

Prediction of human fetal pharmacokinetics using *ex vivo* human placenta perfusion studies and physiologically based models

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WHAT IS ALREADY KNOWN ABOUT THIS SUBJECT

- Pregnant women and fetuses are orphan populations with respect to the safety and efficacy of drugs.
- The cotyledon perfusion experiment is the gold standard that provides an insight into placental transfer. After maternal intake, fetal exposure is estimated by cord blood sampling.
- There is no validated method that is able to predict fetal drug concentrations.

WHAT THIS SUBJECT ADDS

- Transplacental transfer parameters ([i.e. diffusion (D_{cot}), elimination constant (k_{PE}) and placental partition coefficient (K_{pP})] were estimated from the cotyledon perfusion model.
- A novel approach to predict fetal drug exposure quantitatively, incorporating estimated transplacental transfer parameters in p-PBPK models, was proposed and validated by comparing predictions to *in vivo* observations.

AIMS

Pregnant women can be exposed to numerous drugs during the gestational period. For obvious ethical reasons, *in vivo* studies of fetal exposure to drugs are limited. Information about the transplacental transfer of drugs prior to their administration to pregnant women would be highly useful. In the present study, a novel approach was developed quantitatively predict or to predict the fetal exposure to drugs administered to the mother quantitatively.

METHODS

Transplacental parameters estimated from *ex vivo* human placenta perfusion experiments were implemented in pregnancy-physiologically based pharmacokinetic (p-PBPK) models in order to predict fetal PK. Thereafter, fetal PK profiles for two antiretroviral drugs, tenofovir (TFV) and emtricitabine (FTC) were simulated. These predictions were then compared to observed cord blood concentrations, to validate these models.

RESULTS

Parameters obtained from the *ex vivo* experiments enabled a good prediction of observed cord blood concentrations without additional a scaling factor. Moreover, a sensitivity analysis showed that fetal predictions were sensitive to changes in transplacental parameters values obtained *ex vivo*.

CONCLUSION

The integration of *ex vivo* human placental perfusion parameters in a p-PBPK model should be a promising new approach for predicting human fetal exposure to xenobiotics.

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Introduction

Drug prescriptions and over-the-counter medications are common in pregnancy, and the average pregnant patient in the US and Canada uses more than two drugs during the course of their pregnancy [1]. However, pregnant women and fetuses are orphan populations with respect to the safety and efficacy of drugs. Fetal toxicity and efficacy are thought to depend both upon the maternal-to-fetal transfer of drugs [pharmacokinetics (PK)] and intrinsic toxicity [pharmacodynamics (PD)].

Assessing drug transport across the human placental barrier is mandatory in order to guarantee drug safety during pregnancy [2]. However, for obvious ethical reasons, *in vivo* fetal risk assessment studies related to maternal drug exposure remain extremely limited. Some studies have evaluated fetal exposure using cord blood plasma samples. Although the cord-to-maternal concentration ratio is informative as an index of relative fetal drug exposure, it is highly variable due to the various delays between drug administration and blood sampling [3, 4]. Population PK analyses enable fetal PK to be estimated but usually require a large number of exposed patients [5, 6]. To ensure drug safety during pregnancy, information about transplacental transfer prior to administration would be highly desirable. As animal studies may not be helpful for predicting human fetal PK because of interspecies differences in the structural and functional features of the placenta, other models have been developed. The *ex vivo* human placental perfusion model is the gold standard and offers a better insight into the various placental drug transporters, xenobiotic metabolism and tissue binding. Nevertheless, this method cannot directly predict fetal PK profiles.

The present study presents a novel approach predicting drug fetal exposure quantitatively. Transplacental parameter estimated from the *ex vivo* human placenta perfusion model were implemented in pregnancy-physiologically based PK (p-PBPK) models in order to predict fetal PK. Physiologically based PK (PBPK) models are based to a large extent on the actual physiology of the organism, whereas conventional PK models use virtual compartments. PBPK models incorporate both physiological parameters that are important for absorption, distribution, metabolism and excretion processes and drug-specific parameters. These models have already been used to predict PK profiles in specific populations, such as pregnant women [7–11]. In regard to their structure, PBPK models are fully suitable to incorporate *ex vivo* data. The aim of the present study was to evaluate this new method to simulate the fetal PK of two antiretroviral drugs, tenofovir (TFV) and emtricitabine (FTC). These simulations were compared to observed cord plasma concentrations, to validate our models.

Materials and methods

Figure 1 shows the workflow of the present study. Briefly, we had previously developed PBPK models for FTC and TFV. When the models were able to accurately describe the PK for different routes of administration and dosing regimens in nonpregnant adults, we implemented the physiological changes occurring during pregnancy. PK simulations were then compared to observed concentrations from pregnant women. Thereafter, the fetoplacental compartment was used in the model, with the placenta,

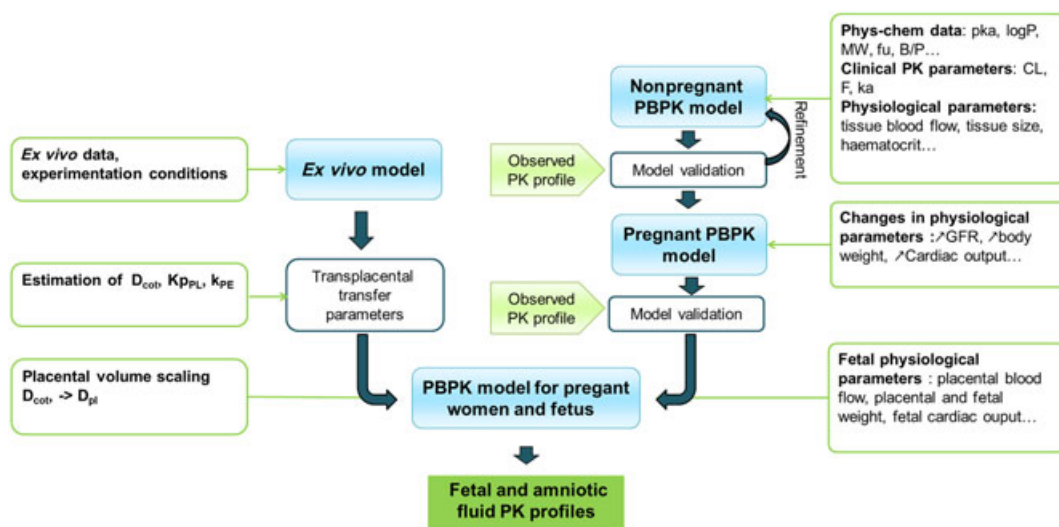


Figure 1

Schematic representation of the workflow of physiologically based pharmacokinetic (PBPK) model development. B/P, blood to plasma ratio; CL, clearance; D_{cot} , diffusion (cotyledon); D_{pl} , diffusion (placenta); F, bioavailability; f_u , free fraction; GFR, glomerular filtration rate; k_a , absorption rate constant; K_{pPL} , placental partition coefficient, k_{PE} , placental elimination; MW, molar mass; Phys-chem, physicochemical; PK, pharmacokinetic

amniotic fluid and fetus considered as separate compartments. From the *ex vivo* experiments on cotyledons, the transplacental transfer parameters [diffusion (D_{cot}), placental elimination constant (k_{PE}) and placental partition coefficient (K_{p_i})] were estimated. After scaling D_{cot} by placental weight, they were implemented in the PBPK models. Finally, simulated fetal and amniotic fluid PK profiles were compared to observed *in vivo* data.

PBPK modelling in the nonpregnant population

In a previous study, we built up whole-body PBPK models for pregnant and nonpregnant adults for FTC and TFV by using the Simcyp® software [12]. To study transplacental transfer, simplified PBPK models had to be coded for the R software [13]. R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <http://www.R-project.org/> [14]. Full PBPK models with first-order rate absorption were used. To code our models in R language, only the physiological parameters were checked. Drug-specific parameters used in the previously developed models in Simcyp® remained unchanged (Table 1). For each step of modelling, simulated PK profiles were compared to observed data and PK profiles previously simulated by Simcyp®. The physiological parameters used for these models are summarized in Table 2 [15–18].

PBPK modelling in pregnant women

Figure 2 shows the PBPK model. The distinct placental and fetal compartments enable *ex-vivo* transplacental transfer parameters to be incorporated. Moreover, as amniotic fluid can affect fetal exposure, this compartment was added to the model. As the two drugs are poorly bound to plasma proteins, the free fractions (f_u) were assumed to be unchanged during pregnancy. Fetal f_u values were assumed to be equal to maternal ones (Table 1). Numerous types of exchange can affect fetal

Table 1

Input values

	Tenofovir	Emtricitabine
MW (g mol ⁻¹)	287.21 [19]	247.25 [20]
pKa	3.7-6.5 [19]	2.65 [20]
log P	-2.21 [21]	-0.43 [20]
F	0.18 [22]	0.93 [20]
ka (h ⁻¹)	0.56 [23]	0.54 [5]
f _u	0.993 [22]	0.96 [20]
B:P ratio	0.58 [24]	1 [20]
Total CL (l h ⁻¹)	14.2 [25]	18 [26]
CL _R (l h ⁻¹)	10.6 [25]	13 [20, 26]

B:P ratio, blood-to-plasma ratio; CL, clearance; CL_R, renal clearance; F, bioavailability; f_u, free fraction; ka, first-order absorption rate; MW, molecular weight; P, partition coefficient; pKa, acid dissociation constant at logarithmic scale.

Table 2

Mean adult physiological parameters

	Men	Women	Pregnant women (GA = 40)
Height (cm)	175	165	165
Body weight (kg)	80	60	74.8
Haematocrit	0.45*	0.39*	0.33*,‡
Glomerular filtration rate (l h ⁻¹)	6.1†	4.8†	6.4†,‡
Organ volumes (l)			
Adipose	20.2*	20.2*	27.9*,‡
Bones	3.6*	2.1*	2.1*
Blood	5.8*	4.2*	6.1*,‡
Lung	1.3*	0.8*	0.8*
Brain	1.4*	1.4*	1.4*
Heart	0.4*	0.3*	0.3*
Kidney	0.3*	0.3*	0.3*
Muscle	32.1*	19.6*	19.6*
Skin	3.8*	3.2*	3.2*
Liver	1.5*	1.2*	1.2*
Spleen	0.1*	0.1*	0.1*
Pancreas	0.1*	0.1*	0.1*
Gut	1.3*	1.1*	1.1*
Blood flows rates (l h ⁻¹)			
Cardiac output	359.8	296.8	381.2
Adipose	24.2*	24.3*	33.5*
Bones	17.5*	10.4*	10.4*
Brain	43.0*	42.5*	42.5*
Heart	15.7*	12.6*	12.6*
Kidney	74.4*	61.6*	62.5*
Muscle	57.8*	35.3*	35.6*
Skin	27.4*	23.0*	23.0*
Liver	90.8*,¶	81.3*,¶	108.7*,¶,§
Spleen	7.8*	7.8*	7.8a
Pancreas	3.8*	3.0*	3.0*
Gut	58.4*	50.8*	78.2*, d

GA, gestational age. *Price et al. [15]. †Peters et al. [16]. ‡Abduljalil et al. [17]. §Clapp et al. [19]. ¶Simcyp®, healthy population.

and amniotic fluid PK. To study fetal-to-placental exchange, the fetal compartment was split into fetal blood and other fetal tissues (Figure 2). This exchange is driven by the blood flow rate between the placenta and the fetus (Q_{plaF}). In late pregnancy, one-fifth of the fetal cardiac output is distributed to the placenta, with the remainder distributed to the rest of the fetal body (Q_{rbf}) [27]. The main exchanges between the amniotic space and the surrounding tissues in late pregnancy come from fetal urine, fetal swallowing (k_{sw}), the intramembranous pathway from the amniotic cavity to the fetal circulation (k_{INT}) and pulmonary excretion (k_L) [28–31]. Fetal renal excretion was indexed on fetal glomerular filtration rate (GFR_f) [32]. The two drugs studied are poorly metabolized, so this elimination pathway was assumed to be negligible for fetus [33]. The maternal physiological changes occurring during pregnancy that can affect the distribution

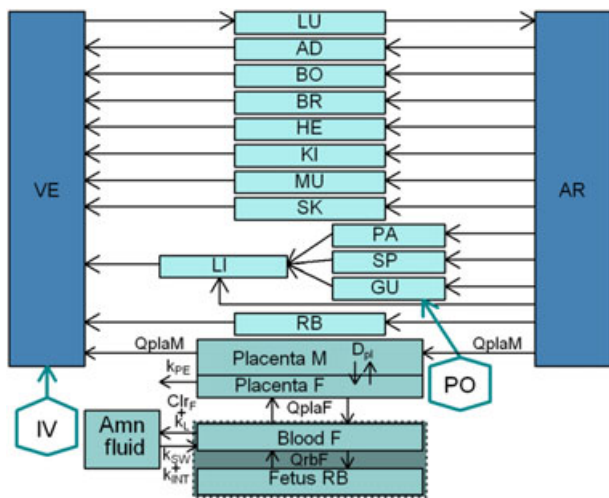


Figure 2

Schematic representation of physiologically based pharmacokinetic (PBPK) models used. AD, adipose; Amn fluid, amniotic fluid; AR, arterial blood; BO, bones; BR, brain; CLRF, fetal urinary excretion; Dpl, diffusion; F, fetus; GU, gut; HE, heart; IV, intravenous; KI, Kidney; k_{int} , intramembranous pathway; k_l , oral, nasal, tracheal and pulmonary secretion constant; k_{pe} , placental elimination; k_{sw} , swallowing constant; LI, liver; LU, lung; M, mother; MU, muscle; PA, pancreas; PO, per os; Qplaf, blood flow from the fetus to the placental tissue; Qplam, blood flow from the mother to the placental tissue; Qrbf, blood flow to the fetal body; RB, rest of the body; SK, skin; SP, spleen; VE, venous blood

and elimination are implemented in this model [12]. All of the basal values of the nonpregnant population, except for portal vein blood flow, were modified according to the gestational age (GA) of the pregnant population, as described by Abduljalil *et al.* (Table 2) [17, 18]. All fetal physiological constants are reported in Table 3. A sensitivity analysis was performed on some parameters to evaluate their impact on fetal and amniotic fluid PK.

Ex vivo model

Placenta tissue collections. Thirty-four placentas from normal pregnancies were obtained from Port Royal Hospital (Paris, France) after uncomplicated vaginal delivery or caesarean section. All mothers were seronegative for HIV infection, were not infected by hepatitis B or C viruses and took no medication other than oxytocin or epidural anaesthesia during labour. All placentas were obtained after a full-term pregnancy (GA from 37 to 41 weeks and 4 days). Written informed consent was obtained for all participants in the study.

Placental perfusion. Placentas were perfused in a recirculating (closed-closed) circuit, according to a method adapted from those of Schneider *et al.* [38] and Forestier *et al.* [39]. Perfusion experiments started within 30 min after delivery. After a visual examination for lack of evident lesions on the chorionic plate, a truncal branch of the chorionic artery and the associated vein were cannulated. The fetal circulation

Table 3

Fetal physiological parameter values

(GA = 40)	Parameters	Ref.
Maternal-to-placental blood flow, Q_{plaM} ($l\ h^{-1}$)	46.5	Abduljalil <i>et al.</i> [17]
Fetal-to-placental blood flow, Q_{plaF} ($l\ h^{-1}$)	14.3	Kiserud <i>et al.</i> [27]
Fetal cardiac output, Q_{ca} ($l\ h^{-1}$)	85.5	Kiserud <i>et al.</i> [27]
Placental weight, V_{pla} (kg)	0.65	Abduljalil <i>et al.</i> [17]
Fetal weight, V_{fo} (kg)	3.56	Abduljalil <i>et al.</i> [17]
Amniotic fluid volume, V_{amf} (l)	0.86	Abduljalil <i>et al.</i> [17]
Fetal blood volume, V_{bloodF} (l)	0.24	Smith <i>et al.</i> [34]
Fetal haematocrit	0.5	Zanardo <i>et al.</i> [35], Chang <i>et al.</i> [36], Eskoka <i>et al.</i> [37]
Fetal glomerular filtration rate, GFR_f ($l\ h^{-1}$)	0.136	Arant <i>et al.</i> [32]
Swallowing volume, k_{sw}	0.8	Underwood <i>et al.</i> [29]
Secretion of oral, nasal, tracheal and pulmonary fluids, k_l (l/day^{-1})	0.126	Underwood <i>et al.</i> [29]
Intramembranous pathway, k_{int} $l\ day^{-1}$	0.35	Underwood <i>et al.</i> [29]

GA, gestational age.

was established at a flow rate of $6\ ml\ min^{-1}$ (Q_f). After confirmation of the absence of vascular leakage, the perfused area progressively whitened and enabled visualization of the selected cotyledon. The perfusion was subsequently initiated by insertion of two catheters into the intervillous space on the maternal side. The maternal circulation was established at a flow rate of $12\ ml\ min^{-1}$ (Q_m). The pHs of maternal and fetal solutions, prepared using Earle medium containing $30\ g\ l^{-1}$ and $40\ g\ l^{-1}$ of human serum albumin, were adjusted to 7.4 ± 0.1 and 7.2 ± 0.1 , respectively. The validation of the cotyledon's viability during the experiment was carried out using antipyrine ($20\ mg\ l^{-1}$). TFV, FTC and antipyrine were perfused into the maternal reservoir. Maternal and fetal reservoir volumes (V_m and V_f , respectively) were 200 ml and 250 ml, respectively. Samples were collected every 10 min during the first half-hour and then every 30 min until 150 min from both the fetal and maternal sides to determine the concentrations (C_f and C_m , respectively). Samples were then stored at $-20^\circ C$ until analysis.

Sample analysis. TFV and FTC concentrations were determined using ultra performance liquid chromatography coupled with tandem mass spectrometry (UPLC-MS/MS), using the Acquity UPLC/TQD (Applied Biosystems, Foster City, CA, USA) [40]. Antipyrine concentrations were determined by high-performance liquid chromatography with ultraviolet detection at 290 nm [41].

The maternal-to-fetal transfer was described by the fetal transfer rate (FTR). It was calculated as follows:

Equation 1:

$$\text{FTR} = (C_f * V_f) * 100 / [(C_f * V_f) + (C_m * V_m)]$$

where C_f and C_m are the drug concentrations in fetal and maternal perfusates, and V_f and V_m are the fetal and maternal perfusate volumes. An FTR of antipyrene $>20\%$ was required to validate each experiment. Clearance index (CLI) was calculated as the ratio of the TFV or FTC FTR to that of antipyrene. The FTR and CLI parameters are only useful for comparison purposes between drugs and are not applied in the PBPK model.

Estimation of transplacental transfer parameters

Drug transfer across the placenta was modelled as a cotyledon split into maternal and fetal compartments (Figure 3). The cotyledon volume averaged 58 ml. Maternal cotyledon volume (V_{mp}) was assumed to be 23 ml [42]. Several transplacental transfer models for TFV and FTC were investigated – i.e. simple diffusion, linear transfer, saturable transfer and addition of placental elimination rate or tissue protein binding. Data were analysed with a nonlinear mixed-effect modelling approach, using NONMEM program version 6.2 (Icon Development Solutions, Ellicott City, MD, USA). Several error models

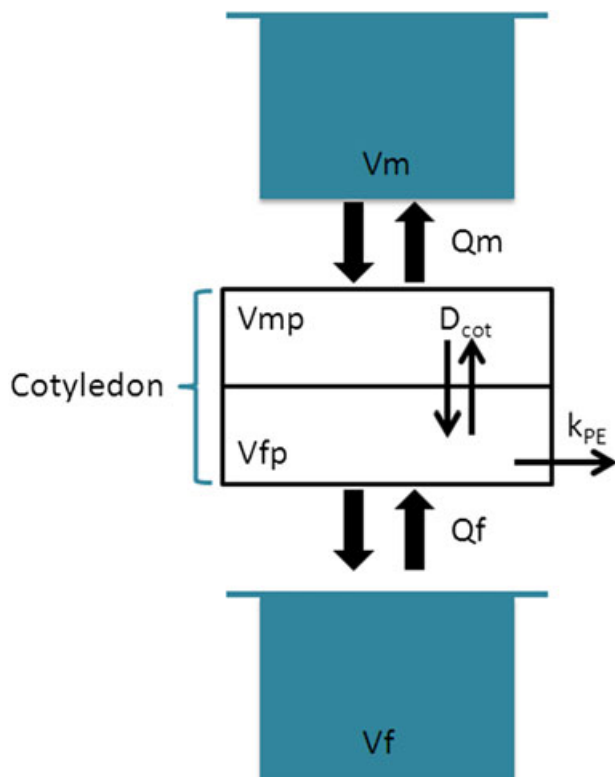


Figure 3

Schematic representation of the *ex vivo* model used for tenofovir (TFV) and emtricitabine (FTC) (recirculating circuit). D_{cot} , diffusion parameter; f, fetus; fp, fetal placenta; k_{PE} , placental elimination constant (h^{-1}); m, mother; mp, maternal placenta; Q, flow rate ($l h^{-1}$); V, volume (l)

(proportional, additive, mixed) were investigated to describe the residual variability (ϵ). The objective function value (OFV) was used to test different hypotheses. A model was kept if the OFV was decreased by at least 3.84 (chi-square test with one degree of freedom). For evaluation of the goodness of fit, the following graphs were performed: observed concentrations vs. predictions, weighted residuals vs. time, and weighted residuals vs. predictions. Diagnostic graphics and distribution statistics were performed using R for Nonmem (RfN) (<http://wfn.sourceforge.net>) from the R program. Simulated concentrations in the maternal and fetal reservoirs were compared to concentrations from the *ex vivo* experiment previously described.

As an example, equations 2–5 were used to describe the *ex vivo* experiment considering passive transfer (D_{cot}) and placental elimination (k_{PE}).

Equation 2: maternal reservoir

$$\frac{dC_m}{dt} = \frac{Q_m}{V_m} * \left(\frac{C_{mp}}{K_{p_{pl}}} - C_m \right)$$

Equation 3: maternal cotyledon

$$\frac{dC_{mp}}{dt} = \frac{(Q_m * (C_m - C_{mp}/K_{p_{pl}}) - D_{cot} * (C_{mp} - C_{fp}))}{V_{mp}}$$

Equation 4: fetal cotyledon

$$\frac{dC_{fp}}{dt} = \frac{(Q_f * (C_f - \frac{C_{fp}}{K_{p_{pl}}}) + D_{cot} * (C_{mp} - C_{fp}) - k_{PE} * C_{fp} * V_{fp})}{V_{fp}}$$

Equation 5: fetal reservoir

$$\frac{dC_f}{dt} = \frac{Q_f}{V_f} * \left(\frac{C_{fp}}{K_{p_{pl}}} - C_f \right)$$

where C denotes a concentration ($mg l^{-1}$), Q a flow rate ($l h^{-1}$) and V a volume (l). Subscripts m, f and p denote mother, fetus and placenta, respectively. $K_{p_{pl}}$ is the placental partition coefficient, D_{cot} is the diffusion parameter ($l h^{-1}$) and k_{PE} is the placental elimination parameter (h^{-1}). $K_{p_{pl}}$, D_{cot} and k_{PE} were estimated.

In vivo fetal and amniotic fluid simulations

The best model describing the transplacental transfer in *ex vivo* experimentation was implemented in the p-PBPK model. Cotyledon volume and experimental flow rates were replaced by placental volume and *in vivo* perfusion rates (Table 3). Transplacental transfer parameters estimated from *ex vivo* experiments were more integrated. The *in vivo* diffusion was related to

the *ex vivo* parameters weighed by their respective volumes as shown in Equation 6.

Equation 6: *in vivo* diffusion parameter

$$D_{pl} = \frac{D_{cot} \times V_{pl}}{V_{cot}}$$

where D_{pl} and D_{cot} stand for the *in vivo* and *ex vivo* diffusion parameters, and V_{pl} and V_{cot} for the placental and cotyledon volumes.

The equations describing the fetoplacental compartments of the p-PBPK models are shown below.

Equation 7: *in vivo* maternal placenta

$$\frac{dC_{plam}}{dt} = \frac{(Q_{plam} * (C_{ab} - \frac{C_{plam}}{K_{ppl}} * B/P) - D_{pl} * (C_{plam} - C_{plaf}))}{V_{plam}}$$

Equation 8: *in vivo* fetal placenta

$$\frac{dC_{plaf}}{dt} = \frac{(Q_{plaf} * (C_{bloodF} - \frac{C_{plaf}}{K_{ppl}} * B/P) + D_{pl} * (C_{plam} - C_{plaf}) - k_{PE} * C_{plaf} * V_{plaf})}{V_{plaf}}$$

Equation 9: fetal blood

$$\begin{aligned} \frac{dC_{bloodF}}{dt} = & \frac{Q_{plaf}}{V_{bloodF}} * \left(\frac{C_{plaf}}{K_{ppl}} * B/P - C_{bloodF} \right) \\ & + \frac{Q_{rbF}}{V_{bloodF}} * \left(\frac{C_{rbF}}{K_{prb}} * B/P - C_{bloodF} \right) \\ & - \left(Cl_r * \frac{GFR_f}{GFR} * \frac{fu}{B/P} - k_L \right) * \frac{C_{bloodF}}{V_{bloodF}} \\ & + (k_{INT} + k_{SW}) * \frac{C_{amf}}{V_{bloodF}} \end{aligned}$$

Equation 10: fetal body

$$\frac{dC_{rbF}}{dt} = \frac{Q_{rbF}}{V_{rbF}} * \left(C_{bloodF} - \frac{C_{rbF}}{K_{prb}} * B/P \right)$$

Equation 11: amniotic fluid

$$\begin{aligned} \frac{dC_{amf}}{dt} = & \left(Cl_r * \frac{GFR_f}{GFR} * \frac{fu}{B/P} - k_L \right) * \frac{C_{bloodF}}{V_{amf}} \\ & - (k_{INT} + k_{SW}) * \frac{C_{amf}}{V_{amf}} \end{aligned}$$

where Q denotes blood flow ($l\ h^{-1}$), V tissue volume (l), B/P the blood-to-plasma concentration ratio, D_{pl} a diffusion parameter ($l\ h^{-1}$), GFR the glomerular filtration rate ($l\ h^{-1}$), Cl_r the maternal renal clearance ($l\ h^{-1}$), fu the free

drug fraction, k_{INT} the intramembranous constant ($l\ h^{-1}$), k_{SW} the swallowing constant ($l\ h^{-1}$) and k_L the oral, nasal, tracheal and pulmonary secretion constant ($l\ h^{-1}$). The subscripts ab , $plam$, $plaf$, $bloodF$, rbF and amf denote maternal blood, maternal placenta, fetal placenta, fetal blood, rest of the fetal body (fetal body – fetal blood) and amniotic fluid.

Women in labour had smaller maximal concentrations (C_{max}) compared to pregnant women, so changes in absorption were assumed to occur during labour for the two drugs. Therefore, absorption rates were reduced by 50% and the FTC bioavailability was fixed at 0.75.

Model validation

Models were validated by comparing maternal, fetal and amniotic fluid concentration simulations to observed data from women in labour. We simulated the administration of a single dose of 400 mg of FTC or 600 mg of TFV to an average 35-year-old patient with a GA of 39 weeks. We compared the simulated maternal and fetal concentrations to the observed concentrations found by Hirt *et al.* [5, 6]. In the Tenofovir/Emtricitabine in Africa and Asia (TEMAA) Agence Nationale de Recherches sur le SIDA et les hépatites virales (ANRS) 12109 study, drugs were given during labour and all women underwent blood sampling for PK analysis at delivery and at 1, 2, 3, 5, 8, 12, and 24 h after drug administration. A cord blood sample was obtained at delivery. We also compared fetal-to-maternal and amniotic fluid-to-maternal concentration ratios for TFV to data obtained by digitalization [43] from another study during labour [4]. Maternal and fetal exposures were compared using the fetal-to-maternal area under the curve (AUC) ratios for a single dose and at steady state. The predicted exposures at the 39th week of gestation are reported in Table 5.

Sensitivity analyses on physiological constants such as the placental maternal blood flow parameter (Q_{plam}), k_{SW} , GFR_f and on parameters obtained *ex vivo* such as diffusion, k_{PE} and K_{ppl} were performed. Each time the reference value was multiplied or divided by 1.3, 2 and 5, and simulated FTC fetal and amniotic fluid PK profiles at steady state were obtained.

Results

Ex vivo model

A total of 26 experiments (TFV = 16, FTC = 10) could be validated for the integrity of the placental membrane and adequate conditions of perfusion. The mean (\pm standard deviation) antipyrine FTR was 53.9% ($\pm 5.9\%$) for TFV and 42.9% ($\pm 3.7\%$) for FTC. The mean FTR and CLI (30–150 min) were $21.0 \pm 5.7\%$ and 0.39 ± 0.11 , respectively, for TFV, and $24.3 \pm 6.4\%$ and 0.53 ± 0.17 , respectively, for FTC.

The transplacental transfer for these two drugs was best described by the diffusion model with estimation of $K_{p_{pl}}$ and k_{PE} (Figure 3).

Figure 4 compares observed to simulated concentrations for the *ex vivo* human perfusion model. Parameter estimates are summarized in Table 4.

In vivo simulations

The parameters estimated *ex vivo* ($K_{p_{pl}}$, k_{PE} , D_{cot}) were applied in the p-PBPK models. PK profiles during labour were simulated for a single dose and compared to observed data [5, 6]. Figure 5 shows that simulated fetal PK profiles using our p-PBPK model were in agreement with observed cord concentrations. As shown in Figure 5, fetal C_{max} values were lower than maternal ones. Table 5 summarizes the simulated fetal-to-maternal plasma and amniotic fluid-to-maternal plasma AUC ratios. A previous population approach estimated fetal AUC_{24}^0 as 10.7 mg h l^{-1} and 1.64 mg h l^{-1} at delivery for a single dose of FTC and TFV, respectively, whereas our model predicted 9.23 mg h l^{-1} and 1.26 mg h l^{-1} , respectively [5, 6].

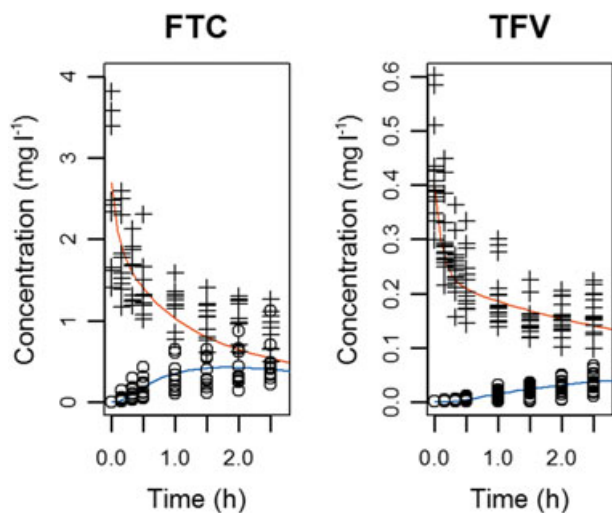


Figure 4

Ex vivo human placental perfusion model for emtricitabine (FTC; left) and tenofovir (TFV; right). Evolution of observed fetal concentration (open circles) and maternal concentration (crosses) compared to fetal (orange) and maternal (blue) simulated profiles in the *ex vivo* human placental model

Table 4

Estimated values for transplacental transfer parameters obtained in the *ex vivo* model, mean (range)

	$D_{cot}(\text{l h}^{-1})$	$K_{PE}(\text{h}^{-1})$	$K_{p_{pl}}$
Emtricitabine	0.104 (0.025–0.395)	1.49*	3.94 (0.92–9.33)
Tenofovir	0.013 (0.003–0.020)	0.443 (0.167–1.2)	7.15 (3.73–1.45)

D_{cot} , diffusion parameter; K_{PE} , placental elimination constant; $K_{p_{pl}}$, coefficient partition.

*No variability was estimated.

PK profiles for a single dose and at steady state obtained after administration of 200 mg of FTC once daily and 300 mg of TFV once daily in 39-week pregnant women were simulated. Figure 6 shows the variations of fetal-to-maternal and amniotic fluid-to-maternal concentration ratios over time. For TFV, the fetal-to-maternal plasma ratio stayed <1 , whereas amniotic fluid accumulation was observed. TFV simulations were close to the observed data obtained by Mirochnick *et al.* [4].

Sensitivity analyses showed that the diffusion and k_{PE} values were strong determinants of simulated fetal and amniotic fluid PK profiles (Figures 7, 8). The modification in $K_{p_{pl}}$ values influenced T_{max} but the effect on AUC was weaker. The daily volume of amniotic fluid swallowed by the fetus and the GFR_f could be multiplied or divided by 5 without significant modification of the simulated fetal PK but with a significant change in amniotic fluid PK. Finally, the change in maternal-to-placental blood flow rate had a negligible effect on both fetal and amniotic fluid PK.

Discussion

We developed a novel approach quantitatively predicting the fetal exposure of drugs administered to the mother. The p-PBPK models which implemented parameters estimated from human placental perfusion *ex vivo* experiments enabled the prediction of the fetal and amniotic fluid PK of FTC and TFV at full term.

This is the first study to report the *ex vivo* transplacental transfer of TFV and FTC. Only one other study has implemented parameters obtained from human *ex vivo* placental perfusion experiments in a mechanistic model [42, 44]. However, these authors used animal data to validate their model because no human data were available for a PK profile comparison.

Data from TEmAA ANRS 12109 enable us to validate our models [5, 6] by comparing simulated with actual fetal PK profiles. The predictions were also compared to other published data. Delays between sampling time and last dose were not available individually for these published data, and as concentrations and ratios are highly variable depending on these delays, these data were used only to obtain an estimate of orders of magnitude.

There are few data on FTC placental transfer; the geometric mean of cord blood concentration was found to be 0.26 mg l^{-1} [90% confidence interval (CI) $0.17\text{--}0.39 \text{ mg l}^{-1}$; $n = 11$] [45]. Moreover, Colbers *et al.* reported that the cord-to-maternal blood ratio 8.5 h (range 0–32 h) after dosing was 1.63 (90% CI $0.46\text{--}1.82$; $n = 10$) [46]. Based on model predictions, the mean ratio was 1.6 and mean fetal concentration was 0.19 mg l^{-1} (delay range: 0–24 h). No amniotic fluid data were available.

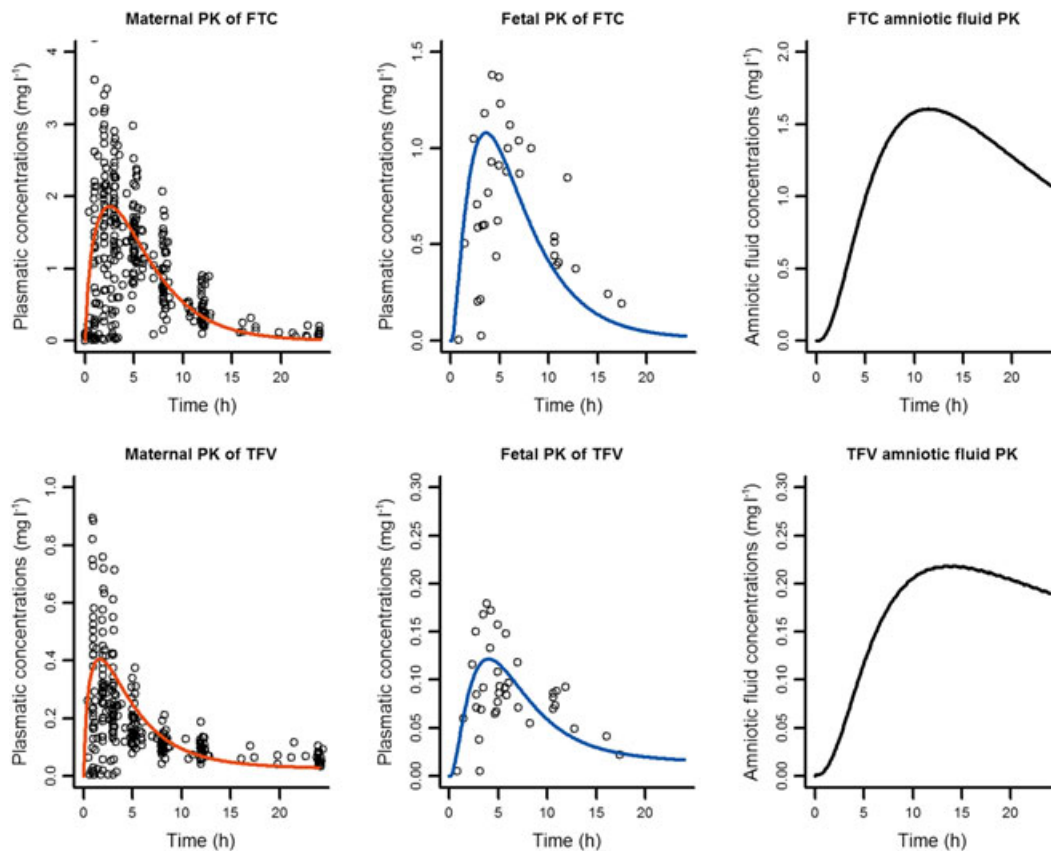


Figure 5

Simulation vs. observation of administration of emtricitabine (FTC) (top) and tenofovir (TFV) (bottom) during labour. Simulation (lines) of mean maternal (left), fetal (middle) and amniotic fluid (right) pharmacokinetic profile for a single dose compared to observations (points) [5, 6]. 'Time' represents the delay between the last dose and sampling time

Table 5

AUC ratio obtained after simulation during labour and during pregnancy at steady state

GA = 39		FTC	TFV
Single dose	Plasma AUC _f /AUC _m	0.63	0.41
	Amniotic fluid AUC/plasma AUC _m	1.93	1.39
Steady state	Plasma AUC _f /AUC _m	0.64	0.45
	Amniotic fluid AUC/plasma AUC _m	3.43	2.76

AUC, area under the curve; f, fetus; FTC, emtricitabine; m, mother; TFV, tenofovir.

TFV placental transfer has been described to be significant. The Pediatric AIDS Clinical Trials Group (PACTG) study observed a median cord blood concentration at delivery of 0.076 mg l⁻¹ (range: 0.000–0.309 mg l⁻¹; n = 10) in the group receiving 600 mg of TFV [47]. Moreover the current study found simulated cord-to-maternal plasma concentration ratios to be close to those measured by Mirochnick *et al.* (Figure 6) [4]. TFV has been reported to accumulate in the amniotic fluid, with highly variable concentrations [4, 48, 49]. Figure 6 shows that the p-PBPK model is able to predict this accumulation [4].

For TFV, the observed amniotic fluid concentrations indicated two features: accumulation and great

variability. This drug is mainly excreted unchanged by the kidney [25, 26]. Accumulation in the amniotic fluid could be explained by a greater excretion rate into amniotic fluid (renal excretion, lung excretion) relative to the absorption rate (swallowing, intramembranous pathway). The high variability of the amniotic fluid concentration can be partially explained by the variability of the volume swallowed by the fetus and the renal excretion volume [28, 29, 32]. Indeed, as showed by sensitivity analyses, modifications of these two phenomena greatly influence amniotic fluid concentrations (Figure 7).

Values for the fetal swallowing constant and renal excretion are not well documented and are difficult to obtain [29, 32, 50, 51]. However, even if the true values are higher or lower than those used in the models, the effect on fetal PK profiles would not be significant (Figure 7). Therefore, uncertainty about these physiological constants was not a major issue. The sensitivity analyses were carried out on one fetal-to-amniotic fluid transfer parameter (GFR_f) and one amniotic fluid-to-fetal transfer parameter (k_{SW}); the impact of analogous parameters (k_L , k_{INT}) on fetal and amniotic fluid PK was similar (data not shown). This sensitivity analysis also suggested that amniotic fluid can change considerably without a

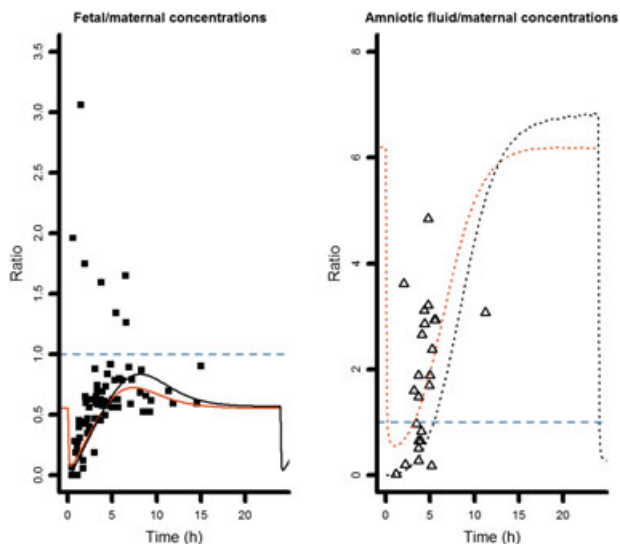


Figure 6

Simulations of fetal-to-maternal concentration ratios and amniotic fluid-to-maternal concentration ratios for tenofovir, for a single dose and at steady state. Evolution of fetal-to-maternal drug concentration ratios (solid lines) and amniotic fluid-to-maternal concentration ratios (dashed lines) for a single dose (black bold lines) and at steady state (thin orange lines). Ratios were compared to the available observed data: fetal-to-maternal drug concentration ratios (solid squares) and amniotic fluid-to-maternal concentration ratios (triangles) obtained by Mirochnick *et al.* [4]

significant change in fetal PK. Therefore, amniotic fluid may not be an accurate surrogate of fetal concentrations. Moreover, our model shows little effect of amniotic fluid concentrations on fetal PK. Thus, if the aim is only to

elucidate the fetal PK profile, this compartment could be ignored. Otherwise, the p-PBPK models provided a good prediction of the fetal PK profiles for these two renally excreted drugs but it might not be suitable for drugs that are mainly metabolized. The placental elimination pathway has not been elucidated for these drugs. In our models, placental elimination (k_{PE}) influenced fetal (and amniotic fluid) PK but did not significantly affect maternal PK.

Protein binding can lead to misleading results with the placental *ex vivo* model, so we added albumin to the media to ensure equivalent protein binding. These drugs are poorly plasma bound, so protein binding should not significantly influence the transplacental transfer. However, for drugs with high protein binding, this potential issue should be considered. Two options are available to decrease the risk of obtaining misleading results: (i) the *ex vivo* free fraction should be the same as the *in vivo* one or (ii) the transplacental transfer should be corrected for the *in vivo* free fraction.

The diffusion and placental elimination parameters obtained from the *ex vivo* experiments enabled good predictions of fetal PK profiles to be obtained. No random scaling factors were needed. Moreover, the sensitivity analyses performed on these parameters showed that any modification had a significant impact on fetal exposure. Thus, the *ex vivo* parameters, integrated into the PBPK model, seem to be a sensitive and accurate method for predicting fetal exposure. However, as the placental structure changes throughout gestation, this approach reflects the placental barrier only at delivery.

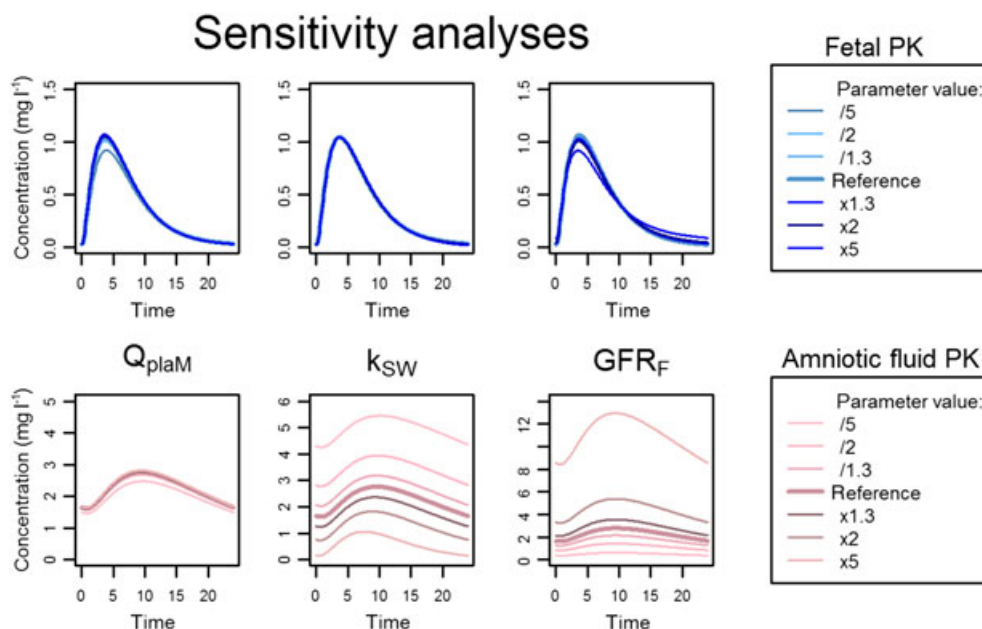


Figure 7

Sensitivity analyses of physiological parameters. Simulation of emtricitabine fetal pharmacokinetic (PK) profile (top) and amniotic fluid PK profile (bottom) after changes in placental perfusion (first column), daily swallowing volume (second column), fetal renal clearance (third column). k_{SW} , swallowing constant; GFR_F , fetal glomerular filtration rate; Q_{plam} placental maternal blood flow parameter

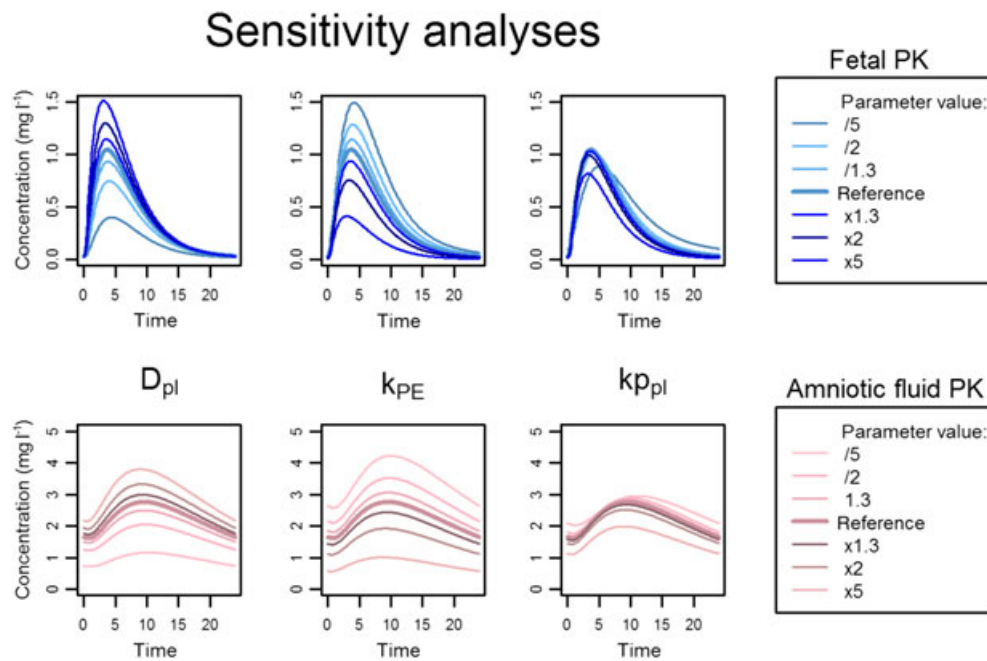


Figure 8

Sensitivity analyses of parameters estimated *ex vivo*. Simulation of emtricitabine fetal pharmacokinetic (PK) profile (top) and amniotic fluid PK profile (bottom) after changes in placental diffusion parameter D_{pl} (first column) and placental elimination constant k_{PE} (second column) and placental coefficient partition, K_{ppl} (third column)

The present approach enables a basic prediction to be made of fetal PK prior to drug administration to the mother. This should be a useful tool for the discovery of drugs targeting the fetus or those that can potentially be used in pregnant women at term. In the future, fetal tissue exposure might also be simulated through a more complex fetal PBPK model.

Conflict of Interest

All authors have completed the Unified Competing Interest form at www.icmje.org/coi_disclosure.pdf (available on request from the corresponding author) and declare: no support from any organization for the submitted work, no financial relationships with any organizations that might have an interest in the submitted work in the previous 3 years, no other relationships or activities that could appear to have influenced the submitted work.

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