

Supplementary Information to “Use of a mechanistic physiologically based kinetic model and quantitative in vitro to in vivo extrapolation to evaluate the tolerable daily intake of zearalenone”

Chrysanthi Pachoulide^{1,*}, Joost Westerhout ², Hans Bouwmeester¹ and Nynke I. Kramer¹

¹Division of Toxicology, Wageningen University and Research, The Netherlands

²The Netherlands and 2Center for Prevention, Lifestyle and Health, National Institute for Public Health and the Environment (RIVM), The Netherlands

Table S1: Current knowledge of the toxicokinetic processes determining the fate of ZEN and its most prominent metabolites in humans and rats, based on in vitro and in vivo studies.

Method	Results	Reference
Absorption		
Intestinal perfusion of an isotonic solution of ZEN at 4 µg/ml; in Wistar rats (6 males).	<u>ZEN:</u> • ka: 9.27(0.69) /h • Intestinal absorption fits first order kinetics • Complete intestinal absorption via passive diffusion	(Ramos et al., 1996)
Exposure to 10 µM ZEN on the mucosal side of the everted small (excluding duodenum) and large intestine of Sprague-Dawley rats (6 males (m); 6 females (f); 6 pregnant females (pf))	<u>ZEN:</u> • Disappearance rate from the mucosal side: (m) 200, (f) 100, (pf) 70 nmol /10cm intestine /h • The rate decreased in the distal small intestine of males and pregnant females <u>α-ZEL:</u> Not detected; completely metabolised to α-ZEL GlcA	(Ieko et al., 2020)
Exposure to 10 µM and 200 µM ZEN on the apical (AP-BL) and basolateral side (BL-AP) of a Caco-2 cell monolayer for 8h.	<u>ZEN:</u> • Papp AP-BL: (10 µM ZEN) 368(43), (200 µM ZEN) 2749(228) pmol/cm²/h • Papp BL-AP: (10 µM ZEN) 137(37), (200 µM ZEN) 2020(184) pmol/cm²/h • Intestinal absorption via passive diffusion • Apparent permeability decreased after 4h of exposure; due to extensive metabolism	(Videmann et al., 2008)
Exposure to 20 µM ZEN or α-ZEL on the apical side (AP-BL) of a Caco-2 cell monolayer for 1h.	<u>ZEN:</u> • Papp AP-BL: 10.4(4.7) × 10⁻⁶ cm/s • α-ZEL: • Papp AP-BL: 5.4(0.5) × 10⁻⁶ cm/s	(Pfeiffer et al., 2011)
Bioavailability		
Single IV bolus and PO administration of 1-8 mg/kg ZEN in intact and bile-duct cannulated male Sprague-Dawley rats (n=16). TK parameters were determined with non-compartmental analysis.	<u>ZEN:</u> • AUC, IV bolus: 181(30), 335(107), 745(386) and 1696(481) ng h/ml at 1, 2, 4 and 8mg/kg ZEN • AUC, PO: 46.1(16.5) (intact), 19.3(0.3) (bile duct-cannulated) ng h/ml • Fa: (intact) 2.7, (bile duct cannulated) 1.1 %	(Shin et al., 2009)

Metabolism

ZEN reduction

Exposure to 40 μ M ZEN on Caco-2 cells for 3h.	22 and 4 % reduced to α -ZEL and β -ZEL respectively	(Pfeiffer et al., 2011)
Exposure to 10 - 200 μ M ZEN on the apical (AP-BL) side of a Caco-2 cell monolayer for 8h.	• Formation/secretion of α-ZEL followed Michaelis-Menten kinetics and was inhibited by 3α-HSD inhibitors • At low ZEN concentrations (<25 μM); secretion was higher in the AP-BL direction; at higher ZEN concentrations; secretion was higher in the BL-AP direction • α-ZEL was the main metabolite in the BL side and cells • Formation/secretion of β-ZEL was higher in the BL-AP direction; followed Michaelis-Menten kinetics and was inhibited by 3β-HSD inhibitors. • β-ZEL was mostly detected in the AP side.	(Ieko et al., 2020)
Incubation of rat S9 (post-mitochondrial) liver fractions with 10-2000 μ M ZEN for 30 minutes.	<u>α-ZEL formation rate:</u> • liver microsomes: 0.9×10^{-6} L/min/mg protein • liver S9: 0.05×10^{-6} L/min/mg protein <u>β-ZEL formation rate:</u> • liver microsomes: 49.5×10^{-6} L/min/mg protein • liver S9: 3.43×10^{-6} L/min/mg protein	(Malekinejad et al., 2006)
Incubation of human microsomal or S9 liver fractions with 1-500 μ M ZEN; for 10 and 5 minutes respectively.	<u>α-ZEL formation rate:</u> • liver microsomes: 10.7×10^{-6} L/min/mg protein • liver S9: 38.7×10^{-6} L/min/mg protein	(Mendez-Catala et al., 2020)
Incubation of human and rat feces (5%; pH 7.4) with 1-250 μ M ZEN for 5h.	<u>α-ZEL formation rate:</u> • Human feces: 6.6×10^{-9} L/min/mg feces • Rat feces: 3.5×10^{-9} L/min/mg feces	

Glucuronidation

Exposure to 40 μ M ZEN on Caco-2 cells for 3h.	• 11 % of ZEN metabolised to ZEN-GlcA • 14.8 % of α-ZEL metabolised to α-ZEL GlcA • 4.9 % of β-ZEL metabolised to β-ZEL GlcA	(Pfeiffer et al., 2011)
Exposure to 10 - 200 μ M ZEN on the apical (AP-BL) side of a Caco-2 cell monolayer for 8h.	• α-ZEL GlcA was formed and secreted at a higher rate than ZEN GlcA • There was no formation of β-ZEL GlcA. • GlcA secretion was saturated and inhibited at high ZEN concentrations.	(Videmann et al., 2008)

Exposure to 40 µM ZEN on Caco-2 cells for 3h.	GlcAs were mostly excreted in the AP side; undetectable in the cells.	
Incubation of rat S9 (post-mitochondrial) liver fractions with 10-2000 µM ZEN for 30 minutes.	14-O-GlcAs were the 97, 89 and 67 % of the total GlcAs formed for ZEN, α-ZEL and β-ZEL respectively.	(Pfeiffer et al., 2011)
Incubation of human and rat S9 fractions with 0.3-150 µM ZEN or α-ZEL; for 7(liver) and 20(intestine) minutes respectively.	The % of Glucuronidation decreased from 89(2.7) to 12(5.0) with an increasing concentration of ZEN (10 to 250 µM) <u>ZEN-GlcA formation rate:</u> Human liver S9: 1.45×10^{-3} L/min/mg protein Human intestinal S9: 0.42×10^{-3} L/min/mg protein Rat liver S9: 1.04×10^{-3} L/min/mg protein - α-ZEL-GlcA formation rate: Human liver S9: 1.24×10^{-3} L/min/mg protein Rat liver S9: 1.24×10^{-3} L/min/mg protein	(Malekinejad et al., 2006) (Mendez-Catala et al., 2021)
Exposure to 10 µM ZEN on the mucosal side of the everted small (excluding duodenum) and large intestine of Sprague-Dawley rats (6 males (m); 6 females (f); 6 pregnant females (pf))	<u>ZEN-GlcA formation rate:</u> Jejunum: (m) 70(10), (f) 50(5), (pf) 80(5) nmol /10cm intestine /h P. ileum: (m) 60(8), (f) 45(5), (pf) 79(12) nmol /10cm intestine /h D. ileum: (m) 52(12), (f) 67(2), (pf) 50(10) nmol /10cm intestine /h Colon: (m) 45(12), (f) 24(6), (pf) 33(7) nmol /10cm intestine /h	(Ieko et al., 2020)
Incubation of hepatic and intestinal rat and human microsomes and individual human UGTs with 50-100 µM ZEN; for up to 1h.	<u>ZEN- and α-ZEL- GlcA formation rate:</u> Human liver: 6(1) nmol/mg protein for both aglycones Human intestine: 5(1) and 4(0) nmol/mg protein for ZEN and α-ZEL respectively Rat liver: 10(1) and 9(1) nmol/mg protein for ZEN and α-ZEL respectively 14-O-GlcAs were 92; 90 and 87 % of the total GlcAs formed in human liver; human intestinal and rat liver microsomes. Amongst UGT isoforms, UGT1A1, 1A3 and 1A8 showed the highest metabolic activities towards both ZEN and α-ZEL	(Pfeiffer et al., 2010)

Sulfation

Exposure to 40 µM ZEN on Caco-2 cells for 3h.	5.2 % of ZEN metabolised to ZEN sulfate 16.0 % of α-ZEL metabolised to α-ZEL sulfate 3.1 % of β-ZEL metabolised to β-ZEL sulfate	(Pfeiffer et al., 2011)
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Distribution

IV infusion of 2.25 mg ZEN/h/kg for 6h; until steady state plasma concentration; 4 male Sprague-Swaley rats.	<u>ZEN:</u> (Shin et al., 2009) • Tissue-Plasma partition coefficients: small intestine 26.26, kidney 3.97, liver 2.91, adipose 2.28, lung 1.31, heart 0.88, stomach 0.88, brain 0.75, spleen 0.66, testis 0.37, muscle 0.37 • Blood: Plasma ratio: 0.75
Elimination	
Single IV bolus and PO administration of 1-8 mg/kg ZEN in intact and bile-duct cannulated male Sprague-Swaley rats (n=16). TK parameters were determined with non-compartmental analysis.	<u>ZEN:</u> (Shin et al., 2009) • t_{1/2}, IV bolus: 0.6(0.3), 1.9(0.9), 1.8(0.7) and 2.8(1.1) h at 1, 2, 4 and 8mg/kg ZEN • t_{1/2}, PO: 16.8(8.4) (intact), 7.0(5.7) (bile duct-cannulated) h • CL systemic, IV bolus: 5.6(1.1), 6.5(2.3), 6.6(3.2) and 5.0(1.4) L/h/kg at 1, 2, 4 and 8mg/kg ZEN • CL apparent, PO: (intact) 189.9(55.9), (bile duct-cannulated) 414.5(6.3) L/h/kg
Single PO administration of 8 mg/kg α -ZEL in female Wistar rats and of 1.23 mg/kg α -ZEL in human healthy volunteers (n=6)	<u>α-ZEL:</u> (Migdalof et al., 1983) • Urinary excretion, rat: 7.5(2.5) % of total dose • Fecal excretion, rat: 73.4(11.2) % of total dose • Fecal excretion; human: 23.4(10.0) % of total dose • 80 % of the blood and urine concentration in conjugated form
Single PO administration of 1 (low) or 100 (high) mg/kg ZEN in female Wistar rats (n=20).	<u>ZEN:</u> (Fitzpatrick et al., 1988) • Urinary excretion, unconjugated: 16.2(2) / 11.5(1.9) (low/high dose) % of total dose • Urinary excretion, conjugated: 2.3(0.7) / 1.9(0.9) (low/high) % of total dose • Fecal excretion: 40.3(4.3) / 54.3(3.6) (low/high) % of total dose <u>α-ZEL:</u> • Urinary excretion, unconjugated: 1.4(0.3) / 1.1(0.2) (low/high dose) % of total ZEN dose • Urinary excretion, conjugated: 0.4(0.1) / 0.2(0.1) (low/high dose) % of total ZEN dose • Fecal excretion: 11.5(1.3) / 9.8(1.1) (low/high dose) % dose

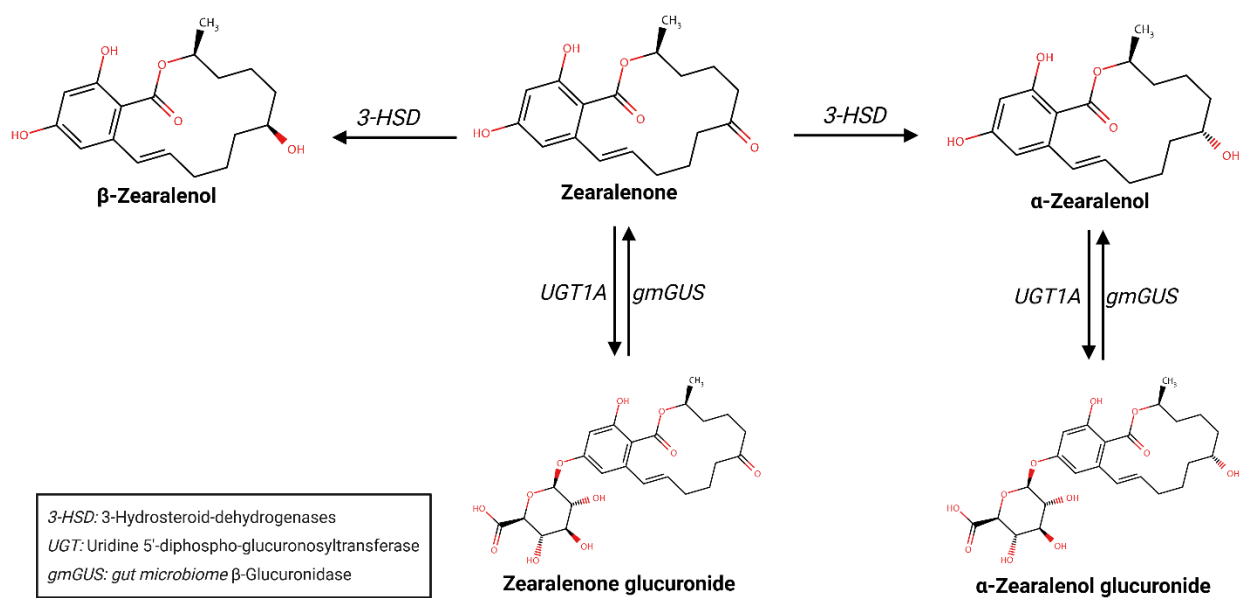


Figure S1: The key pathways of metabolism of Zearalenone in humans and rats, included in this study

Link to github repository: <https://doi.org/10.5281/zenodo.20812373>

PBK model equations

ZEN and α ZEL Aglycones (Agl)

All compartments of the ZEN and α ZEL sub-models are the same except the small intestine and liver, where α ZEL does not undergo reductive metabolism. As α ZEL is only formed in the jejunum, the respective model does not include the duodenum.

Oral dose:

$$\frac{dA_{OD_Agl}}{dt} = -k_{GER} A_{OD_Agl}$$

Small intestinal lumen:

$$\text{Duodenum lumen ZEN: } \frac{dA_{DuoL_ZEN}}{dt} = k_{GER}A_{OD_ZEN} - k_{duo}A_{DuoL_ZEN} - CL_{duo_ZEN}C_{DuoL_ZEN} - ka_{lymph}A_{DuoL_ZEN}$$

$$\text{Jejunum lumen aglycones: } \frac{dA_{JejL_Agl}}{dt} = k_{duo}A_{DuoL_Agl} - k_{jej}A_{JejL_Agl} - CL_{jej_Agl}C_{JejL_Agl} - ka_{lymph}A_{JejL_Agl}$$

$$\text{Ileum lumen aglycones: } \frac{dA_{IleL_Agl}}{dt} = k_{jej}A_{JejL_Agl} - k_{ile}A_{IleL_Agl} - CL_{ile_Agl}C_{IleL_Agl} - ka_{lymph}A_{IleL_Agl} + Vmax_{Hydro_IleL}C_{IleL_GlcA}/Km_{Hydro} + C_{IleL_GlcA}$$

Small intestine:

$$\text{Duodenum ZEN: } \frac{dA_{Duo_ZEN}}{dt} = CL_{duo_ZEN}C_{DuoL_ZEN} + Q_{Duo}(C_{B_ZEN} - CV_{Duo_ZEN})$$

Jejunum ZEN:

$$\begin{aligned} \frac{dA_{Jej_ZEN}}{dt} = & CL_{jej_ZEN}C_{Jej_ZEN} + Q_{Jej}(C_{B_ZEN} - CV_{Jej_ZEN}) - \\ & Vmax_{SI_ZENGlCA}C_{Jej_ZEN}fut_{ZEN}/Km_{SI_ZENGlCA} + C_{Jej_ZEN}fut_{ZEN} - \\ & Vmax_{SI_aZEL}C_{Jej_ZEN}fut_{ZEN}/Km_{SI_aZEL} + C_{Jej_ZEN}fut_{ZEN} - \\ & Vmax_{SI_bZEL}C_{Jej_ZEN}fut_{ZEN}/Km_{SI_bZEL} + C_{Jej_ZEN}fut_{ZEN} \end{aligned}$$

Jejunum α -ZEL:

$$\begin{aligned} \frac{dA_{Jej_aZEL}}{dt} = & C_{Jej_aZEL} + Q_{Jej}(C_{B_aZEL} - CV_{Jej_aZEL}) + \\ & Vmax_{SI_aZEL}C_{Jej_ZEN}fut_{ZEN}/Km_{SI_aZEL} + C_{Jej_ZEN}fut_{ZEN} - \\ & Vmax_{SI_aZELGlCA}C_{Jej_aZEL}fut_{aZEL}/Km_{SI_aZELGlCA} + C_{Jej_aZEL}fut_{aZEL} \end{aligned}$$

Ileum aglycones:

$$\frac{dA_{Ile_ZEN}}{dt} = CL_{ile_ZEN}C_{IleL_ZEN} + Q_{Ile}(C_{B_ZEN} - CV_{Ile_ZEN})$$

Liver ZEN:

$$\begin{aligned} \frac{dA_{L_ZEN}}{dt} = & Q_{Duo}CV_{Duo_ZEN} + Q_{Jej}CV_{Jej_ZEN} + Q_{Ile}CV_{Ile_ZEN} - (Q_{SI} + Q_B)CV_{L_ZEN} - \\ & Vmax_{L_ZENGlCA}C_{L_ZEN}fut_{ZEN}/Km_{L_ZENGlCA} + C_{L_ZEN}fut_{ZEN} - \\ & Vmax_{L_aZEL}C_{L_ZEN}fut_{ZEN}/Km_{L_aZEL} + C_{L_ZEN}fut_{ZEN} - \\ & Vmax_{L_bZEL}C_{L_ZEN}fut_{ZEN}/Km_{L_bZEL} + C_{L_ZEN}fut_{ZEN} \end{aligned}$$

Liver α ZEL:

$$\begin{aligned} \frac{dA_{L_aZEL}}{dt} = & Q_{Duo}CV_{Duo_aZEL} + Q_{Jej}CV_{Jej_aZEL} + Q_{Ile}CV_{Ile_aZEL} - (Q_{SI} + Q_B)CV_{L_aZEL} - \\ & Vmax_{L_aZELGlCA}C_{L_aZEL}fut_{aZEL}/Km_{L_aZELGlCA} + C_{L_aZEL}fut_{aZEL} \end{aligned}$$

Rapidly Perfused Tissue: $\frac{dA_{R_Agl}}{dt} = Q_R(C_{B_Agl} - CV_{R_Agl})$

Slowly Perfused Tissue: $\frac{dA_{S_ZEN}}{dt} = Q_S(C_{B_Agl} - CV_{S_Agl})$

Fat: $\frac{dA_{F_Agl}}{dt} = Q_F(C_{B_Agl} - CV_{F_Agl})$

Blood:

$$\begin{aligned} \frac{dA_{B_Agl}}{dt} = & ka_{Lymph}(A_{DuoL_Agl} + A_{JeJL_Agl} + A_{IleL_Agl}) + CV_{L_Agl}(Q_{SI} + Q_L) + CV_{F_Agl}Q_F \\ & + CV_{R_Agl}Q_R + CV_{S_Agl}Q_S - C_{B_Agl}(Q_{SI} + Q_L + Q_R + Q_S + Q_F) \\ & - GFR C_{B_Agl} fup_{Agl} \end{aligned}$$

Urine: $\frac{dA_{UR_Agl}}{dt} = GFR C_{B_Agl} fup_{Agl}$

Feces: $\frac{dA_{FE_Agl}}{dt} = k_{ile} A_{IleL_Agl}$

β -Zearalenol

$$\begin{aligned} \frac{dA_{\beta ZEL}}{dt} = & Vmax_{SI_BZEL} C_{JeJ_ZEN} ZEN_{fut} / Km_{SI_BZEL} + C_{JeJ_ZEN} ZEN_{fut} \\ & + Vmax_{L_BZEL} C_{L_ZEN} ZEN_{fut} / Km_{L_BZEL} + C_{L_ZEN} ZEN_{fut} \end{aligned}$$

PBK equations for ZEN and α ZEL glucuronides (GlcAs)

Small intestinal lumen:

$$\frac{dA_{Duo_GlcA}}{dt} = CL_{Bile} C_{Bile_GlcA} - k_{duo} A_{DuoL_GlcA}$$

$$\begin{aligned} \frac{dA_{JeJL_GlcA}}{dt} = & k_{duo} A_{DuoL_GlcA} - k_{JeJ} A_{JeJL_GlcA} \\ & + Vmax_{MRP2_IN} C_{JeJ_IC_GlcA} fut_{GlcA} / Km_{MRP2} + C_{JeJ_IC_GlcA} fut_{GlcA} \\ & + Vmax_{BCRP_IN} C_{JeJ_IC_GlcA} fut_{GlcA} / Km_{BCRP} + C_{JeJ_IC_GlcA} fut_{GlcA} \end{aligned}$$

$$\frac{dA_{IleL_GlcA}}{dt} = k_{JeJ} A_{JeJL_GlcA} - k_{ile} A_{IleL_GlcA} - Vmax_{Hydro_IleL} C_{IleL_GlcA} / Km_{Hydro} + C_{IleL_GlcA}$$

Small intestine:

$$\begin{aligned} \frac{dA_{JeJ_IC_GlcA}}{dt} = & Vmax_{SI_GlcA} C_{JeJ_Agl} fut_{Agl} / Km_{SI_GlcA} + C_{JeJ_Agl} fut_{Agl} \\ & - Vmax_{MRP2_IN} C_{JeJ_IC_GlcA} fut_{GlcA} / Km_{MRP2} + C_{JeJ_IC_GlcA} fut_{GlcA} \\ & - Vmax_{BCRP_IN} C_{JeJ_IC_GlcA} fut_{GlcA} / Km_{BCRP} + C_{JeJ_IC_GlcA} fut_{GlcA} \end{aligned}$$

$$\frac{dA_{JeJ_EC_GlcA}}{dt} = Q_{JeJ}(C_{B_GlcA} - CV_{JeJ_GlcA})$$

Liver:

$$\begin{aligned}\frac{dA_{L_IC_GlcA}}{dt} = & Vmax_{L_GlcA} C_{L_Agl} fut_{Agl} / Km_{L_GlcA} + C_{L_Agl} fut_{Agl} \\ & - Vmax_{MRP2_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{MRP2} + C_{L_IC_GlcA} fut_{fut} \\ & - Vmax_{MRP3_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{MRP3} + C_{L_IC_GlcA} fut_{GlcA} \\ & - Vmax_{BCRP_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{BCRP} + C_{L_IC_GlcA} fut_{GlcA}\end{aligned}$$

$$\begin{aligned}\frac{dA_{L_EC_GlcA}}{dt} = & Q_{Jej} CV_{Jej_GlcA} + (Q_L + Q_{DuO} + Q_{Ile}) C_{B_GlcA} - (Q_L + Q_{SI}) CV_{L_GlcA} \\ & + Vmax_{MRP3_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{MRP3} + C_{L_IC_GlcA} fut_{GlcA}\end{aligned}$$

Bile:

$$\begin{aligned}\frac{dA_{Bile_GlcA}}{dt} = & Vmax_{MRP2_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{MRP2} + C_{L_IC_GlcA} fut_{GlcA} \\ & + Vmax_{BCRP_LI} C_{L_IC_GlcA} fut_{GlcA} / Km_{BCRP} + C_{L_IC_GlcA} fut_{GlcA} - CL_{Bile} C_{Bile_GlcA}\end{aligned}$$

$$\text{Rapidly Perfused Tissue: } \frac{dA_{R_GlcA}}{dt} = Q_R (C_{B_GlcA} - CV_{R_GlcA})$$

$$\text{Slowly Perfused Tissue: } \frac{dA_{S_GlcA}}{dt} = Q_S (C_{B_GlcA} - CV_{S_GlcA})$$

$$\text{Fat: } \frac{dA_{F_GlcA}}{dt} = Q_F (C_{B_GlcA} - CV_{F_GlcA})$$

Blood:

$$\begin{aligned}\frac{dA_{B_GlcA}}{dt} = & CV_{L_GlcA} (Q_{SI} + Q_L) + CV_{F_GlcA} Q_F + CV_{R_GlcA} Q_R + CV_{S_GlcA} Q_S + CV_{K_GlcA} Q_K \\ & - C_{B_GlcA} (Q_{SI} + Q_L + Q_R + Q_S + Q_F + Q_K) - GFR C_{B_GlcA} fup_{GlcA}\end{aligned}$$

Kidney:

$$\frac{dA_{K_GlcA}}{dt} = Q_K (C_{B_GlcA} - CV_{K_GlcA}) + GFR C_{B_GlcA} fup_{GlcA} - Vmax_{MRP2_KI} C_{K_GlcA} fut_{GlcA} / Km_{MRP2} + C_{K_GlcA} fut_{GlcA}$$

$$\text{Urine: } \frac{dA_{UR_GlcA}}{dt} = Vmax_{MRP2_KI} C_{K_GlcA} fut_{GlcA} / Km_{MRP2} + C_{K_GlcA} fut_{GlcA}$$

$$\text{Feces: } \frac{dA_{FE_GlcA}}{dt} = k_{ile} A_{IleL_GlcA}$$

Table S2: Active transport and metabolism parameters and their respective in vitro-to in vivo scaling values.

Tissue / Enzyme	Parameter	Rat	Human	Reference (R; H)
Active transport				
Intestine (MRP2)	Vmax	3366e-6	3366e-6	(Li et al., 2019)
	Km	78.9	78.9	
	MGPG	133	133	(Utsey et al., 2020)
	A. in vivo	0.025	3.3	(Couto et al., 2020; Gavins et al., 2023)
	A. in vitro	133.9	133.9	(Li et al., 2019)
	REF	1.8e-4	0.0246	
	f. in-out	0.42	0.42	(Li et al., 2019)
	V. tissue	0.0024	0.2172	
	Scaled Vmax/Km	3.67e-4	162.29	
Intestine (BCRP)	Vmax	2470e-6	2470e-6	(Li et al., 2019)
	Km	676.4	676.4	
	MGPG	133	133	(Utsey et al., 2020)
	A. in vivo	0.167	0.167	(Couto et al., 2020; Gavins et al., 2023)
	A. in vitro	11.4	11.4	(Li et al., 2019)
	REF	0.015	0.0517	
	f. in-out	0.326	0.326	(Li et al., 2019)
	V. tissue	0.0024	0.2172	
	Scaled Vmax/Km	0.050	42.11	
Liver (MRP2)	Vmax	3366e-6	3366e-6	(Li et al., 2019)
	Km	78.9	78.9	
	MGPG	210	180	(Utsey et al., 2020)
	A. in vivo	2.08	0.38	(Fallon et al., 2016)
	A. in vitro	133.9	133.9	(Li et al., 2019)
	REF	0.016	0.0028	
	f. in-out	0.42	0.42	(Li et al., 2019)
	V. tissue	0.0052	0.8423	
	Scaled Vmax/Km*	0.103	62.935	
Liver (MRP3)	Vmax	986.2e-6	986.2e-6	(Li et al., 2019)
	Km	5.7	5.7	
	MGPG	210	180	(Utsey et al., 2020)
	A. in vivo	0.31	0.69	(Fallon et al., 2016)
	A. in vitro	239.4	239.4	(Li et al., 2019)
	REF	0.001	0.0029	
	f. in-out	0.58	0.58	(Li et al., 2019)
	V. tissue	0.0052	0.8423	
	Scaled Vmax/Km	2.63e-2	187.7	
Liver (BCRP)	Vmax	2470e-6	2470e-6	(Li et al., 2019)
	Km	676.4	676.4	
	MGPG	210	180	(Utsey et al., 2020)
	A. in vivo	0.167	0.13	(Fallon et al., 2016)
	A. in vitro	11.4	11.4	(Li et al., 2019)
	REF	0.015	0.0114	
	f. in-out	0.326	0.326	(Li et al., 2019)
	V. tissue	0.0052	0.8423	
	Scaled Vmax/Km*	0.11	2792.2	

Kidney (MRP2)	Vmax	3366e-6	3366e-6	(Li et al., 2019)
	Km	78.9	78.9	
	MGPg	180	170	(Utsey et al., 2020)
	A. in vivo	1.4	0.48	(Fallon et al., 2016)
	A. in vitro	133.9	133.9	(Li et al., 2019)
	REF	0.010	0.004	
	f. in-out	0.42	0.42	(Li et al., 2019)
	V. tissue	0.011	0.1488	
	Scaled Vmax/Km*	1.26	132.64	
Metabolism				
Intestine (3a-HSD)	Vmax	3.2e-5	3.59e-4	(Malekinejad et al., 2006)
	Km	592	9	
	S9	15.5	35.2	(Doerksen et al., 2021; Peters et al., 2016)
	V.tissue	0.0024	0.2172	
	Scaled Vmax/Km	1.20e-4	18.282	
Intestine (3b-HSD)	Vmax	7.2e-5	2.09e-4	(Malekinejad et al., 2006)
	Km	21	23	
	S9	15.5	35.2	(Doerksen et al., 2021; Peters et al., 2016)
	V.tissue	0.0024	0.2172	
	Scaled Vmax/Km	0.008	4.174	
Intestine (UGT1A)-ZEN	Vmax	8.61e-3	4.90e-4	(Mendez-Catala et al., 2021)
	Km	6.20	1.17	
	S9	15.5	35.2	(Doerksen et al., 2021; Peters et al., 2016)
	Alemethicin CF	3	3	(Miners et al., 2021)
	V.tissue	0.0024	0.2172	
	Scaled Vmax/Km	1.034	64.039	
Intestine (UGT1A)-aZEL	Vmax	6.96e-3	6.96e-3	(Mendez-Catala et al., 2021)
	Km	7.43	7.42	
	S9	15.5	35.2	(Doerksen et al., 2021; Peters et al., 2016)
	Alemethicin CF	3	3	(Miners et al., 2021)
	V.tissue	0.0024	0.2172	
	Scaled Vmax/Km	0.697	143.491	
Intestine (gmGUS)	Vmax	4.43	5.35	(Ebuzoeme et al., 2021)
	Km	3.08	0.41	
	S9	4	4	(Mendez-Catala et al., 2021)
	V.tissue	0.0041	0.128	(Mendez-Catala et al., 2021)
	Scaled Vmax/Km	1.41	400.86	
Liver (3a-HSD)	Vmax	3.2e-5	3.59e-4	(Malekinejad et al., 2006)
	Km	592	9	
	S9	46	122	(Mendez-Catala et al., 2021)
	V.tissue	0.0052	0.8423	
	Scaled Vmax/Km	7.75e-4	245.735	
Liver (3b-HSD)	Vmax	7.2e-5	2.09e-4	(Malekinejad et al., 2006)
	Km	21	23	
	S9	46	122	(Mendez-Catala et al., 2021)
	V.tissue	0.0052	0.8423	
	Scaled Vmax/Km	0.049	56.1073	
Liver (UGT1A)-ZEN	Vmax	7.03e-3	2.97e-3	(Mendez-Catala et al., 2021)
	Km	6.75	2.043	
	S9	46	122	(Mendez-Catala et al., 2021)
	Alemethicin CF	3	3	(Miners et al., 2021)
	V.tissue	0.0052	0.8423	
	Scaled Vmax/Km	4.982	2665.554	

Liver (UGT1A)-aZEL	Vmax	6.96e-3	2.98e-3	(Mendez-Catala et al., 2021)
	Km	7.43	2.415	
	S9	46	122	(Mendez-Catala et al., 2021)
	Alemethicin CF	3	3	(Miners et al., 2021)
	V.tissue	0.0052	0.8423	
	Scaled Vmax/Km	4.483	2538.591	

* Empirically scaled by 10-fold increase

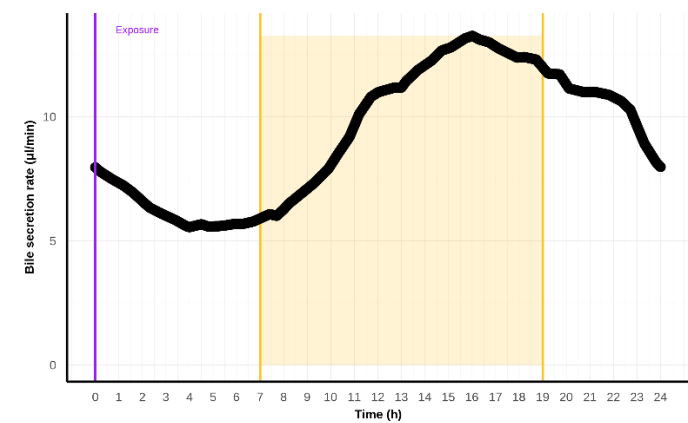


Figure S2: Modelled 24-hour bile secretion rate in rats. Time = 0 corresponds to the start of exposure, which is assumed to be at 8:00 am.

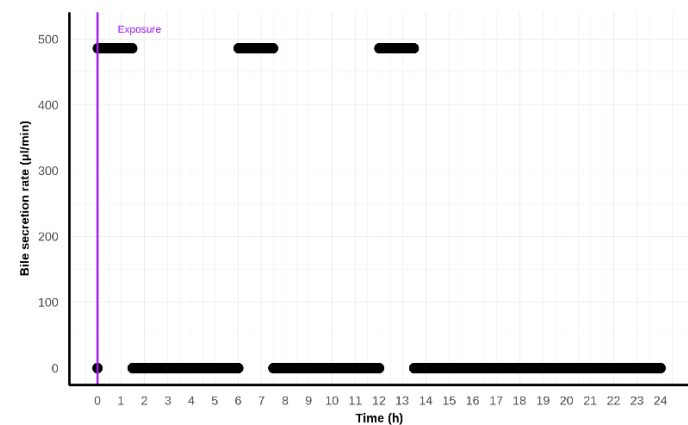


Figure S3: Modelled 24-hour bile secretion rate in humans. Time = 0 corresponds to the start of exposure, which is assumed to be at 8:00 am.

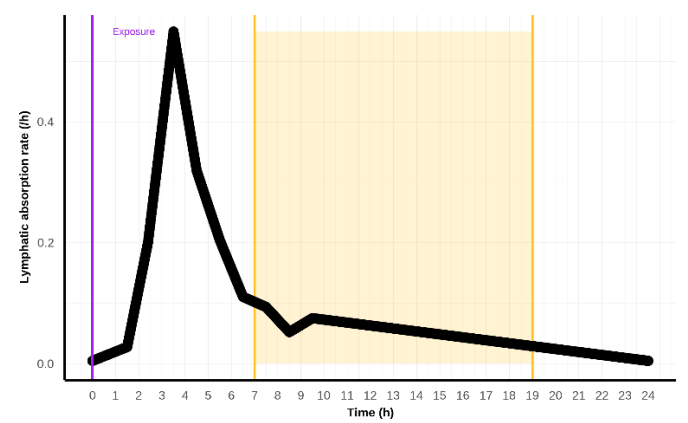


Figure S4: Modelled 24-hour bile secretion rate in humans. Time = 0 corresponds to the start of exposure, which is assumed to be at 8:00 am.

Table S3: In vitro studies and the respective assay, assay medium, derived EC50 and calculated Blood-EC50 used for the QIVIVE.

Assay	Medium	EC50 ZEN (nM)	EC50 αZEL (nM)	Relative Potency Factor	Reference
ERα_CALUX	DMEM/ F12 medium with charcoal stripped serum and non-essential amino-acids	0.490	0.010	51.042	(Ehrlich et al., 2015)
Alkaline phosphatase assay in ishikawa cells	DMEM/F12 supplemented with 5% CD-FBS and 1% P/S	0.359	0.027	13.296	(Grgic et al., 2022)
Alkaline phosphatase assay in ishikawa cells	DMEM/F12 supplemented with 5% CD-FBS stripped of endogenous oestrogen with dextran-coated charcoal	0.058	0.007	8.788	(Le Guevel and Pakdel, 2001)
ERα_CALUX	DMEM/F-12 supplemented with 5% DCC-FCS and 0.5% NEAA	1.580	0.030	52.667	(Mendez-Catala et al., 2020)
MCF-7 bioassay	Phenol red-free DMEM supplemented with 10% dextran-coated charcoal-FBS (10% DCC-FBS)	3.810	0.060	63.500	(Molina-Molina et al., 2014)
MMV-LUC cell line	DMEM and 10% hormone depleted serum	1.600	0.022	72.727	(Frizzell et al., 2011)
MCF-7 bioassay	DMEM containing 10% CD-FCS	1.639	0.050	32.840	(Malekinejad et al., 2006)
MCF-7 bioassay	RPMI medium without phenol red containing 10% DCC-FCS	0.310	0.001	221.429	(Minervini et al., 2005)

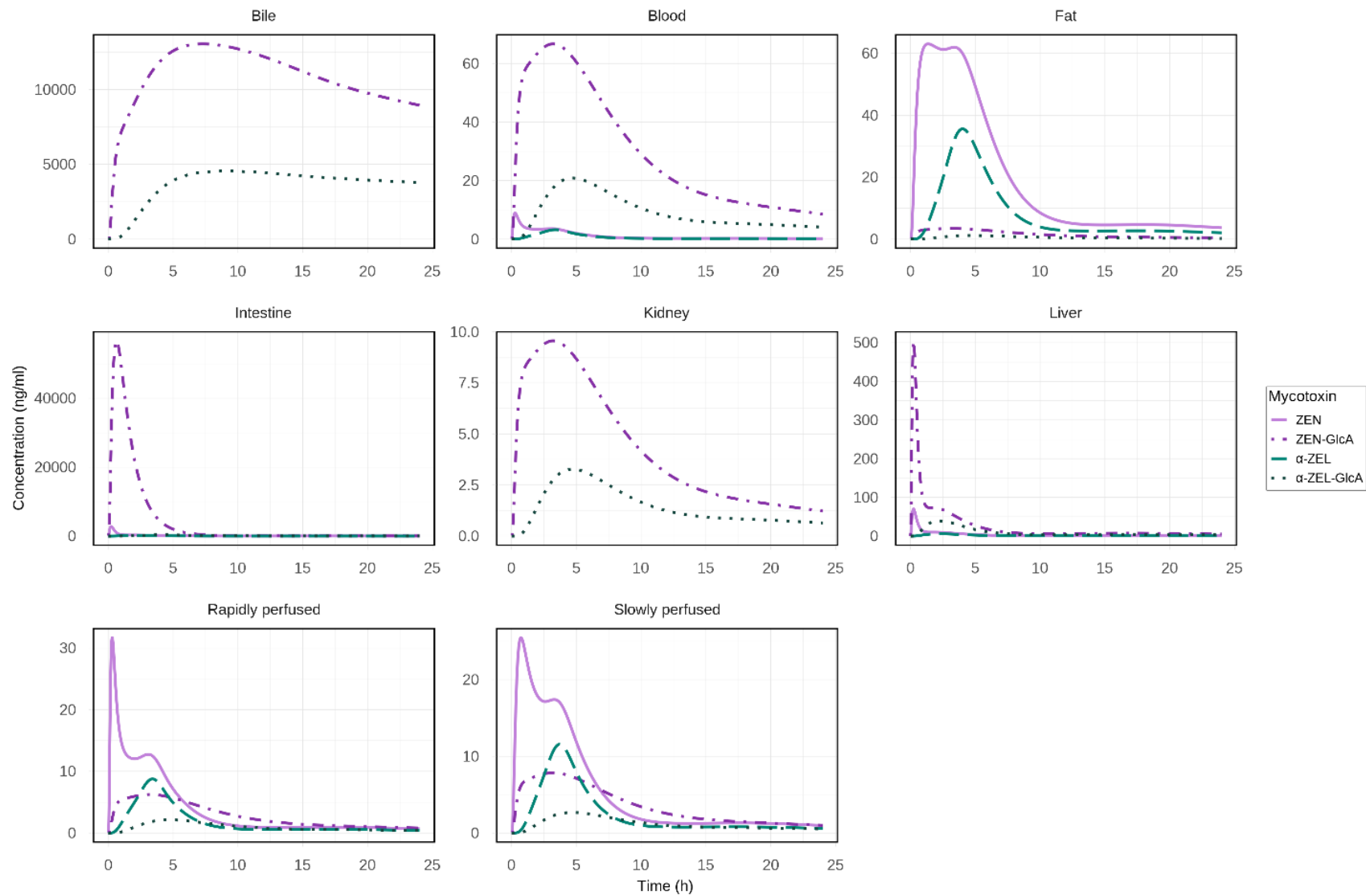


Figure S5: Predicted concentrations of ZEN, α -ZEL, ZEN-GlcA and α -ZEL GlcA in different rat tissues. Green-solid, pink-dashed, dark green-dot-dashed and dark-pink-dotted lines represent ZEN, α -ZEL, ZEN-GlcA and α -ZEL GlcA respectively.

Table S4: Influence of each modelled mechanism on the predicted toxicokinetic parameters of ZEN, ZEN-GlcA, α-ZEL and α-ZEL-GlcA.

	ZEN			ZEN-GlcA			α-ZEL			α-ZEL-GlcA		
	C _{max}	AUC	T _½	C _{max}	AUC	T _½	C _{max}	AUC	T _½	C _{max}	AUC	T _½
<i>mech-PBK</i>	0.185	0.384	11.365	0.067	0.704	12.464	0.034	0.182	11.022	0.015	0.181	14.659
<i>mech-PBK w/o</i>	0.185	0.321	1.222	0.067	0.595	3.306	0.033	0.128	1.137	0.014	0.120	2.783
<i>GlcA biliary excretion</i>												
<i>mech-PBK w/o</i>	0.185	0.237	11.236	0.065	0.465	14.051	0.003	0.042	10.880	0.003	0.046	18.146
<i>GlcA intestinal excretion</i>												
<i>mech-PBK w/o</i>	0.185	0.188	1.102	0.065	0.382	3.290	0.000	0.000	1.071	0.000	0.000	2.765
<i>gmGUS hydrolysis</i>												
<i>mech-PBK w/o</i>	0.185	0.334	11.861	0.067	0.694	12.575	0.014	0.116	11.825	0.014	0.169	15.036
<i>lymphatic absorption</i>												

C_{max}: maximum concentration in blood

AUC: area under the blood concentration over time curve

T_½: half-life

mech-PBK model: the mechanistic-PBK model in rats;

mech-PBK w/o GlcA biliary excretion: the mech-PBK model that does not include the active transport-mediated biliary excretion of GlcAs

mech-PBK w/o GlcA intestinal excretion: the mech-PBK model that does not include the active transport-mediated intestinal excretion of GlcAs

mech-PBK w/o gmGUS hydrolysis: the mech-PBK model that does not include the gmGUS hydrolysis

mech-PBK w/o lymphatic absorption: the mech-PBK model that does not include lymphatic absorption of ZEN and α-ZEL

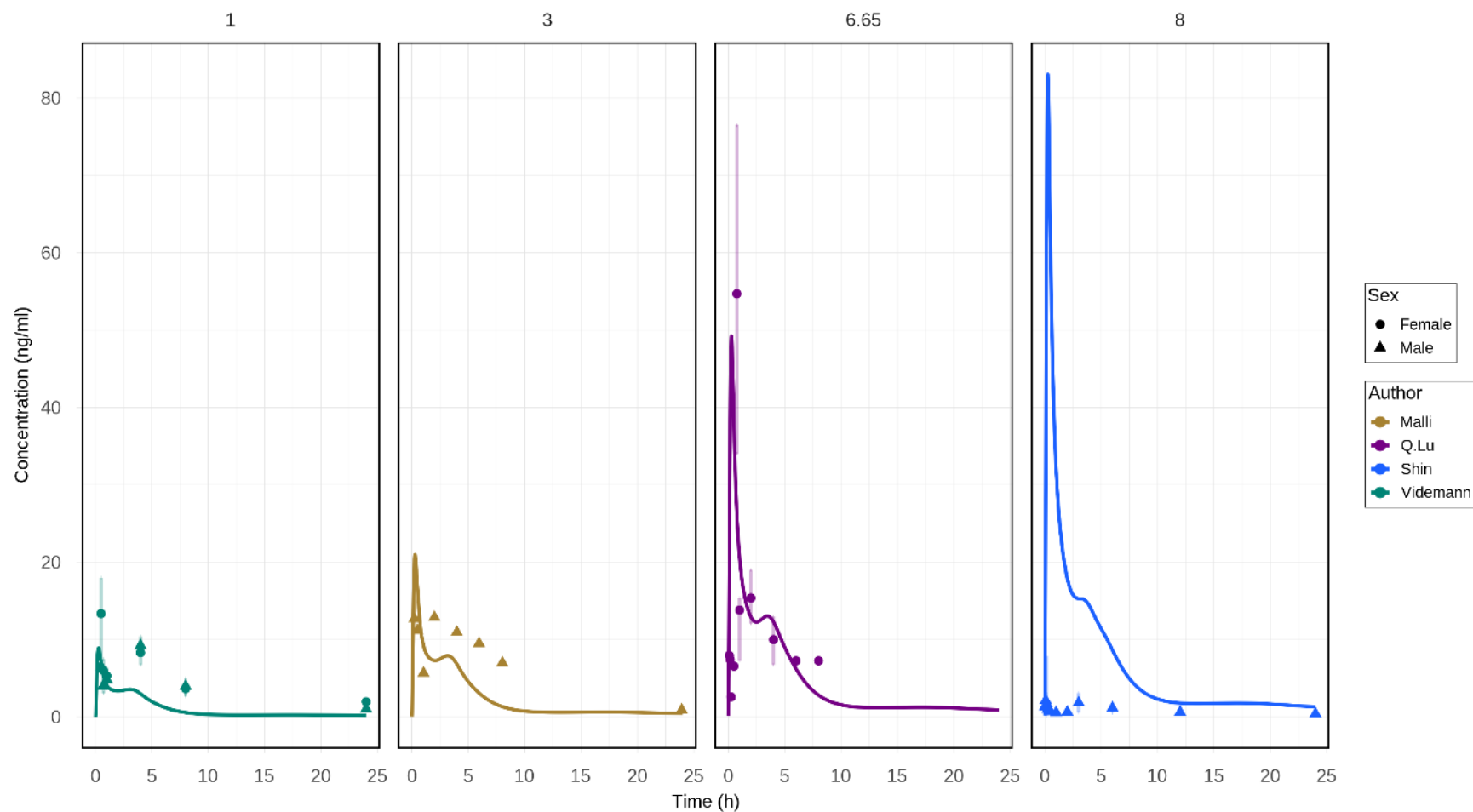


Figure S6: Evaluation of the mechanistic-PBK model against three rat studies. In all studies rats were exposed to ZEN orally. From left to right: (Videmann et al., 2012) in green, dose 1 mg/kg bw; (Mallis et al., 2003) in yellow, dose 3 mg/kg bw; (Lu et al., 2022) in magenta, dose 6.65 mg/kg bw; and (Shin et al., 2009) in blue, dose 8 mg/kg bw. Predicted (solid lines) and observed (points).

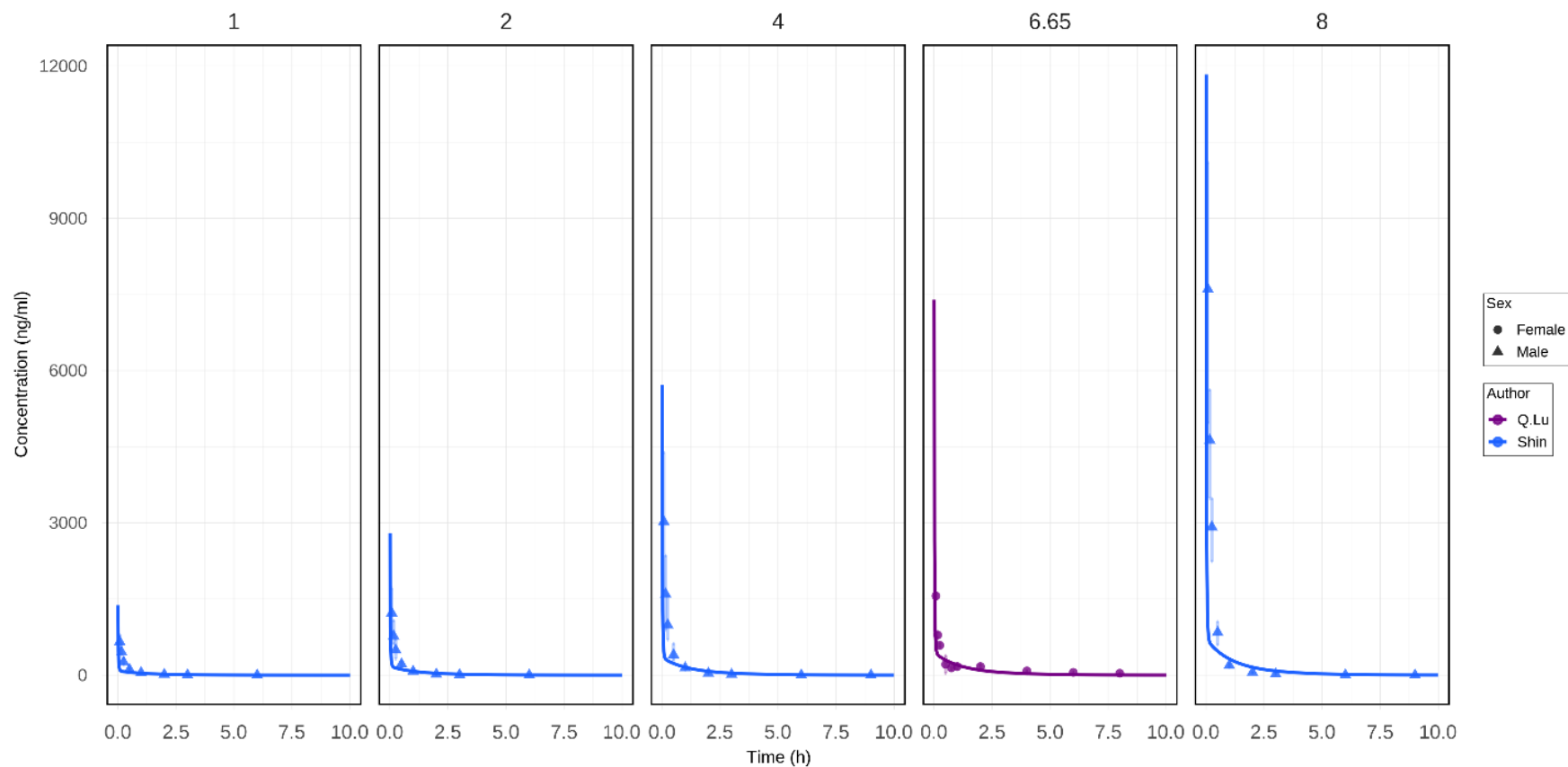


Figure S7: Predicted (solid lines) and observed (points) Zearalenone blood concentration over time curves after IV administration. Data corresponding to the studies of (Lu et al., 2022) and (Shin et al., 2009) are represented in purple, and blue respectively. The panels are separated according to the exposure scheme (1, 2, 4, 6.65 and 8 mg/kg bw/day).

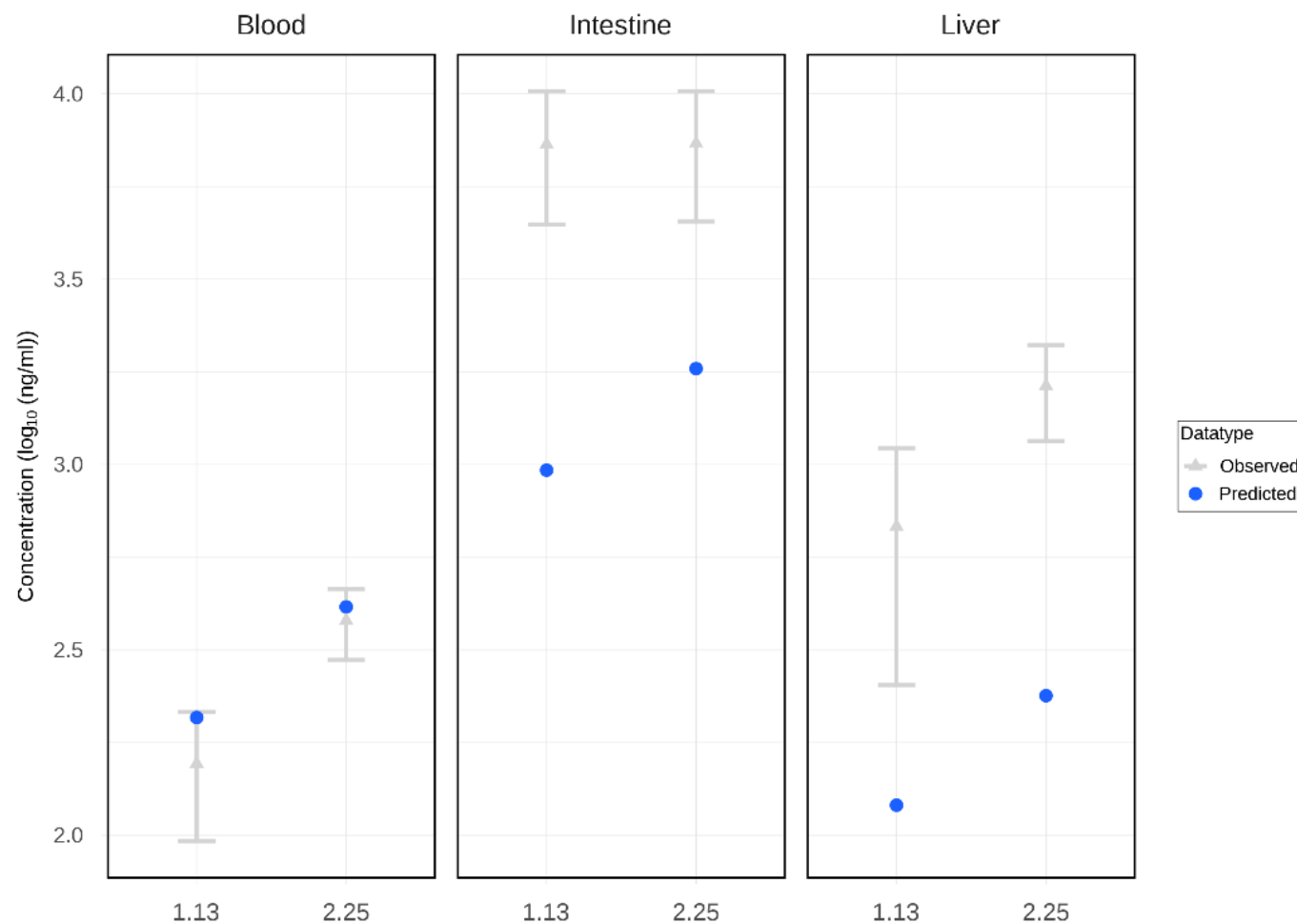


Figure S8: Comparison of the predicted and measured paired blood, liver and intestine concentrations of ZEN, after 6 hours of IV infusion at a low (1.13 mg/h/kg) and high (2.25 mg/h/kg) perfusion rate. The sampling time of 6 hours after continuous perfusion, corresponds to the time needed to reach steady state in the reference study (Shin et al., 2009). Blue circles and grey triangles represent predicted and observed data respectively.

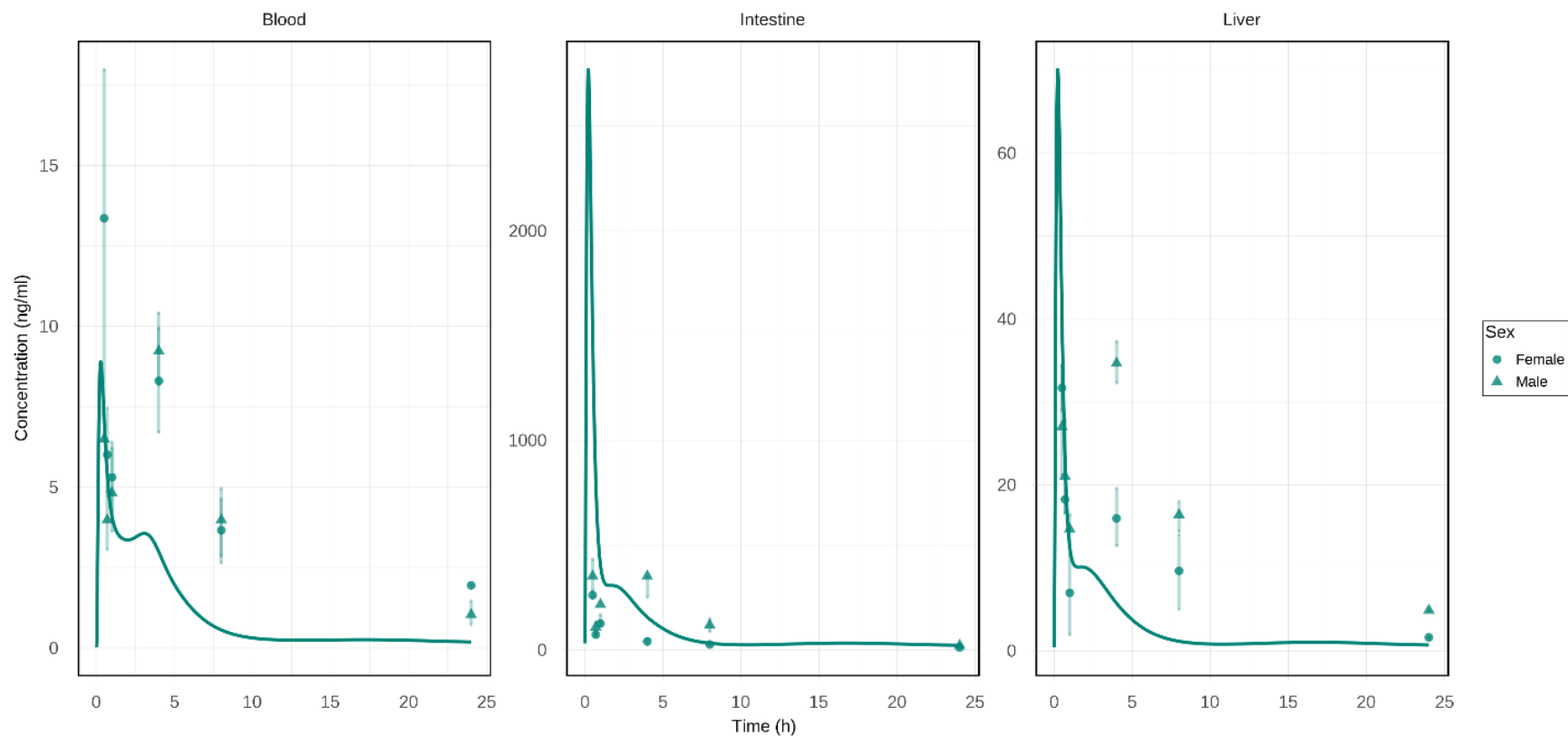


Figure S9: Evaluation of the Mechanistic-PBK model predicted blood, intestinal and liver concentrations of ZEN over time, against those measured in (Videmann et al., 2012). Solid lines and points (circles for females, triangles for males) represent model predictions and measured data respectively.

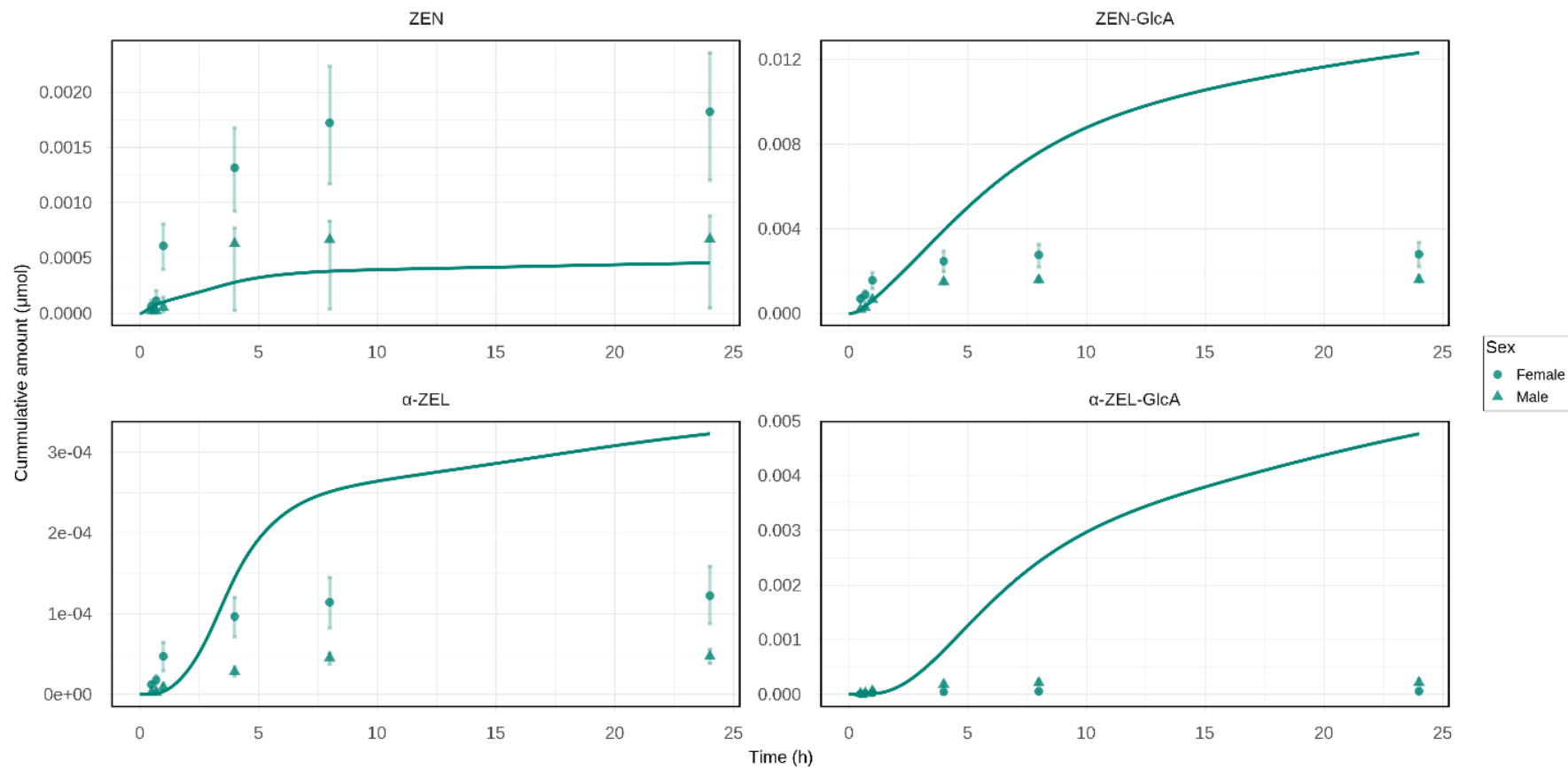


Figure S10: Evaluation of the Mechanistic-PBK model predicted cumulative amounts of ZEN, and its three metabolites in urine over time, against those measured in (Videmann et al., 2012). Solid lines and points (circles for females, triangles for males) represent model predictions and measured data respectively.

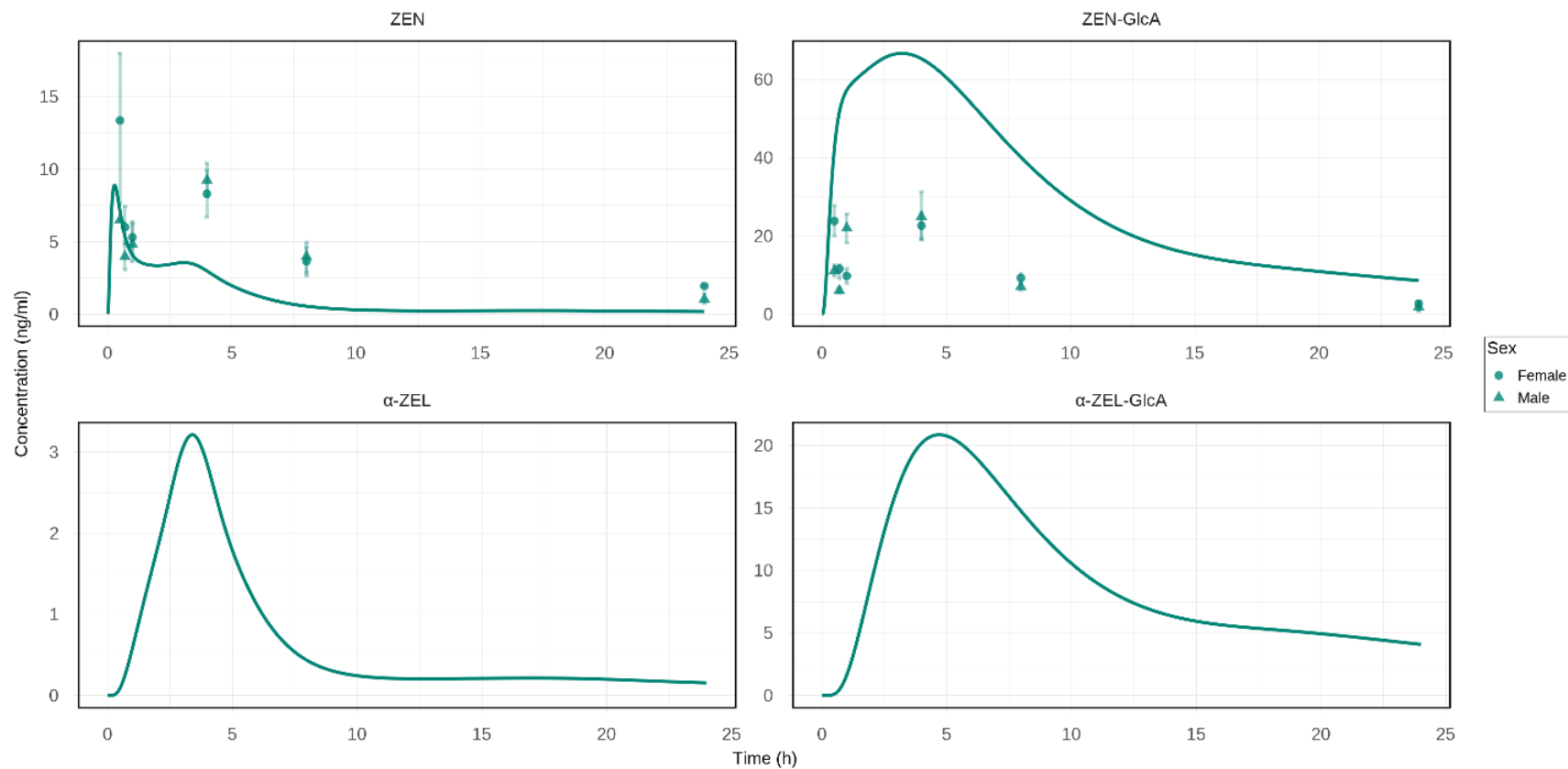


Figure S11: Evaluation of the Mechanistic-PBK model predicted blood concentrations of ZEN, and its three metabolites over time, against those measured in (Videmann et al., 2012). Solid lines and points (circles for females, triangles for males) represent model predictions and measured data respectively. Measured data for αZEL and αZEL-GlcA were below the LOQ and were therefore not available for evaluation.

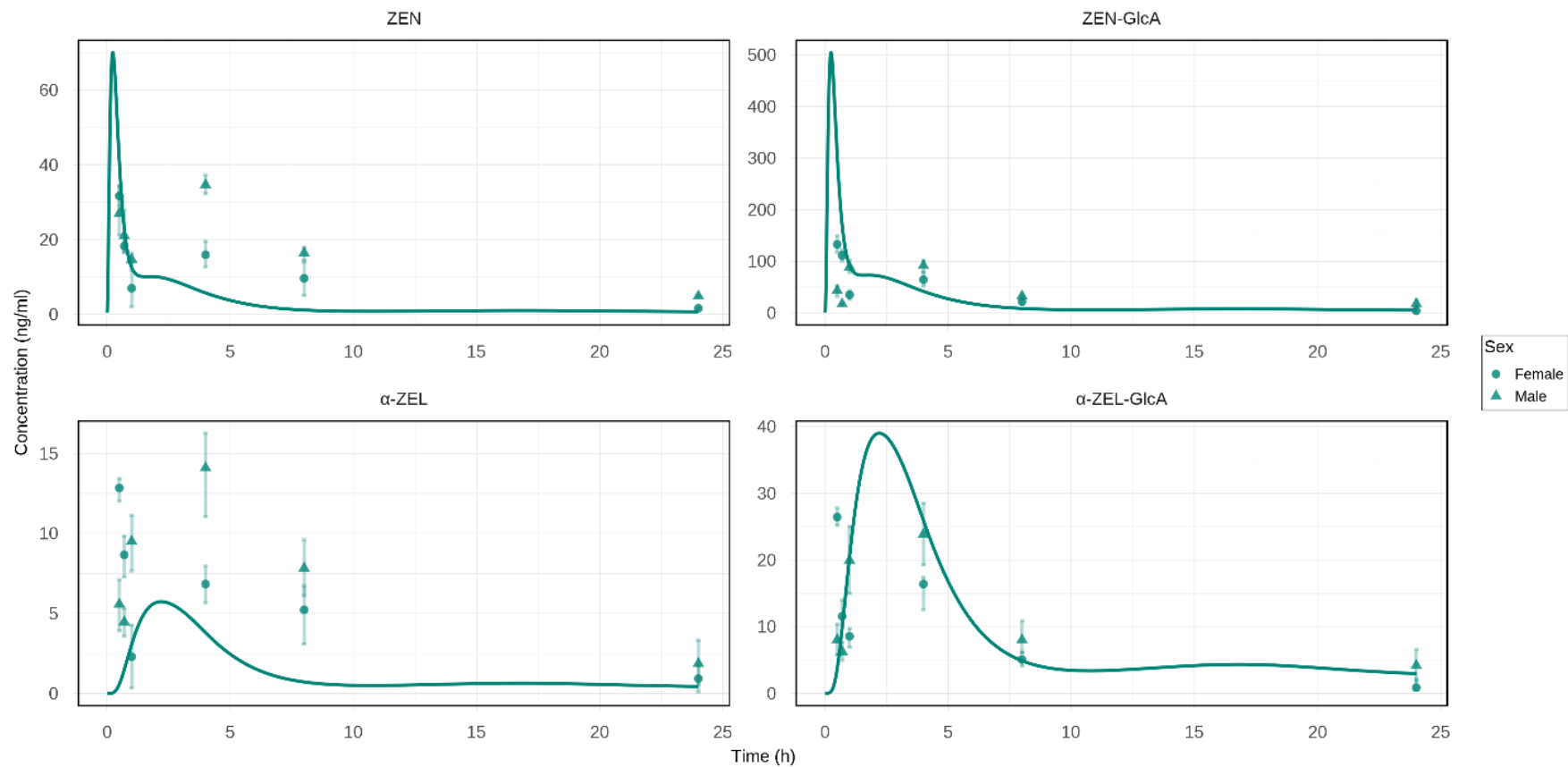


Figure S12: Evaluation of the Mechanistic-PBK model predicted liver concentrations of ZEN, and its three metabolites over time, against those measured in (Videmann et al., 2012). Solid lines and points (circles for females, triangles for males) represent model predictions and measured data respectively.

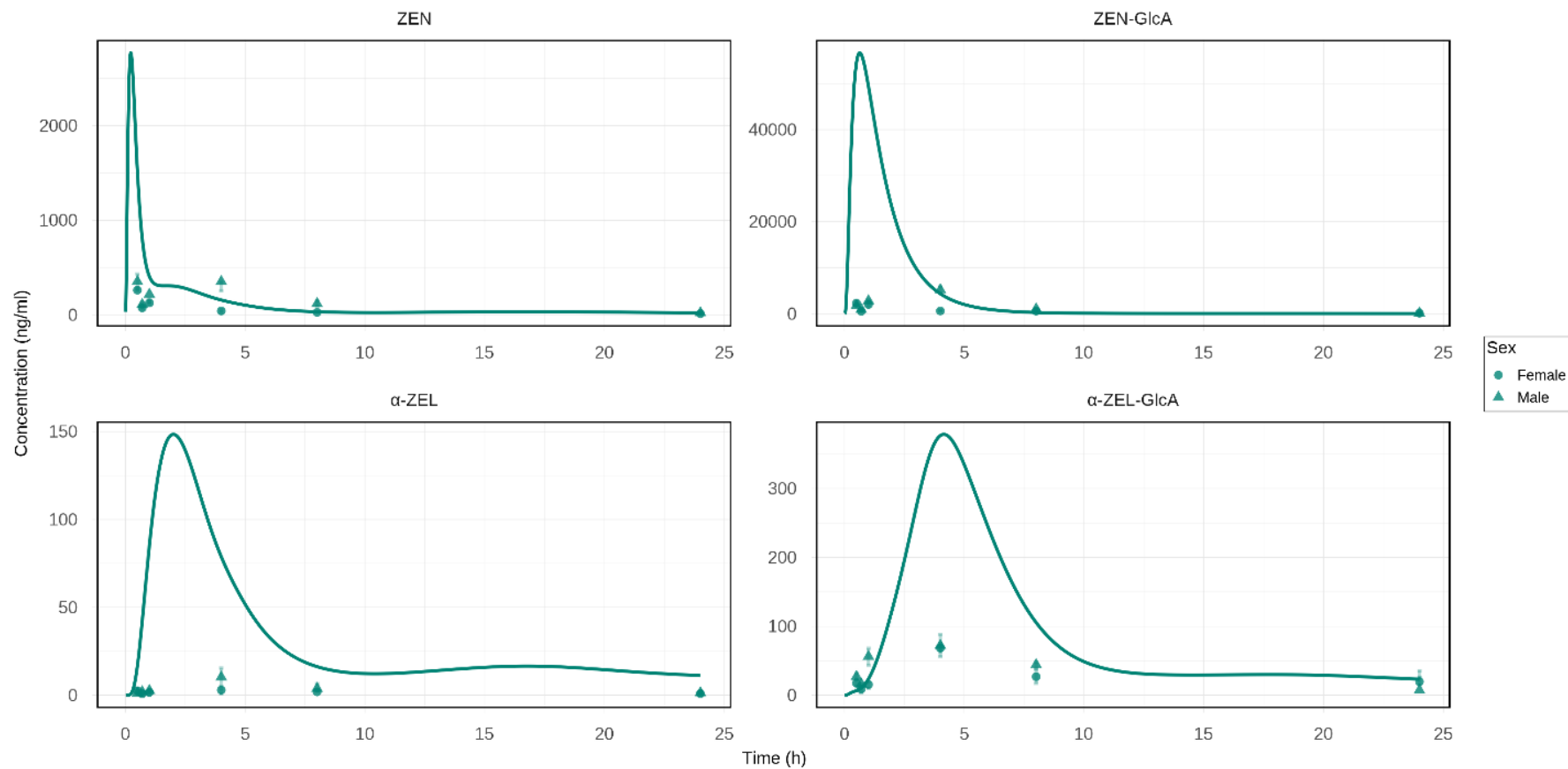


Figure S13: Evaluation of the Mechanistic-PBK model predicted intestinal concentrations of ZEN, and its three metabolites over time, against those measured in (Videmann et al., 2012). Solid lines and points (circles for females, triangles for males) represent model predictions and measured data respectively.

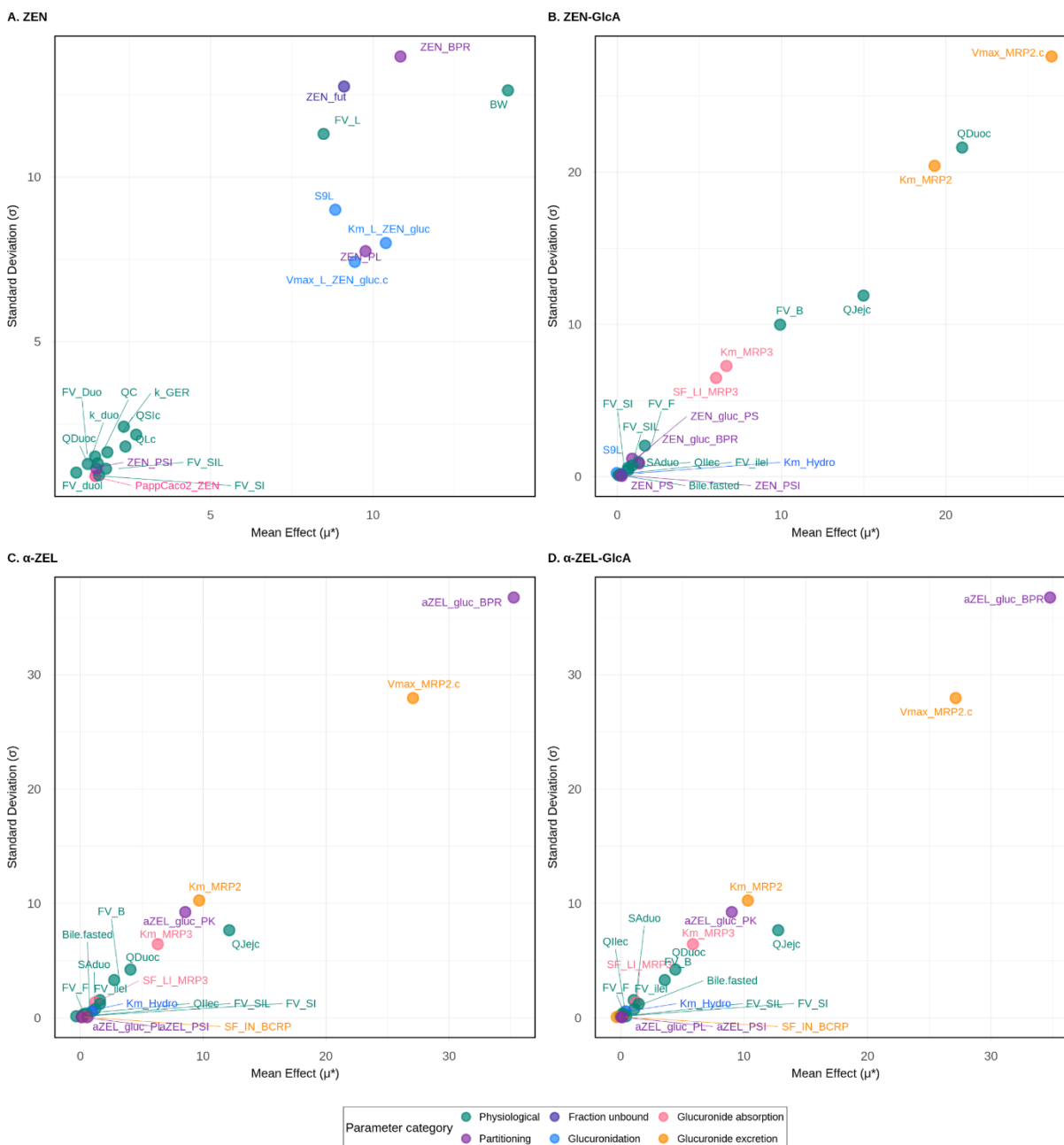
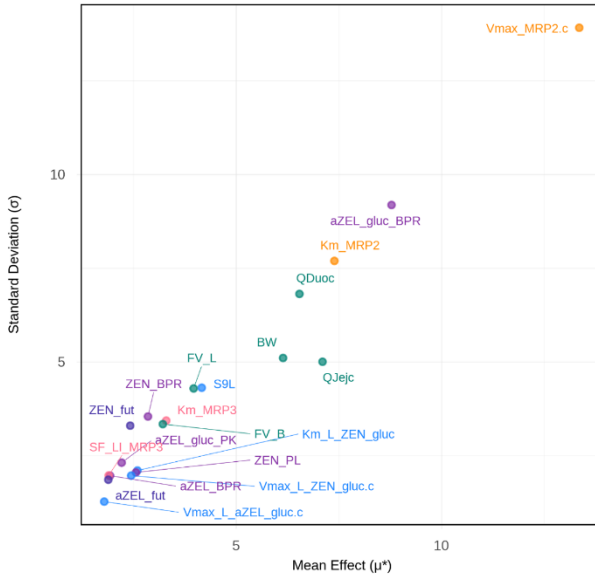
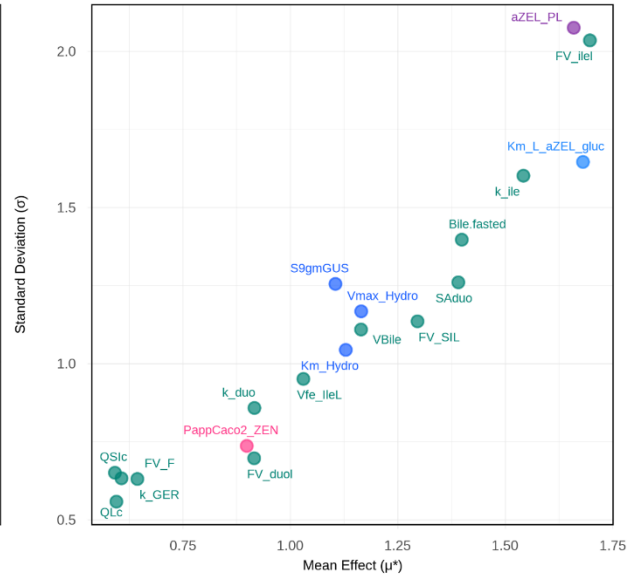


Figure S14: Sensitivity of the rat-PBK model-predicted blood concentration of ZEN (A), ZEN-GlcA (B), α-ZEL (C) and α-ZEL-GlcA (D) according to the Morris Screening test. The graphs show only the 20 most sensitive parameters.

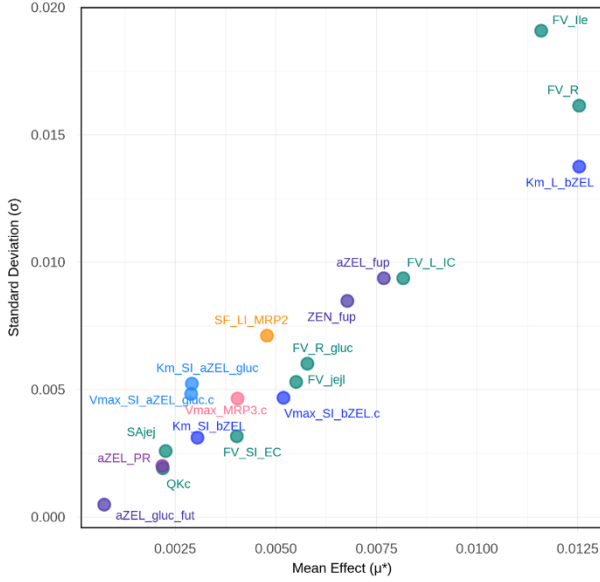
A. Q1



B. Q2



C. Q3



D. Q4

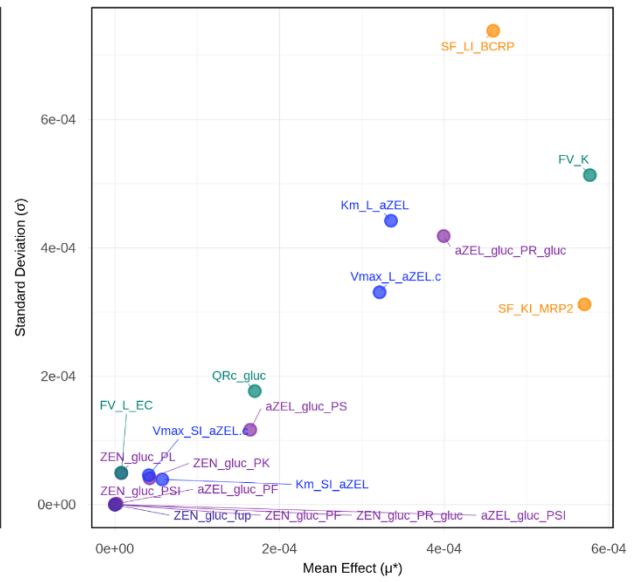


Figure S15: Combined sensitivity of the rat-PBK model-predicted blood concentrations of ZEN, ZEN-GlcA, α -ZEL and α -ZEL-GlcA according to the Morris Screening test, from the most (A.) to the least (D.) sensitive parameters.

Table S5: The most highly ranked parameters towards the predictions of the mech-PBK model in rats, according to the Morris test.

Parameter abbreviation	Maximum μ^*	Occurrence
aZEL_gluc_BPR	35.12044646	1
Vmax_MRP2.c	26.73088629	2
QDuoc	20.82457636	3
Km_MRP2	19.67274002	2
QJc	15.47824976	3
BW	14.32626759	1
ZEN_BPR	11.13329221	1
Km_L_ZEN_gluc	10.04151483	1
ZEN_PL	9.931084223	1
FV_B	9.616027882	2
ZEN_fut	9.486945151	1
Vmax_L_ZEN_gluc.c	9.464797862	1
S9L	8.933429931	3
aZEL_gluc_PK	8.844914369	1
FV_L	8.770934137	1
Km_MRP3	7.006982035	2
SF_LI_MRP3	6.252030603	2
k_GER	2.393179279	1
QLc	2.259402312	1
QSlc	2.252509927	3
FV_SIL	2.109261031	3
FV_F	2.062896934	2
QC	1.672067887	1
FV_ilel	1.491121865	3
PappCaco2_ZEN	1.334865298	3
FV_SI	1.304490067	3
ZEN_PSI	1.268870222	3
k_duo	1.253279443	1
FV_duol	1.211429089	3
FV_Duo	1.210249648	1
SADuo	1.180067452	3
VBile	1.139901465	1
ZEN_gluc_PS	1.139241933	1
Bile.fasted	1.099359542	3
k_ile	1.041715795	1

Maximum μ^* : the maximum mean effect towards the predicted concentration of ZEN, ZEN-GlcA, α -ZEL and α -ZEL-GlcA
Occurrence: occurrence of the parameter within the 40 most highly ranked parameters across all mycotoxins

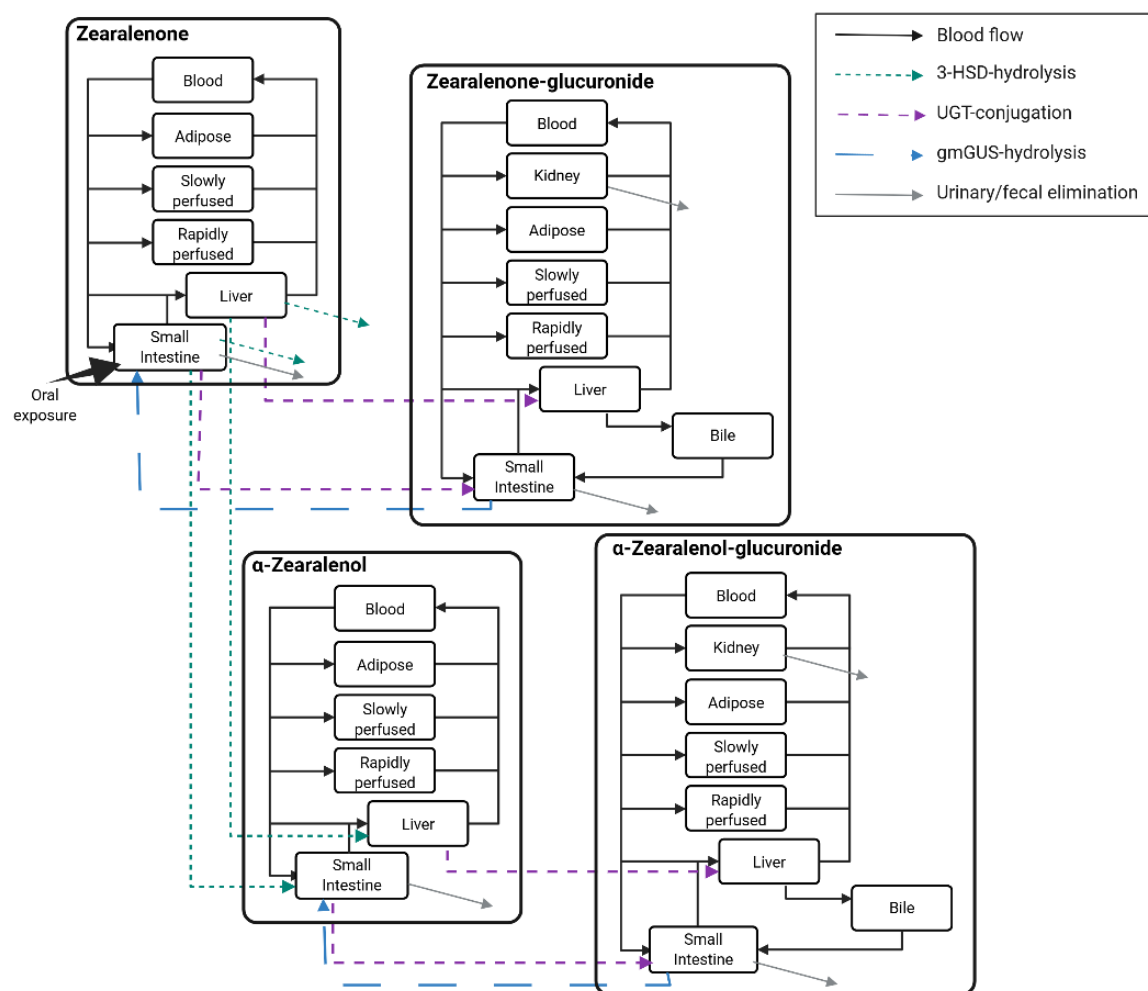


Figure S17. Schematic representation of the human-PBK model for Zearalenone and its metabolites. Solid lines blood and grey lines represent the blood flow and urinary or fecal elimination respectively. Dashed lines represent metabolism; dense-dash green, dash-purple and loose-dash blue lines representing 3-HSD hydrolysis, UGT-conjugation and gmGUS-hydrolysis respectively.

Table S6: Predicted and observed total ZEN (ZEN + ZEN-GlcA¹) urine levels

	Predicted	Observed	Fold-change	Reference
Total daily cumulative amount in urine (μg)	3 165	32 133	10.14	(Mirocha et al., 1981)
Total daily cumulative amount in urine (μg)	1.49	0.94	1.55	(Warth et al., 2013)
Total concentration in urine ($\mu\text{g/L}$) ²	0.032	0.0413 (0.01-0.138) ³	-	(McKeon et al., 2024)

¹99.97% ZEN-GlcA; ² Assuming a total urine volume of 2.42 L; ³ Estimated Daily Intake: 0.0331 $\mu\text{g/kg bw}$

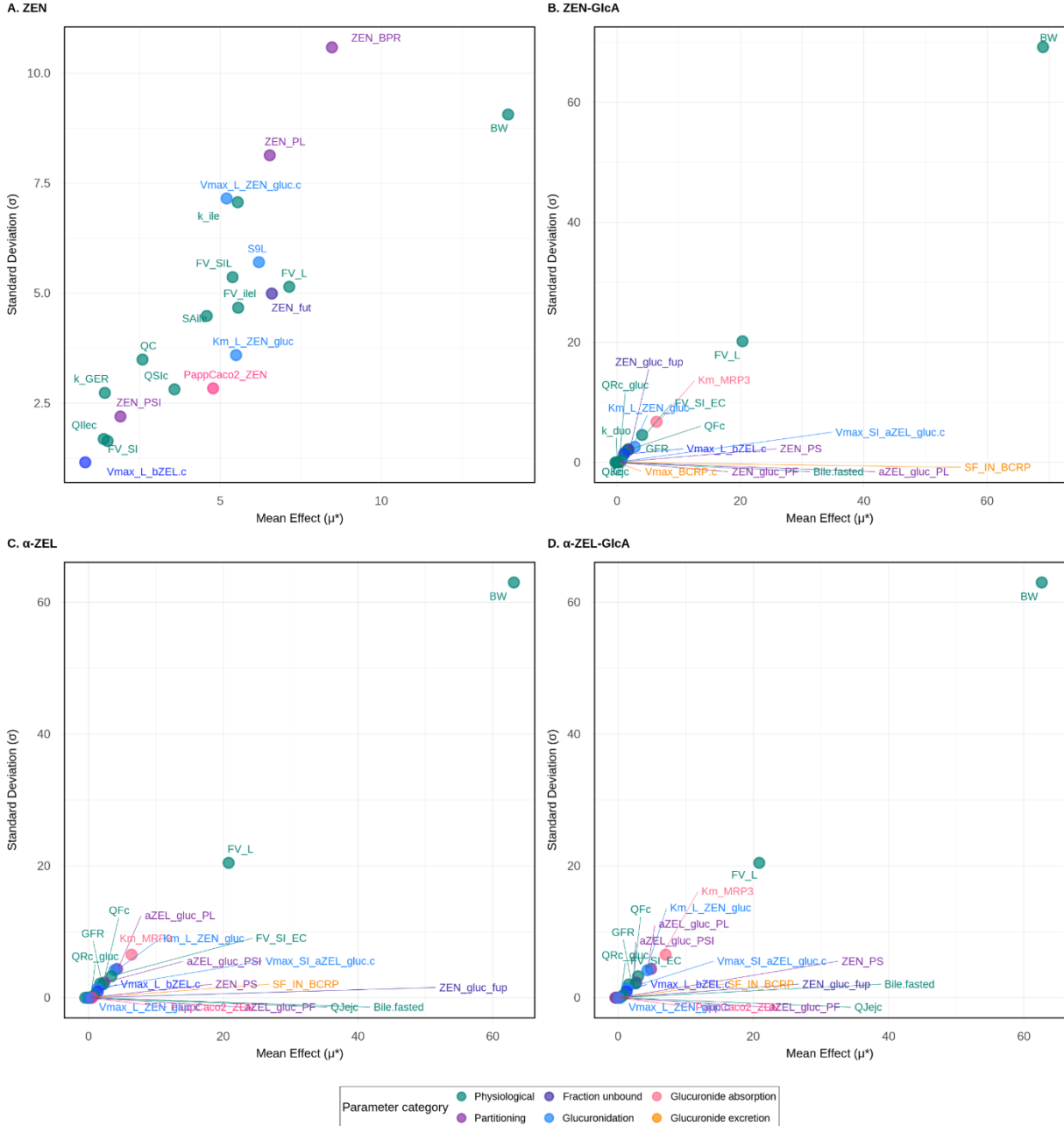


Figure S18: Sensitivity of the rat-PBK model-predicted blood concentration of ZEN (A), ZEN-GlcA (B), α-ZEL (C) and α-ZEL-GlcA (D) according to the Morris Screening test. The graphs show only the 20 most sensitive parameters

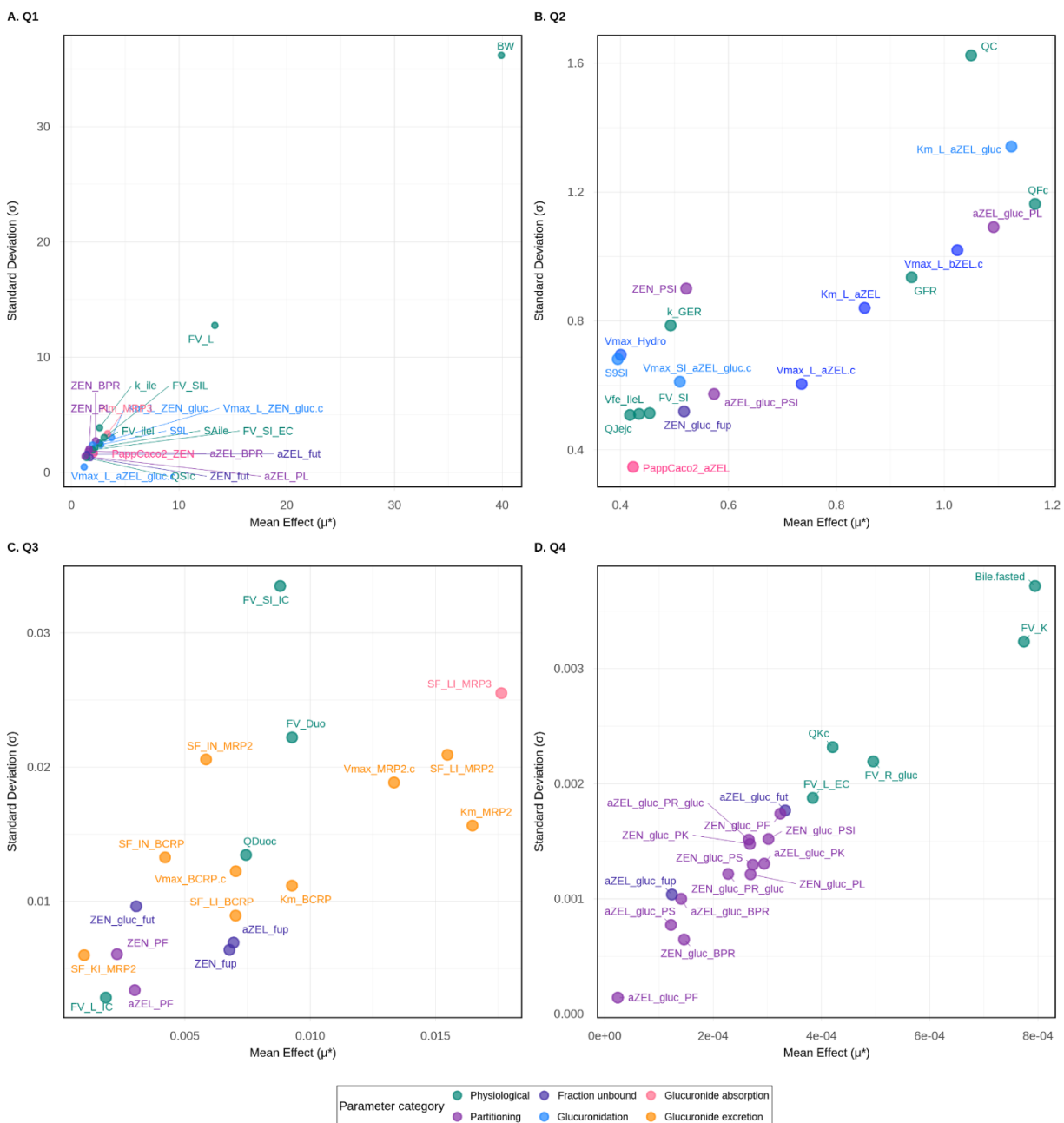


Figure S19: Combined sensitivity of the human-PBK model-predicted blood concentrations of ZEN, ZEN-GlcA, α -ZEL and α -ZEL-GlcA according to the Morris Screening test, from the most (A.) to the least (D.) sensitive parameters.

Table S7: The most highly ranked parameters towards the predictions of the mech-PBK model in humans, according to the Morris test.

<i>Parameter abbreviation</i>	<i>Maximum μ^*</i>	<i>Occurrence</i>
BW	69.19731	3
FV_L	20.45681	3
ZEN_BPR	8.751136	1
ZEN_fut	6.76067	3
Km_MRP3	6.75578	2
S9L	6.577332	1
ZEN_PL	6.561391	3
Km_L_ZEN_gluc	5.585515	3
Vmax_L_ZEN_gluc.c	5.489152	3
FV_SIL	5.441808	1
FV_ilel	5.431249	1
k_ile	5.08075	1
SAile	4.908376	2
PappCaco2_ZEN	4.624308	3
FV_SI_EC	4.560292	2
aZEL_gluc_PL	4.362382	2
QSIc	3.455075	2
QC	2.323755	2
aZEL_gluc_PSI	2.293812	1
QFc	2.184042	2
ZEN_gluc_fup	2.067421	2
GFR	2.019644	2
ZEN_PSI	1.68624	1
k_GER	1.449279	1
Vmax_SI_aZEL_gluc.c	1.414741	2
Vmax_L_bZEL.c	1.367118	3
FV_SI	1.197402	1
Qllec	1.148487	1

Maximum μ^ : the maximum mean effect towards the predicted concentration of ZEN, ZEN-GlcA, α -ZEL and α -ZEL-GlcA*

Occurrence: occurrence of the parameter within the 40 most highly ranked parameters across all mycotoxins

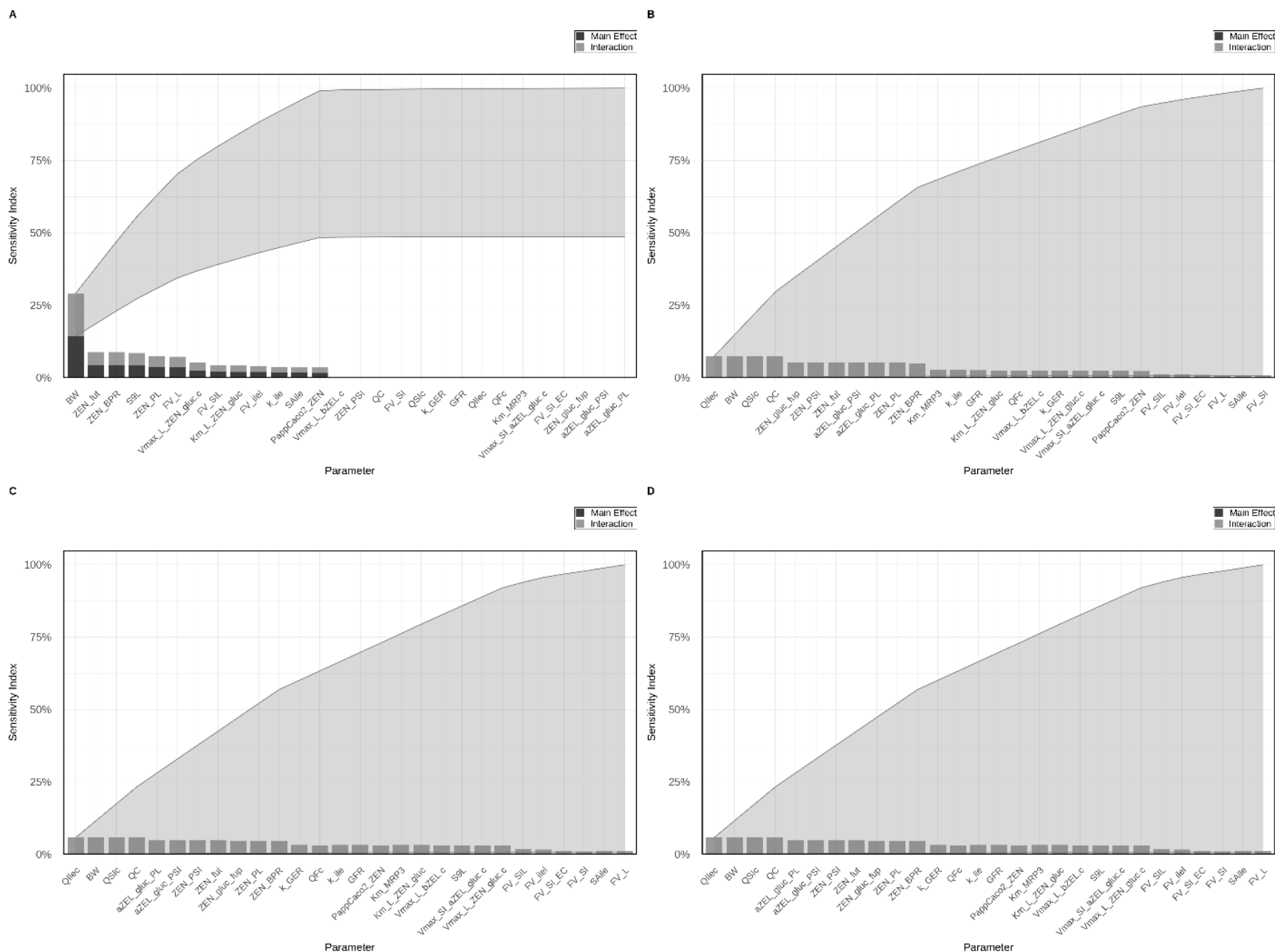


Figure S20: Lowry plots illustrating the results of the eFAST analysis for all model outputs: blood concentration of ZEN (A), ZEN-GlcA (B), α -ZEL (C) and α -ZEL-GlcA (D) of the human-PBK model. The main effect (dark grey) and interaction (light grey) are stacked on top of each other to reflect the total effect of each parameter. The grey ribbon represents the cumulative effect of all parameters to the model output.

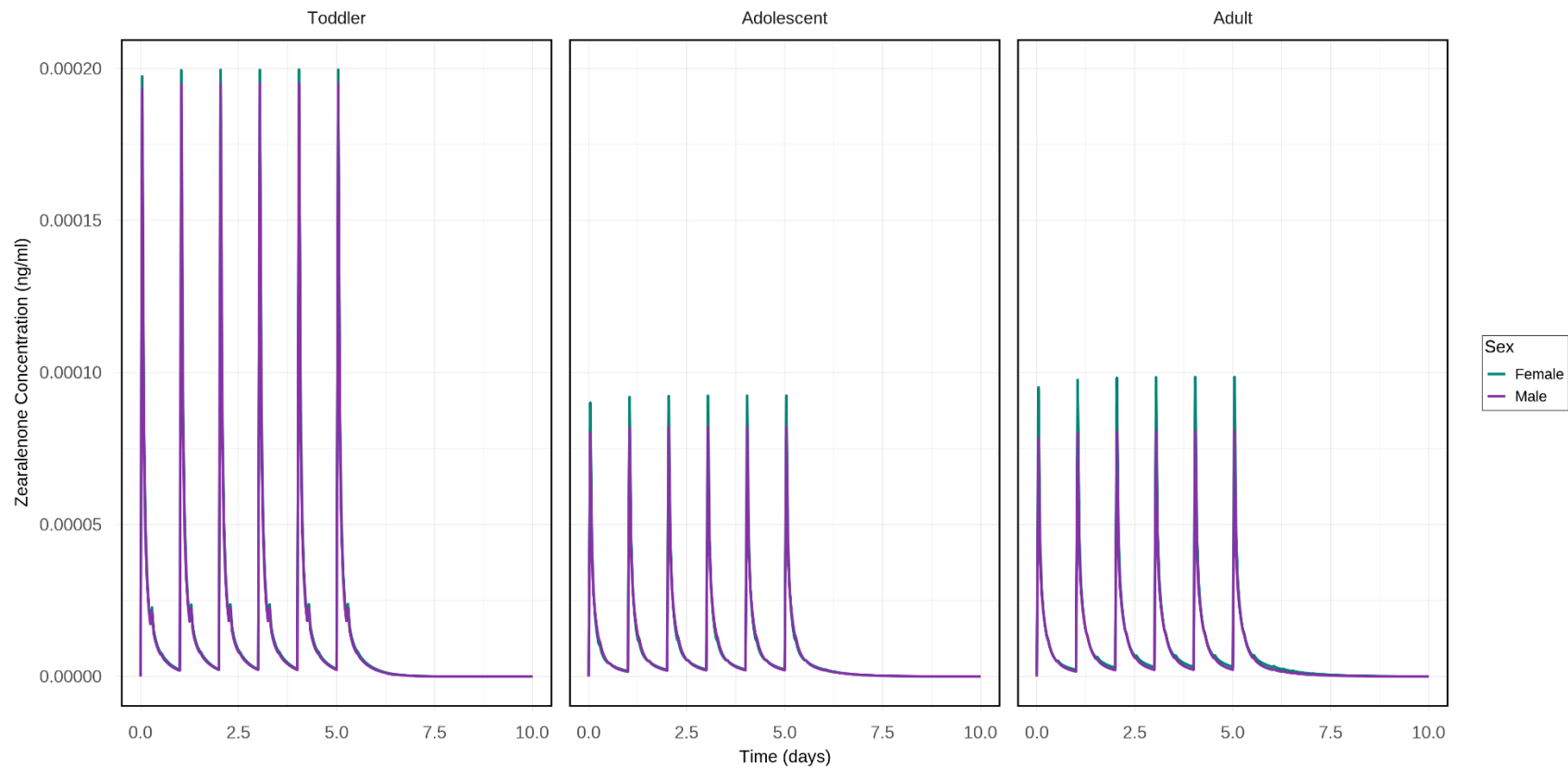


Figure S21: Predicted ZEN blood concentrations in both sexes and different age groups (toddlers, adolescents, adults) after a 6-day-long daily exposure to ZEN of $0.25\mu\text{g/kg bw/day}$, corresponding to the TDI. Purple and green lines represent females and males respectively.

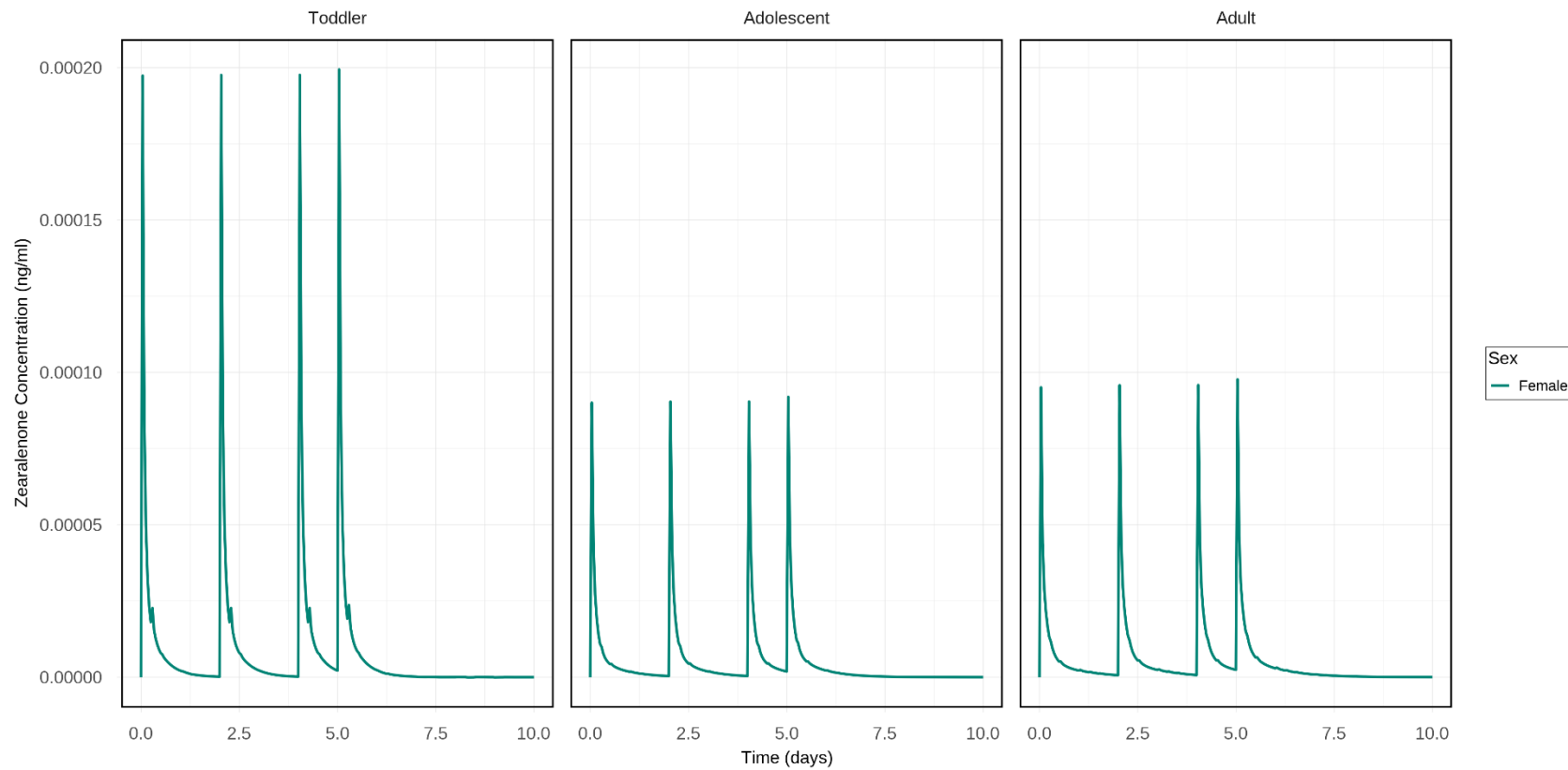


Figure S22: Predicted ZEN blood concentrations in females and different age groups (toddlers, adolescents, adults) after ZEN exposure of 0.25 μ g/kg bw/day at day 1, 3, 4 and 5, corresponding to the TDI.

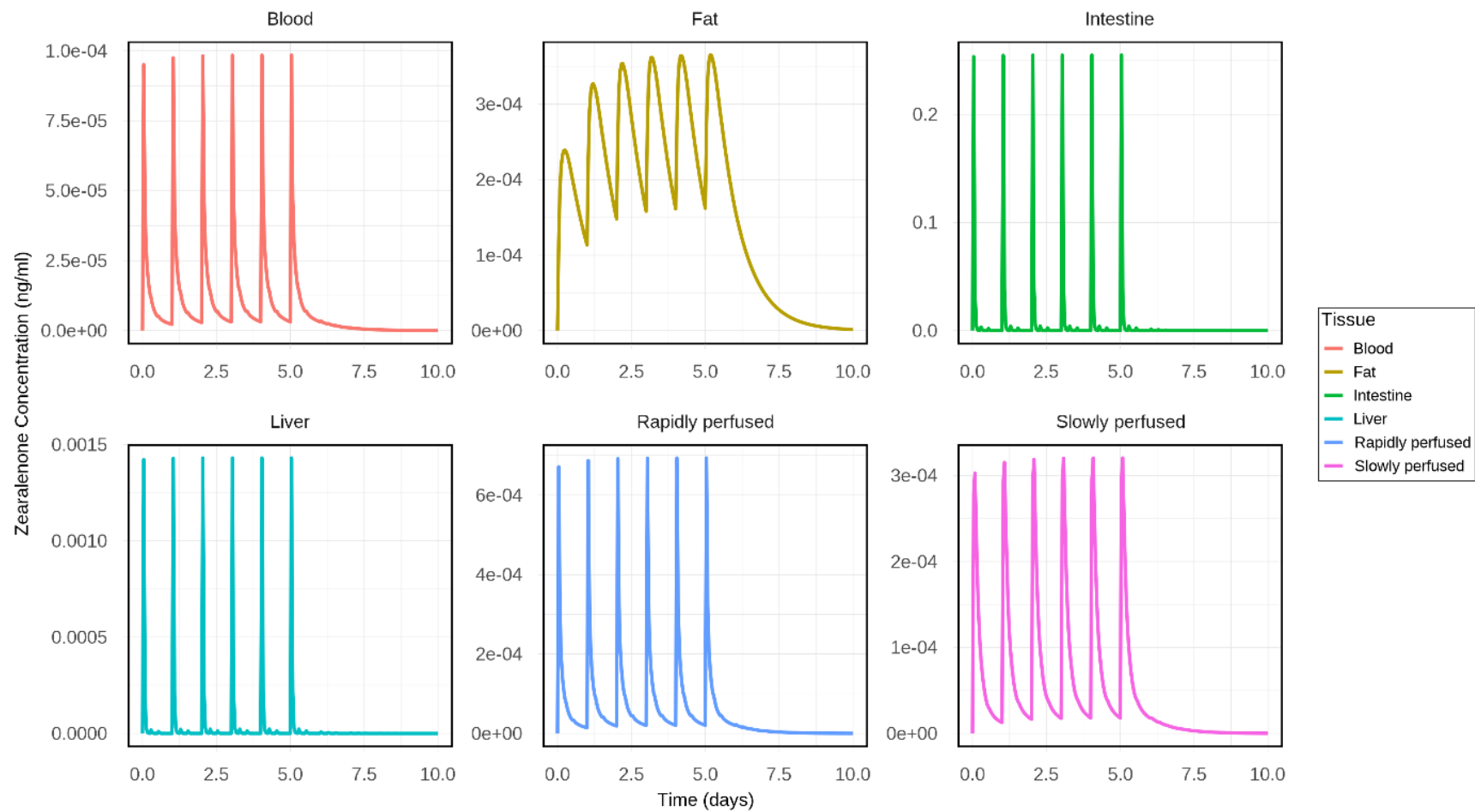


Figure S23: Predicted ZEN tissue concentrations (ng/ml) over time (days), in a 30-year old female. The subject was simulated to have a daily exposure of ZEN of 0.25 ng/kg bw, corresponding to the TDI for 7 days.

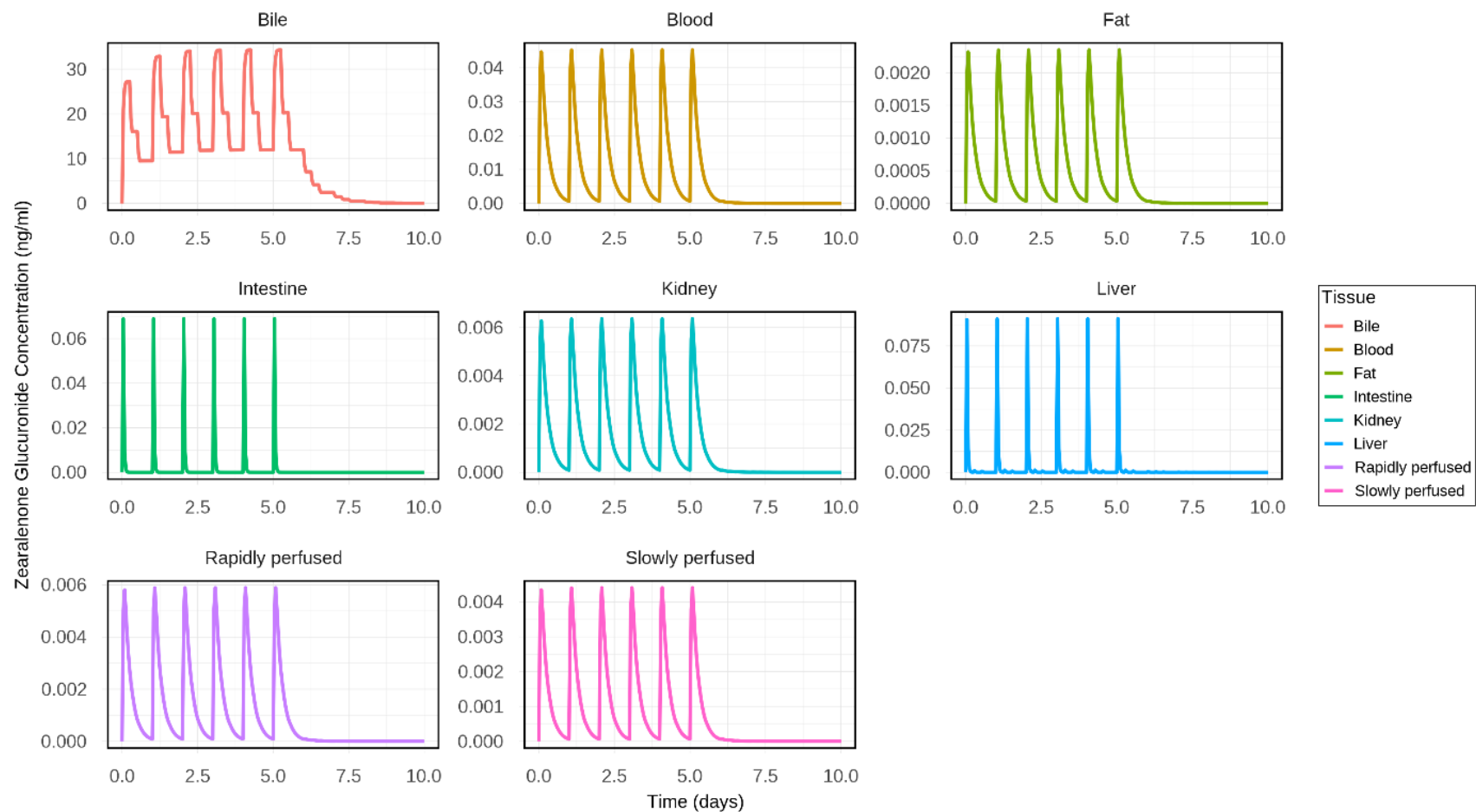


Figure S24: ZEN-GlcA tissue concentrations (ng/ml) over time (days), in a 30-year old female. The subject was simulated to have a daily exposure of ZEN of 0.25 ng/kg bw, corresponding to the TDI for 7 days.

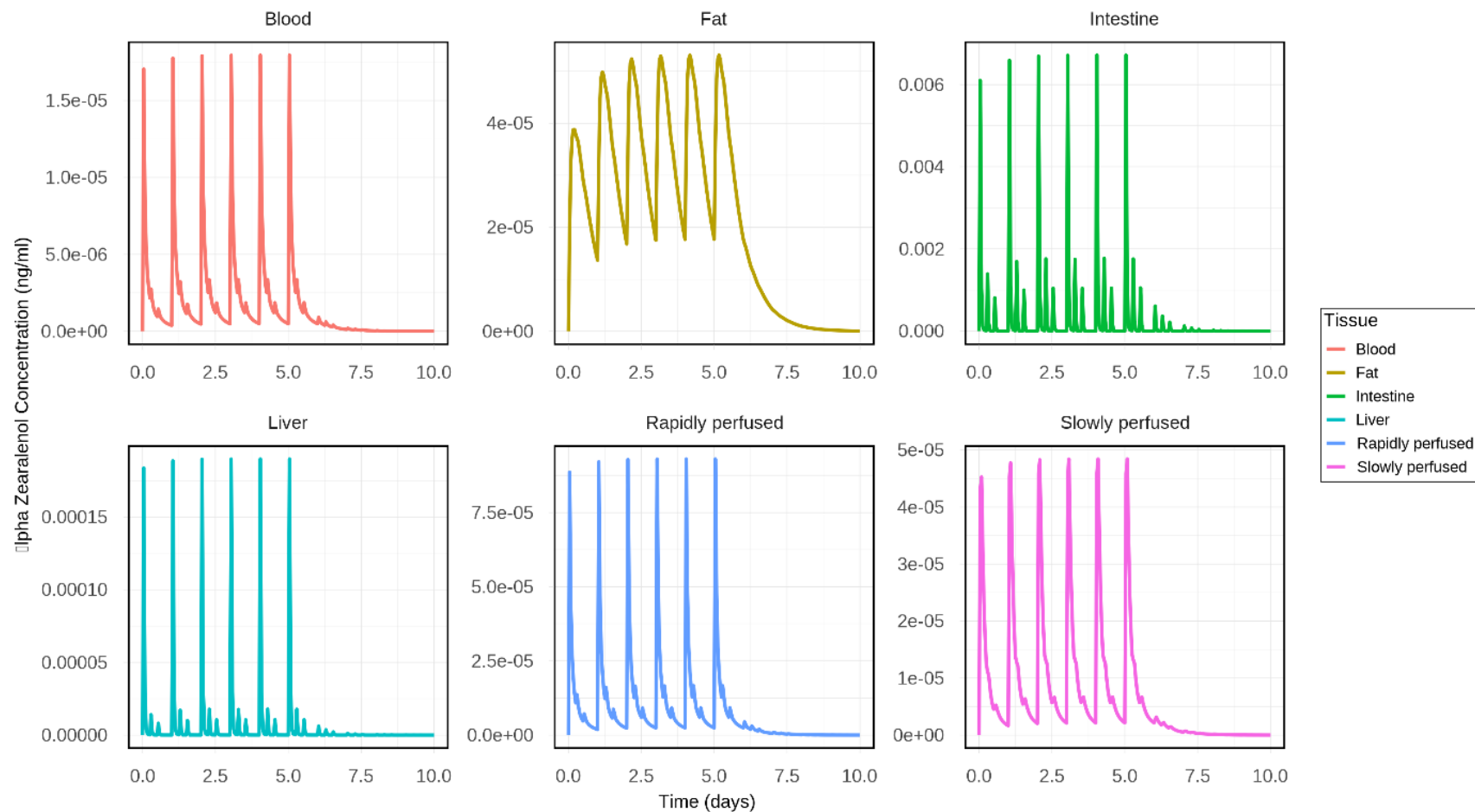


Figure S25: Predicted α -ZEL tissue concentrations (ng/ml) over time (days), in a 30-year old female. The subject was simulated to have a daily exposure of ZEN of 0.25 ng/kg bw, corresponding to the TDI for 7 days.

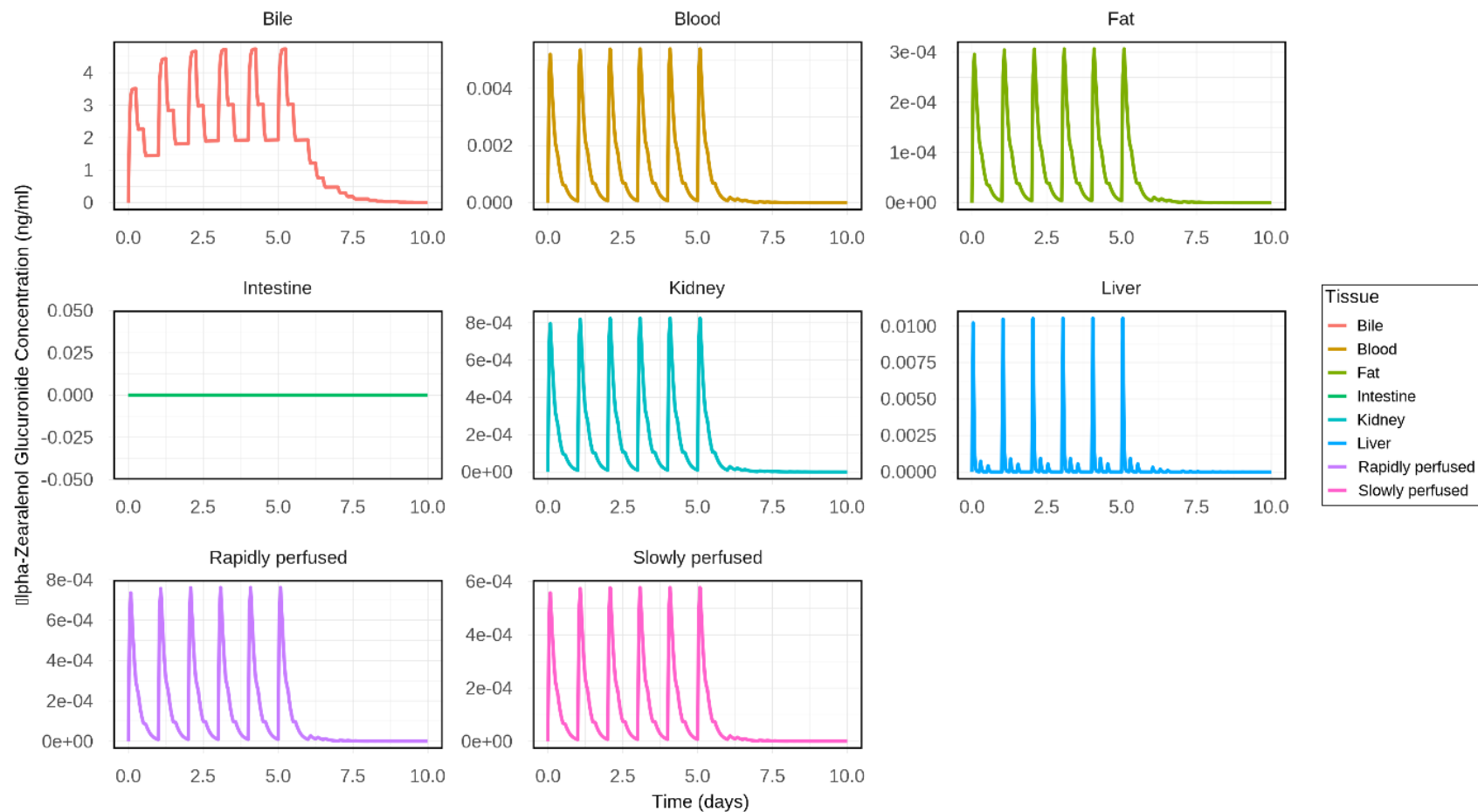


Figure S26: Predicted α -ZEL-GlcA tissue concentrations (ng/ml) over time (days), in a 30-year old female. The subject was simulated to have a daily exposure of ZEN of 0.25 ng/kg bw, corresponding to the TDI for 7 days.

Table S8: QIVIVE results

<i>C_{max}</i> (μM)	<i>Oral Equivalent Dose (μg/kg bw/day)</i>			Initial assay used to derive the EC50	EC50 reference study
	ZEN	ZEN + α-ZEL	ZEN + ZEN-GlcA + α-ZEL + α-ZEL-GlcA		
0.00006	24.48	2.19	3.00E-02	Alkaline phosphatase assay (ishikawa cells)	(Le Guevel and Pakdel, 2001)
0.00031	130.84	11.68	0.15	MCF-7 bioassay	(Minervini et al., 2005)
0.00036	151.52	13.53	0.17	Alkaline phosphatase assay (ishikawa cells)	(Grgic et al., 2022)
0.00049	206.81	18.47	0.23	ERα_CALUX	(Ehrlich et al., 2015)
0.00158	666.86	59.56	0.74	ERα_CALUX	(Mendez-Catala et al., 2020)
0.00160	675.31	60.31	0.75	MMV-LUC cell line	(Frizzell et al., 2011)
0.00164	691.77	61.78	0.77	MCF-7 bioassay	(Malekinejad et al., 2006)
0.00381	1608.07	143.61	1.79	MCF-7 bioassay	(Molina-Molina et al., 2014)

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