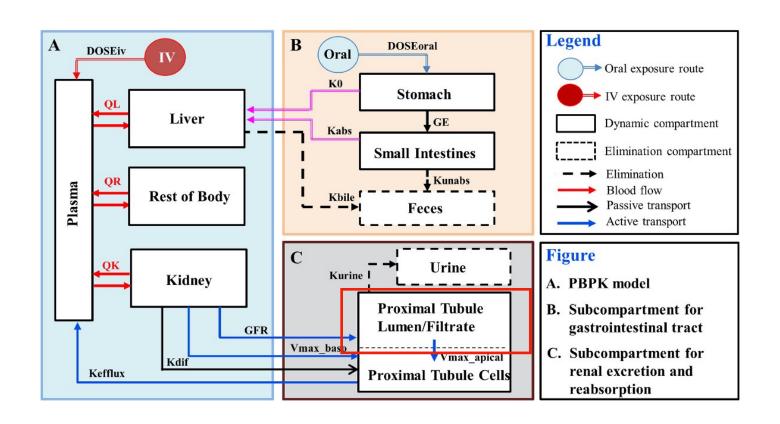
Physiologically based pharmacokinetic model

dxdt AUCCL = CL;

ODE system: Time varying state

```
// Concentrations in the tissues and in the capillary blood of the tissues
double CA free = APlas free/VPlas;
                                        // mg/L, Free PFOSC concentration in the plasma
double CA = CA free/Free;
                                        // mg/L, Concentration of total PFOS in plasma
double CL = AL/VL;
                                        // mg/L, Concentration of parent drug in the tissue of liver
double CVL = CL/PL;
                                        // mg/L, Concentration of parent drug in the capillary blood of liver
double CKb = AKb/VKb;
                                        // mg/L, Concentration of PFOS in Kidney blood
double CVK = CKb/PK;
double CK = CVK*PK;
                                        // mg/L, Concentration of PFOS in Kidney compartment
double CRest = ARest/VRest;
                                        // mg/L, Crest drug concentration in the rest of the body (mg/L)
double CVRest = CRest/PRest;
                                        // mg/L, Concentration of parent drug in the capillary blood of the rest of body
// Kidney compartment plus 2 subcompartment (Proximal Tubule cells: PTC, Filtrate: Fil)
// Concentration in kidney, PTC and fil
double CPTC = APTC/VPTC;
                                        // mg/L, Concentration of PFOS in PTC blood
double Cfil = AFil/Vfil;
                                        // mg/L, Concentration of PFOS in FIL blood
// Virtual compartment:
// Basolateral transport : (baso)
// Diffusion
                        : (dif)
// Apical transport
                        : (apical)
                        : (efflux)
// efflux clearance
// Clearance via glormerular filtration (CL)
double RA baso = (Vmax baso*CKb)/(Km baso+CKb);
                                                          // mg/h, Rate of basolateral transporters
dxdt A baso = RA baso;
                                                           // mg, Amount of basolateral transporters
double RA apical = (Vmax apical*Cfil)/(Km apical + Cfil); // mg/h, Rate of apical transporter
                                                           // mg, Amount of apical transporter
dxdt_A_apical = RA_apical;
double Rdif = Kdif*(CKb - CPTC);
                                                          // mg/h, Rate of diffusion from into the PTC
dxdt Adif = Rdif;
                                                          // mg, Amount moved via glomarular filtration
double RAefflux = Kefflux*APTC;
                                                          // mg/h, Rate of efflux clearance rate from PTC to blood
dxdt Aefflux = RAefflux;
                                                           // mg, Amount of efflux clearance rate from PTC to blood
double RCI = CA*GFR*Free;
                                                    // mg/h, Rate of clearance (CL) to via glormerular filtration (GFR)
                                                           // mg, Amount of clearance via GFR
dxdt ACI = RCI;
```



```
// PFOS in plasma
double RPlas free = (QRest*CVRest*Free)+(QK*CVK*Free)+(QL*CVL*Free)-(QC*CA*Free)+RAefflux; // mg/h, Rate of change in the plasma
double RPlas = (QRest*CVRest)+(QK*CVK)+(QL*CVL)-(QC*CA);
dxdt APlas free = RPlas free;
                                                     // mg, Amount in the plasma
dxdt APlas = RPlas;
                                                     // mg, Amount in the plasma
dxdt AUCCA free = CA free;
                                                     // mg*h/L, Area under curve of PFOS in liver compartment
// Proximal Tubule Cells (PTC)
double RPTC = Rdif + RA apical + RA baso - RAefflux; // mg/h, Rate of change in PTC
dxdt APTC = RPTC;
                                                     // mg, Amount moved in PTC
dxdt AUCCPTC = CPTC;
                                                     // mg*h/L, Area under curve of PFOS in the compartment of PTC
// Proximal Tubule Lumen/ Filtrate (Fil)
double Rfil = CA*GFR*Free - RA apical - AFil*Kurine; // mg/h, Rate of change in Fil
dxdt AFil = Rfil;
                                                     // mg, Amount moved in Fil
dxdt AUCfil = Cfil;
                                                     // mg*h/L, Area under curve of PFOS in the compartment of Fil
// Urine elimination
double Rurine = Kurine*AFil:
                                                     // mg/h, Rate of change in urine
dxdt Aurine = Rurine;
                                                     // mg, Amount in urine
double Curine = Rurine/Kvoid; //
// Kidney compartment
double RKb = QK*(CA-CVK)*Free - CA*GFR*Free - Rdif - RA baso; // mg/h, Rate of change in Kidney compartment
dxdt AKb = RKb;
                                                              // mg, Amount in kidney compartment
                                                              // mg*h/L, Area under curve of PFOS in the Kidney compartment
dxdt AUCKb = CK;
// PFOS in the compartment of rest of body, flow-limited model
double RRest = QRest*(CA-CVRest)*Free;
                                                      // mg/h, Rate of change in rest of body
dxdt ARest = RRest;
                                                      // mg, Amount in rest of body
dxdt AUCCRest = CRest;
                                                      // mg*h/L, Area under curve of PFOS in the compartment of rest of body
// Gastrointestinal (GI) tract
// Stomach compartment
double RST = - K0*AST - GE*AST;
                                                       // mg/h, Rate of chagne in Stomach compartment
                                                       // mg, Amount in Stomach
dxdt AST = RST;
                                                       // mg/h, Rate of absorption in the stomach
double RabsST = K0*AST;
dxdt AabsST = RabsST;
                                                       // mg, Amount absorbed in the stomach
// Small intestine compartment
double RSI = GE*AST - Kabs*ASI - Kunabs*ASI;
                                                       // mg/h, Rate of chagne in Small intestine compartment
dxdt ASI = RSI;
                                                       // mg, Amount in Small intestine
double RabsSI = Kabs*ASI;
                                                       // mg/h, Rate of absorption in the Small intestine
                                                       // mg, Amount absorbed in the Small intestine
dxdt AabsSI = RabsSI;
double Total oral uptake = AabsSI + AabsST;
                                                       // mg, Total oral uptake in the GI
// Biliary excretion
double Abile = Kbile*AL;
                                                       // mg, Amount of PFOS in bile excretion
double amount per gram liver = (AL/ML)*1000;
                                                       // ug/g, Amount of PFOS in liver per gram liver
// Feces compartment
                                                       // mg/h, Rate of change in feces compartment
double Rfeces = Kbile*AL+ Kunabs*ASI;
                                                       // mg, Amount of the feces compartment
dxdt_Afeces = Rfeces;
// PFOS in liver compartment, flow-limited model
double RL = QL*(CA-CVL)*Free - Kbile*AL + Kabs*ASI + K0*AST; // mg/h, Rate of chagne in liver compartment
dxdt AL = RL;
                                                             // mg, Amount in liver compartment
```



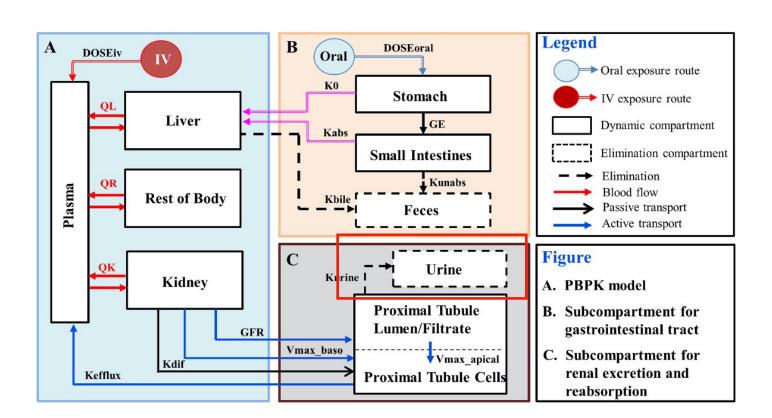
// mg*h/L, Area under curve of PFOS in liver compartment

Physiologically based pharmacokinetic model

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ODE system: Time varying state

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double CL = AL/VL;
                                        // mg/L, Concentration of parent drug in the tissue of liver
double CVL = CL/PL;
                                        // mg/L, Concentration of parent drug in the capillary blood of liver
double CKb = AKb/VKb;
                                        // mg/L, Concentration of PFOS in Kidney blood
double CVK = CKb/PK;
double CK = CVK*PK;
                                        // mg/L, Concentration of PFOS in Kidney compartment
double CRest = ARest/VRest;
                                        // mg/L, Crest drug concentration in the rest of the body (mg/L)
double CVRest = CRest/PRest;
                                        // mg/L, Concentration of parent drug in the capillary blood of the rest of body
// Kidney compartment plus 2 subcompartment (Proximal Tubule cells: PTC, Filtrate: Fil)
// Concentration in kidney, PTC and fil
double CPTC = APTC/VPTC;
                                        // mg/L, Concentration of PFOS in PTC blood
                                        // mg/L, Concentration of PFOS in FIL blood
double Cfil = AFil/Vfil;
// Virtual compartment:
// Basolateral transport : (baso)
// Diffusion
                        : (dif)
// Apical transport
                        : (apical)
                        : (efflux)
// efflux clearance
// Clearance via glormerular filtration (CL)
double RA baso = (Vmax baso*CKb)/(Km baso+CKb);
                                                          // mg/h, Rate of basolateral transporters
dxdt A baso = RA baso;
                                                           // mg, Amount of basolateral transporters
double RA apical = (Vmax apical*Cfil)/(Km apical + Cfil); // mg/h, Rate of apical transporter
                                                           // mg, Amount of apical transporter
dxdt_A_apical = RA_apical;
double Rdif = Kdif*(CKb - CPTC);
                                                          // mg/h, Rate of diffusion from into the PTC
dxdt Adif = Rdif;
                                                          // mg, Amount moved via glomarular filtration
double RAefflux = Kefflux*APTC;
                                                          // mg/h, Rate of efflux clearance rate from PTC to blood
dxdt Aefflux = RAefflux;
                                                           // mg, Amount of efflux clearance rate from PTC to blood
double RCI = CA*GFR*Free;
                                                    // mg/h, Rate of clearance (CL) to via glormerular filtration (GFR)
                                                           // mg, Amount of clearance via GFR
dxdt ACI = RCI;
```



```
// PFOS in plasma
double RPlas free = (QRest*CVRest*Free)+(QK*CVK*Free)+(QL*CVL*Free)-(QC*CA*Free)+RAefflux; // mg/h, Rate of change in the plasma
double RPlas = (QRest*CVRest)+(QK*CVK)+(QL*CVL)-(QC*CA);
dxdt APlas free = RPlas free;
                                                     // mg, Amount in the plasma
dxdt APlas = RPlas;
                                                     // mg, Amount in the plasma
dxdt AUCCA free = CA free;
                                                     // mg*h/L, Area under curve of PFOS in liver compartment
// Proximal Tubule Cells (PTC)
double RPTC = Rdif + RA apical + RA baso - RAefflux; // mg/h, Rate of change in PTC
dxdt APTC = RPTC;
                                                     // mg, Amount moved in PTC
                                                     // mg*h/L, Area under curve of PFOS in the compartment of PTC
dxdt AUCCPTC = CPTC;
// Proximal Tubule Lumen/ Filtrate (Fil)
double Rfil = CA*GFR*Free - RA apical - AFil*Kurine; // mg/h, Rate of change in Fil
dxdt AFil = Rfil;
                                                     // mg, Amount moved in Fil
dxdt AUCfil = Cfil;
                                                     // mg*h/L, Area under curve of PFOS in the compartment of Fil
// Urine elimination
double Rurine = Kurine*AFil:
                                                     // mg/h, Rate of change in urine
                                                     // mg, Amount in urine
dxdt Aurine = Rurine;
double Curine = Rurine/Kvoid; //
// Kidney compartment
double RKb = QK*(CA-CVK)*Free - CA*GFR*Free - Rdif - RA baso; // mg/h, Rate of change in Kidney compartment
dxdt AKb = RKb;
                                                              // mg, Amount in kidney compartment
                                                              // mg*h/L, Area under curve of PFOS in the Kidney compartment
dxdt AUCKb = CK;
// PFOS in the compartment of rest of body, flow-limited model
double RRest = QRest*(CA-CVRest)*Free;
                                                      // mg/h, Rate of change in rest of body
dxdt ARest = RRest;
                                                      // mg, Amount in rest of body
dxdt AUCCRest = CRest;
                                                      // mg*h/L, Area under curve of PFOS in the compartment of rest of body
// Gastrointestinal (GI) tract
// Stomach compartment
double RST = - K0*AST - GE*AST;
                                                       // mg/h, Rate of chagne in Stomach compartment
                                                       // mg, Amount in Stomach
dxdt AST = RST;
                                                       // mg/h, Rate of absorption in the stomach
double RabsST = K0*AST;
dxdt AabsST = RabsST;
                                                       // mg, Amount absorbed in the stomach
// Small intestine compartment
double RSI = GE*AST - Kabs*ASI - Kunabs*ASI;
                                                       // mg/h, Rate of chagne in Small intestine compartment
dxdt ASI = RSI;
                                                       // mg, Amount in Small intestine
double RabsSI = Kabs*ASI;
                                                       // mg/h, Rate of absorption in the Small intestine
dxdt AabsSI = RabsSI;
                                                       // mg, Amount absorbed in the Small intestine
double Total oral uptake = AabsSI + AabsST;
                                                       // mg, Total oral uptake in the GI
// Biliary excretion
double Abile = Kbile*AL;
                                                       // mg, Amount of PFOS in bile excretion
double amount per gram liver = (AL/ML)*1000;
                                                       // ug/g, Amount of PFOS in liver per gram liver
// Feces compartment
                                                       // mg/h, Rate of change in feces compartment
double Rfeces = Kbile*AL+ Kunabs*ASI;
dxdt Afeces = Rfeces;
                                                       // mg, Amount of the feces compartment
// PFOS in liver compartment, flow-limited model
double RL = QL*(CA-CVL)*Free - Kbile*AL + Kabs*ASI + K0*AST; // mg/h, Rate of chagne in liver compartment
dxdt AL = RL;
                                                             // mg, Amount in liver compartment
```



// mg*h/L, Area under curve of PFOS in liver compartment