

Table 3. Model strategies and final pharmacokinetic parameters of included studies.

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between- subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Meagher AK (2003) [44]	ADAPT II (IT2S)	2 CMTs with parallel FO and MM eliminations	V_c	$39.6 \times \text{TBW}/65$	22.7%	11% 0.028 mg/L	GOF	NR	NR
			V_p	$26.3 \times \text{TBW}/65$	41.8%				
			V_{ss}	$65.8 \times \text{TBW}/65$	23.4%				
			Q	$9.09 \times \text{TBW}/65$	14.9%				
			CL_r	$0.269 \times CL_{CR}^a \times \text{TBW}/65$	34.2%				
			CL_i^b	$43.5 \times \text{TBW}/65$	52.5%				
			K_m	1.46	68.1%				
			V_{max}	$53.3 \times \text{TBW}/65$	25.8%				
			T_{lag}	0.371	97.6%				
			K_a	5.73	1.20%				
			AUC	228	584%				
			CL_{tav}	$6.85 \times \text{TBW}/65$	50.3%				
Whitehouse T (2005) [45]	NONMEM (FO)	2 CMT ^c with FO elimination for all doses	CL	$0.0487 \times \text{TBW}$	48.1%	19.0% 2.34 mg/L	GOF	NR	AUC/MIC
			Q	7.48	/				
			V_c	$0.634 \times \text{TBW}$	22.4%				
			V_p	240	146%				
Abe S (2009) [7]	NONMEM (FOCE)	1 CMT with FO absorption and FO elimination	CL	$1.28 \times (\text{TBW}/69.5)^{1.91}$ + $0.0788 \times (110 - \text{AGE})$ (If age < 58) + 0.0788×52 (If age ≥ 58)	46.6%	8.14%	Bootstrap	NR	NR
			V	$47.0 \times (\text{TBW}/69.5)^{0.903}$	25.9%				
			K_a	0.583	181%				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between- subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Swoboda S (2010) [40]	NONMEM (FOCE-I)	2 CMT with FO elimination	CL	$[0.159 \times \text{TBW} + 3.5 \text{ (if dialysis)}]$ $\times 0.4 \text{ (if liver transplantation or resection)}$	51%	4.13% 0.285 mg/L	GOF	NR	NR
			V_c	$0.273 \times \text{TBW}$	21%				
			V_p	$0.271 \times \text{TBW}$	/				
			Q	$0.369 \times \text{TBW}$	/				
Sasaki T (2011) [35]	NONMEM (FOCE-I)	1 CMT with FO elimination	CL	$2.85 \times (\text{CL}_{\text{CR}}^{\text{a}}/60.9)^{0.618}$ $\times 0.472 \text{ (if cirrhosis)}$	35.2%	1.43 mg/L	GOF	NR	Design dosing regimen
			V	$33.6 \times (\text{TBW}/57.9)$	30.8%				
			K_a	0.583 fixed	/				
Tsuji Y (2013) [36]	NONMEM (FO)	1 CMT with FO elimination	CL	$0.00327 \times \text{TBW} \times \text{eGFR}^{0.428}$ $\times \text{HB}^{0.502}$ $\times 0.283 \text{ (if ALT} \geq 100 \text{ IU/L)}^{\text{d}}$	31.3%	21.7%	GOF, Bootstrap	NR	NR
			V	$1.31 \times \text{TBW}$	33.9%				
Matsumoto K (2014) [37]	NONMEM (FOCE)	1 CMT with a FO absorption and FO elimination	K_a	0.583	/	21.4%	GOF, Bootstrap	NR	NR
			CL	$0.0258 \times \text{CL}_{\text{CR}}^{\text{a}} + 2.03;$	30.5%				
			V	27.6	/				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between- subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Luque S (2014) [18]	NONMEM (FOCE-I)	3 CMT with FO elimination	CL	16.6	50.2%	Plasma:	GOF, Bootstrap	NR	AUC _{0-12h} , C _{min} , C _{max}
			V _c	43.2	15.3%	29.8%			
			V _p	58.0	8.2%	0.03 mg/L;			
			V _{CSF}	0.11	/	CSF:			
			Q	3.1	88.9%	36.6%			
Dong HY (2016) [17]	NONMEM (FOCE-I)	1 CMT with FO elimination	Q _{CSF}	0.05	/	0.03 mg/L	Bootstrap	NR	Design dosing regimen
			CL	$6.8 + 0.0134 \times (\text{AGE}-61)$	23.2%	3.6%			
Taubert M (2016) [41]	NONMEM (FOCE-I)	2 CMT with FO absorption and elimination	V	78.6	36.7%	0.227 mg/L	GOF, Bootstrap	NR	NR
			CL	$7.92 \times (\text{lactate}/1.91)^{-0.21}$ $\times (\text{fibrinogen}/13.0)^{0.04}$ $\times 1.82$ (if ARDS)	58%	33.9%			
			Q	65.59	/	0.07 mg/L			
			V _c	$15 \times (\text{TBW}/76)^{1.31}$ $\times 1.53$ (If peritonitis)	37%				
			V _p	26.55	/				
			K _a	1.72	/				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between- subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Minichmayr IK (2017) [11]	NONMEM (FOCE-I)	2 CMT with FO absorption and with concentration and time-dependent inhibition elimination	K _a	1.41 h ⁻¹	118%	PL: 15.7%	GOF, VPC, Bootstrap	NR	AUC
			CL	(7.67 for healthy volunteer, 11.2 for septic patients and 6.35 for diabetic patients) $\times (1 + 0.00835 \times (\text{CL}_{\text{CR}}^{\text{a}} - 80.0))$	40.1%	UF: 12.4%			
						μDA : 33.9%			
						μDM : 27.0%			
						rDA: 19.4%			
						rDM: 19.4%			
			V _c	$22.7 \times (\text{TBW}/69.5)$	36.6%				
			V _p	$19.9 \times (\text{TBW}/69.5)$	39.4%				
			Q	$57.9 \times (\text{TBW}/69.5)^{0.75}$	/				
			F _U	0.88	4.1%				
			TF _A	90%	21.9%				
			TF _M	86.2%	33.8%				
			K _{IC}	0.0017 FIXED	/				
			IC ₅₀	0.48	/				
			TF _{diabetic (%)} ^e	-34.9	/				
			RCLF [%]	51.3	6.45%				
			RR _A [%]	38.5	23.8%				
			RR _M [%]	53.9	18.2%				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between-subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Tsuji Y (2017) [38]	NONMEM (FOCE-I)	2 CMTs with FO absorption and elimination	CL	$(1.86 \times e^{-0.0205 \times (AGE-69)} + 1.44 \times CL_{CR}^{a,f} / 100) \times (TBW / 70)^{0.75}$	36.9%	total: 31.8% 0.251 mg/L;	Bootstrap, pcVPC	NR	NR
			V _c	$22.9 \times (TBW/70)$	142%	free:			
			V _p	$24.7 \times (TBW / 70)$	5%	31.9%			
			Q	$10.9 \times (TBW / 70)^{0.75}$	182%	0.034 mg/L			
			T _{abs}	3.61	/				
			F	0.922	/				
			F _U	0.823	/				
Wicha SG (2017) [22]	NONMEM (FOCE-I)	1 CMT with FO elimination	CL _{nr}	$4.41 \times (TBW/57.9)^{0.75} \times (LiMAX/221.5)^{0.388}$	33.6% (33.3%)	10.0% 0.1 mg/L	GOF, VPC	NR	NR
			CL _r	$0.919 \times [1 + 0.0208 \times (CL_{CR}^a - 45.6)]$	62.2% (56.4%)				
			CL _{dialysis}	1.26	/				
			V	$33.8 \times (TBW/57.9)$	/				
Ide T (2018) [24]	NONMEM (FOCE)	2 CMTs with FO elimination	CL _{preserved}	6.36	66.9%	5.8%	Bootstrap, VPC	NR	Design dosing regimen
			CL _{dysfunction}	2.06	40.1%				
			CL _{CRRT}	2.74	55.5%				
			Q	26.4	44.0%				
			V _c	19.6	56.8%				
			V _p	22.4	35.1%				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between-subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Allegra S (2018) [43]	NONMEM (NR)	1 CMT with FO elimination	CL	7.186 for <i>ABCB1</i> c.3435CT/TT $\times 1.969$ for <i>ABCB1</i> c.3435CC	60.8%	2.59% 1.90 mg/L	GOF, pcVPC	NR	NR
Strydom N (2019) [46]	NONMEM (FOCE)	1 CMT with FO absorption and elimination for plasma model ^g	V	40.520	33.0%	40.4%	VPC	NR	Design dosing regimen
			K _a	2.13	/				
			CL	3.77	57.0%				
Crass RL (2019) [4]	NONMEM (FOCE-I)	1 CMT with FO absorption and elimination	V	145	/	27.1% 1.43 mg/L	GOF, pcVPC	NR	Design dosing regimen
			K _a	1.40	/				
Xie FF (2019) [47]	NONMEM (FOCE-I)	2 CMT with FO elimination	CL	3.43 + 3.49 \times (BSA-1.89) + 1.77 \times (eGFR ^h /80) – 0.0242 \times (AGE-40) (If age > 40)	49.9% ⁱ	15.9%	GOF, VPC, NPDE	NR	Design dosing regimen
			V	42.9 \times e ^{0.901 \times (BSA-1.89)}	17.8% ⁱ				
			CL	7.8 \times (1 – 0.0331 \times (AGE - 60)) \times (TBW/70) ^{0.75}	66.9% ^j				
			V _c	14.3 \times (TBW/70)	43.5% ^j				
			Q	65.1 \times (TBW/70) ^{0.75}	/				
			V _p	23.8 \times (TBW/70)	/				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between-subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Thibault C (2018) [25]	NONMEM (NR)	1 CMT with FO elimination	CL	$0.181 \times (\text{TBW}/1.4)^{0.405} \times (\text{PNA}/0.07)^{0.831}$	38.3%	1.13 mg/L	Bootstrap, GOF, VPC	NR	Design dosing regimen
			V	$1.17 \times (\text{TBW}/1.4)^{0.801}$	100%				
Li SC (2019) [30]	NONMEM (FOCE-ELS)	1 CMT with FO elimination	CL	$1.31 \times (\ln \text{TBW}/2.40)^{0.83} \times (\ln \text{eGFR}^k/4.89)^{0.60}$	39.1%	16.5% 0.02 mg/L	GOF, NPDE, VPC	NR	Design dosing regimen
			V	$4.24 \times (\ln \text{WT}/2.40)^{0.86}$	28.1%				
Garcia-Prats AJ (2019) [48]	NONMEM (FOCE-I)	1 CMT with FO absorption and elimination	K _a	0.77	/	25%	GOF, VPC	NR	Design dosing regimen
			CL/F	$4.73 \times (\text{TBW}/70)^{0.75}$	37%	0.78 mg/L			
			V/F	$54.8 \times (\text{TBW}/70)$	32%				
Ogami C (2019) [29]	NONMEM (FOCE-I)	1 CMT with FO absorption	CL	$5.82 \times (\text{WT}/70)^{0.75} \times [\text{PNA}^{46.0}/(2.06^{46.0} + \text{PNA}^{46.0})]$	52.3%	28.9% and 31.3% for total and unbound concentration respectively	Bootstrap	NR	NR
			V	$41.3 \times (\text{WT}/70)$	121%				
			Tab _s	3.61 h	/				
			F	0.922	/				
			TM50	2.06	/				
			Hill	46.0	/				
			FU	0.811	/				
Wang DD (2019) [31]	NONMEM (NR)	1 CMT with FO elimination	CL	$11.8 \times (\text{PLT}/200)^{0.261}$	29.9%	102.0%	Bootstrap; VPC	NR	NR
			V	209	29.9%				
Alghamdi (2020) [50]	Monolix (SAEM)	1 CMT with FO absorption and elimination	CL	$6.32 \times (\text{CL}_{\text{CR}}^a/97.0)^{0.449}$	36.8%	25.9%	GOF; Bootstrap; VPC	NR	Design dosing regimen
			V	$40.6 \times (\text{WT}/63.0)$	10.9%				
			K _a	1.65	70.7%				
			T _{lag}	0.341	88.2%				

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between- subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Soraluce A (2020) [51]	NONMEM (FOCE-I)	2 CMT with FO elimination	CL	$2.62 + 4.35 \times (\text{CL}_{\text{CR}}^{\text{a}}/44)$ + $\text{Sc} \times \text{Qef}$)	61.5%	15.9%	GOF, Bootstrap, VPC	11	Design dosing regimen
			V ₁	16.2	65.9%	0.266 mg·L ⁻¹			
			Q	71.7	/				
			V ₂	29.0	/				
Zhang SH (2020) [32]	Phoenix NLME (FOCE-ELS)	1 CMT with FO elimination	CL	$2.68 \times (\text{PTA}/48.07)^{0.84}$ $\times (\text{CL}_{\text{CR}}^{\text{a}}/99.3)^{0.36}$	29.82%	18.5%	GOF, Bootstrap, pcVPC	NR	Design dosing regimen
			V	58.34	39.33%				
Ehmann L (2020) [42]	NONMEM (FO)	2 CMT with parallel linear and nonlinear MM clearance	CL _u ¹	3.32	66.7%	C _{p_tot} 4.76%	GOF, Bootstrap, VPC	NR	Design dosing regimen
			V _{max,u}	45.9	/	C _{p_u} 4.56%			
			K _{m,u}	2.93	74.4%	C _{ud} 13.3%			
			V _{1,u}	$17.0 \times (\text{LBW}/51.9)$	42.1%	C _{RD} 1.9%			
			Q _u	$62.4 \times (\text{LBW}/51.9)^{0.75}$	46.8%				
			V _{2,u}	$33.4 \times (\text{LBW}/51.9)$	16.7%				
			f _u	85.6	/				
			TF _{OBE,u}	54.1	TF _u 14.8%				
			TF _{NOBE,u}	69.0					
			ANAE-TF	-13.6	82.2%				
			MAPCL _{Tot,u} %	0.805	/				
			RR _{OBE}	37.5	RR _{intercatheter}				
			RR _{NOBE}	57.5	26.1%				
		RR _{intracatheter}	27.2%						

Table 3 (continued)

Study (publication year)	Software / Algorithm	Structural Model	Fixed effect parameters		Between-subject variability	Residual unexplained variability	Internal validation	External validation (N=number of subjects)	Model application
Wang XP (2020) [33]	NONMEM (FOCE-I)	1 CMT with FO elimination	CL V	$5.6 \times (\text{CL}_{\text{CR}}^{\text{a}}/61)^{0.386}$ 43.4	63.9% 17.6%	36.2% 0.055 mg·L ⁻¹	GOF, VPC, Bootstrap, NPDE	NR	Design dosing regimen
Blackman AL (2021) [49]	Pumas (FOCE)	1 CMT with FO elimination	CL V	6.7×0.289 (if cirrhosis) $\times (\text{TBW}/140)^{1.12}$ $64.3 \times (\text{TBW}/140)^{1.67}$	25.0% 21.0%	14.0%	GOF, QPC, VPC	NR	Design dosing regimen
Yang M (2021) [34]	Phoenix NLME (FOCE-I)	2 CMT with FO elimination	CL Q Vc Vp	$2.34 \times (\text{WT}/15)^{0.8} \times (\text{AST}/45.9)^{(-0.16)}$ $7.14 \times (\text{WT}/15)^{1.09}$ 5.22 28.79	52.51% 53.45% 55.78% /	29.0%	GOF, VPC, Bootstrap	NR	Design dosing regimen

Abbreviations: ALT, alanine amino transferase; AST, aspartate aminotransferase; ARDS, acute respiratory distress syndrome; BSA, body surface area; CIR, liver cirrhosis; CL_{CR}, creatinine clearance; CMT, compartment; CP_{tot}, total plasma concentrations; CP_u, unbound plasma concentrations; CRD, retrodialysate concentration; C_uD, microdialysate concentration; CSF, cerebrospinal fluid; DURA, duration of administration; eGFR, estimated glomerular filtration rate; FO, first order; FOCE, FO conditional estimation; FOCE-ELS, FO estimation-extended least squares method; FOCE-I, FO conditional estimation method with the interaction option; HB, hemoglobin; IT2S, iterative two-stage analysis; LBW, lean body weight, LiMax, the maximal liver function capacity; MAP, mean arterial pressure; MM, Michaelis-Menten; NR, not reported; OBE, obesity; PL, plasma concentration; PLT, platelets; PTA, prothrombin time activity; PNA, postnatal age (in years); Qef: effluent flow; Sc: sieving coefficient; TBW, total body weight

PK parameters: ANAE-TF, anaesthesia effect on TFu; AUC_{0-12 h}, area under the concentration-time curve during 24 hours; CL, total clearance; CL_{CRRT}, clearance for patients on low-dose continuous renal replacement therapy. CL_{dysfunction}, clearance for patients with renal dysfunction; CL_{nr}, non-renal clearance; CL_{preserved}, clearance for patients with preserved renal function; CL_r, renal clearance; CL_{tagv}, the calculated average total clearance of linezolid over the first 7 days of treatment; CL_{Tot,u}, Total clearance of unbound linezolid; CL_u, CL of unbound linezolid; F_u, fraction unbound; IC₅₀, concentration in inhibition compartment yielding 50 % of CL inhibition; K_a, the absorption rate constant; K_{cp}, rate constant for drug distribution from the central to peripheral compartment; K_{el}, linear elimination rate constant. K_{IC}, rate constant for the transfer into inhibition compartment; K_{pc}, rate constant for drug distribution from the peripheral to central compartment; KPL, rate of drug moving from plasma to lesion (KPL) ; K_m,

Michaelis-Menten constant; MAPCLTot,u, effect of MAP on CLTot,u; UF, unbound concentration; Q, intercompartmental (central-peripheral) clearance; Qu, intercompartmental distribution of unbound linezolid; RCLF, remaining fraction of CL at maximum CL inhibition; rDA/rDM, mg/L, retrodialysate of subcutaneous adipose/muscle tissue; RRA, RRM, relative recovery of catheters located in interstitial space fluid of subcutaneous adipose/muscle tissue; RROBE, RRNOBE, relative recovery for obese and non-obese patients; T_{abs}, absorption half-life. TFA, TFM, factors transforming plasma to interstitial space fluid of subcutaneous adipose/muscle tissue concentrations; T_{lag}, the lag time before onset of absorption; TFOBE,u, TFNOBE,u, tissue factor of unbound linezolid for obese and non-obese patients; $\mu D_A / \mu D_M$, microdialysate concentration of subcutaneous adipose/muscle tissue; V, volume of distribution; V_c, volume of distribution of the central compartment; V_{max}, maximum velocity of capacity-limited clearance; V_p, volume of distribution of the central compartment; V_{ss}, volume of distribution at steady state; V1,u, V2,u, volume of distribution parameters of central and peripheral CMTs of unbound linezolid; Vmax,u, maximum elimination rate of unbound linezolid;

^aestimated with Cockcroft-Gault formula.

^bintrinsic clearance, $CL_i \times K_m = V_{max}$

^c 2 CMT was fitted to the multiple-dose data, 1CMT was fitted to single-dose data, but PK parameter in 1CMT was not reported;

^d eGFR was calculated with revised equation for Japanese. $eGFR \text{ (mL / min / 1.73 m}^2\text{)} = 194 \times SCR^{-1.094} \times AGE^{-0.287} \times 0.739$ (if female).

^e Difference in percent compared with healthy volunteers.

^f CL_{CR}/100 was assumed to be 0.5 for the 4 youngest patients (1, 5, 8 and 13 years old).

^g 10 CMTs with FO absorption and elimination for tissue model was developed, and the rates of drug moving from plasma to lesion were shown as follows: lung 0.152 h⁻¹; necrotic nodule 0.229 h⁻¹; caseum closed nodule 0.006 h⁻¹; caseous fibrotic nodule 0.275 h⁻¹; caseum from cavity 0.427 h⁻¹; cavity wall 0.305 h⁻¹; fibrotic tissue 1.29 h⁻¹; small cellular nodule 0.001 h⁻¹; fungal ball NA.

^h calculated using the Chronic Kidney Disease Epidemiology Collaboration equation.

ⁱ The covariance between the variances of CL and V was 0.0603.

^j The covariance between the variances of CL and V_c was 0.23.

^k calculated by the modified Schwartz formula.

^l Clearance was described by the parallel linear (CL_u) and nonlinear concentration-dependent Michaelis-Menten clearance (V_{max} and K_m).