

Building and Evaluation of a PBPK Model for Verapamil in Adults

Version	2.1-OSP12.1
based on <i>Model Snapshot</i> and <i>Evaluation Plan</i>	https://github.com/Open-Systems-Pharmacology/Verapamil-Model/releases/tag/v2.1
OSP Version	12.1
Qualification Framework Version	3.3

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

Verapamil is used for the treatment of high blood pressure, angina (chest pain from not enough blood flow to the heart), and supraventricular tachycardia.

Verapamil is administered as a 1:1 racemate of R- and S-verapamil which are metabolized mainly by CYP3A4 to S- and R-norverapamil. All four entities are mechanism-based inactivators of CYP3A4 and non-competitive inhibitors of P-gp.

The presented verapamil model was established using observed concentration-time profiles of more than 45 clinical studies with doses from 0.1 mg to 250 mg in different verapamil dosing schedules including multiple doses and different routes of administration (intravenous, single and multiple oral administration). It includes enantioselective plasma protein binding, enantioselective metabolism by CYP3A4, non-stereospecific P-gp transport, and passive glomerular filtration. The model building and application has been published by Hanke *et al.* 2020 ([Hanke 2020](#)).

The herein presented model building and evaluation report evaluates the performance of the PBPK model for verapamil in (healthy) adults.

2 Methods

2.1 Modeling Strategy

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

Variability of plasma proteins and CYP enzymes are integrated into PK-Sim® and described in the publicly available PK-Sim® Ontogeny Database Version 7.3 ([PK-Sim Ontogeny Database Version 7.3, Schlender 2016](#)) or otherwise referenced for the specific process.

First, a base mean model was built and adjusted to clinical data including single and multiple dose studies with intravenous (only single dose) and oral applications of verapamil to find an appropriate structure to describe the pharmacokinetics in plasma. The mean PBPK model was developed using a typical European individual adjusted to the demography of the respective study population.

Unknown parameters (see below) 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.

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

In vitro / physico-chemical Data

A literature search was performed to collect available information on physiochemical properties of R- and S-verapamil and R- and S-norverapamil. The obtained information from literature is summarized in the tables below.

R-verapamil

Parameter	Unit	Value	Source	Description
MW	g/mol	454.611	Wishart 2006	Molecular weight
pK _a (base)	-	8.75	Hasegawa 1984	Acid dissociation constant
Solubility (pH 6.54)	g/L	46.0	Vogelpoel 2004	Water solubility
logP		3.79	Hansch 1995	Partition coefficient between octanol and water
fu	%	5.1	Sanaee 2011	Fraction unbound in plasma
CYP3A4 Km -> Norvera	μmol/L	19.59	Wang 2013	CYP3A4 Michaelis-Menten constant for norverapamil formation
CYP3A4 Km -> D617	μmol/L	35.34	Wang 2013	CYP3A4 Michaelis-Menten constant for D617 formation
P-gp Km	μmol/L	1.01	Shirasaka 2008	Pgp Michaelis-Menten constant
CYP3A4 MBI KI	μmol/L	27.63	Wang 2013	Conc. for half-maximal inactivation
CYP3A4 MBI kinact	1/min	0.038	Wang 2013	Maximum inactivation rate
Pgp non-competitive Ki	μmol/L	0.31	Döppenschmitt 1999	Conc. for half-maximal inactivation

S-verapamil

Parameter	Unit	Value	Source	Description
MW	g/mol	454.611	Wishart 2006	Molecular weight
pK _a (base)	-	8.75	Hasegawa 1984	Acid dissociation constant
Solubility (pH 6.54)	g/L	46.0	Vogelpoel 2004	Water solubility
logP		3.79	Hansch 1995	Partition coefficient between octanol and water
fu	%	11	Sanaee 2011	Fraction unbound in plasma
CYP3A4 Km -> Norvera	μmol/L	9.72	Wang 2013	CYP3A4 Michaelis-Menten constant for norverapamil formation
CYP3A4 Km -> D617	μmol/L	23.64	Wang 2013	CYP3A4 Michaelis-Menten constant for D617 formation
P-gp Km	μmol/L	1.01	Shirasaka 2008	Pgp Michaelis-Menten constant
CYP3A4 MBI KI	μmol/L	3.85	Wang 2013	Conc. for half-maximal inactivation
CYP3A4 MBI kinact	1/min	0.034	Wang 2013	Maximum inactivation rate
Pgp non-competitive Ki	μmol/L	0.31	Döppenschmitt 1999 (#5-references)	Conc. for half-maximal inactivation

R-norverapamil

Parameter	Unit	Value	Source	Description
MW	g/mol	440.584	Wishart 2006	Molecular weight
pK _a (base)	-	8.6 - 8.9	Sigma-Aldrich 2013	Acid dissociation constant
fu	%	5.1	assumed (from parent)	Fraction unbound in plasma
CYP3A4 Km -> D620	µmol/L	144	Tracy 1999	CYP3A4 Michaelis-Menten constant for norverapamil degradation
P-gp Km	µmol/L	1.01	assumed (from parent)	Pgp Michaelis-Menten constant
CYP3A4 MBI KI	µmol/L	6.1	Wang 2013	Conc. for half-maximal inactivation
CYP3A4 MBI kinact	1/min	0.048	Wang 2013	Maximum inactivation rate
Pgp non-competitive Ki	µmol/L	0.30	Pauli-Magnus 2000	Conc. for half-maximal inactivation

S-norverapamil

Parameter	Unit	Value	Source	Description
MW	g/mol	440.584	Wishart 2006	Molecular weight
pK _a (base)	-	8.6 - 8.9	Sigma-Aldrich 2013	Acid dissociation constant
fu	%	11	assumed (from parent)	Fraction unbound in plasma
CYP3A4 Km -> D620	µmol/L	36	Tracy 1999	CYP3A4 Michaelis-Menten constant for norverapamil degradation
P-gp Km	µmol/L	1.01	assumed (from parent)	Pgp Michaelis-Menten constant
CYP3A4 MBI KI	µmol/L	2.90	Wang 2013	Conc. for half-maximal inactivation
CYP3A4 MBI kinact	1/min	0.080	Wang 2013	Maximum inactivation rate
Pgp non-competitive Ki	µmol/L	0.30	Pauli-Magnus 2000	Conc. for half-maximal inactivation

Clinical Data

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

Model Building and parameterizing of CYP3A4 interaction

The following studies were used for model building and parameterization of CYP3A4 interaction: If not stated otherwise, the drug was given as a 1:1 racemate of S- and R-verapamil.

Publication	Arm / Treatment / Information used for model building
Eichelbaum 1984	Healthy subjects receiving single intravenous doses of 5, 25 and 50 mg of R-verapamil and 5, 7.5 and 10 mg of S-verapamil
Streit 2005	Healthy subjects receiving single intravenous doses of 5 mg
Barbarash 1988	Healthy subjects receiving single intravenous doses of 10 mg
Abernethy 1993	Healthy subjects receiving single intravenous doses of 20 mg
Maeda 2011	Healthy subjects receiving single oral doses of 0.1, 3 and 16 mg
Blume 1989	Healthy subjects receiving single oral doses of 80 mg
Ratiopharm 1988	Healthy subjects receiving single oral doses of 80 mg
Johnson 2001	Healthy subjects receiving multiple oral doses of 80 mg TID.
Härtter 2012	Healthy subjects receiving single oral doses of 120 mg and multiple oral doses of 120 mg BID
Hla 1987	Healthy subjects receiving multiple oral doses of 120 mg BID

Model verification

The following studies were used for model verification:

Publication	Arm / Treatment / Information used for model building
Mooy 1985	Healthy subjects receiving single intravenous doses of 3 mg and single oral doses of 80 mg
Johnston 1981	Healthy subjects receiving single intravenous doses of 0.1 mg/kg and single oral doses of 120 mg
Abernethy 1985	Healthy subjects receiving single intravenous doses of 10 mg and single oral doses of 120 mg
Barbarash 1988	Healthy subjects receiving single oral doses of 120 mg
Wing 1985	Healthy subjects receiving single intravenous doses 10mg and single oral doses of 80 mg
McAllister 1982	Healthy subjects receiving single intravenous doses of 10 mg
Smith 1984	Healthy subjects receiving single intravenous doses of 10 mg and single oral doses of 120 mg
Freedman 1981	Healthy subjects receiving single intravenous doses of 13.1 mg
Vogelsang 1984	Healthy subjects receiving single oral doses of 250mg R-verapamil
Blume 1983	Healthy subjects receiving single oral doses of 40 mg
Blume 1990	Healthy subjects receiving single oral doses of 40 mg
John 1992	Healthy subjects receiving single oral doses of 40 mg
Sawicki 2002	Healthy subjects receiving single oral doses of 40 mg
Choi 2008	Healthy subjects receiving single oral doses of 60 mg
Wing 1985	Healthy subjects receiving single oral doses of 80 mg
Maeda 2011	Healthy subjects receiving single oral doses of 80 mg
Ratiopharm 1989	Healthy subjects receiving single oral doses of 80 mg
Boehringer 2018	Healthy subjects receiving single oral doses of 120 mg
Blume 1987	Healthy subjects receiving single oral doses of 120 mg
Johnston 1981	Healthy subjects receiving single oral doses of 120 mg
Mikus 1990	Healthy subjects receiving single oral doses of 160 mg
van Haarst 2009	Healthy subjects receiving multiple oral doses of 180 mg BID
Blume 1994	Healthy subjects receiving single oral doses of 240 mg QD
Joergenson 1988	Healthy subjects receiving multiple oral doses of 120 mg BID
Shand 1981	Healthy subjects receiving multiple oral doses of 120 mg TID
Karim 1995	Healthy subjects receiving single oral doses of 240 mg

2.3 Model Parameters and Assumptions

Absorption

Verapamil is transported by P-gp. The model includes non-stereospecific P-gp transport.

Distribution

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.

Metabolism, Elimination and Inhibition

Verapamil is metabolized by CYP3A4 and transported by P-gp. The model includes enantioselective metabolism by CYP3A4, non-stereospecific P-gp transport. Additionally passive glomerular filtration was integrated.

Mechanism-based inactivation of CYP3A4 and non-competitive inhibition of P-gp by all four entities (S-verapamil, R-verapamil, S-norverapamil and R-norverapamil) was taken into account. The CYP3A4 MBI KI and kinact values were taken from literature, the KI values for P-gp inhibition were optimized.

Automated Parameter Identification

The parameter identification tool in PK-Sim has been used to estimate selected model parameters by adjusting to PK data of the clinical studies that were used in the model building process.

The result of the final parameter identification is shown in the tables below:

R-verapamil

Model Parameter	Optimized Value	Unit
logP	2.84	
CYP3A4 kcat -> Norvera	34.94	1/min
CYP3A4 kcat -> D617	43.98	1/min
P-gp kcat	12.60	1/min
Pgp non-competitive Ki	0.038	µmol/L
Cellular permeability	9.94E-02	cm/min
Intestinal permeability	3.54E-06	cm/min
SR tablet Weibull time	155.24	min
SR tablet Weibull shape	2.37	

S-verapamil

Model Parameter	Optimized Value	Unit
logP	2.84	
CYP3A4 kcat -> Norvera	26.17	1/min
CYP3A4 kcat -> D617	56.42	1/min
P-gp kcat	12.60	1/min
Pgp non-competitive Ki	0.038	µmol/L
Cellular permeability	9.94E-02	cm/min
Intestinal permeability	3.54E-06	cm/min
SR tablet Weibull time	155.24	min
SR tablet Weibull shape	2.37	

R-norverapamil

Model Parameter	Optimized Value	Unit
logP	2.84	
CYP3A4 kcat -> D620	145.64	1/min
P-gp kcat	3.39	1/min
Pgp non-competitive Ki	0.038	µmol/L
Cellular permeability	9.94E-02	cm/min
Intestinal permeability	3.54E-06	cm/min

S-norverapamil

Model Parameter	Optimized Value	Unit
logP	2.84	
CYP3A4 kcat -> D620	41.10	1/min
P-gp kcat	3.39	1/min
Pgp non-competitive Ki	0.038	µmol/L
Cellular permeability	9.94E-02	cm/min
Intestinal permeability	3.54E-06	cm/min

3 Results and Discussion

The PBPK model for verapamil was developed and evaluated using publically available, clinical pharmacokinetic data from studies listed in [Section 2.2](#).

The next sections show:

1. the final model 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 Final input parameters

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

Compound: R-Verapamil

Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	46 mg/ml		Measurement	True
Reference pH	6.54		Measurement	True
Lipophilicity	2.8407448658 Log Units	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	logP	True
Fraction unbound (plasma, reference value)	5.1 %	Publication-In Vivo	Measurement	True
Permeability	0.0994098912 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Specific intestinal permeability (transcellular)	3.5447164381E-06 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Is small molecule	Yes			
Molecular weight	454.611 g/mol			
Plasma protein binding partner	Unknown			

Calculation methods

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

Processes

Systemic Process: Glomerular Filtration-GFR

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	1	

Transport Protein: P-gp-Paper

Molecule: P-gp

Parameters

Name	Value	Value Origin
In vitro Vmax/transporter	0.00057724 pmol/min/pmol transporter	
Km	1.01 μmol/l	
kcat	12.5970868779 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: P-gp-Non-competitive

Molecule: P-gp

Parameters

Name	Value	Value Origin
Ki	0.0383697779 μmol/l	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: CYP3A4-MBI

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
kinact	0.038 1/min	
K_kinact_half	27.63 μmol/l	

Metabolizing Enzyme: CYP3A4-Norverapamil

Molecule: CYP3A4

Metabolite: R-Norverapamil

Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	1.27 nmol/min/mg mic. protein	
Km	19.59 μmol/l	
kcat	34.9352466212 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Metabolizing Enzyme: CYP3A4-D617

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	1.17 nmol/min/mg mic. protein	
Km	35.34 μmol/l	
kcat	43.9812289146 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Compound: S-Verapamil

Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	46 mg/ml		Measurement	True
Reference pH	6.54		Measurement	True
Lipophilicity	2.8407448658 Log Units	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	logP	True
Fraction unbound (plasma, reference value)	11 %	Publication-In Vivo	Measurement	True
Permeability	0.0994098912 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Specific intestinal permeability (transcellular)	3.5447164381E-06 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Is small molecule	Yes			
Molecular weight	454.611 g/mol			
Plasma protein binding partner	Unknown			

Calculation methods

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

Processes

Systemic Process: Glomerular Filtration-GFR

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	1	

Transport Protein: P-gp-Paper

Molecule: P-gp

Parameters

Name	Value	Value Origin
In vitro Vmax/transporter	0.00057724 pmol/min/pmol transporter	
Km	1.01 μmol/l	
kcat	12.5970868779 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: P-gp-Non-competitive

Molecule: P-gp

Parameters

Name	Value	Value Origin
Ki	0.0383697779 μmol/l	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: CYP3A4-MBI

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
kinact	0.034 1/min	
K_kinact_half	3.85 μmol/l	

Metabolizing Enzyme: CYP3A4-Norverapamil

Molecule: CYP3A4

Metabolite: S-Norverapamil

Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	1.02 nmol/min/mg mic. protein	
Km	9.72 μmol/l	
kcat	26.1743386639 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Metabolizing Enzyme: CYP3A4-D617

Parameters

Name	Value	Value Origin
In vitro Vmax for liver microsomes	0.86 nmol/min/mg mic. protein	
Km	23.64 $\mu\text{mol/l}$	
kcat	56.4245798193 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Compound: R-Norverapamil

Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	46 mg/ml		Measurement	True
Reference pH	6.54		Measurement	True
Lipophilicity	2.8407448658 Log Units	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	logP	True
Fraction unbound (plasma, reference value)	5.1 %	Other-Assumption	Measurement	True
Permeability	0.0994098912 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Specific intestinal permeability (transcellular)	3.5447164381E-06 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Is small molecule	Yes			
Molecular weight	440.584 g/mol			
Plasma protein binding partner	Unknown			

Calculation methods

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

Processes

Metabolizing Enzyme: CYP3A4-D620

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
In vitro Vmax/recombinant enzyme	9 pmol/min/pmol rec. enzyme	
Km	144 μmol/l	
kcat	145.6385399671 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Systemic Process: Glomerular Filtration-GFR

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	1	

Transport Protein: P-gp-Paper

Molecule: P-gp

Parameters

Name	Value	Value Origin
In vitro Vmax/transporter	0.00057724 pmol/min/pmol transporter	
Km	1.01 μmol/l	
kcat	3.3916609583 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: P-gp-Non-competitive

Molecule: P-gp

Parameters

Name	Value	Value Origin
Ki	0.0383697779 µmol/l	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: CYP3A4-MBI

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
kinact	0.048 1/min	
K_kinact_half	6.1 µmol/l	

Compound: S-Norverapamil

Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	46 mg/ml		Measurement	True
Reference pH	6.54		Measurement	True
Lipophilicity	2.8407448658 Log Units	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	logP	True
Fraction unbound (plasma, reference value)	11 %	Other-Assumption	Measurement	True
Permeability	0.0994098912 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Specific intestinal permeability (transcellular)	3.5447164381E-06 cm/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23	Fitted	True
Is small molecule	Yes			
Molecular weight	440.584 g/mol			
Plasma protein binding partner	Unknown			

Calculation methods

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

Processes

Metabolizing Enzyme: CYP3A4-D620

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
In vitro Vmax/recombinant enzyme	6.5 pmol/min/pmol rec. enzyme	
Km	36 μmol/l	
kcat	41.0994047535 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Systemic Process: Glomerular Filtration-GFR

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	1	

Transport Protein: P-gp-Paper

Molecule: P-gp

Parameters

Name	Value	Value Origin
In vitro Vmax/transporter	0.00057724 pmol/min/pmol transporter	
Km	1.01 μmol/l	
kcat	3.3916609583 1/min	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: P-gp-Non-competitive

Molecule: P-gp

Parameters

Name	Value	Value Origin
Ki	0.0383697779 µmol/l	Parameter Identification-Parameter Identification-Value updated from '30b - final' on 2019-12-30 13:23

Inhibition: CYP3A4-MBI

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
kinact	0.08 1/min	
K_kinact_half	2.9 µmol/l	

Formulation: Solution

Type: Dissolved

Formulation: Retard Tablet Verapamil (Knoll)

Type: Weibull

Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	155.2445399403 min	Parameter Identification-Parameter Identification-Value updated from '240 mg retard (Isoptin RR) QD vs Verabeta 240 RR' on 2019-12-31 11:13
Lag time	0 min	
Dissolution shape	2.3662989419	Parameter Identification-Parameter Identification-Value updated from '240 mg retard (Isoptin RR) QD vs Verabeta 240 RR' on 2019-12-31 11:13
Use as suspension	Yes	

3.2 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](#).

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
Intravenous administration - norverapamil	1.66
Intravenous administration - R-verapamil	1.25
Intravenous administration - S-verapamil	1.31
Intravenous administration - verapamil	1.45
Oral administration - norverapamil	1.29
Oral administration - R-norverapamil	1.14
Oral administration - R-verapamil	1.33
Oral administration - S-norverapamil	1.16
Oral administration - S-verapamil	1.31
Oral administration - verapamil	1.42
All	1.36

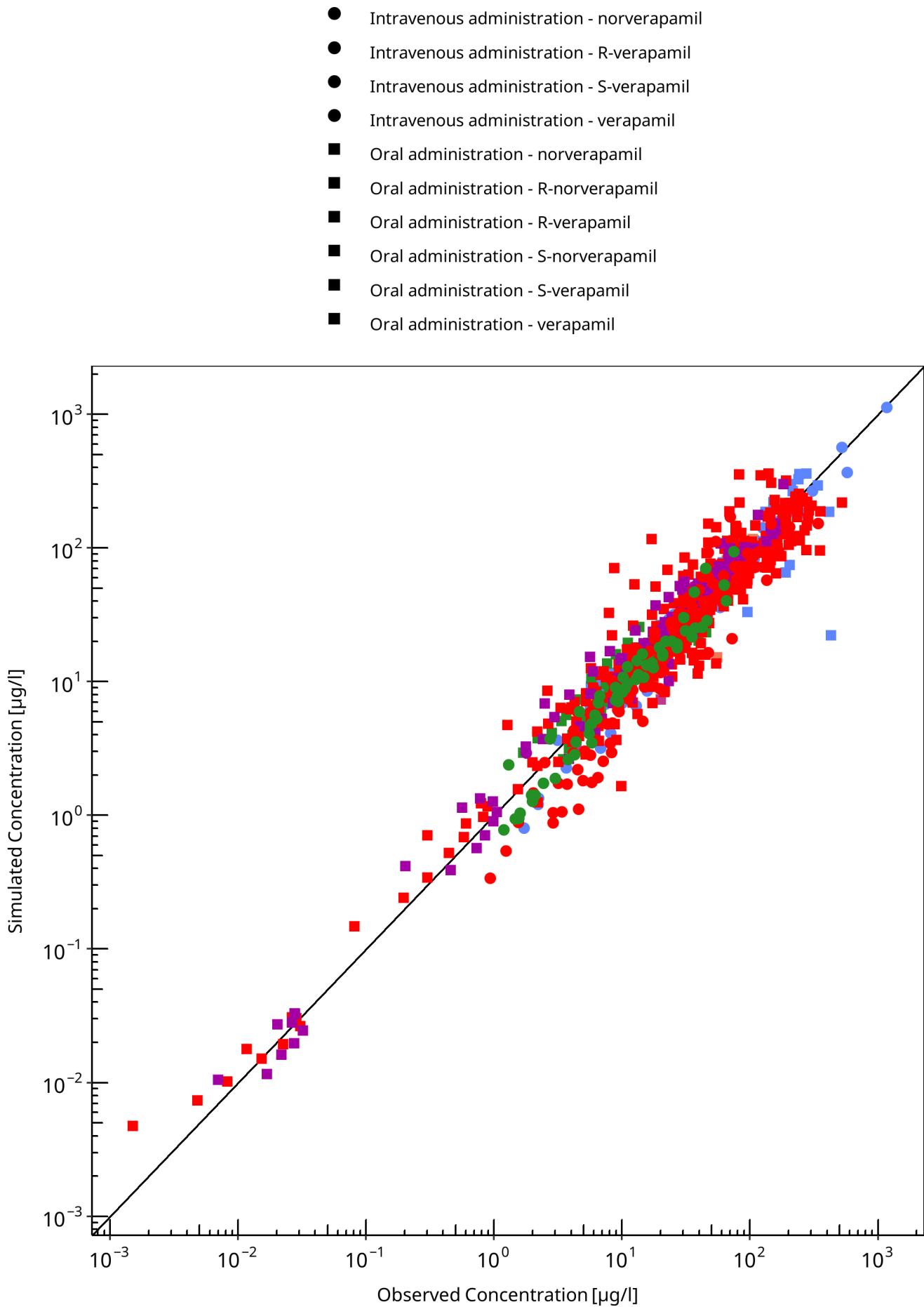


Figure 3-1: Goodness of fit plot for concentration in plasma

- Intravenous administration - norverapamil
- Intravenous administration - R-verapamil
- Intravenous administration - S-verapamil
- Intravenous administration - verapamil
- Oral administration - norverapamil
- Oral administration - R-norverapamil
- Oral administration - R-verapamil
- Oral administration - S-norverapamil
- Oral administration - S-verapamil
- Oral administration - verapamil

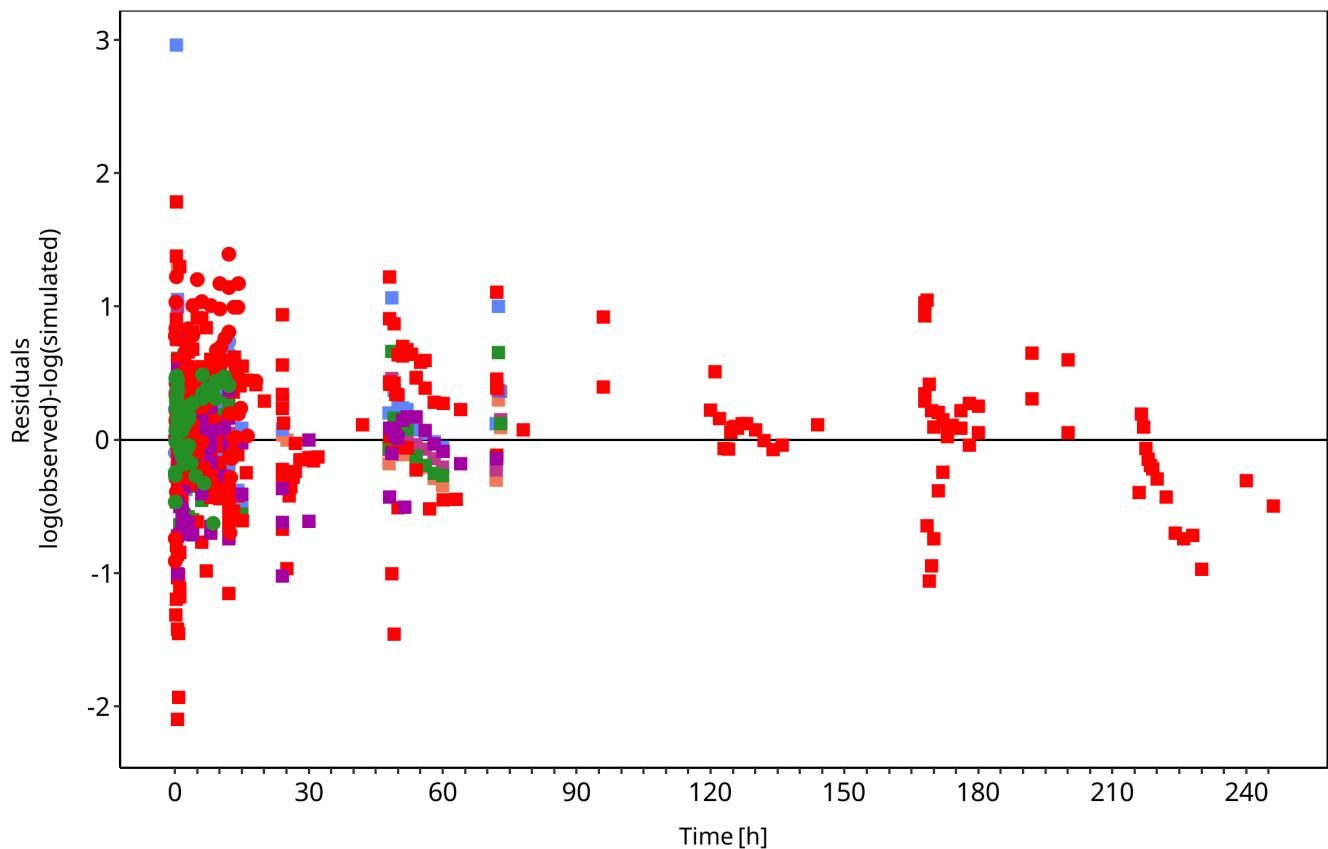


Figure 3-2: Goodness of fit plot for concentration in plasma

3.3 Concentration-Time Profiles

Simulated versus observed concentration-time profiles of all data listed in [Section 2.2](#) are presented below.

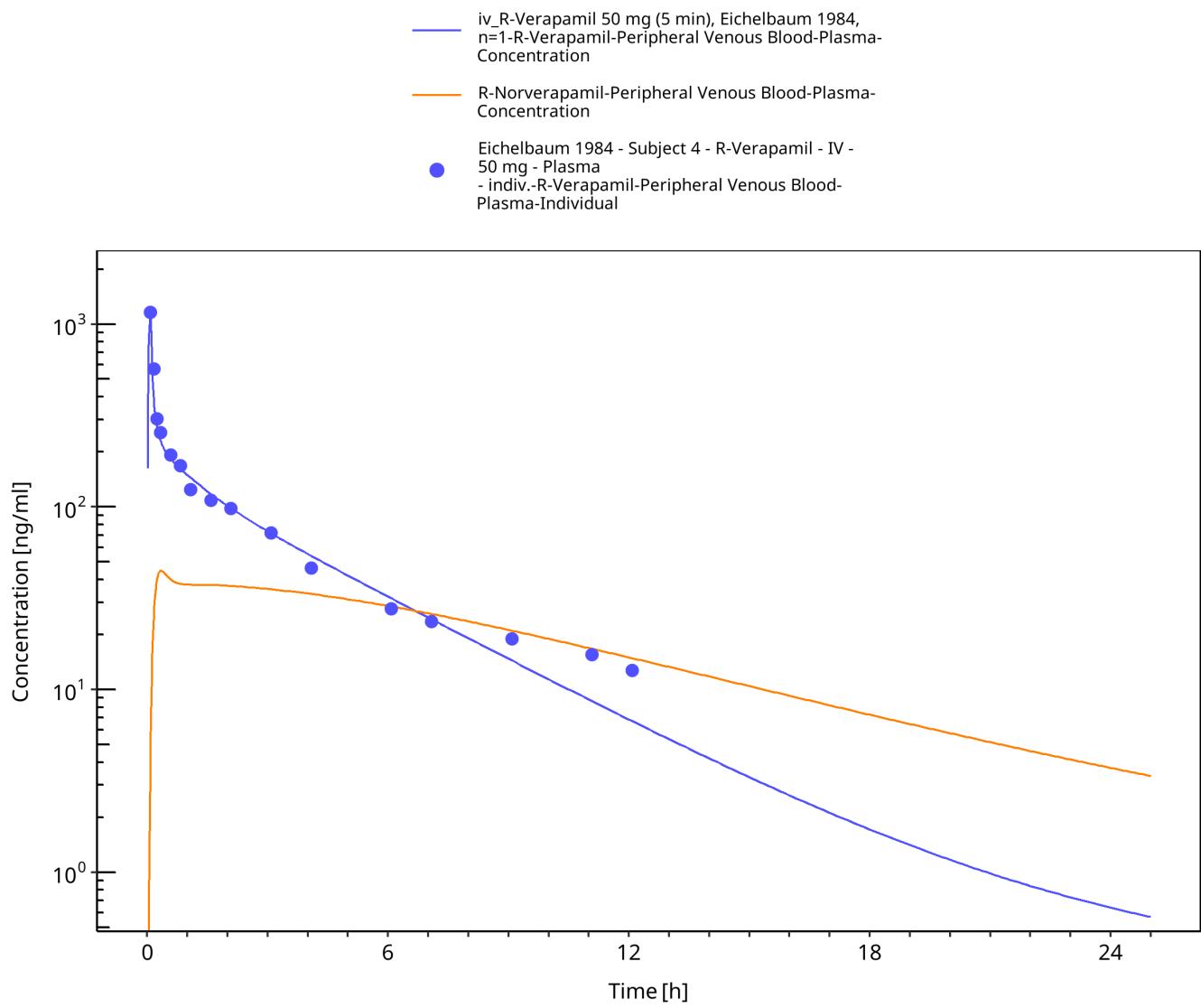


Figure 3-3: Time Profile Analysis

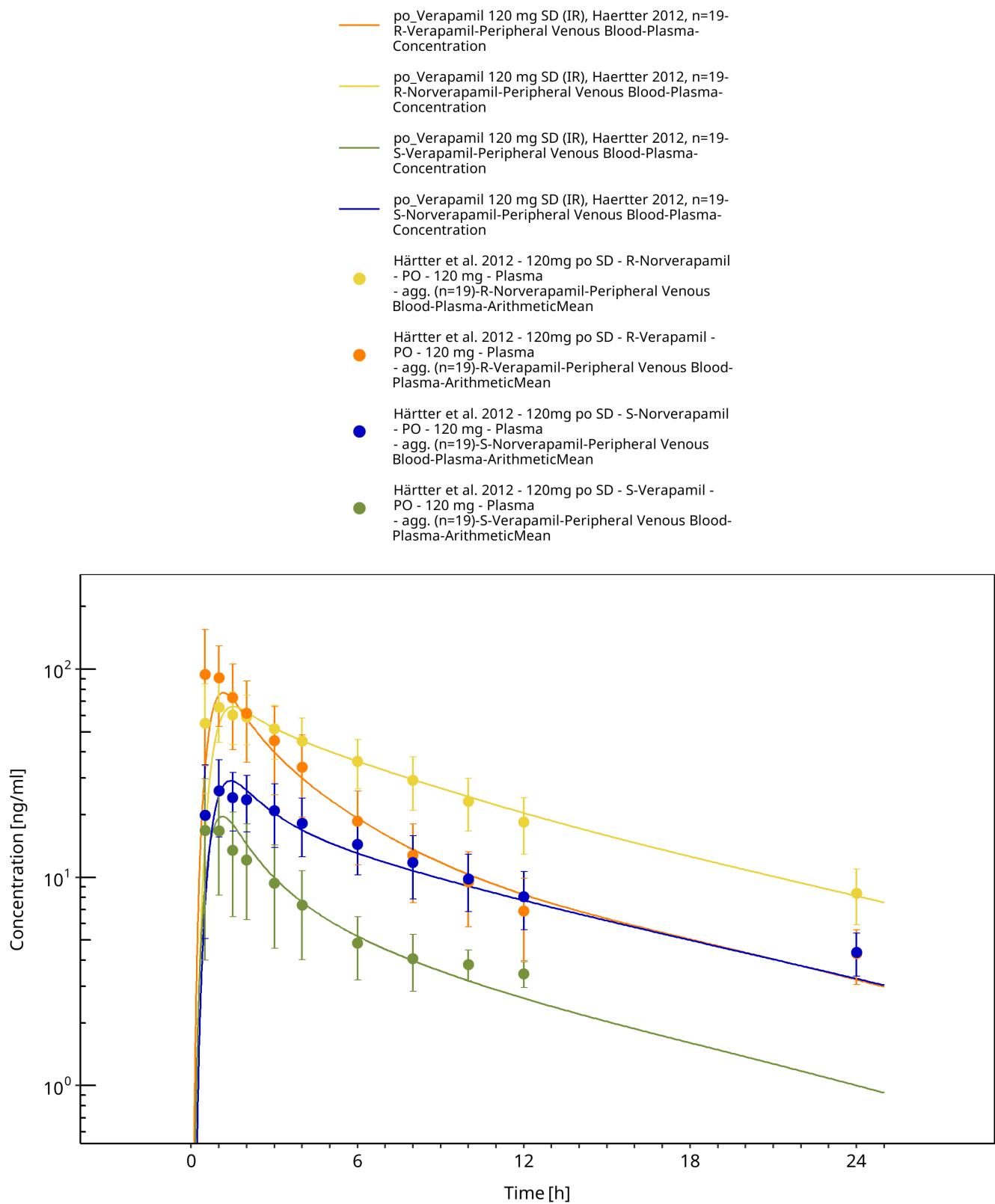


Figure 3-4: Time Profile Analysis

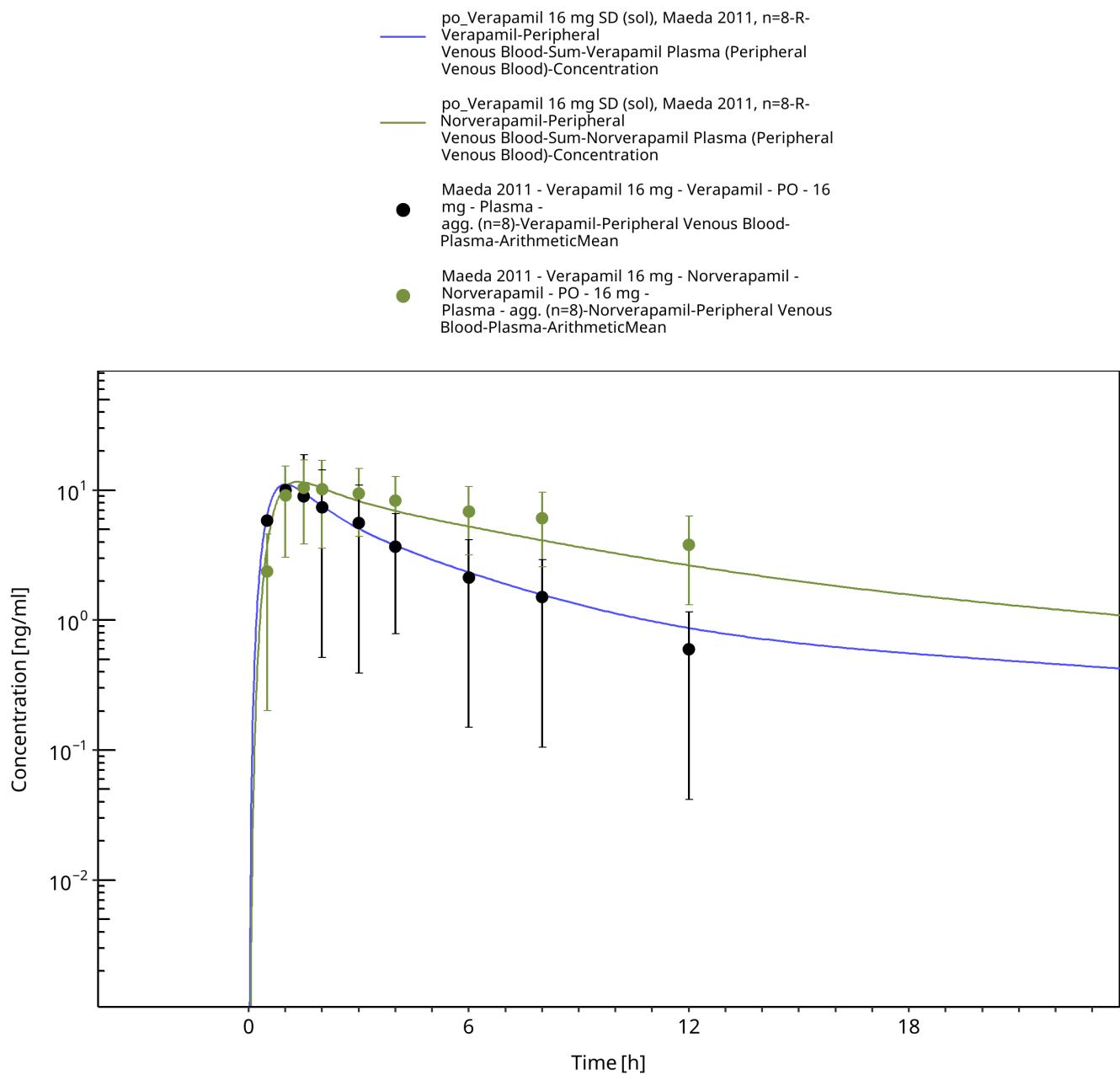


Figure 3-5: Time Profile Analysis

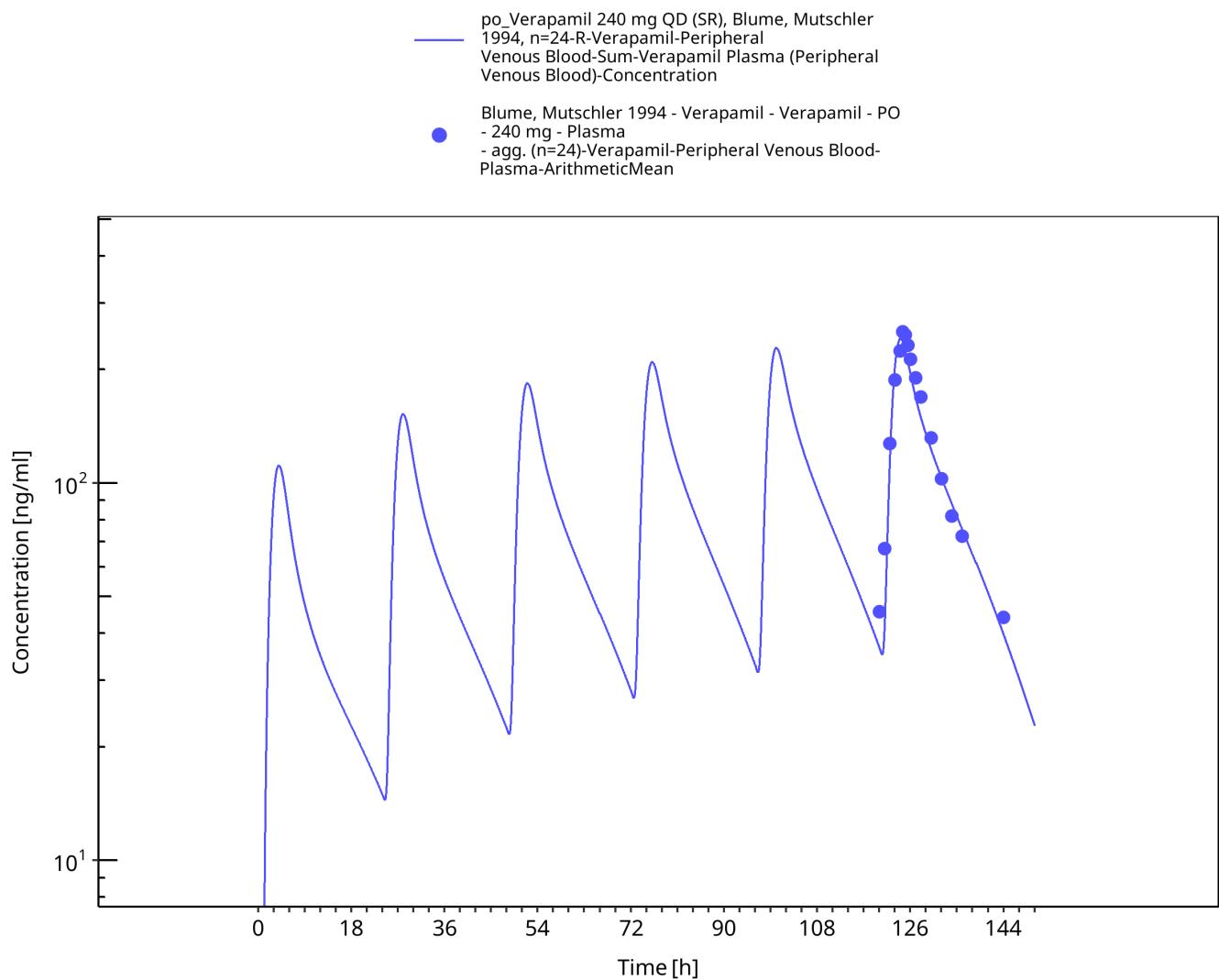


Figure 3-6: Time Profile Analysis

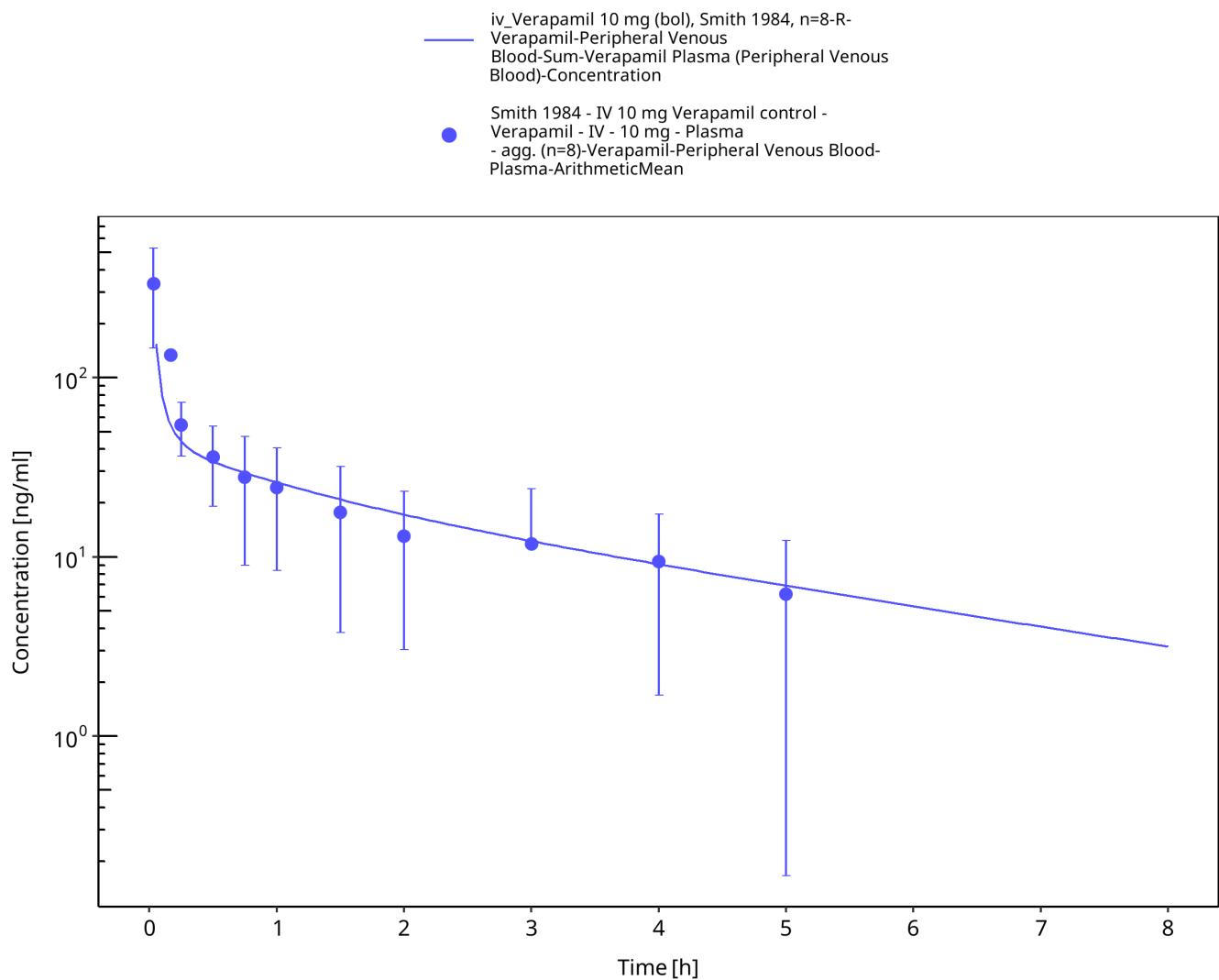


Figure 3-7: Time Profile Analysis

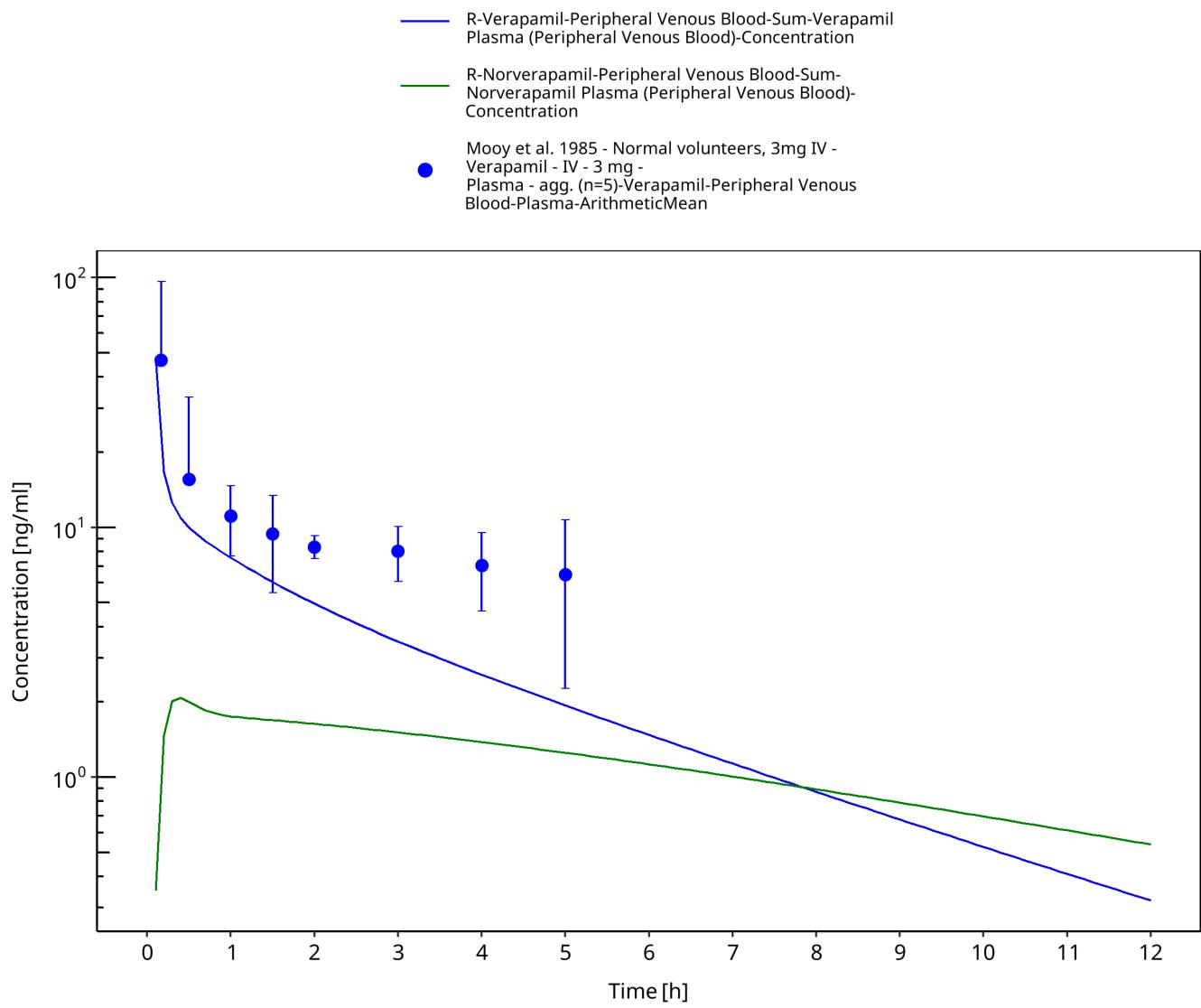


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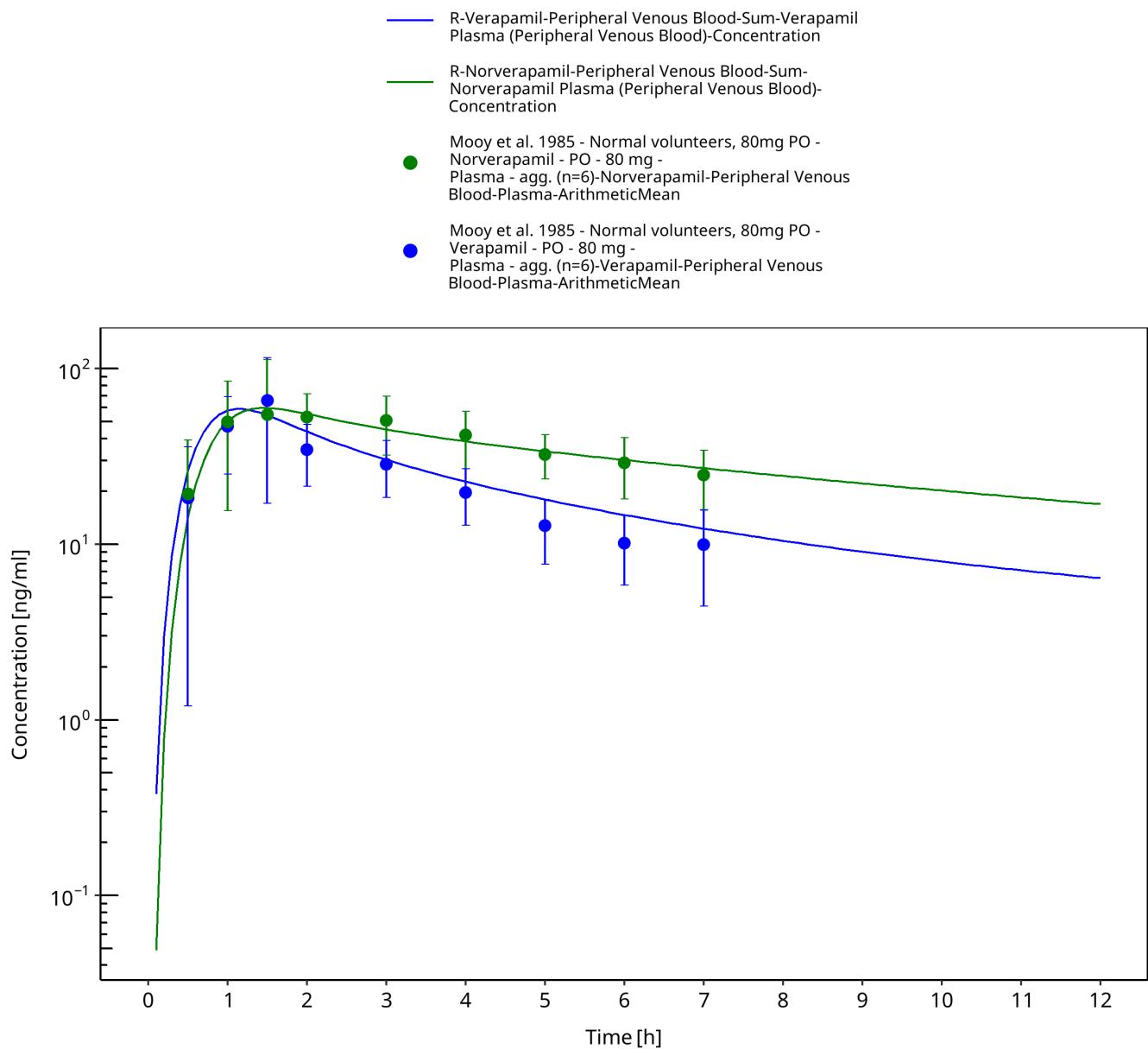


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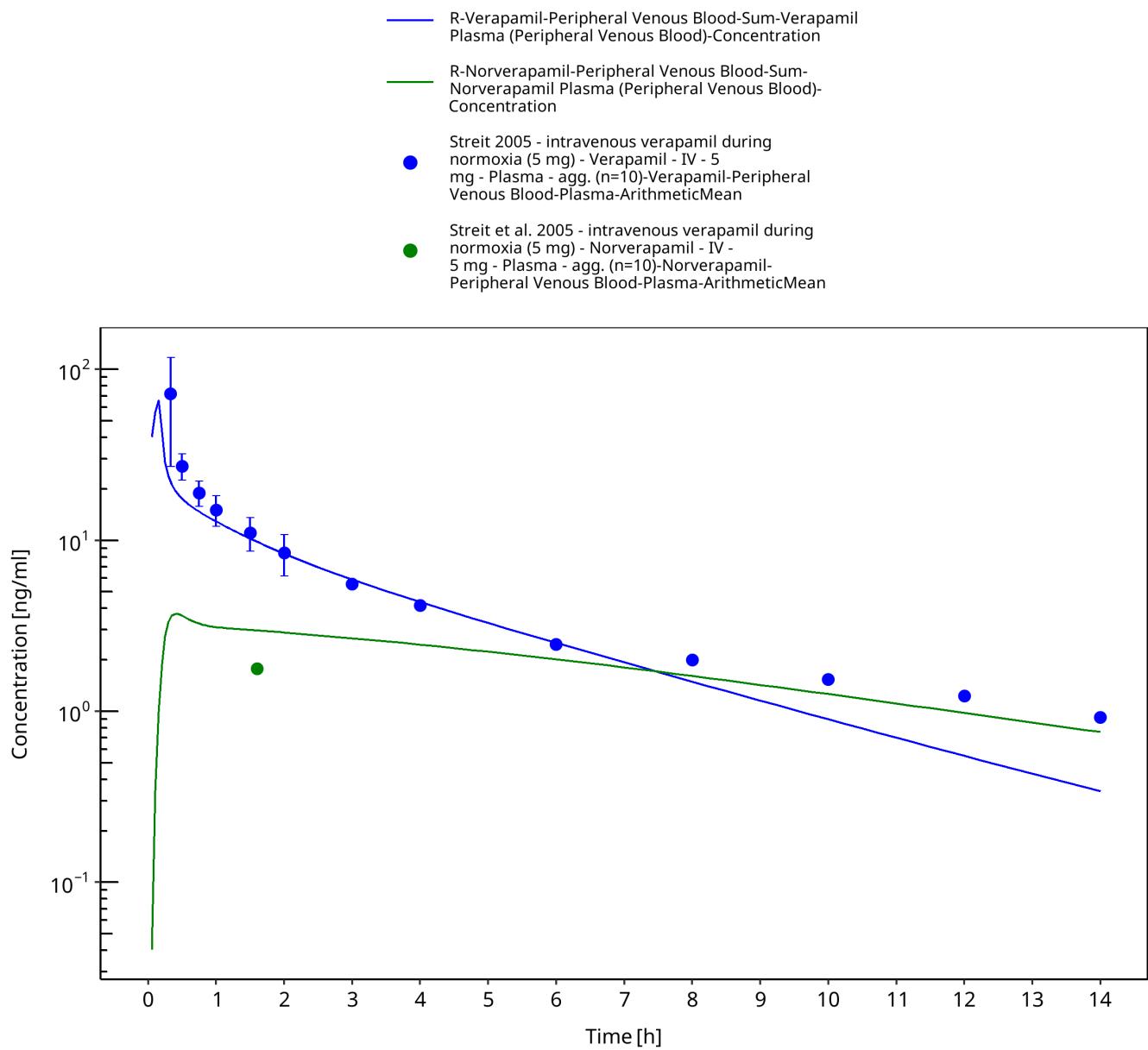


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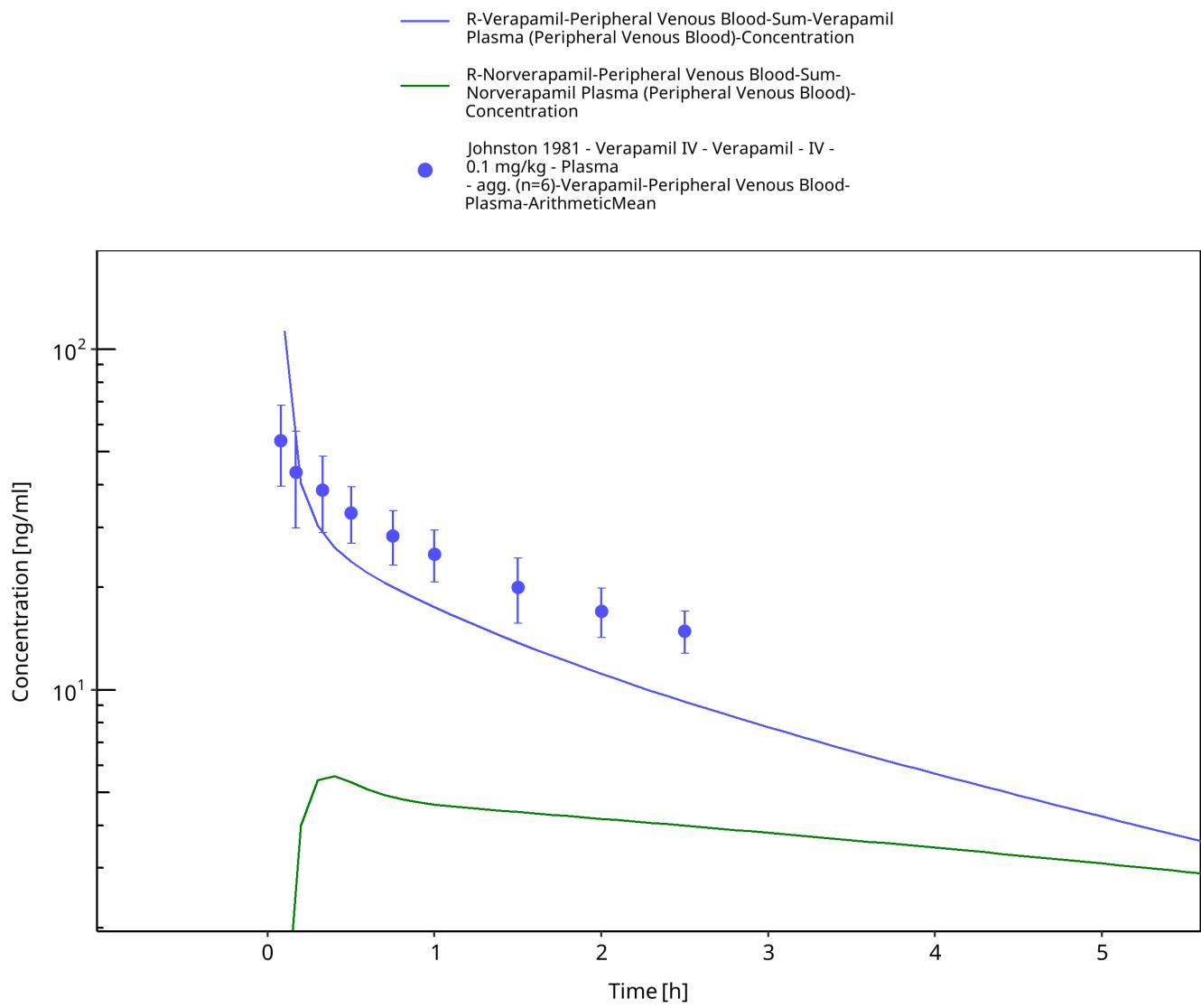


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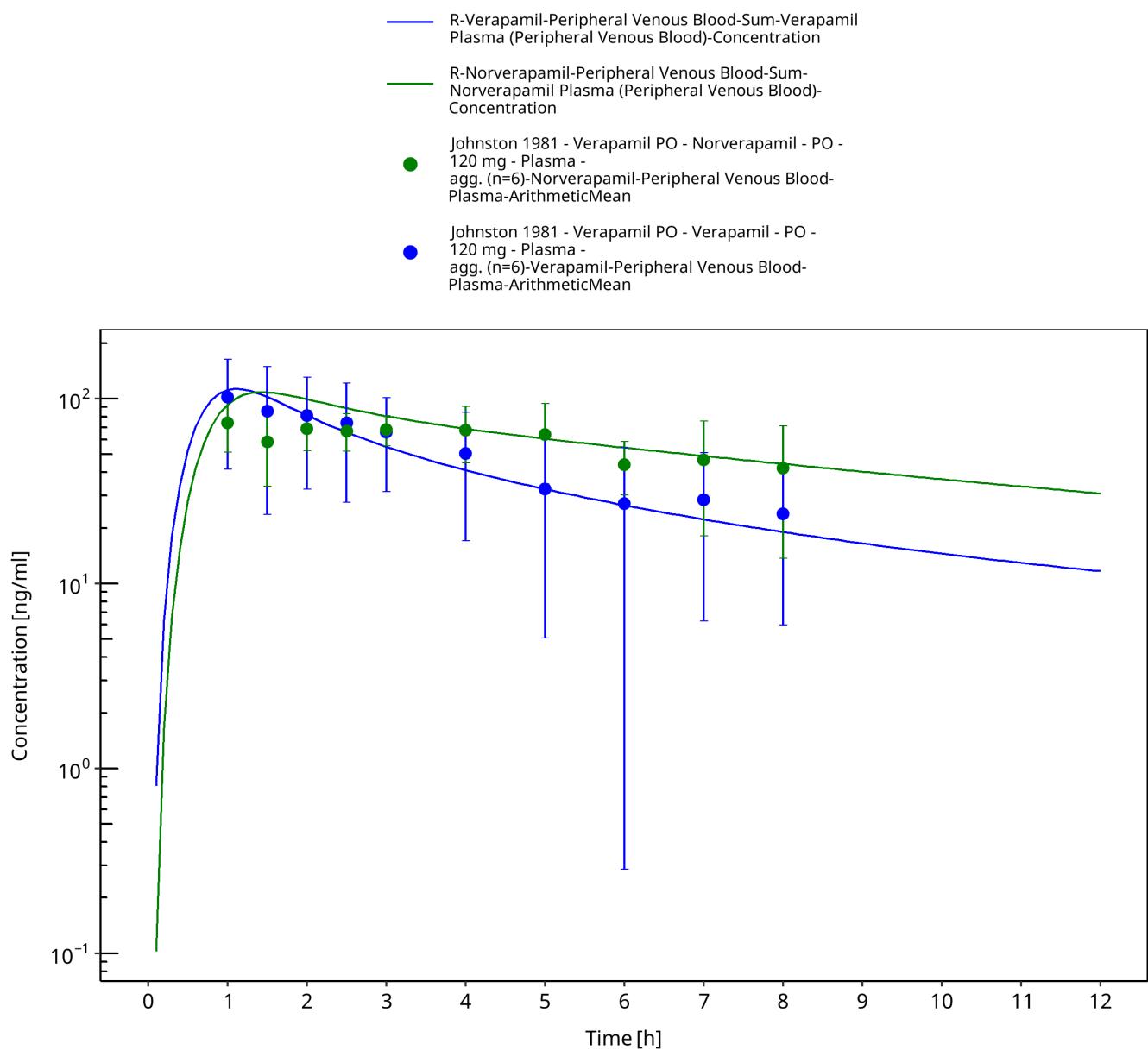


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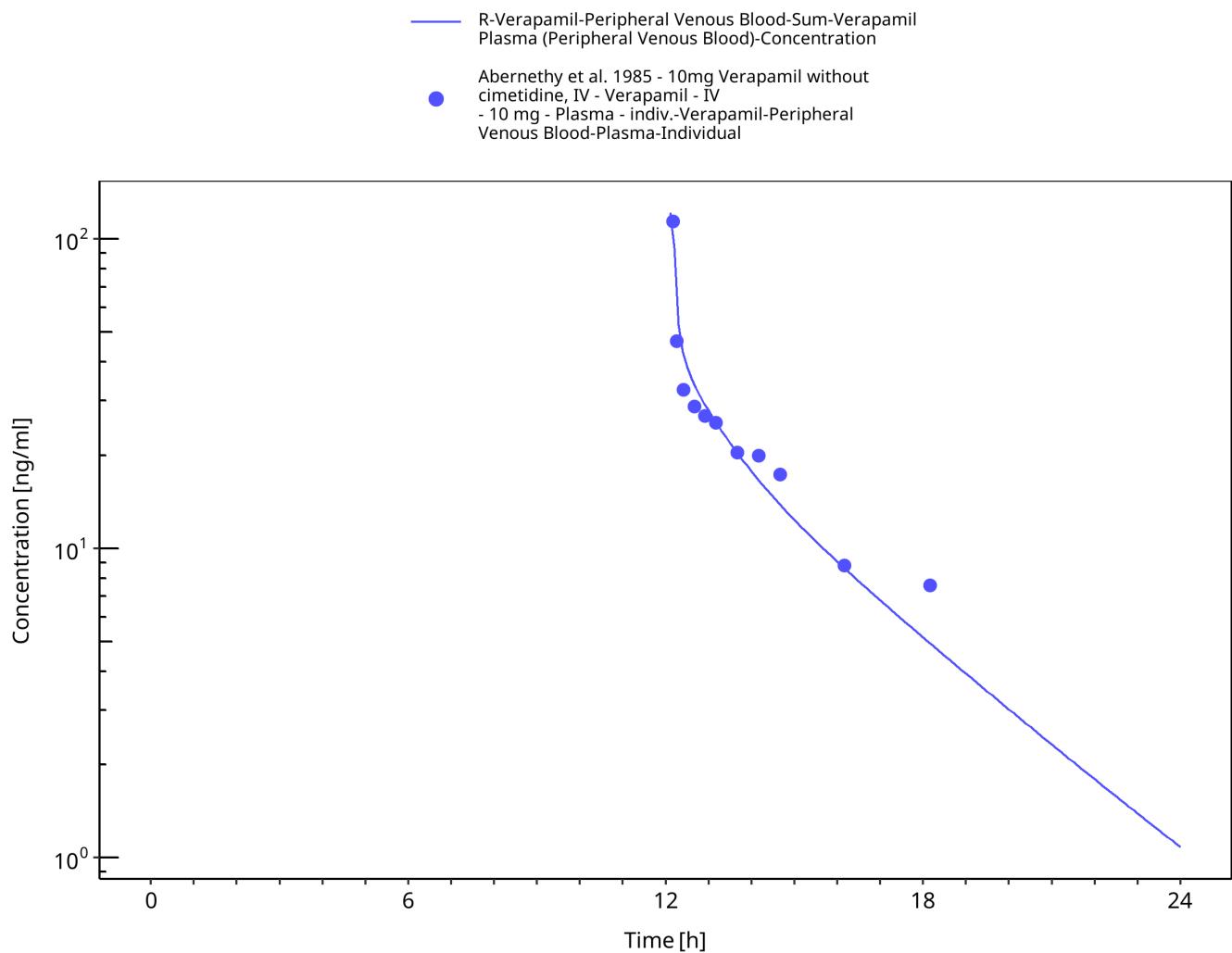


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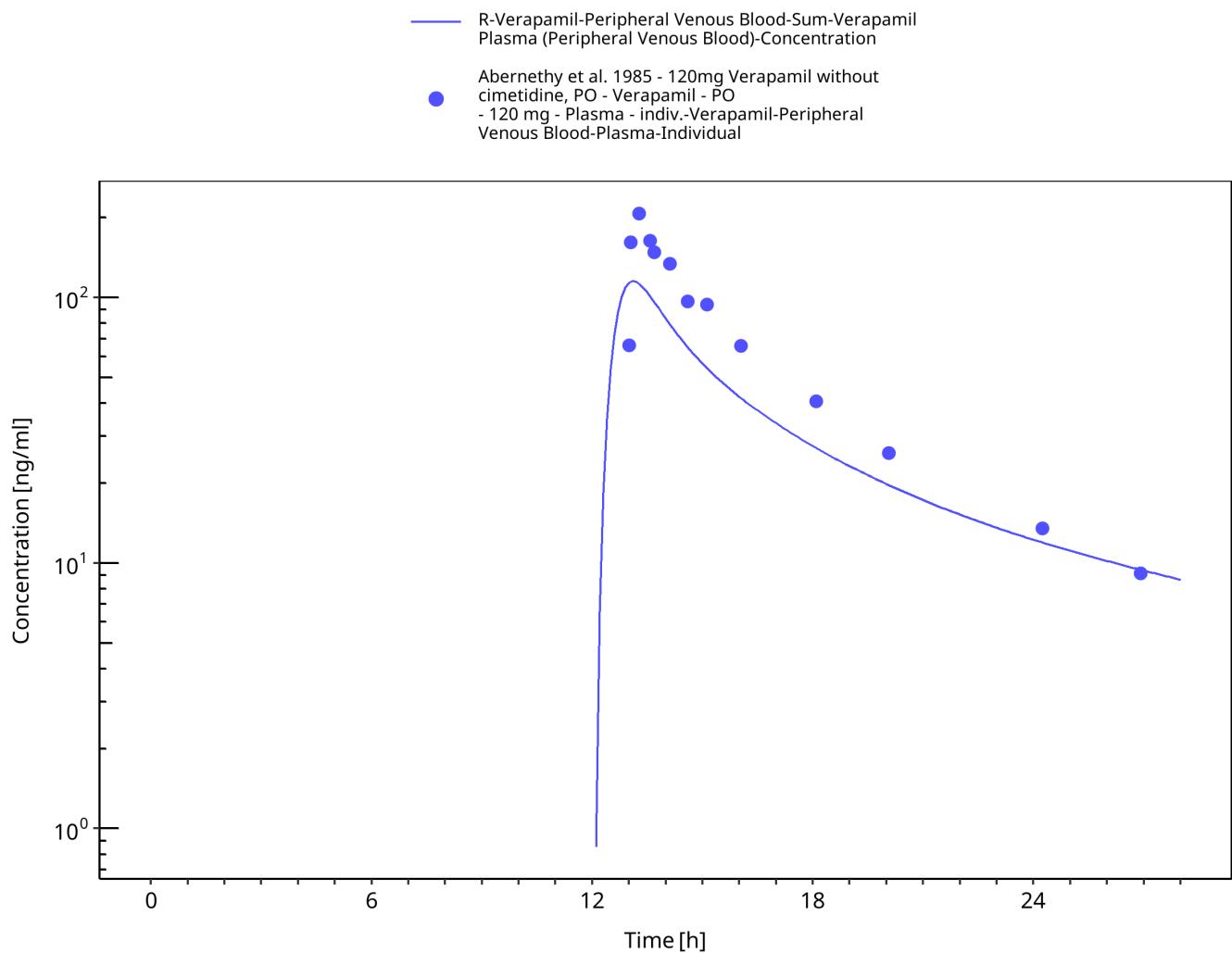


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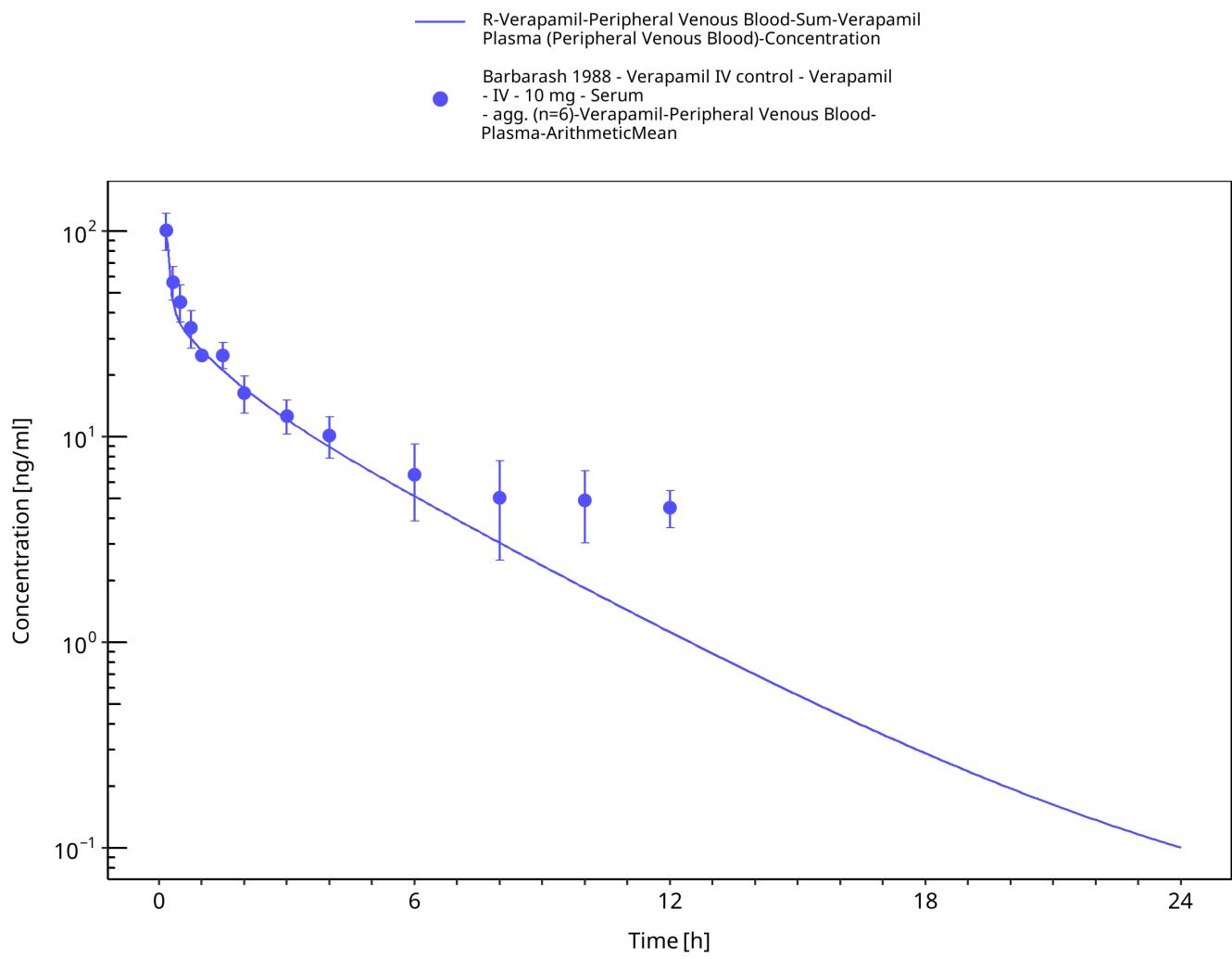


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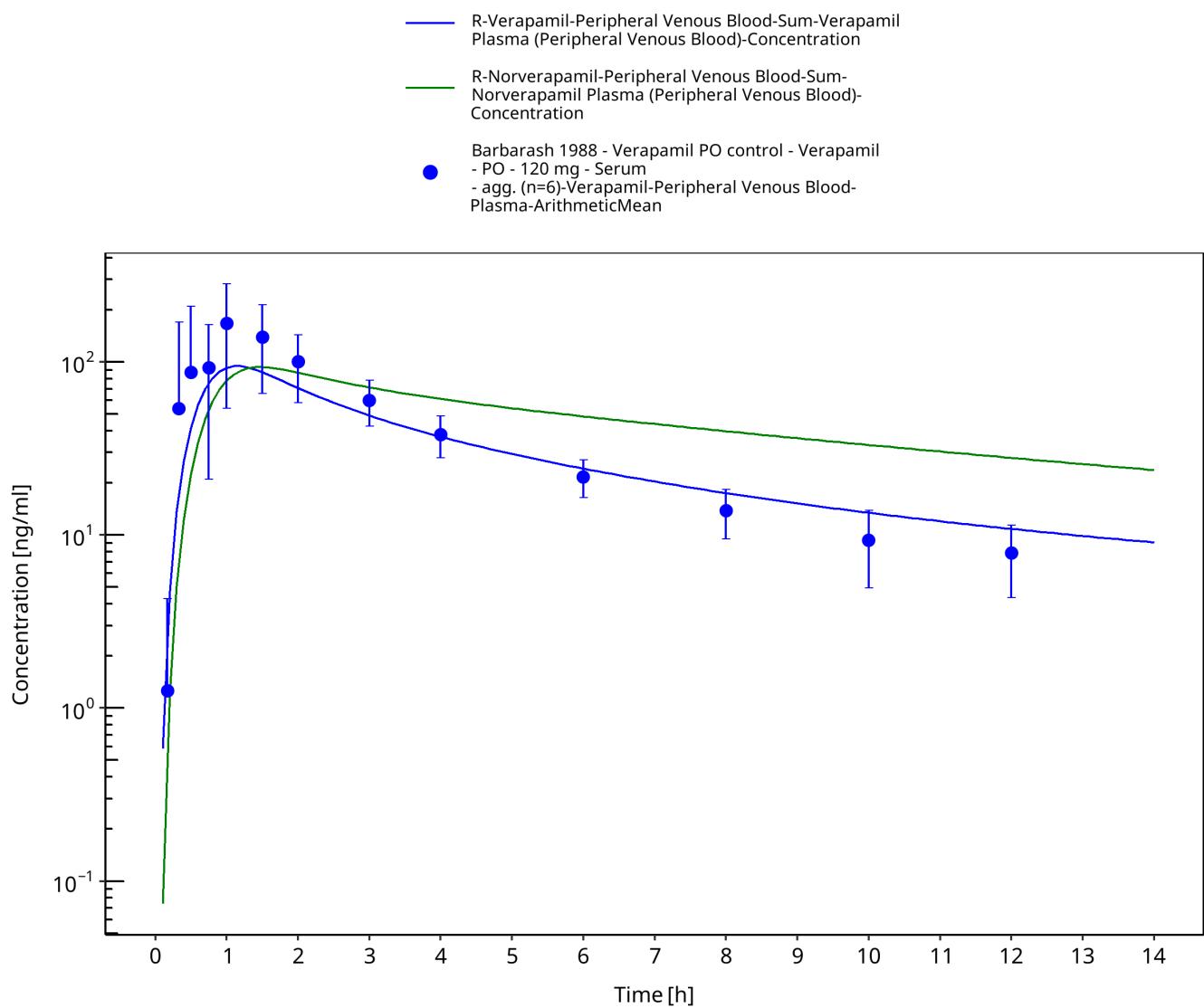


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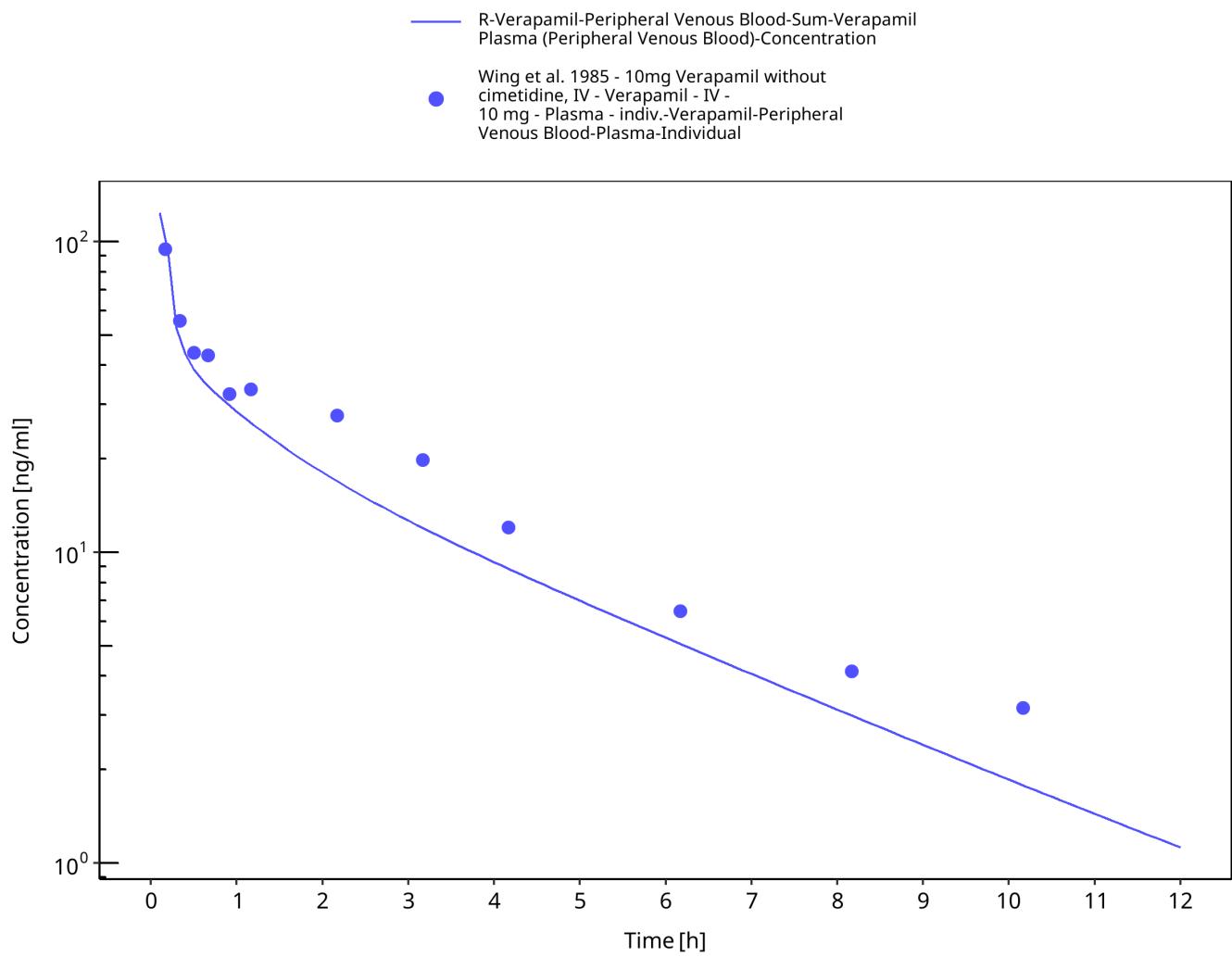


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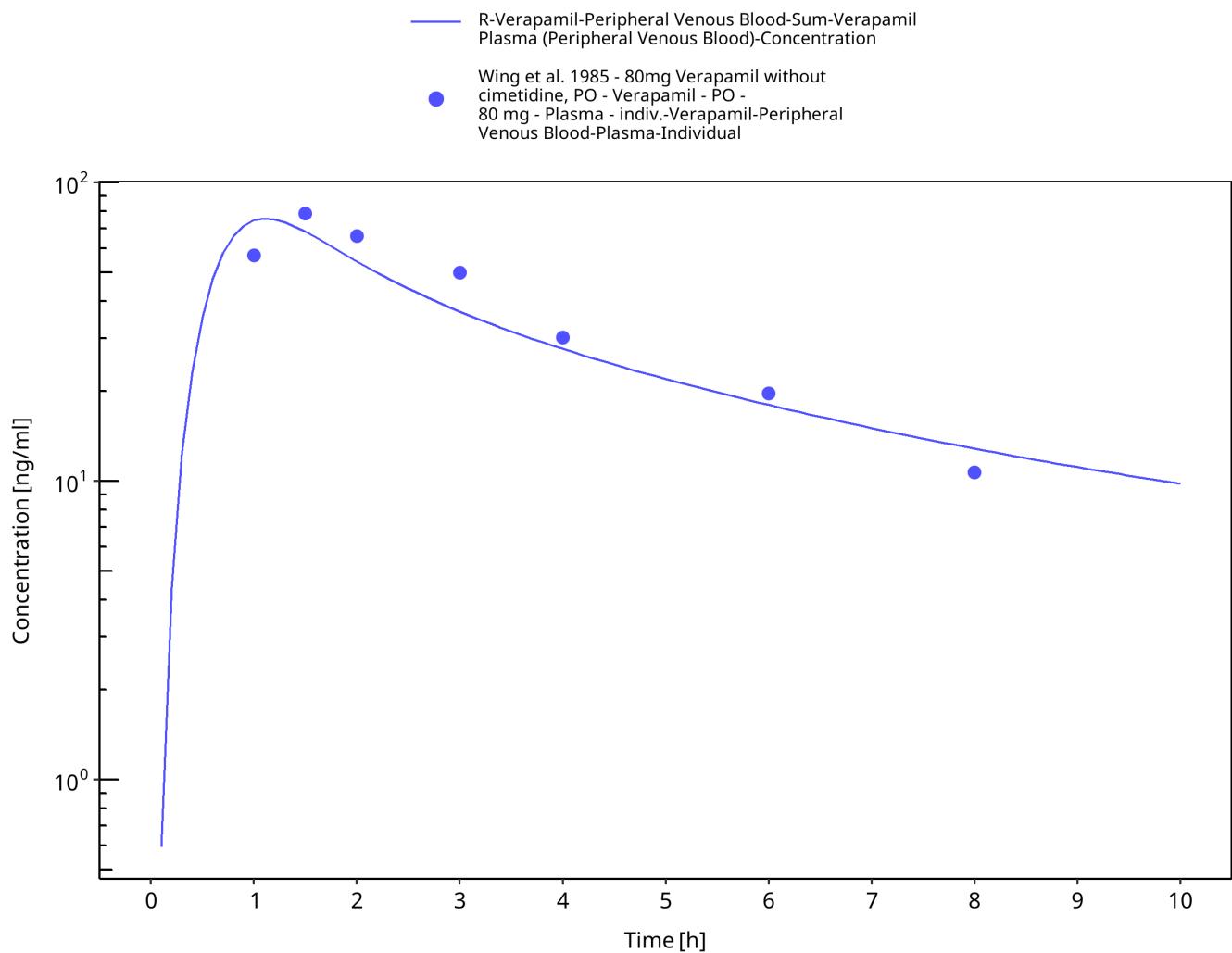


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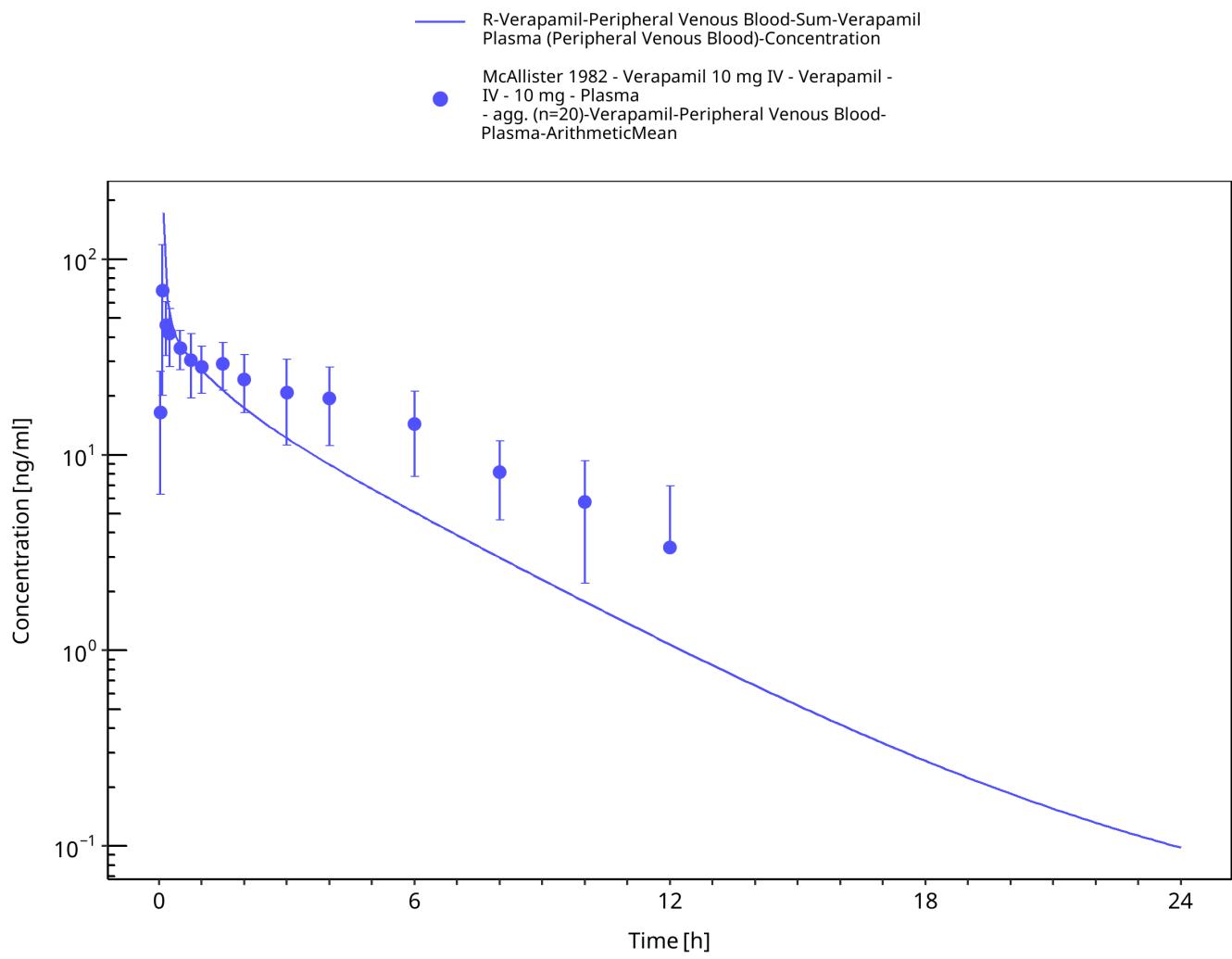


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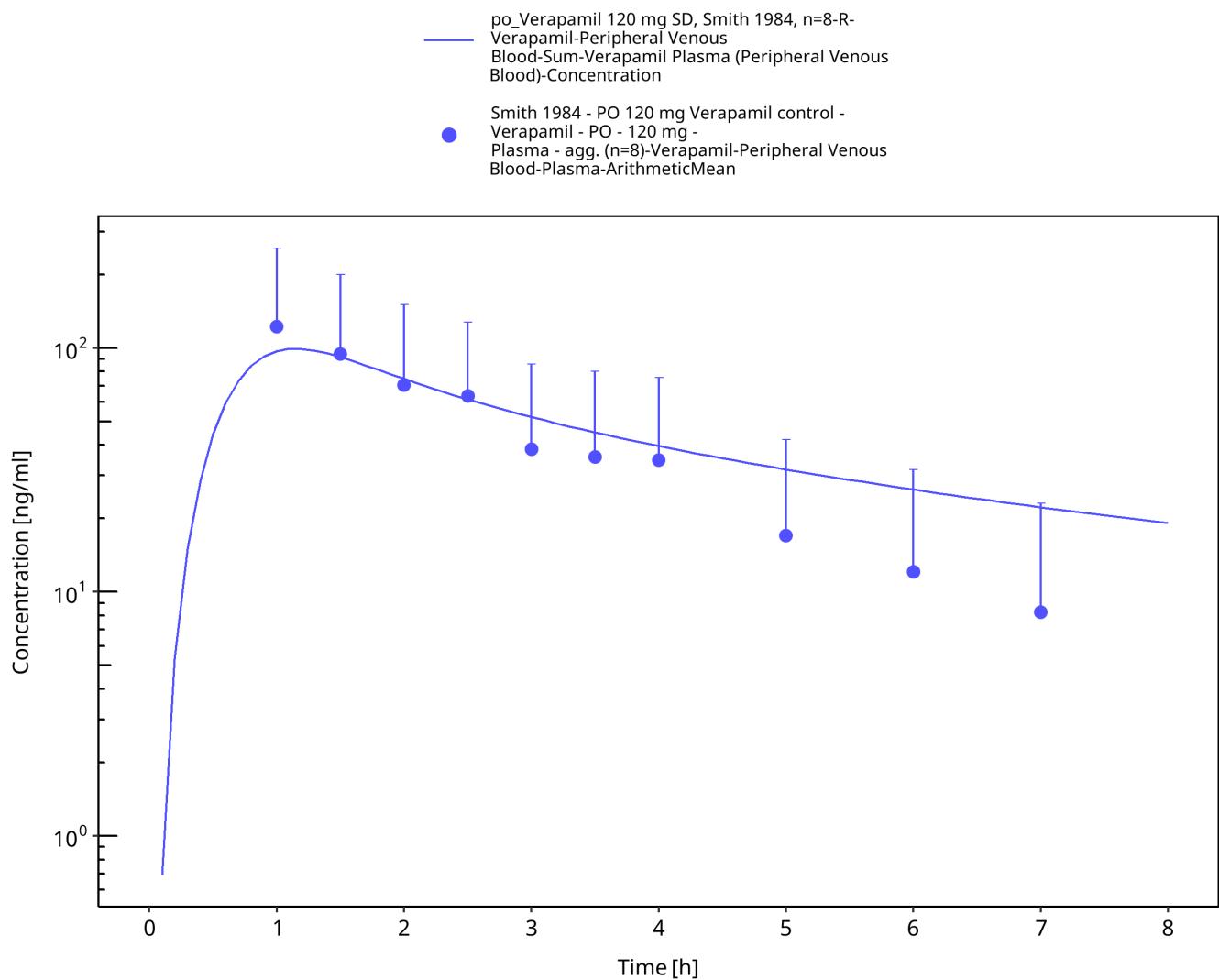


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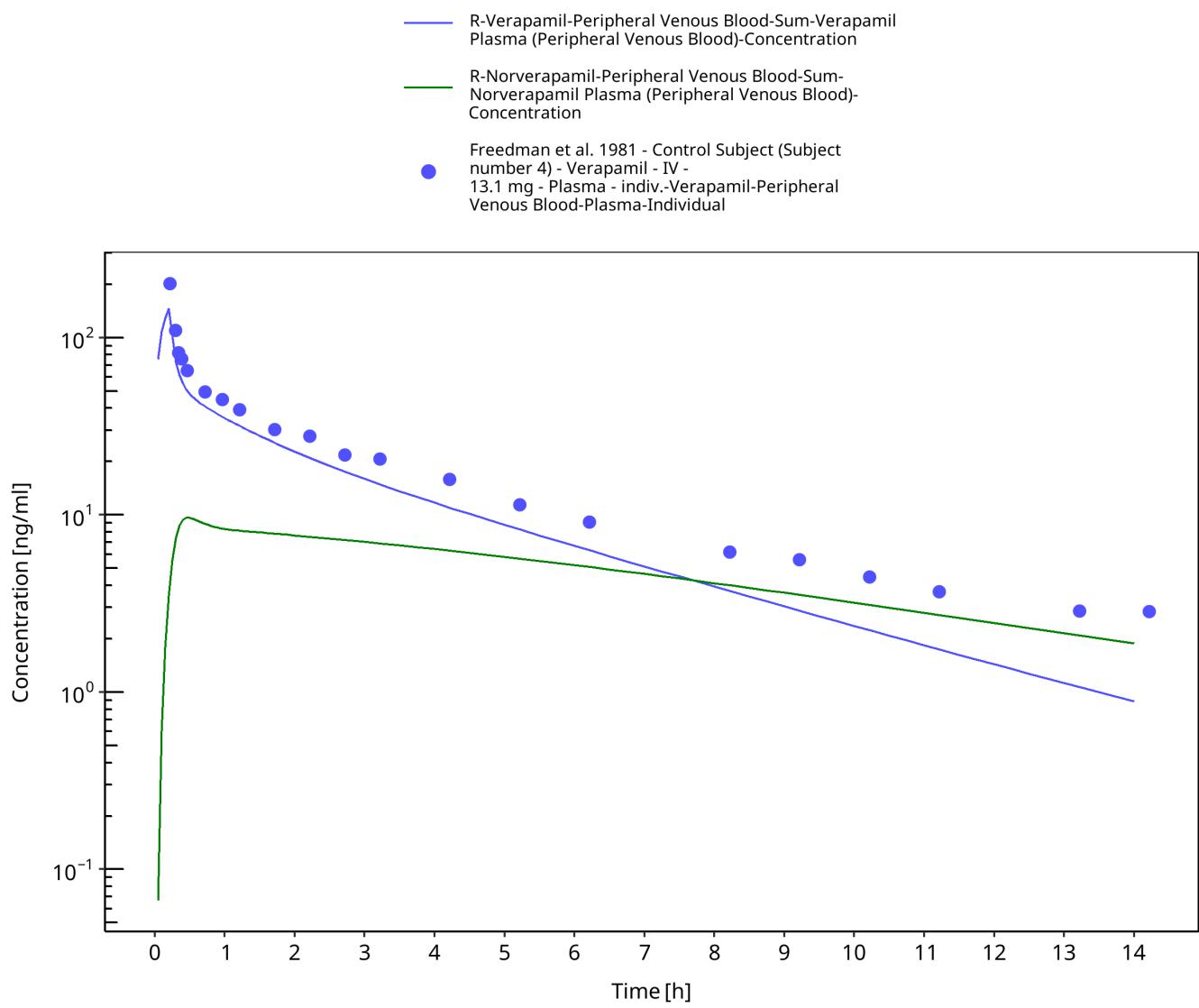


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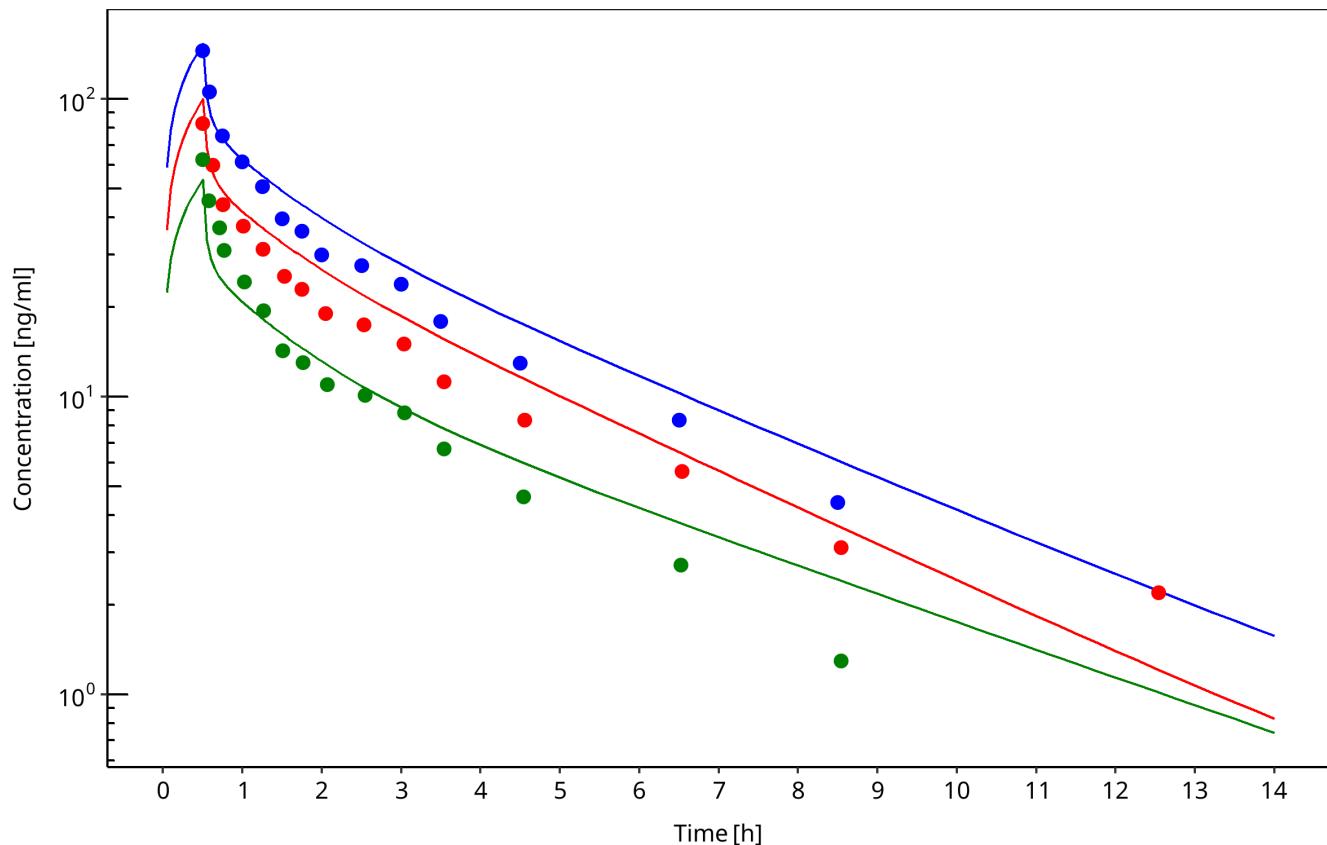
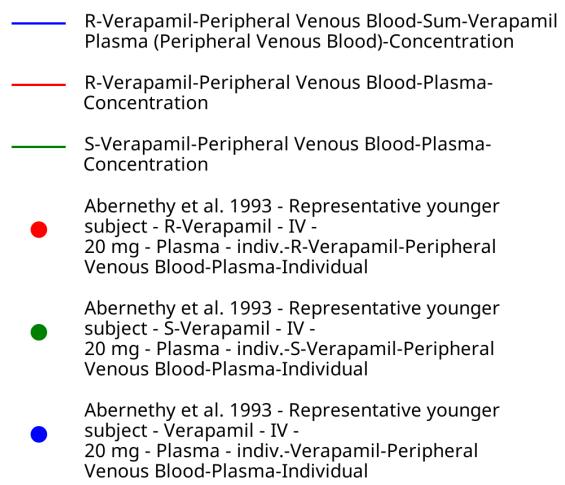


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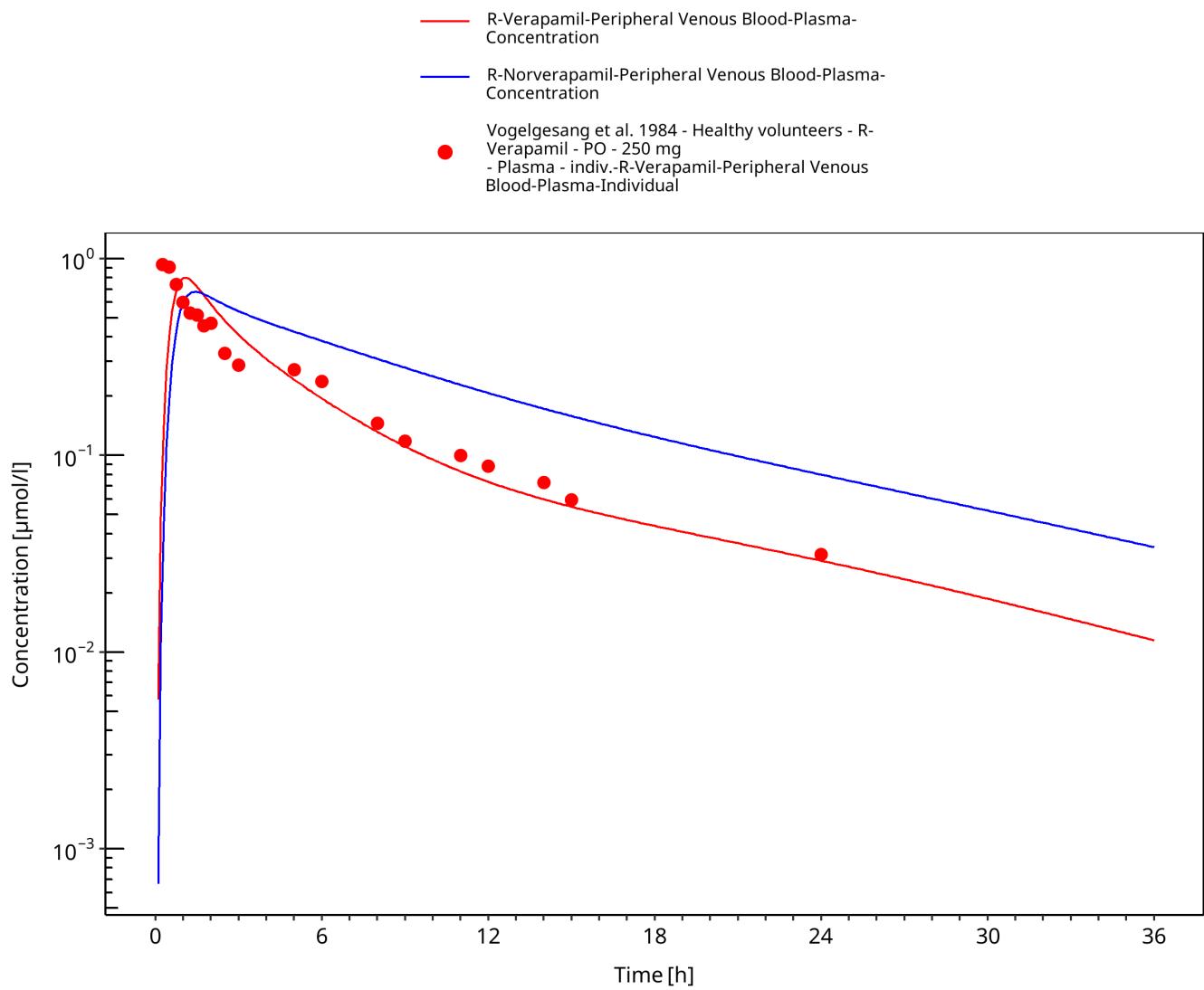


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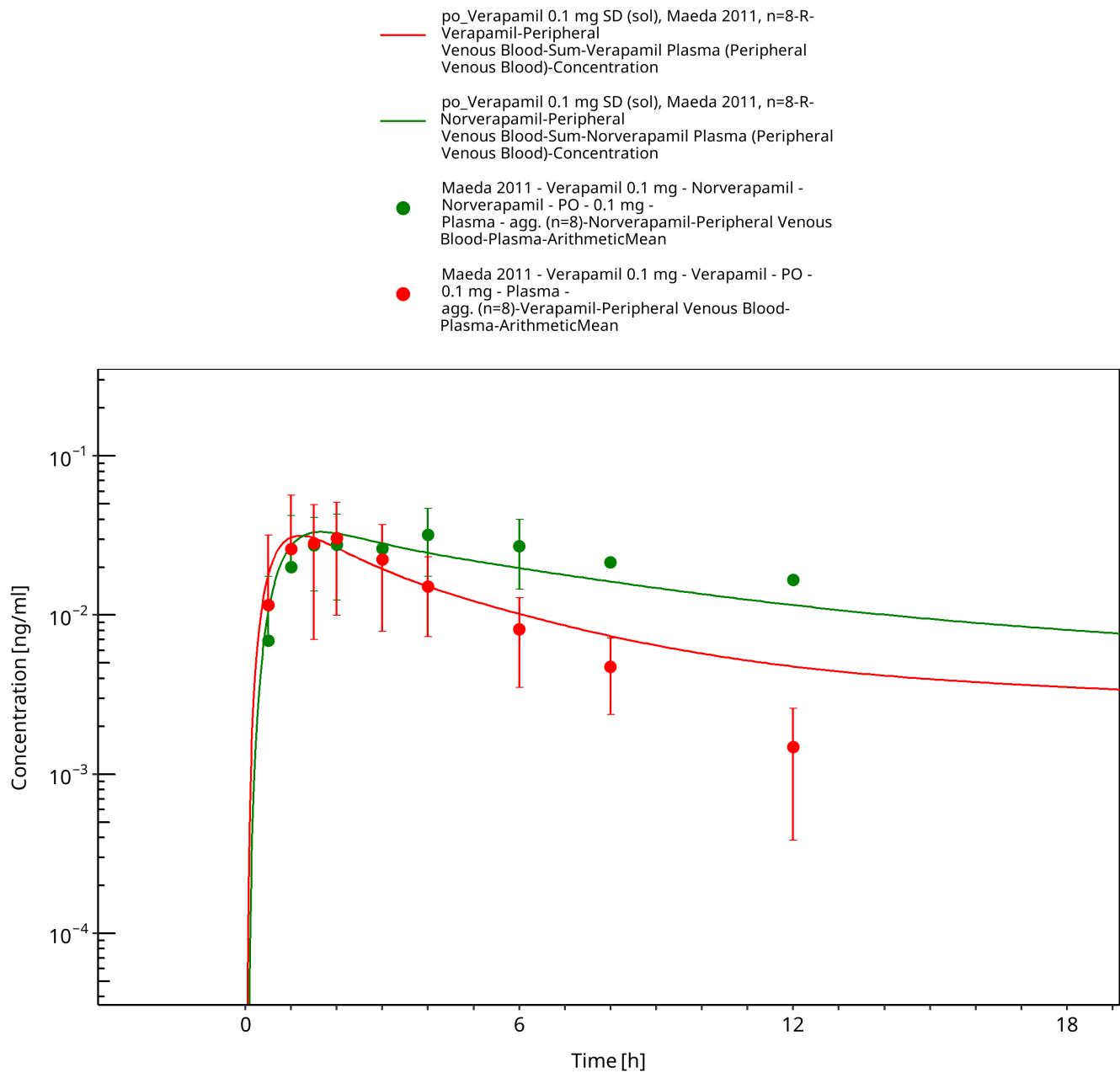


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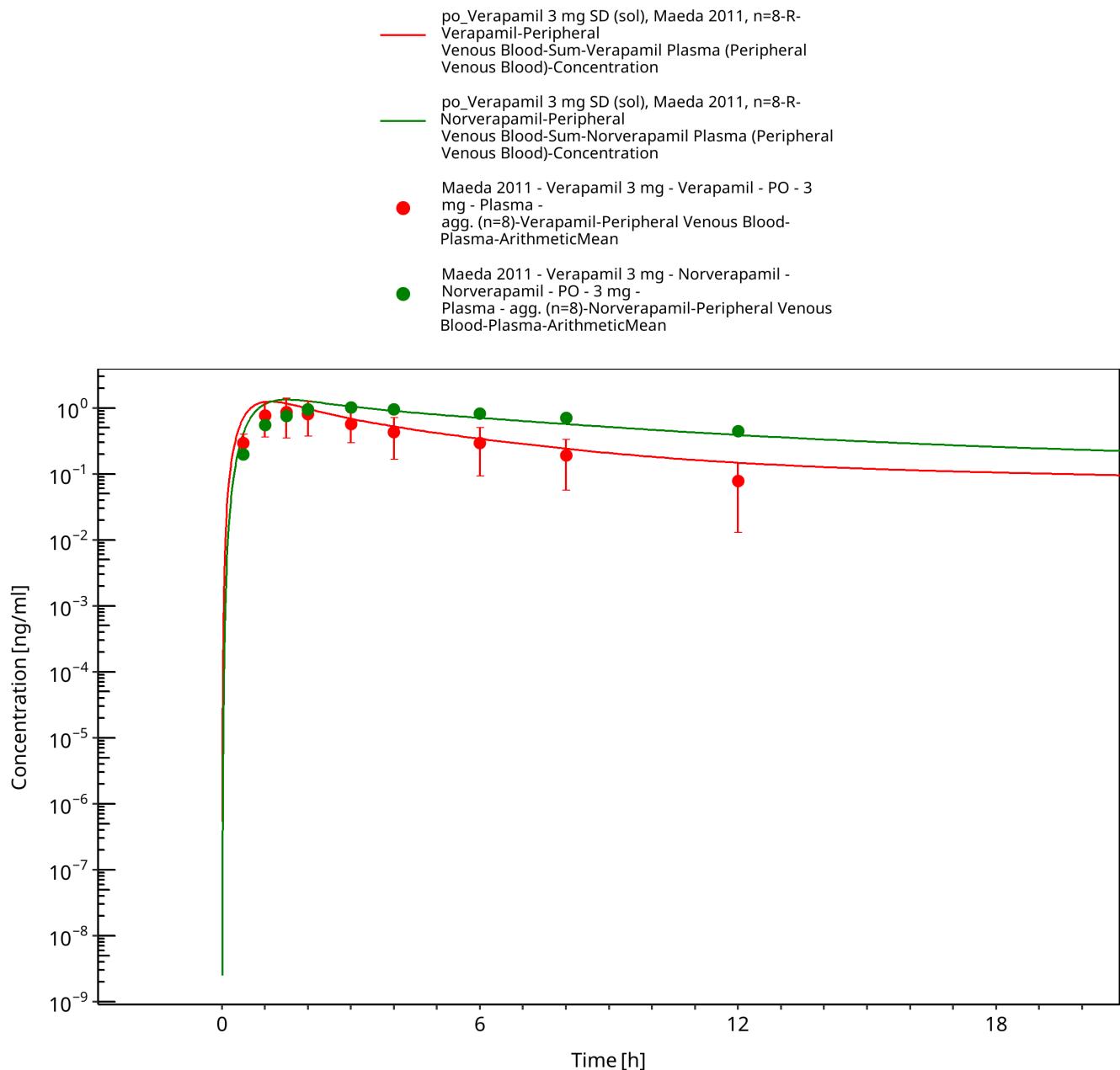


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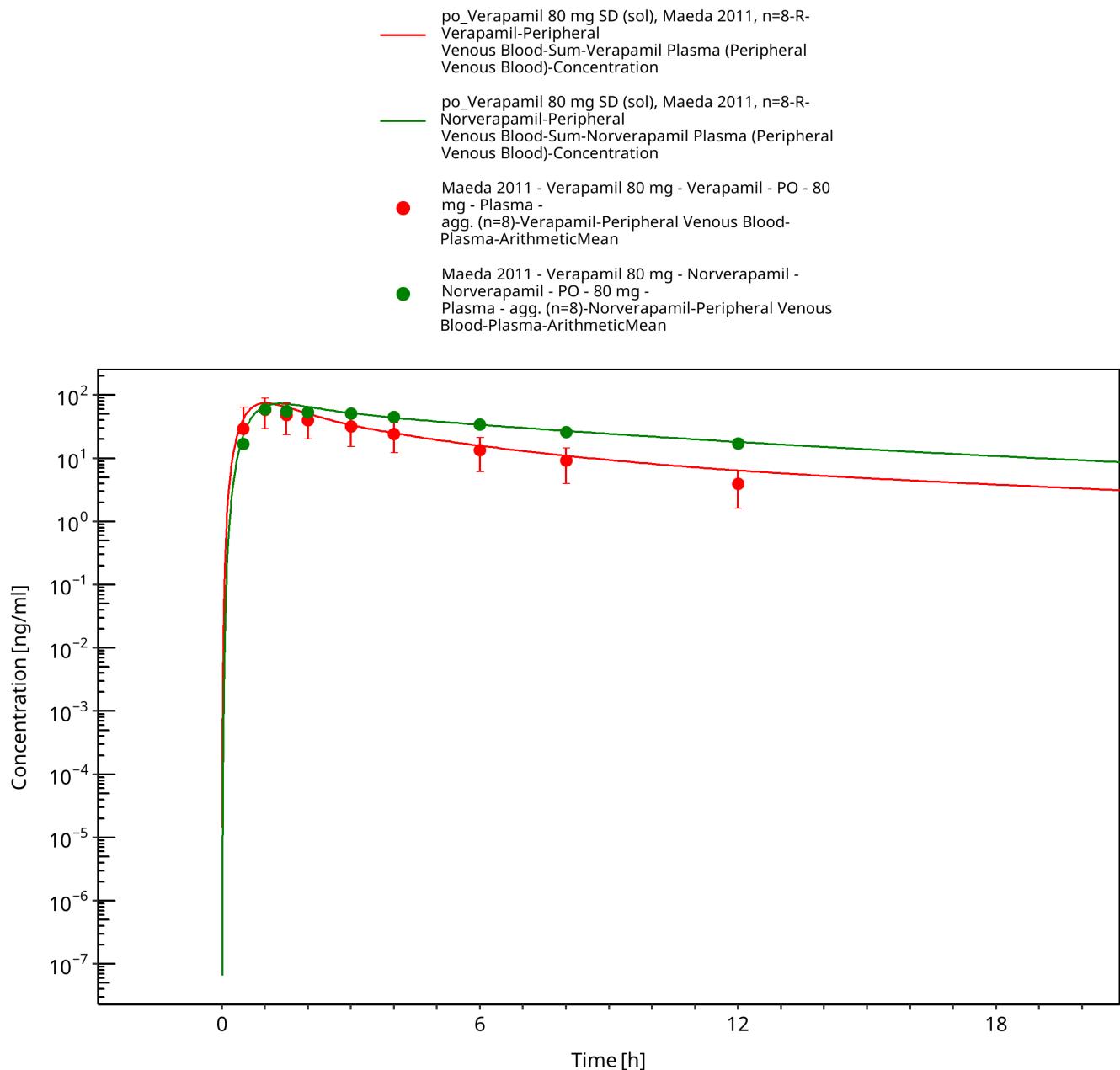


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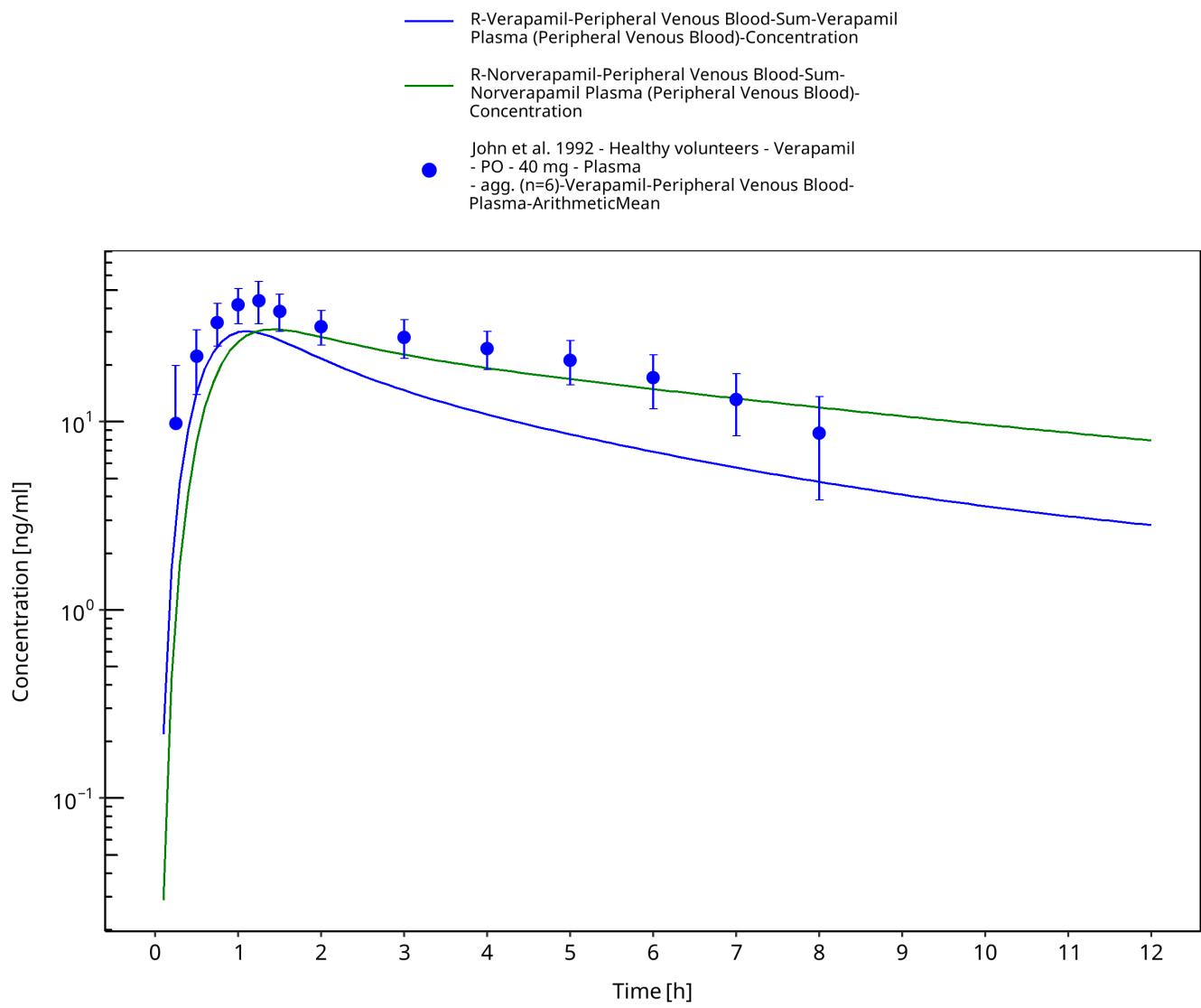


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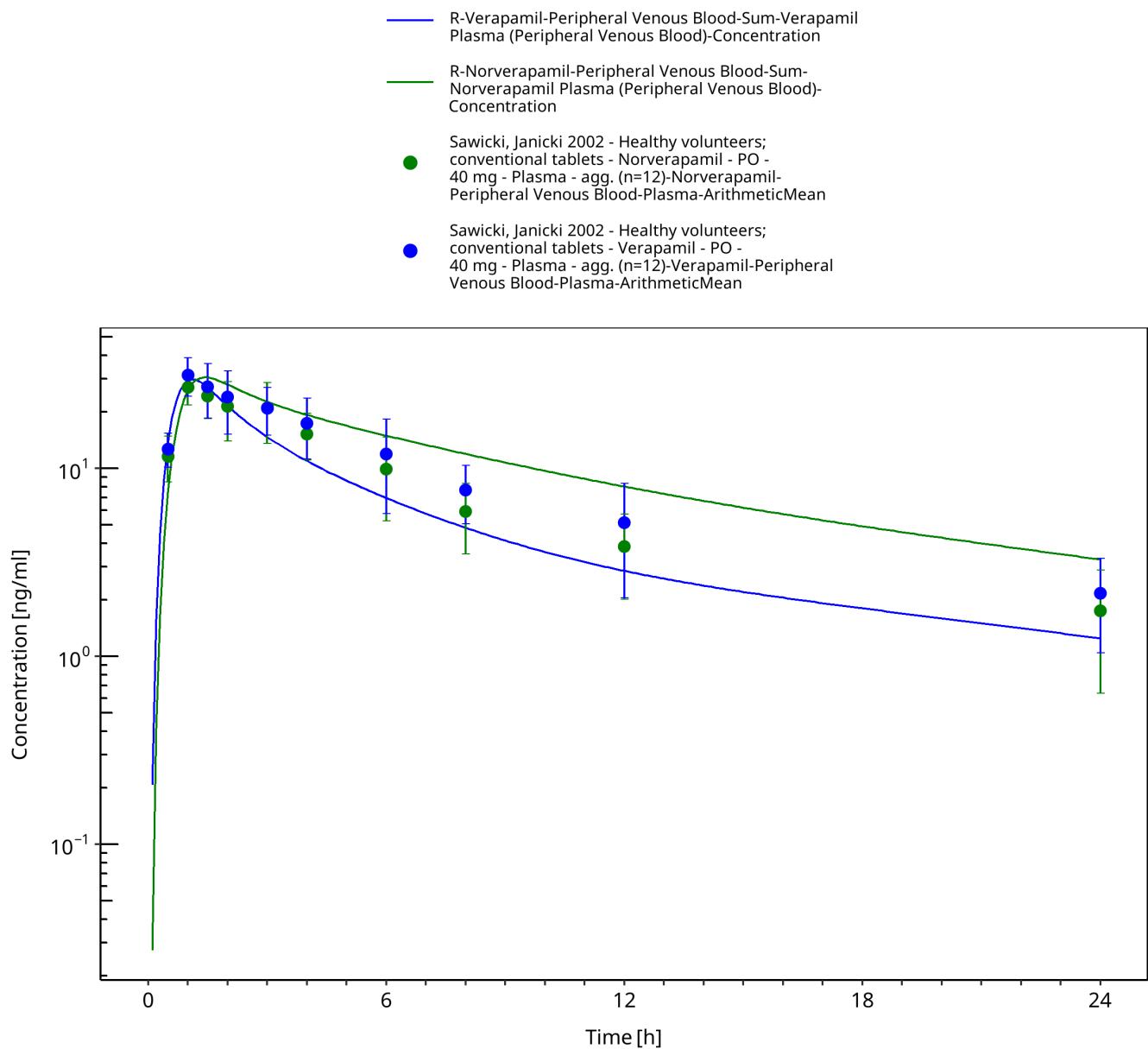


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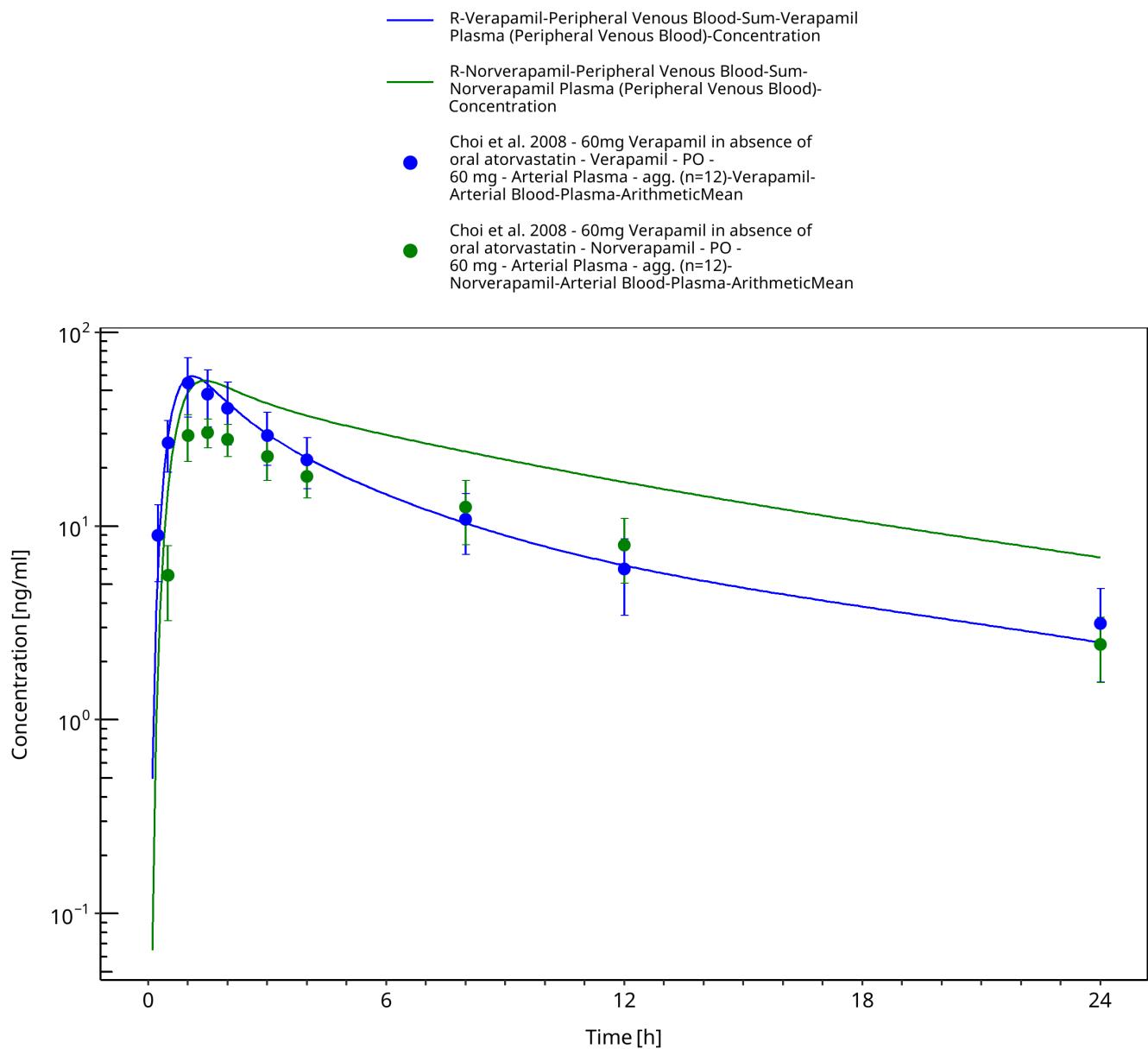


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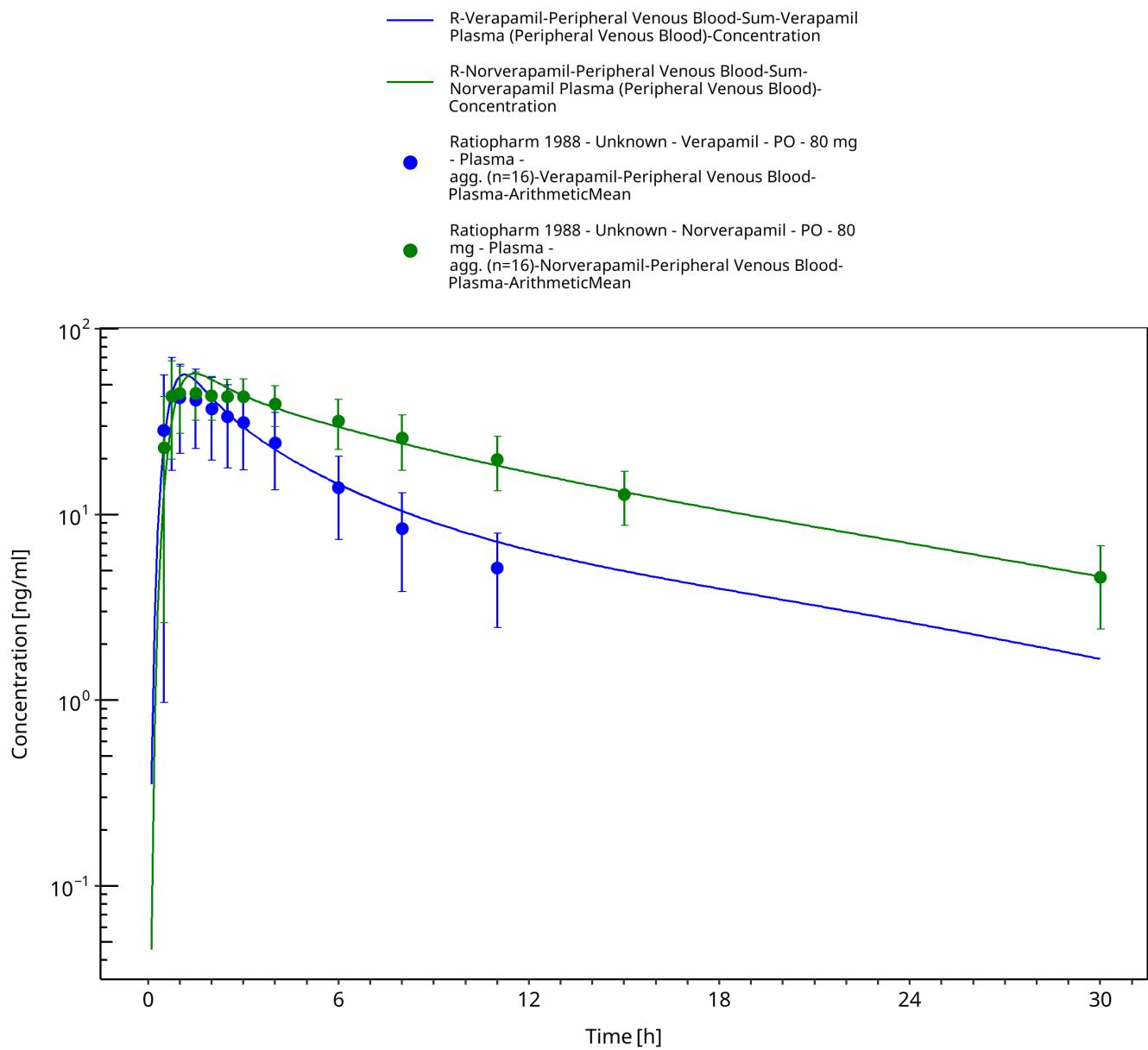


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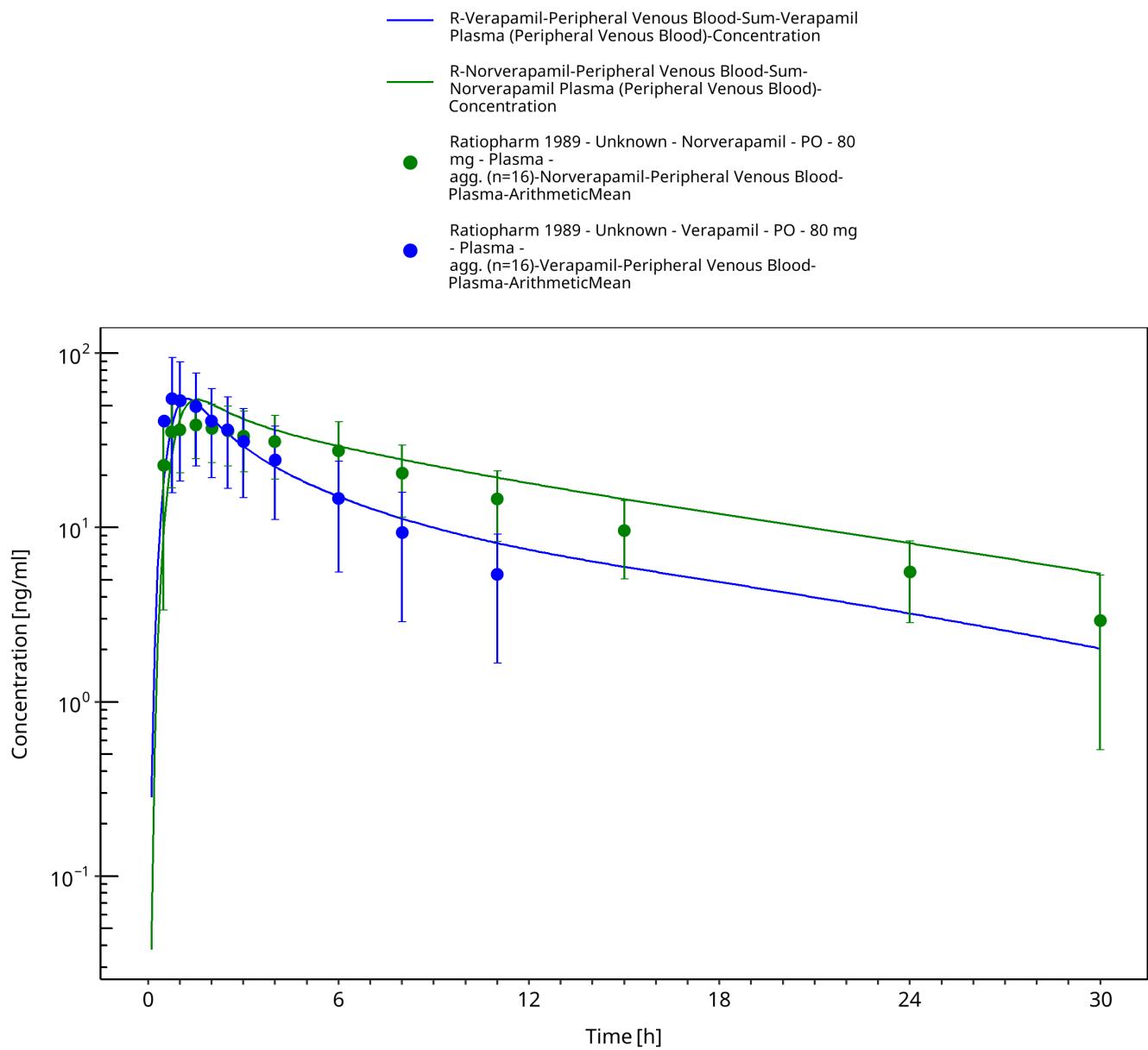


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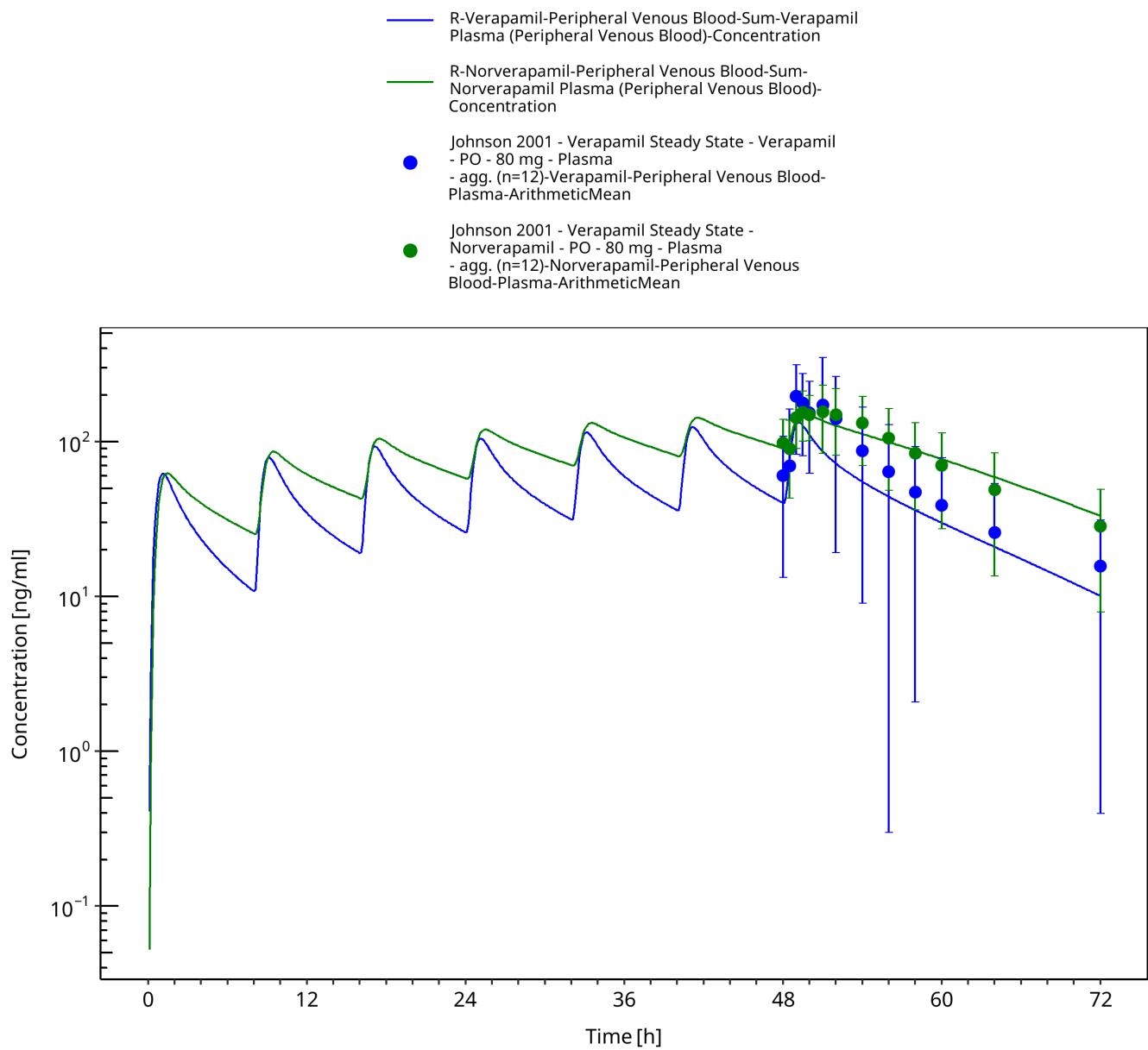


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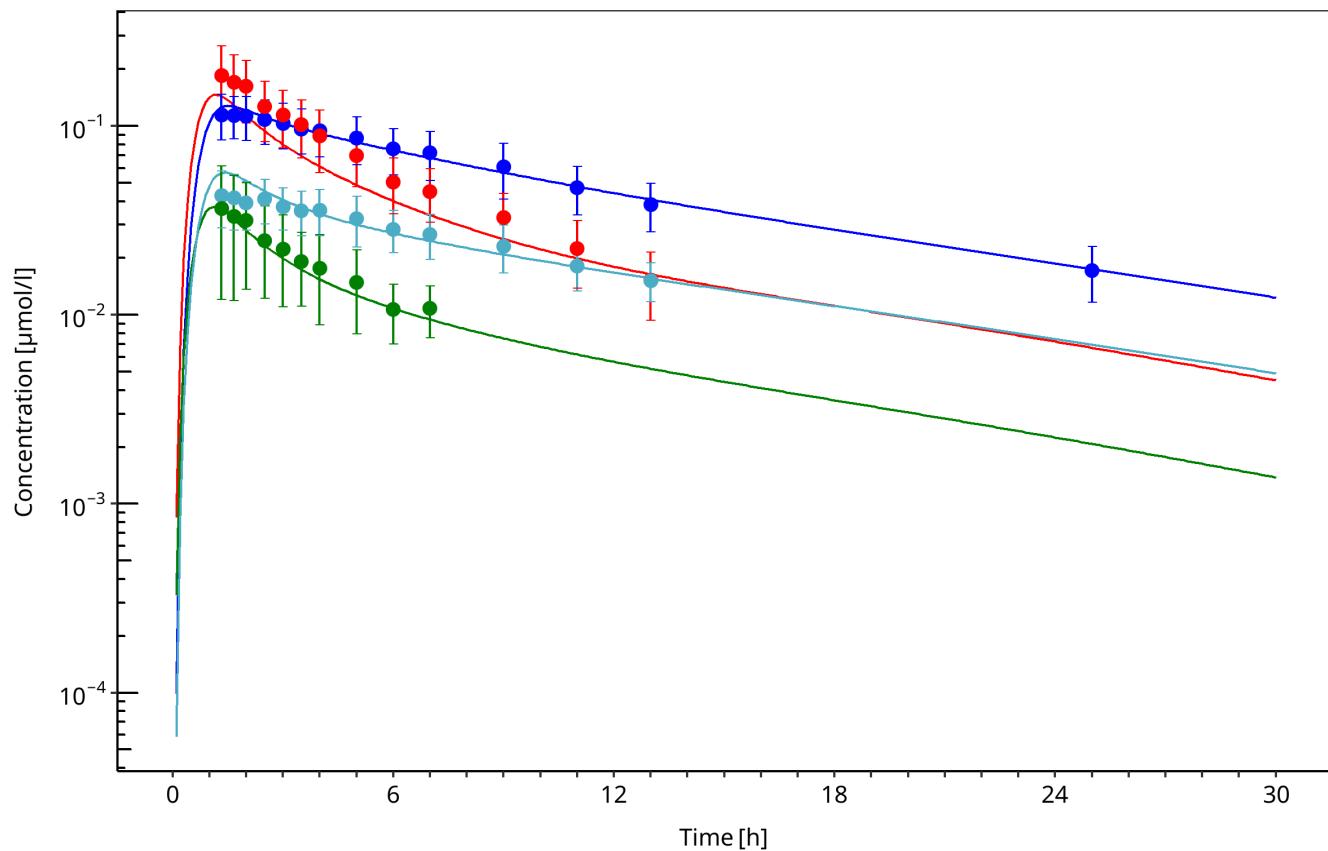
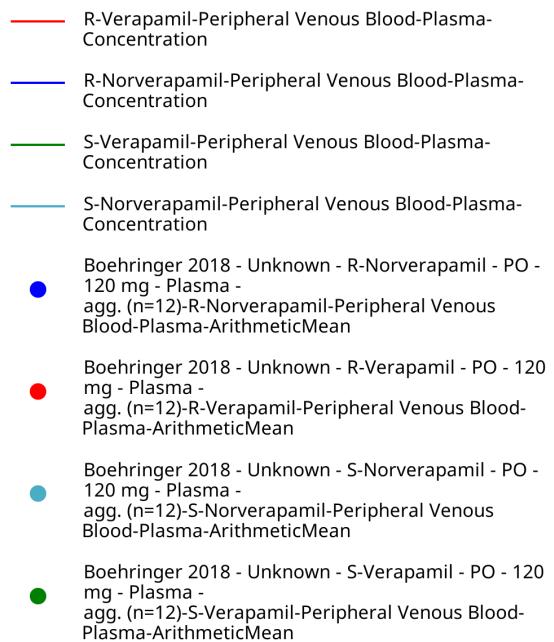


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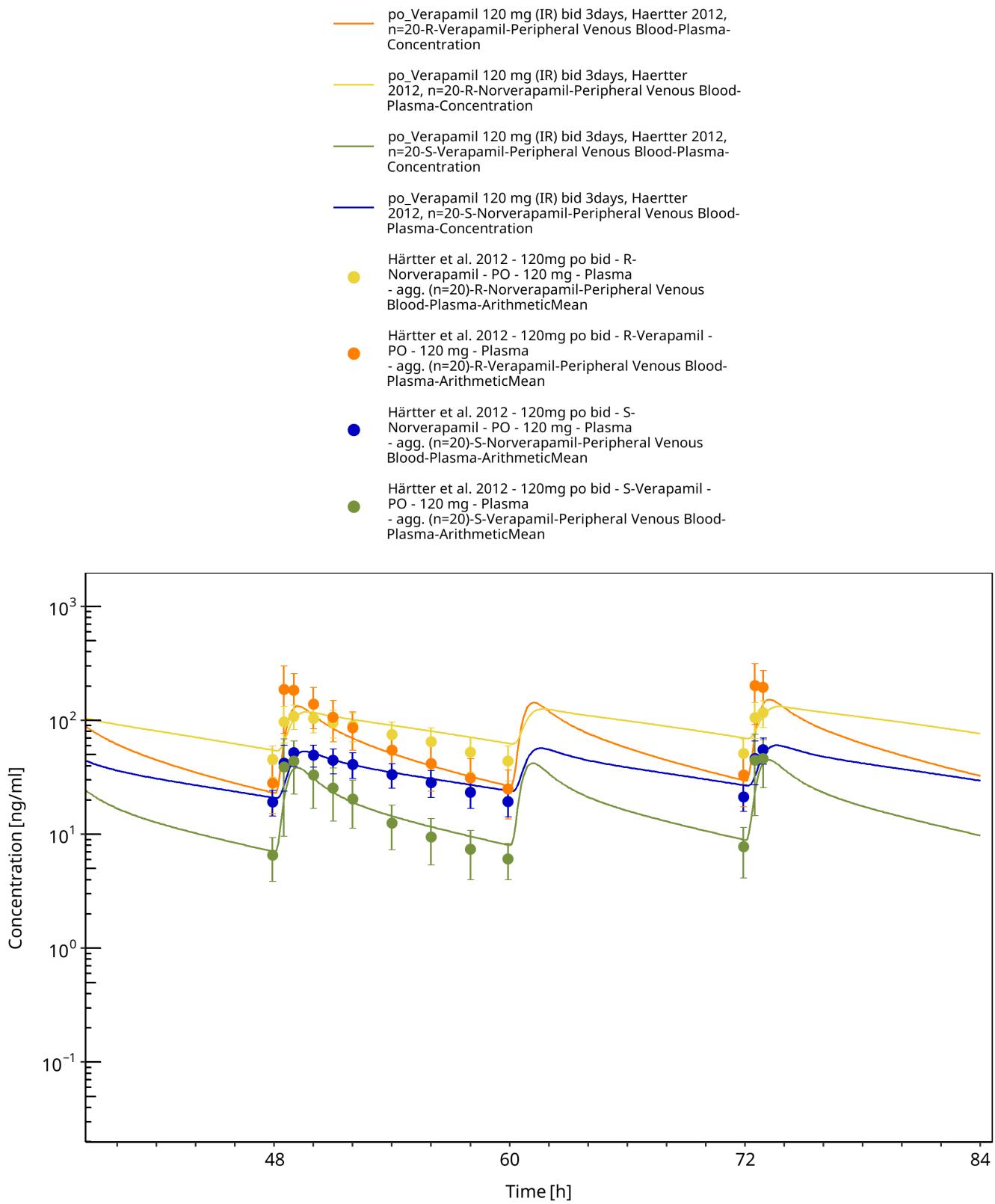


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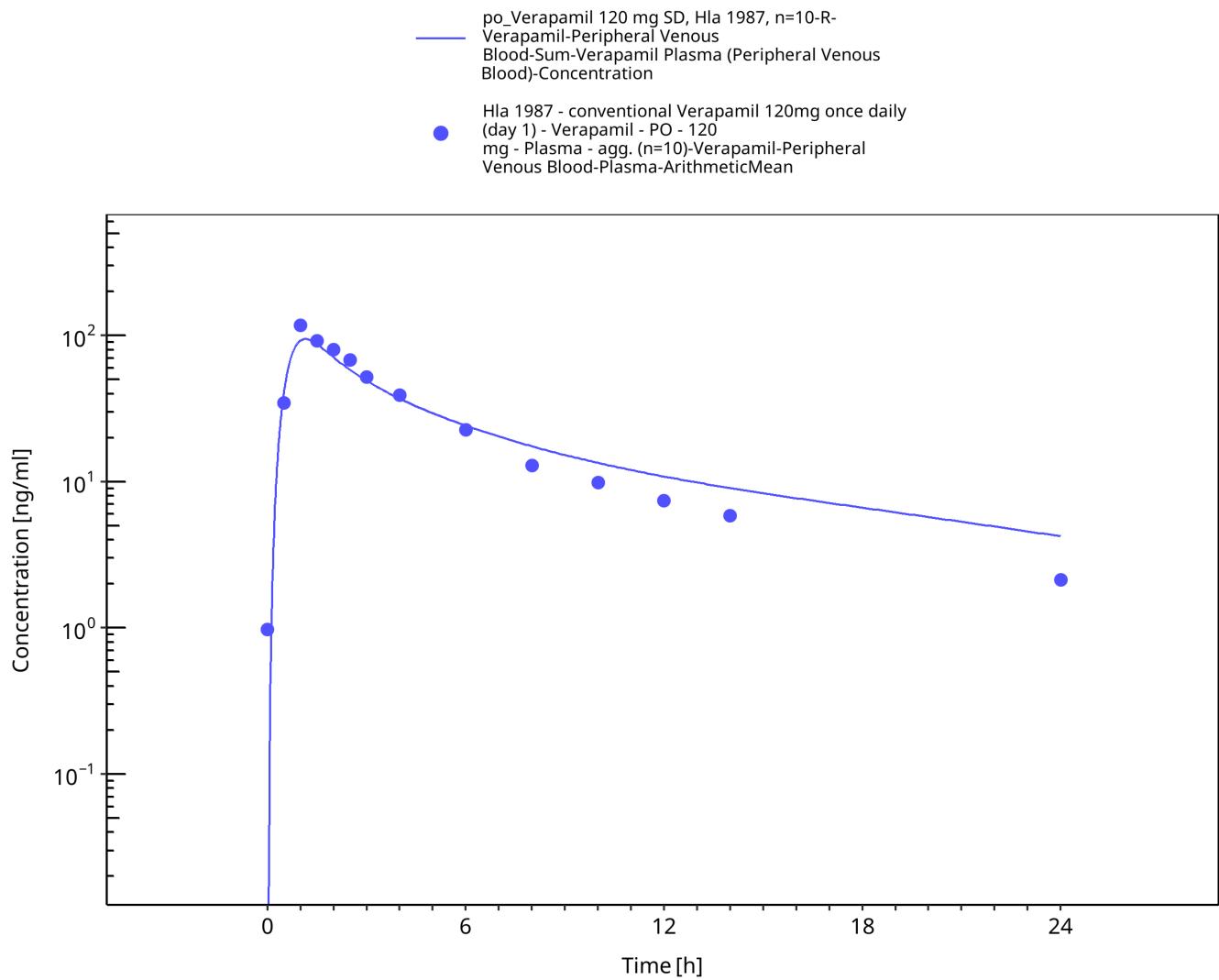


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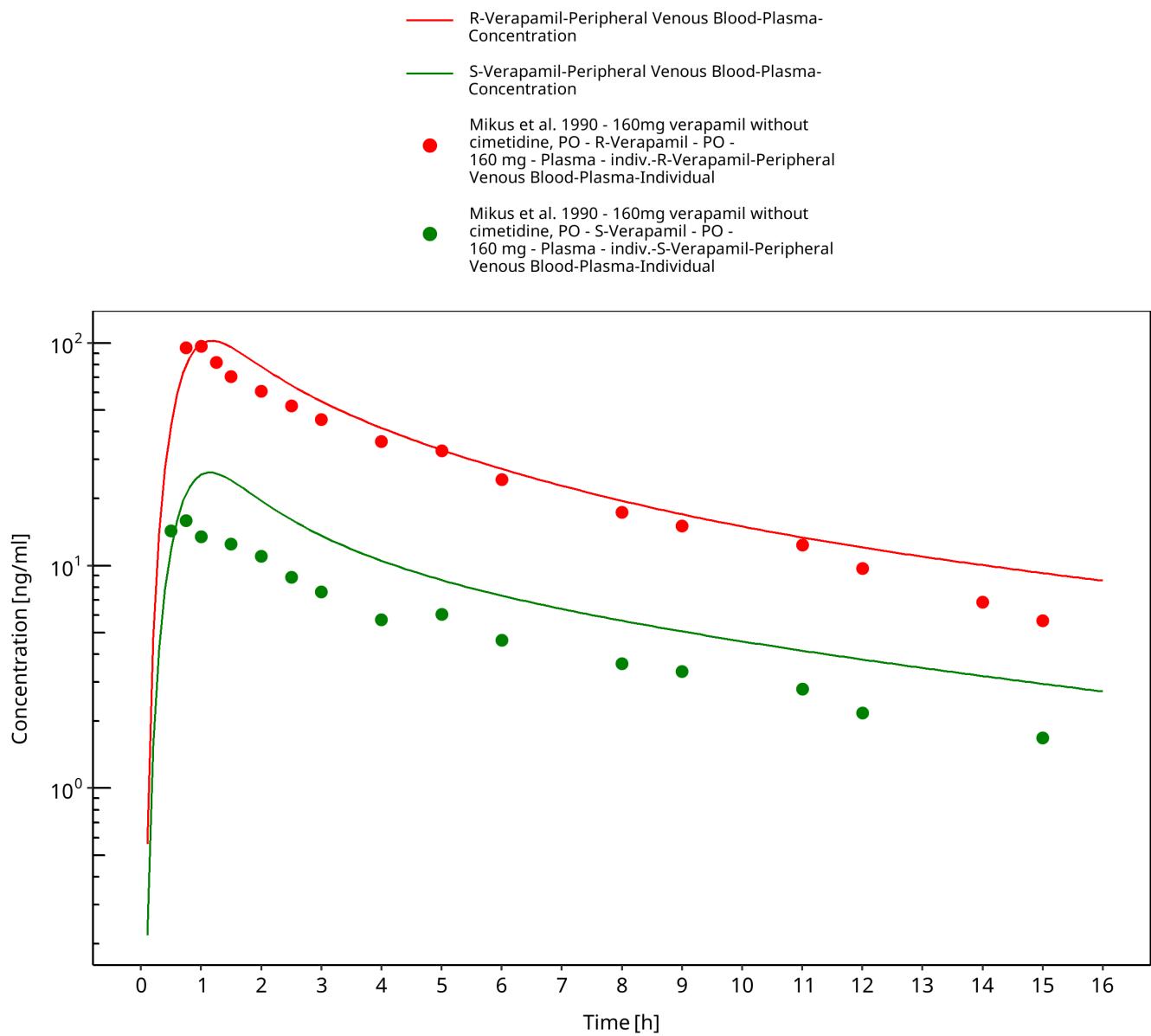


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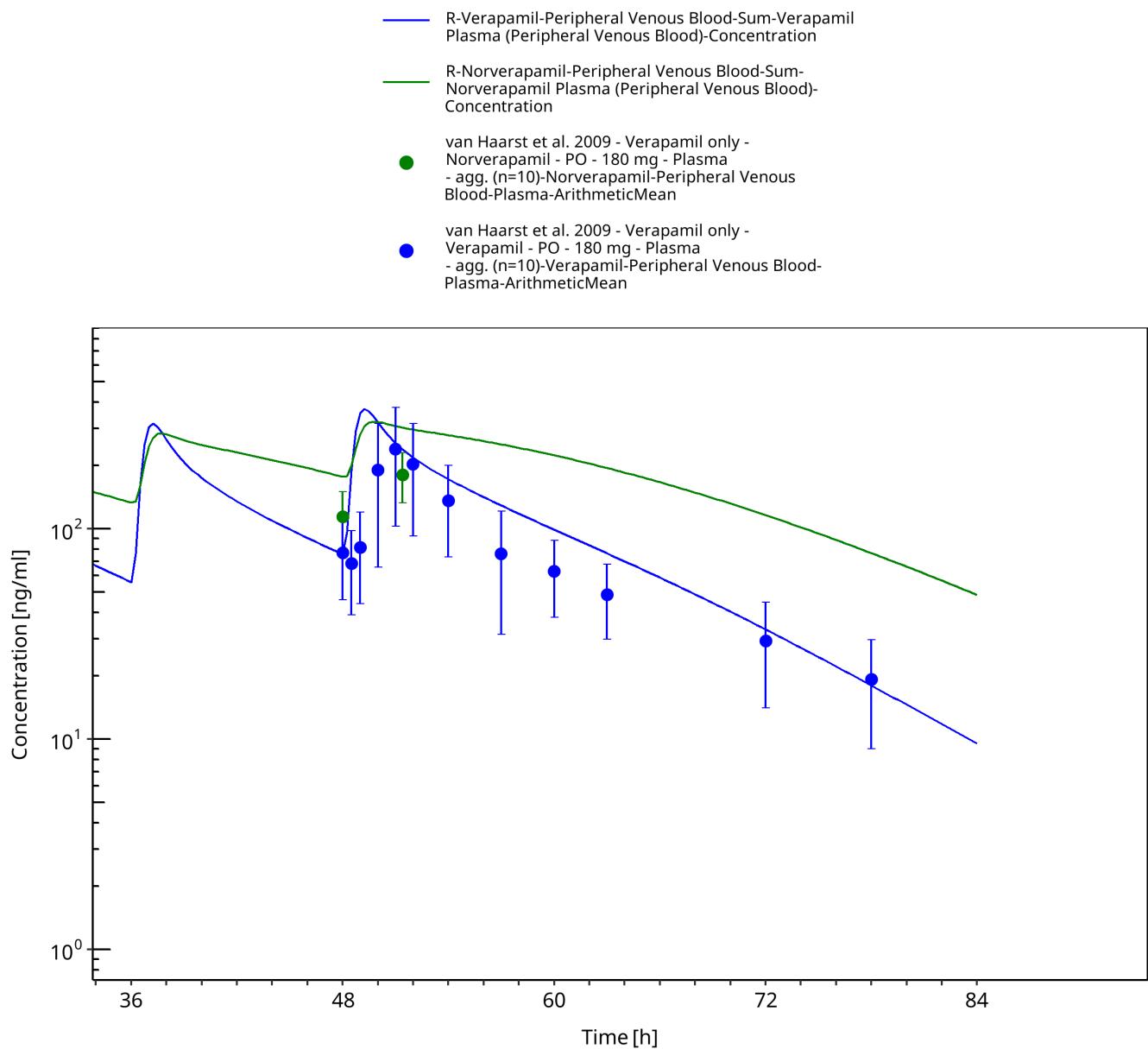


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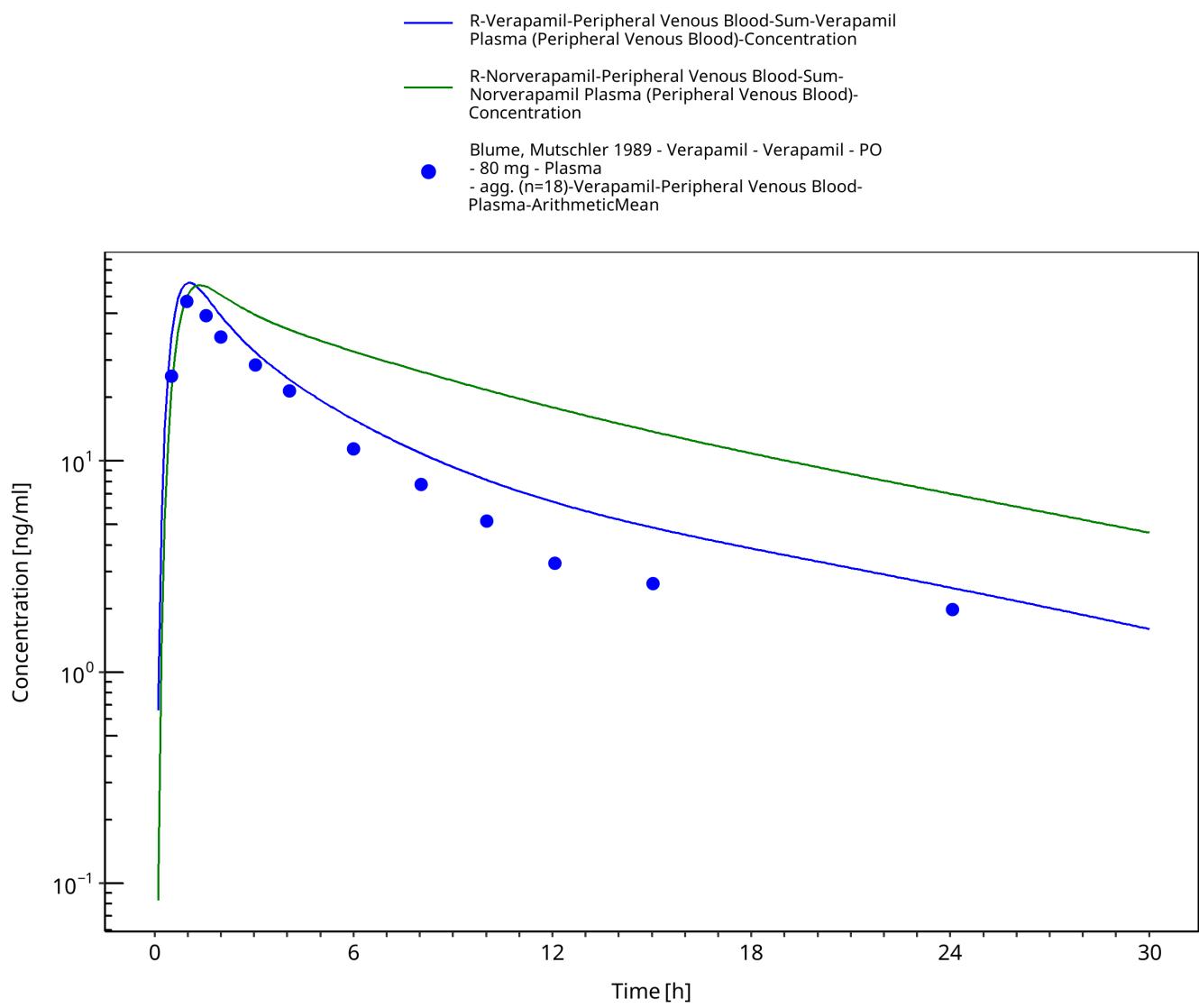


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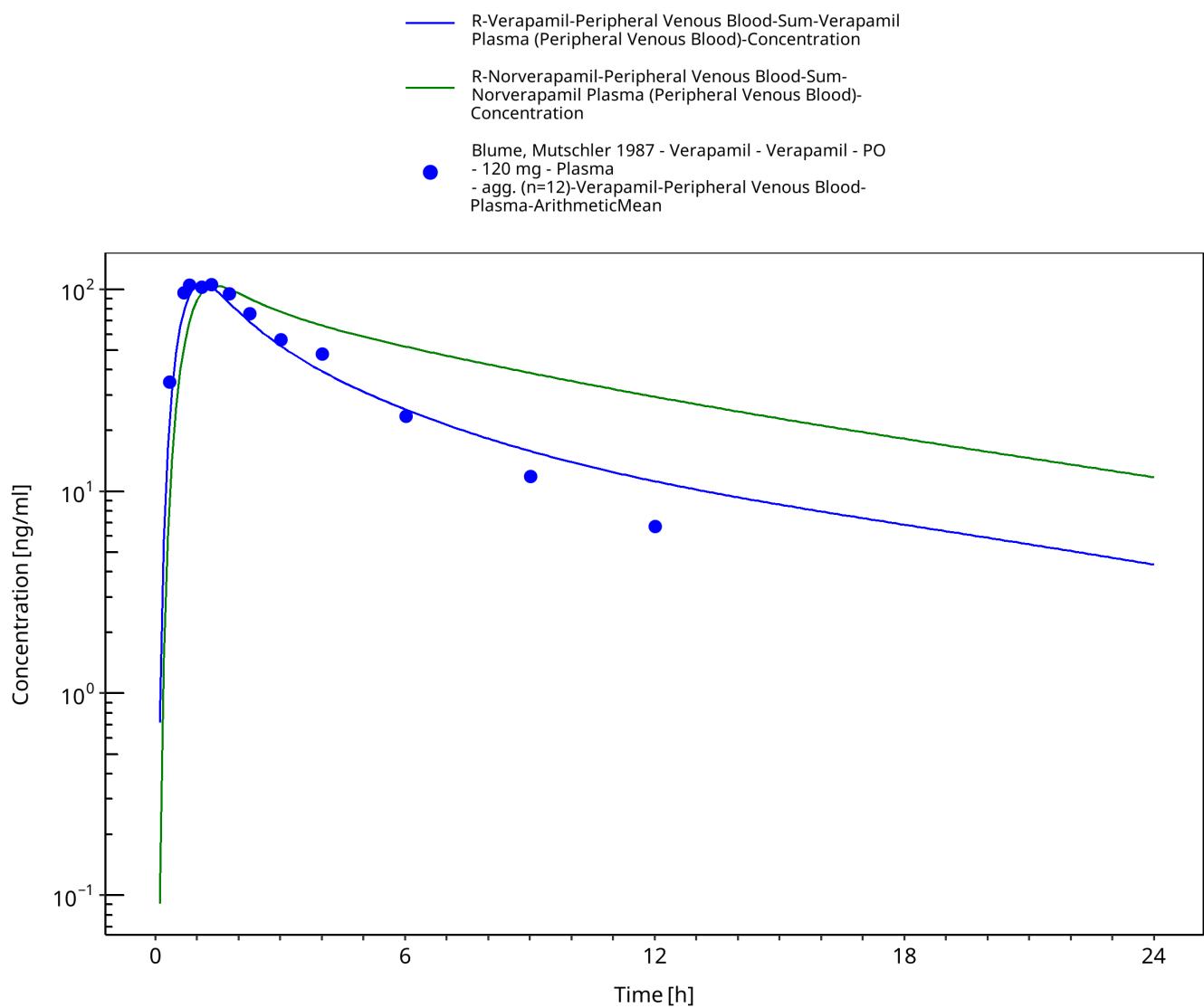


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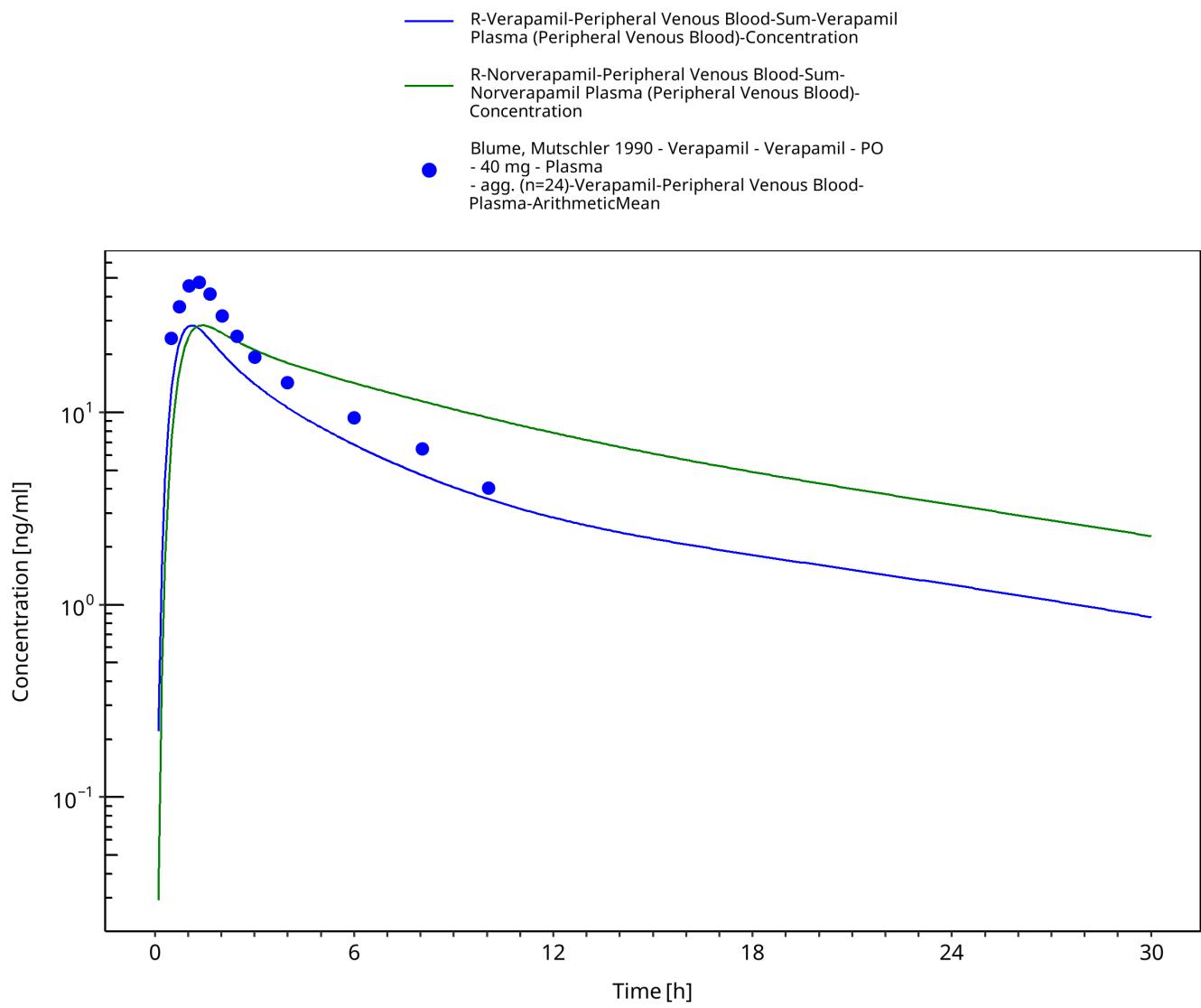


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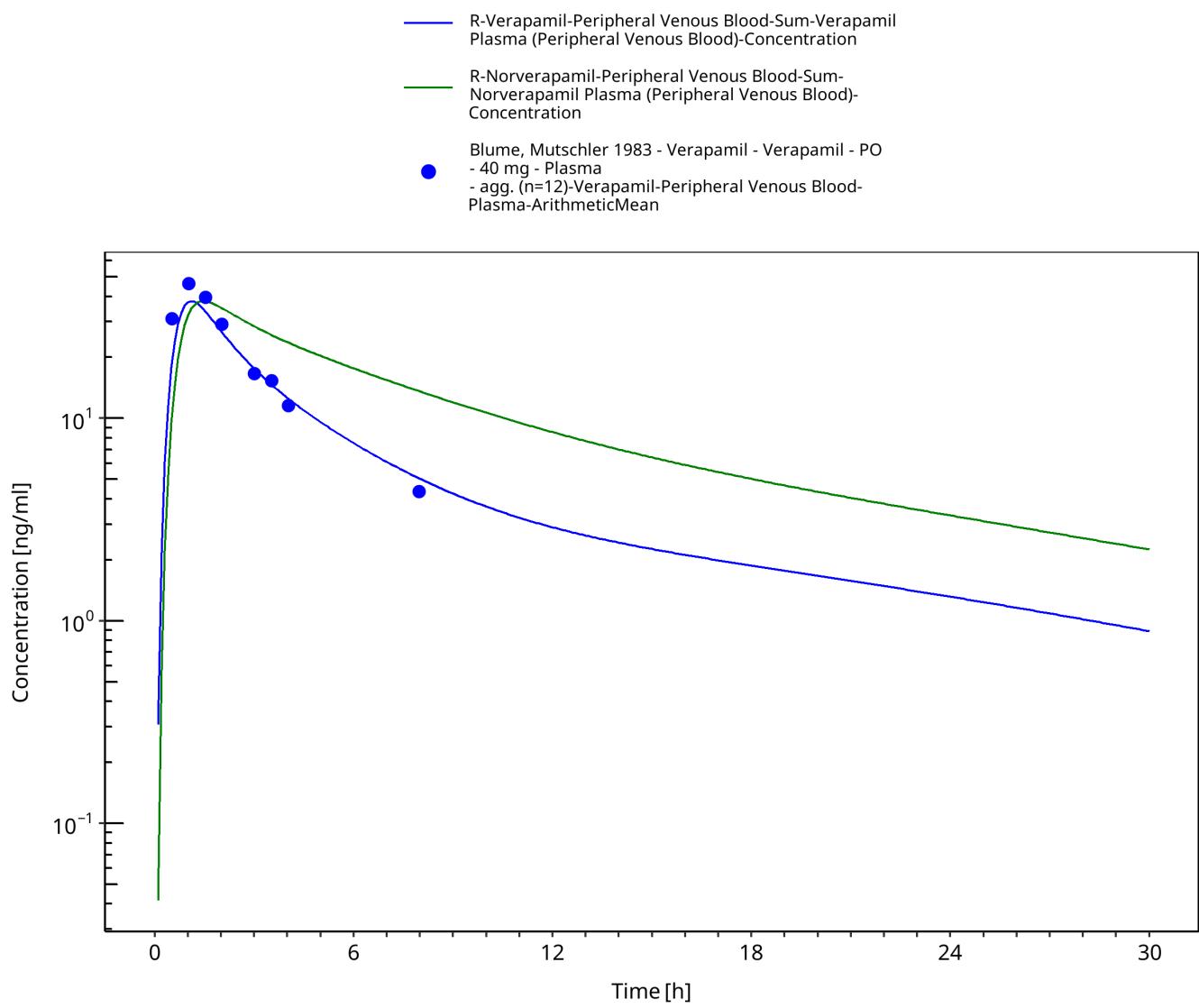


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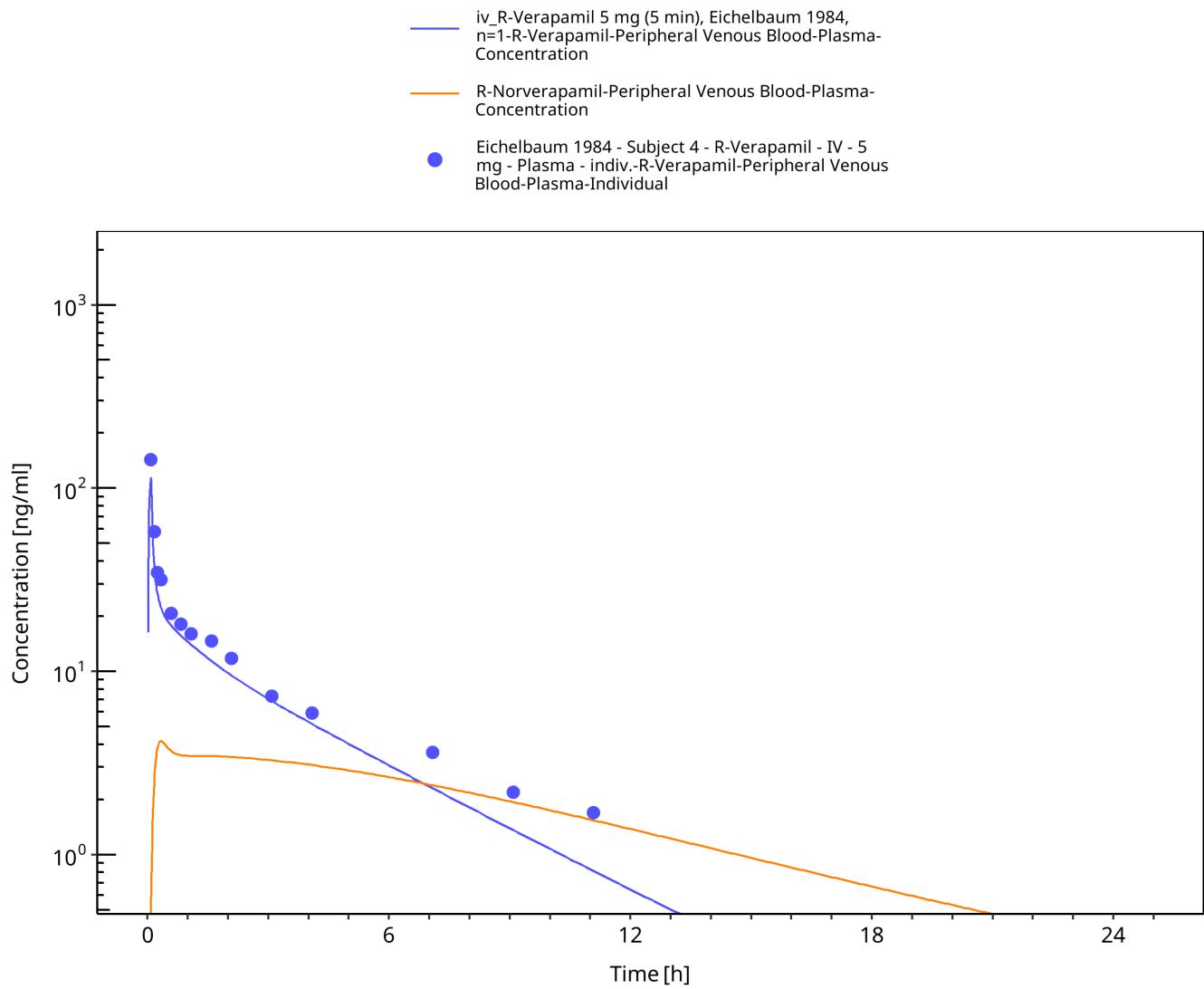


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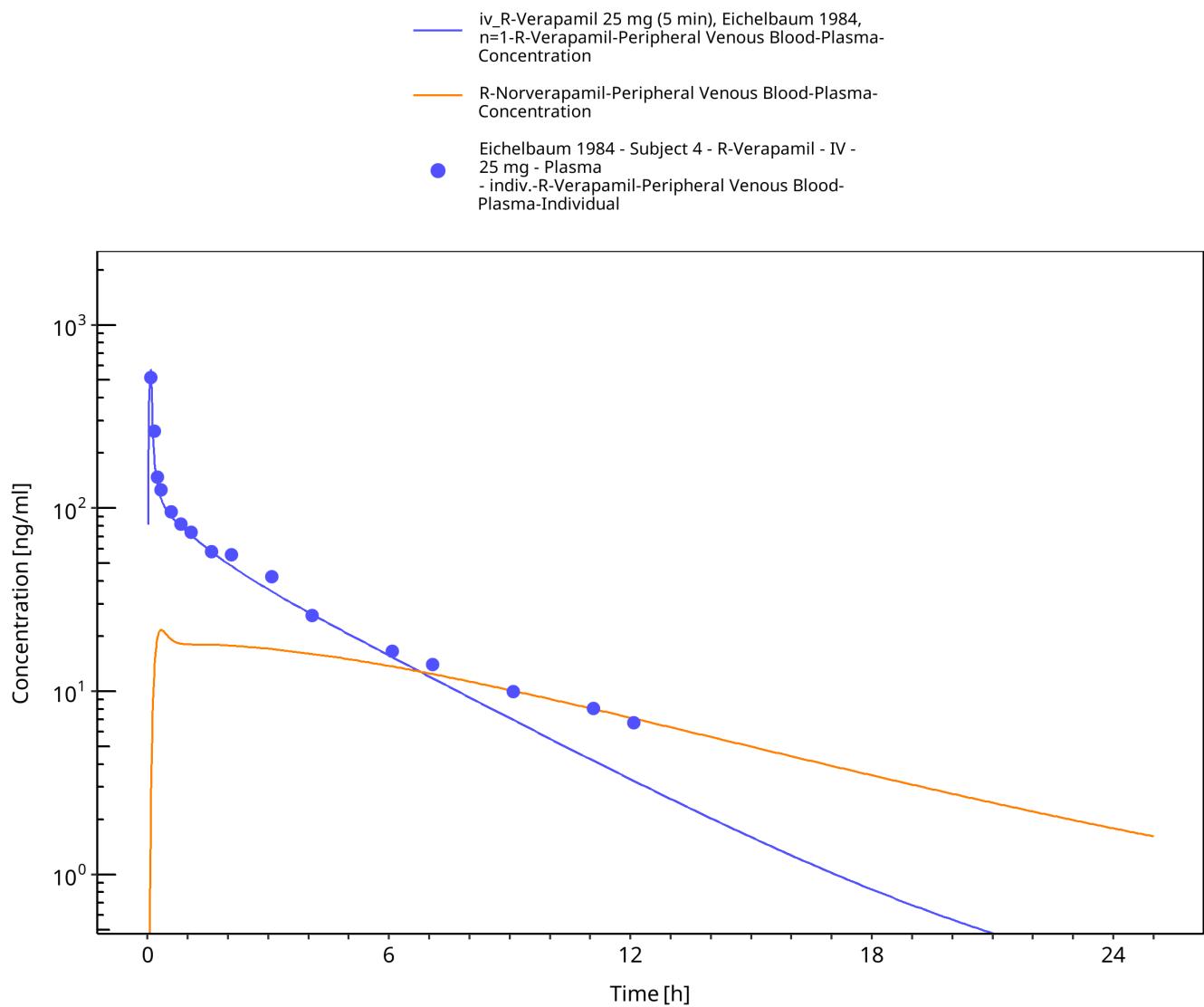


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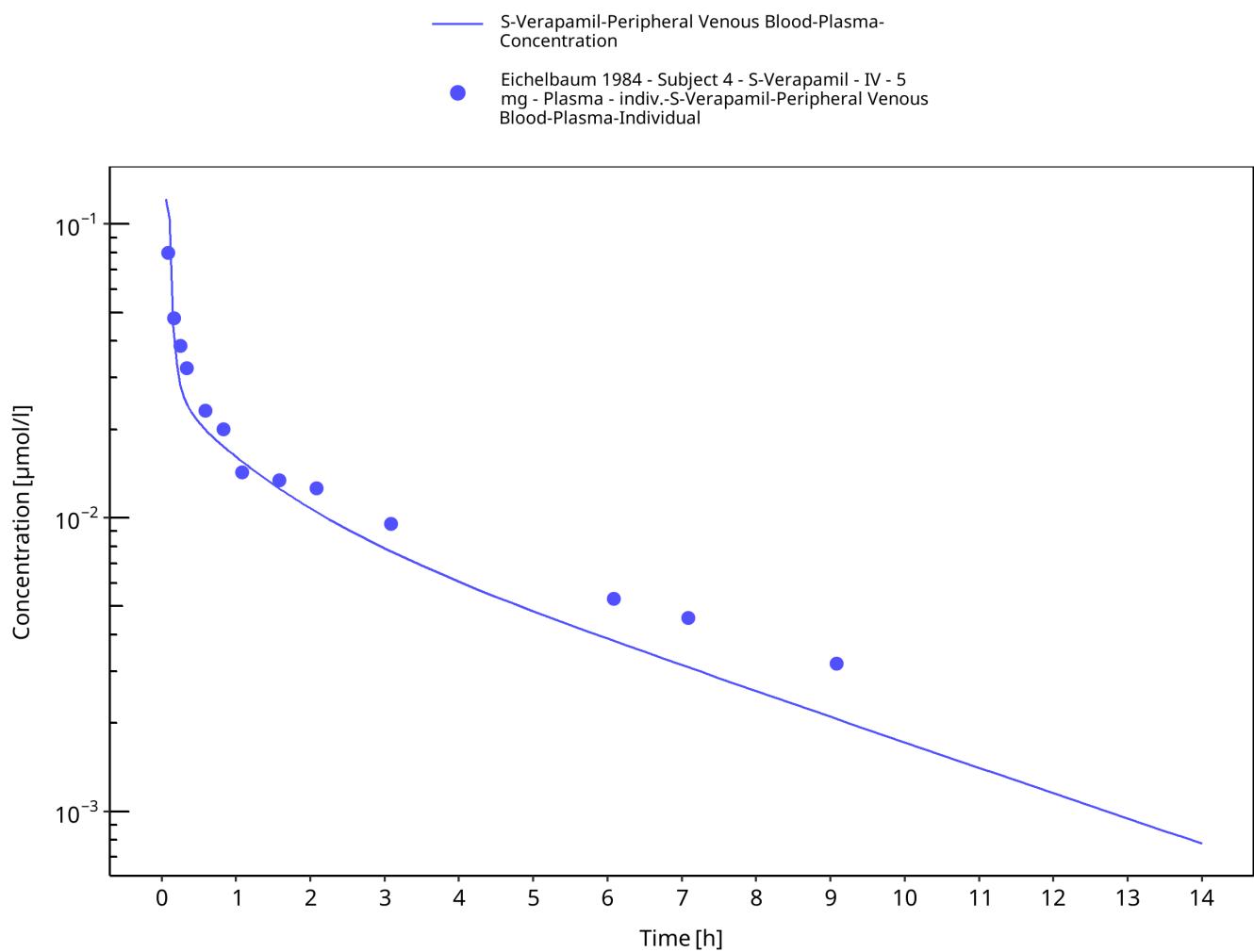


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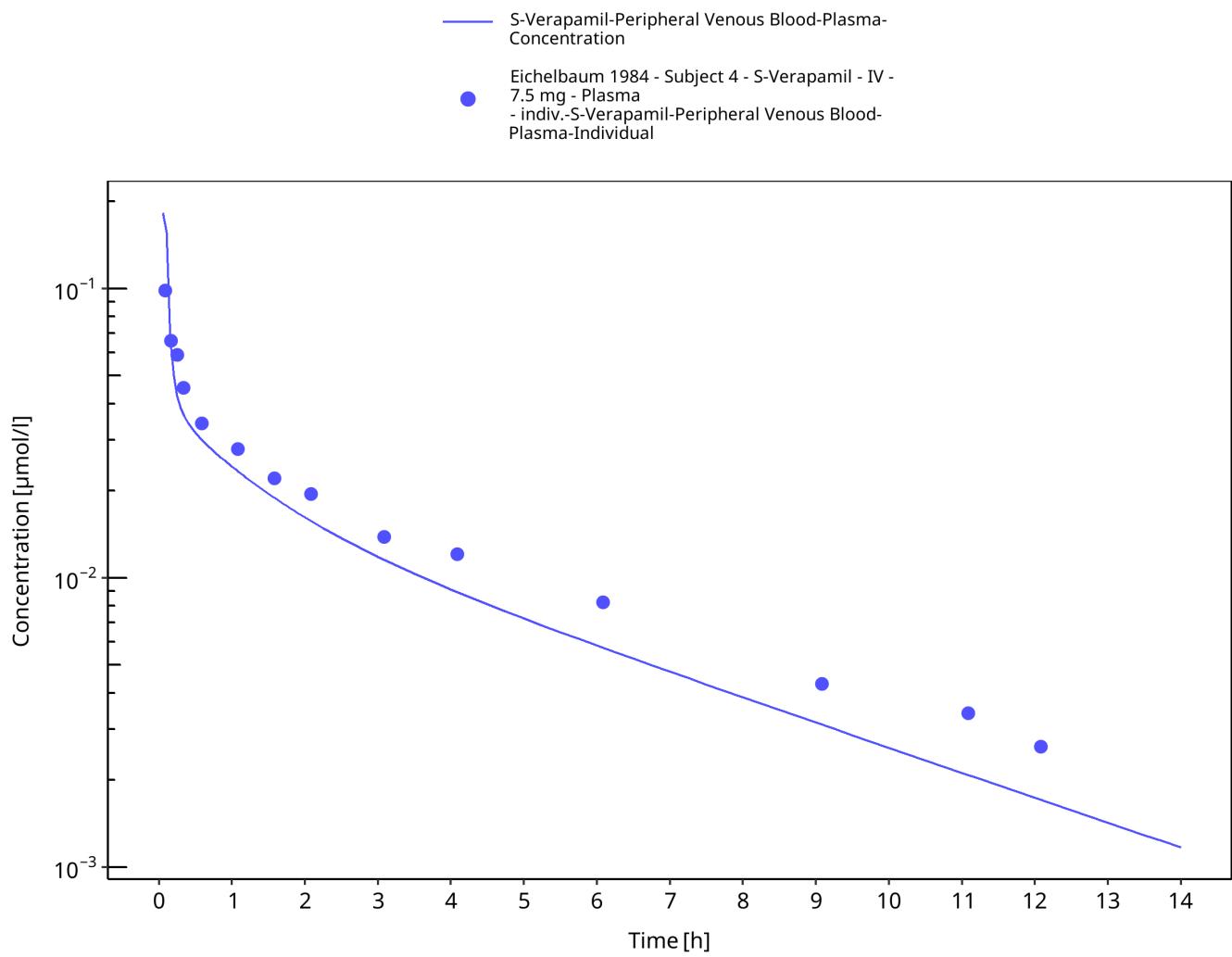


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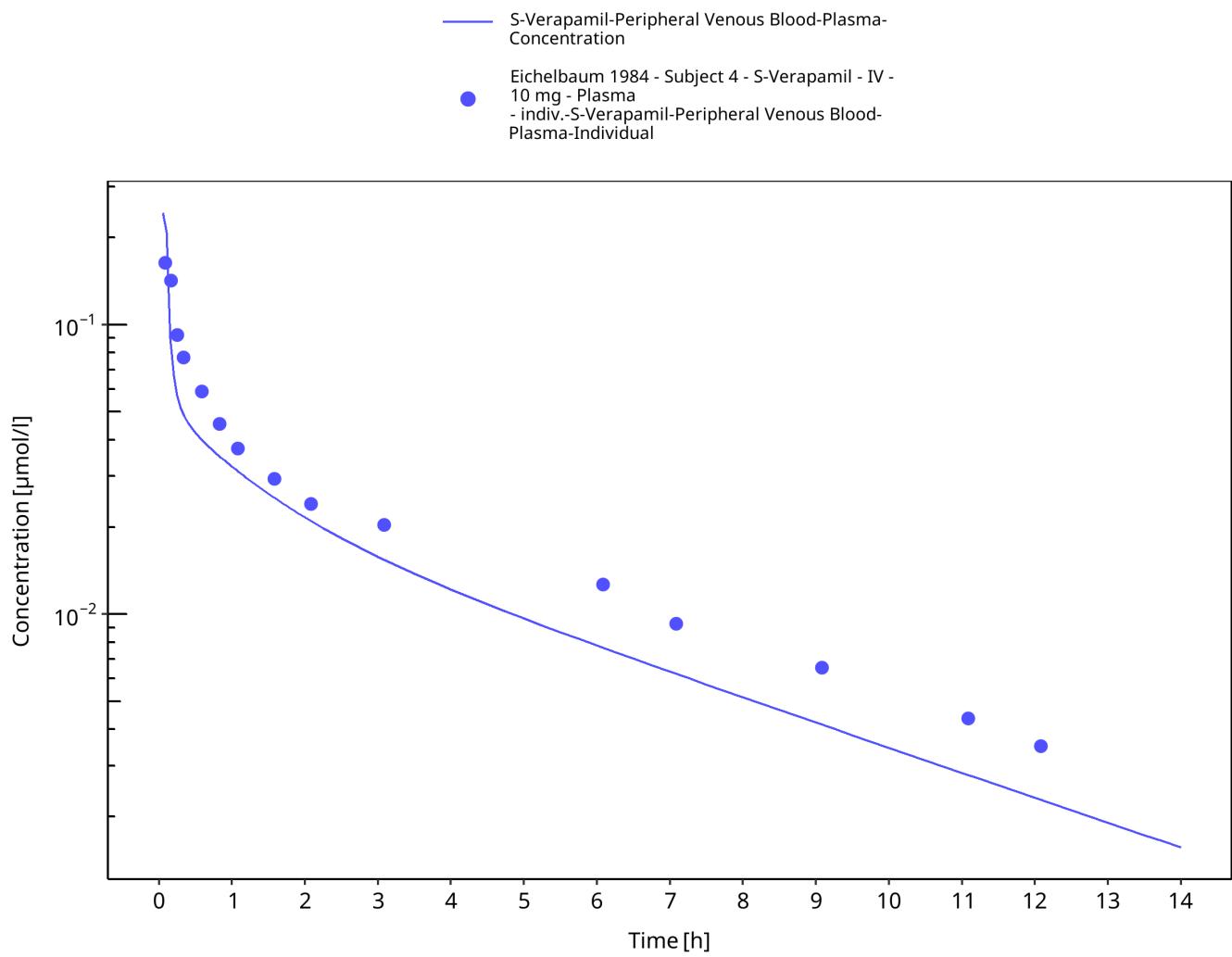


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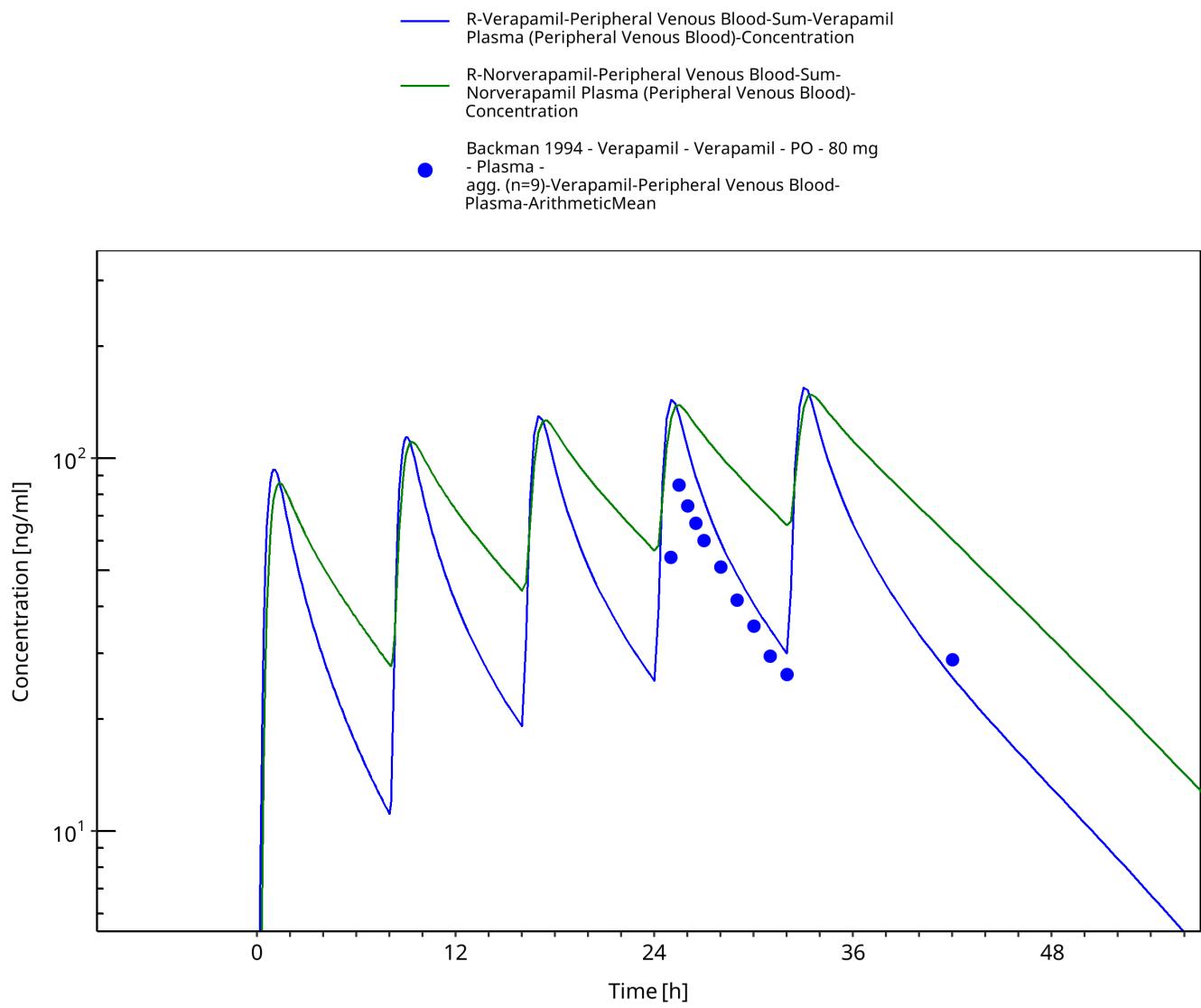


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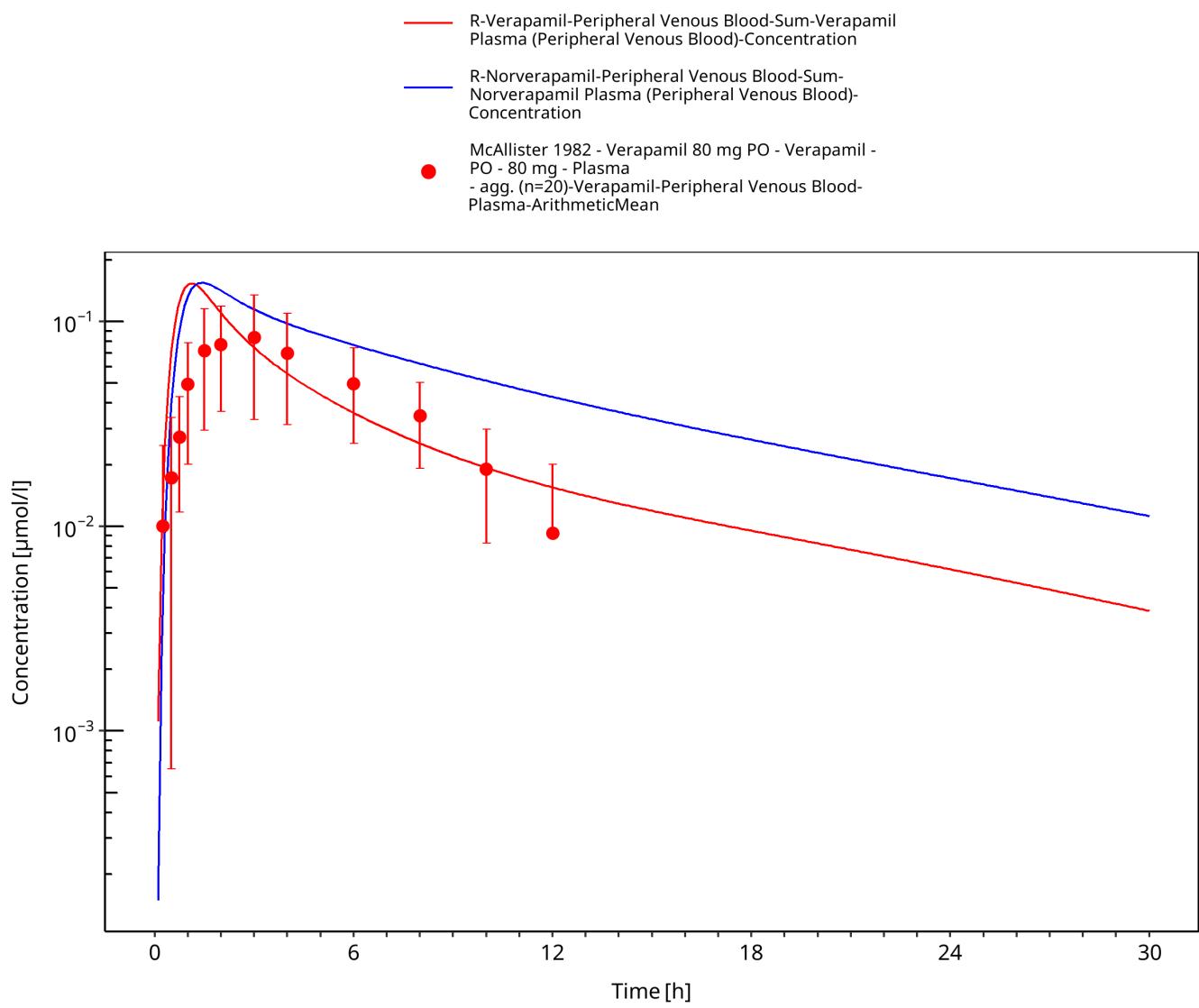


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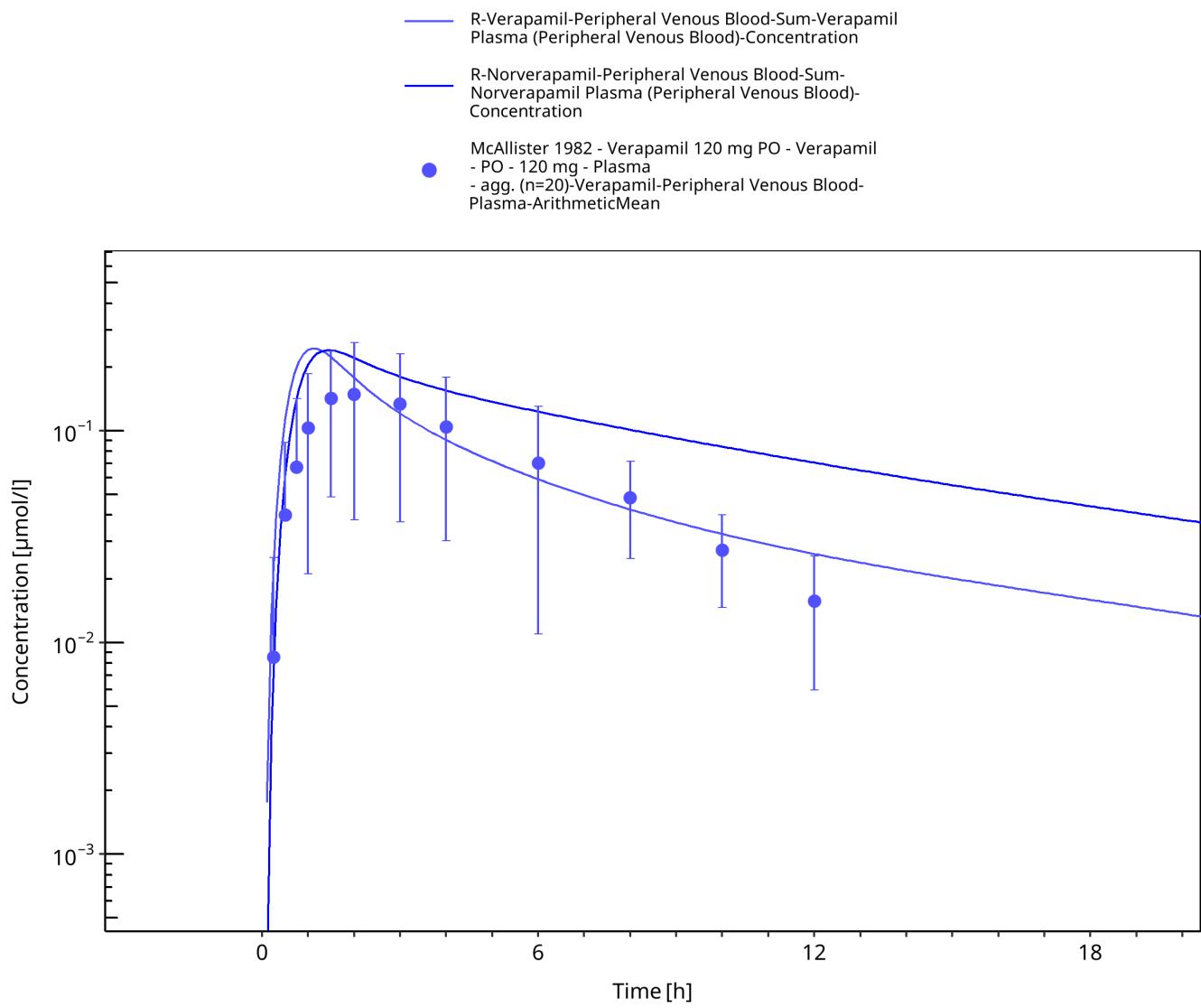


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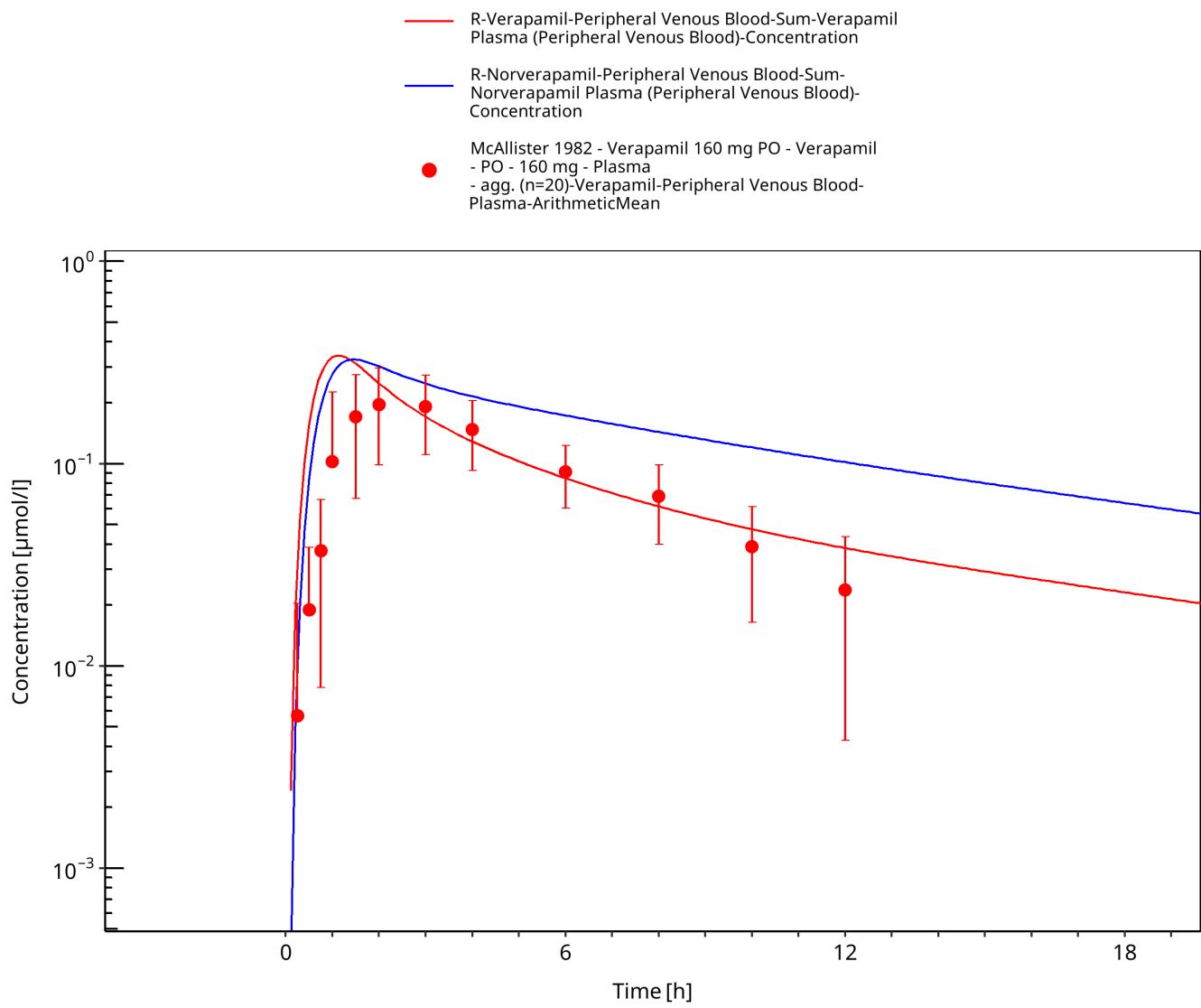


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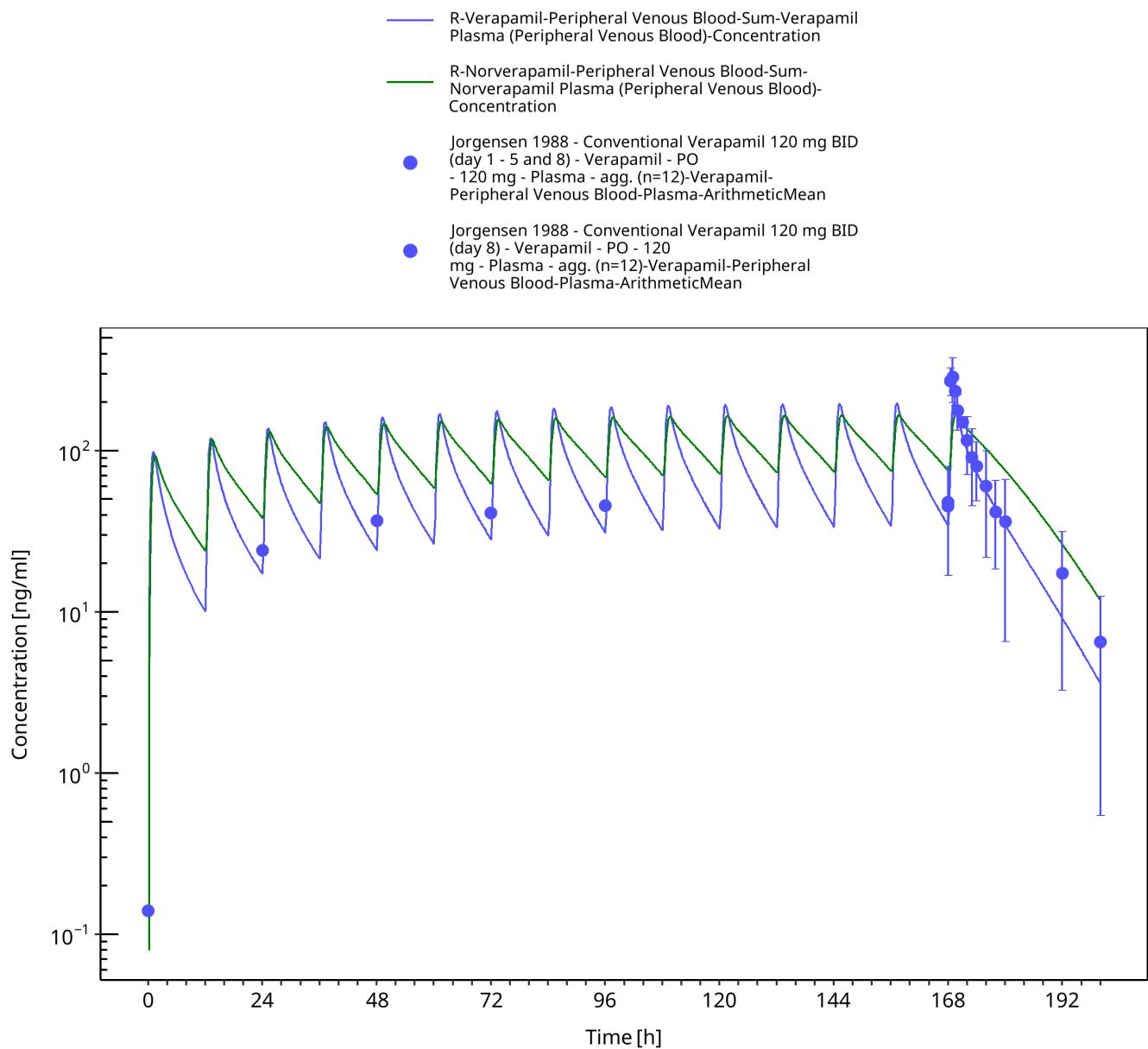


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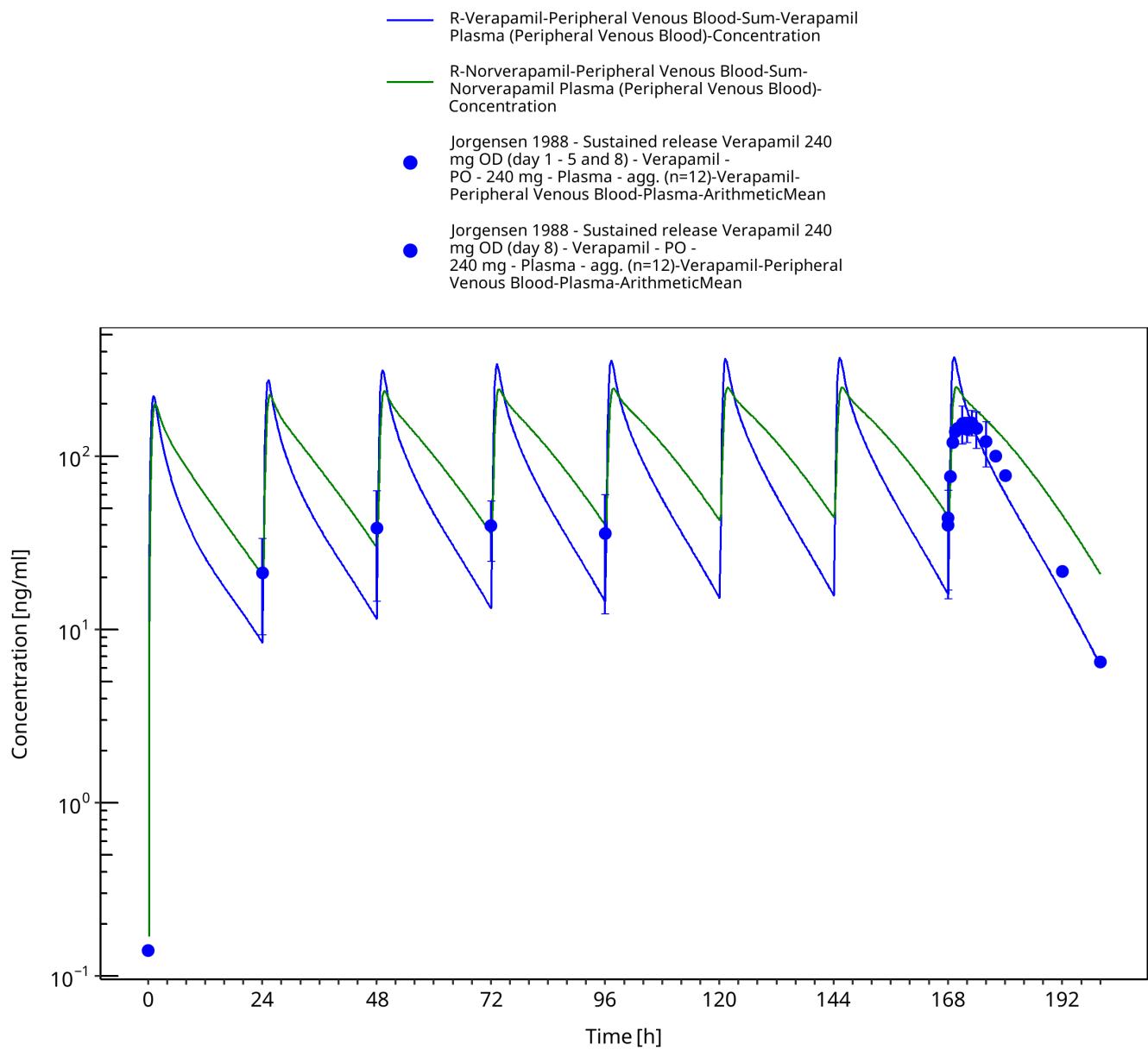


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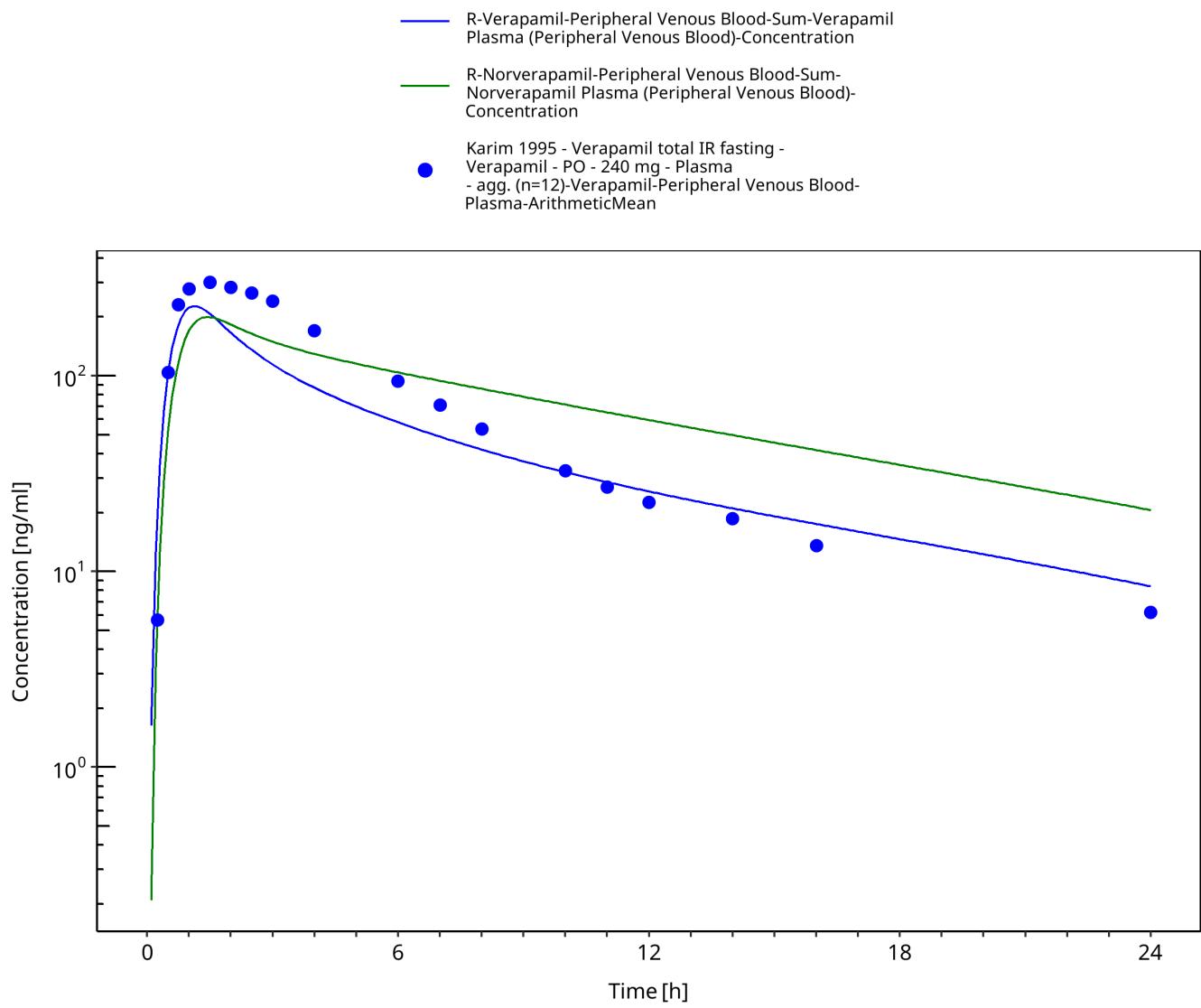


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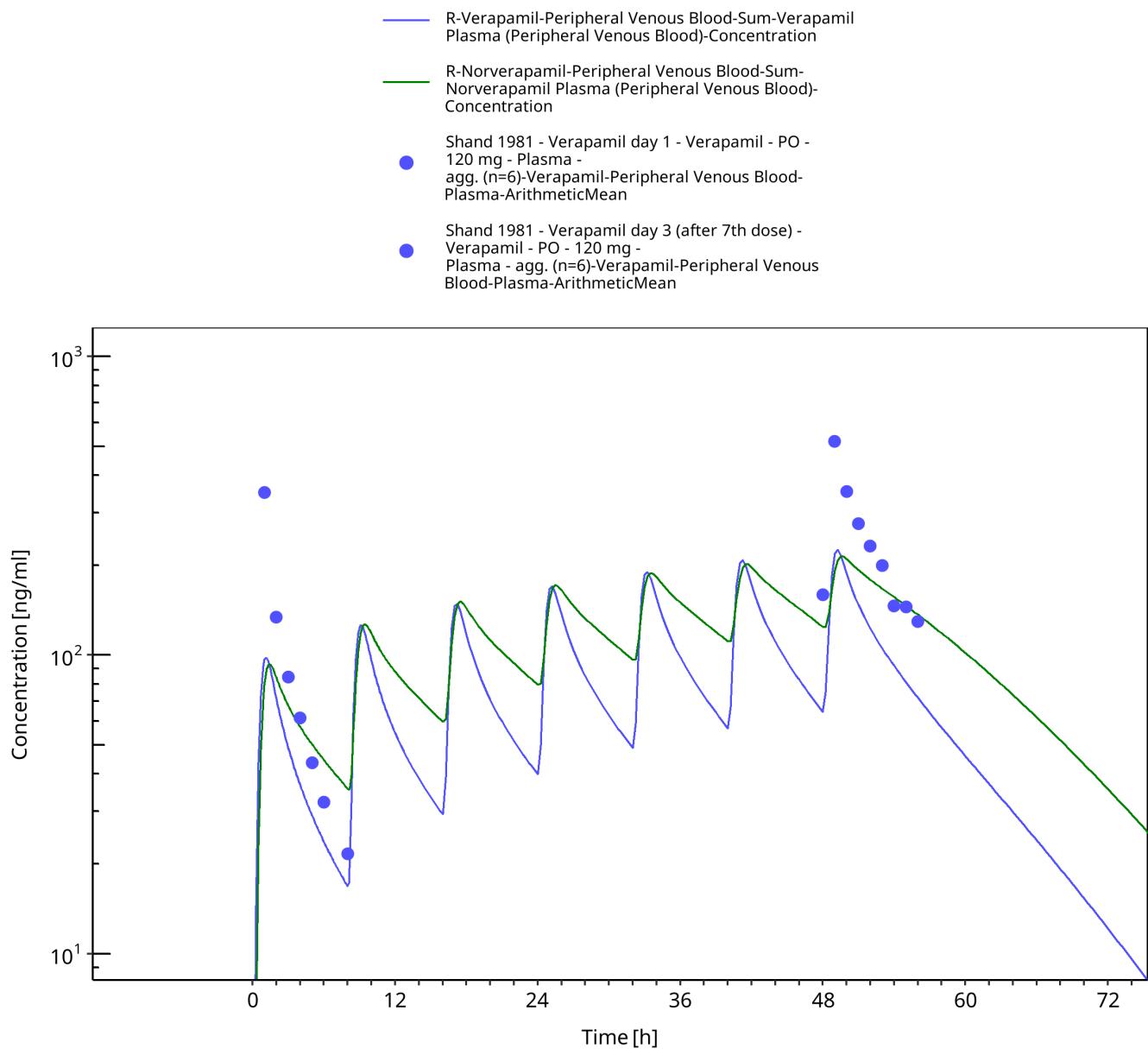


Figure 3-54: Time Profile Analysis

4 Conclusion

The herein presented PBPK model adequately describes the pharmacokinetics of R-verapamil, S-verapamil, R-norverapamil and S-norverapamil after single and multiple administration of a variety of doses to healthy adults. Furthermore, mechanism-based CYP3A4 (auto-) inactivation on verapamil itself can be described well with the optimized parameterization.

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