

MATE1 Deficiency Exacerbates Dofetilide-Induced Proarrhythmia

Muhammad Erfan Uddin ¹, Eric D. Eisenmann ¹, Yang Li ¹, Kevin M. Huang ¹, Dominique A. Garrison ¹, Zahra Talebi ¹, Alice A. Gibson ¹, Yan Jin ¹, Mahesh Nepal ¹, Ingrid M. Bonilla ², Qiang Fu ¹, Xinxin Sun ¹, Alec Millar ³, Mikhail Tarasov ³, Christopher E. Jay ⁴, Xiaoming Cui ⁵, Heidi J. Einolf ⁵, Ryan M. Pelis ⁵, Sakima A. Smith ^{6,7}, Przemysław B. Radwański ^{3,7}, Douglas H. Sweet ⁴, Jörg König ⁸, Martin F. Fromm ⁸, Cynthia A. Carnes ^{3,7,9}, Shuiying Hu ³ and Alex Sparreboom ^{1,*}

¹ Division of Pharmaceutics and Pharmacology, College of Pharmacy and Comprehensive Cancer Center, The Ohio State University, Columbus, OH 43210, USA; uddin.33@osu.edu (M.E.U.); eisenmann.11@buckeyemail.osu.edu (E.D.E.); li.10991@osu.edu (Y.L.); huang.2834@buckeyemail.osu.edu (K.M.H.); garrison.220@buckeyemail.osu.edu (D.A.G.); talebi.9@osu.edu (Z.T.); gibson.972@osu.edu (A.A.G.); jin.1134@osu.edu (Y.J.); nepal.17@osu.edu (M.N.); pkpdfu@gmail.com (Q.F.); xinxin224@gmail.com (X.S.)

² Department of Physiology and Cell Biology, College of Medicine, The Ohio State University, Columbus, OH 43210, USA; ingrid.bonilla4@upr.edu

³ Division of Outcomes and Translational Sciences, College of Pharmacy, The Ohio State University, Columbus, OH 43210, USA; alec.miller@midwestern.edu (A.M.); tarasov.4@osu.edu (M.T.); radwanski.2@osu.edu (P.B.R.); carnes.4@osu.edu (C.A.C.); hu.1333@osu.edu (S.H.)

⁴ Department of Pharmaceutics, School of Pharmacy, Virginia Commonwealth University, Richmond, VA 23298, USA; jayce@mymail.vcu.edu (C.E.J.); dsweet@vcu.edu (D.H.S.)

⁵ Novartis Institute for Biomedical Research, East Hanover, NJ 07936, USA; xiaoming.cui@novartis.com (X.C.); heidi.einolf@novartis.com (H.J.E.); ryan.pelis@novartis.com (R.M.P.)

⁶ OSU Wexner Medical Center, Department of Internal Medicine, Division of Cardiovascular Medicine, The Ohio State University, Columbus, OH 43210, USA; sakima.smith@osumc.edu

⁷ Dorothy M. Davis Heart and Lung Research Institute, The Ohio State University, Columbus, OH 43210, USA

⁸ Institute of Experimental and Clinical Pharmacology and Toxicology, Friedrich-Alexander-Universität Erlangen-Nürnberg, 91054 Erlangen, Germany; joerg.koenig@fau.de (J.K.); martin.fromm@fau.de (M.F.F.)

⁹ Division of Pharmacy Practice and Science, College of Pharmacy, The Ohio State University, Columbus, OH 43210, USA

* Correspondence: sparreboom.1@osu.edu

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Table S1. Pharmacokinetic parameters of dofetilide after i.v. administration (2.5 mg/kg) in mice.

Group	C _{max} (ng/ml)	C _{max} (Fold-change)	AUC _{last} (ng×h/ml)	AUC _{last} (Fold-change)
Female				
Wild-type	1640 ± 170		673 ± 28.0	
OCT1/2 ^{-/-}	1960 ± 126	1.29	756 ± 47.0	1.19
MATE1 ^{-/-}	1520 ± 149	0.93	635 ± 61.0	0.94
Male				
Wild-type	1540 ± 131		587 ± 29.4	
OCT1/2 ^{-/-}	1120 ± 113	0.81	461 ± 23.0	0.76
MATE1 ^{-/-}	1380 ± 112	0.90	603 ± 28.5	1.03

Data represent mean ± SEM.

Abbreviations: C_{max}, peak plasma concentration; AUC_{last}, area under the plasma concentration-time curve to the last measurable concentration.

Table S2. Influence of contraindicated drugs on the pharmacokinetic parameters of dofetilide (5 mg/kg) in female mice.

Concurrent treatment	C _{max} (ng/ml)	C _{max} (Fold-change)	AUC _{last} (ng×h/ml)	AUC _{last} (Fold-change)
None				
Wild-type	191 ± 13.0		329 ± 24.5	
OCT1/2 ^{-/-}	218 ± 18.0	1.14	362 ± 25.0	1.10
MATE1 ^{-/-}	276 ± 45.4	1.44	346 ± 37.0	1.05
Bictegravir (30 mg/kg)				
Wild-type	365 ± 72.0*	1.91	595 ± 194*	1.81
OCT1/2 ^{-/-}	282 ± 132	1.47	354 ± 46.4	1.08
MATE1 ^{-/-}	383 ± 86.4*	2.00	614 ± 77.0*	1.87
Cimetidine (100 mg/kg)				
Wild-type	656 ± 81.3***	3.43	1400 ± 183***	4.25
OCT1/2 ^{-/-}	420 ± 89.4**	2.20	1270 ± 150***	3.86
MATE1 ^{-/-}	625 ± 92.2***	3.27	1400 ± 89.5***	4.27
Ketoconazole (50 mg/kg)				
Wild-type	315 ± 34.0*	1.64	588 ± 64.0**	1.79
OCT1/2 ^{-/-}	475 ± 61.5**	2.48	1090 ± 128***	3.31
MATE1 ^{-/-}	455 ± 32.0***	2.38	938 ± 181***	2.85
Trimethoprim (100 mg/kg)				
Wild-type	264 ± 27.0	1.38	428 ± 25.4	1.30
OCT1/2 ^{-/-}	220 ± 38.0	1.15	440 ± 76.0	1.34
MATE1 ^{-/-}	212 ± 45.0	1.11	345 ± 29.3	1.05
Verapamil (10 mg/kg)				
Wild-type + Verapamil	518 ± 47.0***	2.71	602.0 ± 35.0*	1.83
OCT1/2 ^{-/-} + Verapamil	519 ± 42.4***	2.71	669.0 ± 42.0**	2.04
MATE1 ^{-/-} + Verapamil	384 ± 58.0**	2.01	638.0 ± 26.5**	1.94

Data represent mean \pm SEM. *P < 0.05, **P < 0.01, ***P < 0.001 versus single agent dofetilide.

Abbreviations: C_{max}, peak plasma concentration; AUC_{last}, area under the plasma concentration-time curve to the last measurable concentration.

Table S3. Influence of ketoconazole or verapamil on the pharmacokinetics of dofetilide (5 mg/kg) in male mice.

Concurrent treatment	C _{max} (ng/ml)	C _{max} (Fold-change)	AUC _{last} (ng×h/ml)	AUC _{last} (Fold-change)
None				
Wild-type	151 \pm 8.80		278 \pm 12.3	
OCT1/2 ^{-/-}	191 \pm 23.4	1.26	351 \pm 36.0	1.26
MATE1 ^{-/-}	235 \pm 50.0*	1.55	407 \pm 32.6**	1.46
DDI with Ketoconazole (50 mg/kg)				
Wild-type	415 \pm 43.0***	2.75	728 \pm 55.4***	2.62
OCT1/2 ^{-/-}	453 \pm 65.0***	3.00	1050 \pm 142***	3.77
MATE1 ^{-/-}	509 \pm 62.2***	3.36	894 \pm 92.0***	3.21
DDI with Verapamil (10 mg/kg)				
Wild-type	181 \pm 32.0	1.20	259 \pm 18.3	0.93
OCT1/2 ^{-/-}	148 \pm 32.0	0.98	328 \pm 52.0	1.18
MATE1 ^{-/-}	246 \pm 38.4	1.63	406 \pm 38.0	1.46

Data represent mean \pm SEM. *P < 0.05, **P < 0.01, ***P < 0.001 vs single agent dofetilide.

Abbreviations: C_{max}, peak plasma concentration; AUC_{last}, area under the plasma concentration-time curve to the last measurable concentration.

Table S4. Observed and predicted pharmacokinetic parameters of dofetilide with the presence and absence of cimetidine and ketoconazole.

	Mean C _{max} ± SD (CV%) (ng/mL)		Mean AUC _{0-last} ± SD (CV%) (ng×h/mL)		Mean CL ± SD (CV%) (L/h)	
	Predicted	Observed	Predicted	Observed	Predicted	Observed
Dofetilide (0.5 mg BID, p.o.)	2.48 ± 0.58 (23) PE = +32.66%	1.87 ± 0.65	18.0 ± 3.38 (19) PE = +2.91%	17.5 ± 3.2	28.8 ± 6.05 (21) PE = +30.8%	22.0
Dofetilide (0.5 mg i.v. infusion over 90 min)	4.28 ± 0.52 (12) PE = +38.06%	3.1 ± 0.50	20.4 ± 3.92 (19) PE = -3.92%	21.2 ± 4.5	25.4 ± 4.94 (19) PE = +40.48%	18.1
DDI with Cimetidine 400 mg BID (p.o.)						
Dofetilide (0.5 mg, p.o.)	2.84 ± 0.53 (19) PE = +25.7%	2.26	19.6 ± 3.66 (19) PE = +12.6%	17.4	26.4 ± 5.36 (20) PE = +28.2%	20.6
Dofetilide + Cimetidine	3.98 ± 0.70 (18) PE = +15.7%	3.44	32.4 ± 6.76 (21) PE = +17.9%	27.5	16.1 ± 3.79 (23) PE = +39.1%	11.6
% Change	40.1% ↑	52.2% ↑	65.5% ↑	58.0% ↑	38.9% ↓	43.7% ↓
DDI with Ketoconazole 400 mg QD (p.o.)						
Dofetilide (0.5 mg, p.o.)	2.87 ± 0.52 (18) PE = +27.0%	2.26	19.9 ± 3.69 (19) PE = +14.3%	17.4	26.0 ± 5.20 (20) PE = +26.3%	20.6
Dofetilide + Ketoconazole	3.70 ± 0.73 (20)	-	28.1 ± 6.17 (22)	-	18.7 ± 4.68 (25)	-
% Change	28.9% ↑	53.0% ↑	41.1% ↑	41.0% ↑	28.0% ↓	-

Abbreviations: p.o., per oral; BID, twice per day; QD, once a day, C_{max}, peak plasma concentration; AUC_{0-last}, area under the plasma concentration-time curve between time zero and the last measurable concentration; CL, clearance; PE, calculated prediction error (%) = [(predicted value - observed value)/observed value] × 100.

Table S5. Input parameters of dofetilide in PBPK model.

Input parameters	Description	Units	Value	Reference
1. Physicochemical and binding properties				
MW	Molecular weight	g/mol	441.57	PubChem
Log P	Octanol-water partition	-	2.1	
Compound type	Acid,base or neutral	-	Monoprotic base	
pK _a	Acid dissociation constant	-	7.89	UWDIDB
B/P profile	Blood to plasma ratio dependent on drug concentration	-	1.048	Simcyp predicted
f _u	Fraction unbound in plasma	-	0.36	UWDIDB
2. Absorption				
Absorption model	First order absorption model			
f _a	Fraction available from dosage form	-	1	Default
CV f _a	Coefficient of variation f _a	%	30	
k _a	Absorption rate constant	1/h	0.9	
CV k _a	Coefficient of variation k _a	%	30	Default
f _{u,gut}	Unbound fraction in enterocytes	-	0.039	Simcyp predicted
Q _{gut}	Nominal flow in gut model	L/h	4.068	Simcyp predicted
CV Q _{gut}	Coefficient of variation Q(gut)	%	30	Default
P _{eff,man}	Effective permeability in man	10 ⁻⁴ cm/s	12	Simcyp predicted
Permeability assay	Passive + active permeability		Physicochemical	
PSA	Polar surface area	Å ²	104.81	Drug central
HBD	Hydrogen bond donor		2	Drug central
3. Distribution				
Distribution model	Full PBPK model			
Prediction method			Method 2	
K _p scalar	Tissue to plasma partition coefficient		0.61	Optimized

Input parameters	Description	Units	Value	Reference
V_{ss}	Volume of distribution at steady-state	L/kg	2.8	Smith et al 1992 [16]
CV V_{ss}	Coefficient of variation V_{ss}	%	30	Default
4. Elimination				
Elimination model	Enzyme kinetics			
CYP, recombinant			CYP3A4	Smith et al 1992 [16]
CL_{int} (HLM)	<i>In vitro</i> clearance (human liver microsomes)	$\mu\text{L}/\text{min}/\text{mg}$ protein	0.25	Smith et al 1992 [16]
f_{umic}	Fraction unbound <i>in vitro</i>		0.81	
CL_r	Renal clearance in 20-30yr healthy male	L/h	18.06	Smith et al 1992 [16]
5. Transport				
Organ/Tissue			Kidney	
Transporter			SLC22A2 (OCT2)	
Location			Basolateral	
Function			EGD model	
J_{max}		$\text{pmol}/\text{min}/10^6$ cells	506.35	Measured
K_m	Michaelis-Menten constant	μM	347.2	Measured
System			User	
RAF/REF	Relative activity factor/relative expression factor		1	Optimized
CL_{PD} basal	Passive diffusion clearance	$\text{mL}/\text{min}/10^6$ cells	0	
Transporter			SLC47A1 (MATE1)	
Location			Apical	
Function			Efflux	
CL_{int}	Intrinsic clearance	$\mu\text{L}/\text{min}/10^6$ cells	22.2	Measured
System			User	
RAF/REF	Relative activity factor/relative expression factor		0.25	Optimized

Input parameters	Description	Units	Value	Reference
CL _{PD} apical	Passive diffusion clearance	mL/min/10 ⁶ cells	0	

Drug central, <https://drugcentral.org/drugcard/942>

UWDIDB, University of Washington Drug Interaction Database

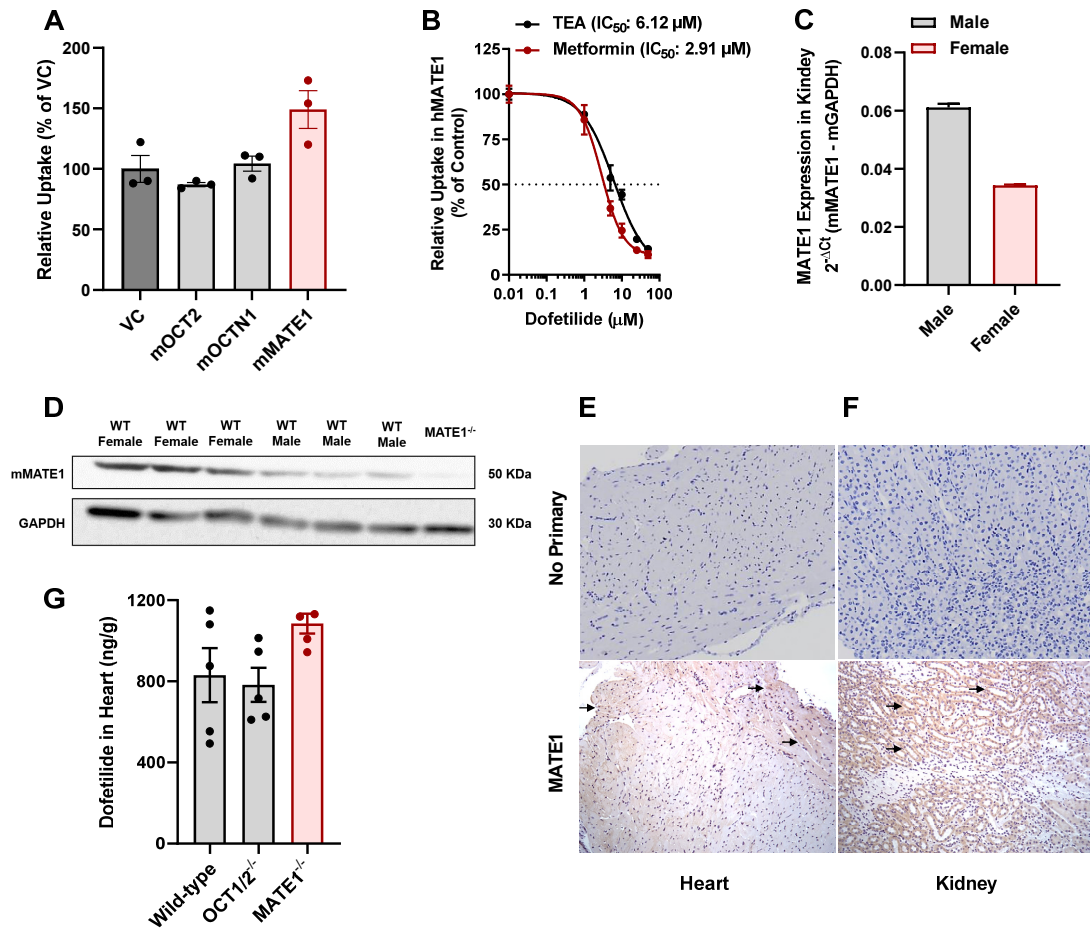


Figure S1. Influence of MATE1 on the transport of dofetilide. (A) Transport of [3 H] dofetilide (1 μ M) in cells overexpressing mouse (m) OCT2, OCTN1, or MATE1. Relative uptake is expressed as percentage change compared with empty vector controls (n = 3). (B) Uptake of [14 C] TEA (2 μ M) and [14 C] metformin (5 μ M) in HEK293 cells overexpressing human (h) MATE1 after preincubation with dofetilide at various concentrations (1-50 μ M) for 15 min, followed by the co-incubation with TEA and metformin for 15 min. Data represent the mean \pm SEM and are expressed as a percentage over vector control. (C) Gene expression of MATE1 in the kidney isolated from untreated male and female wild-type mice (n = 4 per group). (D) Protein expression of MATE1 in the heart isolated from untreated wild-type (WT) and MATE1^{-/-} mice (n=3). Immunohistochemical detection of MATE1 (bottom panels) in the wild-type mouse heart (E) and kidney (F). Tissue staining lacking the primary antibody was performed as a negative control for each tissue group (top panels). Representative images were taken at x40 magnification. (G) Concentration of dofetilide in whole heart tissue from male wild-type, OCT1/2-deficient, and MATE1-deficient mice 15 min after a single i.v. injection of dofetilide at a dose of 2.5 mg/kg (n = 4-5 per group).

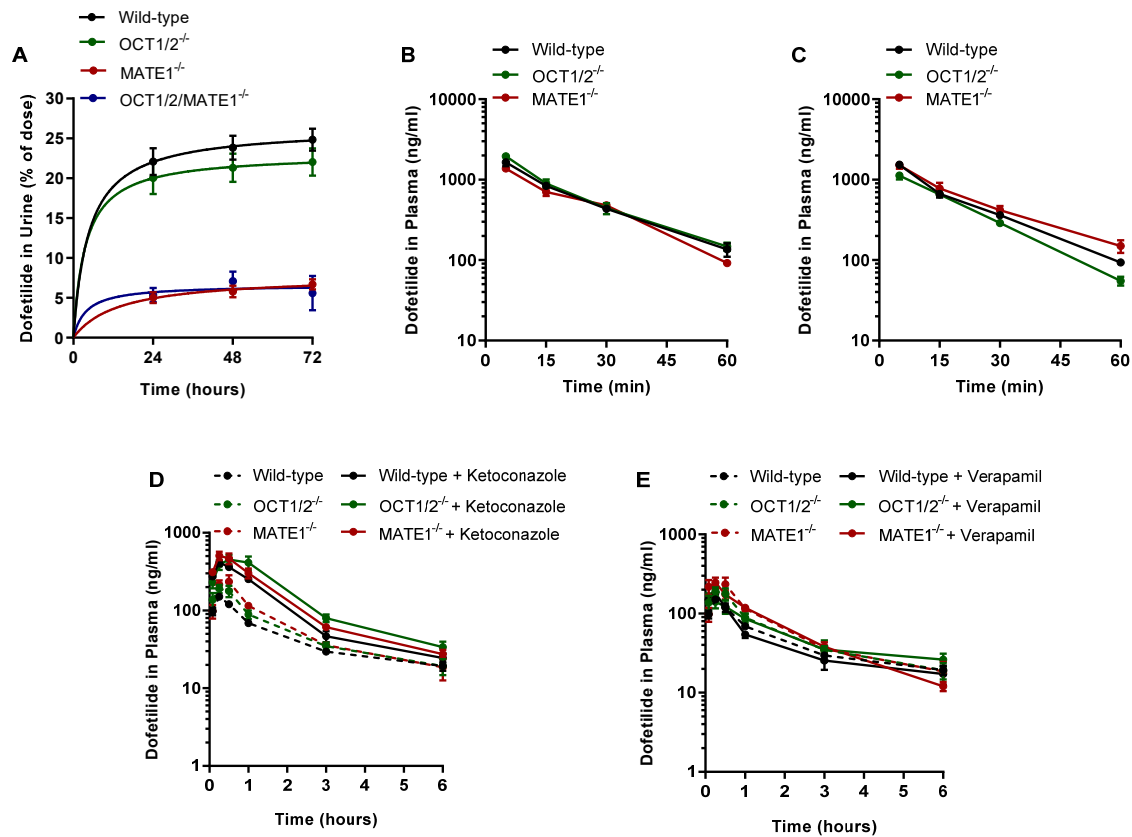


Figure S2. Influence of mouse genotype and ketoconazole or verapamil treatment on the pharmacokinetics of dofetilide. (A) Urinary excretion of dofetilide in male wild-type, OCT1/2-deficient, MATE1-deficient, and OCT1/2/MATE1-deficient mice (n = 5) following an i.v. dose of dofetilide (2.5 mg/kg). Plasma concentration time profiles of dofetilide in female (B) and male (C) wild-type, OCT1/2-deficient, and MATE1-deficient mice (n = 5 per group) receiving a single i.v. dose of dofetilide (2.5 mg/kg). Plasma concentration time profiles of dofetilide in wild-type, OCT1/2-deficient, and MATE1-deficient male mice (n = 5 to 38 per group) receiving an oral dose of dofetilide (5 mg/kg) 30 min after the administration of ketoconazole (50 mg/kg) (D) or verapamil (10 mg/kg) (E).

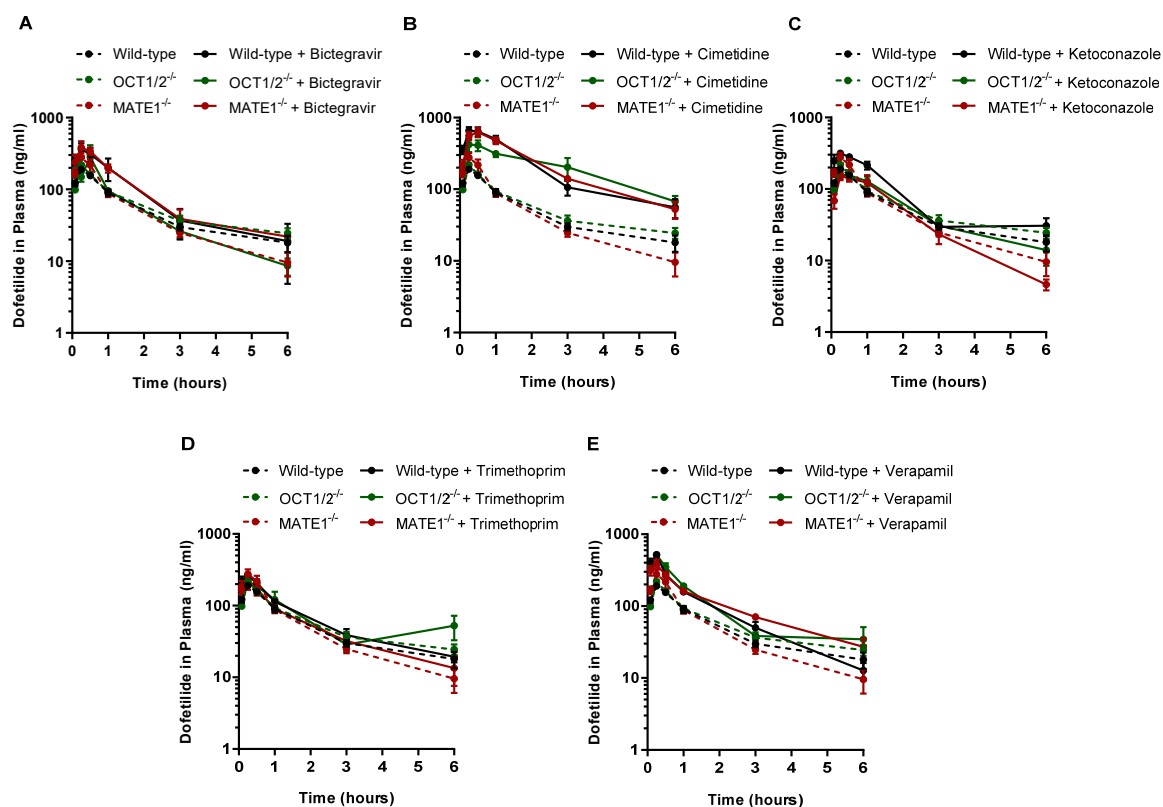


Figure S3. Drug-drug interactions between dofetilide and contraindicated drugs. Plasma concentration time profiles of dofetilide in female wild-type, OCT1/2-deficient, and MATE1-deficient mice (n = 5 to 29 per group) receiving a single oral dose of dofetilide (5 mg/kg) 30 min after the administration of bicitegravir (30 mg/kg) (A), cimetidine (100 mg/kg) (B), ketoconazole (50 mg/kg) (C), trimethoprim (100 mg/kg) (D), or verapamil (10 mg/kg) (E).

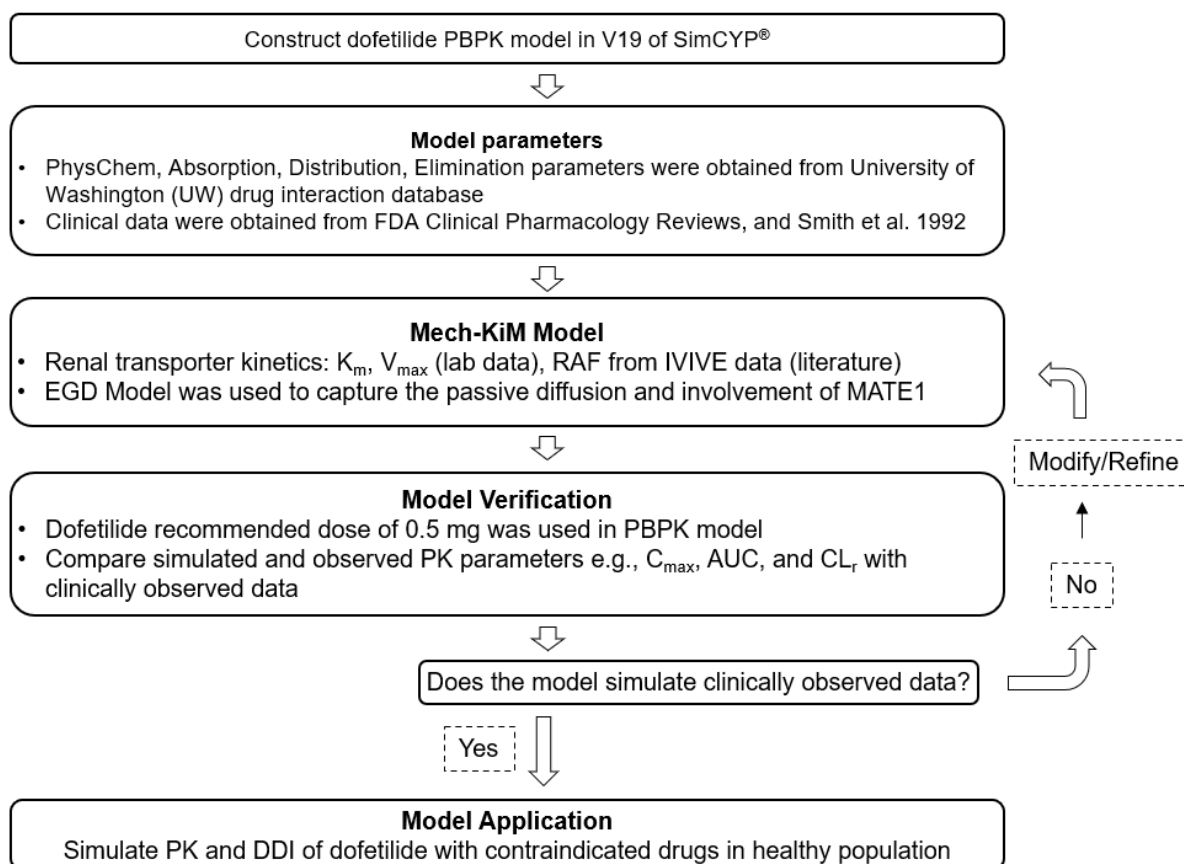


Figure S4. Workflow of the development of physiologically based pharmacokinetic (PBPK) model.

Abbreviations: AUC, area under the curve; C_{max} , peak plasma concentration; CL_r , clearance; FDA, U.S. Food and Drug Administration; IVIVE, *in vitro* to *in vivo* extrapolation; K_m , Michaelis-Menten constant; RAF, relative activity factor; PK, pharmacokinetics; V_{max} , maximum velocity.