided to estimate the Pint from *in vivo* clinical data instead. Nevertheless, for plausibility check, a theoretical Pint was calculated using the aforementioned functions without correction, resulting in a range of Pint from 4.64E-04 to 1.47E-09 cm/min. The finally estimated (based on *in vivo* data) Pint falls within this range.

**Table 2.** Reported Caco-permeability and calculated theoretical effective permeability (intestinal transcellular permeability, Peff) values for raltegravir via different reported functions, lacking a reference compound for correcting inter-study variability.

Reference publication of reported function	pH apical to basolateral	Peff apical to basolateral (cm/min)	Reference compound available for correcting Inter study variability
Raltegravir Caco permeability (Moss 2012 )	7.4	6.60E-6	-
Raltegravir Caco permeability (Moss 2012)	6.5	9.20E-6	-
Parrot 2002	7.4	2.14E-04	Not available
Thelen 2010	7.4	1.47E-09	Not available
Sjögren 2013	7.4	1.03606E-6	Not available
Sun 2002	7.4	2.77E-4	Not available
Sun 2002	6.5	2.86E-4	Not available
Simcyp (*)	6.5	4.62E-4	Not available

\*Not published as paper, Simcyp applied an adapted version of Sun et al 2002

## 2.3.2 Distribution

Laufer et al. ([Laufer 2009](#)) published a fu in humans to be 0.17. [Barau et al 2013](#) reported that raltegravir binds to serum albumin, and not alpha glycoprotein, which is built-in as such in the PBPK model.

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 cell permeability calculation by PK-Sim standard. Specific organ permeability normalized to surface area was automatically calculated by PK-Sim.

## 2.3.3 Metabolism and Elimination

Kassahun et al. ([Kassahun 2007](#)) studied the absorption, metabolism, and excretion of raltegravir in healthy volunteers after a single oral dose of 200 mg (200Ci) of [14C] raltegravir. Human liver microsomal incubations confirmed the dominant role of UGT metabolism for raltegravir. Additionally, data from incubations using cDNA-expressed UGTs indicate that the major mechanism of metabolism of raltegravir in humans is UGT1A1-mediated glucuronidation. Raltegravir was in

particular converted by UGT1A1 and 1A9. The apparent arithmetic mean  $K_m$  values for the glucuronidation of raltegravir by UGT1A1 and UGT1A9 were 99 (standard deviation (SD): 16) and 296 (SD: 55)  $\mu\text{M}$ , respectively. The corresponding  $V_{\text{max}}$  values (nmol/min/mg) were 0.89 (SD: 0.05) for UGT1A1, and 0.53 (SD: 0.06) for UGT1A9.

Based on this information, the reported *in vitro*  $K_m$  values for UGT1A1 and 1A9 were included in the model. Reported  $V_{\text{max}}$  values were in units nmol/min/mg protein and thus not directly transferable into the PBPK model. Therefore, a joint scaling factor  $f_{\text{UGT}}$  on the *in vitro*  $V_{\text{max}}$  values was estimated to match observed *in vivo* data, and keeping the relative relationship between those *in vitro* values (0.89 and 0.53 nmol/min/mg) for UGT1A1 and UGT1A9 fixed according to:

$$V_{\text{max,UGT1A1}} = f_{\text{UGT}} * V_{\text{max,in-vitro,UGT1A1}}$$

$$V_{\text{max,UGT1A9}} = f_{\text{UGT}} * V_{\text{max,in-vitro,UGT1A9}}$$

It is especially important to fix the relative contribution of both enzymes as a ratio to ensure that, when translating to other populations (e.g. children where both UGTs undergo a different ontogeny pattern, or patients who have differently reduced amounts of UGT1A1 vs 1A9) the relative contributions can be adequately scaled. Note that the estimated scaling factor  $f_{\text{UGT}}$  will be directly implemented into the final *in vivo*  $V_{\text{max}}$  values (only  $V_{\text{max,UGT1A1}}$  and  $V_{\text{max,UGT1A9}}$  will be reported in [section 3](#))

Finally, as ~9% of the dose is excreted in human urine as unchanged parent compound, GFR is introduced in the raltegravir PBPK model.

## 3 Results and Discussion

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The PBPK model **raltegravir** was developed with clinical pharmacokinetic data covering 4 different oral formulation and a dose range of 10-1600mg, including single dose (SD) as well as multiple dose (MD) clinical data.

As there were 4 different oral formulations available for model evaluation, all formulations require an estimation of the dissolution kinetics via a Weibull function. This function requires the estimation of 2 parameters, the dissolution time (time where 50% of the drug is dissolved), and dissolution shape (shape parameter of the Weibull function). Therefore, to minimize the amount of parameters for fitting, as a first step, the PK study data (lactose formulation) by Iwamoto et al. ([Iwamoto 2007](#)) was fitted which includes SD escalation and has a broad dose-range (10mg-1600mg) to capture dose (non-) linearity. During the model-fitting, the following parameters were estimated (all other parameters were fixed to reported values):

- $V_{\max}$  (as unique scaling factor  $f_{\text{UGT}}$ , as described in [section 2.3.3](#))
- Weibull function parameters: Dissolution time and dissolution shape
- Specific intestinal permeability (transcellular)

The fit resulted in an adequate description of all data. As there is no iv data available, it was not possible to clearly distinguish between clearance and absorption, resulting in a considerable correlation between  $V_{\max}$  and dissolution shape (Weibull). An attempt to fix  $V_{\max}$  to reported in vitro values, and only estimating absorption (lipophilicity and intestinal transcellular permeability) resulted in an underprediction of the clearance, and clearly indicated a need for increase in clearance. As described above, no reported intestinal permeability was found other than Caco2-permeability. Caco2-permeability could not be translated to effective intestinal permeability without a reference compound. Therefore it was decided to continue with the model where both  $P_{\text{int}}$  and  $V_{\max}$  were estimated.

As a second step, clinical study data for all other formulations summarised in [section 2.2.2](#) were included for model fitting, including film-coated tablets (100-400mg MD, 200-400mg SD), chewable tablets (400mg fasted + fed) and oral granules in suspension (400mg). In this step, only the Weibull functions were estimated with all other parameters fixed based on the first step. Finally, as the parameters of the Weibull functions were highly correlated (as expected), only dissolution shape was estimated as a last step. The model results show that the PBPK model of raltegravir adequately described the data for all formulations and doses available.

### 3.1 Raltegravir final input parameters

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

#### Compound: Raltegravir

##### Parameters



Name	Value	Value Origin	Alternative	Default
Solubility table	40 mg/l	Publication-In Vitro-Moss 2013 Table 2	Moss 2013	True
Lipophilicity	0.58 Log Units	Publication-Moss 2012	Moss 2012	True
Fraction unbound (plasma, reference value)	0.17	Publication-In Vitro-Laufer 2009	Measurement	True
Specific intestinal permeability (transcellular)	2.8481843854E-07 cm/s	Parameter Identification	Fit	True
F	1	Publication-Other-Drugbank.ca		
Is small molecule	Yes			
Molecular weight	444.4163 g/mol	Publication-Other-Drugbank.ca		
Plasma protein binding partner	Albumin			

## Calculation methods

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

## Processes

### Systemic Process: Glomerular Filtration-Kassahun 2007

Species: Human

#### Parameters

Name	Value	Value Origin
GFR fraction	1	Publication-In Vitro-Kassahun 2007

### Metabolizing Enzyme: UGT1A1-Kassahun 2007

Molecule: UGT1A1

#### Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	2.7351231632 nmol/min/mg mic. protein	Parameter Identification
Km	99 µM	Publication-In Vitro-Kassahun 2007

Metabolizing Enzyme: UGT1A9-Kassahun 2007

Molecule: UGT1A9

Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	1.6287812095 nmol/min/mg mic. protein	Parameter Identification
Km	296 µM	Publication-In Vitro-Kassahun 2007

Formulation: filmcoated tablet (original Merck formulation)

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	500 min	Parameter Identification
Lag time	0 min	
Dissolution shape	0.03536656	Parameter Identification
Use as suspension	Yes	

Formulation: chewable tablet

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	1.0000049774E-05 min	Parameter Identification
Lag time	0 min	
Dissolution shape	0.050078869	Parameter Identification
Use as suspension	Yes	

Formulation: Weibull (lactose formulation)

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	2.30152527E-10 min	Parameter Identification
Lag time	0 min	
Dissolution shape	0.0389537131	Parameter Identification
Use as suspension	Yes	

Formulation: Weibull (granules)

Type: Weibull

Parameters

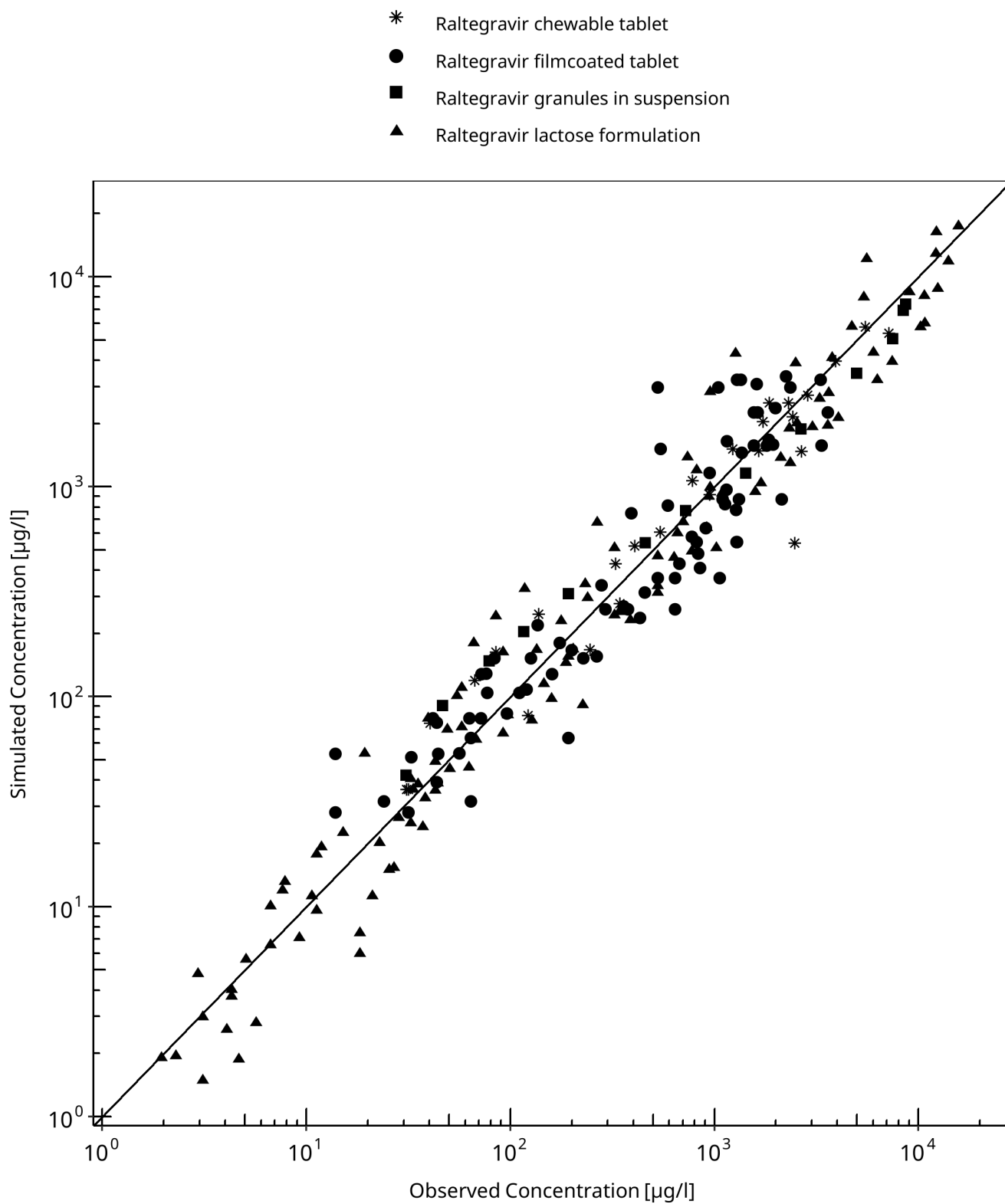
Name	Value	Value Origin
Dissolution time (50% dissolved)	0.00010000047426 min	Parameter Identification
Lag time	0 min	
Dissolution shape	0.0654456264	Parameter Identification
Use as suspension	Yes	

3.2 Raltegravir Diagnostics Plots

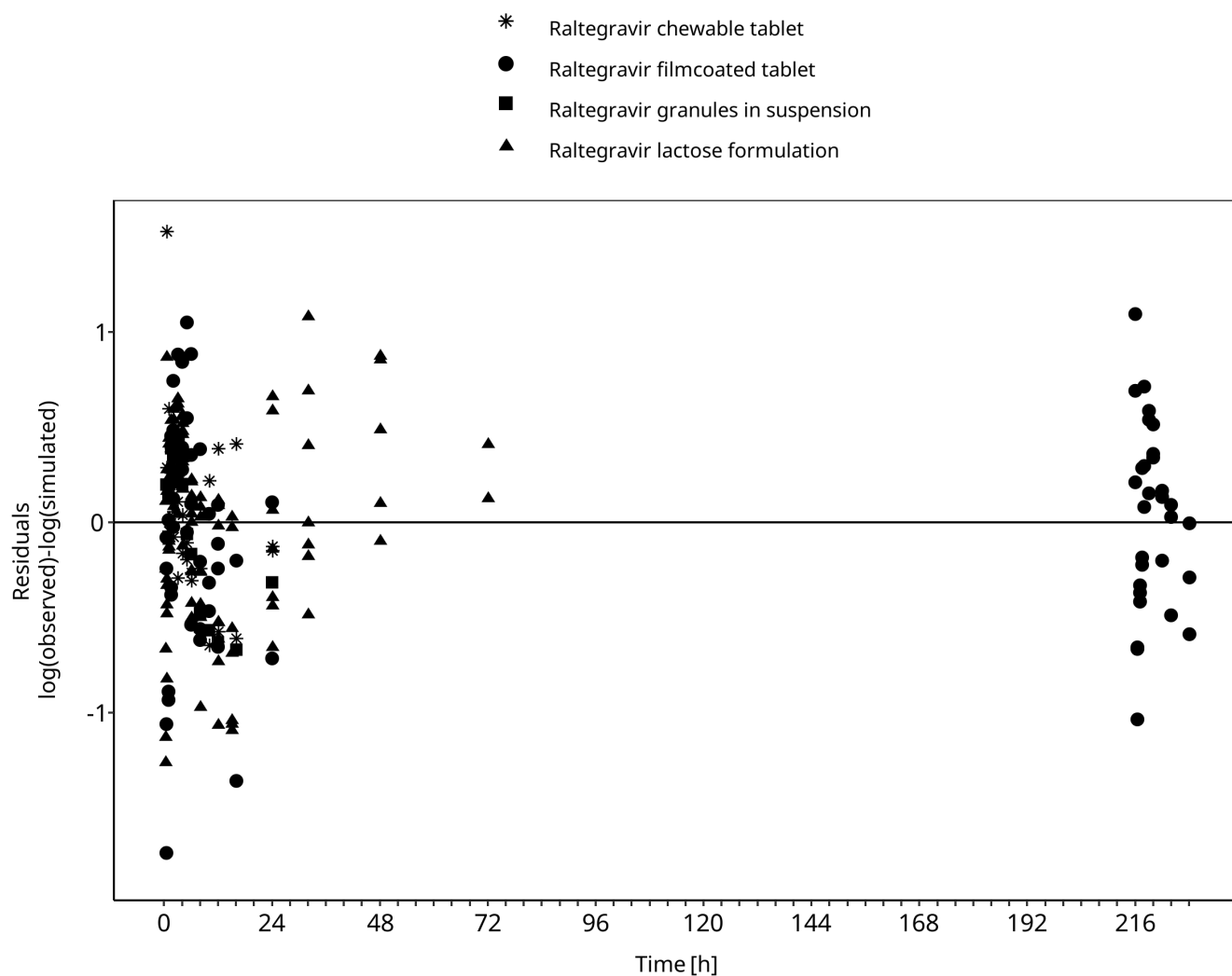
Below you find the goodness-of-fit visual diagnostic plots for raltegravir PBPK model performance (observed versus individually simulated plasma concentration and weighted residuals versus time, including the geometric mean fold error (GMFE)) of all data used for model building.

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

Group	GMFE
Raltegravir chewable tablet	1.37
Raltegravir filmcoated tablet	1.56
Raltegravir granules in suspension	1.42
Raltegravir lactose formulation	1.48
All	1.49



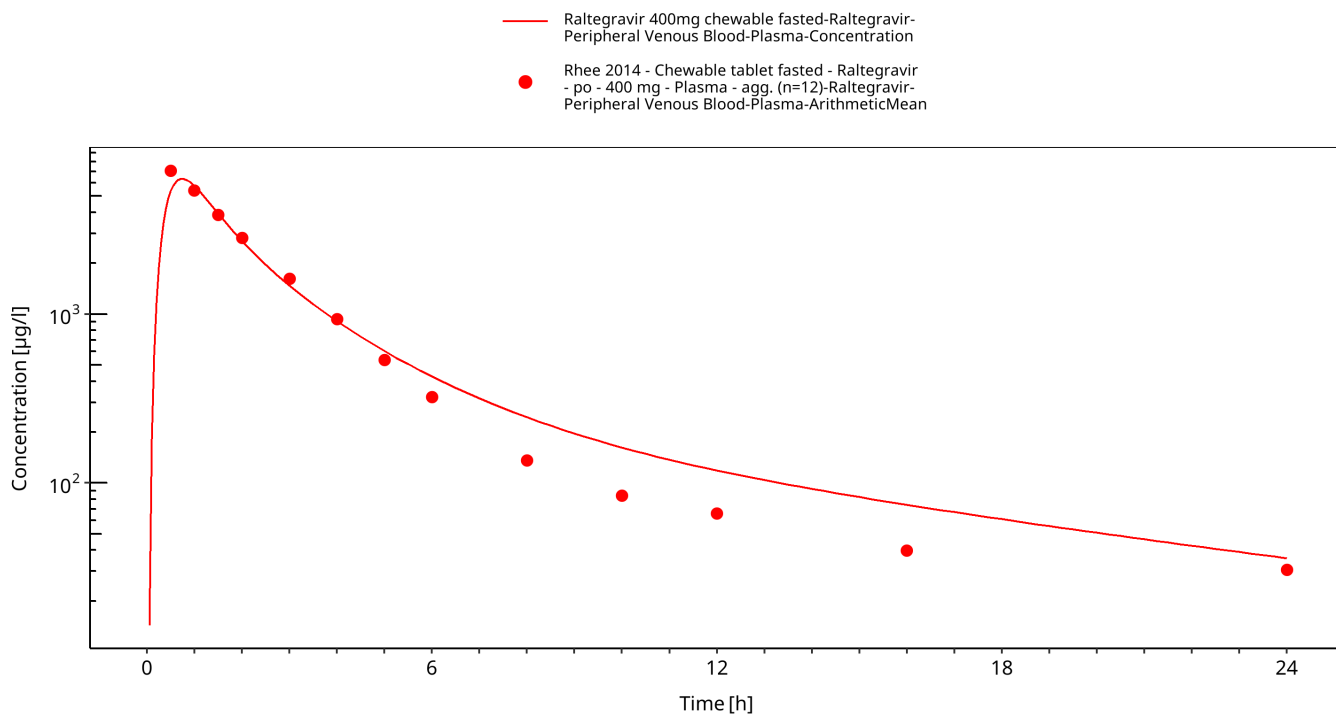
**Figure 3-1: Goodness of fit plot for concentration in plasma.**



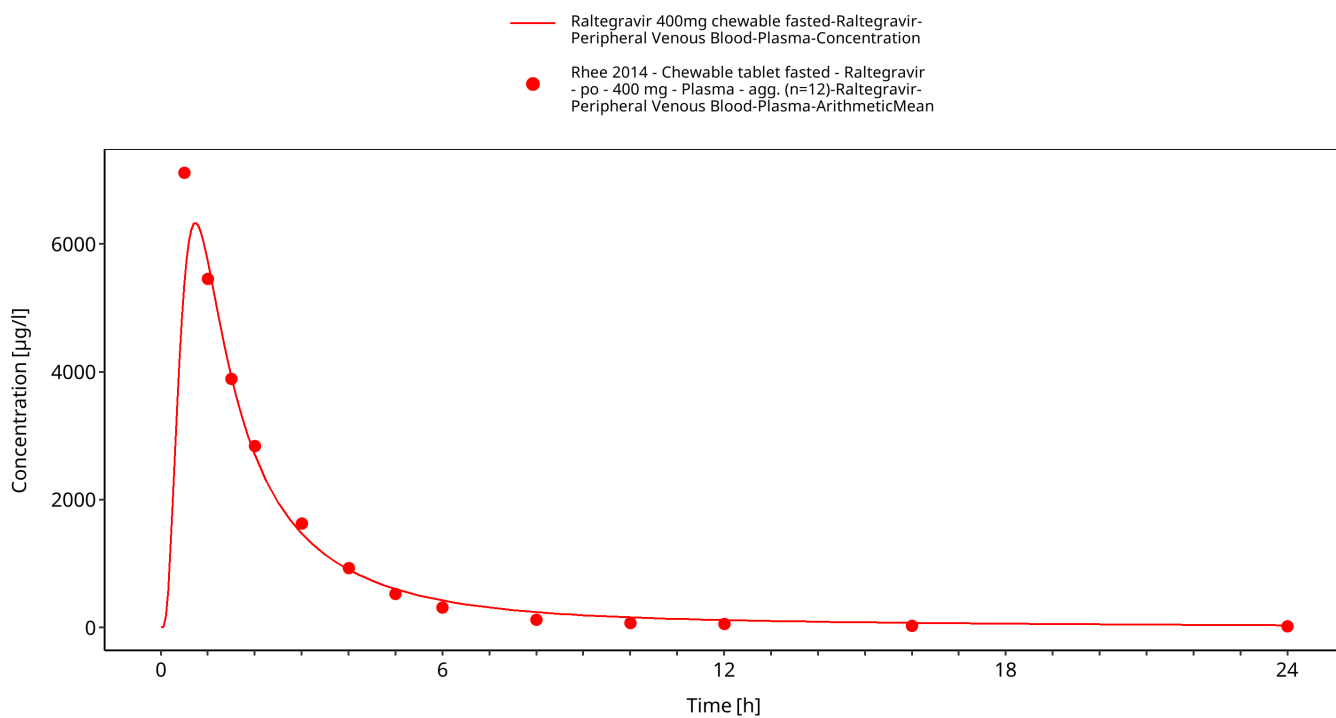
**Figure 3-2: Goodness of fit plot for concentration in plasma.**

### 3.3 Raltegravir Concentration-Time profiles

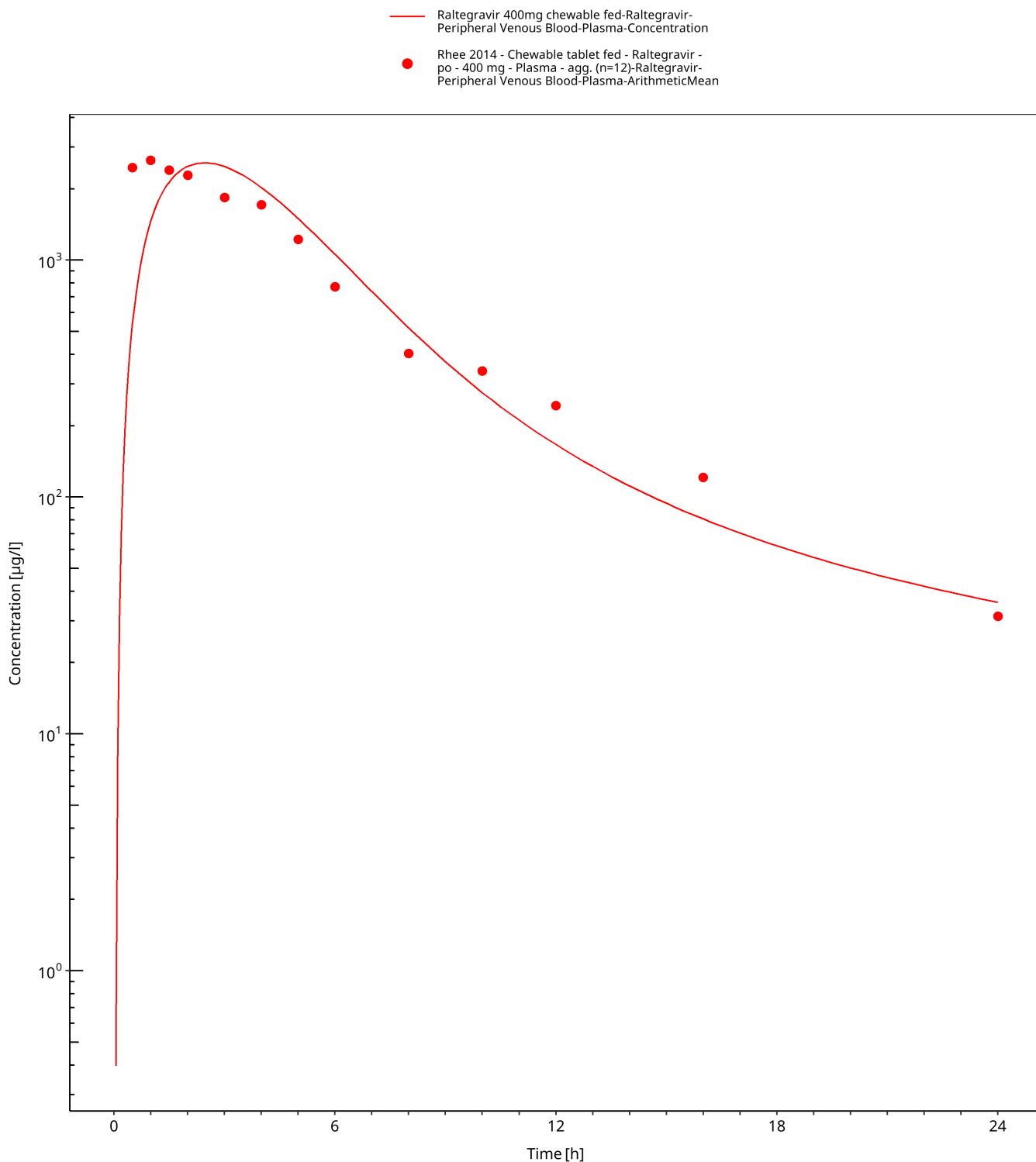
Simulated versus observed plasma concentration-time profiles of all data are listed below.



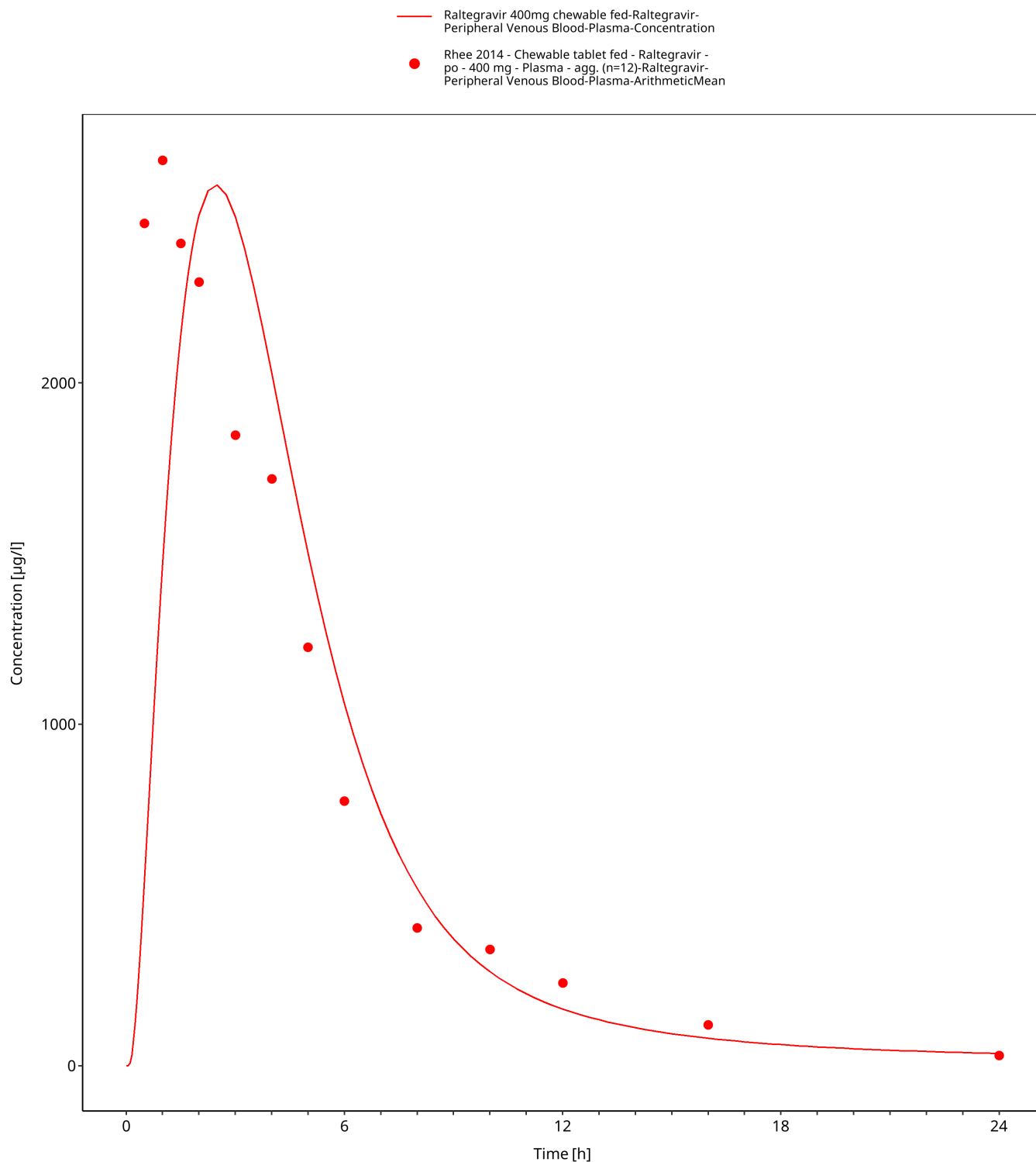
**Figure 3-3: Time Profile Analysis**



**Figure 3-4: Time Profile Analysis 1**

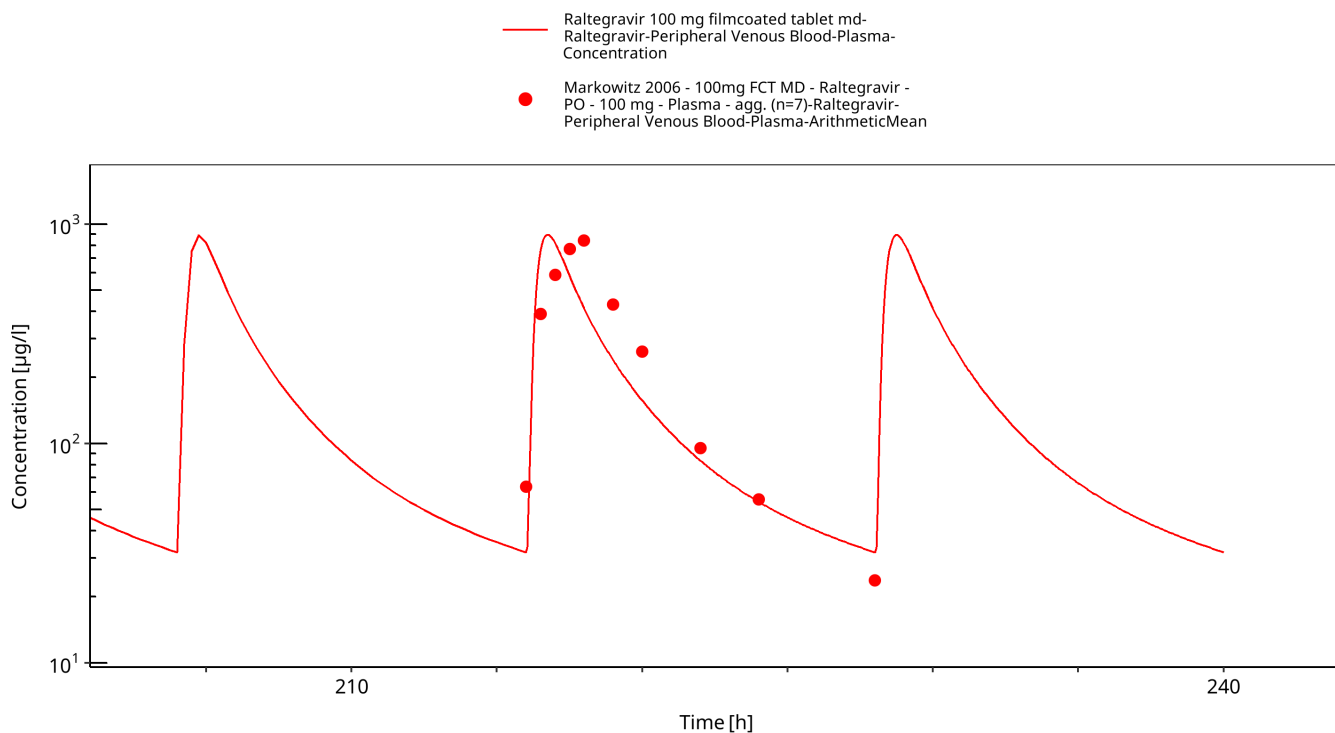


**Figure 3-5: Time Profile Analysis**

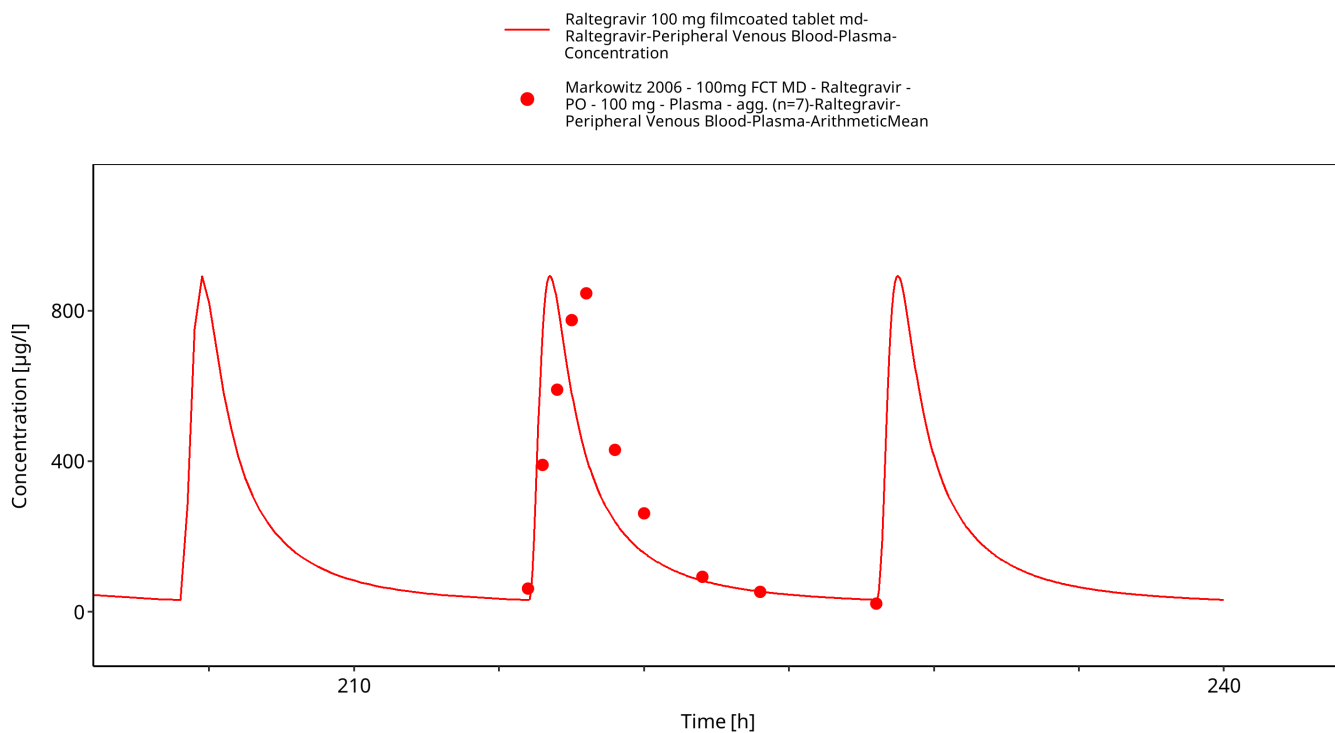


**Figure 3-6: Time Profile Analysis 1**

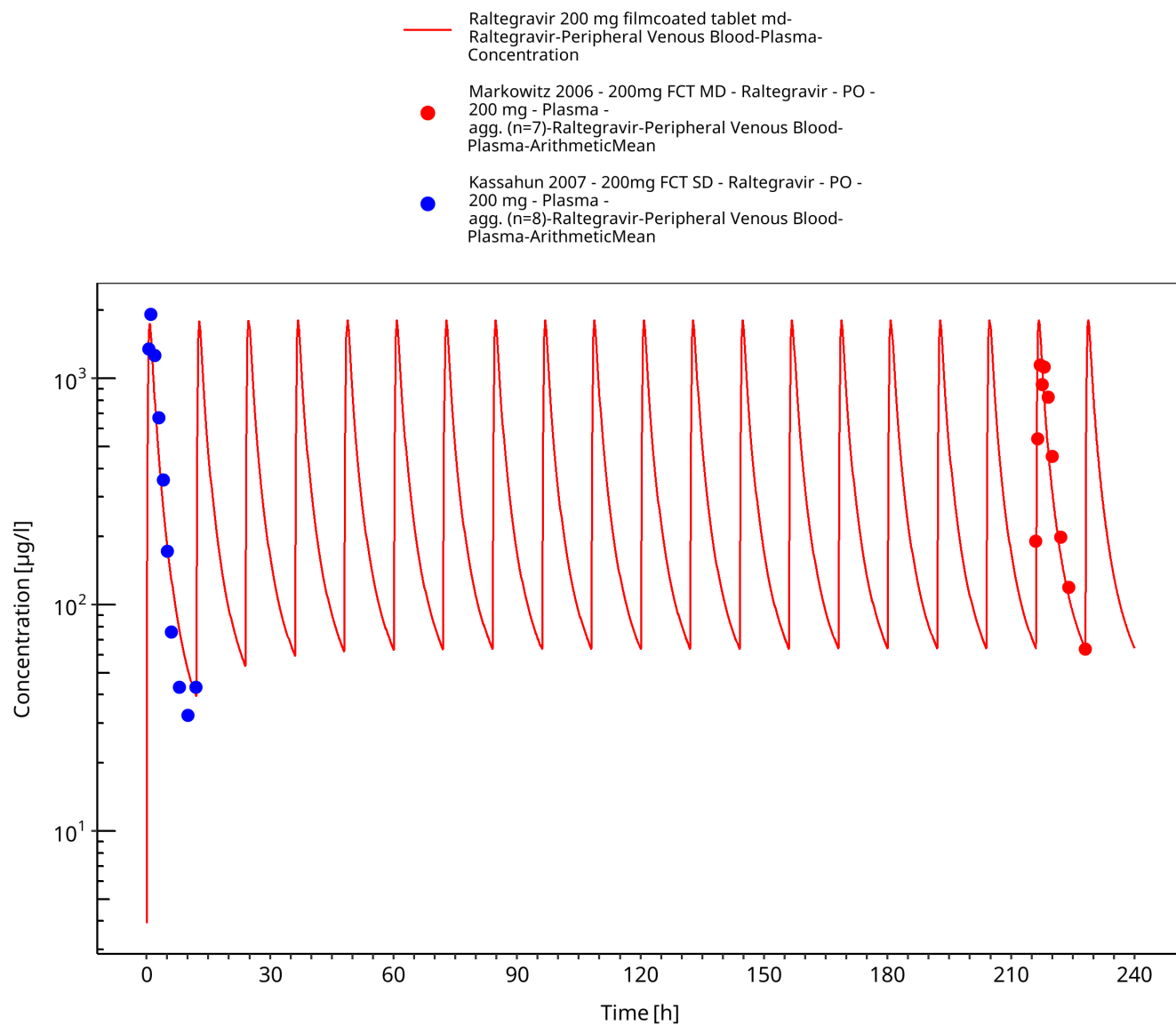




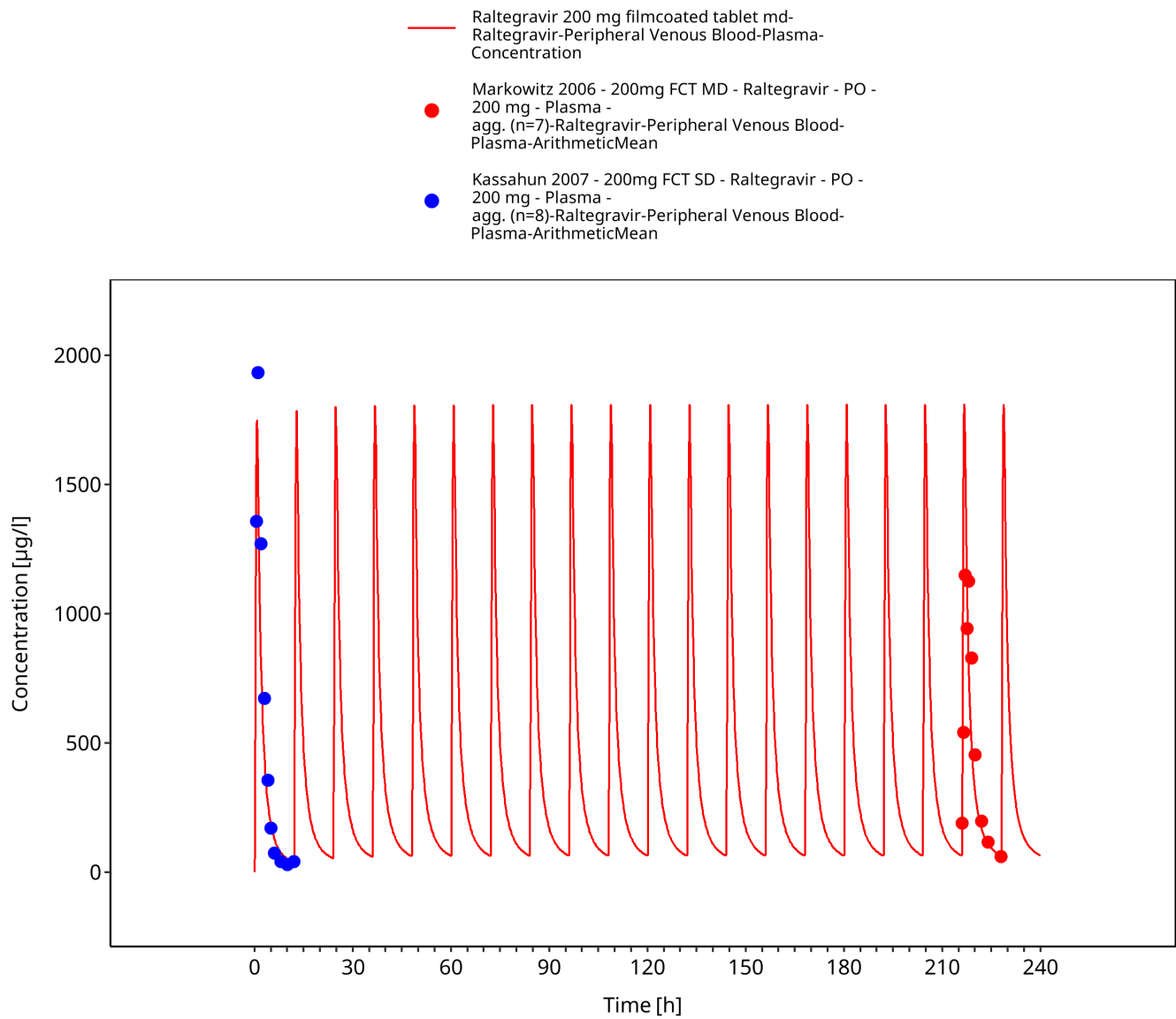
**Figure 3-7: Time Profile Analysis**



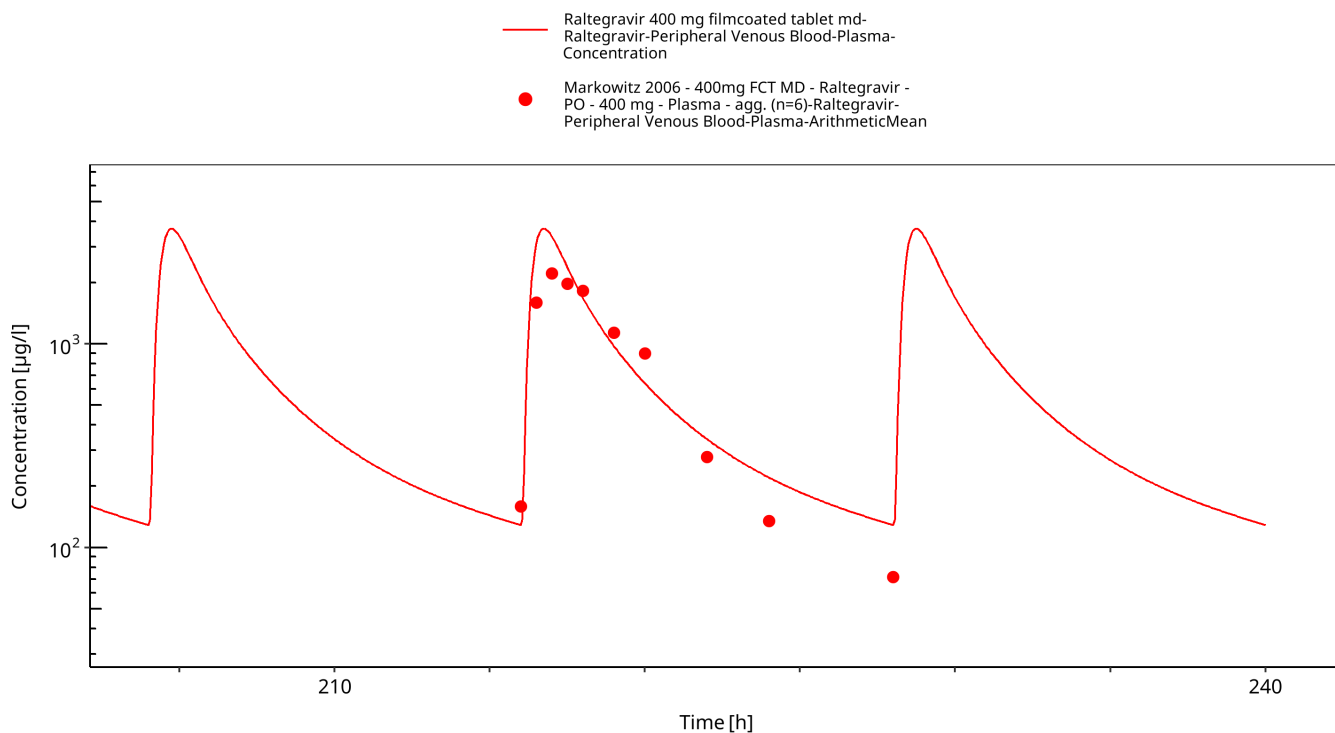
**Figure 3-8: Time Profile Analysis 1**



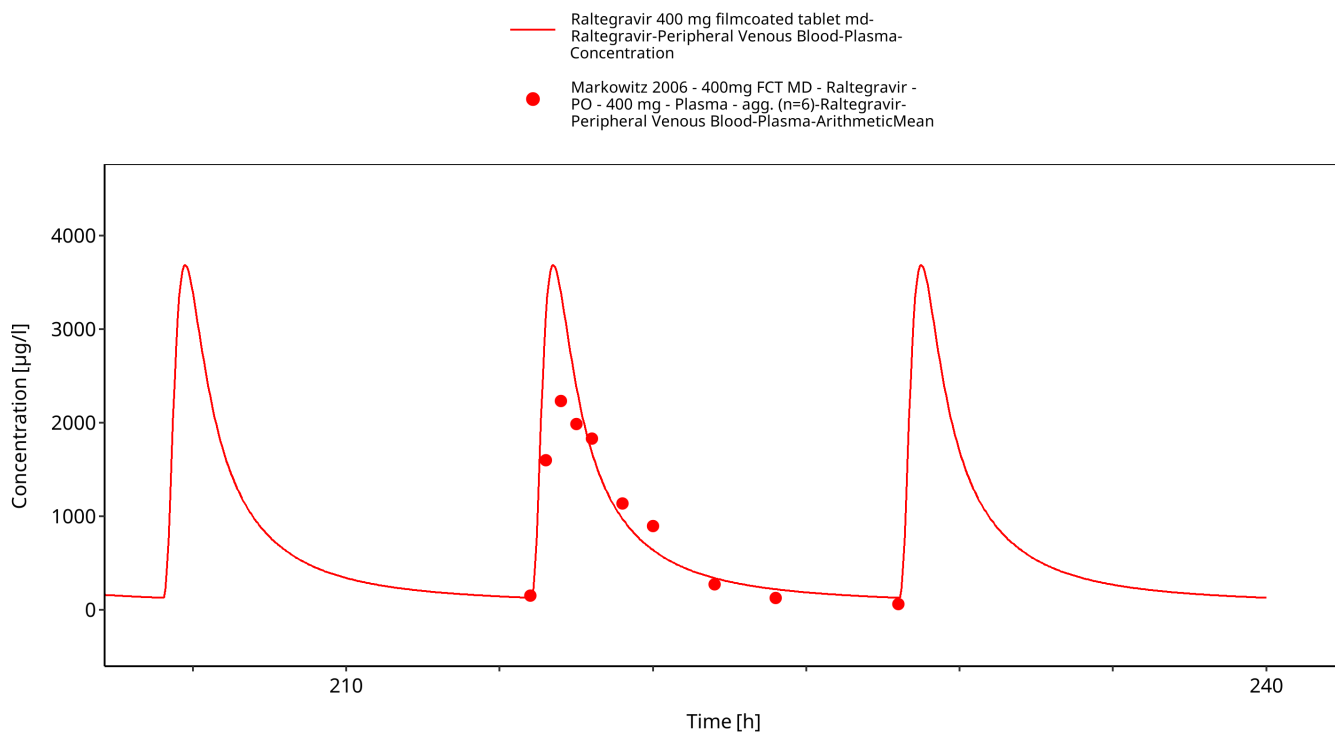
**Figure 3-9: Time Profile Analysis**



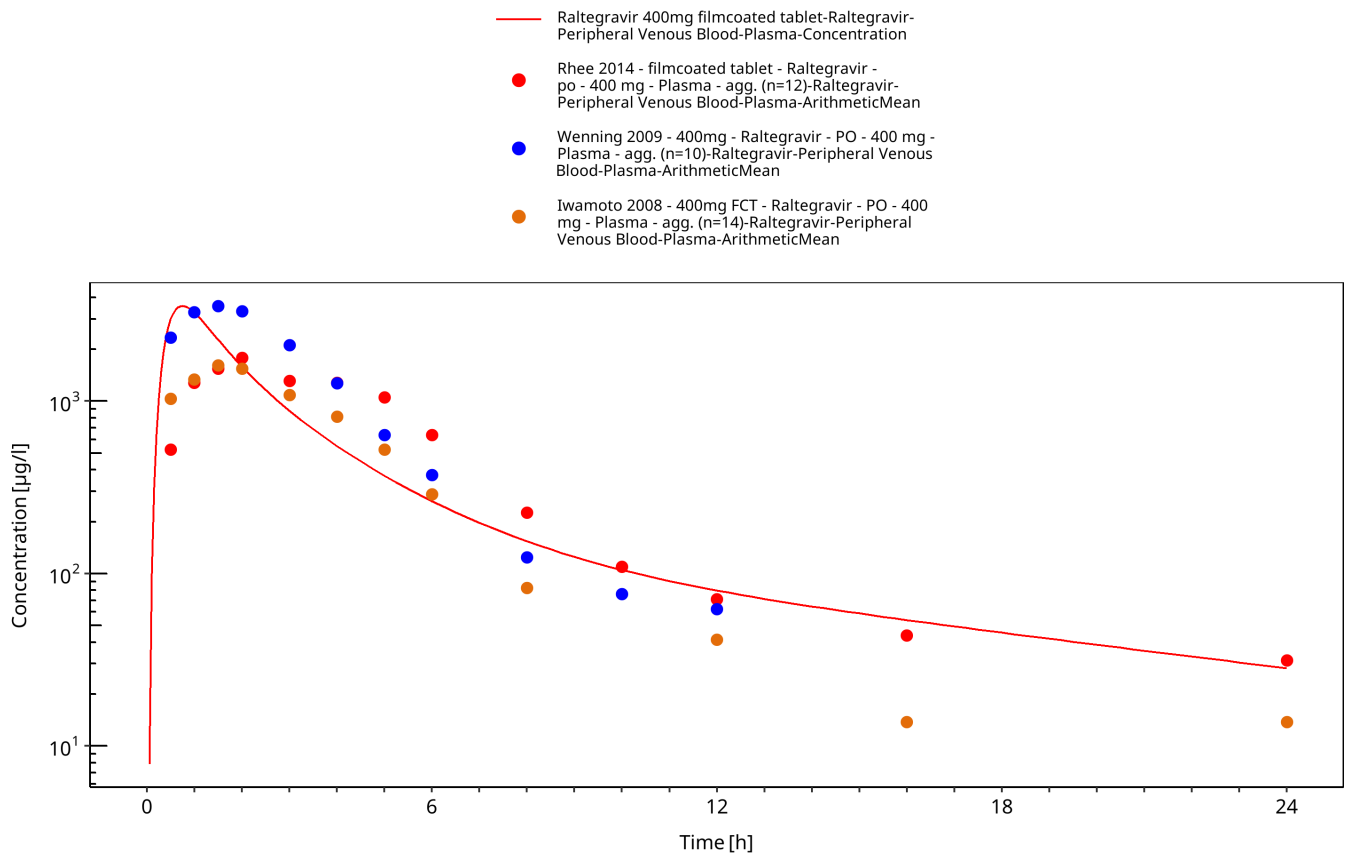
**Figure 3-10: Time Profile Analysis 1**



**Figure 3-11: Time Profile Analysis**



**Figure 3-12: Time Profile Analysis 1**



**Figure 3-13: Time Profile Analysis**

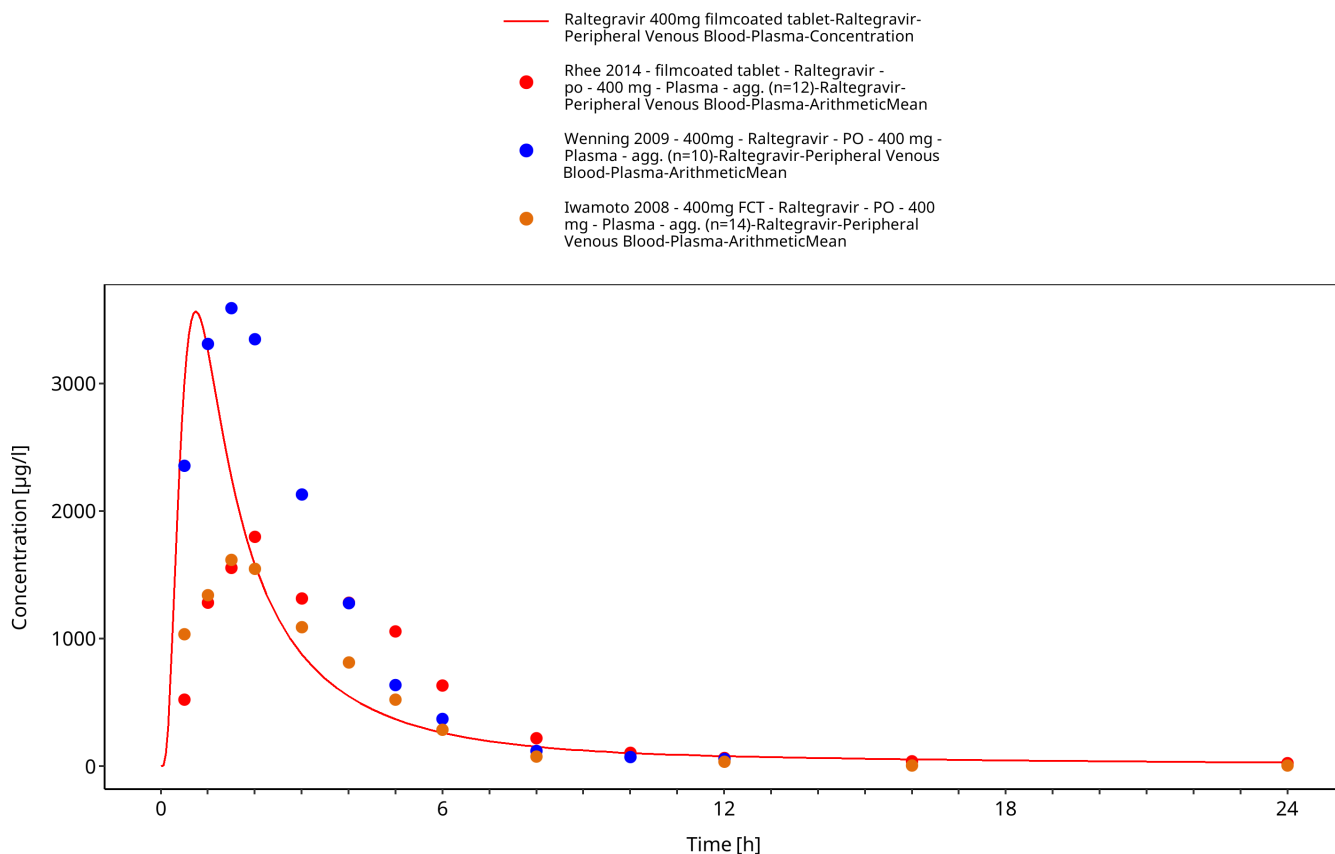


Figure 3-14: Time Profile Analysis 1

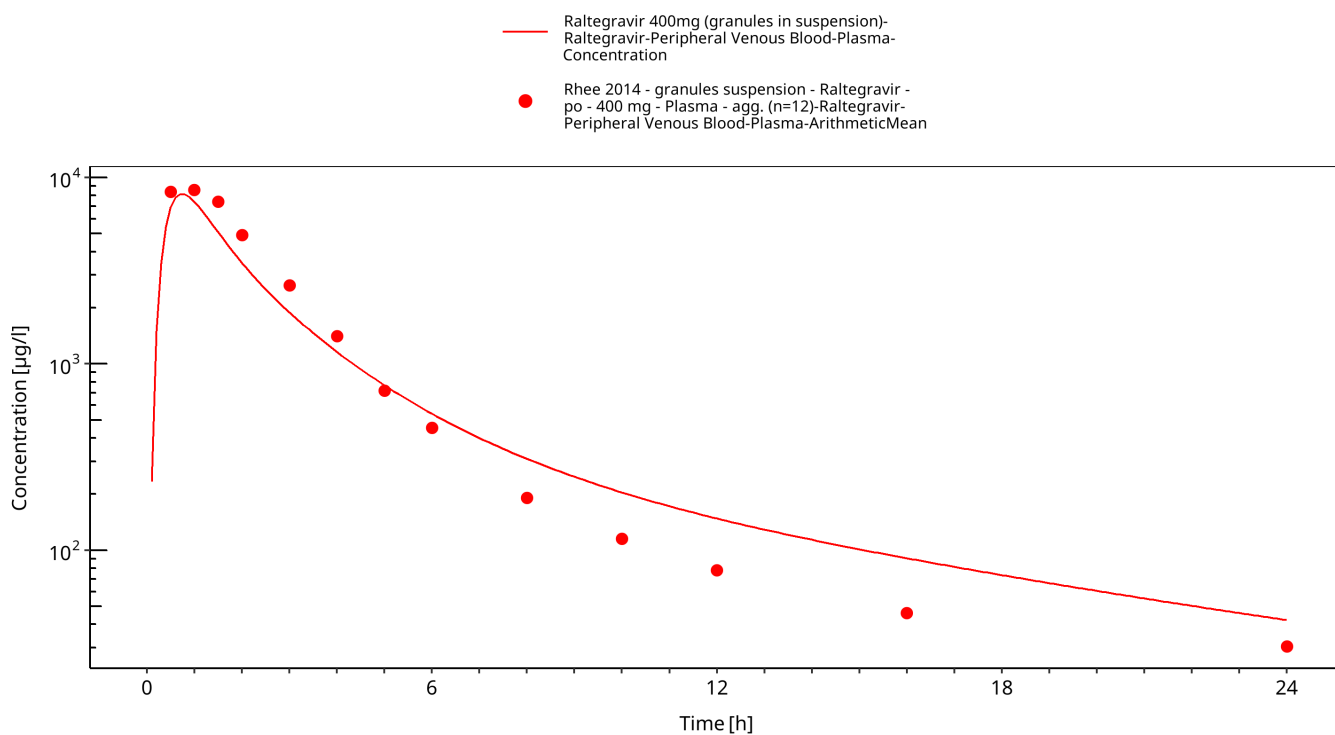


Figure 3-15: Time Profile Analysis

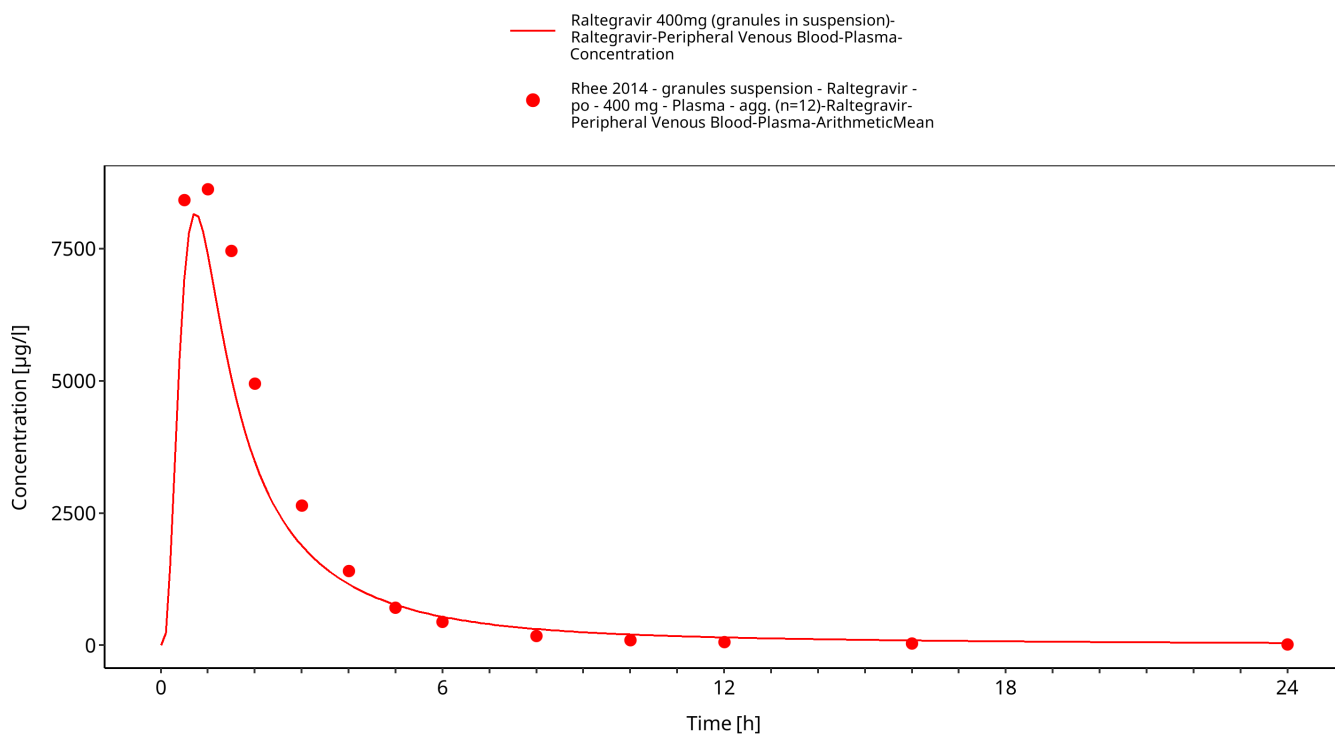


Figure 3-16: Time Profile Analysis 1

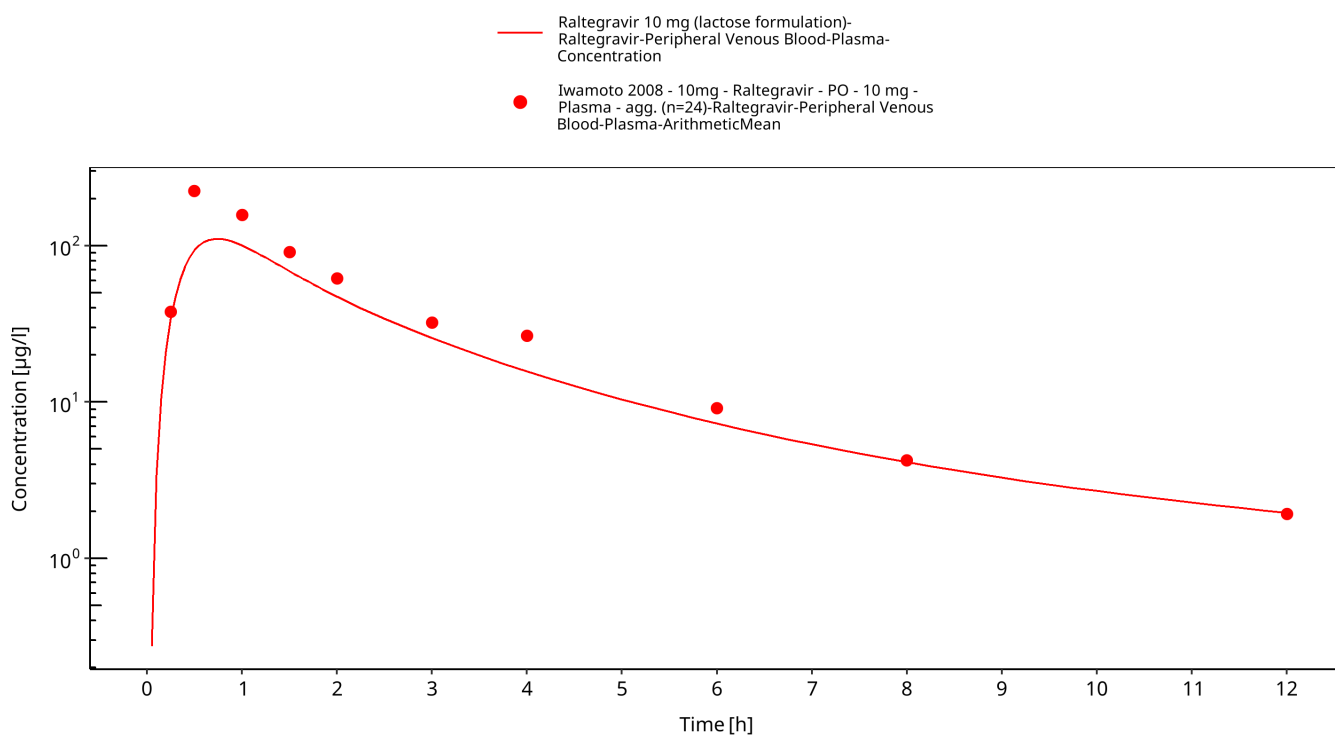
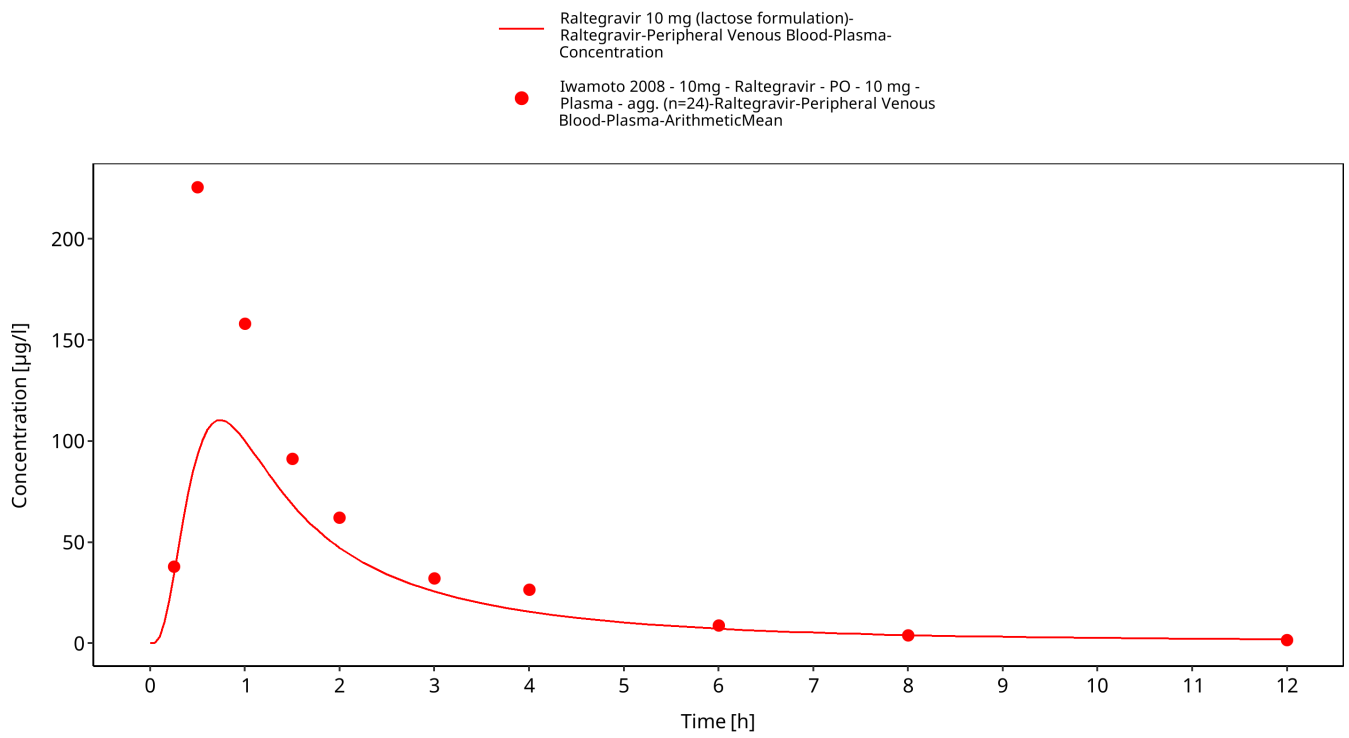
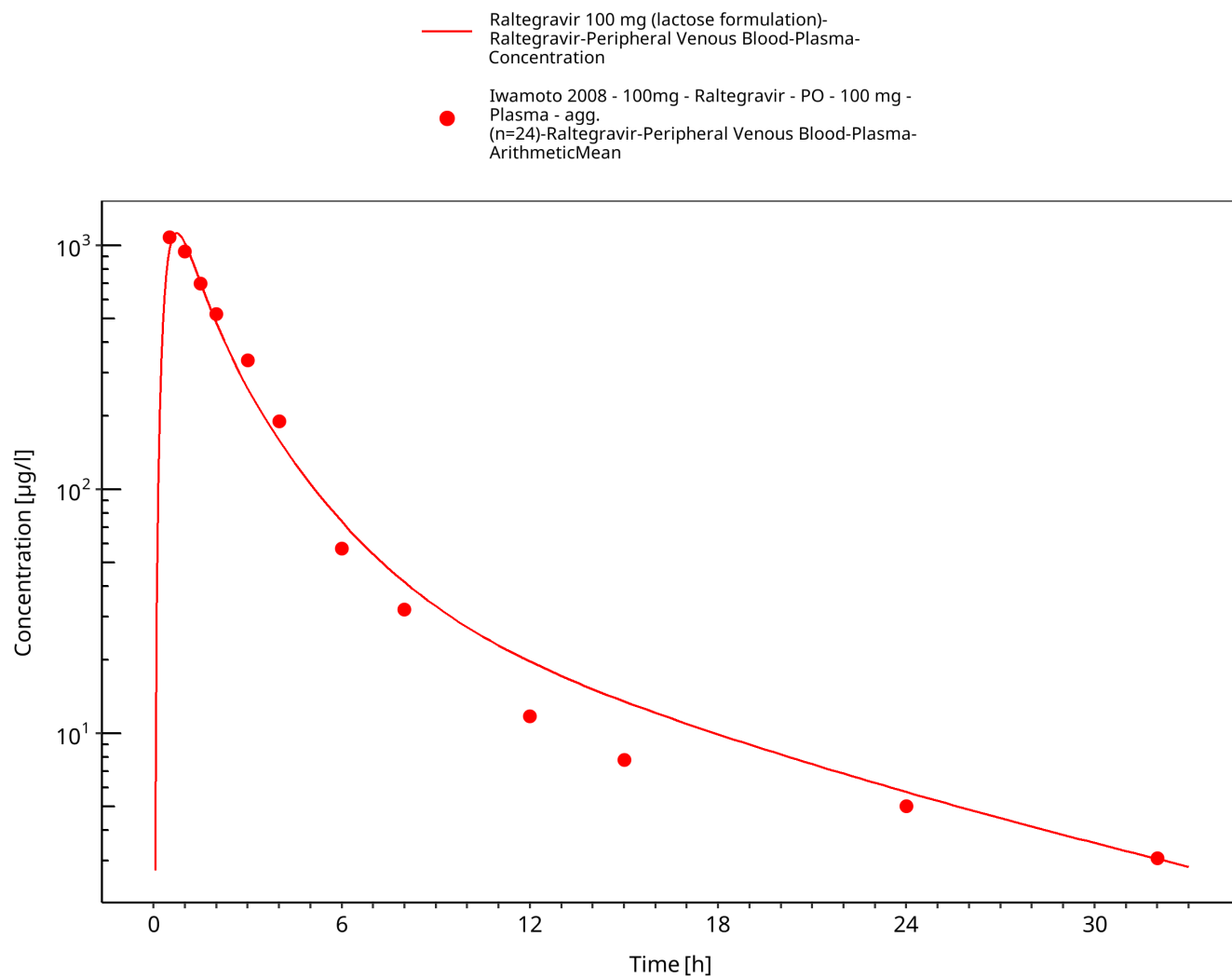


Figure 3-17: Time Profile Analysis



**Figure 3-18: Time Profile Analysis 1**





**Figure 3-19: Time Profile Analysis**

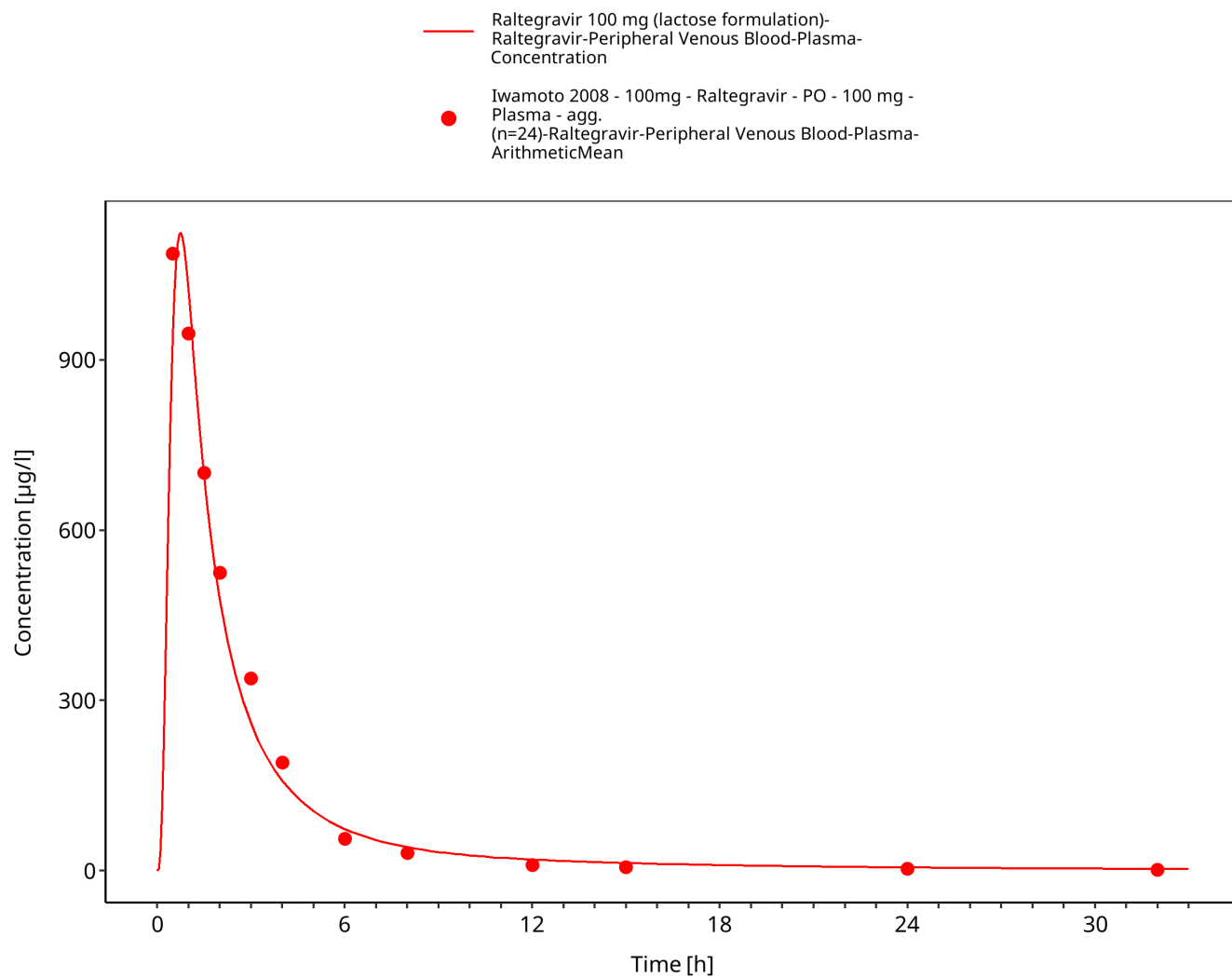


Figure 3-20: Time Profile Analysis 1

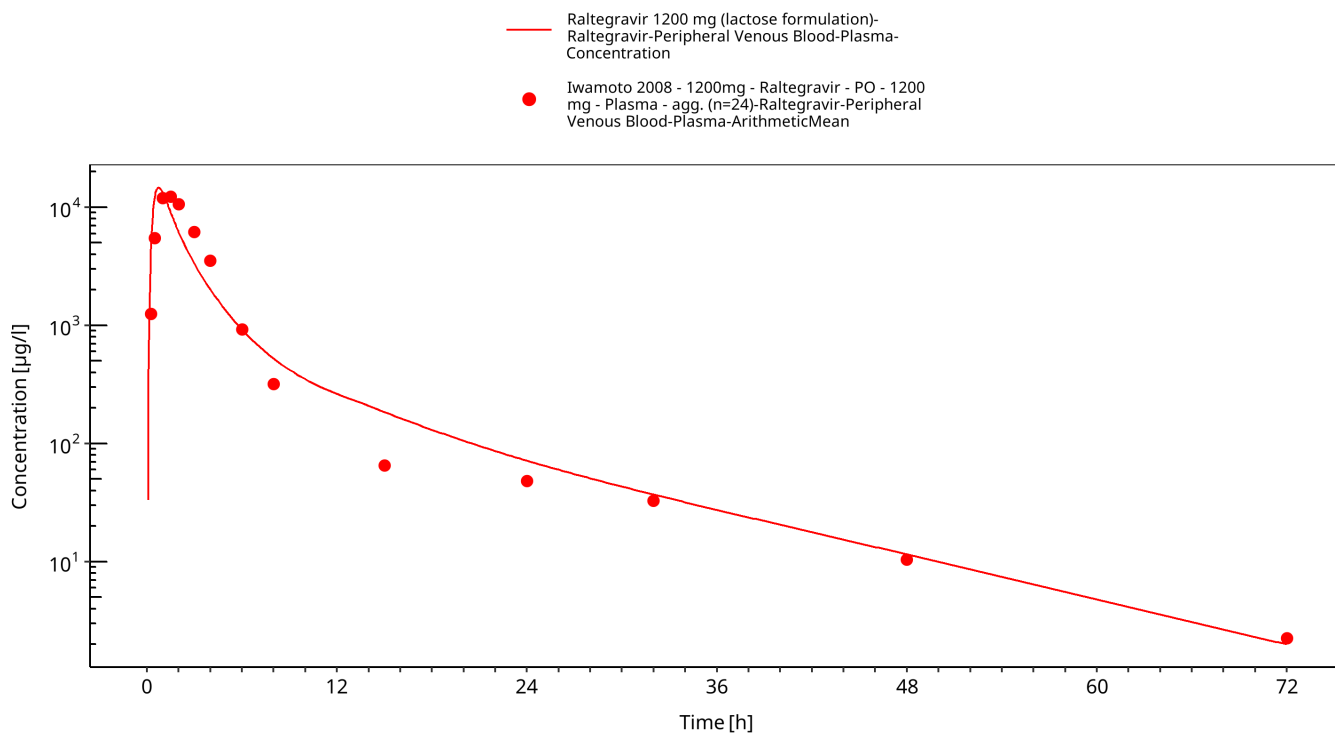


Figure 3-21: Time Profile Analysis

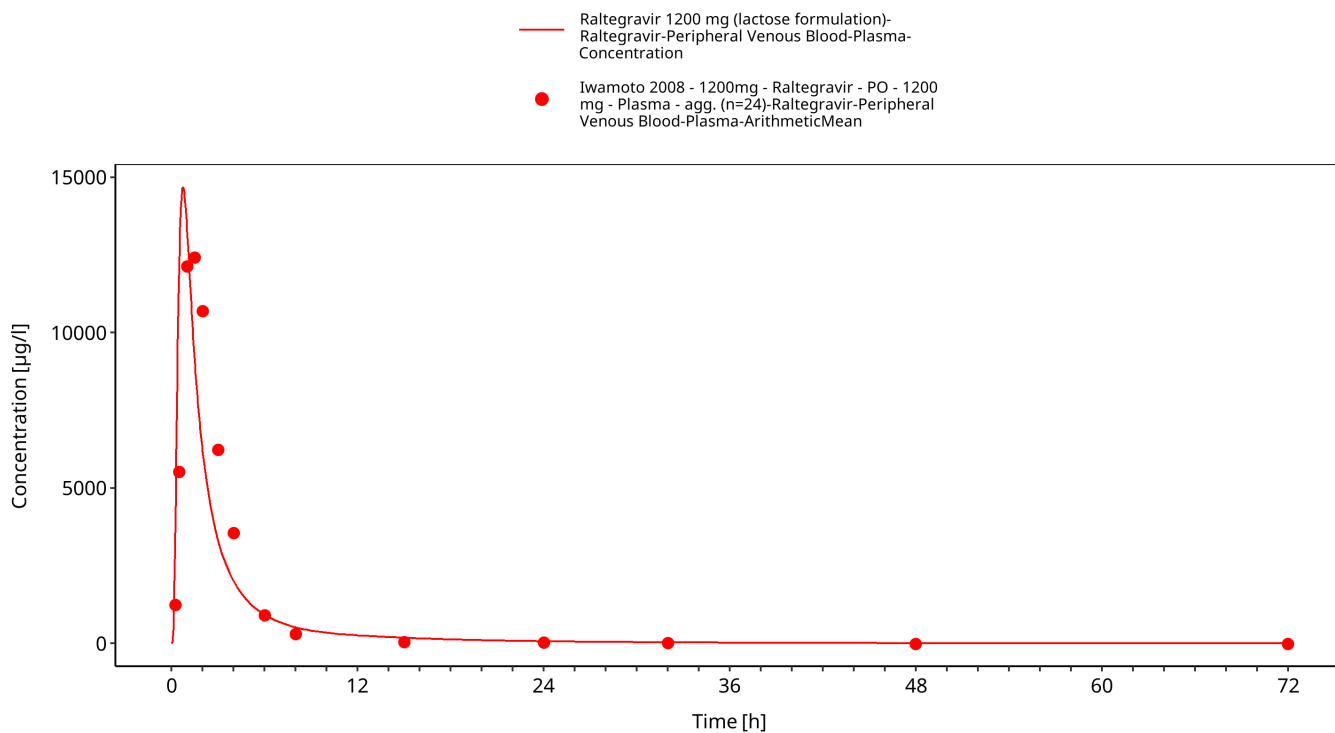


Figure 3-22: Time Profile Analysis 1

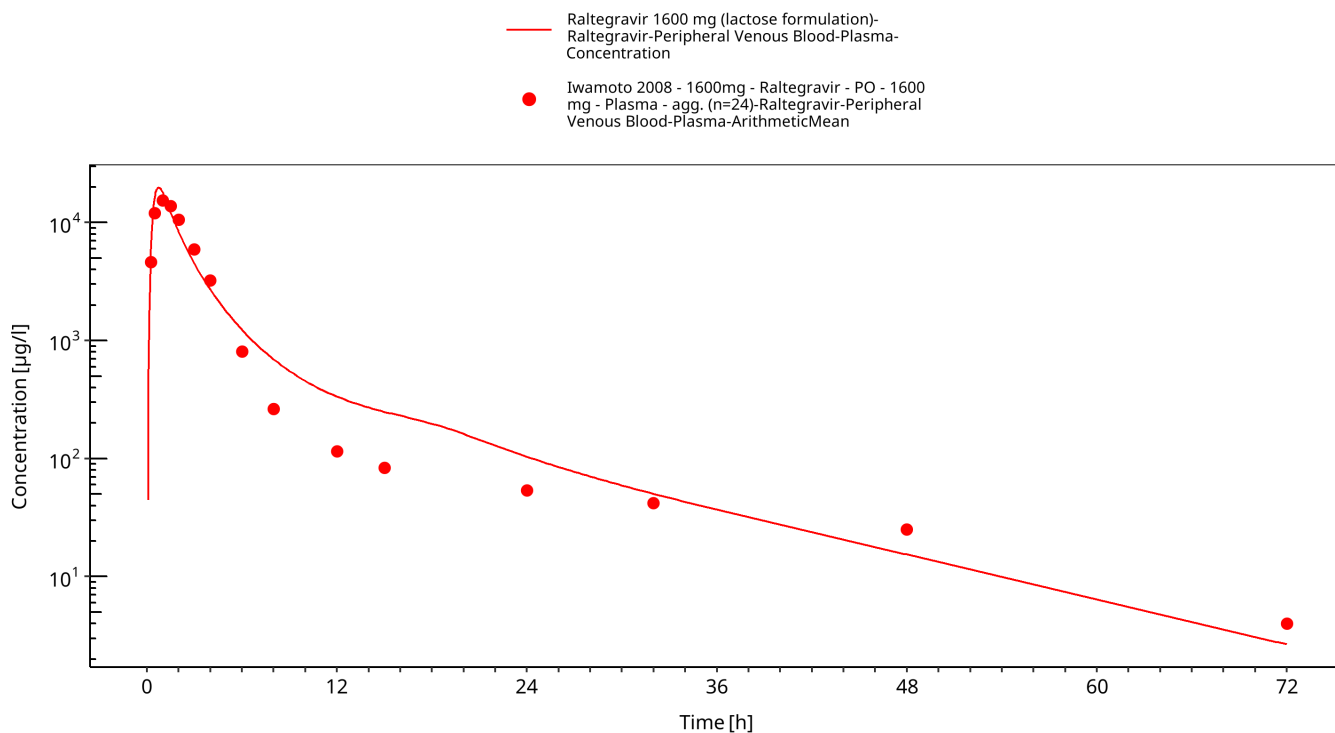


Figure 3-23: Time Profile Analysis

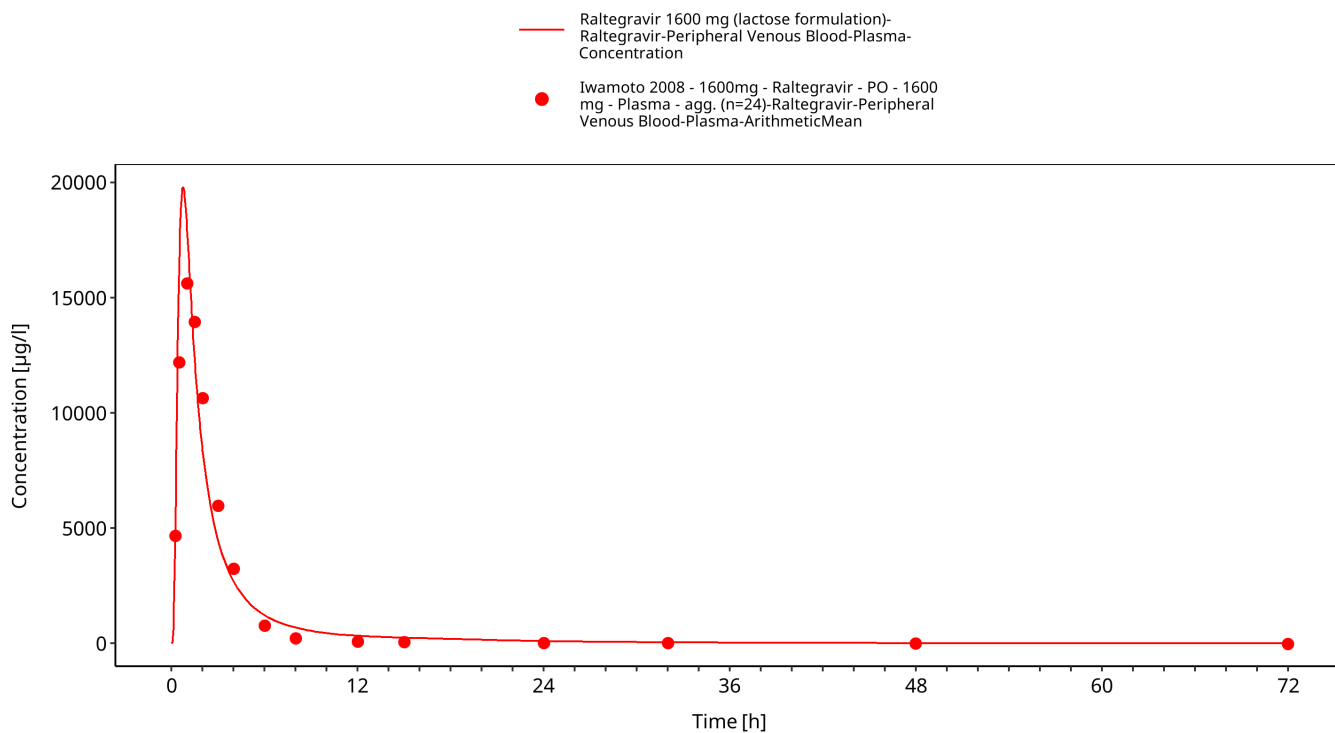
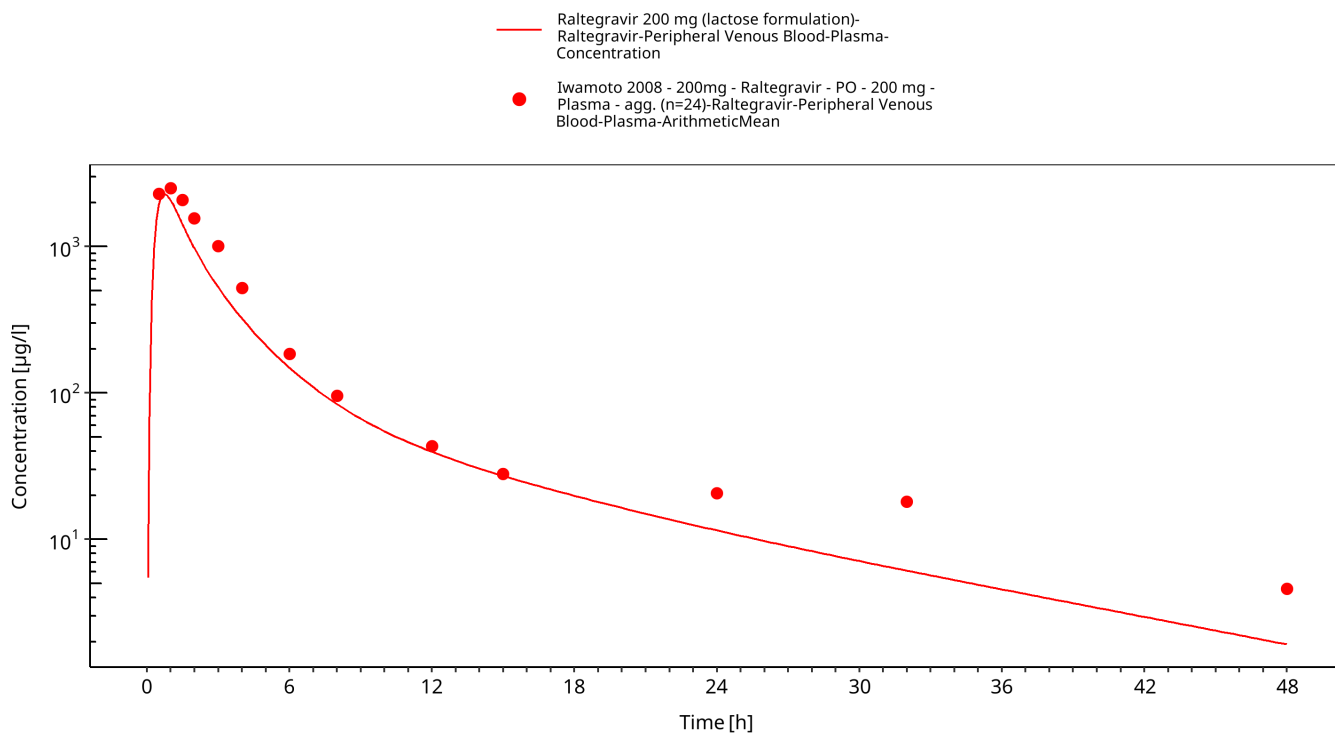
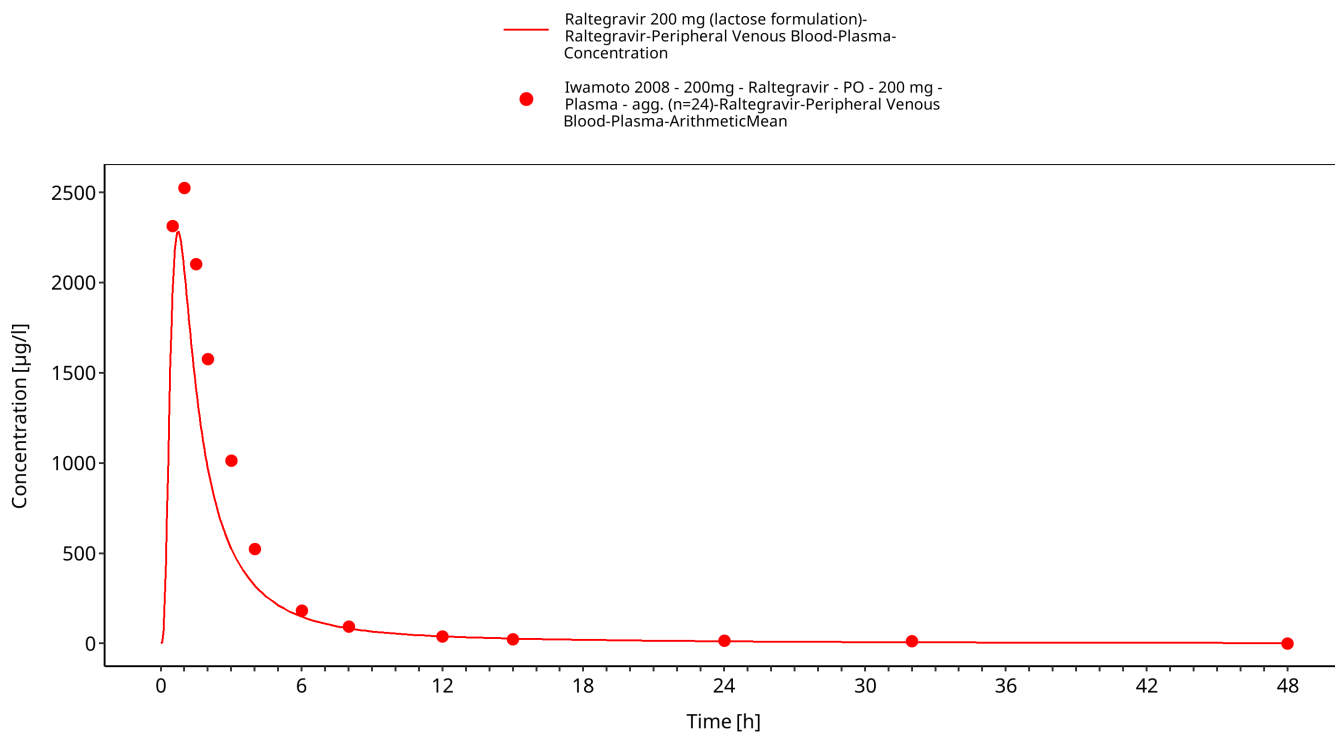


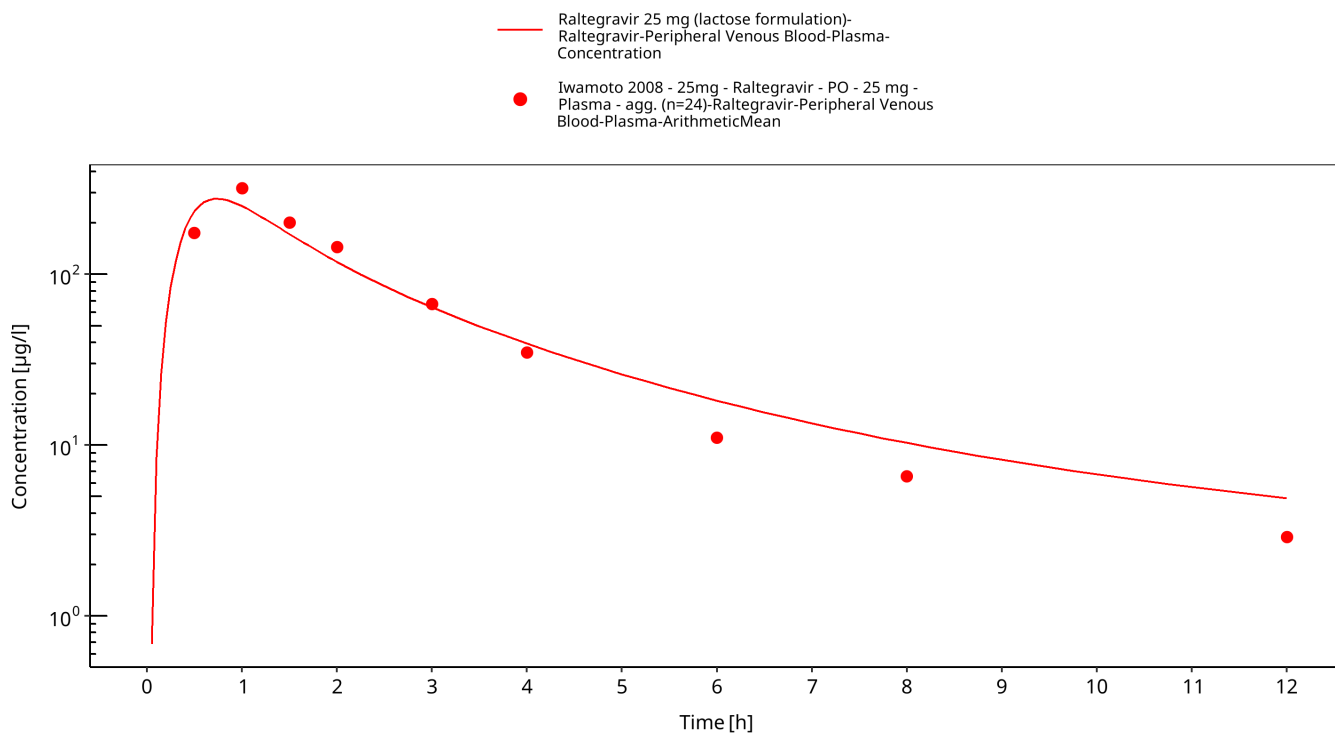
Figure 3-24: Time Profile Analysis 1



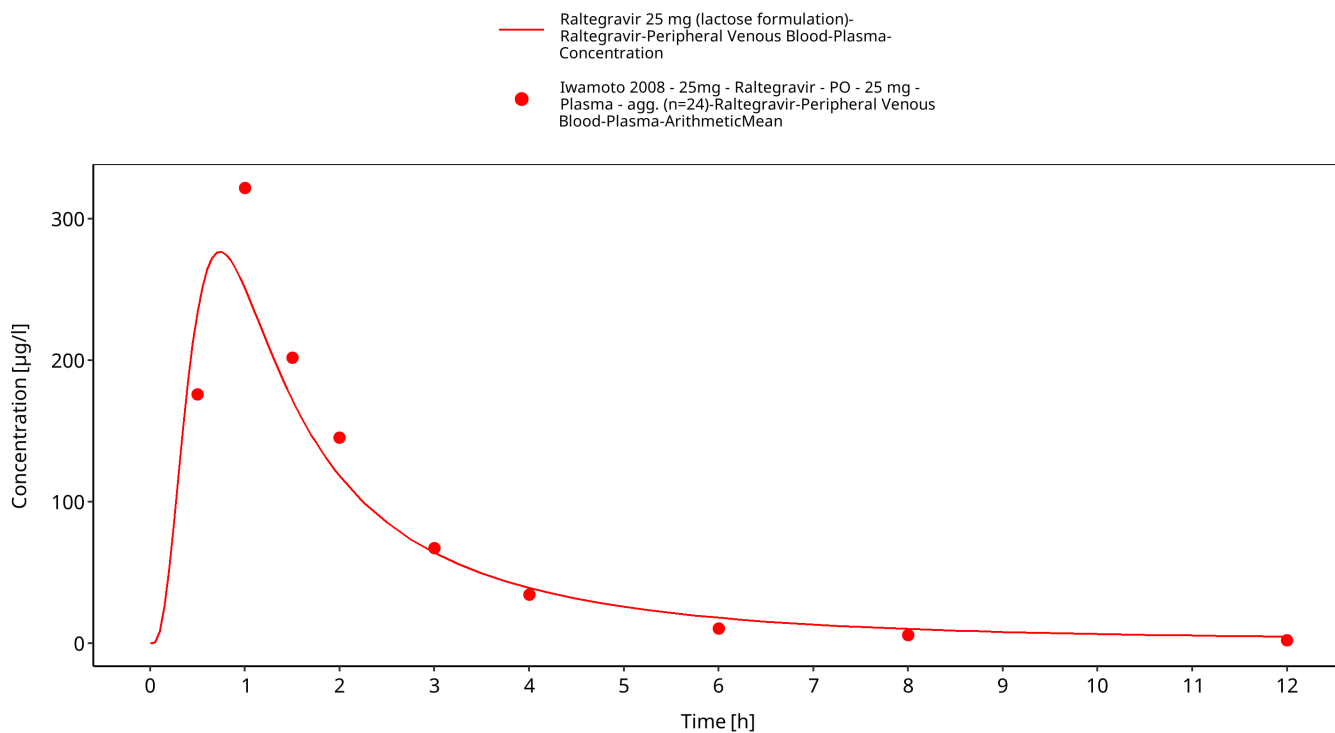
**Figure 3-25: Time Profile Analysis**



**Figure 3-26: Time Profile Analysis 1**



**Figure 3-27: Time Profile Analysis**



**Figure 3-28: Time Profile Analysis 1**

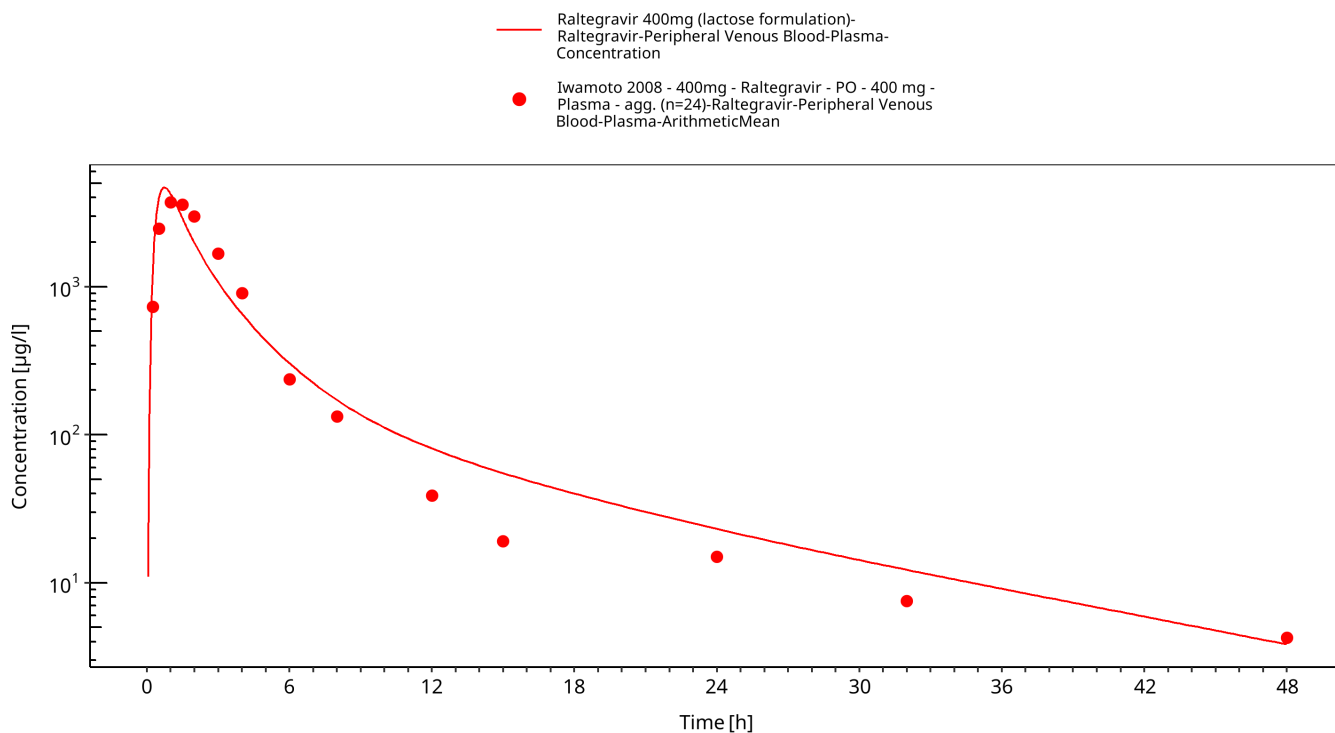


Figure 3-29: Time Profile Analysis

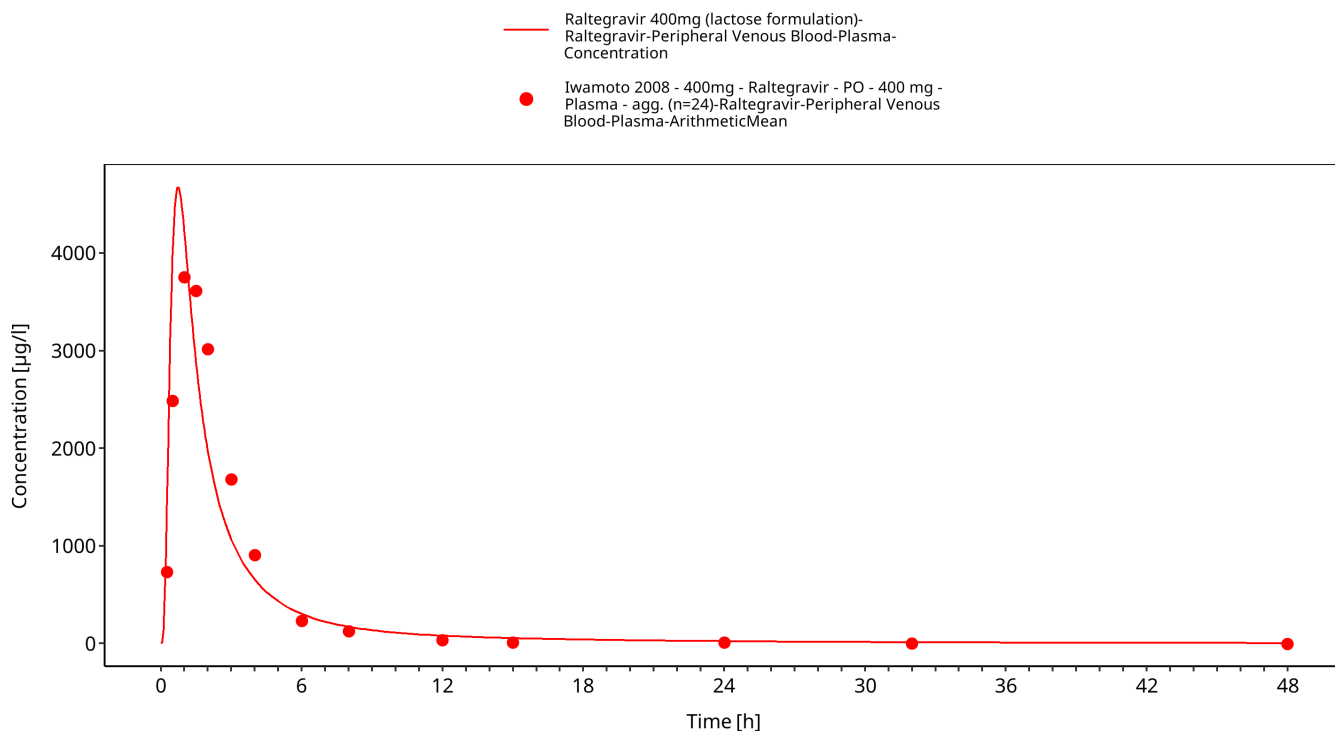
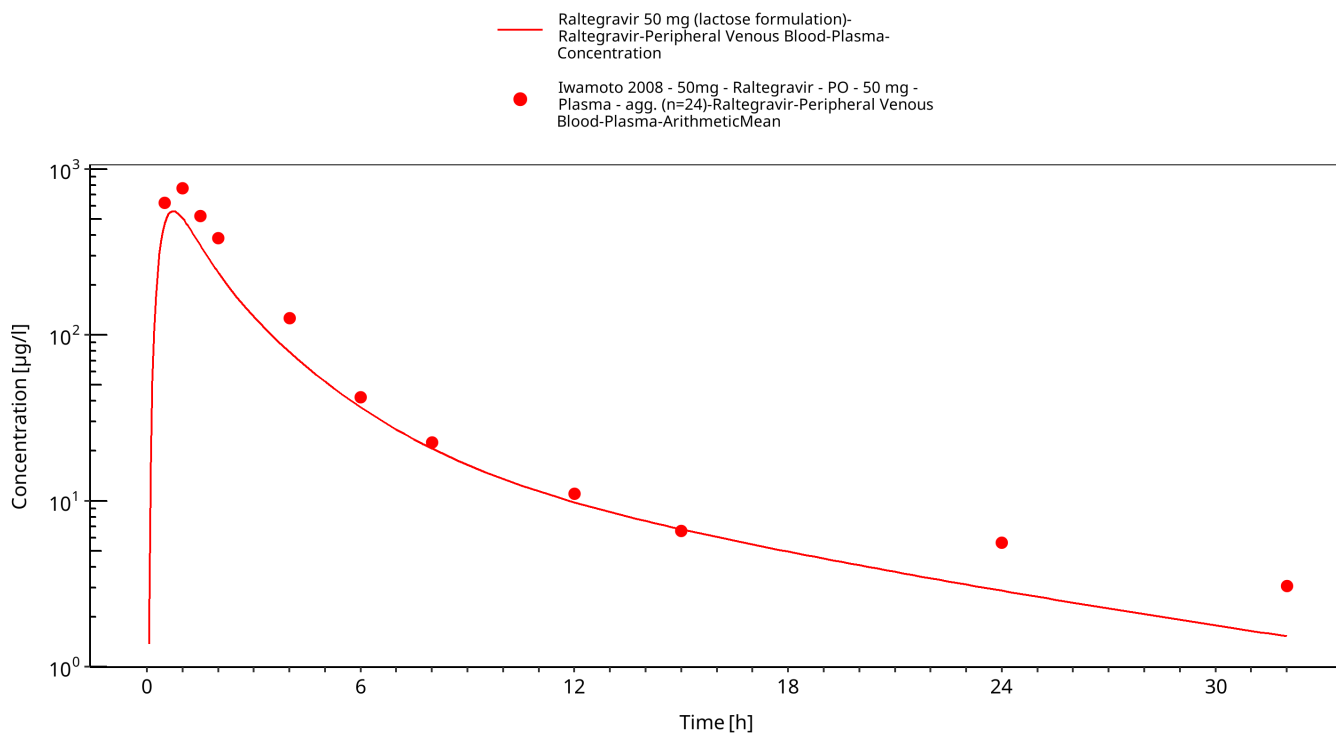
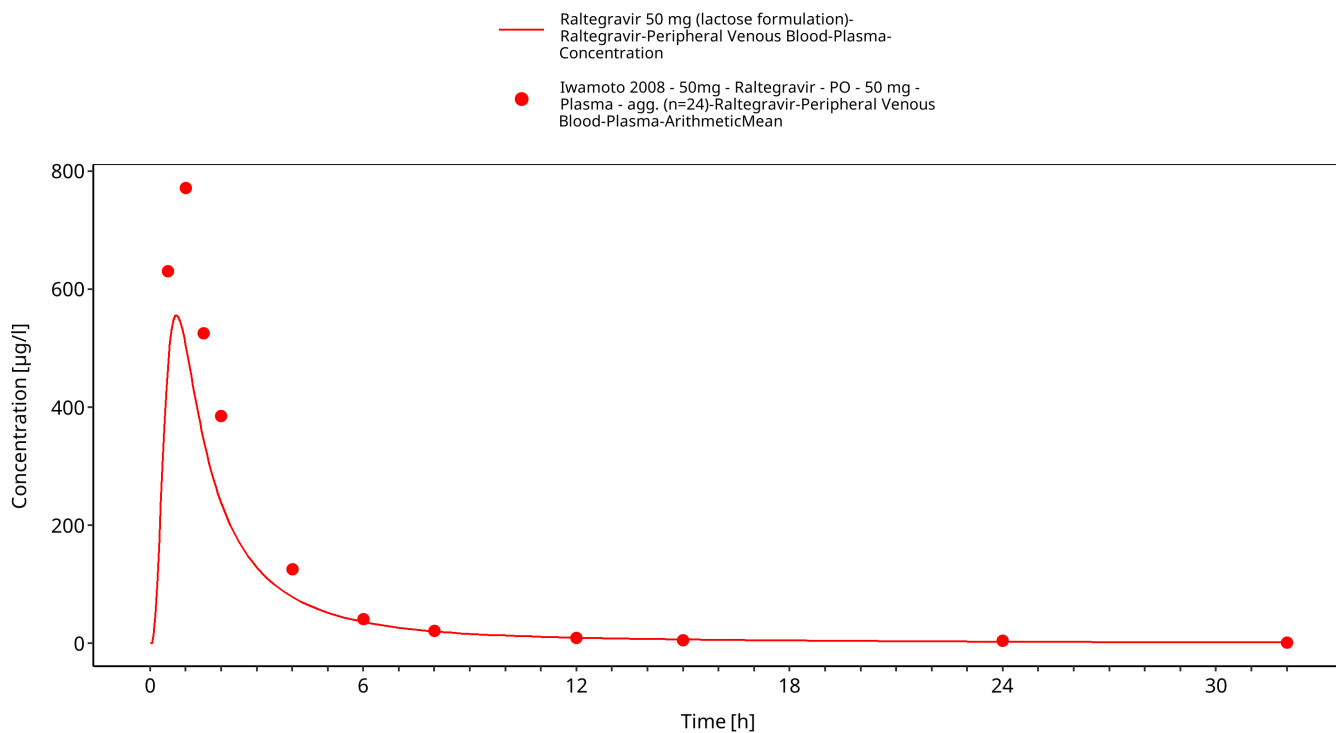


Figure 3-30: Time Profile Analysis 1



**Figure 3-31: Time Profile Analysis**



**Figure 3-32: Time Profile Analysis 1**



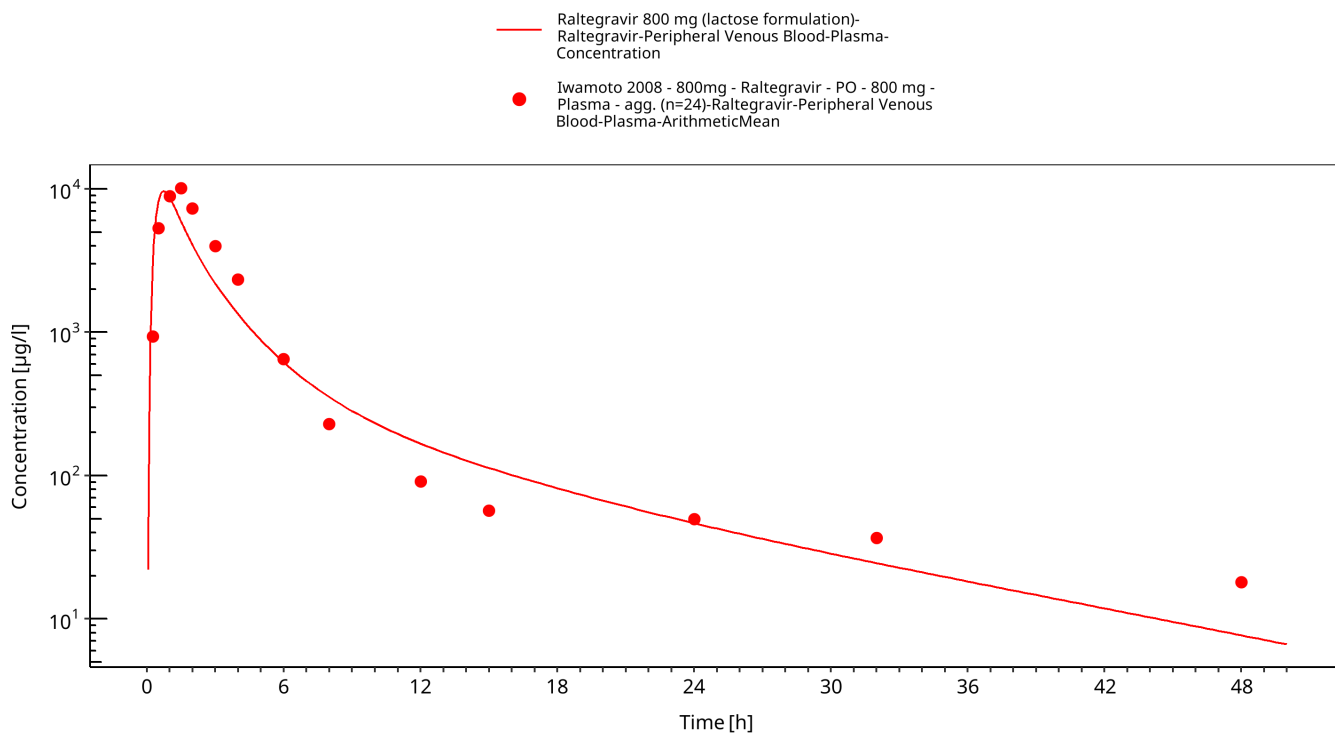


Figure 3-33: Time Profile Analysis

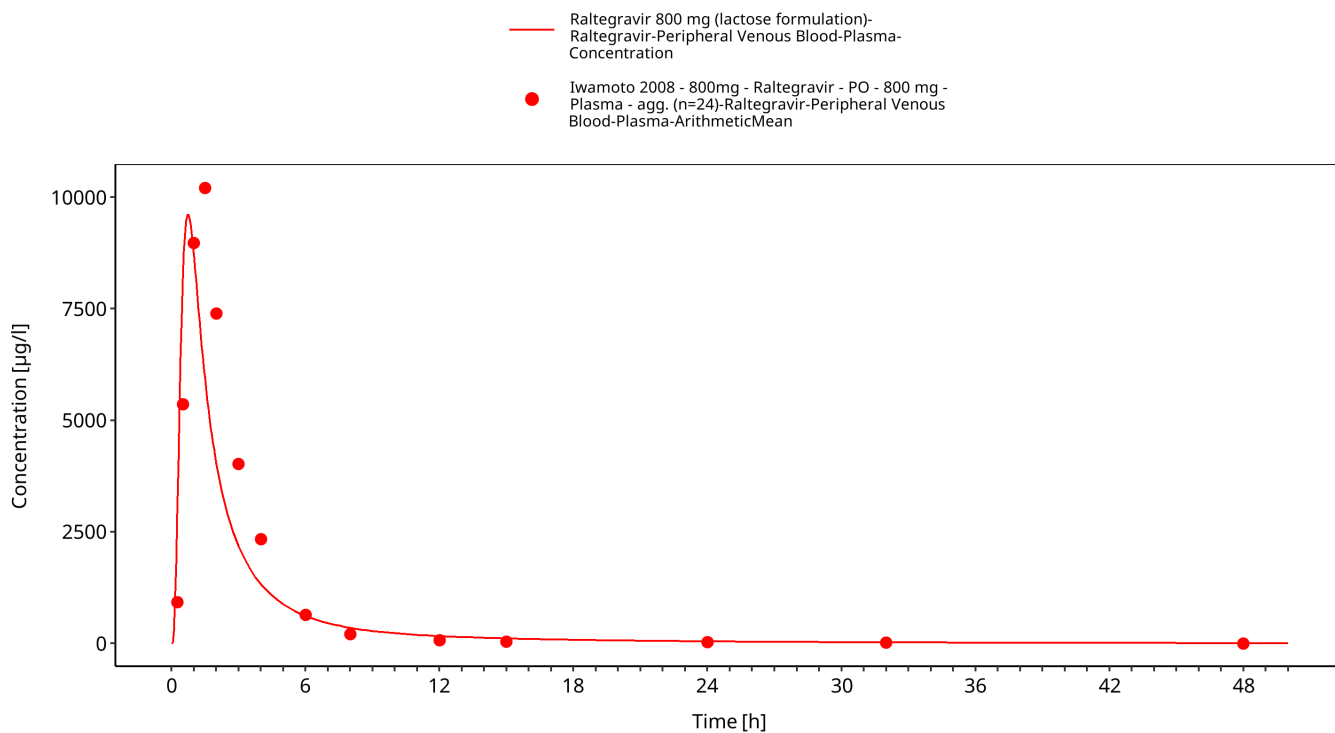


Figure 3-34: Time Profile Analysis 1

## 4 Conclusion

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The final raltegravir PBPK model applies metabolism by UGT1A1, UGT1A9 and glomerular filtration and adequately describes the pharmacokinetics of raltegravir in adults receiving SD, MD of Raltegravir ranging from 10mg to 1600mg, including four different oral formulations.

This model could be applied for the investigation of drug-drug interactions (DDI), and translation to special populations such as pediatrics with regard to UGT1A1 and 1A9 metabolism.

## 5 References

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