

# Compound Name: Midazolam

Property	Value
Compound type:	Substrate
Prefix:	Sim
Species:	Human
File data last updated:	V9: Permeability and UGT data added V15: SAC added to minimal PBPK
	V18: Compound type, Monoprotic base
	V19: ADAM absorption option added (not default) Km updated for CYP3A5 formation of 4-OH UGT data changed to HLM
Population used for verification:	Sim-Healthy Volunteer

## Simcyp Version this document relates to: V19r1

## Prepared: March 2025

The Sim-Midazolam model within the Simcyp Compound Database has been developed as a probe substrate of CYP3A4.

This document provides:

- 1. Examples of model performance
- 2. A summary of the key pharmacokinetic features of Sim-Midazolam considered within the model

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# 1. MODEL PERFORMANCE.

#### A. Table 1. Sim-Midazolam Model Summary

Parameter	Model	Input		
Absorption	1st order absorption	User inputs for fa and Ka		
Distribution	Minimal PBPK model with SAC	Fitted SAC: $K_{in} K_{out}$ and $V_{sac}$ . User input $V_{ss}$ (based on clinical data)		
Elimination	Enzyme kinetics	Recombinant data for CYP3A4 and CYP3A5; HLM data for UGT1A4 Renal clearance from clinical studies		

#### **OPTIMISED PARAMETERS**

- k<sub>in</sub>
- $\bullet \ k_{out}$
- $V_{sac}$

For V15 a single adjusting compartment (SAC) was added to the minimal PBPK model. The relevant parameters ( $k_{in}$ ,  $k_{out}$  and  $V_{sac}$ ) were fitted using the parameter estimation (PE) module in V14 to recover the PK data following an single IV dose from Kupferschmidt *et al.*, 1995.

Drug characteristics based on the DIDB drug monograph DDI summary Monographs - Certara Drug Interaction Solutions

CYP3A FDA clinical index substrate CYP3A sensitive substrate



### Oral administration, Single dosing



**Figure 1.** Simulated (black line) and observed (data points) mean plasma concentration-time profiles of midazolam after a single oral dose of (A) 2 mg (10 trials of 15 subjects, 18-55 years, 19% female) (B) 5 mg (10 trials of 12 subjects, 18-56 years, 40% female), (C) 7.5 mg (10 trials of 10 subjects, 19-44 years, 50% female) and (D) 15 mg (10 trials of 10 subjects, 19-37 years, 20% female). Observed data were extracted from (A) Krishna *et al.*, 2012 (blue), Stoch *et al.*, 2009 (red), Templeton *et al.*, 2010 (green) and Wandel *et al.*, 2000 (yellow); (B) Lee *et al.*, 2002 (red) and Ozdemir *et al.*, 2006 (green); (C) Abel *et al.*, 2008 (red), Ahonen *et al.*, 1997 (yellow), and Mandema *et al.*, 1992 (green); (D) Allonen *et al.*, 1981 (red), Backman *et al.*, 1996 (yellow), and Kupferschmidt *et al.*, 1995 (green). The grey lines represent the predictions from individual trials. The dash lines represent the 5<sup>th</sup> and 95<sup>th</sup> percentiles of the total virtual population.



Oral administration, Single dosing (log scale)



**Figure 2.** Simulated (black line) and observed (data points) mean plasma concentration-time profiles on a semi-logarithmic scale of midazolam after a single oral dose of (A) 2 mg (10 trials of 15 subjects, 18-55 years, 19% female), (B) 5 mg (10 trials of 12 subjects, 18-56 years, 40% female), (C) 7.5 mg (10 trials of 10 subjects, 19-44 years 50% female) and (D) 15 mg (10 trials of 10 subjects, 19-37 years, 20% female). Observed data were extracted from (A) Krishna *et al.*, 2012 (blue), Stoch *et al.*, 2009 (red), Templeton *et al.*, 2010 (green), and Wandel *et al.*, 2000 (yellow); (B) Lee *et al.*, 2002 (red) and Ozdemir *et al.*, 2006 (green); (C) Abel *et al.*, 2008 (red), Ahonen *et al.*, 1997 (yellow), and Mandema *et al.*, 1992 (green); (D) Allonen *et al.*, 1981 (red), Backman *et al.*, 1996 (yellow), and Kupferschmidt *et al.*, 1995 (green). The grey lines represent the predictions from individual trials. The dash lines represent the 5<sup>th</sup> and 95<sup>th</sup> percentiles of the total virtual population.



Intravenous administration, Single dosing



**Figure 3.** Simulated (black line) and observed (data points) mean plasma concentration-time profiles of midazolam after an intravenous dose of (A) 0.15 mg/kg (10 trials of 6 subjects, 22-27 years, 50% female), (B) 1 mg (10 trials of 15 male subjects, 24-44 years), (C) 2 mg (10 trials of 11 subjects, 19-43 years 16% female) and (D) 5 mg (10 trials of 8 subjects, 22-30 years, 50% female). Observed data was extracted from (A) Heizmann *et al.*, 1983 (red), (B) Wandel *et al.*, 2000 (green), (C) Farkas *et al.*, 2007 (yellow), Tsunoda *et al.*, 1999 (blue), and (D) Schwagmeier *et al.*, 1998 (orange). The grey lines represent the predictions from individual trials. The dash lines represent the 5<sup>th</sup> and 95<sup>th</sup> percentiles of the total virtual population.

Intravenous administration, Single dosing (log scale)



**Figure 4.** Simulated (black line) and observed (data points) mean plasma concentration-time profiles on a semi-logarithmic scale of midazolam after an intravenous dose of (A) 0.15 mg/kg (10 trials of 6 subjects, 22-27 years, 50% female), (B) 1 mg (10 trials of 15 male subjects, 24-44 years), (C) 2 mg (10 trials of 11 subjects, 19-43 years 16% female) and (D) 5 mg (10 trials of 8 subjects, 22-30 years, 50% female). Observed data was extracted from (A) Heizmann *et al.*, 1983 (red), (B) Wandel *et al.*, 2000 (green), (C) Farkas *et al.*, 2007 (yellow), Tsunoda *et al.*, 1999 (blue) and (D) Schwagmeier *et al.*, 1998 (orange). The grey lines represent the predictions from individual trials. The dash lines represent the 5<sup>th</sup> and 95<sup>th</sup> percentiles of the total virtual population.

## 2. MODEL OVERVIEW.

#### ABSORPTION

Oral midazolam is absorbed rapidly and almost completely from the gastrointestinal tract. Peak plasma concentrations are reached within 1 hour (Figure 5).





**Figure 5.** Simulated (open circles; mean  $\pm$  SD ) (10 trials of 10 subjects, 19-55 years, 50% female) and observed (closed circles; mean  $\pm$  SD) values of Tmax for midazolam after a single oral dose of 7.5 mg. Predicted mean ( $\pm$  SD) T<sub>max</sub> values (n=100) are 0.66  $\pm$  0.16 hours. Observed data were reported by Abel *et al.*, 2008 (11), Eap *et al.*, 2004 (12), and Mandema *et al.*, 1992 (13).





Figure 6. Simulated (open circles; mean  $\pm$  SD) (10 trials of 10 subjects, 19-55 years, 50% female) and observed (closed circles; mean $\pm$  SD) values of C<sub>max</sub> for midazolam after a single oral dose of 7.5 mg. Predicted mean ( $\pm$  SD) maximal plasma concentrations (n = 100) are 37.59  $\pm$  23.51 ng/mL. Observed data were reported by Abel *et al.*, 2008 (11), Eap *et al.*, 2004 (12), and Mandema *et al.*, 1992 (13).

#### DISTRIBUTION

Following intravenous administration, midazolam is rapidly and widely distributed. In the V15 update to the Sim-Midazolam file, distribution was described using a SAC compartment within the minimal PBPK model. The parameters were fitted using IV data reported by Kupferschmidt *et al.*, 1995. Values of  $k_{in} = 0.2$  1/h,  $k_{out} = 0.25$  1/h,  $V_{sac}$  0.23 L/kg and  $V_{ss} = 0.88$  L/kg are used in the Sim-Midazolam file. Midazolam is significantly bound to plasma protein (fu 0.032).





Figure 7. Simulated (open circles; mean  $\pm$  SD) (10 trials of 6 male subjects, 22-27 years) and observed (closed circles; mean $\pm$  SD) values of midazolam V<sub>ss</sub> after an IV dose of 0.15 mg/kg. Observed data were reported by Heizmann *et al.*, 1983 (11). The mean of the virtual population (n= 60) was 0.92  $\pm$  0.24 L/kg.

#### ELIMINATION

Less than 1% midazolam is excreted unchanged in the urine (Thummel *et al.*, 1996). The drug is cleared virtually entirely by CYP3A4 and CYP3A5 mediated metabolism in the gastrointestinal tract and liver (Emoto *et al.*, 2006; 2007). The primary metabolite is 1-hydroxy midazolam. Both CYP3A4 and CYP3A5 also form 4-hydroxy midazolam but to a much lesser extent. UGT1A4 has a minor (< 5%) contribution to the metabolic elimination of midazolam (Hyland *et al.*, 2009; Figure 8).





Figure 8. Predicted mean contribution of metabolic and renal clearance to the systemic elimination of midazolam (10 trials of 10 subjects, 20-50 years, 50% female).

The predicted systemic and oral clearance values are consistent with clinical data as shown in Figure 9 and Figure 10, respectively.



**Figure 9.** Simulated (open circles; mean  $\pm$  SD) (10 trials of 12 subjects, 18-45 years, 35% female) and observed (closed circles; mean  $\pm$  SD) values of midazolam systemic clearance after an iv dose of 2 mg. Observed data were reported by Farkas *et al.*, 2007 (11), Majumdar *et al.*, 2007 (12), and Tsunoda *et al.*, 1999 (13). The simulated population mean value was  $25.13 \pm 7.97$  L/h.





Figure 10. Simulated (open circles; mean  $\pm$  SD) (10 trials of 10 subjects, 21-54 years, 22% female) and observed (closed circles; mean  $\pm$  SD) values of midazolam oral clearance after an oral dose of 2 mg. Observed data were reported by Thummel *et al.*, 1996 (11) and Wandel *et al.*, 2000 (12). The simulated population mean value was 111.84  $\pm$  99.49 L/h.





Figure 11. Simulated (open circles; mean  $\pm$  SD) (10 trials of 9 subjects, 33% female 19-41 years) and observed (closed circles; mean  $\pm$  SD) values of midazolam half-life after an iv dose of 2 mg. Observed data was reported by Tsunoda *et al*, 1999. (11). The simulated population mean value is  $4.50 \pm 0.91$  h.



# BIOAVAILABILITY

Simulated F, Fg and Fh are comparable to observed data. Overall bioavailability (F)  $0.29 \pm 0.05$  Fraction escaping gut metabolism (Fg)  $0.51 \pm 0.1$  Fraction escaping hepatic metabolism (Fh)  $0.58 \pm 0.09$  (Taken from Galetin *et al.*, 2008)



Figure 12. Simulated (open circles; mean  $\pm$  SD) (10 trials of 10 subjects, 50% female 20-50 years) and observed (closed circles; mean  $\pm$  SD) values of midazolam Fg values. Observed data were obtained from Galetin *et al.*, 2008 and represent the mean  $\pm$  SD values reported. The simulated population mean values are 0.59.





Figure 13. Simulated (open circles; mean  $\pm$  SD) (10 trials of 10 subjects, 50% female 20-50 years) and observed (closed circles; mean  $\pm$  SD) values of midazolam F<sub>h</sub> values. Observed data were obtained from Galetin *et al.*, 2008 and represent the mean  $\pm$  SD values reported. The simulated population mean values are 0.57.



Figure 14. Simulated (open circles; mean  $\pm$  SD) (10 trials of 10 subjects, 50% female 20-50 years) and observed (closed circles; mean  $\pm$  SD) values of midazolam overall F values. Observed data were obtained from Galetin *et al.*, 2008 and represent the mean  $\pm$  SD values reported. The simulated population mean values are 0.29.

## **INTERACTION**

#### Sim-Midazolam as a victim

Investigation of Midazolam as a victim allows verification of the fm CYP3A4 used in the file. To verify the contribution of CYP3A4 within the Sim-Midazolam file, simulated drug-drug interaction (DDI) studies with the CYP3A4 inhibitors were compared to observed studies. The trial designs used were based on the clinical studies and the dosing regimen for each study is shown in Table 1. Simulated and observed AUC and  $C_{max}$  ratios are shown in Table 2.



## Table 2. Dosing regimens for DDI studies

Study	Reference	Inhibitor	Midazolam
1	Ahonen <i>et al.</i> , 1997	Fluconazole 400 mg SD	7.5 mg SD (1 hr after fluconazole)
2	Olkkola <i>et al.</i> , 1994	Itraconazole 200 mg QD for 4 days	$\overline{7.5 \text{ mg SD (Day 4)}}$
3	Yeates <i>et al.</i> , 1996	Clarithromycin 250 mg BID for 5 days	15  mg SD (Day 5)
4	Backman <i>et al.</i> , 1994	Diltiazem 60 mg TID for 2 days (5 doses)	15 mg SD (Day 2) (1h after diltiazem dose 4)
5	Backman <i>et al.</i> , 1994	Verapamil 80 mg TID for 2 days (5 doses)	15  mg SD (Day 2) (1h after verapamil dose 4)
6	Tsunoda <i>et al.</i> , 1999	Ketoconazole 200 mg BID for 2 days (3 doses)	6 mg SD (12 hrs after first KTZ dose)
7	Tsunoda <i>et al.</i> , 1999	Ketoconazole 200 mg BID for 2 days (3 doses)	2 mg iv SD (12 hrs after first KTZ dose)
8	Olkkola <i>et al.</i> , 1994	Ketoconazole 400 mg QD for 4 days	$\overline{7.5 \text{ mg SD (Day 4)}}$
9	Stoch <i>et al.</i> , 2009	Ketoconazole 400 mg QD for 5 days (5 doses)	2  mg SD (Day 5)
10	Salonen <i>et al.</i> , 1986	Cimetidine 400 mg SD	15 mg SD PO, 2h after cimetidine
11	Elliott <i>et al.</i> , 1984	Cimetidine 200 mg TID with 400 mg at night on day 1. 200 mg on day 2	15 mg SD PO, day2, 2.5 h after cimetidine
12	Olkkola <i>et al.</i> , 1993	Erythromycin 500 mg TID for 6 days, day 6 2 hrs before midazolam	15 mg oral SD day 6
13	Lam <i>et al.</i> , 2003	Fluvoxamine 50 mg (36.65 mg free base) BID days 1-6, 100 mg (73.33 mg free base) BID days 7-12	10 mg SD on day 12 (1h after fluvoxamine)
14	Ahonen <i>et al.</i> , 1995	Itraconazole 100 mg QD for 4 days	$\overline{7.5 \text{ mg SD}}$ on day 4
15	Backman <i>et al.</i> , 1998	Itraconazole 200 mg QD for 4 days	$\overline{7.5 \text{ mg SD on}} \text{ day } 4$
16	Stoch <i>et al.</i> , 2009	Ketoconazole 400 mg oral SD	2 mg oral SD, 5 min after ketoconazole dosing



Study	Reference	Inhibitor	Midazolam
17	Eap <i>et al.</i> , 2004	Ketoconazole 200 mg BID for 2 days	75 ug SD Day 3
18	Martinez <i>et al.</i> , 1999	Cimetidine 800 mg SD	7.5 mg SD
19	Chen <i>et al.</i> , 2006	Fluvoxamine 150 mg QD oral for 28 days	0.025  mg/kg SD IV on day 28



**Table 3** Observed and predicted mean  $C_{max}$  and AUC ratios for CYP3A4 inhibitors. Predicted values show mean and trial range from 10 simulated trials matching the clinical study design.

		Obse	rved	Pred	licted	Predicte	d/Observed
Study	Dose	AUC Ratio	$\begin{array}{c} \mathrm{C}_{\mathrm{max}} \\ \mathrm{Ratio} \end{array}$	AUC Ratio	C <sub>max</sub> Ratio	AUC Ratio	C <sub>max</sub> Ratio
1	Fluconazole 400 mg SD	2.30	3.73	1.91 (1.79 to 2.16)	3.03 (2.77 to 3.39)	0.83	0.81
2	Itraconazole 200 mg QD	3.41	10.77	2.82 (2.37  to  3.57)	9.59 (8.62  to  11.76)	0.83	0.89
3	Clarithromycin 250 mg BID	2.44	3.57	2.07 (1.88  to  2.50)	4.13 (3.32  to  5.70)	0.85	1.16
4	Verapamil 80 mg TID	2.05	3.75	1.55 (1.44  to  1.65)	2.03 (1.88  to  2.27)	0.76	0.54
5	Verapamil	1.96	2.92	2.07 (1.87  to  2.23)	3.50 (3.11  to  4.04)	1.06	1.20
6	Ketoconazole 200 mg BID	4.24	15.10	3.56 (2.98  to  4.49)	12.56 (10.03 to 16.16)	0.84	0.83
7	Ketoconazole 200 mg BID	NA	4.80	1.00 (1.00  to  1.00)	4.67 (4.11  to  5.34)	NA	0.97
8	Ketoconazole 400 mg QD	4.09	15.90	3.78 (3.06 to 5.04)	12.50 (9.73 to 16.10)	0.92	0.79
9	Ketoconazole 400 mg QD*	5.42	13.96	3.76 (3.05  to  5.04)	12.12 (9.41  to  15.56)	0.69	0.87
10	Cimetidine 400 mg SD†	1.36	1.37	1.51 (1.41  to  1.65)	$1.42 \ (1.34 \text{ to } 1.52)$	1.11	1.03
11	Cimetidine 200 mg $MD$	2.02	2.38	$1.22 \ (1.15 \ \text{to} \ 1.27)$	1.19 (1.13 to 1.23)	0.60	0.50
12	Erythromycin 500 mg MD	4.42	2.70	8.74 (6.52 to 10.6)	3.01 (2.28 to 3.78)	1.98	1.11
13	Fluvoxamine 50 mg BID	1.39	1.38	1.37 (1.29 to 1.48)	$1.25 \ (1.21 \text{ to } 1.32)$	0.99	0.91
14	Itraconazole 100 mg QD	5.75	2.56	5.11 (3.77  to  6.41)	$2.38 \ (2.15 \ {\rm to} \ 2.69)$	0.89	0.93
15	Itraconazole 200 mg QD†	5.17	2.91	7.15~(5.87  to  9.32)	2.82 (2.43 to 3.53)	1.38	0.97
16	Ketoconazole 400 mg SD	10.31	5.01	13.34 (11.16  to  15.87)	4.07 (3.17  to  5.05)	1.29	0.81
17	Ketoconazole 200 mg BID	6.47	3.74	4.9 (3.07 to 8.45)	2.56 (1.94 to 3.6)	0.76	0.68
18	Cimetidine 800 mg	1.50	NA	2 (1.86 to 2.15)	1.63 (1.55 to 1.74)	1.33	NA
19	Fluvoxamine 150 mg QD	1.49	NA	1.76 (1.63 to 1.86)	1.56 (1.48  to  1.62)	1.18	NA

Note:

AUC calulated from zero to infinity unless otherwise stated.

 $^*$  Geometric mean

 $^\dagger$  AUC last



## WHAT IS NOT CURRENTLY CONSIDERED IN THE MODEL:

. The ability to model the concentration time profile of the primary (active) metabolite 1-OH midazolam

. The ability to model the pharmacodynamics of midazolam and it's metabolites



## Table 4 Input parameters for SIM-Midazolam

Parameter	Value	Method/Reference
Molecular weight (g/mol)	325.8	Pubchem
Log P	3.53	Calculated in ALOGPS 2.1
Compound Type	Monoprotic base	Simcyp data archive
pKa	6.0	Meta-analysis (Dollery, 1999; Orive <i>et al.</i> , 1989; Andersin <i>et al.</i> , 1991; Walser <i>et al.</i> , 1978;Vire <i>et al.</i> , 1986)
B/P	0.603	Meta-analysis (Allonen <i>et al.</i> , 1981; Heizmann <i>et al.</i> , 1983; Simcyp data archive)
fu	0.032	Meta-analysis (Greenblatt <i>et al.</i> , 1984; Vinik <i>et al.</i> , 1983; Thummel <i>et al.</i> , 1996)
Main binding protein	Human serum albumin	Boman, 1973
fa	1	Assumed
$\overline{\text{ka (1/h)}}$	3	Allonen et al., 1981
fu <sub>gut</sub>	1	Assumed
$\overline{\mathrm{Q}_{\mathrm{gut}}~(\mathrm{L/h})}$	16.184	Predicted (Yang <i>et al.</i> , 2007)
Distribution Model	Minimal PBPk model	
$\overline{\mathrm{V}_{\mathrm{ss}}~(\mathrm{L/kg})}$	0.88	Kupferschmidt <i>et al.</i> , 1995
$k_{in} (1/h)$	0.2	Optimised to recover PK data following single IV dose (Kupferschmidt <i>et al.</i> , 1995)
Kout $(1/h)$	0.25	Optimised to recover PK data following single IV dose (Kupferschmidt <i>et al.</i> , 1995)
$\overline{\rm V_{SAC}}$ (L/kg)	0.23	Optimised to recover PK data following single IV dose (Kupferschmidt <i>et al.</i> , 1995)
Metabolism	Enzyme kinetics	
Enzyme	CYP3A4	
Pathway	1-OH	



$V_{max}$ 5.23       Meta-analysis (Carr et al., 2006; Emoto et al., 2007;Galetin et al., 2007;Galetin et al., 2004; Williams et al., 2003; Soars et al., 2006; Walsky et al., 2004; Nakajima et al., 2004; Vacaver et al., 2003; Emoto et al., 2006; Emoto et al., 2003; Carr et al., 2006; Emoto et al., 2007;Galetin et al., 2007; Enoto et al., 2007;Galetin et al., 2007; Huang et al., 2007;Galetin et al., 2007; Emoto et al., 2007;Galetin et al., 2007; Huang et al., 2004; Nakajima et al., 2002;         Enzyme       CYP3A5         Pathway       1-OH         V <sub>max</sub> 19.7         Meta-analysis (Emoto et al., 2006; Eunoto et al., 2004; Soars et al., 2006; Walsky et al., 2004; Williams et al., 2002)         K <sub>m</sub> (\muM)       4.16         Meta-analysis (Emoto et al., 2004; Soars et al., 2006; Walsky et al., 2004; Williams et al., 2002)         Finzyme       CYP3A4         Pathway       4-OH         V <sub>max</sub> Meta-analysis (Emoto et al., 2007; Galetin et al., 2007; Emoto et al., 2004; Williams et al., 2002; Williams et al., 2002)         K <sub>m</sub> (\muM)       31.8         Meta-analysis (Emoto et al., 2004; Williams et al., 2004; Williams et al., 2002)	Parameter	Value	Method/Reference
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	V <sub>max</sub> (pmol/min/pmol)	5.23	Meta-analysis (Carr <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2003; Emoto <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2007;Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Nakajima <i>et al.</i> , 2002; Soars <i>et al.</i> , 2006; Walsky <i>et al.</i> , 2004; Weaver <i>et al.</i> , 2003; Williams <i>et al.</i> , 2002)
Enzyme       CYP3A5         Pathway       1-OH $V_{max}$ 19.7         (pmol/min/pmol)       19.7         Meta-analysis (Emoto et al., 2007; Galetin et al., 2006; Emoto et al., 2004; Williams et al., 2002) $K_m(\mu M)$ 4.16         Enzyme       CYP3A4         Pathway       4-OH         Vmax       (pmol/min/pmol)         Enzyme       CYP3A4         Pathway       4-OH         Vmax       5.2         (pmol/min/pmol)       31.8         Meta-analysis (Emoto et al., 2004; Williams et al., 2002)         K_m(\mu M)       31.8         Enzyme       CYP3A5         Pathway       4-OH	$\overline{\mathrm{K}_{\mathrm{m}}(\mu\mathrm{M})}$	2.16	Meta-analysis (Carr <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2003; Emoto <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2007;Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Nakajima <i>et al.</i> , 2002; Soars <i>et al.</i> , 2006; Walsky <i>et al.</i> , 2004; Weaver <i>et al.</i> , 2003; Williams <i>et al.</i> , 2002)
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$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Pathway	1-ОН	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	V <sub>max</sub> (pmol/min/pmol)	19.7	Meta-analysis (Emoto <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2007;Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Soars <i>et al.</i> , 2006; Walsky <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002)
EnzymeCYP3A4Pathway4-OH $V_{max}$ (pmol/min/pmol)5.2Meta-analysis (Emoto et al., 2007; Emoto et al., 2006; Galetin et al., 2004; Huang et al., 2004; Williams et al., 2002) $K_m(\mu M)$ 31.8Meta-analysis (Emoto et al., 2007; Emoto et al., 2006; Galetin et al., 2004; Huang et al., 2007; Emoto et al., 2006; Galetin et al., 2004; Huang et al., 2007; Emoto et al., 2006; Galetin et al., 2004; Huang et al., 2004; Williams et al., 2002)EnzymeCYP3A5Pathway4-OH	$\overline{\mathrm{K}_{\mathrm{m}}(\mu\mathrm{M})}$	4.16	Meta-analysis (Emoto <i>et al.</i> , 2006; Emoto <i>et al.</i> , 2007; Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Soars <i>et al.</i> , 2006; Walsky <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002)
Pathway4-OH $V_{max}$ (pmol/min/pmol)5.2Meta-analysis (Emoto <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002) $K_m(\mu M)$ 31.8Meta-analysis (Emoto <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2004; Huang <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2004; Huang <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2004; Huang <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Huang <i>et al.</i>	Enzyme	CYP3A4	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Pathway	4-OH	
$K_m(\mu M)$ 31.8Meta-analysis (Emoto et al., 2007; Emoto et al., 2006; Galetin et al., 2004; Huang et al., 2004; Williams et al., 2002)EnzymeCYP3A5Pathway4-OH	V <sub>max</sub> (pmol/min/pmol)	5.2	Meta-analysis (Emoto <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002)
Enzyme CYP3A5 Pathway 4-OH	$\overline{\mathrm{K}_{\mathrm{m}}(\mu\mathrm{M})}$	31.8	Meta-analysis (Emoto <i>et al.</i> , 2007; Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002)
Pathway 4-OH	Enzyme	CYP3A5	
	Pathway	4-OH	



Parameter	Value	Method/Reference
$\overline{V_{max}}$ (pmol/min/pmol)	4.03	Meta-analysis (Emoto $et al.$ , 2006; Galetin $et al.$ , 2004; Huang $et al.$ , 2004; Williams $et al.$ , 2002)
$\overline{\mathrm{K}_{\mathrm{m}}(\mu\mathrm{M})}$	38.4	Meta-analysis (Emoto <i>et al.</i> , 2006; Galetin <i>et al.</i> , 2004; Huang <i>et al.</i> , 2004; Williams <i>et al.</i> , 2002)
Enzyme	UGT1A4	
Pathway	Pathway 1	
$V_{max}$ (pmol/min/pmol)	445	Hyland <i>et al.</i> , 2009
$\overline{K_{m}(\mu M)}$	40.3	Hyland <i>et al.</i> , 2009
$\overline{\mathrm{CL}_{\mathrm{R}}~(\mathrm{L/h})}$	0.085	Thummel et al., 1996



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