

Qualification of CKD Populations

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

<https://github.com/Open-Systems-Pharmacology/Qualification-CKD>

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1 Introduction

The presented qualification report performs simulations of drug disposition and pharmacokinetics in subjects with chronic kidney disease (CKD) to qualify the parameterization of the foundational anatomy and physiology parameters for virtual populations with CKD in the Open Systems Pharmacology (OSP) Suite.

CKD is defined as a reduction in kidney function as approximated by the estimated glomerular filtration rate (eGFR) below 60 mL/min/1.73m² for at least 3 months. CKD is a systemic and multifaceted disease that alters many body systems beyond the glomerular filtration rate. As summarized by Malik et al. (2020),¹ the disease can affect anatomy and physiology parameters such as kidney volume, kidney blood flow, plasma protein binding, hematocrit, gastric emptying time, small intestinal transit time, and colonic transit time. These quantitative changes have been implemented into the Open Systems Pharmacology Suite for the purpose of building mechanistic models of drug disposition, termed physiologically based pharmacokinetic (PBPK) models. The present implementation considers subjects with CKD from Stage 3 to Stage 5 (end-stage renal disease) who have not yet begun treatment with dialysis.

Beyond foundational anatomy and physiology parameters, the disease can also affect the activities or expression of metabolic enzymes (such as CYP3A4) and active drug transporters (such as the organic anion transporters, OAT1 and OAT3). Because of the varied and conflicting literature on the effects of the disease on numerous enzymes and transporters, the present parameterization is concerned only with foundational anatomy and physiology parameters that would be consistently applicable to all mechanistic simulations in CKD. When building a PBPK model for subjects with CKD, the OSP user is recommended to consult literature specific to the effect of disease on enzymes and transporters that are most relevant to the disposition of the drug in question and adapt the model parameters (e.g., activity of the enzyme) accordingly.

This report replicates in part the evaluation of the parameterization done by Malik et al. Section 2 describes the algorithm used to generate CKD populations. Section 3 evaluates the predicted effects of CKD on pharmacokinetics for three compounds: Gabapentin, Acebutolol, Atenolol. Gabapentin is selected as an index compound that is predominantly eliminated by glomerular filtration. Atenolol additionally undergoes active tubular secretion through the organic cation transporter (OCT) system, one system whose function is largely conserved in the disease. Acebutolol also undergoes secretion through the OCT system, but has a non-renal component (hepatic metabolism) that is suggested to be unaffected by the disease. Evaluation for each compound is split into two steps: an initial model calibration against a healthy population, followed by an extension to populations of subjects with relevant staging of CKD.

2 CKD Population Development

In order to generate a virtual population of subjects with CKD, the user must specify a target range for glomerular filtration rate (GFR) (minimum to maximum). PK-Sim will first generate a healthy population using the standard algorithm² - including the effects of aging as appropriate - and modify the physiological parameters of the simulated individuals to replicate a realistic CKD phenotype. The details of this process are outlined in **Figure 2-2**, with references to **Figure 2-1**, **Table 2-1** and **Table 2-2**. For a complete description, see Malik et al.¹

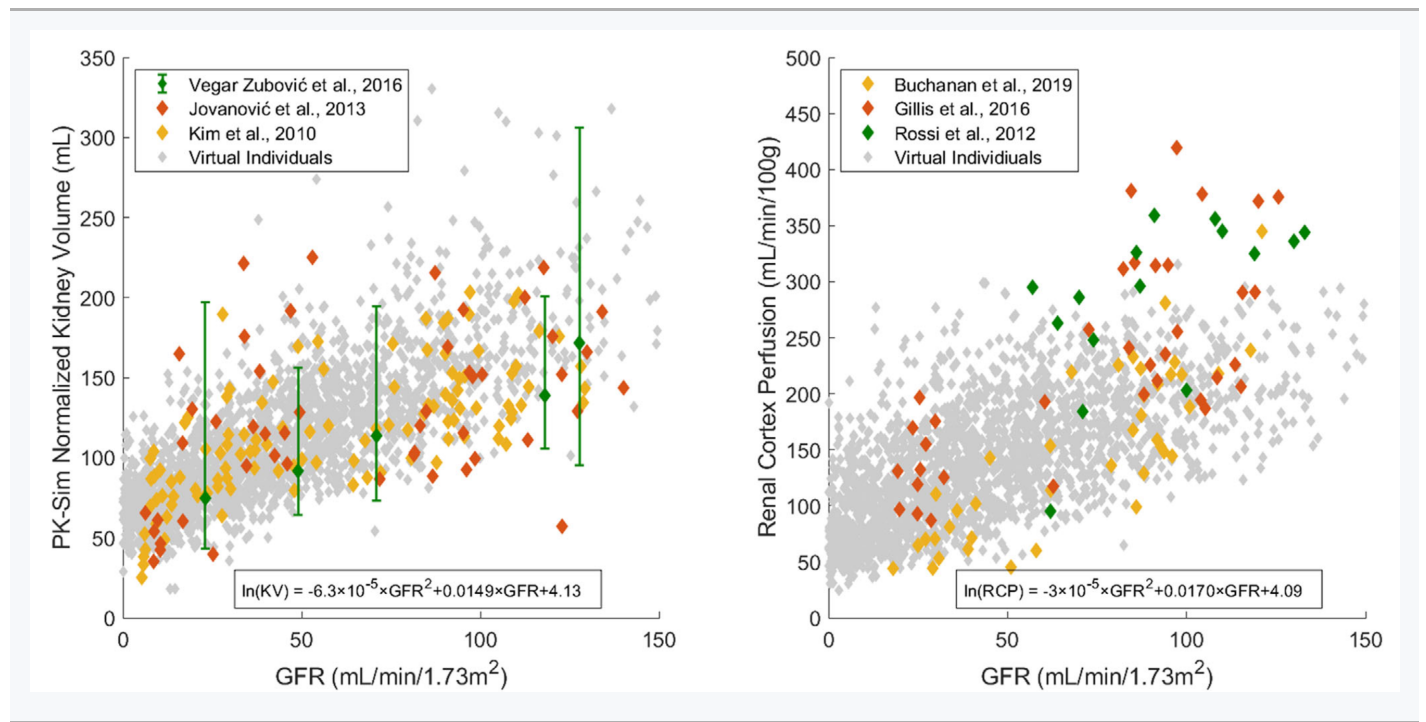


Figure 2-1: Simulated kidney volumes (left) and simulated renal cortex perfusion rates (right) in a population of adults aged 30-70 years with varying degrees of renal impairment compared with observed data from the literature.³⁻⁸ Diamonds represent individual data, whereas error bars represent the range in a study. In order to assign realistic physiological parameters to virtual individuals, quadratic equations for kidney volume and renal cortex perfusion rates were optimized to log-transformed data.

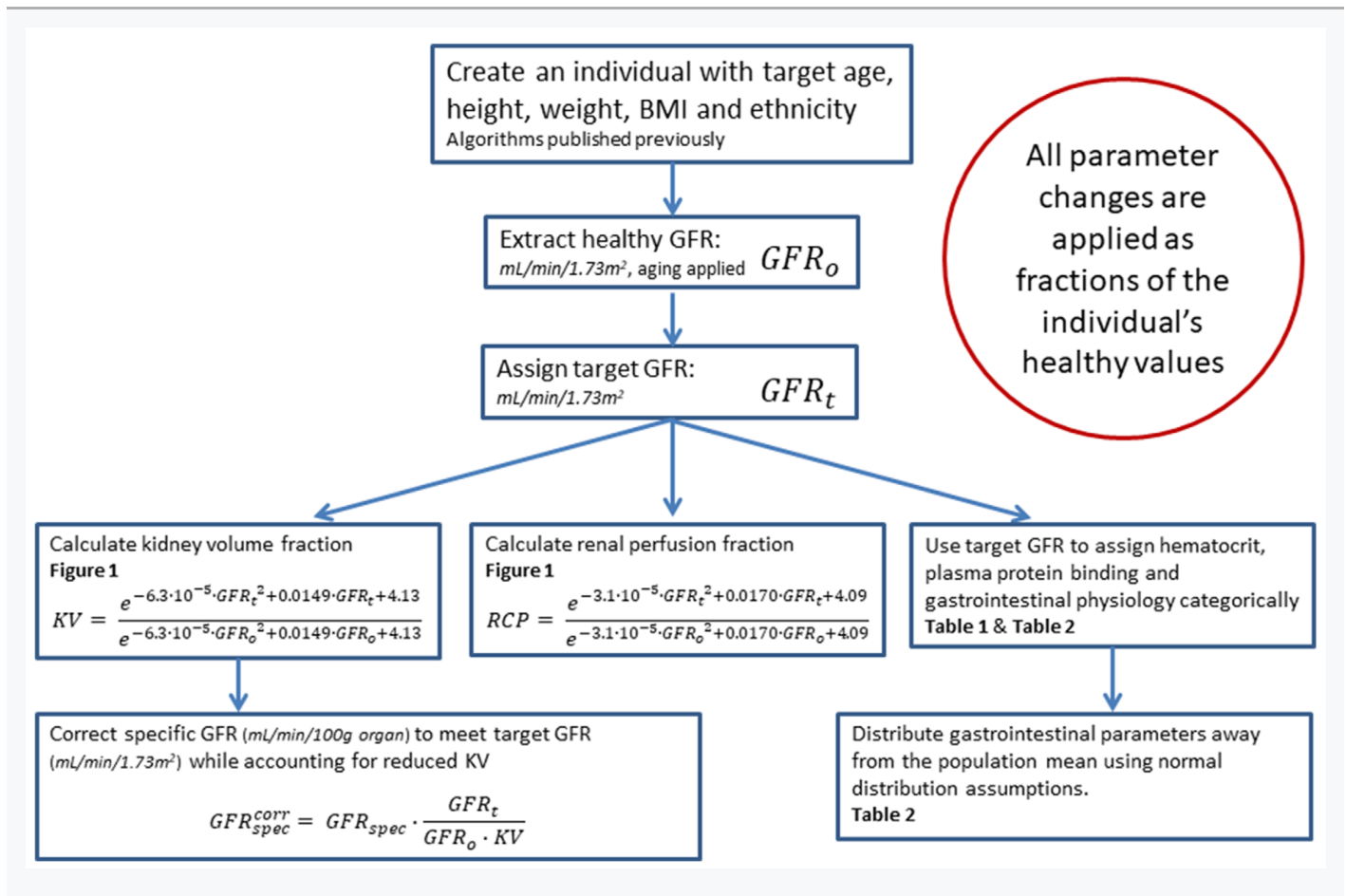


Figure 2-2: Algorithm for generation of virtual individuals with CKD while accounting for the effects of aging

Table 2-1. Hematocrit in Patients With Chronic Kidney Disease^{9, 10, 11}

Creatinine Clearance (mL/min/1.73 m ²)	Men HCT (%)	Women HCT (%)
Healthy	45.5	40
60<CrCl≤70	45.2	39.9
50<CrCl≤60	44.8	39.6
40<CrCl≤50	43.3	39.7
30<CrCl≤40	42.6	38.4
20<CrCl≤30	41.7	37.4
CrCl≤20	34.3	33.5
Hemodialysis	31	29

Table 2-2. Fraction of Healthy Values (Normal Coefficient of Variation %) in Chronic Kidney Disease Patients by Stage

Parameter	Stage 3 (30-60 mL/min/1.73 m ²)	Stage 4 (15-30 mL/min/1.73 m ²)	Stage 5 (<15 mL/min/1.73 m ²)	Dialysis
Fraction unbound in plasma	1.07 ¹²	1.16 ¹²	1.55 ¹²	1.55 ¹²
Gastric emptying time	1.0	1.6 (25%) ¹⁴	1.6 (25%) ^{13,14}	1.6 (30%) ¹⁵⁻¹⁸
Small intestinal transit time	1.0	1.4 (25%) ¹⁴	1.4 (25%) ¹⁴	1.8 (30%) ¹⁹
Colonic transit time	1.0	1.0	1.0	1.8 (50%) ²⁰

3 Compounds

The method to generate virtual populations with CKD was qualified by testing its predictive performance when used in combination with PBPK models for healthy subjects in order to predict the pharmacokinetics of drugs in patients with CKD from literature.

The method was evaluated using three compounds and first considers subjects with CKD (Stage 3 to Stage 5, end-stage renal disease) who are not on dialysis. The three compounds evaluated for the first qualification are gabapentin, acebutolol, and atenolol. This evaluation replicates in part the work published by Malik et al., 2020. The objective is to determine success of the population generation method by comparing the predicted pharmacokinetics in CKD to clinical pharmacokinetic data from literature.

Each subsection that follows will contain an introduction to the test compound, a description of the healthy PBPK model development, and the subsequent translation to a population with CKD.

3.1 Gabapentin

Gabapentin is structurally related to the neurotransmitter, gamma aminobutyric acid (GABA). In adults, it is indicated for the management of post-herpetic neuralgia; and in adults and children 3 years and older with epilepsy, it is used as an adjunctive therapy in treating partial onset seizures with and without secondary generalization.

Gabapentin is available as a capsule, tablet, and oral solution. It is a BCS class 3 drug that is absorbed from the intestines by a process that is mediated by the active transporter, large neutral amino acid transporter 1 (LAT1, SLC7A5). This process is known to be saturable, with the bioavailability of gabapentin to be inversely proportional to the administered dose. The absolute oral bioavailability of a three times per day (tid) dosing regimen of 900, 1200, 2400, 3600, 4800 mg per day was 60, 47, 34, 33 and 27%, respectively.²¹ The absolute bioavailability of the lowest dose of gabapentin tested (100 mg Q8h) was 80%.²²

Gabapentin is not metabolized and completely renally cleared as unchanged drug by glomerular filtration at the maximal rate.²³ While gabapentin is a substrate of the OCT2 influx transporter in kidney, co-administration in humans with the OCT2 inhibitor cimetidine only reduced oral clearance of gabapentin by 14%.²¹ This suggests that renal clearance is likely best described by the passive process of glomerular filtration only. Gabapentin has a volume of distribution of 58 ± 6 L (mean \pm SD) with a half-life of 5-7 hours.²¹ Additionally, it is 97% unbound in plasma.

3.1.1 Gabapentin in a Healthy Population

Table 3-1 presents the drug-specific parameters of gabapentin and the values used for the oral administration model. Parameter optimization was carried out in PK-Sim using a Monte Carlo approach for exploring the parameter space, using the datasets summarized in **Table 3-2**.

Table 3-1. Physicochemical properties and ADME of gabapentin for the final oral model

Physicochemical properties	
Octanol:water coefficient (logP)	-0.08 Log Units
Fraction unbound in plasma (f_u)	0.97 ²⁴
Molecular weight (MW)	171.20 g/mol ²⁵
pKa	3.68, 10.70 ²⁶
Water solubility	100.00 mg/mL ²⁷
ADME	
Partition coefficient	Rodgers and Rowland
Cell permeability	PK-Sim Standard
Total apparent clearance	100 mL/min ²⁸
GFR fraction	1.0
Oral absorption parameters	
Formulation dissolution	Immediately dissolved ^a
LAT1 concentration	log-normally distributed with mean 1.0 μ M and geometric SD 1.40
LAT1 K_m	8630.97 μ M
LAT1 V_{max}	763.59 μ M/min
Specific intestinal permeability	2.09E-7 cm/min

^a T_{max} of solution ²⁹ and tablets ³⁰ equivalent; all oral formulations modeled as dissolved.

Gabapentin was assumed to be immediately dissolved as a solution or IR formulation.

LAT1 was added as an influx transporter and its relative expression throughout the organs of the body was defined by RT-PCR within the PK-Sim database query. Based on cell line work showing no colonic permeability, colonic LAT1 was removed.²²

Table 3-2 presents the gabapentin datasets used for building the oral model. Lipophilicity was optimized to -0.08 Log Units. Cell line K_m values of LAT1 were found in the range of 200-500 μ M³¹ and this value was optimized in order to accurately recreate the nonlinear absorption of gabapentin. V_{max} was also optimized to the oral datasets.

Table 3-2. Pharmacokinetic datasets for gabapentin oral model construction

Study	Dose and administration	Cohort	N	Age (years) ^a	Weight (kg) ^a
Boyd 1999 ³²	400 mg	White American females	18	49.1 ± 16.5	65.7 ± 8.5
Boyd 1999 ³²	400 mg	White American males	18	49.9 ± 19.8	79.4 ± 10.8
Gidal 1998 ³⁰	400 mg tid	European males	1	30 ^b	73 ^b
Gidal 1998 ³⁰	800 mg tid	European males	1	30 ^b	73 ^b
Gidal 1998 ³⁰	1200 mg tid	European males	1	30 ^b	73 ^b
Gidal 1998 ³⁰	1600 mg tid	European males	1	30 ^b	73 ^b
Tjandrawinata 2014 ³³	300 mg	Asian males (65%) and females	37	30 [19 – 54] ^c	60 ^c

^aMean ± SD reported, or range in square brackets if SD not reported.

^bAge and weight of subject estimated from a BMI of approximately 24.

^cApproximated based on the reported range of BMI, 18.03 – 24.99 kg/m².

Figure 3-1 demonstrates the simulated oral model PK profiles in a healthy population compared against observed Blum 1994 study data.³⁴

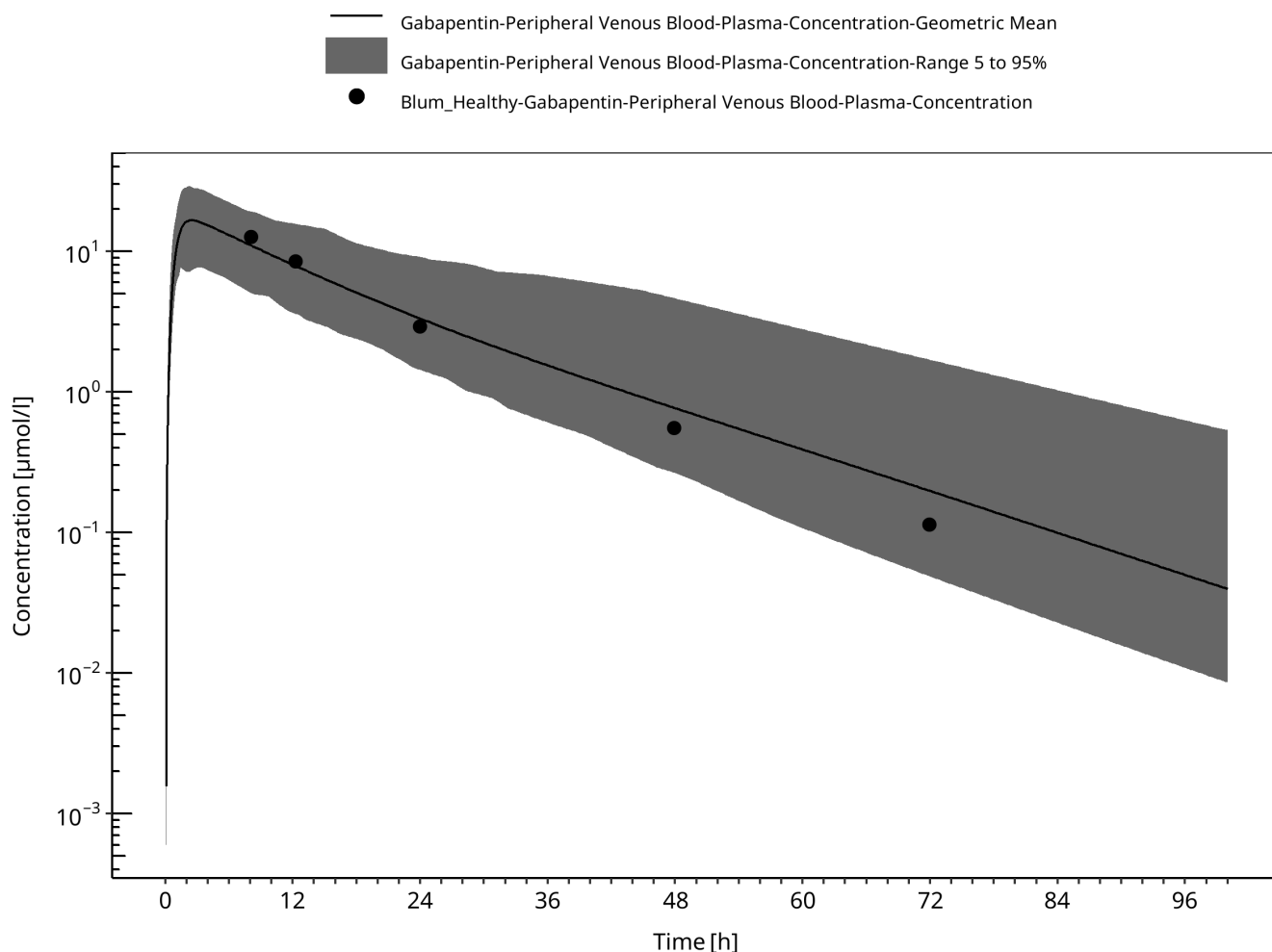


Figure 3-1: Simulation of the pharmacokinetics of Gabapentin after a single dose of 400 mg in healthy subjects. Observed data (circles) presented from Blum.

3.1.2 Gabapentin in a CKD Population

With the drug-specific parameters fixed, the healthy PBPK model was translated to a CKD PBPK model CKD according to the defined method.

A population with Stage 3 CKD was created according to the demographic parameters of the target population presented in the study of Blum 1994³⁴ (eGFR 30-59 mL/min/1.73m²). Simulation of the pharmacokinetics of gabapentin after a single oral dose of 400 mg in this target population is presented in **Figure 3-2**

A population with Stage 4-5 CKD was created according to the demographic parameters of the target population presented in the study of Blum 1994³⁴ (eGFR 1-30 mL/min/1.73m²). Simulation of the pharmacokinetics of gabapentin after a single oral dose of 400 mg in this target population is presented in **Figure 3-3**

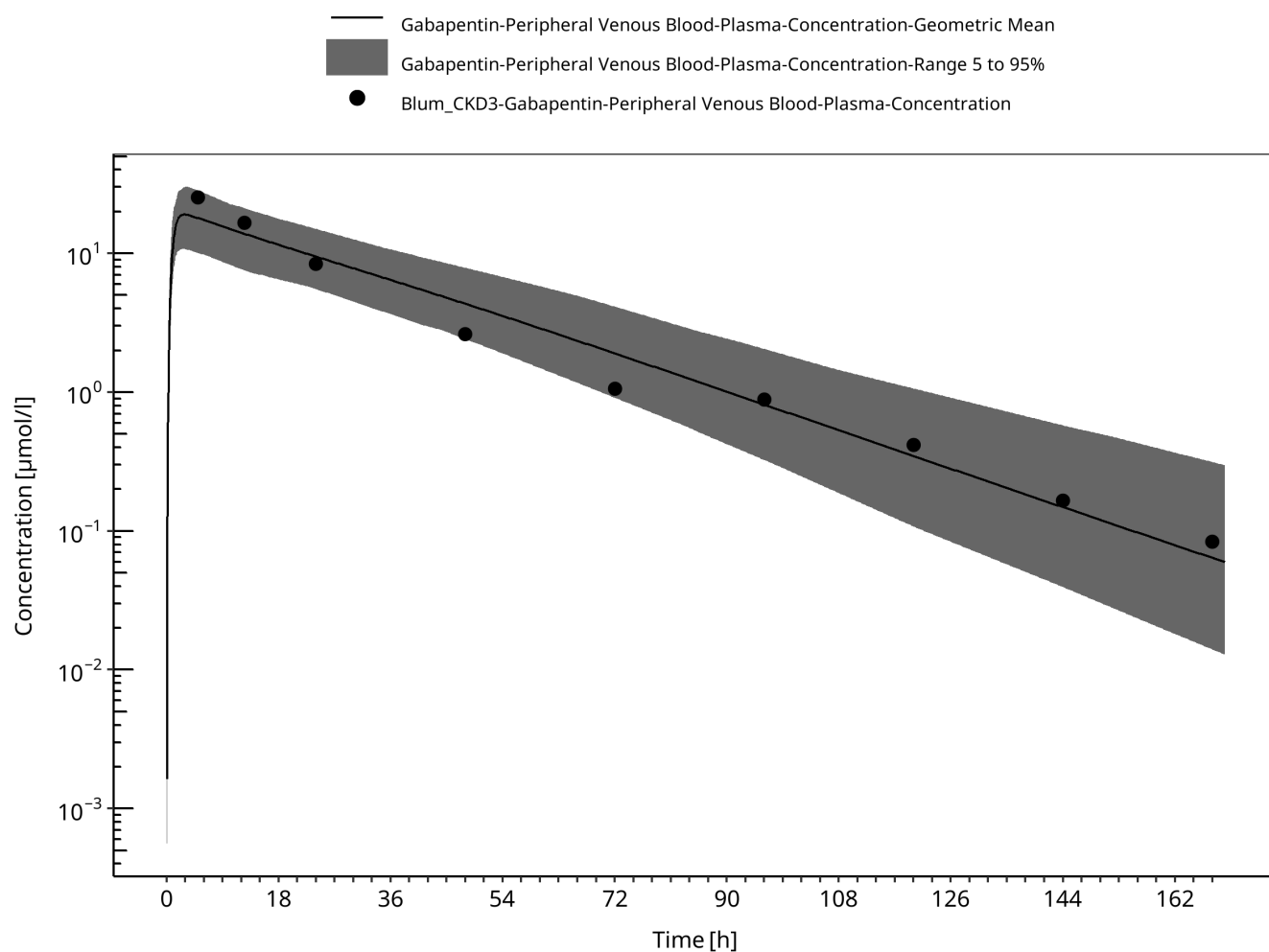


Figure 3-2: Simulation of the pharmacokinetics of Gabapentin after a single dose of 400 mg in subjects with CKD. Observed data (circles) presented from Blum.

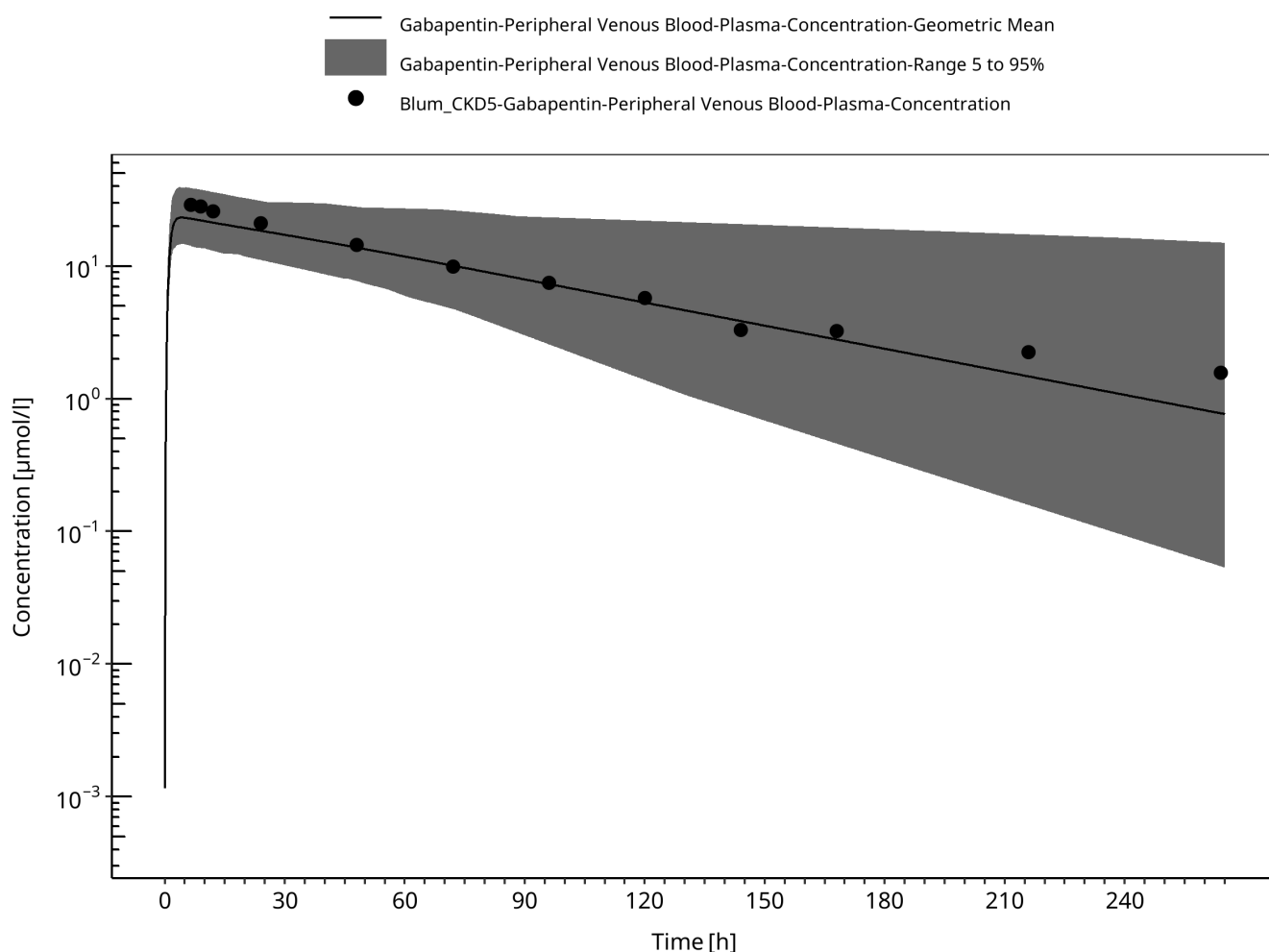


Figure 3-3: Simulation of the pharmacokinetics of Gabapentin after a single dose of 400 mg in subjects with CKD. Observed data (circles) presented from Blum.

3.2 Acebutolol

Acebutolol is a cardioselective beta-blocker with mild intrinsic sympathomimetic activity indicated for hypertension and ventricular arrhythmias. It is administered orally and classified as a BCS Class 3 drug. Acebutolol is well absorbed from the gastrointestinal tract with an absolute bioavailability of approximately 40%. Its elimination half life is 3–4 hours.³⁵

Acebutolol undergoes extensive first pass metabolism in the liver by the two enzymes carboxylesterase 2 (CES2) and N-Acetyltransferase 2 (NAT2) to form the active metabolite diacetolol.³⁶ Other metabolic pathways in the liver that are largely undefined also contribute. Approximately 35% of acebutolol is renally cleared³⁵ and undergoes tubular secretion by the kidneys as demonstrated by a renal clearance of 167 mL/min.³⁷ It has a fraction unbound of 0.85 to plasma proteins.

3.2.1 Acebutolol in a Healthy Population

Table 3-3 presents the drug-specific parameters of acebutolol and the values used for the combined IV-oral model. Parameter optimization was carried out in PK-Sim using a Monte Carlo approach for exploring the parameter space, using the datasets summarized in **Table 3-4**.

Table 3-3. Physicochemical properties and ADME of acebutolol for the final IV-oral model

Physicochemical properties	
Octanol:water coefficient (logP)	1.71 Log Units ³⁸
Fraction unbound in plasma (f_u)	0.85 ³⁹
Molecular weight (MW)	336.43 g/mol ³⁹
pKa	9.40 ⁴⁰
Water solubility	200 mg/mL ⁴⁰
ADME	
Partition coefficient	Rodgers and Rowland
Cell permeability	PK-Sim Standard
Total clearance	615 ± 59 mL/min ⁴¹
CYP concentration	log-normally distributed with mean 1.0 μ M and geometric SD 1.40
CYP specific clearance	0.68 1/min
OCT concentration	normally distributed with mean 1.0 μ M and SD 0.20 μ M ⁴⁶
OCT K_m	100 μ M
OCT In vitro V_{max} /transporter	35.31 μ M/min
GFR fraction	1.0
Oral absorption parameters	
Dissolution half-time	10 minutes
Dissolution profile shape	0.92
Intestine 1 concentration	1.00 μ M
Intestine 1 K_m	5000 μ M
Intestine 1 V_{max}	5000 μ M/min
Intestine 2 concentration	normally distributed with mean 1.0 μ M and SD 0.5 μ M ⁴⁵
Intestine 2 K_m	5000 μ M
Intestine 2 V_{max}	157.34 μ M/min
Specific intestinal permeability	1.48E-6 cm/min (PK-Sim calculated)

Since acebutolol is significantly metabolized by the liver, the expression of the non-specific enzyme processes, referred to as CYP throughout this report, were added with a first order intrinsic clearance process.

Acebutolol undergoes renal transportation via MATE proteins (MATE1, MATE2/2-K) and OCT proteins (OCT2/SLC22A2). The OCT2 proteins draw acebutolol through the basolateral side of the proximal tubule cells and the MATE proteins excrete the drug into the urine from the apical side of the proximal tubule cells. The kinetics of the two transporters are difficult to identify individually as there is no in vitro data. However, it was assumed that acebutolol's efflux is rate limited by MATE proteins and that the Permeability x Surface Area product was sufficiently fast enough to populate acebutolol in the renal epithelium. Therefore, the unknown kinetics of transport proteins were simplified into one average efflux transport protein which was represented on the apical side of the kidney. This simplified process of the OCT transport system is referred to as OCT throughout this report. The process followed active transport Michaelis-Menten kinetics and the K_m of OCT was fixed at 100 μM .

The acebutolol oral formulation was developed assuming high solubility and fast dissolution based on its hydrophilicity (see **Table 3-3**). A Weibull function was used to describe the dissolution profile. The intestinal transporter-mediated uptake of acebutolol is likely driven by the influx of a transporter system, referred to as Intestine 2, that is located on the apical membrane of the caecum. Segment-dependent absorption was modeled by adding a fast efflux transporter to the basolateral side of the caecum, referred to as Intestine 1. Intestine 2 was added to the apical membrane of the caecum and its K_m was fixed at 5000 μM to allow for linear kinetics while V_{max} was optimized.

Table 3-4 presents the acebutolol datasets used for building the combined IV-oral model. The optimized CYP specific clearance, OCT V_{max} , Intestine 2 V_{max} , and specific intestinal permeability values are presented in **Table 3-3**.

The estimated fraction excreted to urine of 12% approximated the observed value of 15% measured at 70 hours after oral administration.^{37,42,43} In contrast, the estimated fraction excreted to urine of 21% slightly underestimated the observed value of 35% measured at 48 hours after IV bolus administration.^{35,41}

Table 3-4. Pharmacokinetic datasets for acebutolol IV-oral model construction

Study	Dose and administration	Cohort	N	Age (years) ^a	Weight (kg) ^a
Gulaid 1981 ⁴²	400 mg PO	European males	8	24 [20 – 28]	66.2 ^b
Roux 1980 ³⁷	200 mg PO	European males	10	26 \pm 4	68 \pm 9
Roux 1983 ⁴³	400 mg PO	European males	12	23.5 \pm 2.1	69.3 \pm 7.5
Roux 1983 ⁴⁴	0.35 mg/kg IV bolus	European males	5	23.4 \pm 1.7	76 \pm 3.9
Roux 1983 ⁴⁴	400 mg PO	European males	5	23.4 \pm 1.7	76 \pm 3.9

^aMean \pm SD reported, or range in square brackets if SD not reported.

^bAverage weight not reported in study. Estimated based on an average BMI of 22 kg/m².

Figure 3-4 shows the simulated oral model PK profiles in a healthy population compared against observed Roux 1980 study data.³⁷

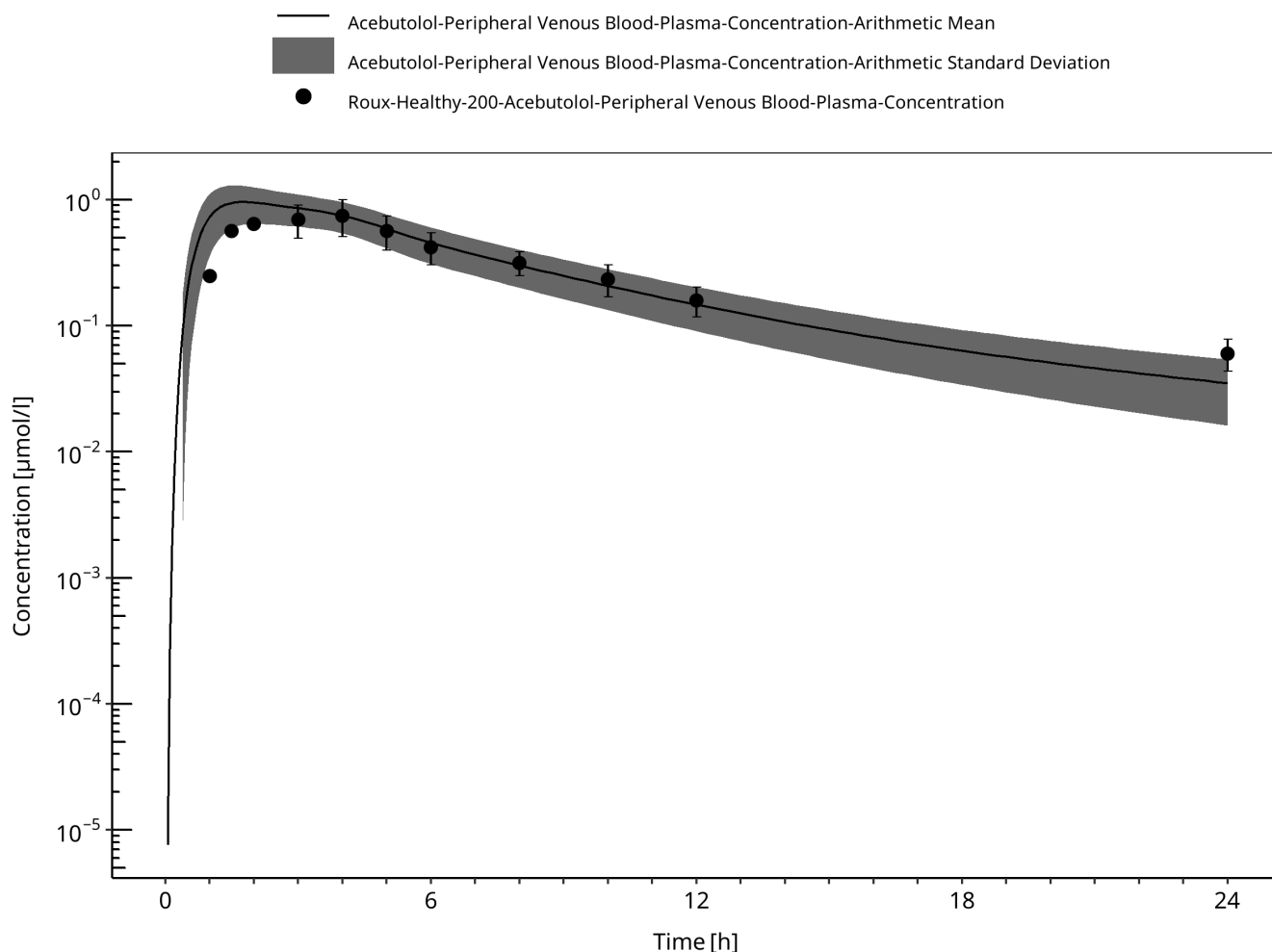


Figure 3-4: Simulation of the pharmacokinetics of Acebutolol after a single dose of 200 mg in healthy subjects. Observed data (circles) presented from Roux.

3.2.2 Acebutolol in a CKD Population

With the drug-specific parameters fixed, the healthy PBPK model was translated to a CKD PBPK model CKD according to the defined method.

A population with Stage 3-5 CKD was created according to the demographic parameters of the target population presented in the study of Roux 1980 (eGFR 6-56 mL/min/1.73m²). Simulation of the pharmacokinetics of acebutolol after a single oral dose of 200 mg in this target population is presented in **Figure 3-5**

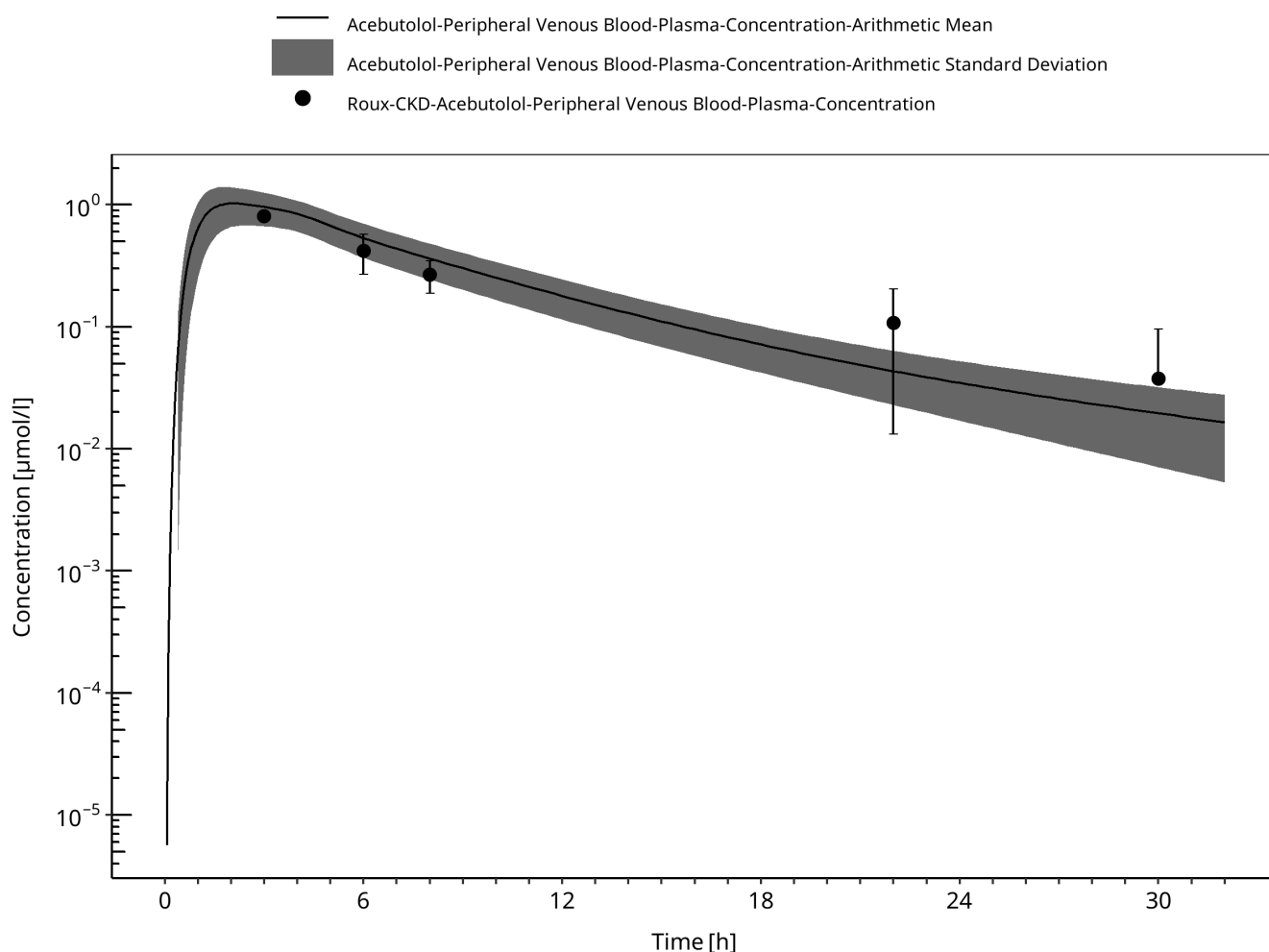


Figure 3-5: Simulation of the pharmacokinetics of Acebutolol after a single dose of 200 mg in subjects with CKD. Observed data (circles) presented from Roux.

3.3 Atenolol

Atenolol is a beta-selective beta-adrenergic receptor blocking agent. It is indicated for hypertension, angina pectoris due to coronary atherosclerosis, and acute myocardial infarction. Atenolol is 6-16% bound to proteins in plasma.⁴⁷ The elimination half-life of atenolol from plasma is 6-7 hours in healthy adults.

Taken orally, atenolol is classified as a BCS Class III drug (low permeability, high solubility) with incomplete absorption in humans. It undergoes little or no metabolism by the liver, and the portion that is absorbed is mainly eliminated by renal excretion with an active tubular secretion component.⁴⁸ Following oral administration, 50-60% of the dose is recovered in urine.⁴⁸ These findings suggested that atenolol is not fully absorbed in the gastrointestinal tract and has a bioavailability of approximately 50%.⁴⁸

Atenolol is a substrate of the organic cation transporter 2 (OCT2/SLC22A2) on the basolateral side of renal tubular epithelial cells and is a substrate for the multidrug and toxic compound extrusion proteins (MATE1, MATE2/2-K) on the apical side.⁴⁹

3.3.1 Atenolol in a Healthy Population

Table 3-5 and **Table 3-7** present the drug-specific parameters of atenolol and the values used for the IV and Oral administration models. Parameter optimization was carried out in PK-Sim using a Monte Carlo approach for exploring the parameter space, using the datasets summarized in **Table 3-6** and **Table 3-8**.

IV model

Table 3-5. Physicochemical properties and ADME of atenolol for the final IV model

Physicochemical properties	
Octanol:water coefficient (logP)	-0.23 Log Units
Fraction unbound in plasma (f_u)	0.89 ⁴⁷
Molecular weight (MW)	266.30 g/mol
pKa	9.60 ⁴⁷
Water solubility	13.30 mg/mL ⁴⁷
ADME	
Partition coefficient	PK-Sim Standard
Cell permeability	PK-Sim Standard
Total clearance	97.3–176.3 mL/min ⁴⁸
OCT concentration	normally distributed with mean 1.0 μ M and SD 0.20 μ M ⁴⁶
OCT K_m	200 μ M ⁵⁰
OCT V_{max}	18.99 μ M/min
GFR fraction	1.0

Atenolol undergoes renal transportation via MATE proteins (MATE1, MATE2/2-K) and OCT proteins (OCT2/SLC22A2).⁴⁹ The OCT2 proteins draw atenolol through the basolateral side of the proximal tubule cells and the MATE proteins excrete the drug into the urine from the apical side of the proximal tubule cells. The kinetics of the two transporters are difficult to identify individually. However, it was assumed that atenolol's efflux is rate limited by MATE proteins and that the Permeability x Surface Area product was sufficiently fast enough to populate atenolol in the renal epithelium. Therefore, the unknown kinetics of transport proteins were simplified into one net efflux transport protein which was represented on the apical side of the kidney. This simplified process of the OCT transport system is referred to as OCT throughout this report. To allow the process to follow linear kinetics, the K_m of OCT was fixed at a high value (200 μ M).⁵⁰

The PK-Sim Standard and Rodgers and Rowland methods to calculate partition coefficients were evaluated with logP and OCT V_{max} for optimization to the IV datasets describing atenolol disposition. The observed PK data were best described by using the PK-Sim Standard method for partition coefficient. The PK-Sim Standard method was also used for the calculation of cell permeability.

Table 3-6 presents the atenolol datasets used for building the IV model. The optimized values are presented in **Table 3-5**. A proportional error model was chosen (i.e., log scaling).

Table 3-6. Pharmacokinetic datasets for atenolol IV model construction

Study	Dose and administration	Cohort	N	Age (years) ^a	Weight (kg) ^a
Brown 1976 ⁵¹	Single 10 mg IV bolus	European males	4	30.5 [25 – 36]	74
Brown 1976 ⁵¹	Single 20 mg IV bolus	European males	4	30.5 [25 – 36]	74
Brown 1976 ⁵¹	Single 50 mg IV bolus	European males	4	30.5 [25 – 36]	74
Brown 1976 ⁵¹	Single 80 mg IV bolus	European males	4	30.5 [25 – 36]	74
Kirch 1980 ⁵²	Single 5 mg IV bolus	European males	4	47.5 [26 – 69]	74.1 ^b
Kirch 1981 ⁵³	Single 5 mg IV bolus	European males (71%) and females	7	28.7 ± 5.9	67.9 ± 13.4
Mason 1979 ⁵⁴	50 mg IV infusion over 12 min	White American males	12	24.5 [21 – 28]	71.6 [63.6 – 79.5]
Wan 1979 ⁵⁵	Single 50 mg IV bolus	White American males	6	24.5 [22 – 27]	72.9 [64.5 – 79.9]

^aMean ± SD reported, or range in square brackets if SD not reported.

^bNot reported in the study. Approximated to be 74.1 kg based on cohort composition and a BMI of 24.

Oral model

The oral PBPK model for the atenolol tablet was developed using literature values for the dissolution profile and solubility (see **Table 3-7**). A Weibull function was used to describe the dissolution profile.

Table 3-7. Oral absorption parameters for the final atenolol oral model

Dissolution half-life	5 minutes ⁵⁶
Dissolution profile shape	0.92
Water solubility	7.90 mg/mL ⁴⁷
Intestine 1 concentration	1.00 µM
Intestine 1 K _m	5000 µM
Intestine 1 V _{max}	5000 µM/min
PMAT concentration	normally distributed with mean 1.0 µM and SD 0.5 µM ⁴⁵
PMAT K _m	5000 µM
PMAT V _{max}	214.7 µM/min
Specific intestinal permeability	1.15E-7 cm/min (PK-Sim calculated)

Atenolol is predominantly absorbed in the ileum.⁵⁷⁻⁶⁰ The transporter-mediated uptake of atenolol is likely driven by the influx Plasma Membrane Monoamine Transporter (PMAT/SLC29A4) located on the apical membrane.^{61,62} Segment-dependent absorption was modeled by adding a fast efflux transporter to the basolateral side of the lower ileum, referred to as Intestine 1. PMAT was added to the apical membrane of the lower ileum and its K_m was fixed at 5000 μM to allow for linear kinetics while V_{max} was optimized. Optimization of PMAT V_{max} was carried out using a Monte Carlo approach to explore the parameter space.

Table 3-8. Pharmacokinetic datasets for atenolol oral model construction

Study	Dose and administration	Cohort	N	Age (years) ^a	Weight (kg) ^a
Brown 1976 ⁵¹	50 mg PO	European males	4	30.5 [25 – 36]	74
Conway 1976 ⁶³	100 mg PO	European males	5	38 [34 – 45]	69 [64 – 73]
Frost 2017 ⁶⁴	100 mg PO	White American males (87%) and females	14	29 \pm 8	80.9 \pm 10.4
Kirch 1981 ⁵³	100 mg PO daily	European males (71%) and females	7	28.7 \pm 5.9	67.9 \pm 13.4
Mason 1979 ⁵⁴	25 mg PO solution	White American males	12	24.5 [22 – 27]	72.9 [64.5 – 79.9]
Mason 1979 ⁵⁴	100 mg PO solution	White American males	12	24.5 [22 – 27]	72.9 [64.5 – 79.9]
Sassard 1977 ⁶⁵	100 mg PO	European Males	12	30 ^b	69 \pm 4 ^c
Wan 1979 ⁵⁵	50 mg PO	White American males	12	28.5 [21 – 36]	68.6 [64.5 – 87.6]
Wan 1979 ⁵⁵	100 mg PO	White American males	12	28.5 [21 – 36]	68.6 [64.5 – 87.6]

^aMean \pm SD reported, or range in square brackets or another noted measure if SD not reported.

^bAge not reported. Estimated to be 30 years old based on cohort composition.

^cMean \pm SE reported.

Figures 3-6, 3-7, and 3-8 demonstrate the simulated oral model PK profiles in a population compared against observed data from Wan 1979, Kirch 1981, and Sassard 1977, respectively.^{53,55,65}

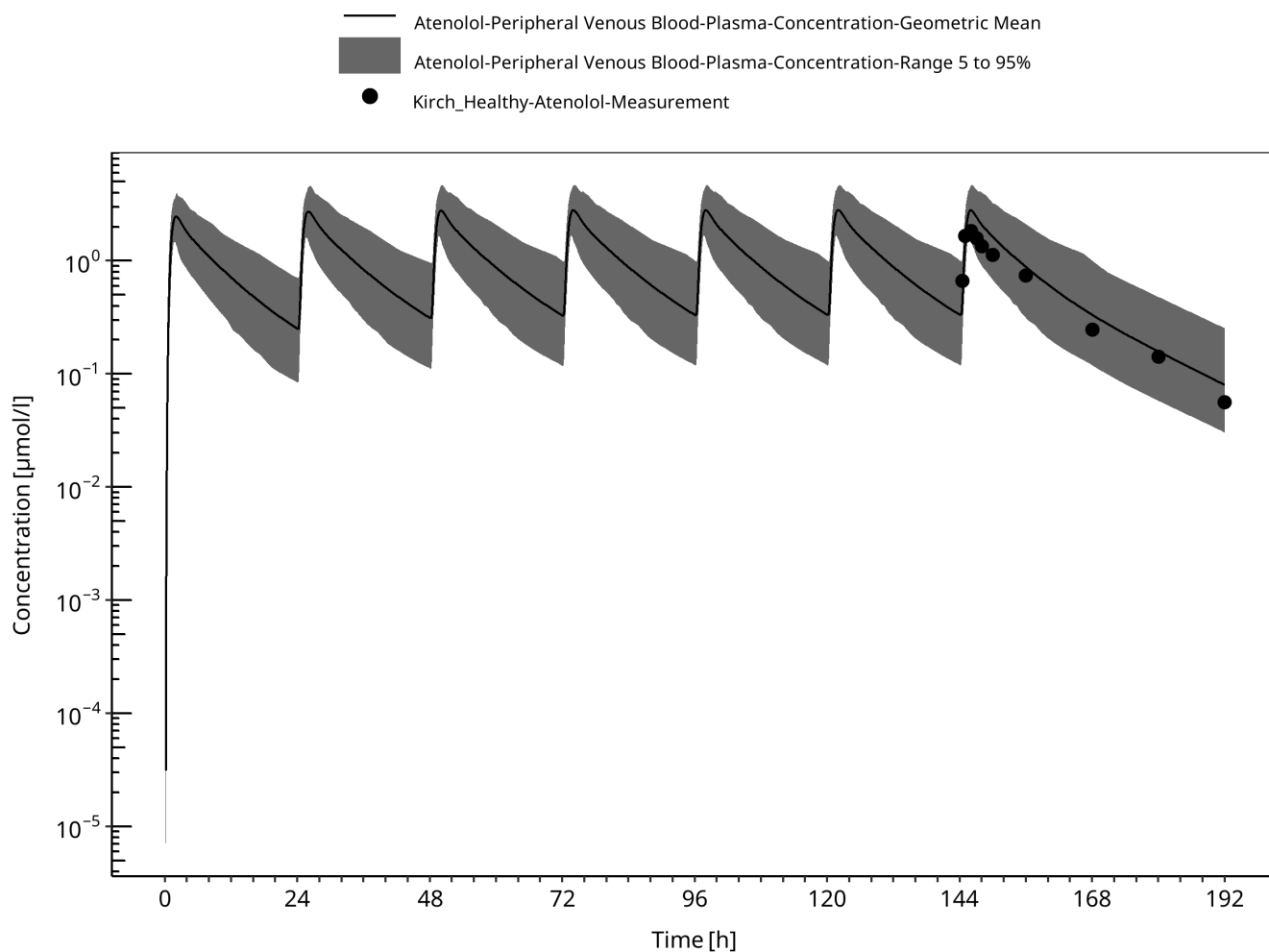


Figure 3-6: Simulation of the pharmacokinetics of Atenolol after multiple doses of 100 mg once daily in healthy subjects. Observed data (circles) presented from Kirch.

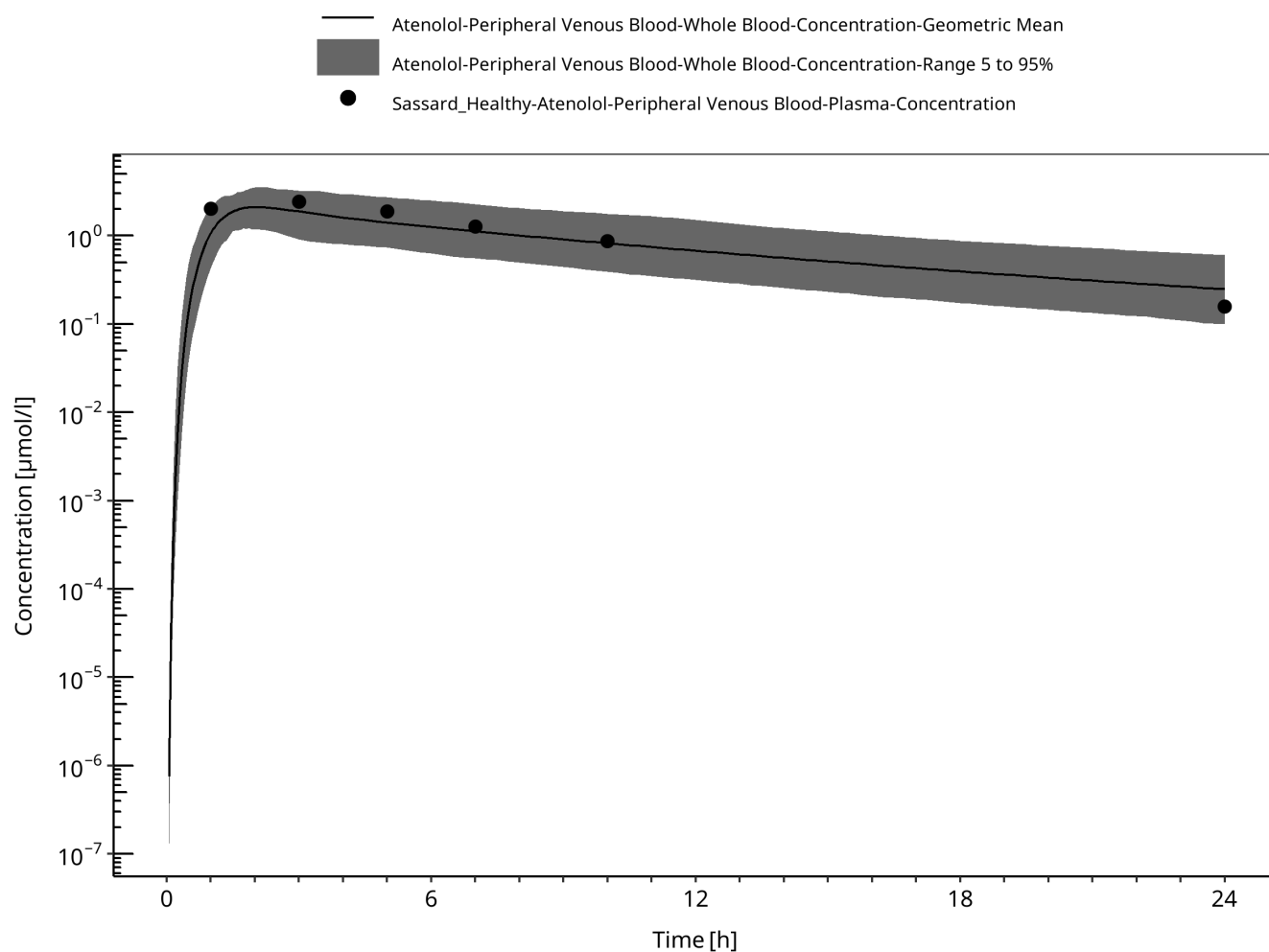


Figure 3-7: Simulation of the pharmacokinetics of Atenolol after a single dose of 100 mg in healthy subjects. Observed data (circles) presented from Sassard.

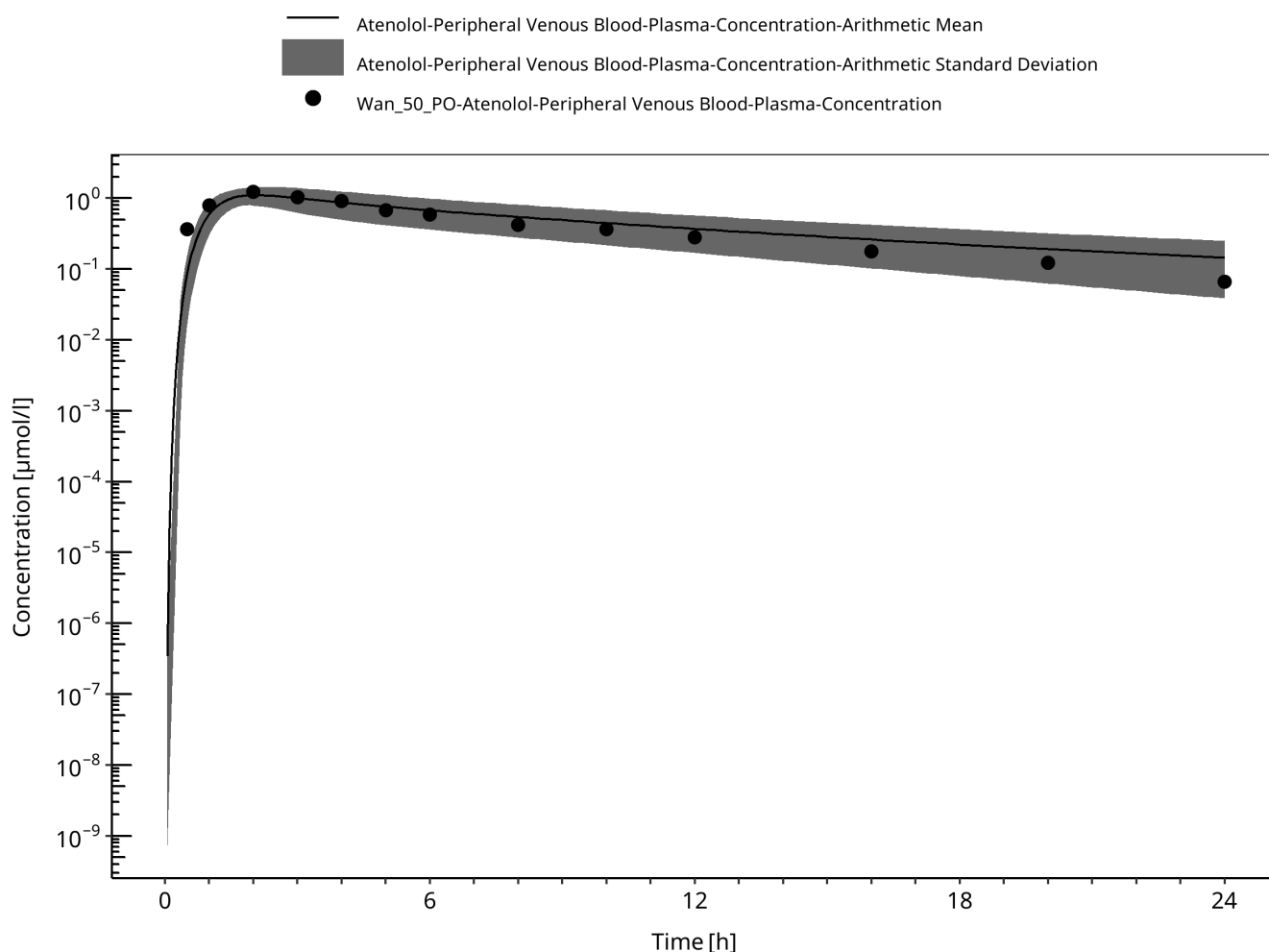


Figure 3-8: Simulation of the pharmacokinetics of Atenolol after a single dose of 50 mg in healthy subjects. Observed data (circles) presented from Wan.

3.3.2 Atenolol in a CKD Population

With the drug-specific parameters fixed, the healthy PBPK model was translated to a CKD PBPK model CKD according to the defined method.

A population with Stage 3 CKD was created according to the demographic parameters of the target population presented in the study of Kirch 1981⁵³ (eGFR 37-60 mL/min/1.73m²). Simulation of the pharmacokinetics of atenolol after multiple oral doses of 100 mg once daily for 7 days in this target population is presented in **Figure 3-9**.

A population with Stage 4 CKD was created according to the demographic parameters of the target population presented in the study of Kirch 1981⁵³ (eGFR 17-30 mL/min/1.73m²). Simulation of the pharmacokinetics of atenolol after multiple oral doses of 100 mg once daily for 7 days in this target population is presented in **Figure 3-10**.

A population with Stage 5 CKD was created according to the demographic parameters of the target population presented in the study of Kirch 1981⁵³ (eGFR 5-9 mL/min/1.73m²). Simulation of the pharmacokinetics of atenolol after multiple oral doses of 100 mg once daily for 7 days in this target population is presented in **Figure 3-11**.

A population with Stage 3 CKD was created according to the demographic parameters of the target population presented in the study of Sassard 1977⁶⁵ (eGFR 46-60 mL/min/1.73m²). Simulation of the whole blood pharmacokinetics of atenolol

after a single oral dose of 100 mg in this target population is presented in Figure 3-12.

A population with Stage 4-5 CKD was created according to the demographic parameters of the target population presented in the study of Sassard 1977⁶⁵ (eGFR 8-24 mL/min/1.73m²). Simulation of the whole blood pharmacokinetics of atenolol after a single oral dose of 100 mg in this target population is presented in Figure 3-13.

A population with Stage 3-4 CKD was created according to the demographic parameters of the target population presented in the study of Wan 1979⁵⁵ (eGFR 15-42 mL/min/1.73m²). Simulation of the pharmacokinetics of atenolol after a single oral dose of 50 mg in this target population is presented in Figure 3-14.

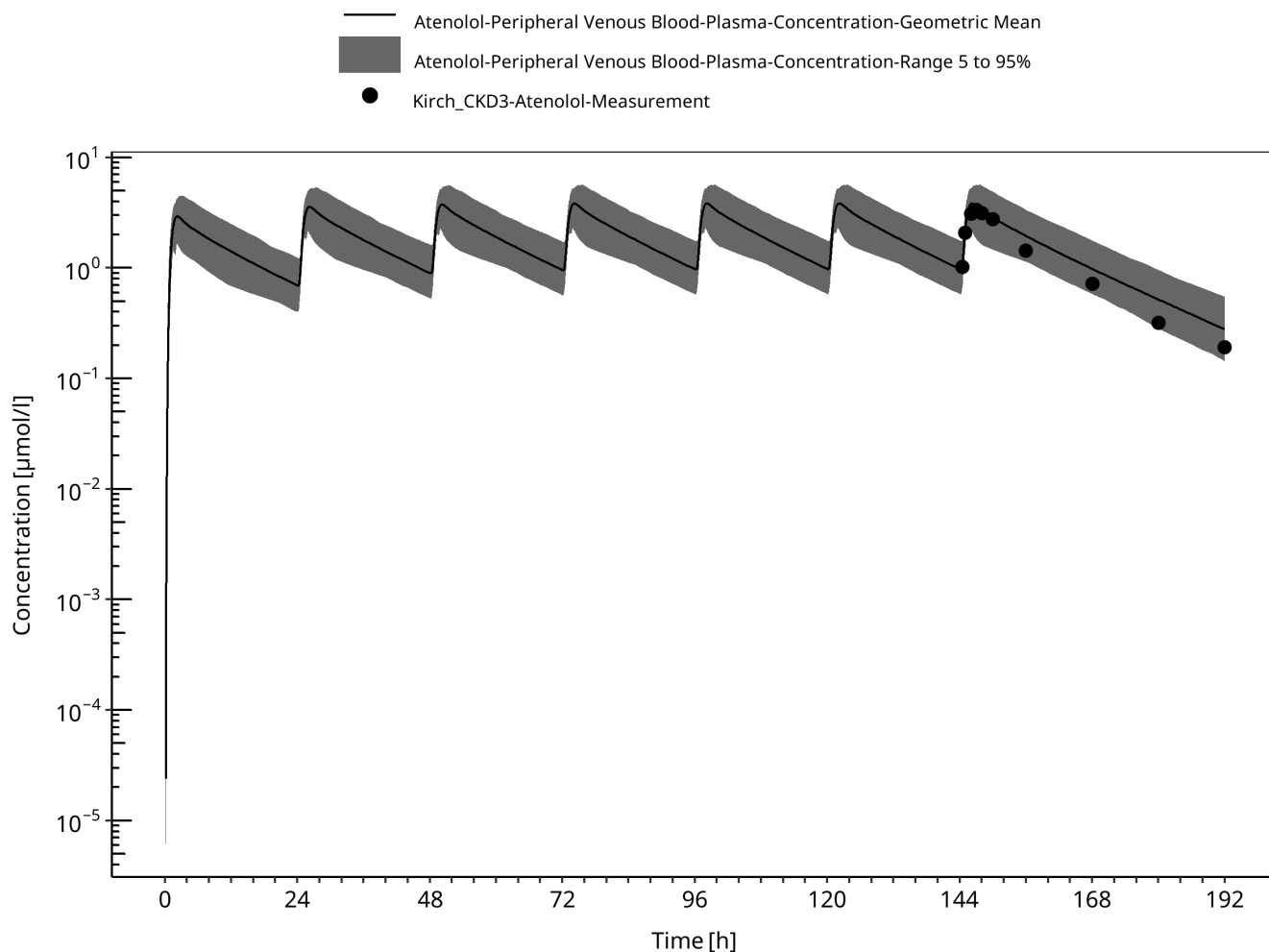


Figure 3-9: Simulation of the pharmacokinetics of Atenolol after multiple doses of 100 mg once daily in subjects with CKD. Observed data (circles) presented from Kirch.

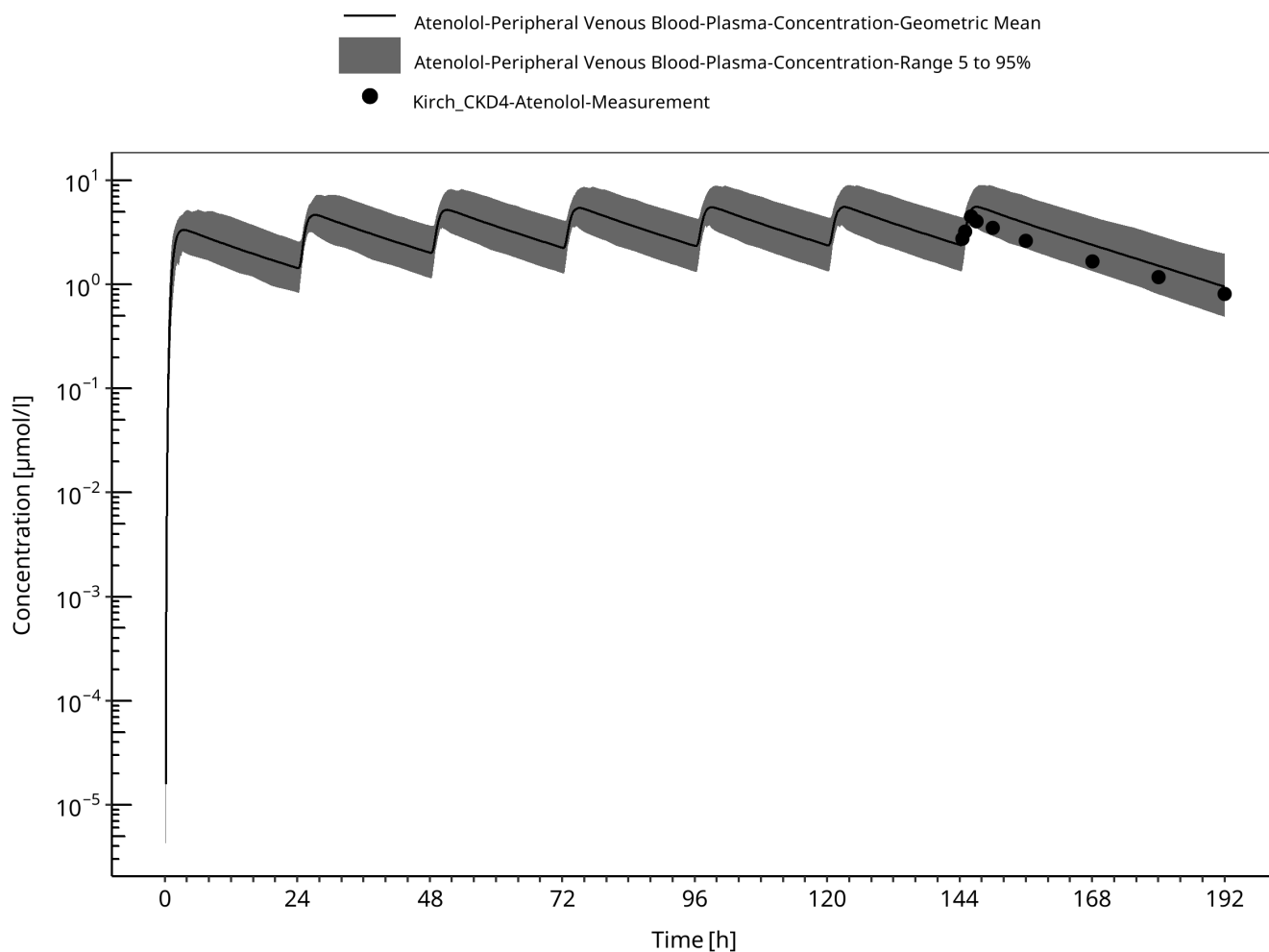


Figure 3-10: Simulation of the pharmacokinetics of Atenolol after multiple doses of 100 mg once daily in subjects with CKD. Observed data (circles) presented from Kirch.

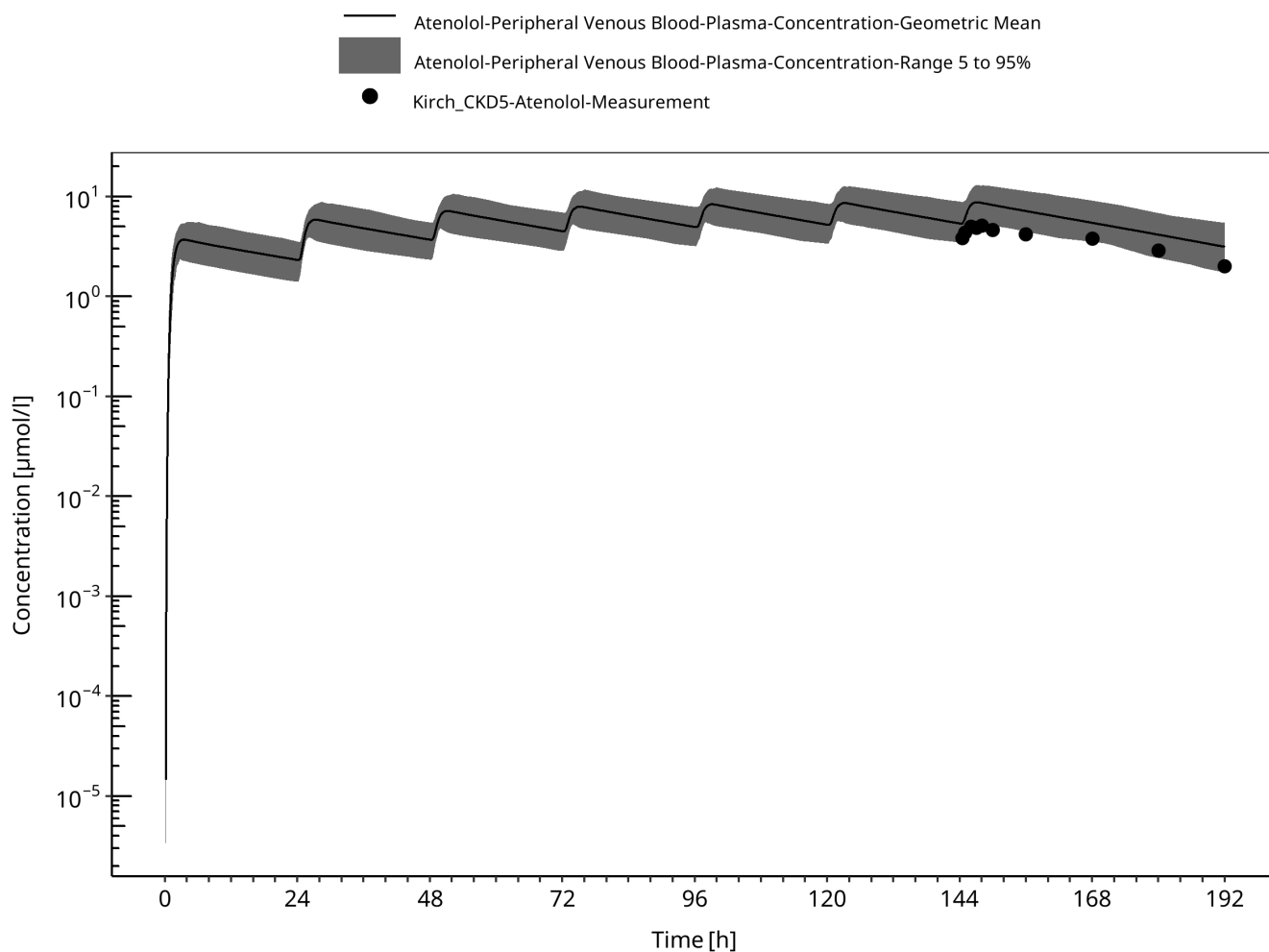


Figure 3-11: Simulation of the pharmacokinetics of Atenolol after multiple doses of 100 mg once daily in subjects with CKD. Observed data (circles) presented from Kirch.

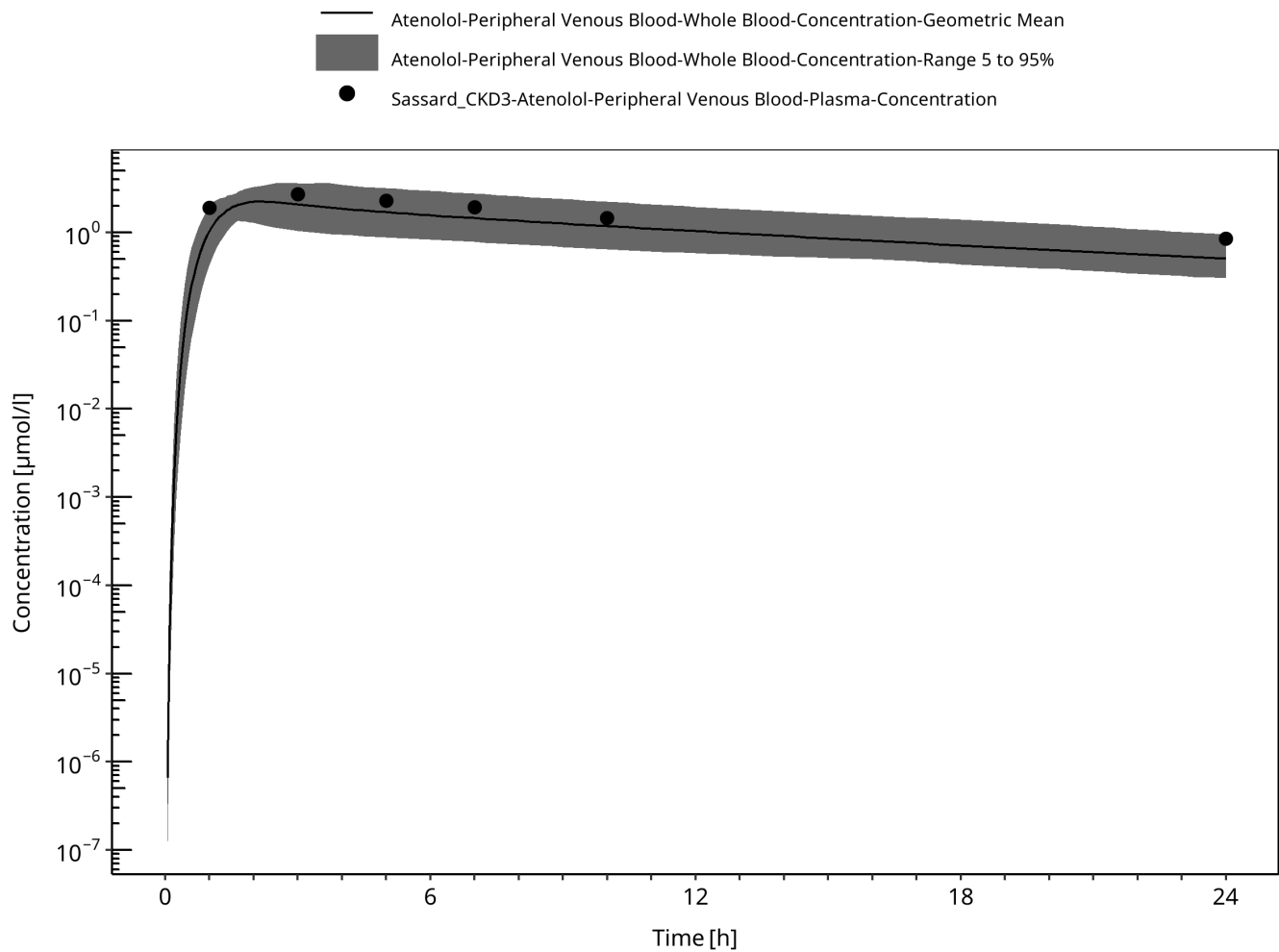


Figure 3-12: Simulation of the pharmacokinetics of Atenolol after a single dose of 100 mg in subjects with CKD. Observed data (circles) presented from Sassard.

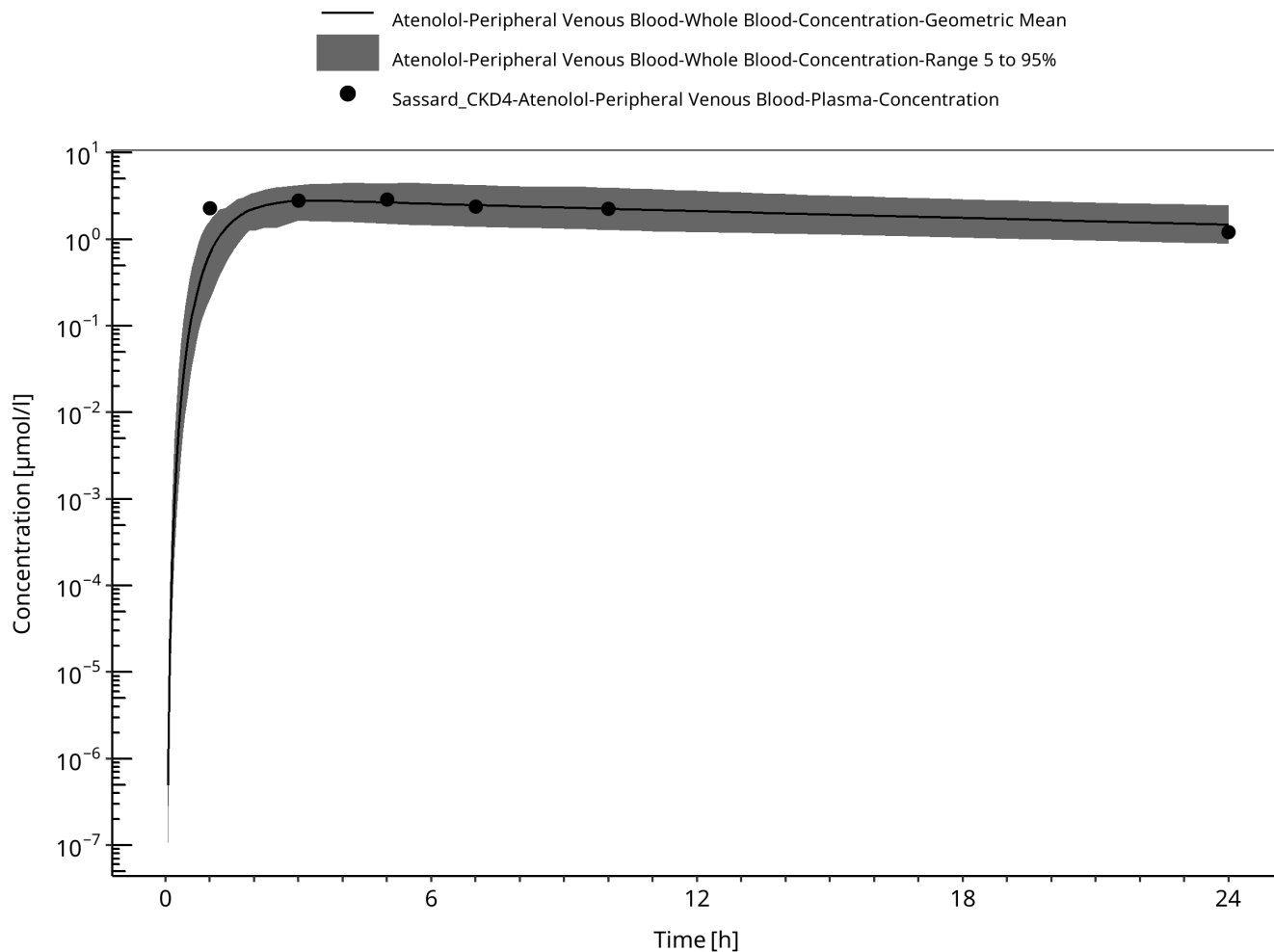


Figure 3-13: Simulation of the pharmacokinetics of Atenolol after a single dose of 100 mg in subjects with CKD. Observed data (circles) presented from Sassard.

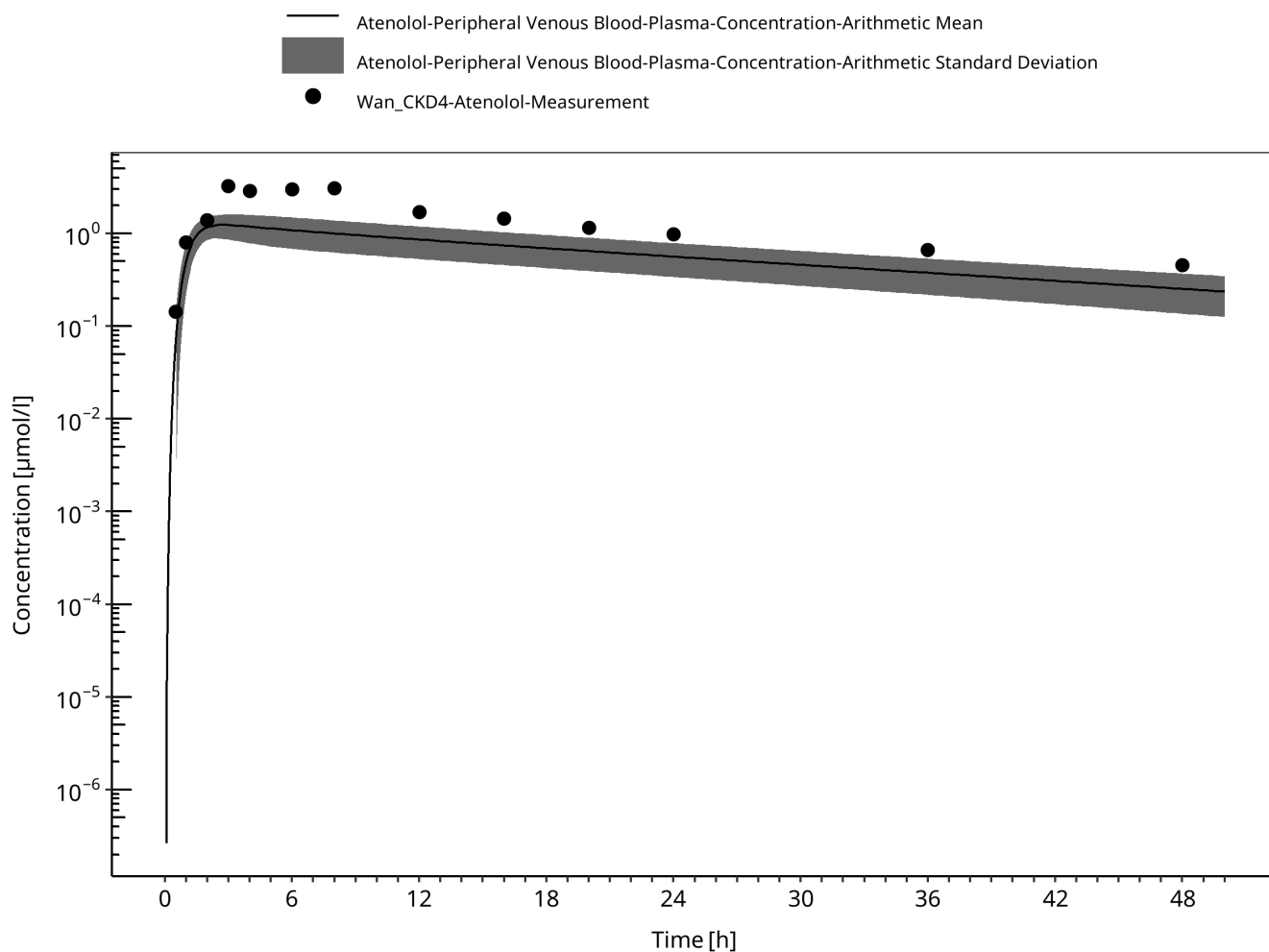


Figure 3-14: Simulation of the pharmacokinetics of Atenolol after a single dose of 50 mg in subjects with CKD. Observed data (circles) presented from Wan.

4 Conclusion

Success of the virtual population with the test models for predicting drug pharmacokinetics in subjects with CKD qualifies the method to support the use of the population for PBPK-CKD modeling.

The evaluation was performed for compounds with significant renal elimination, for compounds subject to active transport through the OCT system, and compounds with non-renal elimination pathways that are largely conserved in subjects with CKD. The population parameters are intended to account for the underlying anatomy and physiology changes that occur in subjects with varying stages of CKD. Specific parameterizations from the user may be necessary for drug-specific cases where non-renal elimination pathways are affected by the disease (e.g., a metabolic enzyme) or where a transport pathway is significantly compromised (e.g., the OAT system).

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