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

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1 Introduction

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

Erythromycin a macrolide antibiotic used for the treatment of a number of bacterial infections, including respiratory tract infections, skin infections, chlamydia infections, and others. Erythromycin is available in the form of various salts and formulations, for example as:

- erythromycin lactobionate for injection
- erythromycin base in enteric-coated capsules or tablets for oral administration
- erythromycin stearate in filmcoated tablets for oral administration
- erythromycin ethylsuccinate in suspension or in filmcoated tablets for oral administration

In its free form as base, erythromycin is easily hydrolyzed in acidic aqueous solution (Mordi 2000). Therefore, orally administered erythromcyin is given in the form of enteric-coated formulations or as more acid-stable salts or esters of erythromycin (e.g. erythromycin ethylsuccinate). Once in the small intestine, erythromycin is rapidly absorbed displaying a highly variable bioavailability (Chun 1977, Mather 1981). Erythromycin diffuses in most tissues and accumulates in leukocytes and phagocytes (Miller 1984, Carlier 1987). About 70% of erythromycin is bound to plasma proteins (Barre 1987). Erythromycin has been shown to be a substrate for various transporters including P-gp and OATP1B1. The latter has been shown to critically affect erythromycin disposition (Lancaster 2012). Erythromycin is extensively metabolized through N-demethylation catalyzed by CYP3A. Metabolism via CYP4F11 has also been suggested (Kalsotra 2004). Biliary excretion also appears to play an important role in erythromycin clearance (Acocella 1968, Chelvan 1979), but tits contribution to total elimination remains unknown. The dose fraction excreted unchanged in urine is minimal and highly variable; reported fractions range from 0.018 ± 0.005 to 0.171 ± 0.11 (mean ± SD) (Pasic 1987, Austin 1980). Erythromycin is a moderate inhibitor of CYP3A4 and listed by the FDA as clinical index inhibitor for CYP3A.

The presented erythromycin PBPK model as well as the respective evaluation plan and evaluation report are provided open-source (https://github.com/Open-Systems-Pharmacology/Erythromycin-Model).

2 Methods

2.1 Modeling Strategy

The general workflow for building an adult PBPK model has been described by Kuepfer et al. (Kuepfer 2016). Relevant information on the anthropometry (height, weight) was gathered from the respective clinical study, if reported. Information on physiological parameters (e.g. blood flows, organ volumes, hematocrit) in adults was gathered from the literature and has been incorporated in PK-Sim® as described previously (Willmann 2007). The applied activity and variability of plasma proteins and active processes that are integrated into PK-Sim® are described in the publicly available 'PK-Sim® Ontogeny Database Version 7.3' (PK-Sim Ontogeny Database Version 7.3).

The PBPK model was developed based on clinical data of healthy adult subjects obtained from the literature, covering different formulation types and erythromycn salts. Multiple doses and dosing schedules following intravenous (IV) and oral (PO) administration were included in model building. Mass balance information on urinary excretion of unchanged erythromycin after IV administration was also accounted for during the model building process.

Unknown parameters were simultaneously optimized using all available PK data, in particular:

- 4 data sets following single IV administration of 4 different doses of erythromycin (125 mg, 250 mg, 300 mg, 500 mg) as erythromycin lactobionate
- 6 data sets following single and multiple PO administration of 3 different doses of erythromycin (250 mg, 500 mg, 1000 mg) as film-coated tablets containing erythromycin stearate
- 2 data sets following single PO administration of 500 mg erythromycin as enteric-coated tablets containing erythromycin as base
- 2 data sets following single and multiple PO administration of 2 different doses of erythromycin (250 mg, 500 mg) as enteric-coated capsules containing pellets of erythromycin as base

Structural model selection was mainly guided by visual inspection of the resulting description of data and biological plausibility. The following parameters were identified using the Parameter Identification module provided in PK-Sim® and MoBi® (<u>Open Systems Pharmacology</u> <u>Documentation</u>):

- Dissolution shape (for the film-coated tablet containing erythromycin stearate)
- Dissolution time (50% dissolved) (for the film-coated tablet containing erythromycin stearate)
- Dissolution lag time (for the enteric-coated tablet containing erythromycin as base)
- Dissolution shape (for the enteric-coated tablet containing erythromycin as base)
- Dissolution time (50% dissolved) (for the enteric-coated tablet containing erythromycin as base)
- Solubility at reference pH (for the enteric-coated tablet containing erythromycin as base)
- Dissolution lag time (for the enteric-coated capsule containing pellets of erythromycin as base)
- Dissolution shape (for the enteric-coated capsule containing pellets of erythromycin as base)
- Dissolution time (50% dissolved) (for the enteric-coated capsule containing pellets of erythromycin as base)

- Specific intestinal permeability (transcellular)
- Transport Protein OATP1B1 kact
- Metabolizing Enzyme CYP3A4 kact
- Total Hepatic Clearance Specific clearance
- GFR fraction

Details about input data (physicochemical, *in vitro* and clinical) can be found in <u>Section 2.2</u>.

Details about the structural model and its parameters can be found in <u>Section 2.3</u>.

2.2 Data

2.2.1 In vitro / physicochemical data

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

Parameter	Unit	Literature	Description
Molecular weight	g/mol	733.9 (drugbank.ca)	Molecular weight
pK _a (basic)		8.8 (<u>Lien 1974</u>); 8.88 (<u>McFarland 1997</u>)	Acid dissociation constant
logP		2.48 (<u>Lien 1974</u>); 2.92 (<u>Capobianco</u> 1994); 3.06 (<u>McFarland 1997</u>)	Partition coefficient between octanol and water
f _u		0.27 ± 0.05 ^a (<u>Sun 2010</u>); 0.28 ± 0.04 ^a (<u>Iliopoulou 1982</u>); 0.305 ± 0.028 ^a (<u>Barre 1987</u>); 0.326 (<u>Xu 2009</u>)	Fraction unbound in human plasma
Water solubility (erythromycin lactobionate)	mg/L	200000 (<u>U.S. Patent 2,761,859</u>)	Solubility of erythromycin lactobionate in water
Water solubility (erythromycin stearate)	mg/L	182 (<u>Jones 1969</u>)	Solubility of erythromycin stearate in water
Water solubility (erythromycin base)	mg/L	2100 (<u>Manna 1998</u>)	Solubility of free erythromycin as base in water

^a mean ± SD

2.2.2 In vitro data on mechanism-based inhibition of CYP3A

A literature search was carried out to collect quantitative information the meachnism-based inhibition of CYP3A by erythromycin, in particular measured values for K_l and k_{inact} . The gathered information is summarized in the table below and is used for model building.

Victim compound	In vitro system	Enzyme	Κ _Ι [μΜ]	k _{inact} [min ⁻¹]	Concentration range of erythromycin [µM]	Concentration of victim compound [µM]	Reference
testosterone	rec cDNA CYP3A4	CYP3A4	1.04	0.0293	0 - 20	100	Akiyoshi 2013
testosterone	rec cDNA CYP3A4	CYP3A4	1.21	0.0164	0 - 20	100	Akiyoshi 2013
testosterone	rec cDNA CYP3A4	CYP3A4	0.415	0.0159	0 - 20	100	Akiyoshi 2013
testosterone	rec cDNA CYP3A4	CYP3A4	2.24	0.0293	0 - 20	100	Akiyoshi 2013
testosterone	rec cDNA CYP3A4	CYP3A4	0.753	0.0248	0 - 20	100	Akiyoshi 2013
midazolam	rec cDNA CYP3A4	СҮРЗА	9.5	0.16	50-fold range	10	Atkinson 2005
midazolam	rec cDNA CYP3A4	СҮРЗА	8.82	0.12	50-fold range	10	Atkinson 2005
midazolam	rec cDNA CYP3A4	СҮРЗА	8.3	0.09	50-fold range	10	Atkinson 2005
testosterone	rec cDNA CYP3A4	СҮРЗА	9.5	0.06	4 - 200	16.7	Atkinson 2005
triazolam	HLM	СҮРЗА	5.4	0.069		400	Aueviriyavit 2010
testosterone	rec cDNA CYP3A4	CYP3A4	5.3	0.12	5 - 20	200	<u>Chan 2000</u>
midazolam	cryopreserved HEP	СҮРЗА	60	0.081	3 - 300	30	<u>Chen 2011</u>
midazolam	cryopreserved HEP	СҮРЗА	67.9	0.079	3 - 300	30	<u>Chen 2011</u>
midazolam	rec cDNA CYP3A4	CYP3A4	0.762	0.0648	0.1 - 30	4	Ishikawa 2017
testosterone	rec cDNA CYP3A4	CYP3A4	1.00	0.0604	0.1 - 30	150	Ishikawa 2017
nifedipine	rec cDNA CYP3A4	CYP3A4	0.794	0.0766	0.1 - 30	6	Ishikawa 2017
triazolam	HLM	CYP3A4	15.9	0.062	3 - 100	300	Kanamitsu 2000
triazolam	HLM	CYP3A4	17.4	0.055	3 - 100	300	Kanamitsu 2000
triazolam	rec cDNA CYP3A4	CYP3A4	19.1	0.173	3 - 100	300	Kanamitsu 2000
triazolam	rec cDNA CYP3A4	CYP3A4	18.9	0.097	3 - 100	300	Kanamitsu 2000
testosterone	HLM	CYP3A4	29.4	0.0271		250	<u>Kosaka</u> 2017
testosterone	HLM	CYP3A4	30	0.040	0.3 - 300		Kozakai 2013
midazolam	HLM	CYP3A4	12	0.035	0.3 - 300		Kozakai 2013
midazolam	HLM	CYP3A4	20	0.033	0.3 - 300		Kozakai 2013
midazolam	cryopreserved HEP	СҮРЗА	25.15	0.08	0.13 - 100	30	<u>Mao 2011</u>

Victim compound	In vitro system	Enzyme	Κ _Ι [μΜ]	k _{inact} [min ⁻¹]	Concentration range of erythromycin [µM]	Concentration of victim compound [µM]	Reference
midazolam	HLM	СҮРЗА	10.8	0.032			<u>Mao 2016</u>
midazolam	cryopreserved HEP	СҮРЗА	30.7	0.05	0 - 300	20	<u>Mao 2016</u>
midazolam	cryopreserved HEP	СҮРЗА	59.2	0.062	0 - 100	5	<u>Mao 2016</u>
midazolam	cryopreserved HEP	СҮРЗА	80.3	0.052	0 - 100	20	<u>Mao 2016</u>
midazolam	rec cDNA CYP3A4	CYP3A4	7.47	0.042	2 - 50	8	<u>McConn</u> 2004
midazolam	HLM	CYP3A4	10.9	0.046	2 - 100	8	<u>McConn</u> 2004
midazolam	primary HEP	СҮРЗА	11	0.07	0.1 - 10	3	McGinnity 2006
midazolam	HLM	CYP3A4	10	0.036	0 - 25		<u>Obach</u> 2007
testosterone	HLM	CYP3A4	9.8	0.039	0 - 25	500	<u>Obach</u> 2007
testosterone	rec cDNA CYP3A4	CYP3A4	0.92	0.058	5 - 100	250	Polasek 2006
testosterone	HLM	CYP3A4	12.8	0.037	5 - 100	250	Polasek 2006
midazolam	rec cDNA CYP3A4	CYP3A4	5.1	0.30	0.5 - 50	100	Ring 2005
testosterone	rec cDNA CYP3A4	CYP3A4	0.92	0.058			<u>Teng 2010</u>
testosterone	HLM	CYP3A4	4.579	0.0115			<u>Teng 2010</u>
domperidone	HLM	CYP3A4	18.4	0.022	2.5 - 200	500	<u>Ung 2009</u>
domperidone	rec cDNA CYP3A4	CYP3A4	4.1	0.026	2.5 - 200	500	<u>Ung 2009</u>
midazolam	HLM	CYP3A4	12.1	0.0215	0 - 100	25	Watanabe 2007
nifedipine	HLM	CYP3A4	11.3	0.0295	0 - 100	50	Watanabe 2007
testosterone	HLM	CYP3A4	10.9	0.0352	0 - 100	200	Watanabe 2007
midazolam	HLM	СҮРЗА	1.48	0.017	0.5 - 500	20	Xu 2009
midazolam	cryopreserved HEP	СҮРЗА	20.0	0.016	0.5 - 500	20	<u>Xu 2009</u>
midazolam	cryopreserved HEP	СҮРЗА	109	0.055	0.5 - 500	20	<u>Xu 2009</u>
midazolam	HLM	СҮРЗА	81.8	0.0665	20 - 400	10	<u>Yamano</u> <u>2001</u>
midazolam	HLM	СҮРЗА	15.7	0.1			<u>Zhang</u> <u>2006</u>
testosterone	rec cDNA CYP3A4	CYP3A4	5	0.34	1 - 50	200	<u>Zhang</u> <u>2009</u>
testosterone	HLM	CYP3A4	15.7	0.09	1 - 50	200	<u>Zhang</u> <u>2009</u>

Victim compound	In vitro system	Enzyme	Κ _Ι [μΜ]	k _{inact} [min ⁻¹]	Concentration range of erythromycin [µM]	Concentration of victim compound [µM]	Reference
midazolam	HLM	CYP3A4	26.5	0.041	2.5 - 50	10	Zimmerlin 2011

Note: Abbreviations: HEP: human hepatocytes; HLM: human liver microsomes; rec cDNA CYP3A4: human recombinant c-DNA CYP3A4 enzymes (e.g. supersomes, baculovirus-insect cell system, E. coli transfected cells)

2.2.3 Clinical data

A literature search was carried out to collect available PK data on erythromycin in healthy adults.

The following publications were found and used for model building and evaluation:

Publication	Study description
Austin 1980	IV administration of 125 mg, 250 mg, 500 mg, and 900 mg as erythromycin lactobionate; single dose
<u>Barre 1987</u>	IV administration of 500 mg as erythromycin lactobionate; single dose
<u>Huppertz</u> 2011	IV administration of 1000 mg as erythromycin lactobionate; single dose
<u>Pasic 1987</u>	IV administration of 300 mg as erythromycin lactobionate; single dose
<u>Sun 2010</u>	IV administration of 125 mg and PO administration of 250 mg (salt and formulation type not specified); single dose
<u>Iliopoulou</u> 1982	PO administration of 500 mg as film-coated tablet containing erythromycin stearate; multiple dose
Josefsson 1982	PO administration of 500 mg as film-coated tablet containing erythromycin stearate and 250 mg, 500 mg, 1000 mg as enteric coated capsules containing pellets of erythromycin base; single dose
<u>Kavi 1988</u>	PO administration of 500 mg as film-coated tablet containing erythromycin stearate; single dose
Malmborg 1978	PO administration of 500 mg as film-coated tablet containing erythromycin stearate; multiple dose
Miglioli 1990	PO administration of 1000 mg erythromycin stearate; multiple dose
<u>Posti 1983</u>	PO administration of 500 mg as film-coated tablets containing erythromycin stearate, as enteric-coated tablets containing erythromycin base, and as enteric-coated tablets containing erythromycin stearate; single dose
Schreiner 1984	PO administration of 500 mg as film-coated tablets containing erythromycin stearate and as enteric-coated capsule containing pellets of erythromycin base; single dose
Yakatan 1980	PO administration of 250 mg as film-coated tablet containing erythromycin stearate, as enteric-coated tablet containing erythromycin base, and as enteric-coated capsules containing erythromycin estolate; single and multiple dose
Birkett 1990	PO administration of 250 mg as enteric-coated capsules containing pellets of erythromycin base; single dose
Kivistö 1997	PO administration of 500 mg as enteric-coated capsules containing erythromycin base; multiple dose
<u>DiSanto</u> 1981	PO administration of 500 mg as unprotected tablets containing erythromycin base, as film-coated tablets containing erythromycin base, as enteric-coated tablets containing erythromycin base and as film-coated tablets containing erythromycin stearate; single and multiple dose
Kroboth 1982	PO administration of 500 mg as enteric-coated tablets containing erythromycin base; single dose

2.3 Model Parameters and Assumptions

2.3.1 Dissolution and absorption

Development of an adequate absorption model for erythromycin was complicated by the large intersubject variability in the absorption kinetics of erythromycin (<u>Chun 1977</u>, <u>Mather 1981</u>). Additionally, multiple formulation types are available entailing different absorption kinetics (<u>Chun 1977</u>, <u>Yakatan 1980</u>, <u>Mather 1981</u>). The herein presented model was developed for the following oral dosage forms:

- film-coated tablets containing erythromycin stearate
- enteric-coated tablets containing erythromycin as free base
- enteric-coated capsules containing pellets of erythromycin as free base

It was assumed that the solubility of erythromycin as stearate in the film-coated tablets and as free base pellets in enteric-coated capsules is high and was therefore fixed to a value of 500 mg/L. Solubility of erythromycin base in enteric-coated tablets was optimized to capture the relatively lower C_{max} value of this formulation observed in the herein used *in vivo* PK studies. For all three formulations types, dissolution kinetics was described by a Weibull function with different parameter values for each formulation type. Optimized parameters were the Dissolution shape and Dissolution time (50% dissolved). Additionally, for the enteric-coated formulations, a lag time was applied in the model to account for the passage through the stomach, while the film-coated tablet was assumed to dissolve immediately upon entering the stomach, i.e. no lag time was applied here. Specific intestinal permeability (transcellular) was the same for all formulations and simultaneously optimized with the other parameters.

2.3.2 Distribution

With an average fraction unbound in human plasma of approximately 0.30, erythromycin is moderately protein-bound. In the developed model, the fraction unbound (plasma, reference value) was set to 0.305 which is the value reported by Barre 1987. Lipophilicity was fixed to a value of 2.82 which is the average of three experimentally measured values reported in the literature (Lien 1974, Capobianco 1994, McFarland 1997). The observed PK data were found to be best described using the model for estimating intracellular-to-plasma partition coefficients according to the method by Rodgers and Rowland (Rodgers 2005, Rodgers 2006). Cellular permeabilities were automatically calculated using the method Charge dependent Schmitt (Open Systems Pharmacology Documentation). Active transfer of erythromycin by OATP1B1 was modeled as Michaelis-Menten kinetics using an Influx transporter type; the Km value of 13.2 µM was taken from the literature (Lancaster 2012) and kcat was optimized. The gene expression profile of OATP1B1 (default symbol for the gene: LST-3TM12) was loaded from the internal PK-Sim® database using the expression data quantified by RT-PCR (Open Systems Pharmacology Documentation).

2.3.3 Elimination

Erythromycin is extensively metabolized via N-demethylation catalyzed by CYP3A. Kinetics of this biotransformation was described by a Michaelis-Menten process with kcat being optimized and km fixed to a value of 70 μ M, the average of reported values in the literature (Riley 1997, Wang 1997). The gene expression profile of CYP3A4 was loaded from the internal PK-Sim® database using the expression data quantified by RT-PCR (Open Systems Pharmacology Documentation).

Although it has also been observed that erythromycin is metabolized via CYP4F11 (Kalsotra 2004), this elimination pathway was not accounted for in the model because its contribution to overall elimination was assumed to be low. In humans, CYP4F11 is mainly expressed in the liver and to a much lesser extent in the kidney (Cui 2000) and the CYP4F family makes up approximately 15% of all hepatic CYP enzymes (Michaels 2014). The K_m and V_{max} values for the CYP4F11-mediated biotransformation reported by Kalsotra 2004 are similar to those measured for CYP3A4 (Riley_1997, Wang 1997), suggesting that the relative mass balance of these two metabolism pathways mainly depends on the absolute amount of each enzyme in the liver. While no information on total CYP4F11 in the human liver could be found in the literature, CYP4F11 expression in the liver of cynomolgus monkeys was observed to be approximately 6-fold lower than that of CYP3A4 (Uehara 2015). Hence, it was assumed that CYP4F11-mediated metabolism of erythromycin can be neglected in humans.

Additional elimination pathways suggested for erythromycin are acid-catalyzed degradation (hydrolysis) the acidic milieu of the stomach (Mordi 2000) and biliary excretion (Acocella 1968, Chelvan 1979), but no quantitative information on the mass balance of these pathways could be found in the literature. Additionally, mechanism-based inhibition of CYP3A4 by erythromycin might constitute another clearance process which was neither considered in the model. However, a total hepatic clearance process was implemented in the model which could at least partly account for other elimination pathways not explicitly accounted for in the model.

The reported dose fractions of erythromycin undergoind unchanged renal excretion after IV administration range from 0.018 ± 0.005 to 0.171 ± 0.11 (mean \pm SD) (Pasic 1987, Austin 1980). This information was accounted for in the model by implementing a glomerular filtration process and optimizing the GFR fraction to match the observed dose fractions excreted unchanged in urine.

2.3.2 Autoinhibition

There is abundant evidence from *in vitro* studies that erythromycin irreversibly inhibits CYP3A (e.g. <u>Larrey 1983</u>) and the FDA lists erythromycin as moderate index inhibitor for CYP3A. Ample data quantifying the mechanism-based inhibition of CYP3A4 by erythromycin was collected from the literature (see section <u>2.2.2</u>). The data listed in the Table in section <u>2.2.2</u> can be statistically summarized as follows:

Parameter [unit]	Min	Q1	Geometric mean	Median	Arithmetic mean	Q3	Max
Κ _Ι [μΜ]	0.420	4.89	8.71	10.9	18.4	19.3	109
k _{inact} [min ⁻	0.0115	0.0314	0.0504	0.0535	0.0664	0.0772	0.340

In the PBPK model, [Ki] and kinact were fixed to the geometric mean values tabulated above.

3 Results and Discussion

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

The next sections show:

- 1. the final model parameters for the building blocks: <u>Section 3.1</u>.
- 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: <u>Section 3.3</u>.

3.1 Final input parameters

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

Compound: Erythromycin

Name	Value	Value Origin	Alternative	Default
Solubility at reference pH	500 mg/l	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Erythromycin stearate film- coated tablet	True
Reference pH	7	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Erythromycin stearate film- coated tablet	True
Solubility at reference pH	500 mg/l	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Erythromycin base enteric coated pellets	False
Reference pH	7	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Erythromycin base enteric coated pellets	False
Solubility at reference pH	200 mg/ml	Publication-In Vitro-Hoffhine, Jr Charles E. "Aqueous soluble salts of erythromycin." U.S. Patent 2,761,859, issued September 4, 1956.	Erythromycin lactobionate	False

Name	Value	Value Origin	Alternative	Default
Reference pH	7	Publication-In Vitro-Hoffhine, Jr Charles E. "Aqueous soluble salts of erythromycin." U.S. Patent 2,761,859, issued September 4, 1956.	Erythromycin lactobionate	False
Solubility at reference pH	2.5002520571 mg/l	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Erythromycin base enteric coated tablet	False
Reference pH	7		Erythromycin base enteric coated tablet	False
Lipophilicity	2.82 Log Units	Publication-In Vitro-Average of reported values in the literature (PMID: 9135031, PMID: 18611609, and: Lien, E. J., Kuwahara, J., & Koda, R. T. (1974). Diffusion of drugs into prostatic fluid and milk. Drug Intelligence & Clinical Pharmacy, 8(8), 470-475. http:// journals.sagep ub.com/doi/ab s/10.1177/1060 028074008008 03)	Literature (average value)	True

Name	Value	Value Origin	Alternative	Default
Fraction unbound (plasma, reference value)	0.305	Publication-In Vivo-PMID: 3606934	Barre 1987	True
Specific intestinal permeability (transcellular)	0.00057882652836 cm/min	Parameter Identification- Parameter Identification- Value updated from '001' on 2019-10-01 13:46	Fitted	True
Is small molecule	Yes			
Molecular weight	733.927 g/mol	Internet- drugbank.ca		
Plasma protein binding partner	Unknown			

Calculation methods

Name	Value
Partition coefficients	Rodgers and Rowland
Cellular permeabilities	Charge dependent Schmitt

Processes

Metabolizing Enzyme: CYP3A4-Biotransformation_fitted

Molecule: CYP3A4

Parameters

Name	Value	Value Origin	
In vitro Vmax for liver microsomes	345 pmol/min/mg mic. protein	Publication-In Vitro-PMID: 9566442	
Km	70 μM	Publication-In Vitro-Average of reported values in the literature (PMID: 9107550 and PMID: 9566442)	
kcat	6.477728756 1/min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019- 10-01 13:46	

Systemic Process: Glomerular Filtration-fitted

Species: Human

Parameters

Name	Value	Value Origin
GFR fraction	1.3151947529	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:46

Inhibition: CYP3A4-MBI

Molecule: CYP3A4

Parameters

Name	Value	Value Origin
kinact	0.05044312 1/min	Parameter Identification-Parameter Identification- Value updated from '001' on 2019-10-01 13:46
K_kinact_half	8.705192 µmol/l	Parameter Identification-Parameter Identification- Value updated from '001' on 2019-10-01 13:46

Transport Protein: OATP1B1-fitted

Molecule: OATP1B1

Name	Value	Value Origin
Transporter concentration	1 μmol/l	
Vmax	11.66 pmol/ml/min	Publication-In Vitro-PMID: 22990751
Km	13.2 μmol/l	Publication-In Vitro-PMID: 22990751
kcat	18.3753550107 1/min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019- 10-01 13:46

Systemic Process: Total Hepatic Clearance-fitted

Species: Human

Name	Value	Value Origin	
Fraction unbound (experiment)	0.305		
Lipophilicity (experiment)	2.48 Log Units		
Plasma clearance	0 ml/min/kg		
Specific clearance	3.1542129297 1/min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019- 10-01 13:46	

Formulation: Erythromycin_Weibull_enteric-coated-pellets

Type: Weibull

Name	Value	Value Origin
Dissolution time (50% dissolved)	3.4504159922 min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Lag time	107.7187470594 min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Dissolution shape	1.1678641251	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Use as suspension	Yes	

Formulation: Erythromycin_Weibull_enteric-coated-tablet

Type: Weibull

Name	Value	Value Origin
Dissolution time (50% dissolved)	121.1893502595 min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Lag time	66.1803696499 min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Dissolution shape	1.5167999313	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Use as suspension	Yes	

Formulation: Erythromycin_Weibull_filmtablet

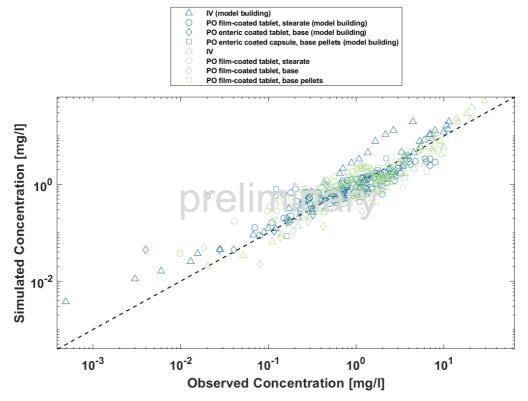
Type: Weibull

Name	Value	Value Origin
Dissolution time (50% dissolved)	27.6449832276 min	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Lag time	0 min	Other-Assumption
Dissolution shape	0.3779018493	Parameter Identification-Parameter Identification-Value updated from '001' on 2019-10-01 13:29
Use as suspension	Yes	

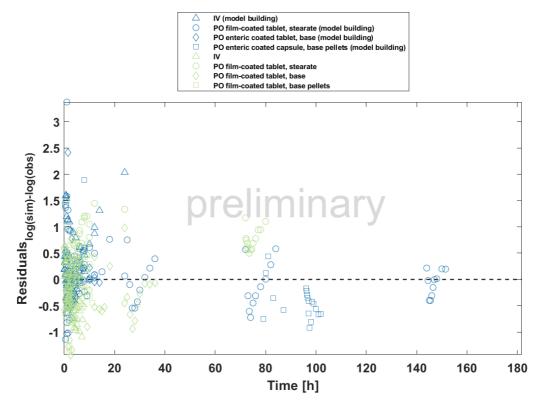
3.2 Diagnostics Plots

Below you find the goodness-of-fit visual diagnostic plots for the PBPK model performance of all data used presented in <u>Section 2.2.2</u>.

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



Goodness of fit plot for concentration in plasma



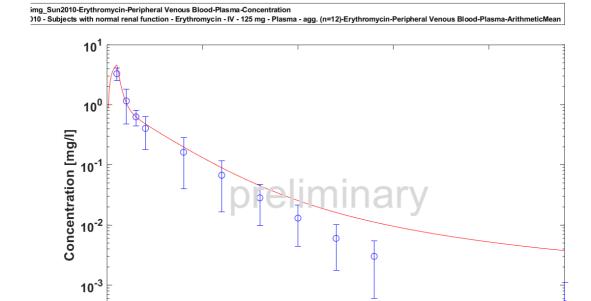
Goodness of fit plot for concentration in plasma

GMFE = 1.579291

3.3: Concentration-Time Profiles

Simulated versus observed concentration-time profiles of all data listed in <u>Section 2.2.2</u> are presented below.

3.3.1 Model Building



10

Time [h]

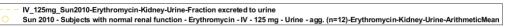
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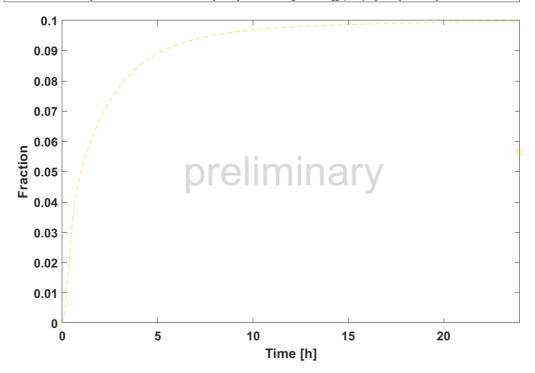
20

Time Profile Analysis

10⁻⁴

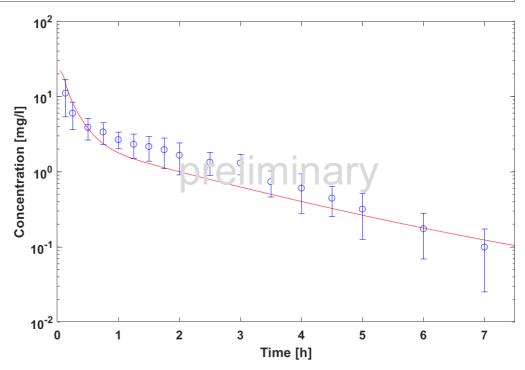
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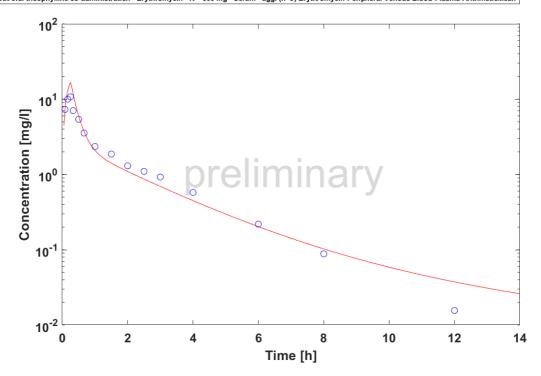




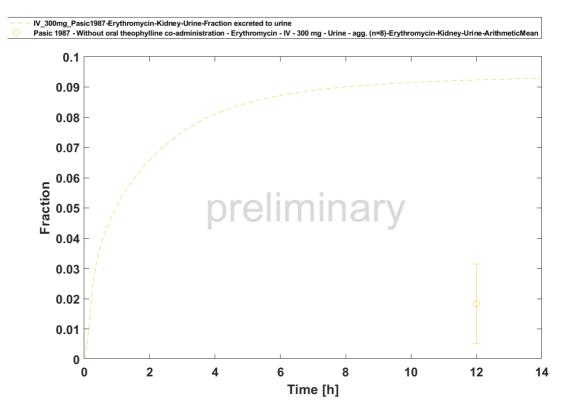
Time Profile Analysis 1

250mg_3min_Austin1980-Erythromycin-Peripheral Venous Blood-Plasma-Concentration stin 1980 - 250 mg over 3 min (Regimen 1) - Erythromycin - IV - 250 mg - Serum - agg. (n=24)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

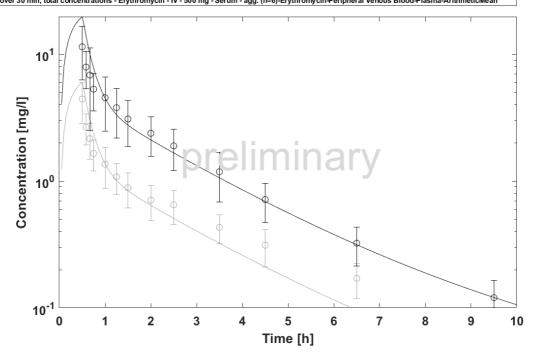




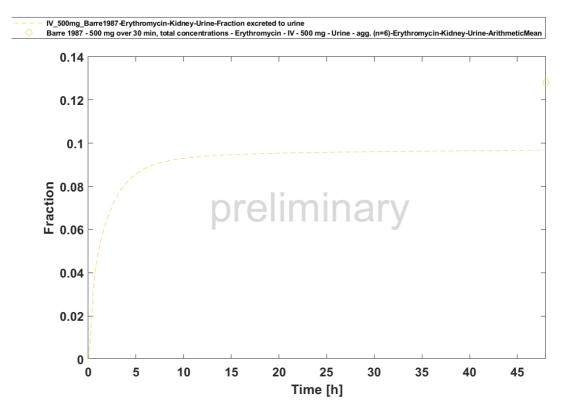
Time Profile Analysis



e1987-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
e1987-Erythromycin-Peripheral Venous Blood-Plasma Unbound-Concentration
) mg over 30 min, unbound concentrations - Erythromycin - IV - 500 mg - Serum - agg. (n=6)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMea
) mg over 30 min, total concentrations - Erythromycin - IV - 500 mg - Serum - agg. (n=6)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

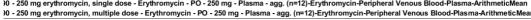


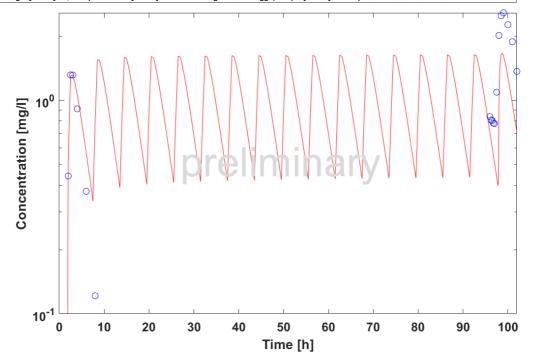
Time Profile Analysis



rkett1990-Erythromycin-Peripheral Venous Blood-Plasma-Concentration

30 - 250 mg erythromycin, single dose - Erythromycin - PO - 250 mg - Plasma - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

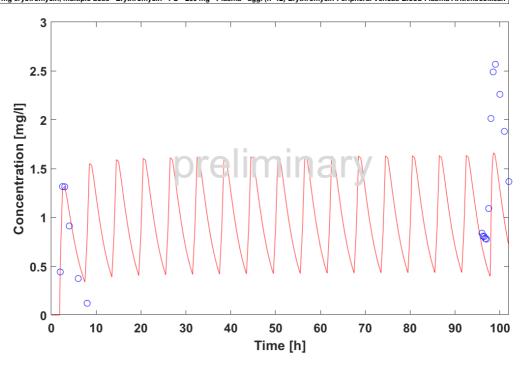


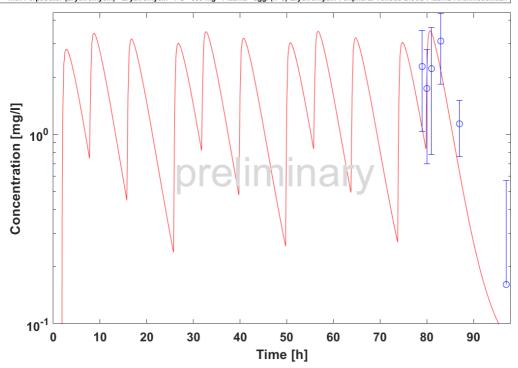


Time Profile Analysis

rkett1990-Erythromycin-Peripheral Venous Blood-Plasma-Concentration

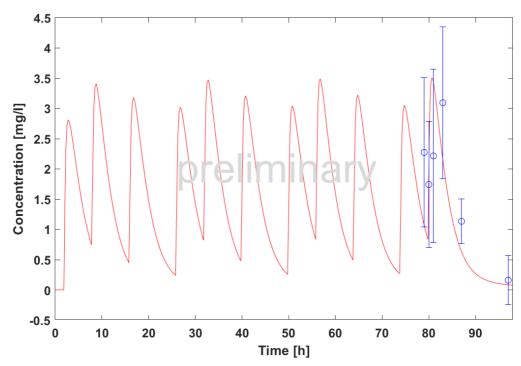
30 - 250 mg erythromycin, single dose - Erythromycin - PO - 250 mg - Plasma - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean 30 - 250 mg erythromycin, multiple dose - Erythromycin - PO - 250 mg - Plasma - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



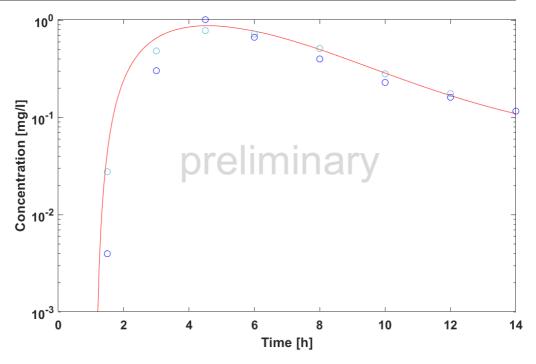


Time Profile Analysis

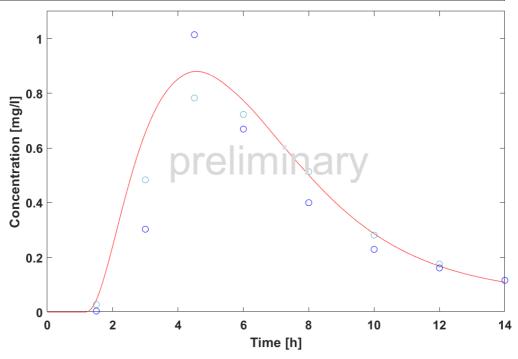


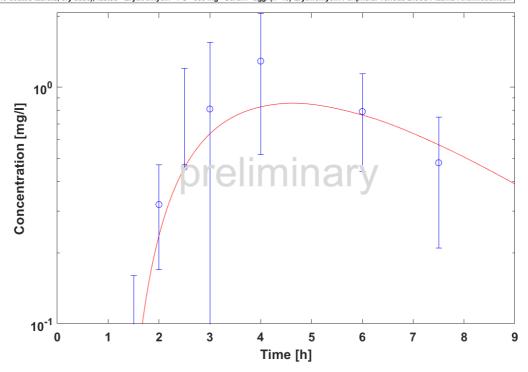


-Peripheral Venous Blood-Plasma-Concentration ric-coated erythromycin tablets (study 1) - Erythromycin - PO - 500 mg - Serum - agg. (n=21)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean ed erythromycin tablets (study 2) - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

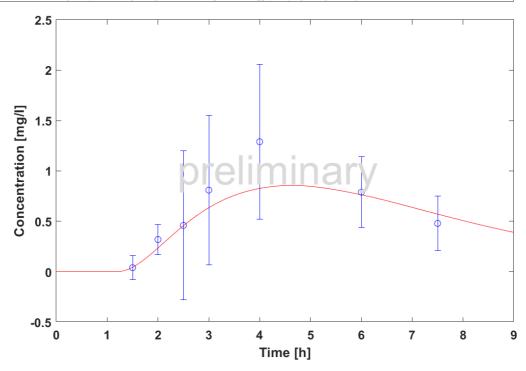


-Peripheral Venous Blood-Plasma-Concentration ric-coated erythromycin tablets (study 1) - Erythromycin - PO - 500 mg - Serum - agg. (n=21)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean ed erythromycin tablets (study 2) - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

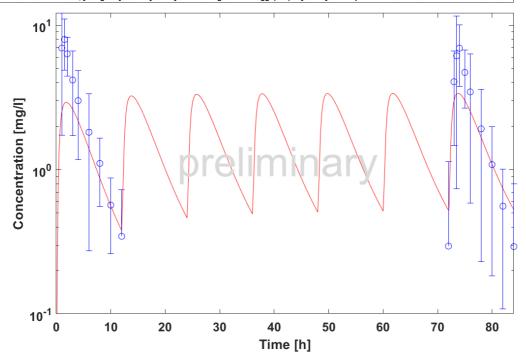




omycin-Peripheral Venous Blood-Plasma-Concentration (enteric-coated tablets, ery base), fasted - Erythromycin - PO - 500 mg - Serum - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

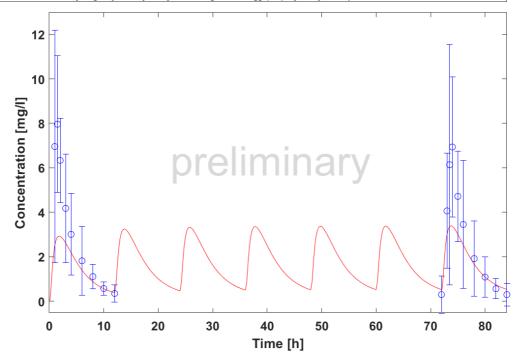


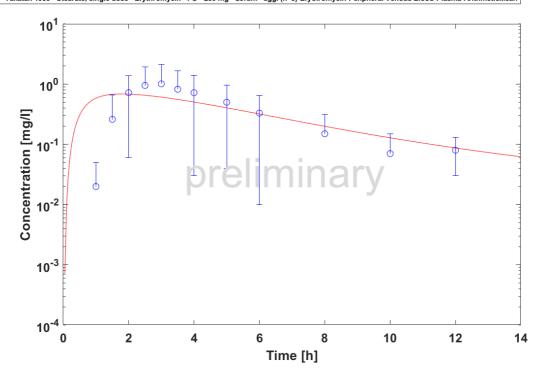
1000mg_Miglioli1990_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
Miglioli 1990 - First dose, young subjects - Erythromycin - PO - 1 g - Serum - agg. (n=8)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean
Miglioli 1990 - Seventh dose, young subjects - Erythromycin - PO - 1 g - Serum - agg. (n=8)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



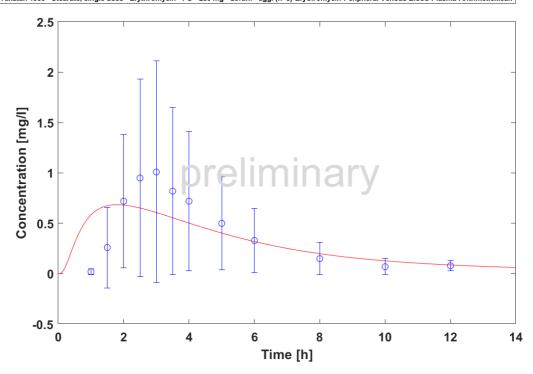
Time Profile Analysis

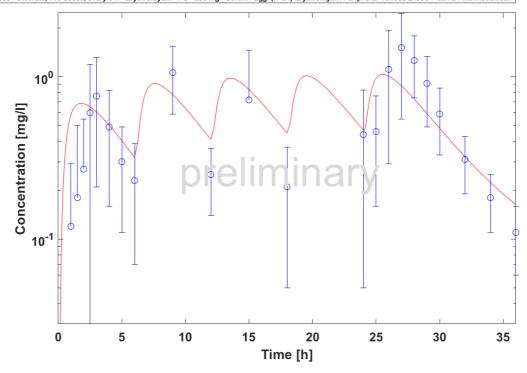
1000mg_Miglioli1990_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
Miglioli 1990 - First dose, young subjects - Erythromycin - PO - 1 g - Serum - agg. (n=8)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean
Miglioli 1990 - Seventh dose, young subjects - Erythromycin - PO - 1 g - Serum - agg. (n=8)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



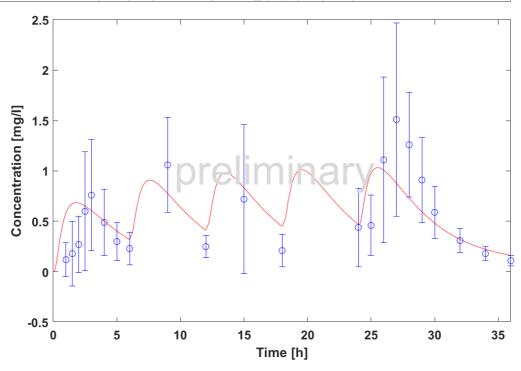


250mg_Yakatan1980_stearate-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
Yakatan 1980 - Stearate, single dose - Erythromycin - PO - 250 mg - Serum - agg. (n=8)-Erythromycin-Peripheral Venous Blood-Plasma-Arithmetich

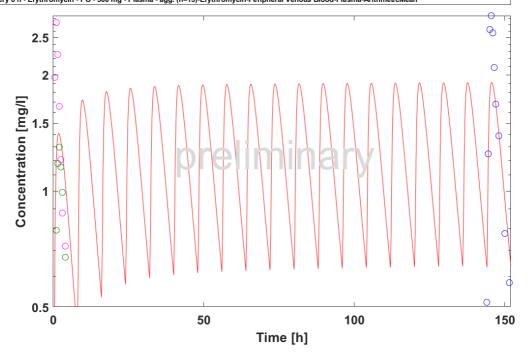




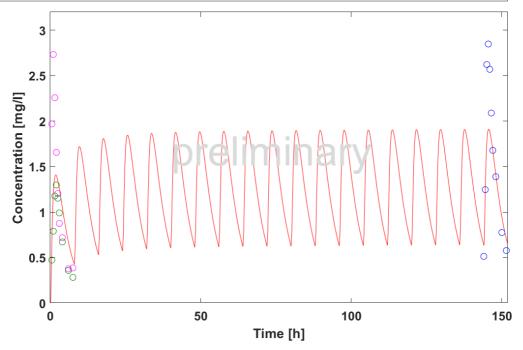
ng_Yakatan1980_stearate_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
tan 1980 - Stearate, five doses, every 6 h - Erythromycin - PO - 250 mg - Serum - agg. (n=24)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMea



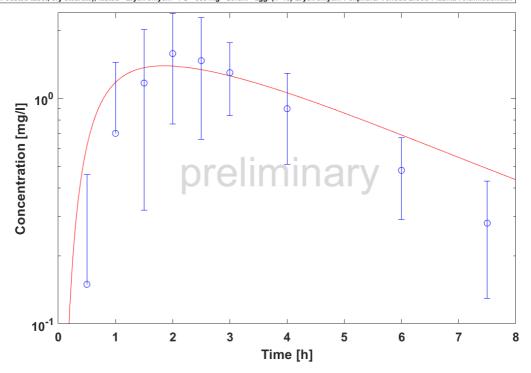
178_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration
ngle dose before standardized breakfast - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMea
ngle dose after overnight fast - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean
y 7, every 8 h - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



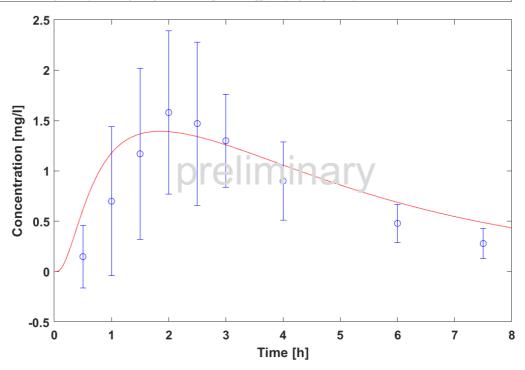
178_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration to the dose before standardized breakfast - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMeatigle dose after overnight fast - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean y 7, every 8 h - Erythromycin - PO - 500 mg - Plasma - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



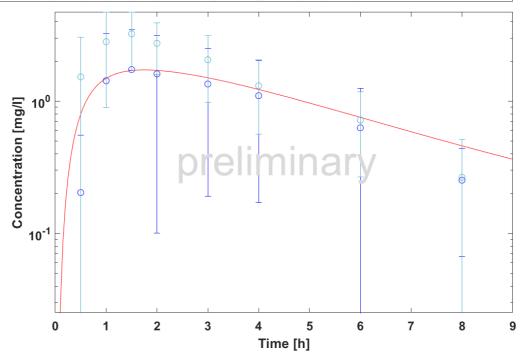




hromycin-Peripheral Venous Blood-Plasma-Concentration
1 (film-coated tabet, ery stearate), fasted - Erythromycin - PO - 500 mg - Serum - agg. (n=15)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

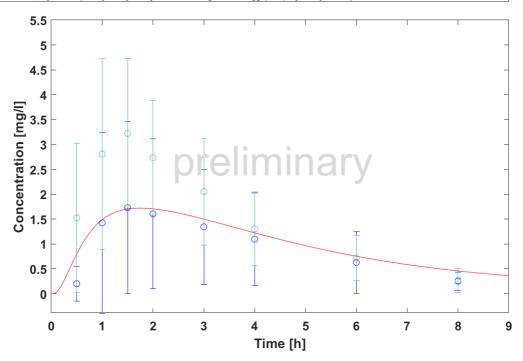


ythromycin-Peripheral Venous Blood-Plasma-Concentration
(film-coated tabet, ery stearate), study 2 - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean
(film-coated tabet, ery stearate), study 2 - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



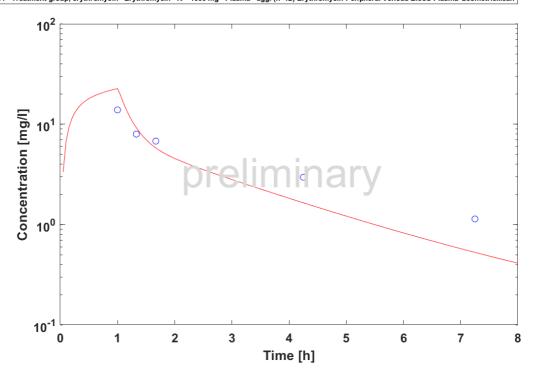
Time Profile Analysis

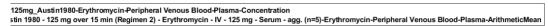
ythromycin-Peripheral Venous Blood-Plasma-Concentration
(film-coated tabet, ery stearate), study 2 - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean
(film-coated tabet, ery stearate), study 2 - Erythromycin - PO - 500 mg - Serum - agg. (n=12)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean

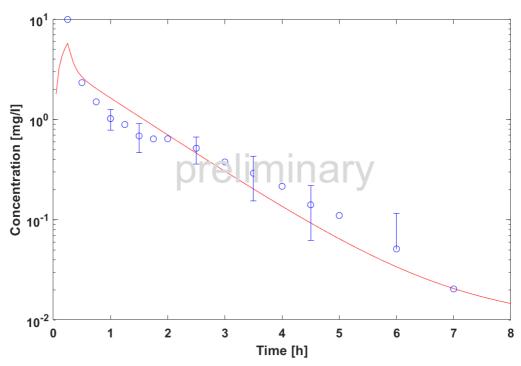


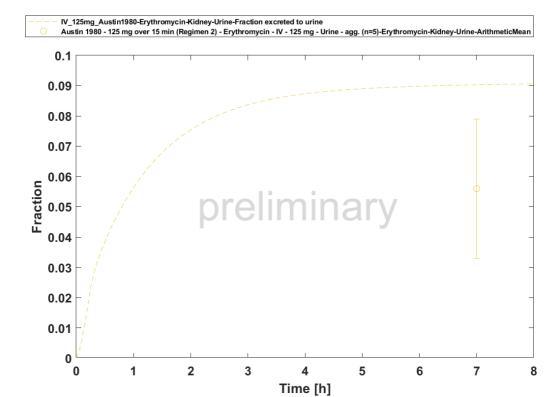
Time Profile Analysis 1

3.3.2 Model Verification

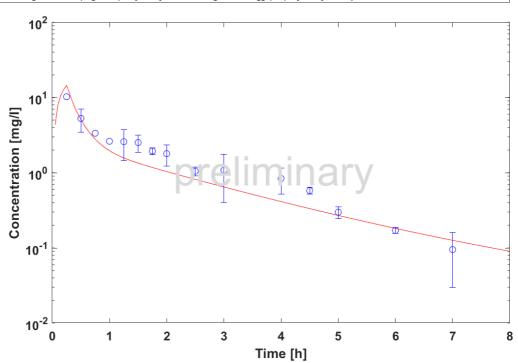


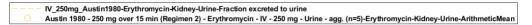


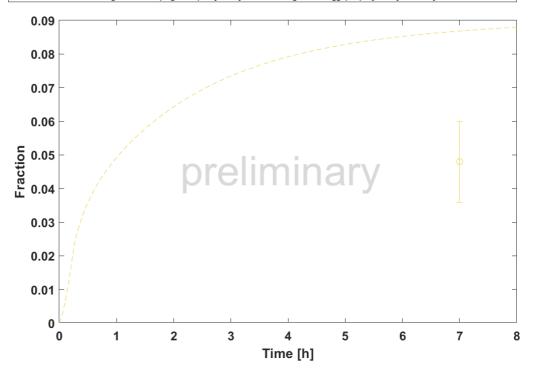


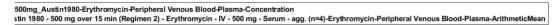


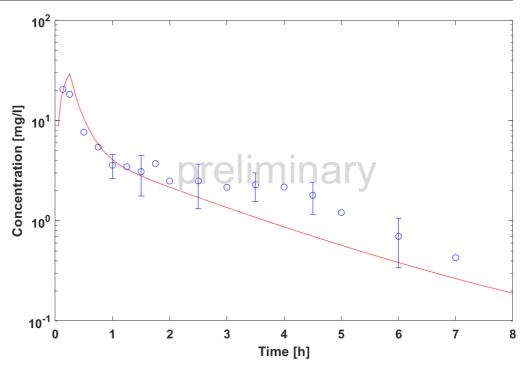


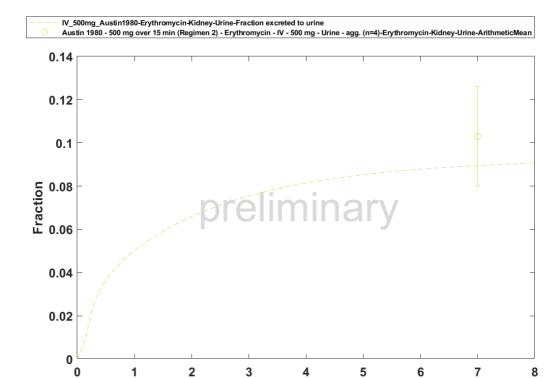






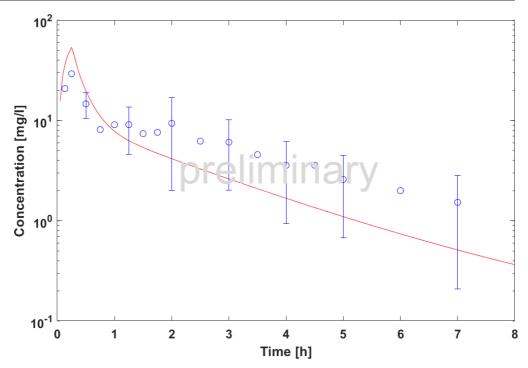


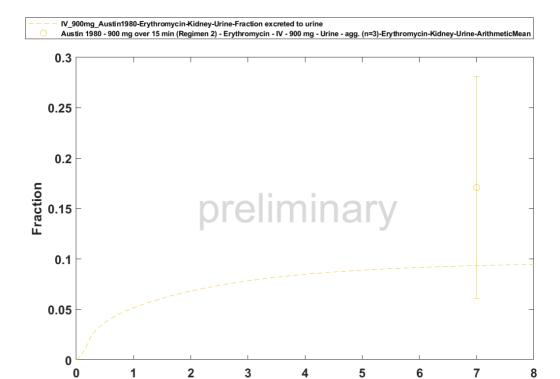






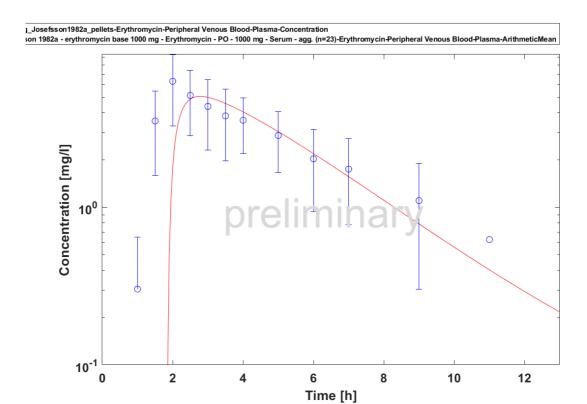
Time [h]

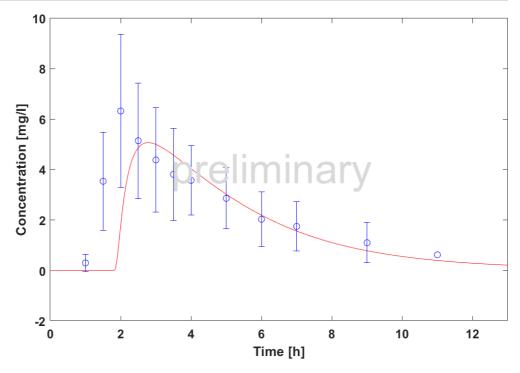




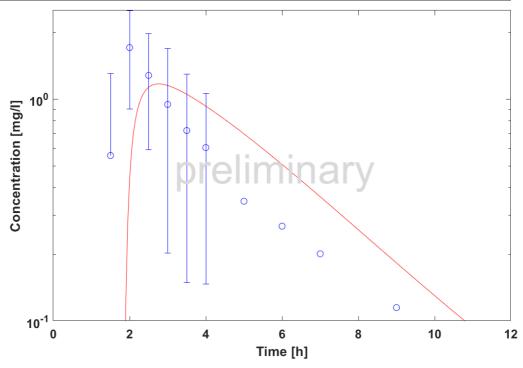
Time [h]

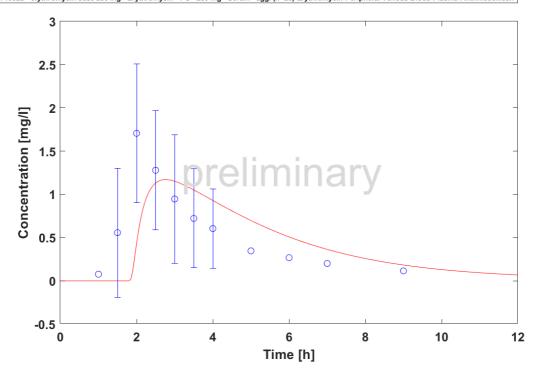
Time Profile Analysis 1

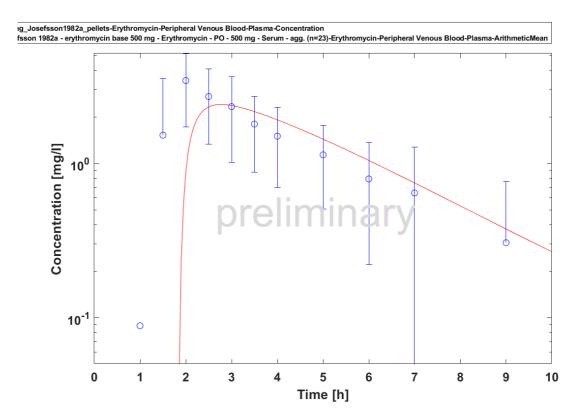


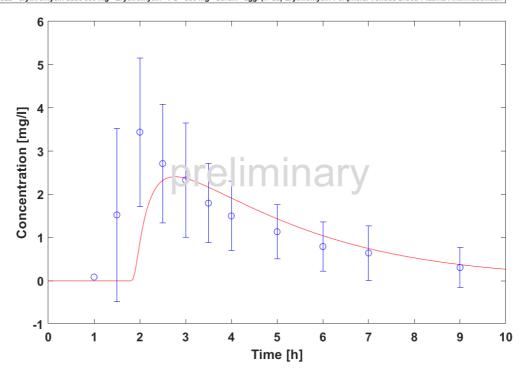




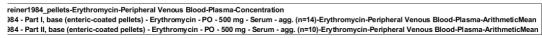


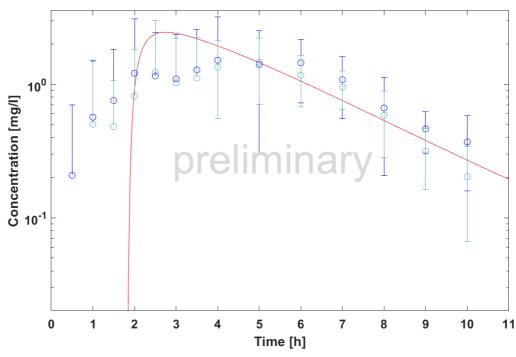


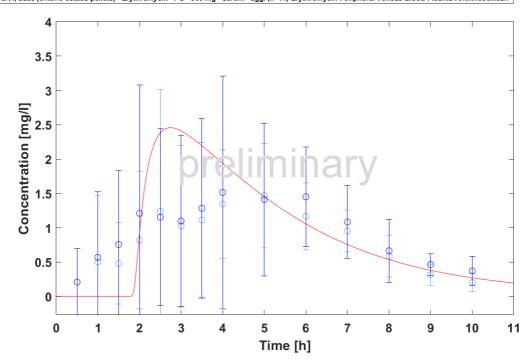


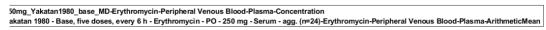


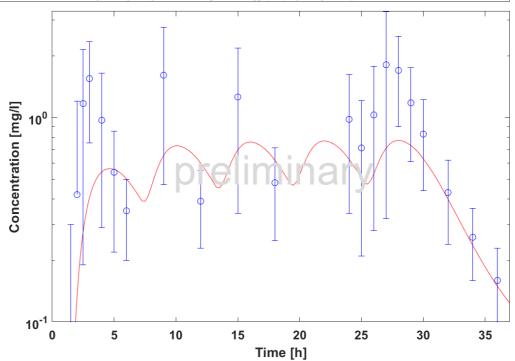
Time Profile Analysis 1

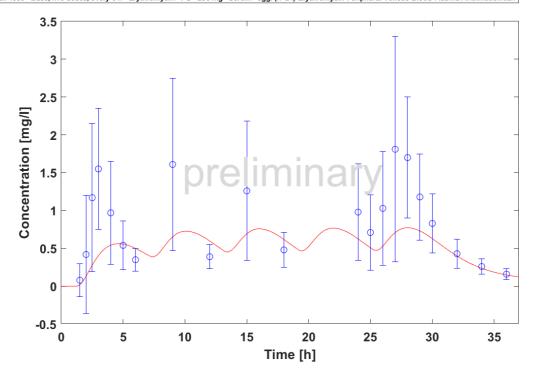


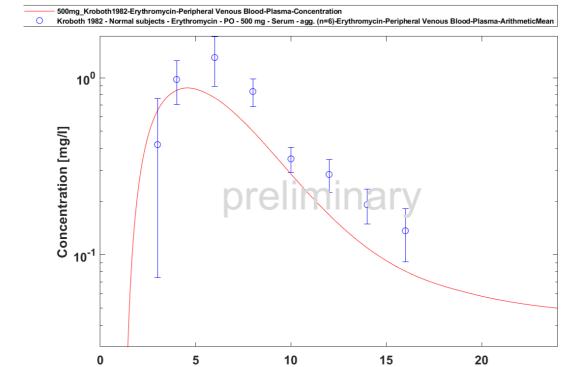




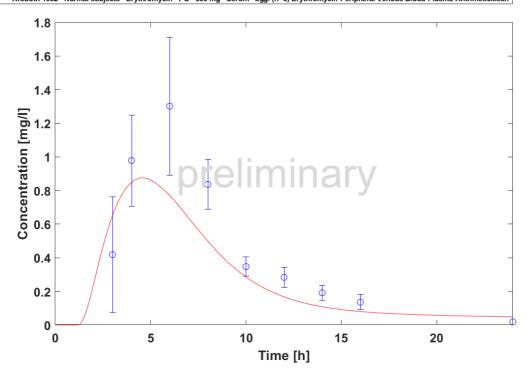


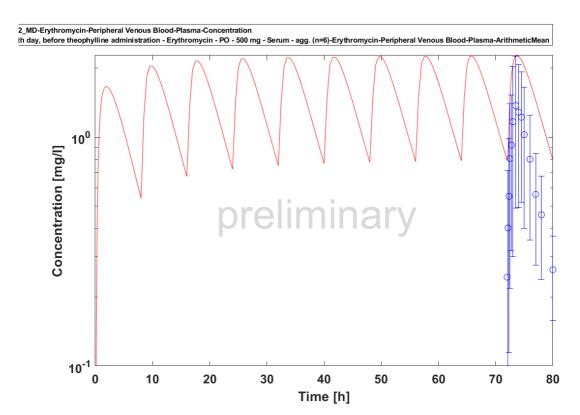




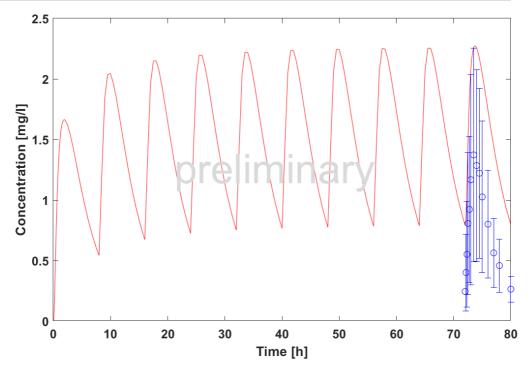


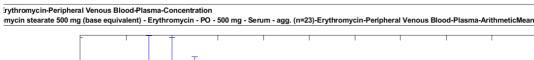
Time [h]

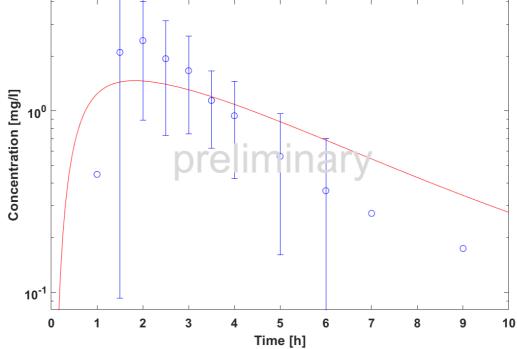


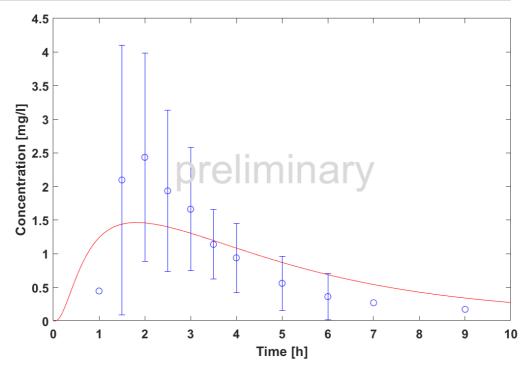


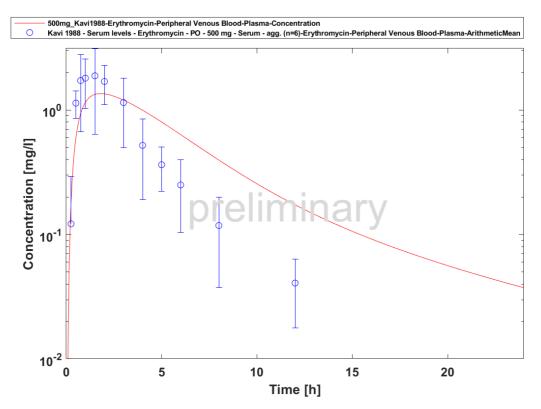
2_MD-Erythromycin-Peripheral Venous Blood-Plasma-Concentration th day, before theophylline administration - Erythromycin - PO - 500 mg - Serum - agg. (n=6)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMea

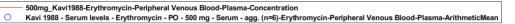


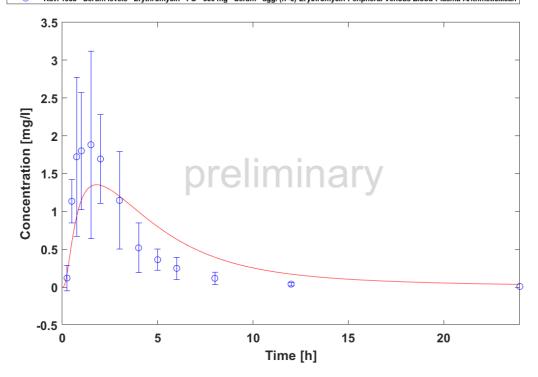




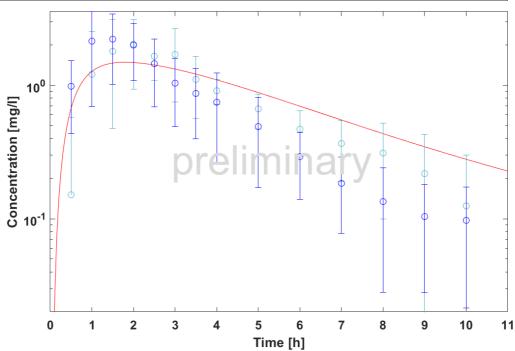




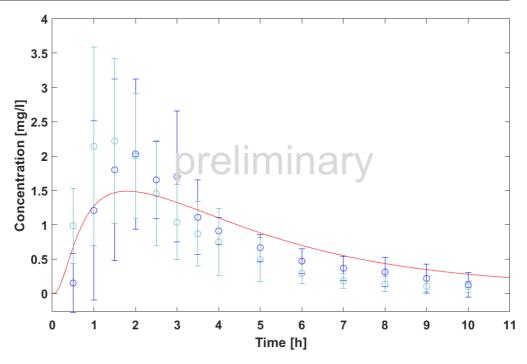








– 500mg_Schreiner1984_stearate-Erythromycin-Peripheral Venous Blood-Plasma-Concentration Schreiner 1984 - Part I, stearate tablet - Erythromycin - PO - 500 mg - Serum - agg. (n=14)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean Schreiner 1984 - Part II, stearate tablet - Erythromycin - PO - 500 mg - Serum - agg. (n=10)-Erythromycin-Peripheral Venous Blood-Plasma-ArithmeticMean



4 Conclusion

The final erythromycin PBPK model applies metabolism by CYP3A4, hepatic plasma clearance, glomerular filtration and mechanism-based inhibition of CYP3A4. Overall, the model adequately describes the oral pharmacokinetics of erythromycin in healthy adults receiving different single or multiple doses of several oral dosage forms.

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