

# Building and evaluation of a PBPK model for cimetidine in healthy adults

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Version	1.1-OSP11.1
based on <i>Model Snapshot and Evaluation Plan</i>	<a href="https://github.com/Open-Systems-Pharmacology/Cimetidine-Model/releases/tag/v1.1">https://github.com/Open-Systems-Pharmacology/Cimetidine-Model/releases/tag/v1.1</a>
OSP Version	11.1
Qualification Framework Version	2.3

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

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

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# 1 Introduction

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Cimetidine is a histamine H<sub>2</sub> receptor antagonist that inhibits stomach acid production. It is mainly used as an antacid for the treatment of gastric and duodenal ulcers, Zollinger-Ellison syndrome and esophageal reflux.

The herein presented model was developed and published by Hanke et al. ([Hanke 2020](#)).

Cimetidine is mainly excreted unchanged via the kidneys (40–80% of the dose) with a high renal clearance of 400 ml/min. Metabolism is reported to account for 25–40% of the total elimination of cimetidine, with less than 2% of the dose excreted unchanged with the bile. Cimetidine inhibits several transporters and CYP enzymes and it is recommended by the FDA as strong inhibitor of OCT2/MATE and as weak inhibitor of CYP3A4 and CYP2D6 for the use in clinical DDI studies and drug labeling.

The cimetidine model was established using 27 clinical studies, covering a dosing range from 100 to 800 mg. The final model applies active uptake of cimetidine into the liver by OCT1, uptake into the kidney by OAT3 and secretion from the kidney into the urine by MATE1, as well as an unspecific hepatic clearance and passive renal glomerular filtration.

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

## 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 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 ([Willmann 2007](#)). 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](#)) or otherwise referenced for the specific process. The final model applies active uptake of cimetidine into the liver by OCT1, uptake into the kidney by OAT3 and secretion from the kidney into the urine by MATE1, as well as an unspecific hepatic clearance and passive renal glomerular filtration. The transporters were integrated into the PBPK model using the ([PK-Sim Ontogeny Database Version 7.3](#)) and is described in detail in [Hanke 2020](#).

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

Oral administration of cimetidine in the fasted state frequently produces two plasma concentrations peaks. These double peaks are probably caused by the phasic gastrointestinal motility that controls gastric emptying in the fasted state. To describe the very different shapes of the observed mean cimetidine plasma profiles, split dose administration protocols for all studies of cimetidine administered orally in the fasted state were optimized in a NONMEM analysis (see [Hanke 2020](#)). The resulting split dose administration protocols were then implemented and used for the PBPK modeling of the respective cimetidine studies.

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

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

A literature search was performed to collect available information on physiochemical properties of cimetidine. The obtained information from literature is summarized in the table below.

Parameter	Unit	Value	Source	Description
MW	g/mol	252.34	<a href="#">Wishart 2006</a>	Molecular weight
pK <sub>a1</sub>	6.93	(base)	<a href="#">Avdeef 2001</a>	Acid dissociation constant
pK <sub>a2</sub>	13.38	(acid)	<a href="#">Wishart 2006</a>	Acid dissociation constant
Solubility (pH)	mg/L	24.00 (6.8)	<a href="#">Avdeef 2001</a>	Water solubility
logP		0.48	<a href="#">Avdeef 2001</a>	Partition coefficient between octanol and water
f <sub>u</sub>	%	78.00	<a href="#">Taylor 1978</a>	Fraction unbound in plasma
B/P ratio		0.98	<a href="#">Somogyi 1983</a>	Blood to plasma ratio
OCT1 K <sub>m</sub>	μmol/l	2600	<a href="#">Umehara 2007</a>	Michaelis-Menten constant
OAT3 K <sub>m</sub>	μmol/l	149	<a href="#">Tahara 2005</a>	Michaelis-Menten constant
MATE1 K <sub>m</sub>	μmol/l	8.0	<a href="#">Ohta 2005</a>	Michaelis-Menten constant
OCT1 K <sub>i</sub>	μmol/l	104	<a href="#">Ito 2012</a>	Inhibition constant for competitive inhibition
OCT2 K <sub>i</sub>	μmol/l	124	<a href="#">Ito 2012</a>	Inhibition constant for competitive inhibition
MATE1 K <sub>i</sub>	μmol/l	3.8	<a href="#">Ito 2012</a>	Inhibition constant for competitive inhibition
CYP3A4 K <sub>i</sub>	μmol/l	268	<a href="#">Wrighton 1994</a>	Inhibition constant for competitive inhibition

## 2.2.2 Clinical Data

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

### 2.2.2.1 Model Building

The following studies were used for model building:

Publication	Arm / Treatment / Information used for model building
<a href="#">Bodemar 1981</a>	Peptic ulcer patients receiving a single intravenous dose of 200 mg and oral doses of 200, 400 and 800 mg
<a href="#">Morgan 1983</a>	Peptic ulcer patients receiving a single intravenous dose of 200 mg (5 min infusion)
<a href="#">Bodemar 1979</a>	Healthy subjects receiving single oral doses of 200 and 400mg (tablet)
<a href="#">Walkenstein 1978</a>	Healthy subjects receiving a single oral dose of 300mg (solution)
<a href="#">D'Angio 1986</a>	Healthy subjects receiving a single oral dose of 300mg (tablet)

### 2.2.2.2 Model verification

The following studies were used for model verification:

Publication	Arm / Treatment / Information used for model verification
<a href="#">Grahnen 1979</a>	Healthy subjects receiving a single intravenous dose of 100 mg and a single oral dose of 400 mg (tablet)
<a href="#">Larsson 1982</a>	Peptic ulcer patients receiving a single intravenous dose of 200 mg
<a href="#">Mihaly 1984</a>	Peptic ulcer patients receiving a single intravenous and a single oral dose of 200 mg
<a href="#">Morgan 1983</a>	Peptic ulcer patients receiving a single intravenous dose of 200 mg (30 min infusion)
<a href="#">Lebert 1981</a>	Healthy subjects receiving a single intravenous dose of 300 mg (2 min infusion)
<a href="#">Walkenstein 1978</a>	Healthy subjects receiving a single intravenous dose of 300 mg (2 min infusion) and a single oral dose of 300 mg (tablet)
<a href="#">Kanto 1981</a>	Healthy subjects receiving a single oral dose of 200 mg
<a href="#">Burland 1975</a>	Healthy subjects receiving single oral doses of 200 mg solution and capsule
<a href="#">Bodemar 1979</a>	Peptic ulcer patients receiving a single oral dose of 200 mg (tablet)
<a href="#">Bodemar 1981</a>	Peptic ulcer patients receiving single oral doses of 800 mg and multiple oral doses of 200 and 400 mg
<a href="#">Barbhaiya 1995</a>	Healthy subjects receiving multiple oral doses of 300 mg (tablet)
<a href="#">Somogyi 1981</a>	Healthy subjects receiving a single oral dose of 400 mg (tablet)
<a href="#">Tiseo 1998</a>	Healthy subjects receiving multiple oral doses of 800 mg (tablet)

## 2.3 Model Parameters and Assumptions

### 2.3.1 Absorption

Absorption observed in clinical studies can be fully explained by passive absorption.

### 2.3.2 Distribution

Cimetidine is reported to be actively taken up into the liver by OCT1 ([Umehara 2007](#)), into the kidney by OAT3 ([Tahara 2005](#)) and secreted from the kidney into the urine by MATE1 ([Ohta 2010](#)).

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 method by `Rodgers` and `Rowland` and cellular permeability calculation by `PK-Sim Standard`.

A `Lipophilicity` of 1.66 was back-calculated from the blood-to-plasma ratio of 0.98 ([Somogyi 1983](#), [Hanke 2020](#)).

### 2.3.3 Metabolism, Elimination and Inhibition

Cimetidine is mainly excreted unchanged via the kidneys. Additionally, 25 to 40 % is hepatically metabolized via an unknown pathway.

Cimetidine inhibits several enzymes such as CYP3A4 and CYP2D6 as well as transporters such as OCT2, OCT2 and MATE.

## 2.3.4 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 (see [Section 2.2](#)).

All values were reestimated in PK-Sim Version 10, and, therefore, do not correspond to the original values published by [Hanke 2020](#). The result of the final parameter identification is shown in the table below:

Model Parameter	Optimized Value	Unit
Specific intestinal permeability	5.26E-06	cm/min
CL <sub>hep</sub>	0.12	1/min
k <sub>cat</sub> OCT1	14098.32	1/min
k <sub>cat</sub> OAT3	2522831.10	1/min
k <sub>cat</sub> MATE1	159.47	1/min

## 3 Results and Discussion

The PBPK model for efavirenz was developed and evaluated using publicly available clinical pharmacokinetic data from studies listed in [Section 2.2.2](#).

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: Cimetidine

##### Parameters

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	24 mg/ml	Publication-Avdeef 2001	Measurement	True
Reference pH	6.8	Publication-Avdeef 2001	Measurement	True
Lipophilicity	1.655 Log Units	Parameter Identification	Measurement	True
Fraction unbound (plasma, reference value)	0.78	Publication-Taylor 1978	Measurement	True
Specific intestinal permeability (transcellular)	5.2554004942E-06 cm/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification' on 2021-09-13 17:00	Fit	True
Is small molecule	Yes			
Molecular weight	252.34 g/mol	Database-Drugbank		
Plasma protein binding partner	Unknown			

##### Calculation methods

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



## Processes

### Systemic Process: Total Hepatic Clearance-Somogyi 1983

Species: Human

#### Parameters

Name	Value	Value Origin
Fraction unbound (experiment)	0.78	
Lipophilicity (experiment)	1.655 Log Units	
Plasma clearance	0 ml/min/kg	
Specific clearance	0.1209722937 1/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification' on 2021-09-13 17:00

### Transport Protein: MATE1-Paper

Molecule: MATE1

#### Parameters

Name	Value	Value Origin
Transporter concentration	1 $\mu\text{mol/l}$	
Vmax	0 $\mu\text{mol/l/min}$	
Km	8 $\mu\text{mol/l}$	Parameter Identification
kcat	159.4749627996 1/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification' on 2021-09-13 17:00

### Transport Protein: OAT3-Paper

Molecule: OAT3

#### Parameters

Name	Value	Value Origin
Transporter concentration	1 $\mu\text{mol/l}$	
Vmax	0 $\mu\text{mol/l/min}$	
Km	149 $\mu\text{mol/l}$	Publication-Tahara 2005
kcat	2522831.1016 1/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification' on 2021-09-13 17:00

## Transport Protein: OCT1-Paper

Molecule: OCT1

### Parameters

Name	Value	Value Origin
Transporter concentration	1 $\mu\text{mol/l}$	
Vmax	0 $\mu\text{mol/l/min}$	
Km	2600 $\mu\text{mol/l}$	Publication-Umehara 2007
kcat	14098.3224931732 1/min	Parameter Identification-Parameter Identification-Value updated from 'Parameter Identification' on 2021-09-13 17:00

## Systemic Process: Glomerular Filtration-GFR

Species: Human

### Parameters

Name	Value	Value Origin
GFR fraction	1	

## Inhibition: OCT1-Ito 2012

Molecule: OCT1

### Parameters

Name	Value	Value Origin
Ki	104 $\mu\text{mol/l}$	Publication-Ito 2012

## Inhibition: OCT2-Ito 2012

Molecule: OCT2

### Parameters

Name	Value	Value Origin
Ki	124 $\mu\text{mol/l}$	Publication-Ito 2012

## Inhibition: MATE1-Ito 2012

Molecule: MATE1

### Parameters

Name	Value	Value Origin
Ki	3.8 $\mu\text{mol/l}$	Other-NBI measurement

## Inhibition: CYP3A4-Wrighton 1994

Molecule: CYP3A4

### Parameters

Name	Value	Value Origin
Ki	268 µmol/l	

## Formulation: Tablet

Type: Weibull

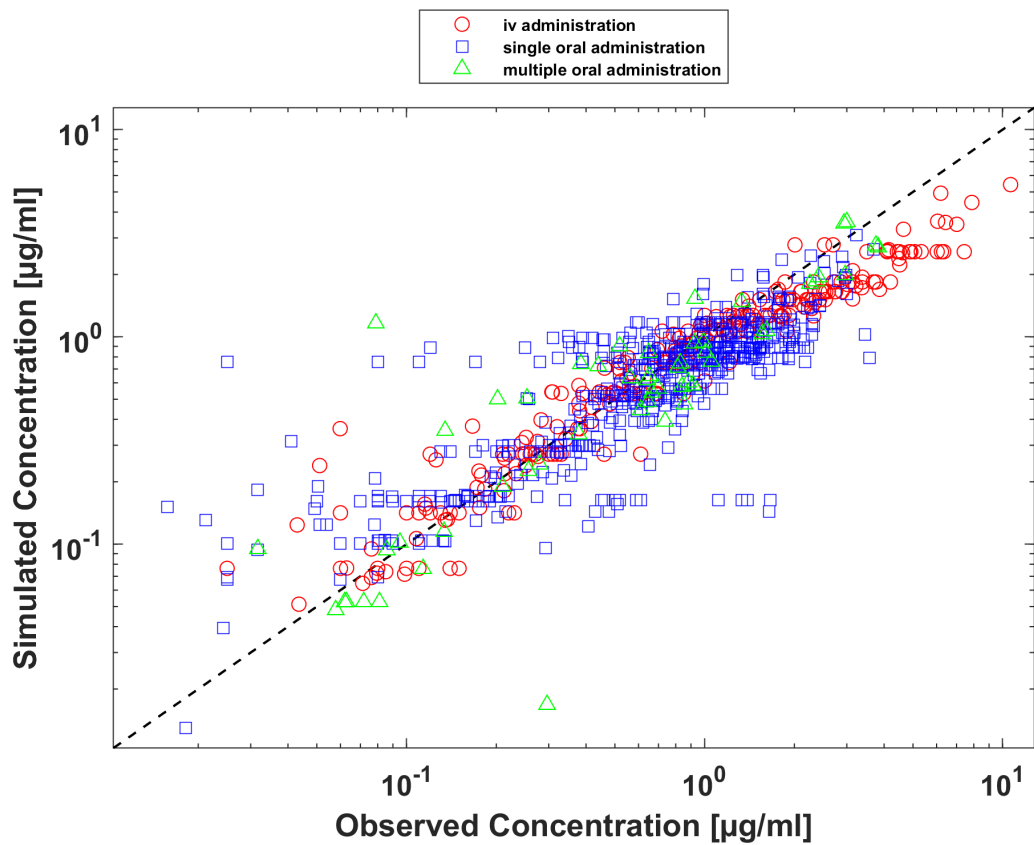
### Parameters

Name	Value	Value Origin
Dissolution time (50% dissolved)	1 min	
Lag time	0 h	
Dissolution shape	10	
Use as suspension	Yes	

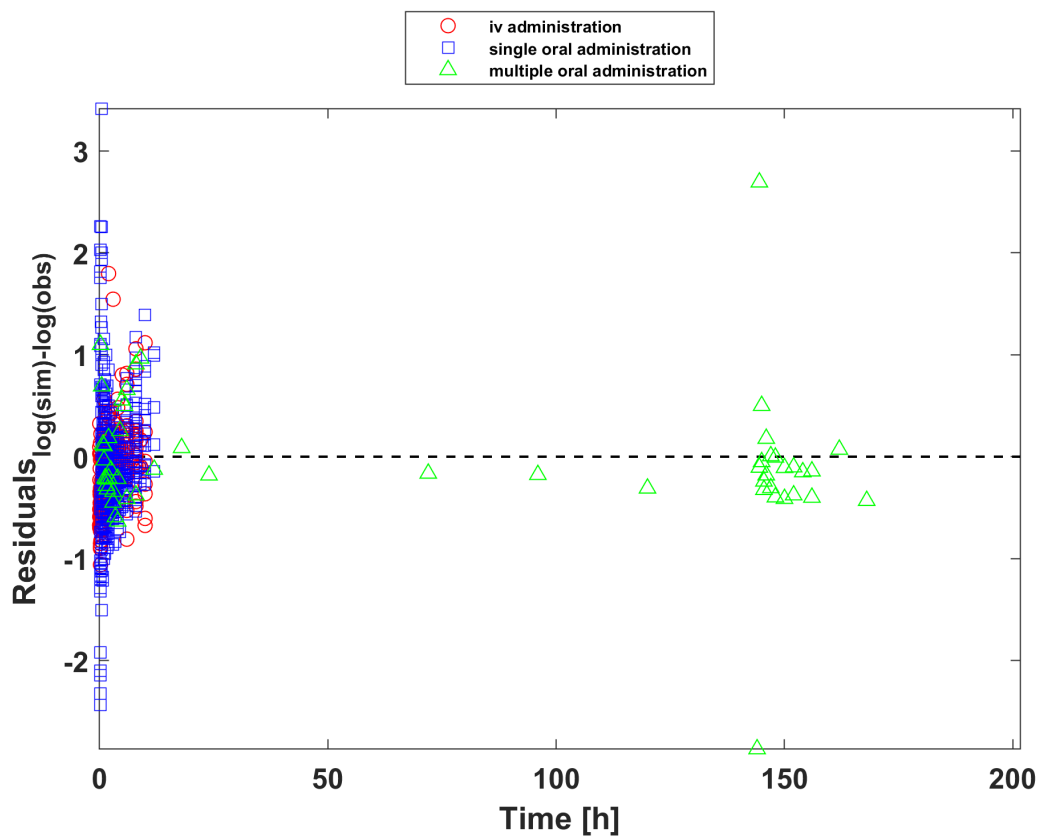
## 3.2 Diagnostics Plots

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

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



Goodness of fit plot for concentration in plasma



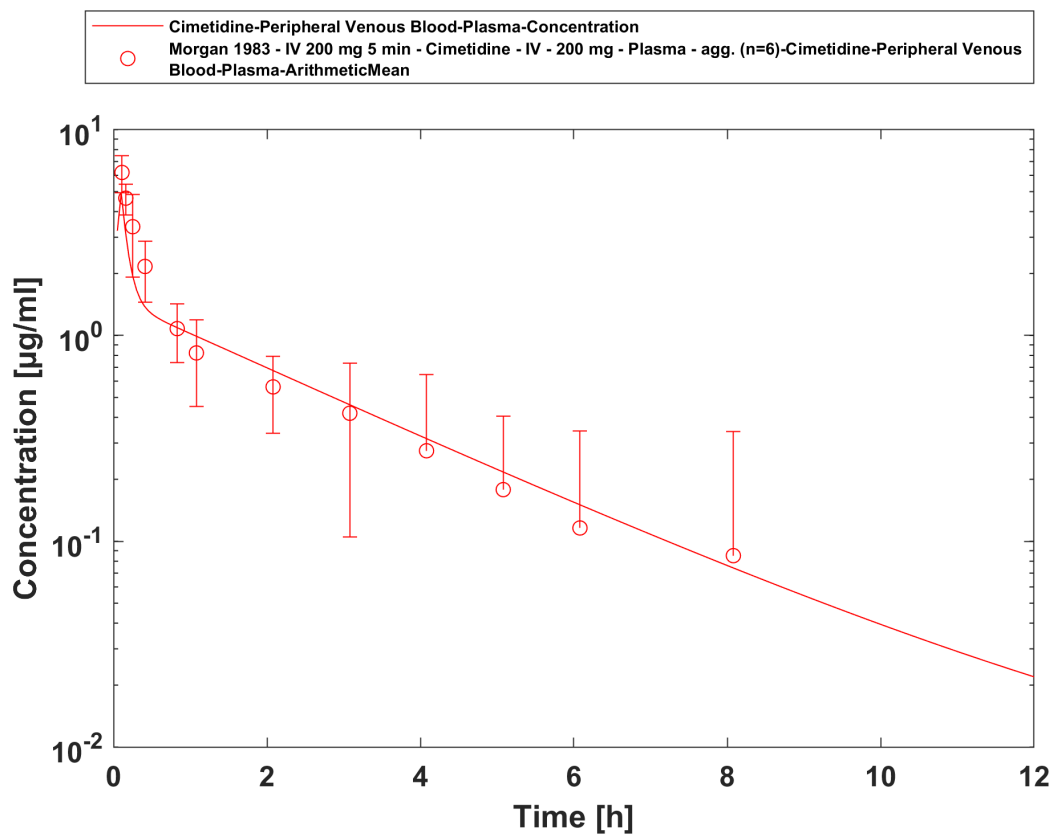
Goodness of fit plot for concentration in plasma

GMFE = 1.472437

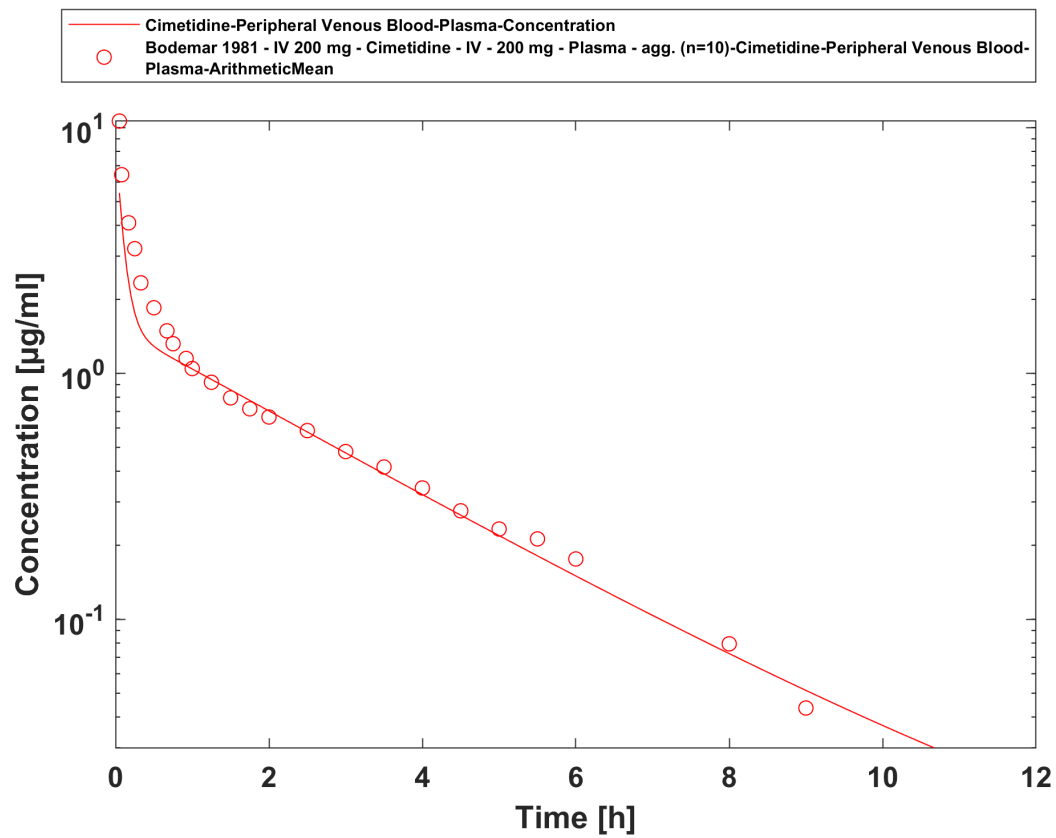
### 3.3: Concentration-Time Profiles

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

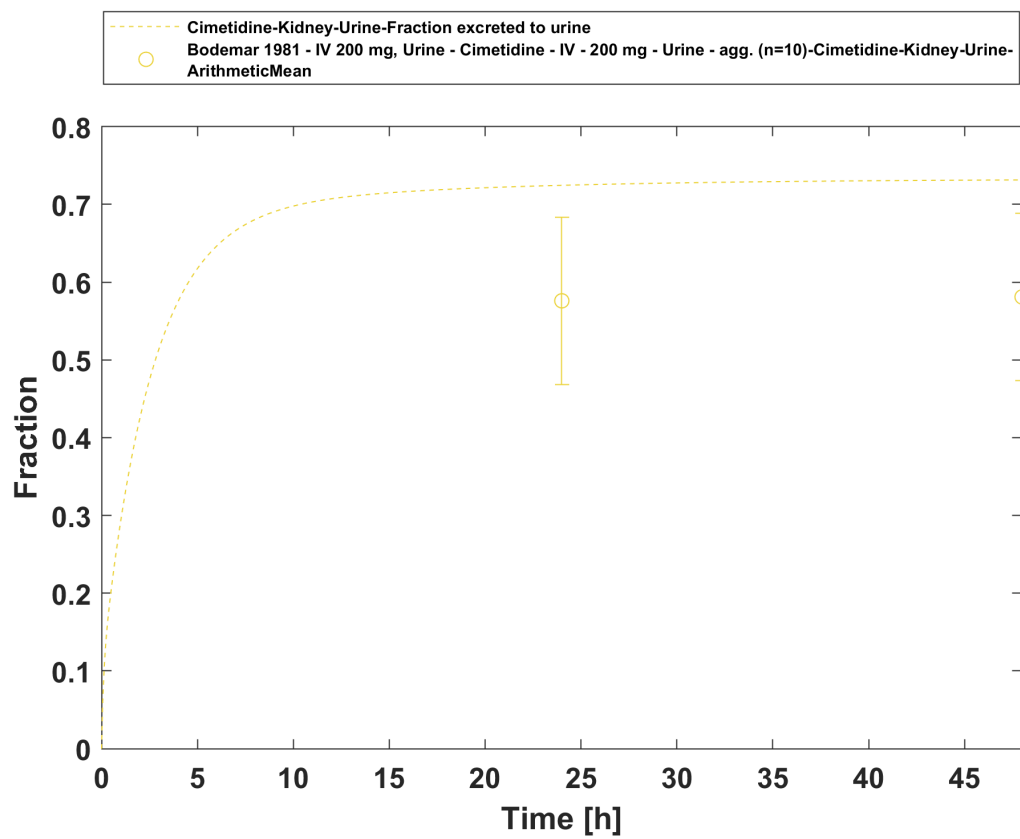
### 3.3.1 Model Building



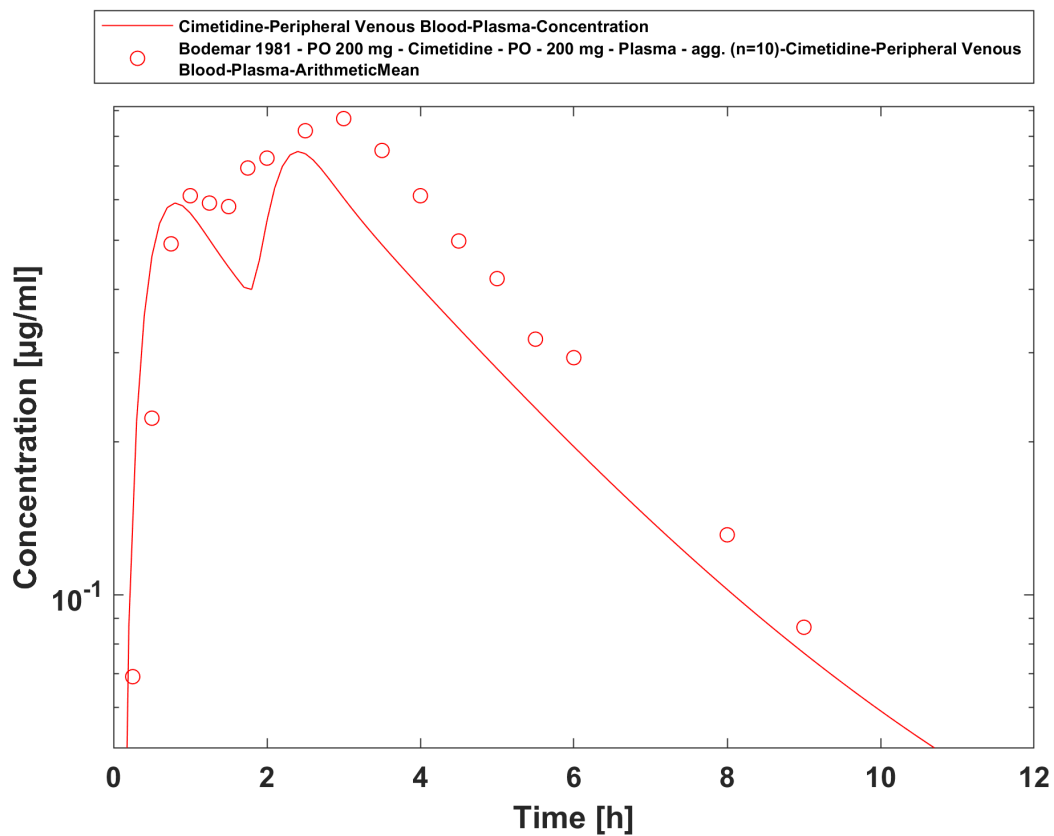
iv 200 mg (5 min),Morgan 1983, n=6



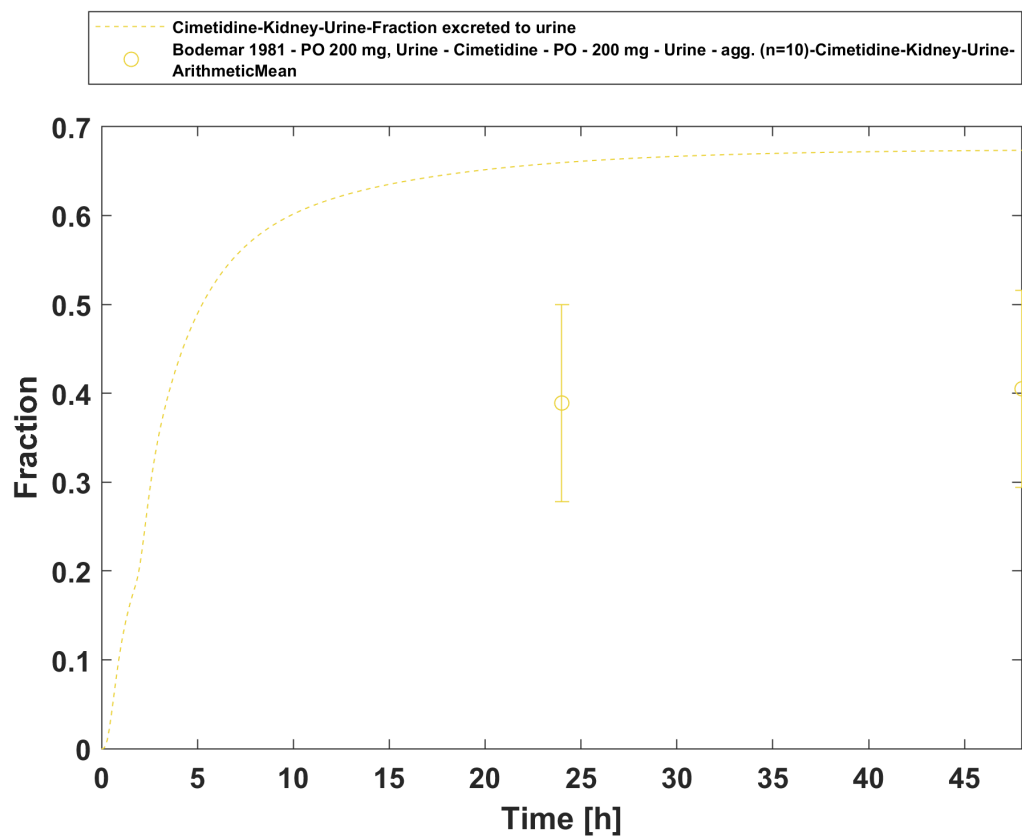
iv 200 mg, Bodemar 1981, n=10



iv 200 mg, Bodemar 1981, n=10, urine



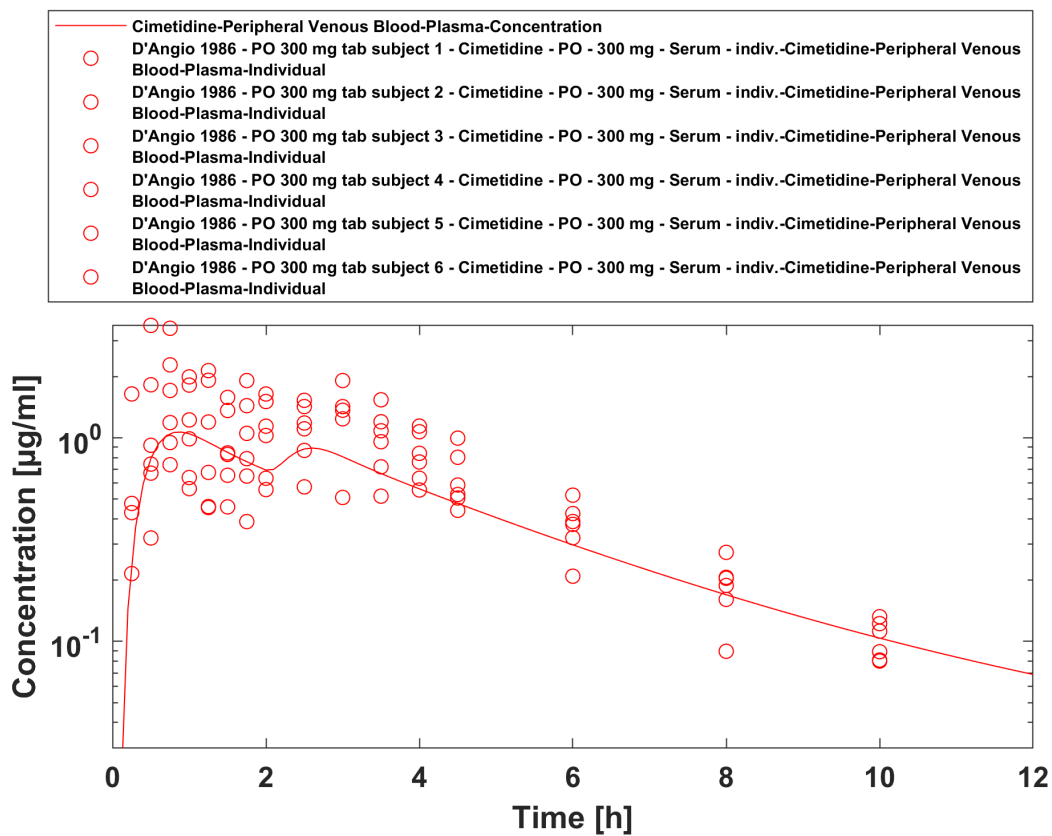
po 200 mg, Bodemar 1981, n=10



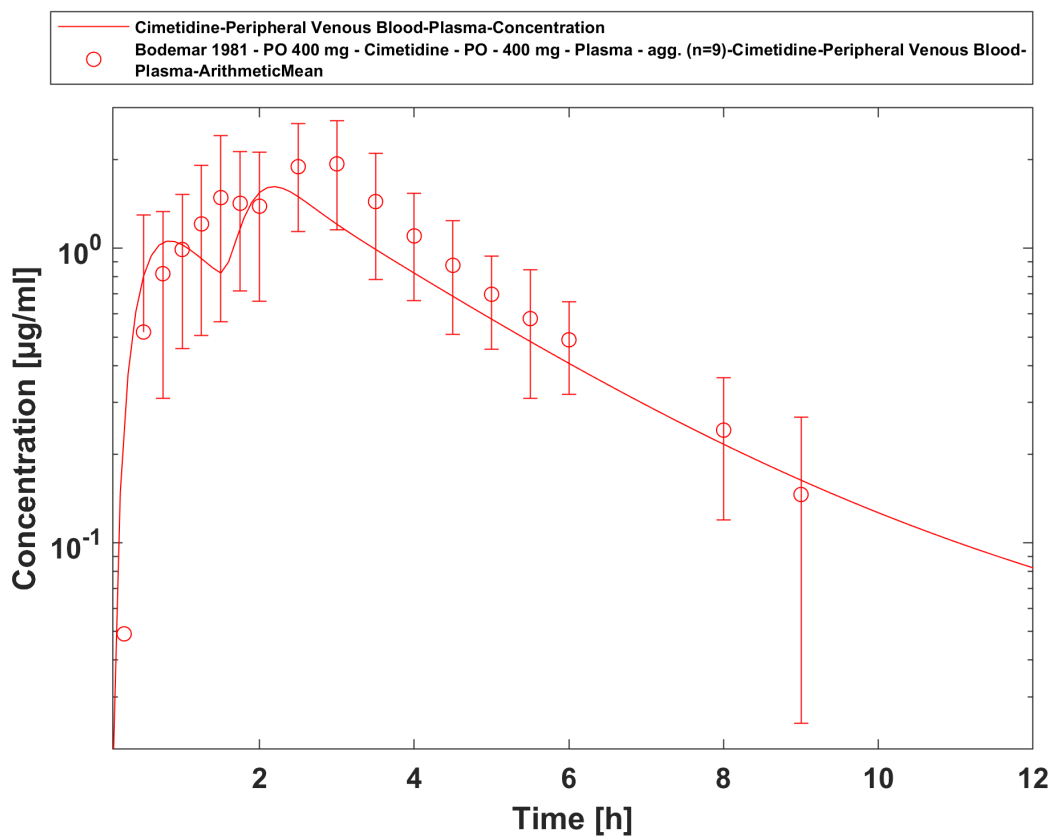
po 200 mg, Bodemar 1981, n=10, urine



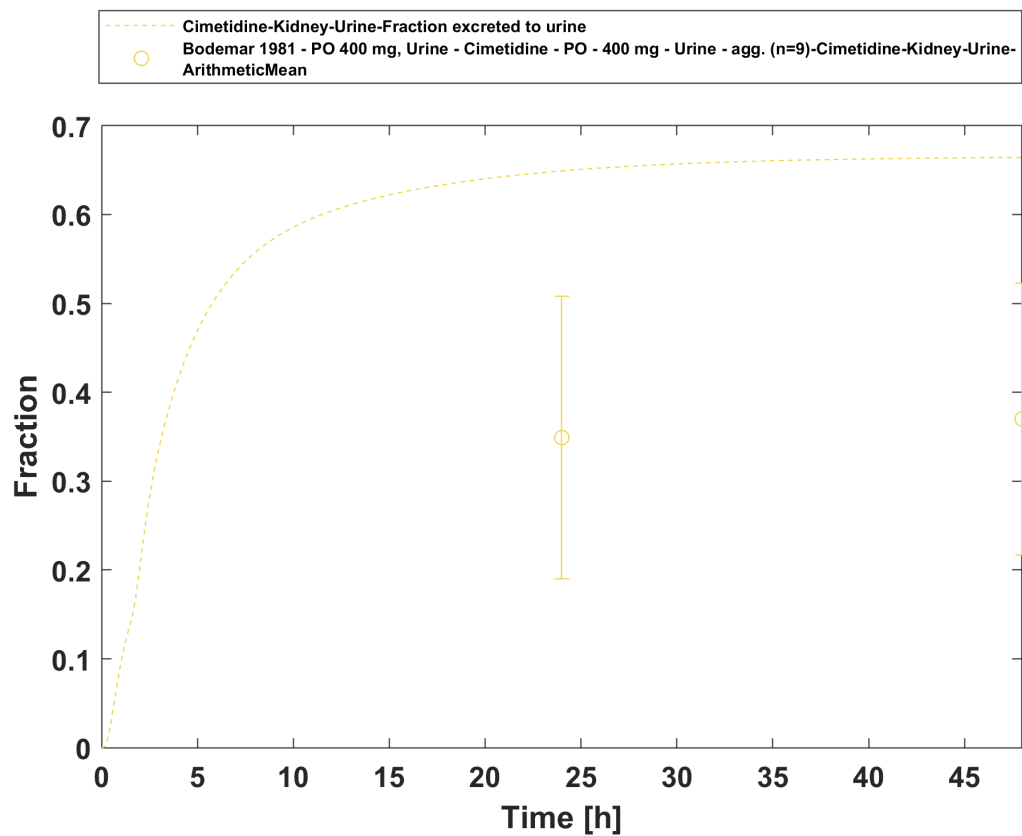




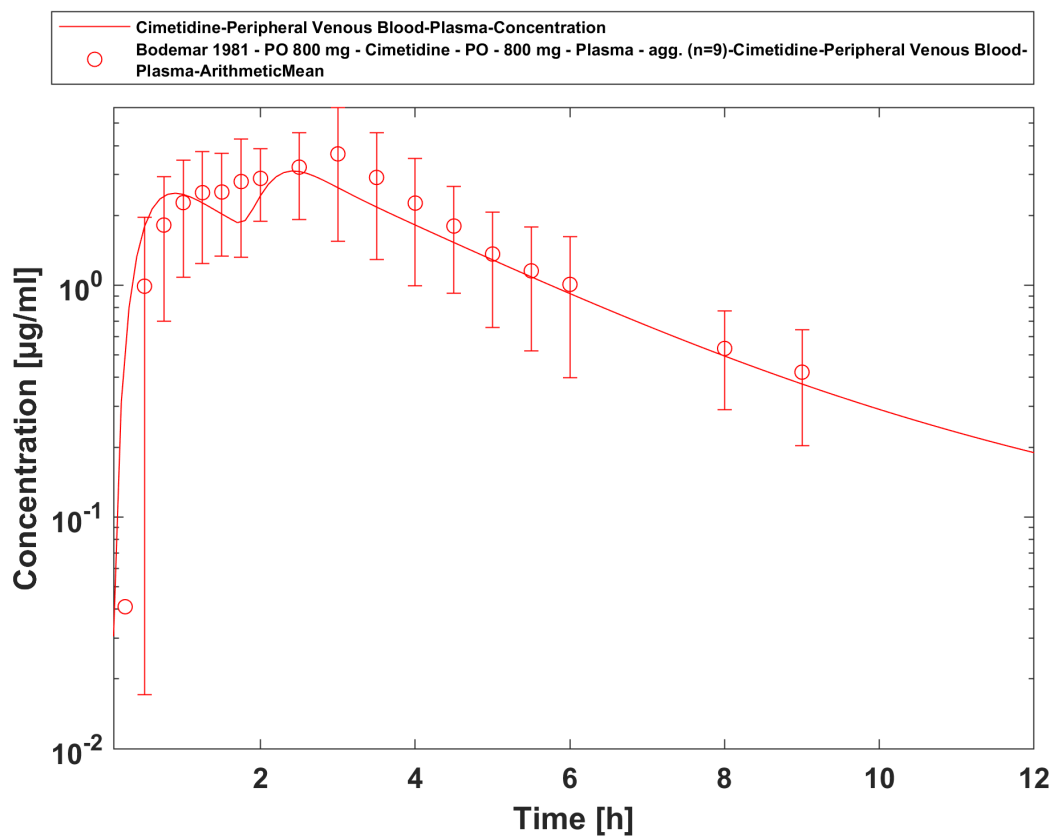
po 300 mg (tab), D'Angio 1986, n=6



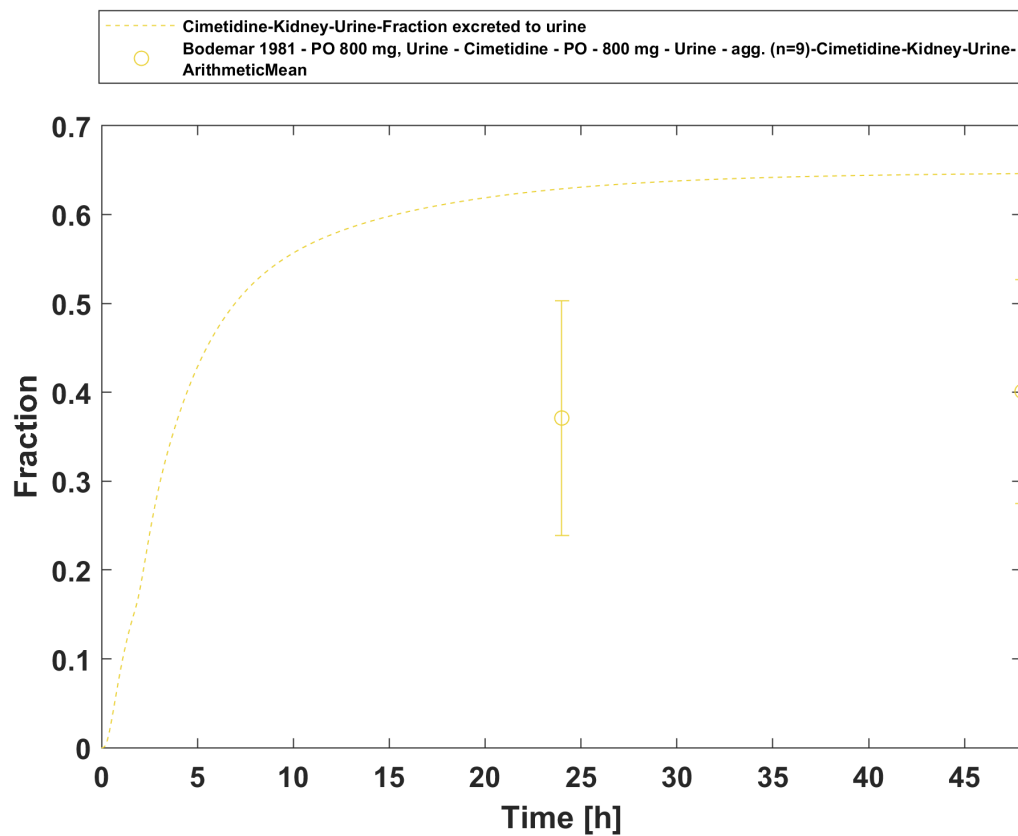
po 400 mg, Bodemar 1981, n=9



po 400 mg, Bodemar 1981, n=9, urine

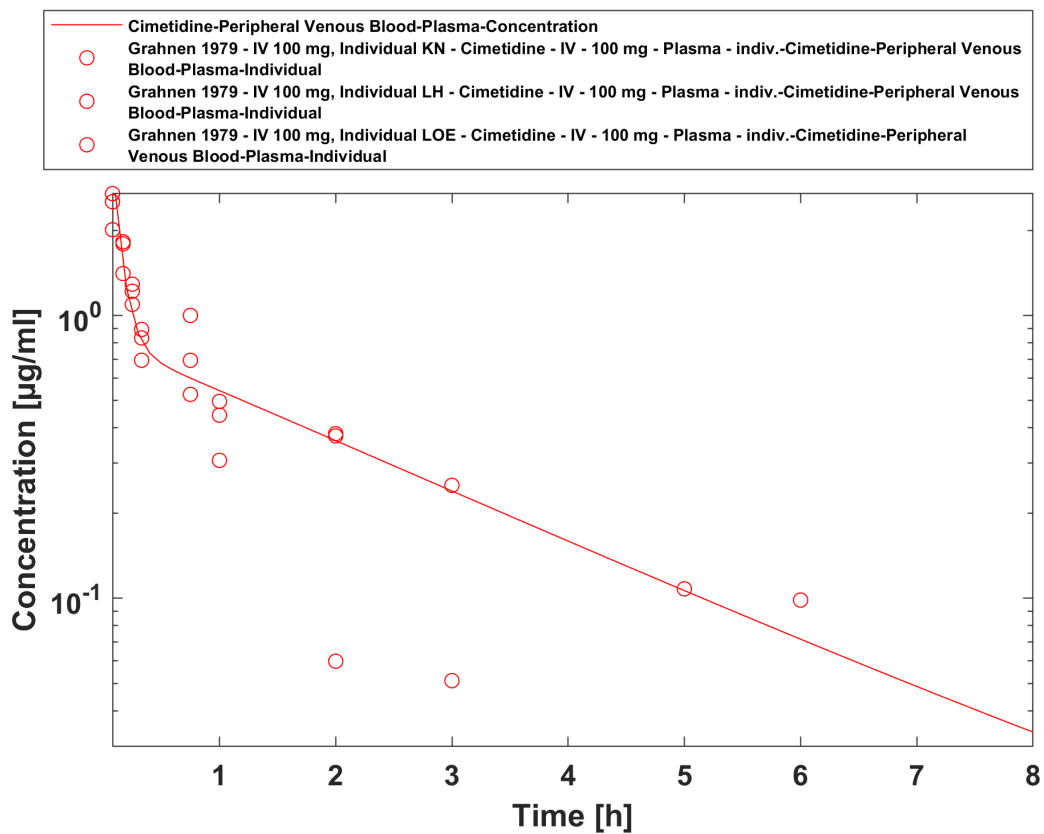


po 800 mg, Bodemar 1981, n=9

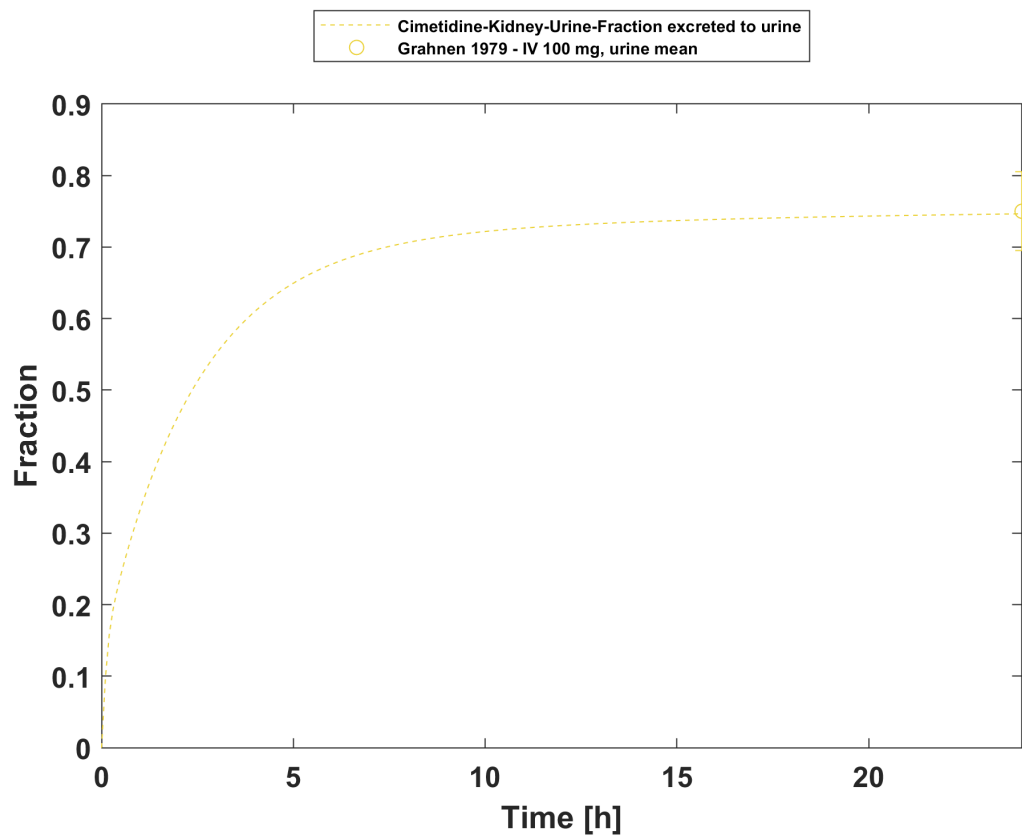


po 800 mg, Bodemar 1981, n=9, urine

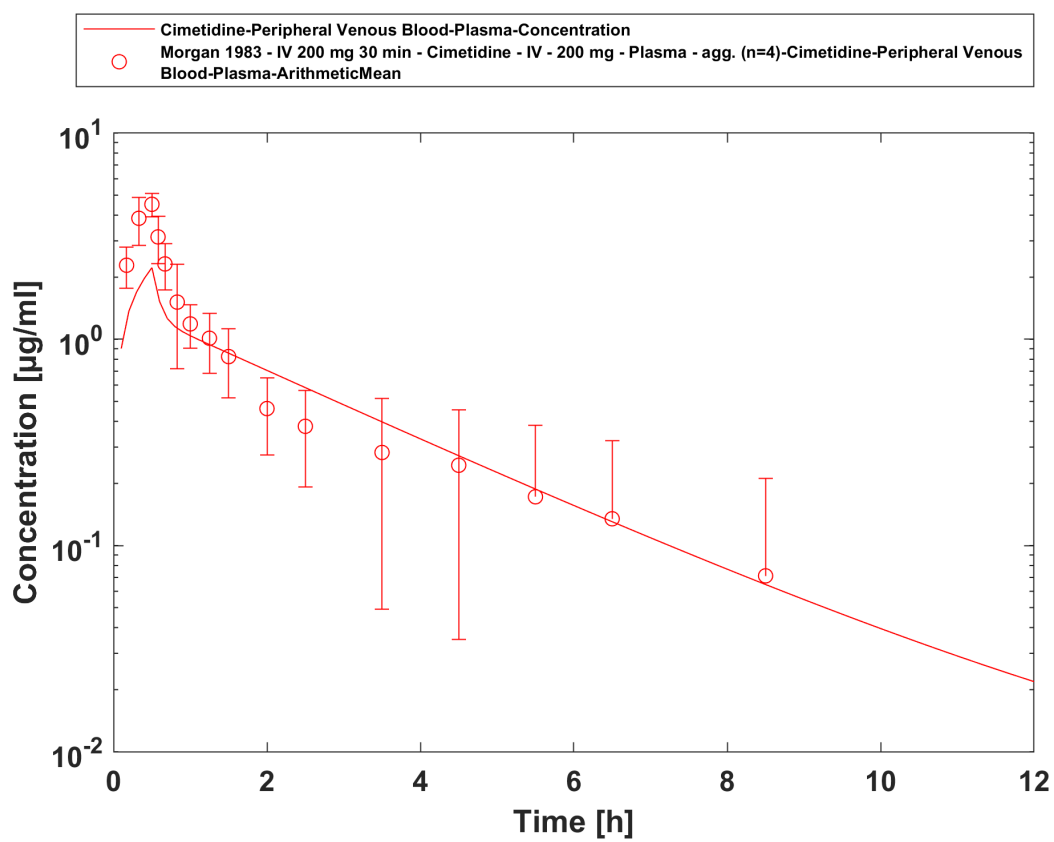
### 3.3.2 Model Validation



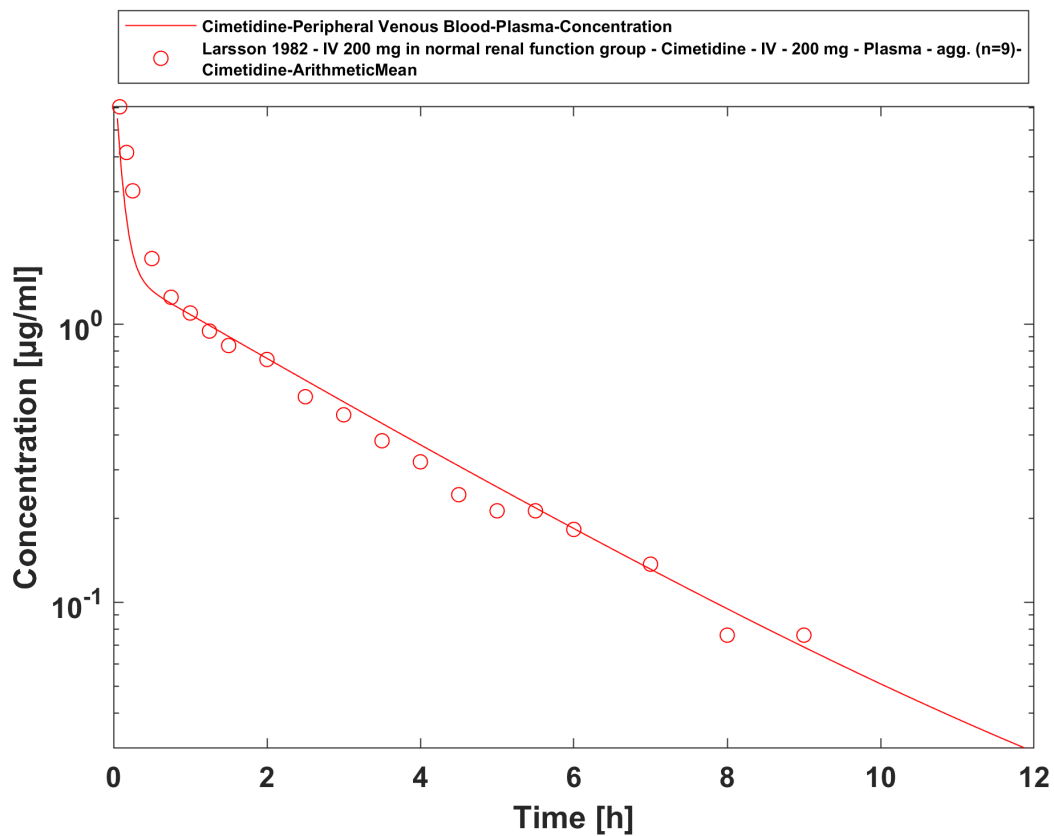
iv 100 mg (5 min), Grahnén 1979, n=3



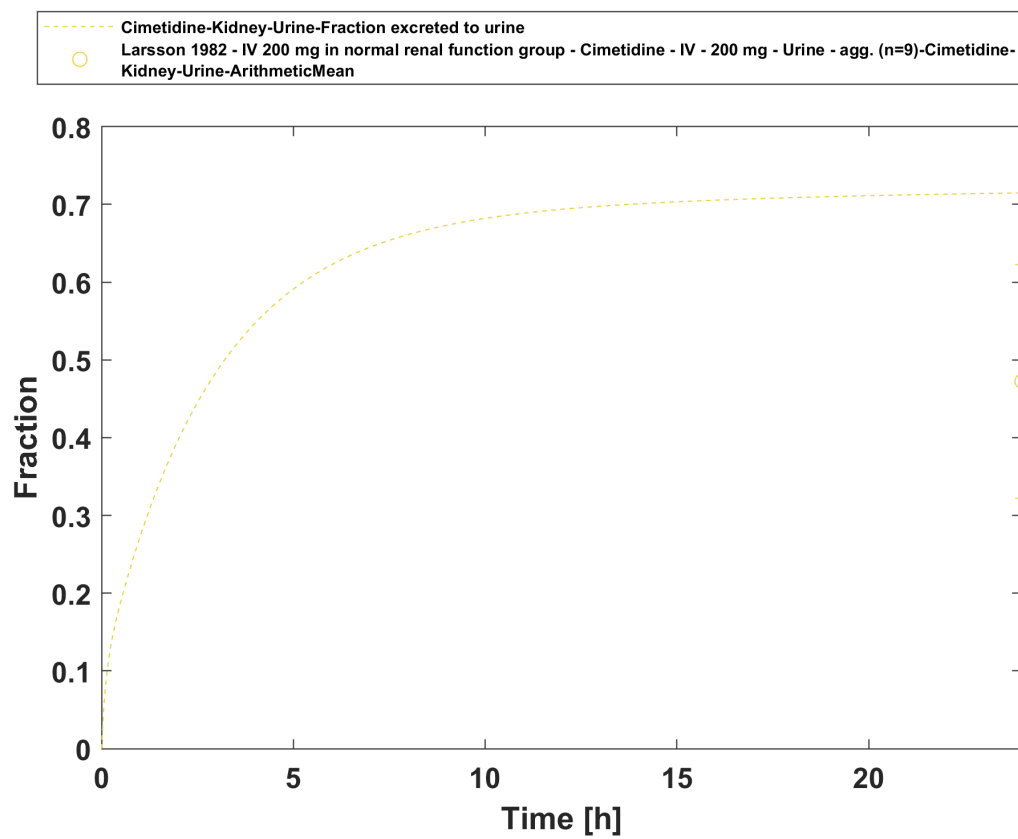
iv 100 mg (5 min), Grahnén 1979, n=3, urine



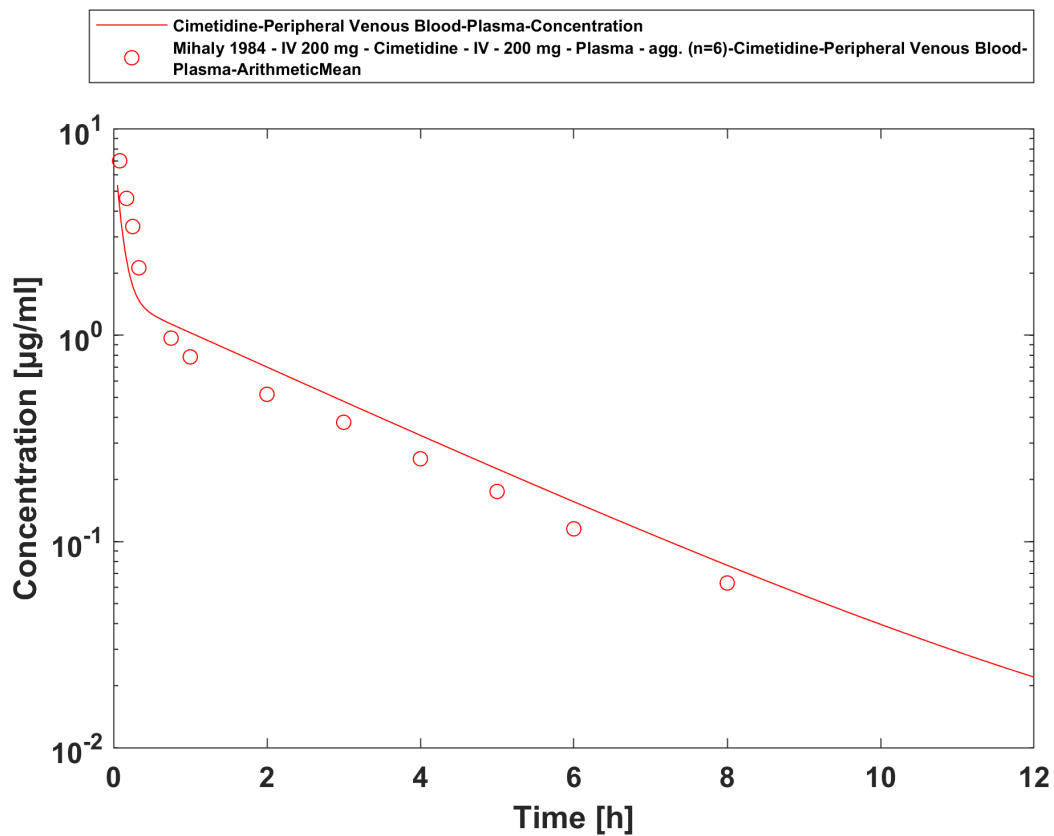
iv 200 mg (30 min), Morgan 1983, n=4



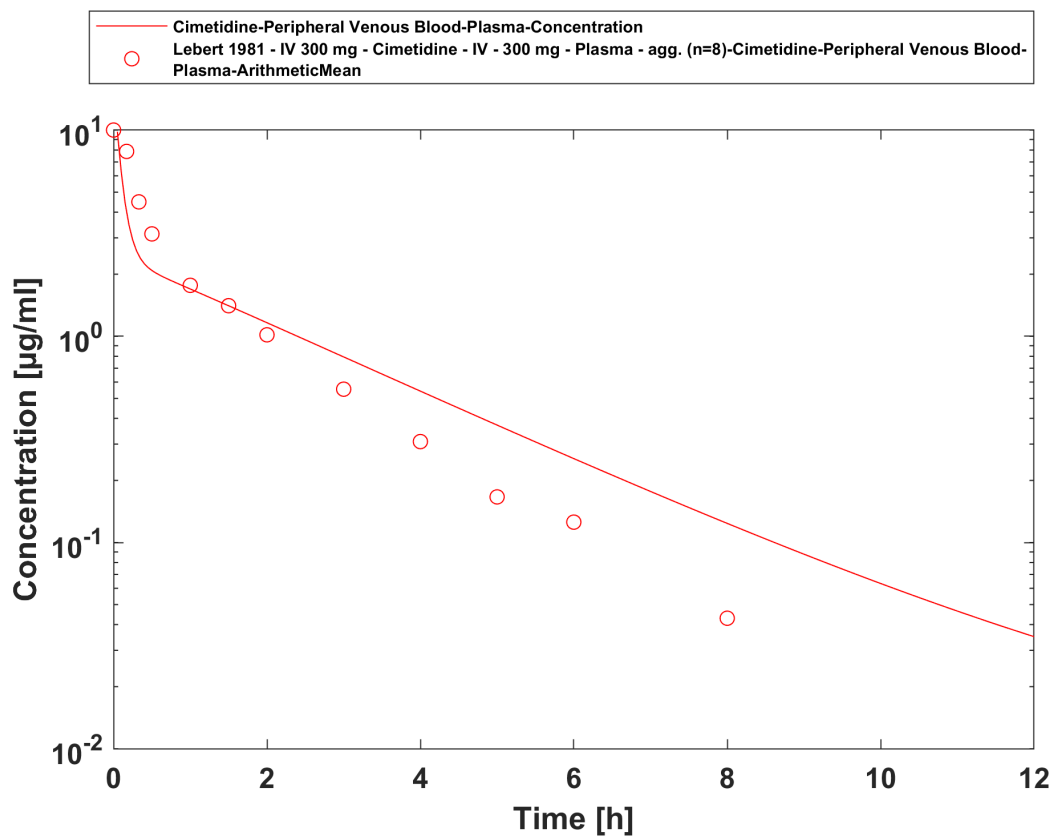
iv 200 mg, Larsson 1982, n=9



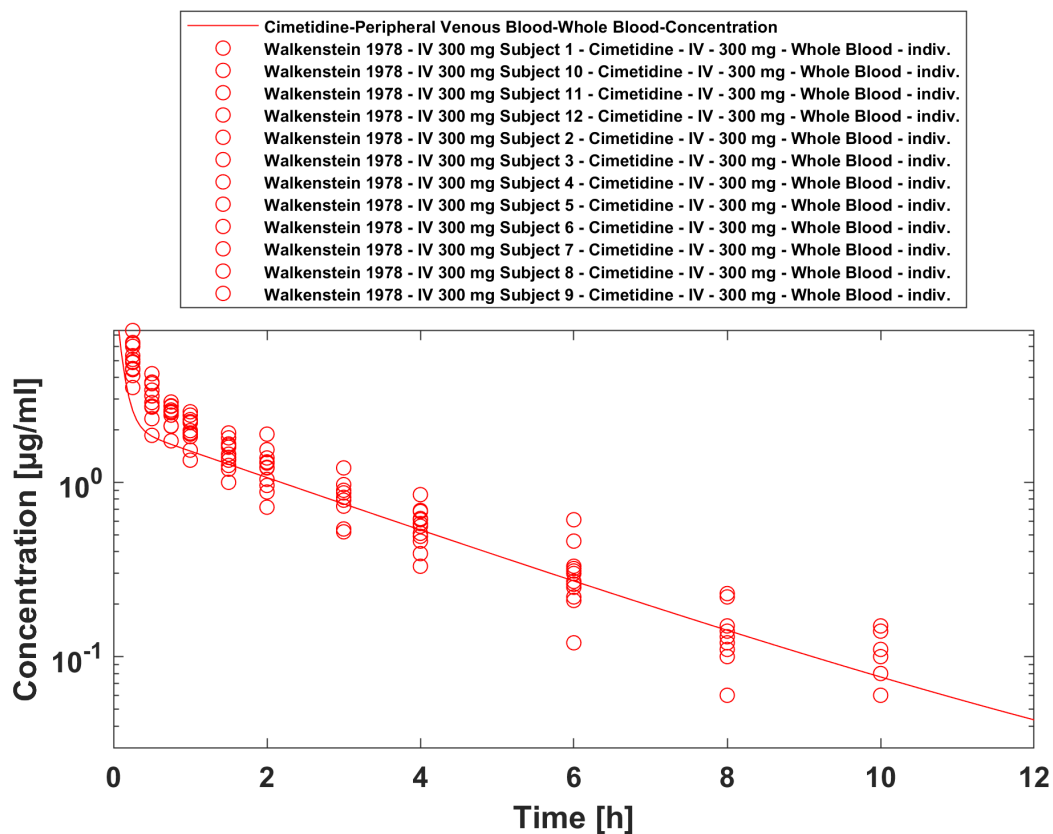
iv 200 mg, Larsson 1982, n=9, urine



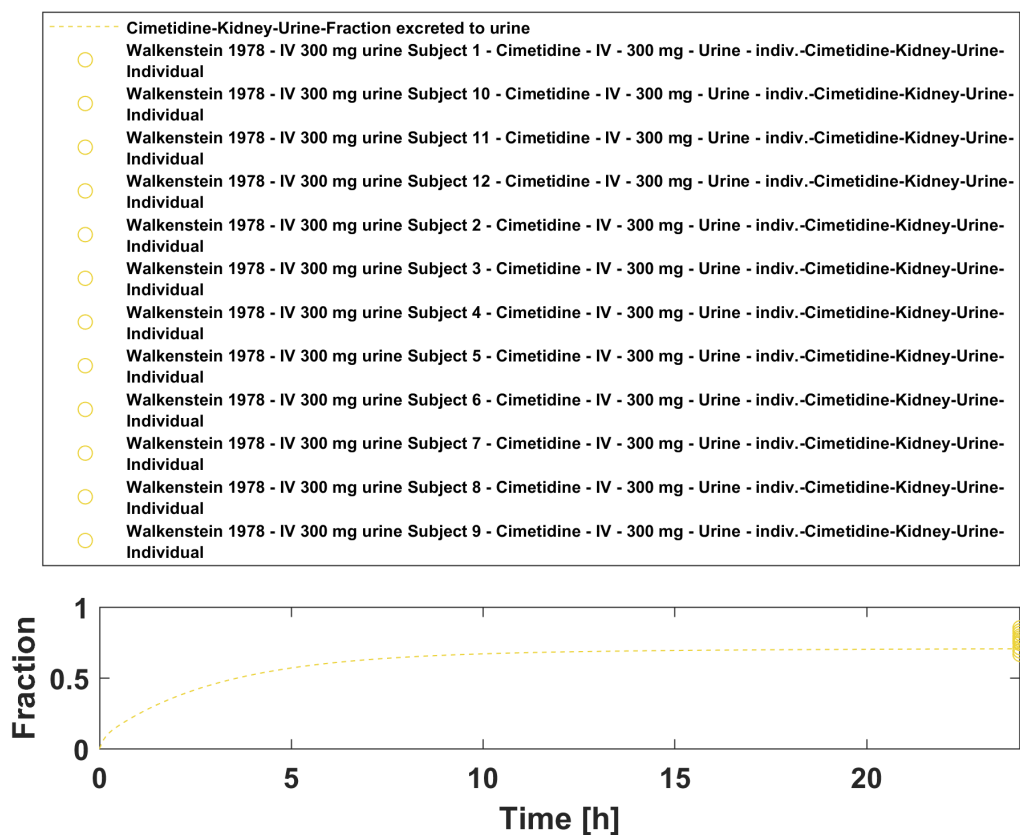
iv 200 mg, Mihaly 1984, n=6



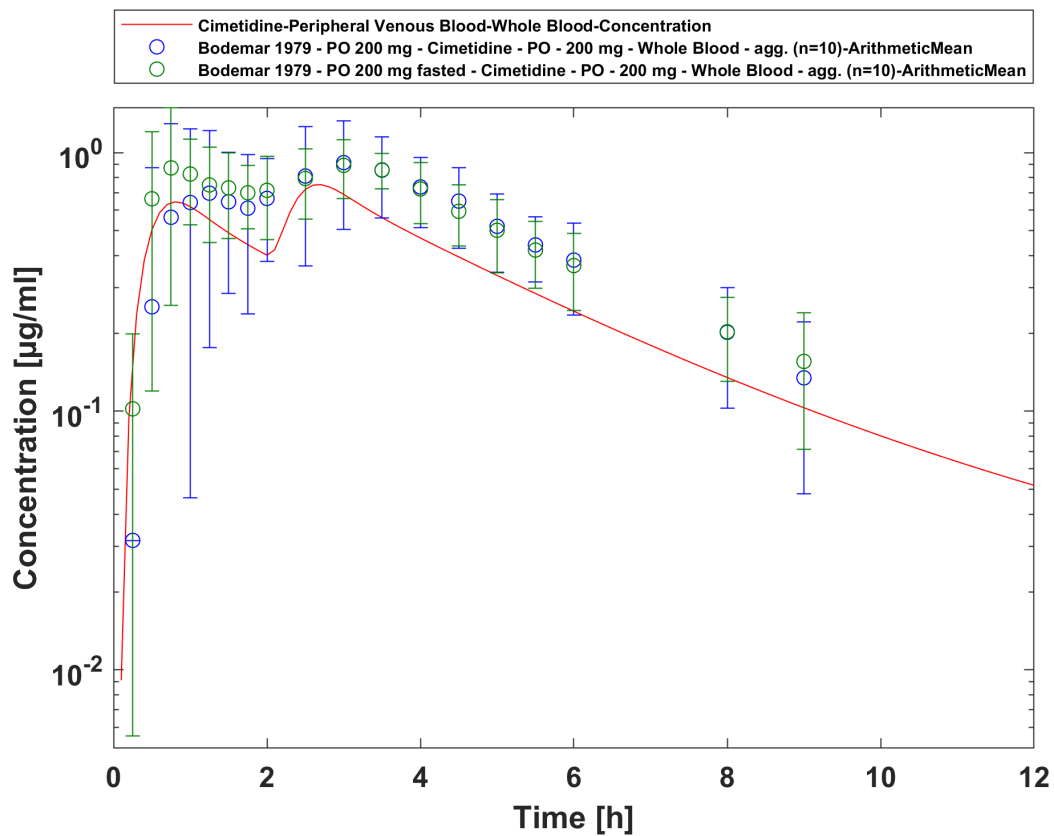
iv 300 mg (2 min), Lebert 1981, n=1



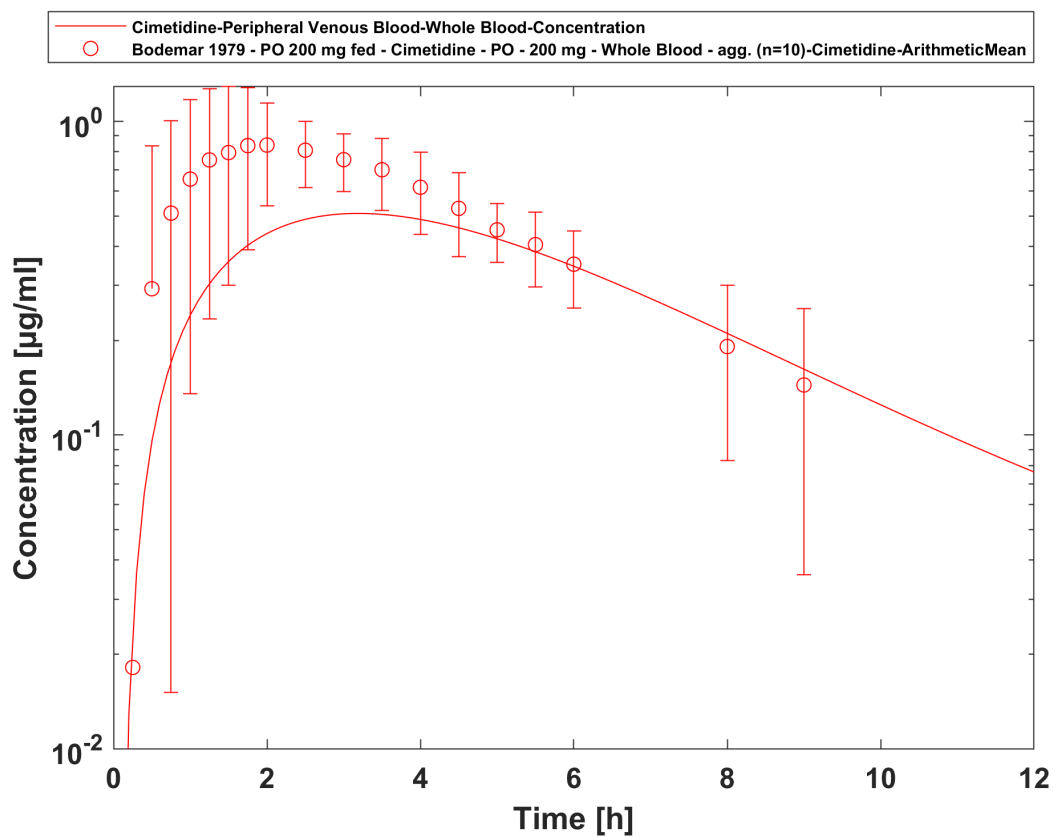
iv 300 mg (2 min), Walkenstein 1978, n=12



iv 300 mg (2 min), Walkenstein 1978, n=12

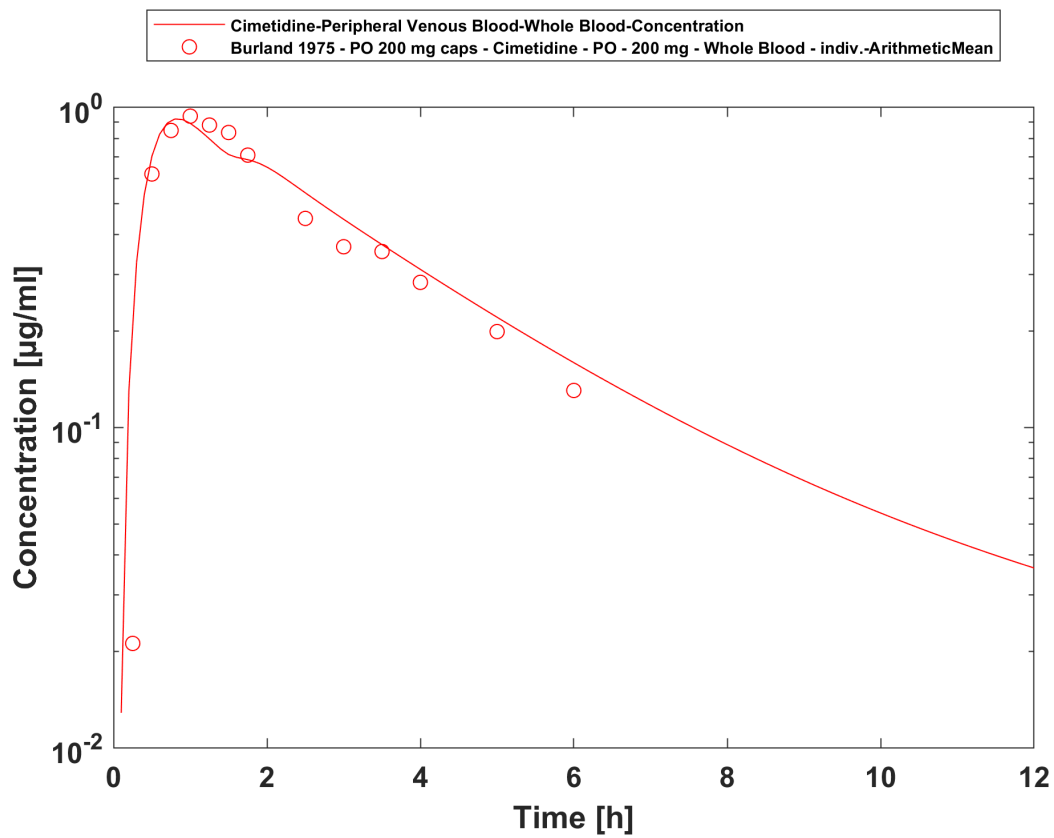


po 200 mg (tab), Bodemar 1979 (fasted)

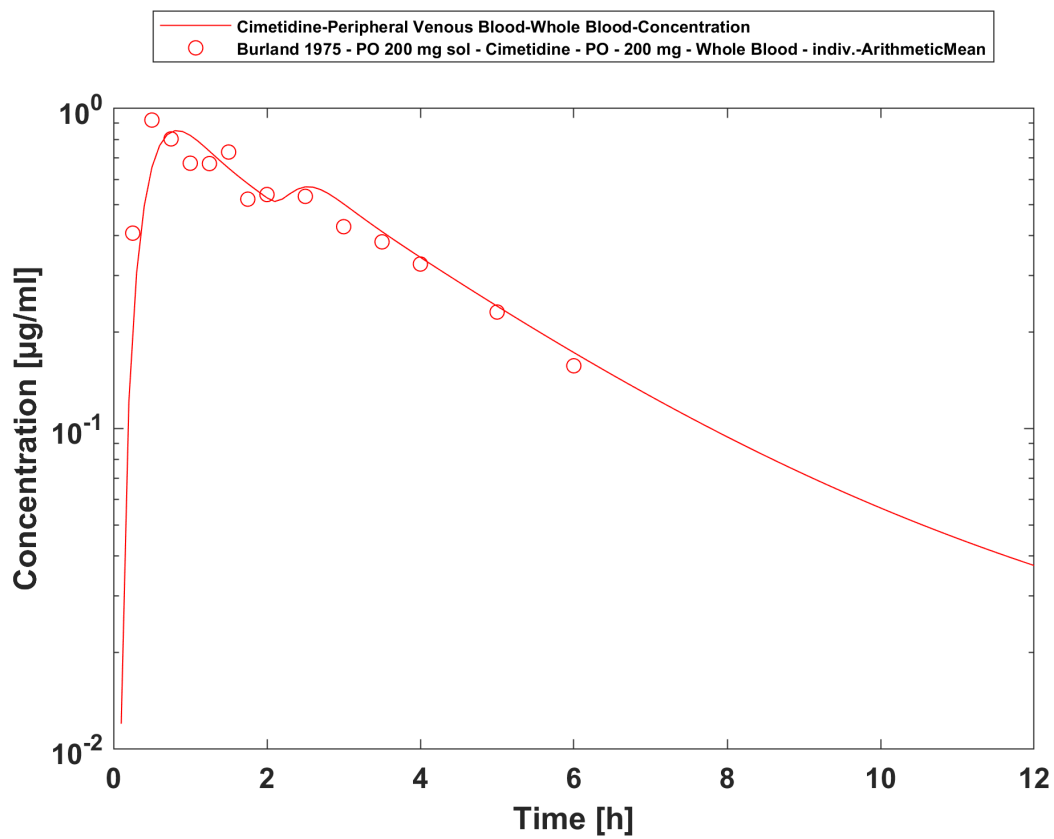


po 200 mg (tab), Bodemar 1979 (fed)

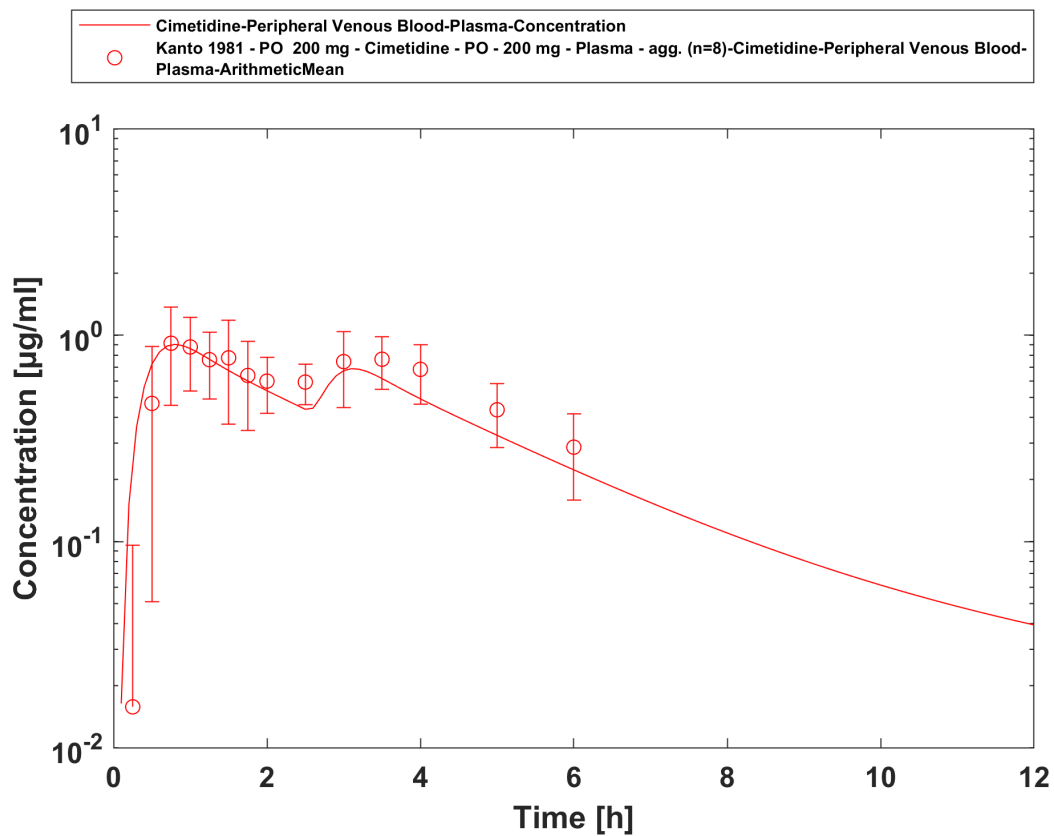




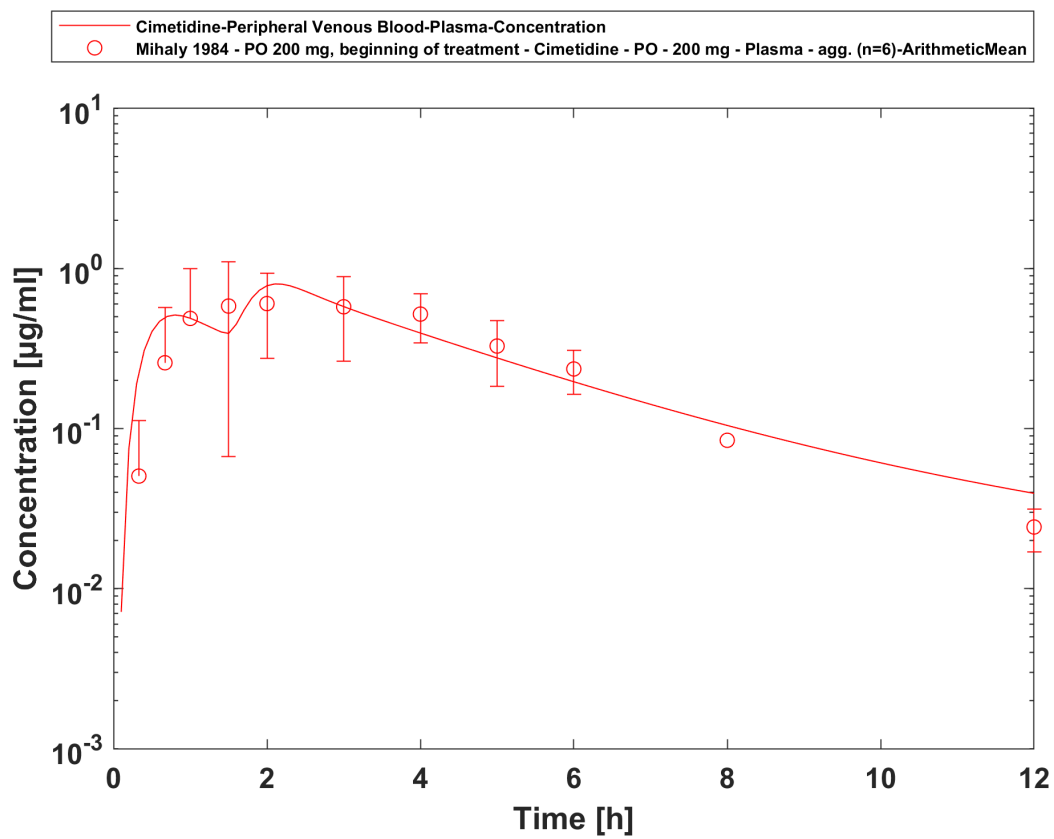
po 200 mg, Burland 1975, caps



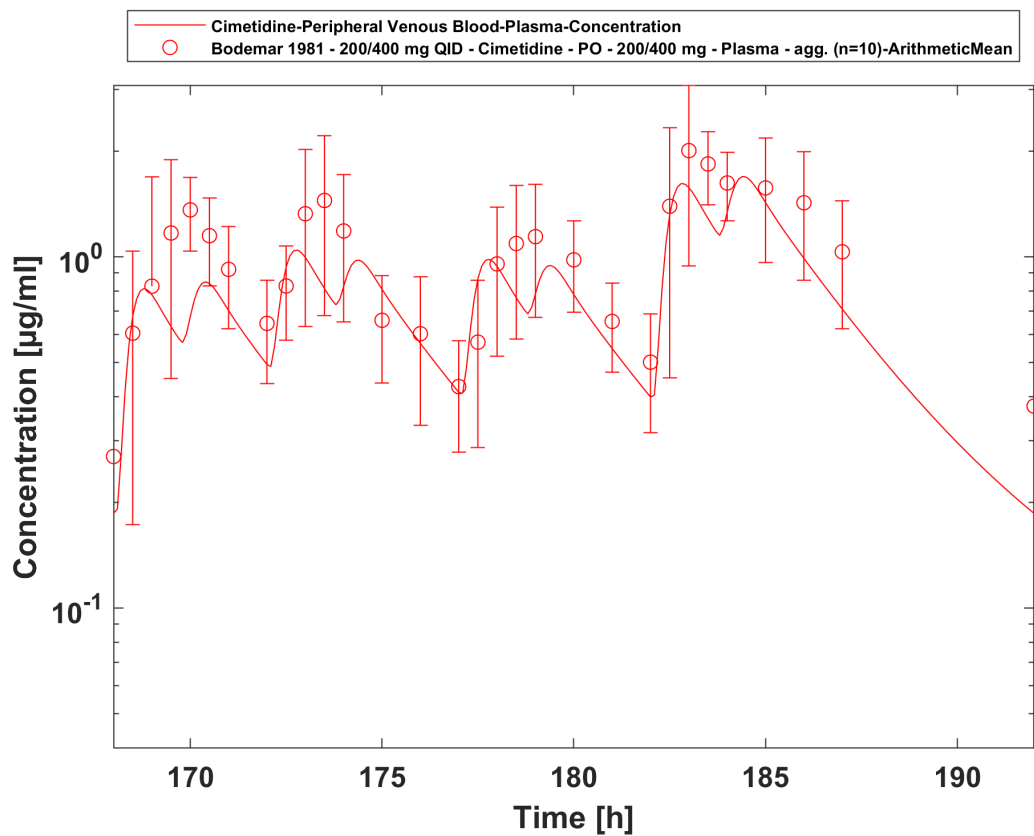
po 200 mg, Burland 1975, sol



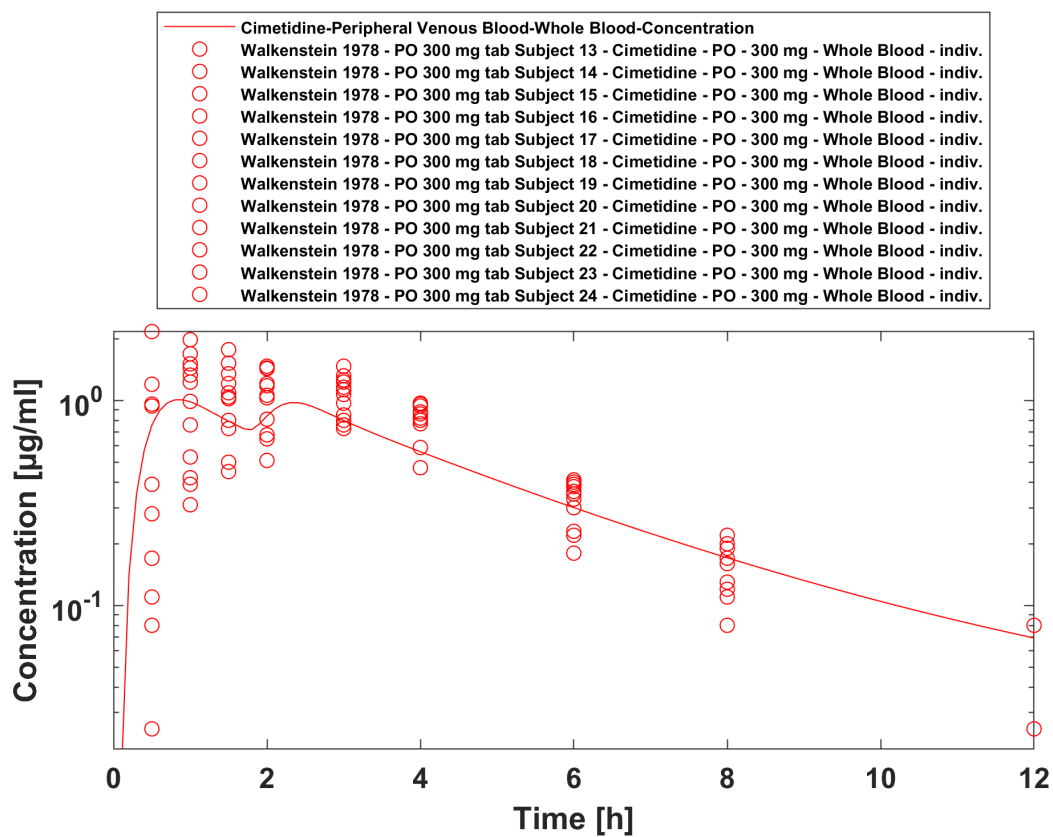
po 200 mg, Kanto 1981, n=8



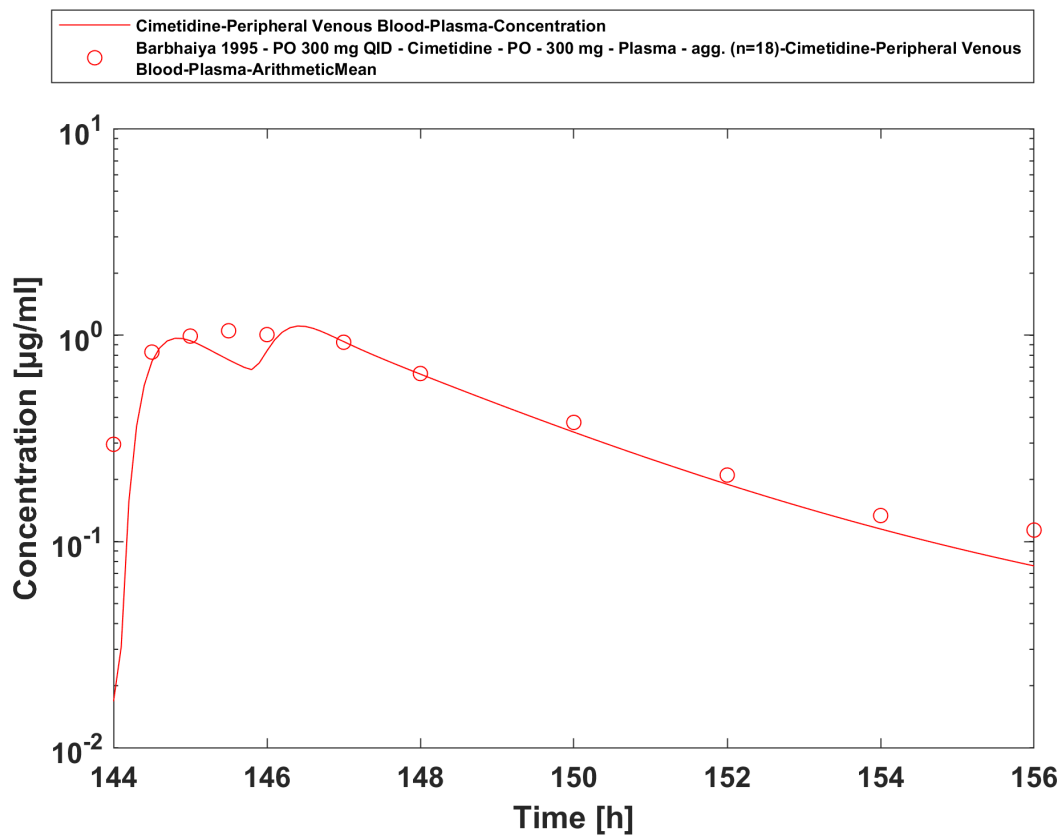
po 200 mg, Mihaly 1984, n=8



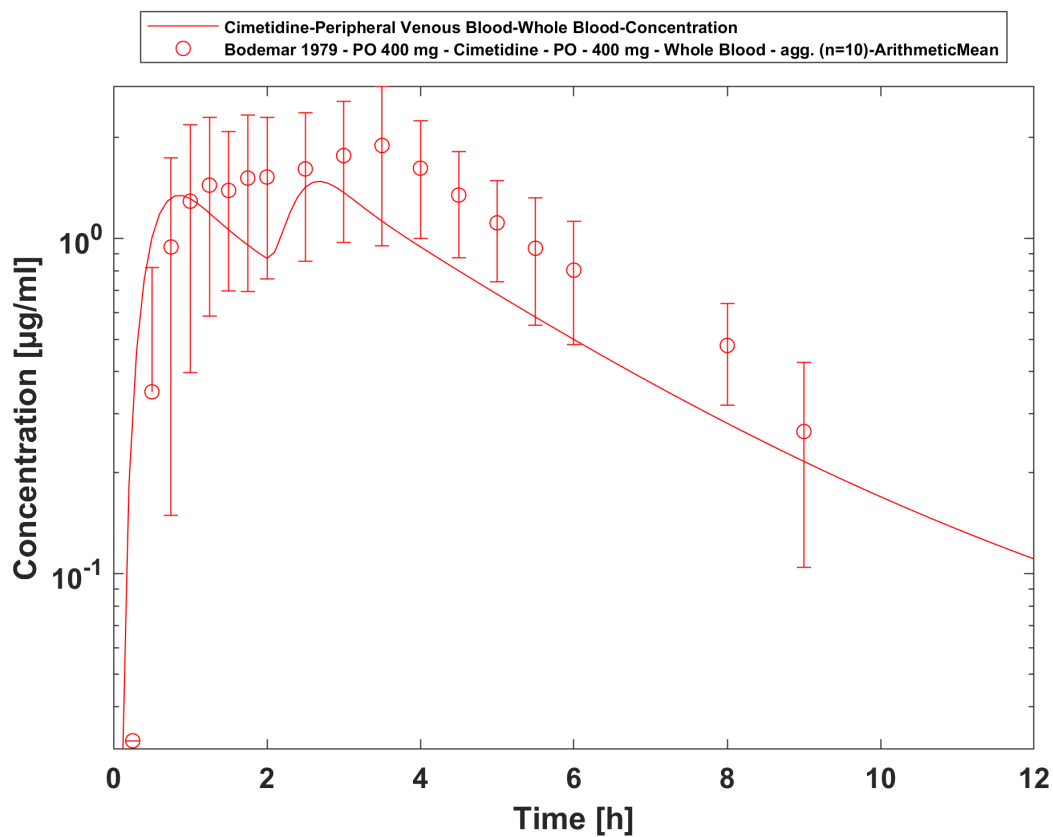
po 200/400 mg QID, Bodemar 1981 (fasted)



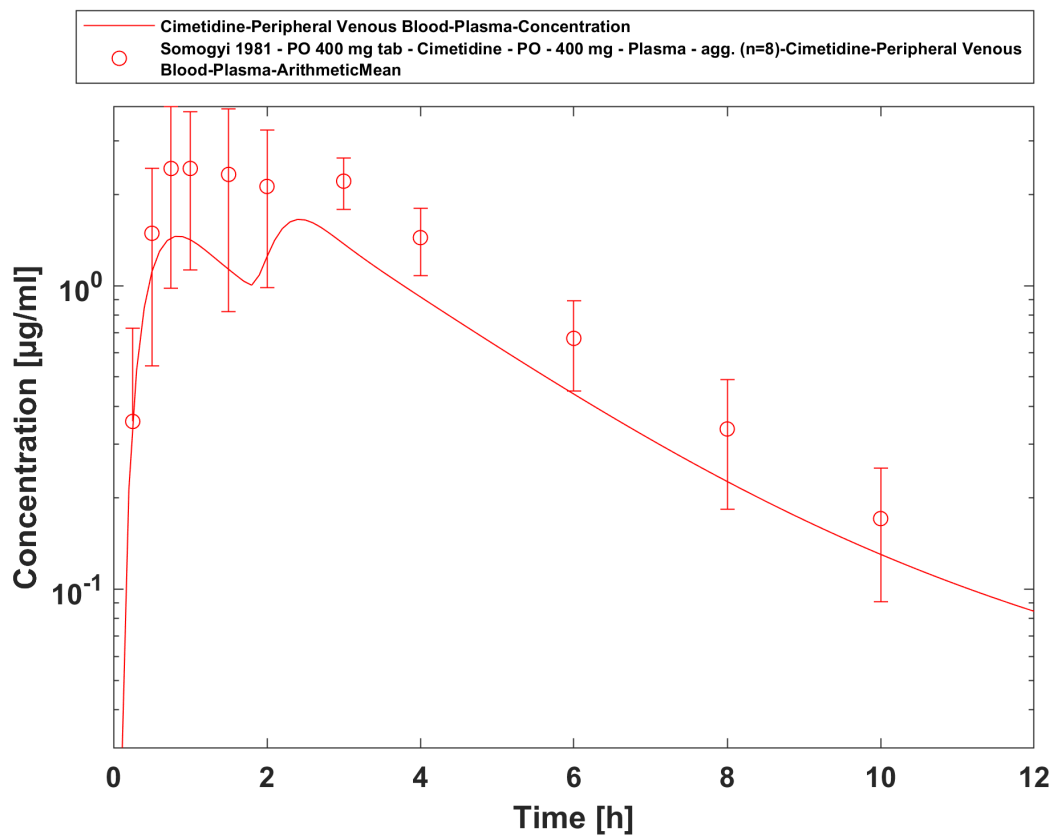
po 300 mg (tabl), Walkenstein 1978, n=12



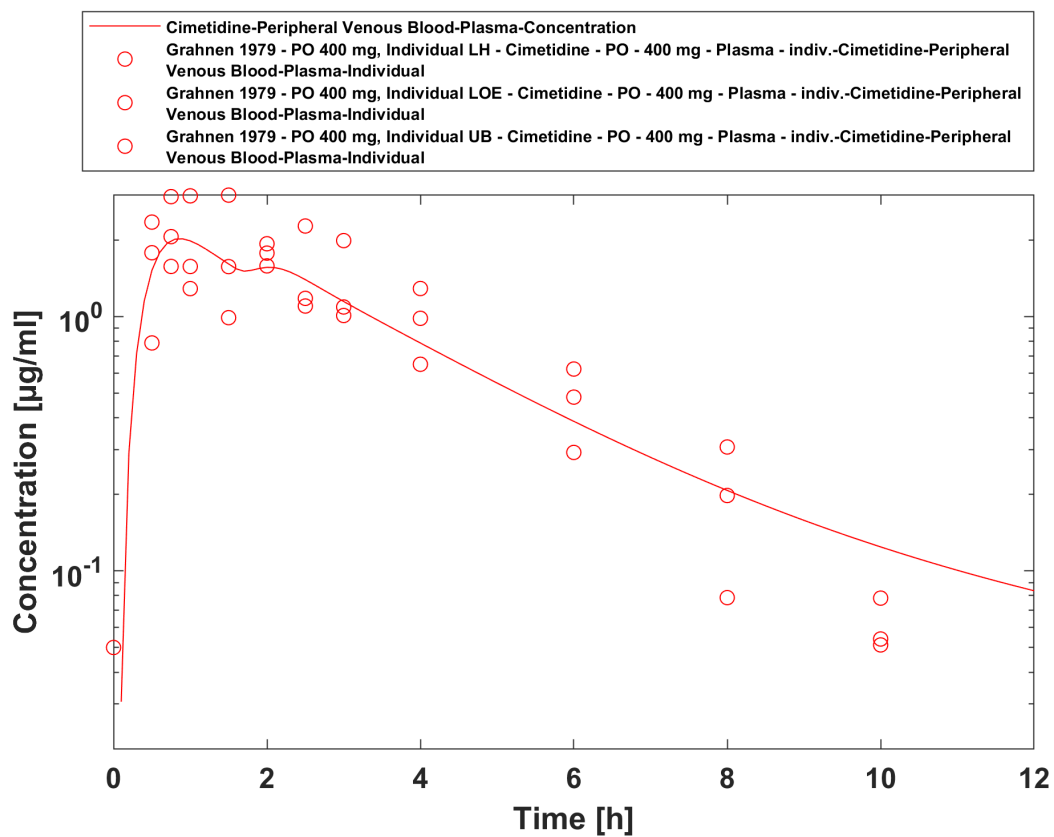
po 300 mg QID (sol), Barbhaiya 1995, n=18



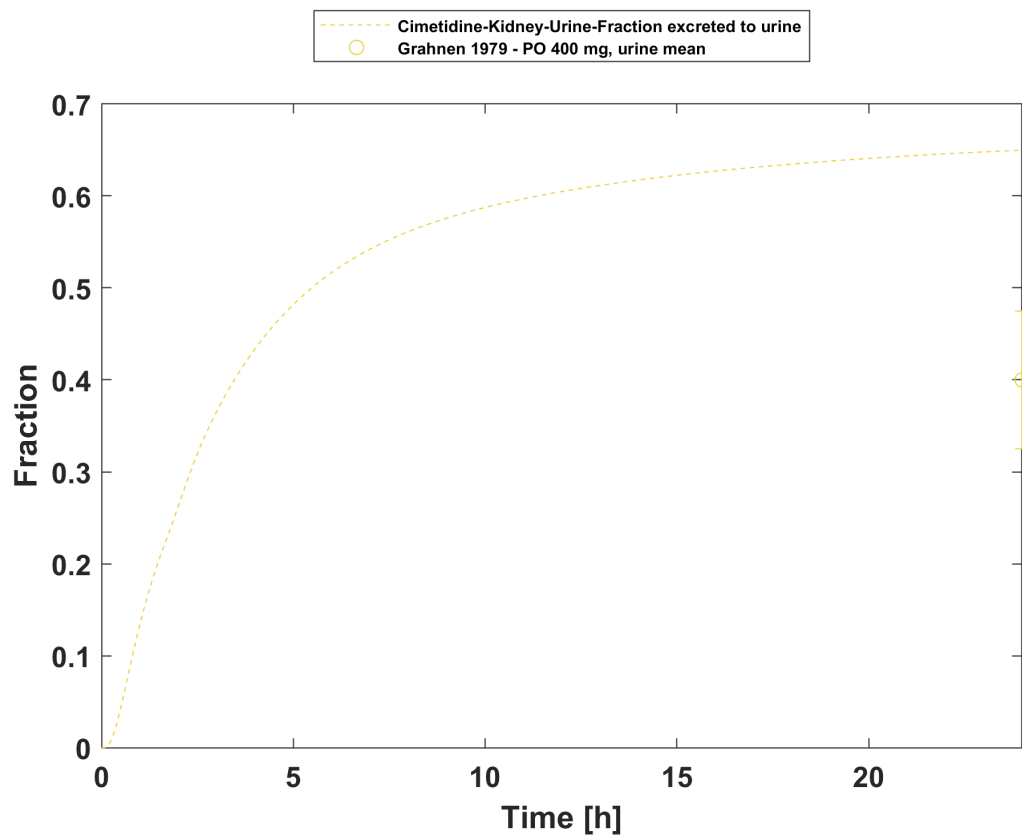
po 400 mg (tab), Bodemar 1979, n=10



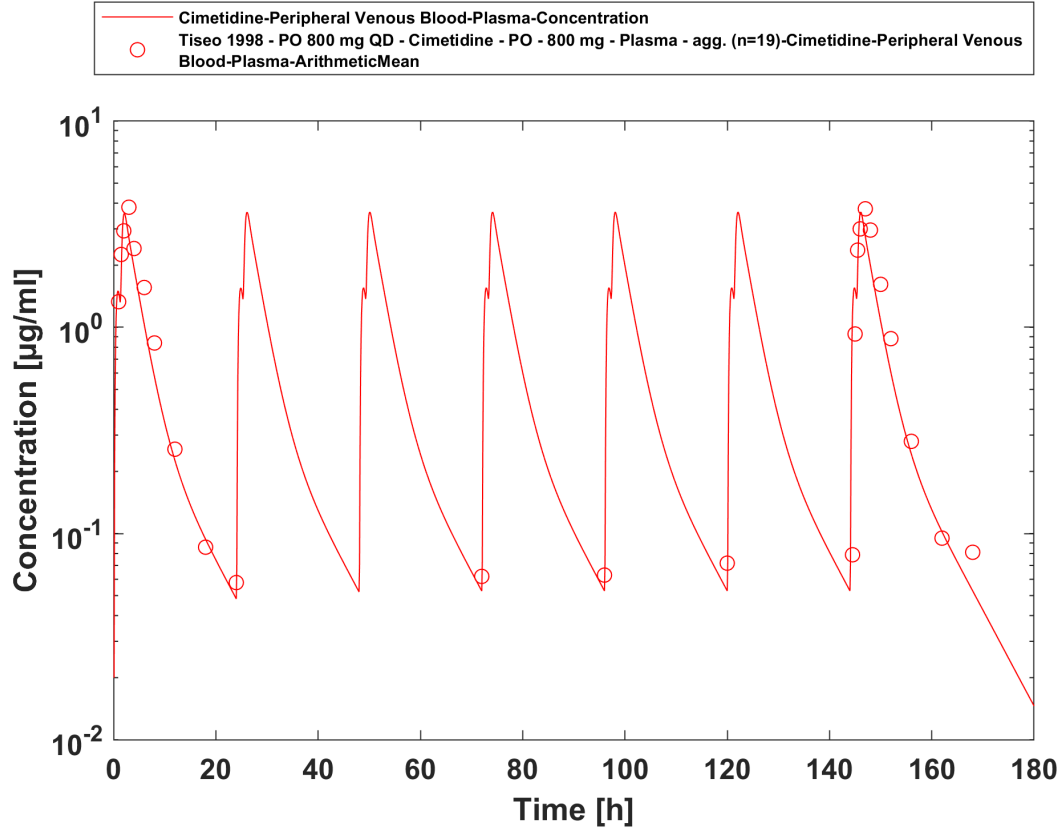
po 400 mg (tab), Somogyi 1981, n=8



po 400 mg (tab), Grahnen 1979, n=3



po 400 mg (tab),Grahnen 1979, n=3, urine



po 800 mg (tab) qd, Tiseo 1998, n=18

## 4 Conclusion

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The herein presented PBPK model adequately describes the pharmacokinetics of cimetidine after intravenous and oral administration of single and multiple doses to healthy adults and peptic ulcer patients covering a broad dosing range from 100 to 800 mg. The established cimetidine PBPK model is verified for the use as a mild inhibitor of CYP3A4 drug in drug-drug interaction simulations.

## 5 References

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- Avdeef 2001** Avdeef A, Berger CM. pH-metric solubility. 3. Dissolution titration template method for solubility determination. *Eur J Pharm Sci.* 2001 Dec;14(4):281-91. doi: 10.1016/s0928-0987(01)00190-7. PMID: 11684402.
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