Population pharmacokinetics of cefazolin in maternal and umbilical cord plasma, and simulated exposure in term neonates

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Background: Intra-partum cefazolin is used to prevent group B Streptococcus (GBS) vertical transmission in mothers allergic to penicillin without a history of anaphylaxis.

Objectives: To investigate the maternal cefazolin dose–exposure relationship and subsequent maternal and neonatal target attainment at delivery.

Methods: Data were obtained from 24 healthy, GBS-colonized pregnant women (20–41 years), undergoing vaginal delivery (gestational age ≥37 weeks). During labour, all women received a 2 g cefazolin IV infusion. Eight hours later, eight women received another 1 g in the event of delayed (>8 h) delivery. Next to maternal plasma concentrations (up to 10 per dosing interval, until delivery), venous and arterial umbilical cord concentrations were determined at delivery. Target attainment in maternal/neonatal plasma was set at 1 mg/L for 60% of the dosing interval (unbound cefazolin, worst-case clinical breakpoint). A population pharmacokinetic (popPK) model was built (NONMEM 7.4). ClinicalTrials.gov Identifier: NCT01295606.

Results: At delivery, maternal blood and arterial umbilical cord unbound cefazolin concentrations were >1 mg/L in 23/24 (95.8%) and 11/12 (91.7%), respectively. The popPK of cefazolin in pregnant women was described by a two-compartment model with first-order elimination. Two additional compartments described the venous and arterial umbilical cord concentration data. Cefazolin target attainment was adequate in the studied cohort, where delivery occurred no later than 6.5 h after either the first or the second dose. PopPK simulations showed adequate maternal and umbilical cord exposure for 12 h following the first dose.

Conclusions: PopPK simulations showed that standard pre-delivery maternal cefazolin dosing provided adequate target attainment up to the time of delivery.

Introduction

Group B Streptococcus (GBS) bacteria is one of the leading causes of maternal infections during pregnancy and post-partum, as well as in the neonate.1,2 GBS is a colonizer of the gastrointestinal or genitourinary tract in about 20% of pregnant women.3 Vertical GBS transmission from the mother to the neonate occurs during labour or birth in 36% of colonized mothers, which may lead to meningitis, sepsis and other infections in the neonate.3 Therefore, antibiotic regimens are administered in mothers, to provide intra-partum GBS prophylaxis. The antimicrobial of choice for prophylaxis against GBS infections is penicillin G or amoxicillin. Both are deemed equally effective.4 However, for women that have penicillin allergy without a history of anaphylaxis, cefazolin is a safe and effective alternative in this setting.4,5

Cefazolin is a first-generation cephalosporin that works by inhibiting bacterial cell wall synthesis. It is only available for IV and intramuscular administration. Besides GBS prophylaxis, cefazolin is a standard prophylactic antimicrobial agent for various surgical interventions, including delivery-related interventions like cesarean section, owing to its broad-spectrum activity against most Gram-positive bacteria and some Gram-negative bacteria.5 It is highly bound to serum albumin (75%–85%), with unbound cefazolin available for pharmacological activity.6–11 The efficacy of cefazolin for GBS prophylaxis depends on the time that the unbound
Several population pharmacokinetic (popPK) models of cefazolin have been developed in different clinical settings. Two models described the PK of cefazolin during pregnancy. Three other models described the cefazolin PK during caesarean delivery. There are some data suggesting that labour influences the PK of antibiotics. However, no model described the PK of cefazolin during labour and subsequent vaginal delivery. Also, the subsequent post-delivery early neonatal exposure is unknown. Therefore, a popPK model will be of relevance, providing insight into the intra-partum neonatal protection against GBS infections.

The objectives of our work were: (i) to build a popPK model of cefazolin based on maternal and umbilical cord plasma concentrations obtained from women undergoing vaginal delivery; (ii) to investigate target attainment [fT>MIC] for GBS, for 60%–70% of the dosing interval; and (iii) to investigate the early post-partum cefazolin exposure in the newborn.

Patients and methods

Patients and data collection

Data were obtained from a prospective, exploratory study. The goal of the study was to obtain insight into target attainment and the PK of cefazolin in women undergoing vaginal delivery. Healthy singleton pregnant women were recruited. All women were colonized with GBS and were admitted for vaginal delivery. During labour, the women received a 2 g cefazolin IV infusion over 30 min. If no delivery occurred 8 h after infusion, a second 1 g cefazolin infusion was given over 15 min. Maternal blood samples were obtained following a fixed sampling scheme with maximally 10 samples per dosing interval (until delivery), plus an additional sample right after delivery. Besides maternal blood samples, arterial and venous umbilical cord blood samples were collected at delivery. Additional data collected were gestational age, maternal age, maternal bodyweight and birthweight. The two parents’ informed consents were collected according to the Ethics Committee approval of the Centre Hospitalier Universitaire de Liége before sample collection (ClinicalTrials.gov Identifier: NCT01295606).

Analytical procedures

All heparinized blood samples were centrifuged immediately after sampling (10 min, 1200 g, ambient temperature) and plasma was stored at −80°C until further analysis. Cefazolin total plasma concentrations were measured using HPLC coupled with a diode array detector (HPLC-DAD) with a lower limit of quantification of 0.5 mg/L. The assay intra-day precision (expressed as relative standard deviation) varies from 0.935% to 4.396% depending on concentration (repeatability), while the assay inter-day precision varies from 1.483% to 5.426% depending on concentration (intermediate precision).

PK/pharmacodynamic target

An unbound cefazolin concentration above the MIC throughout at least 60% of the dosing period (i.e. 60% fT>MIC) was used as an optimal PK/pharmacodynamic target. In the absence of an MIC of cefazolin for GBS, we use the 0.25 mg/L ECOFF of amoxicillin for *Streptococcus agalactiae*. To avoid selecting resistant bacteria, a worst-case target of at least 60% fT>MIC was assumed.

Unbound cefazolin concentrations in maternal plasma were calculated from the measured total concentrations assuming 75% protein binding (lower limit of the 75%-85% interval, because albumin is known to decrease in the later stage of pregnancy).

PopPK modelling

Software

Dataset formatting and exploration were performed using R (version 3.5.1, R Core Team, Vienna, Austria) using custom scripts based on packages including dplyr and ggplot2. PopPK analysis was performed using NONMEM (version 7.4; ICON Development Solutions, Goithersburg, MD, USA), with a GNU Fortran 95 compiler and the Perl-speaks-NONMEM (PsN; version 4.7.0) toolkit on the interface software Pirana (version 2.9.7; Certara, Inc., Princeton, NJ, USA).

Model development

Parameter estimation was performed using first-order conditional estimation with interaction (FOCE-I) and differential equation solver ADVAN 13. The precision of the parameter estimates was evaluated based on the root squared error (RSE).

A base popPK model was developed. Different structural models with varying numbers of compartments were explored. Individual PK parameters were assumed to be log-normally distributed, which was achieved using an exponential function. Magnitudes of differences of individual parameters from the typical value (interindividual variability) and differences between observed values and model predicted values (residual variability) were quantified. The parametrization was performed using an exponential formula. For example, the clearance parametrization followed this equation:

\[
\text{CL}_i = \text{TVCL} \times \exp(\eta_i) \text{ with } \eta_i \sim N(0, \omega_i^2) \quad \text{(Eqn 1)}
\]

with CL<sub>i</sub> the clearance of subject i, TVCL the typical population value of the clearance, \( \eta_i \) the individual subject’s deviation from the typical value and \( \omega_i^2 \) the variance of inter-subject variability.

The base model was selected based on objective function value (OFV) comparisons (difference > 3.84 points; P < 0.050), Akaike information criterion (difference > 3.84 points; P < 0.050), plausibility and precision of parameter estimates, and goodness-of-fit plots. A final model, including covariate effects, was built via a two-way stepwise covariate modelling procedure (forward inclusion \( \alpha = 0.010 \); backward elimination \( \alpha = 0.001 \)). The tested covariates were gestational age and maternal bodyweight.

Final model evaluation

The final model was evaluated using a prediction-corrected visual predictive check (pcVPC; \( n = 1000 \) simulated replicates of the original dataset). Furthermore, bootstrapping was performed to obtain non-parametric estimates of uncertainty in parameter estimates (\( n = 2000 \) bootstraps).

Simulations in newborns

Neonatal exposure after birth in our study cohort was predicted using Monte Carlo simulations. These simulations were based on the popPK model-predicted arterial umbilical cord concentrations at birth to enable simulations in all 24 cases. Subsequent cefazolin exposure in the newborns was predicted from the cefazolin popPK model of De Cock et al. The cefazolin PK was simulated 1000 times for each newborn. Missing neonatal serum albumin was multiply imputed (\( n = 1000 \)) with values sampled from a uniform distribution within the range reported by Smits et al. (28.2–43.7 g/L). Postnatal age was set to zero and current weight was set equal to the registered birthweight in the current cohort of mother–newborn data.
**Maternal and arterial umbilical cord simulations**

PK simulations were performed during standard pre-delivery maternal cefazolin dosing (see the Patients and data collection section). The maternal exposure and the PTA in arterial umbilical cord plasma were calculated up to 12 h after the first infusion. Arterial umbilical cord plasma concentrations were considered fetal plasma concentrations, since arterial cord blood originates directly from the fetal circulation. Based on previous literature, protein binding was assumed to be 75% in maternal plasma (see the PK/pharmacodynamic target section) and 60% in neonatal plasma.10,11,26

**Results**

**Patient characteristics**

Data were obtained from 24 healthy pregnant women (20–41 years and bodyweight 70.5–89.7 kg), colonized with GBS, undergoing vaginal delivery (gestational age ≥37 weeks). A summary of participant characteristics is presented in Table 1. During labour, all women received a 2 g cefazolin IV infusion over 30 min for prophylaxis against GBS. In 16 women, delivery occurred within 6.5 h after the infusion [median (IQR) = 4.6 (3.3–5.6) h]. Eight hours after the first administration, the remaining eight women received a second cefazolin infusion (1 g) over 15 min.

Altogether, the women contributed 245 plasma samples to the analysis, 24 of which were matched venous and arterial umbilical cord samples from 12 women (Figure 1). At the time of delivery, the maternal unbound cefazolin plasma concentrations were above the 1 mg/L target in 23/24 (95.8%) women [5.82 (1.89–5.86) mg/L] (Figure S1, available as Supplementary data at JAC Online). The unbound 1 mg/L target was achieved in 11/12 (91.6%) arterial umbilical cord samples [5.12 (3.56–5.72) mg/L].

**PopPK model**

**Model development**

A two-compartment popPK model with linear elimination best described the maternal cefazolin concentration data (Table S1). Two additional compartments were added to represent the venous and arterial umbilical cord concentrations, with blood flowing from the maternal central compartment to the venous umbilical cord compartment, then to the arterial umbilical cord compartment, and then back to the maternal central compartment with the placenta as a barrier between both (Figure 2).

Interindividual variability was estimated on the maternal clearance [43% coefficient of variation (CV)], central volume of distribution (67%CV) and peripheral volume of distribution (68%CV). A mixed error model was selected to describe the residual variability based on goodness-of-fit plots and the OFV.

Gestational age and body weight were tested as covariates, but did not significantly improve the OFV or the parameter estimation accuracy. Therefore, these covariates were not retained in the final model (Table S2). Parameter estimates from the final model are shown in Table 2.

**Model evaluation**

Goodness-of-fit plots showed that the final model adequately described the observed data (Figure S2 and Figure S3). A pcVPC showed the good predictive capacity of our model, where the median values and the 95% CIs of the prediction-corrected observed data fell within the 95% prediction intervals of the model (Figure 3). Median values of the non-parametric bootstrap were in good agreement with point estimates (Table 2).

**Simulations in newborns**

The PK simulations showed adequate neonatal antimicrobial exposure for protection against GBS in neonates at birth in our study cohort [3.58 (2.50–4.84) mg/L] (n = 24) (Figure 4). Unbound cefazolin is predicted to remain detectable for at least 24 h after birth.

**Maternal and arterial umbilical cord simulations**

The PK simulations showed that the median maternal unbound cefazolin plasma is above the therapeutic target of 1 mg/L for the entire 12 h simulation period (Figure 5a). The PTA of neonatal (arterial) unbound cefazolin concentrations slowly increases and reaches 90% or more approximately 1 h after the start of infusion (Figure 5b).
Discussion

In this study, we performed a popPK modelling and simulation study of cefazolin in healthy women undergoing vaginal delivery and their newborns. We obtained a better understanding of the pre-delivery maternal cefazolin dose–exposure relationship and subsequent post-delivery duration of neonatal target attainment. Although PK variability is large, fixed 2 g cefazolin dosing—followed by 1 g cefazolin 8 h later if delivery had not occurred yet—provided adequate unbound cefazolin exposure in the mother and in the neonate at the time of birth.

A two-compartment model with linear elimination best described the maternal and umbilical cord concentration data. To the best of our knowledge, no popPK model of cefazolin in women in labour and subsequently undergoing vaginal delivery has been published before. However, there are two popPK models describing the cefazolin PK in women during pregnancy (gestational age range 17–40 weeks, fetal surgery or caesarean).\(^{10,11}\) Additionally, three popPK models are available describing cefazolin PK in women undergoing caesarean delivery.\(^{16,18}\) Two of the previous models (both caesarean delivery models) used a one-compartment model to describe the cefazolin popPK.\(^{16,17}\) This structural difference could be due to the sparse sampling schemes in these studies, as well as the physiological difference between vaginal and caesarean deliveries (different volume of distribution and clearance). In the previously published models, the most common covariates were maternal bodyweight, gestational age and maternal albumin. Bodyweight and gestational age were tested,
Table 2. Population parameter estimates

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Estimate (%RSE) [%shrinkage]</th>
<th>Bootstrap median (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Typical values</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$CL_{\text{maternal}}$ (L/h)</td>
<td>7.15 (10)</td>
<td>6.99 (3.41–9.03)</td>
</tr>
<tr>
<td>$V_c$ (L)</td>
<td>6.23 (19)</td>
<td>6.25 (3.81–8.82)</td>
</tr>
<tr>
<td>$Q_1$ (L/h)</td>
<td>26.1 (19)</td>
<td>27.75 (18.56–45.95)</td>
</tr>
<tr>
<td>$V_p$ (L)</td>
<td>6.95 (23)</td>
<td>7.05 (4.04–9.71)</td>
</tr>
<tr>
<td>$Q_2$ (L/h)</td>
<td>2.61 (16)</td>
<td>2.77 (1.04–6.86)</td>
</tr>
<tr>
<td>$V_{\text{venous umbilical cord}}$ and $V_{\text{arterial umbilical cord/neoatal}}$ (L)</td>
<td>6.97 (15)</td>
<td>8.87 (2.57–34.7)</td>
</tr>
<tr>
<td>$Q_3$ (L/h)</td>
<td>4.48 (12)</td>
<td>4.85 (1.84–12.48)</td>
</tr>
<tr>
<td>$Q_4$ (L/h)</td>
<td>5.47 (14)</td>
<td>5.49 (2.27–10.91)</td>
</tr>
<tr>
<td>$CL_{\text{neonatal}}$ (L/h)</td>
<td>0 FIXED</td>
<td>0 FIXED</td>
</tr>
<tr>
<td><strong>Interindividual variability model</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>on $CL$ (%CV)</td>
<td>42.9 (16) [10]</td>
<td>45.8 (26.3–80.8)</td>
</tr>
<tr>
<td>on $V_c$ (%CV)</td>
<td>66.6 (25) [8]</td>
<td>65.3 (17.9–114.1)</td>
</tr>
<tr>
<td>on $V_p$ (%CV)</td>
<td>67.9 (33) [26]</td>
<td>71.8 (22.5–158.3)</td>
</tr>
<tr>
<td><strong>Residual variability</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>additive residual variability (mg/L) (maternal)</td>
<td>0.082 (30) [9]</td>
<td>0.075 (0.034–0.124)</td>
</tr>
<tr>
<td>proportional residual variability (%CV) (maternal)</td>
<td>14.7 (43) [9]</td>
<td>19 (5.7–69.9)</td>
</tr>
<tr>
<td>additive residual variability (mg/L) (venous)</td>
<td>0.021 (58) [23]</td>
<td>0.014 (2.44 × 10^{–6}–0.043)</td>
</tr>
<tr>
<td>proportional residual variability (%CV) (venous)</td>
<td>0 FIX</td>
<td>0 FIX</td>
</tr>
<tr>
<td>additive residual variability (mg/L) (arterial)</td>
<td>0 FIX</td>
<td>0 FIX</td>
</tr>
<tr>
<td>proportional residual variability (%CV) (arterial)</td>
<td>9.16 (47) [5]</td>
<td>8.21 (0.43–17.81)</td>
</tr>
</tbody>
</table>

$CL$, clearance; $CV$, coefficient of variation calculated as $\sqrt{\exp(\omega^2) – 1} \times 100\%$; $V_c$, central volume of distribution; $V_p$, peripheral volume of distribution.

Figure 3. pcVPC of the final model. The observed prediction-corrected total cefazolin concentrations are represented by circles. The black continuous and broken lines represent the median and the 2.5th and 97.5th percentiles, respectively, of the prediction-corrected observations. The shaded areas indicate the 90% prediction intervals of the median and the 2.5th and 97.5th percentiles of the simulated values. This figure appears in colour in the online version of JAC and in black and white in the print version of JAC.
but not retained in our model, while serum albumin concentrations were not available in our study dataset.

Muller et al.\textsuperscript{19} suggested that physiological changes during labour and subsequent vaginal delivery may influence the maternal PK of antibiotics (mainly distribution characteristics), as well as their placental transfer and subsequent fetal and neonatal exposure. Comparing our cefazolin PK parameters during labour with those reported by others during pregnancy and caesarean delivery, no clear differences could be observed.\textsuperscript{17,27} However, a future popPK model meta-based analysis is warranted to obtain further insight into potential differences.

Besides maternal PK, Muller et al.\textsuperscript{19} also indicated the need to document fetal and neonatal exposure. Our dataset included samples from the umbilical cord (venous and arterial) at the time of cord clamping. Unfortunately, no postnatal observations in the newborns of the mothers recruited were available in our study. However, similar to Muller et al.,\textsuperscript{28} we postulated that the arterial umbilical cord concentrations reflect the neonatal concentrations at birth. Starting from these model-predicted arterial umbilical cord concentrations, subsequent neonatal exposure was simulated with the De Cock et al.\textsuperscript{26} model. Unbound cefazolin was predicted to remain above 0.5 mg/L for at least 24 h after birth. Consequently, a blood culture collected within this time frame, for example because of suspected early-onset sepsis in the newborn, is more likely to remain false negative.

Although target attainment was adequate in our study cohort, delivery occurred no later than 6.5 h after the first infusion and 4 h after the second infusion in all women. Therefore, we performed PK simulations, calculating the PTA at delivery across the 8 h after the first infusion and the 4 h after the second infusion (total 12 h period). These simulations showed that the unbound cefazolin concentration in the maternal plasma remains above the therapeutic target for the entire 12 h period of the simulations performed. Furthermore, the simulations also illustrated the slow increase in neonatal (arterial umbilical cord) unbound cefazolin concentrations, such that 90% PTA is reached approximately 1 h after the start of the first infusion.

Our study had several strengths. First, our dataset included rich sampling, which enabled the accurate capture of the structural elements of the popPK model. Second, our data included umbilical cord samples, which enabled us to have insight into the neonatal target attainment. Our study had some limitations. First, we were limited with regard to covariate exploration, because our dataset had only 24 pregnant women, which limited the ability of our model to identify covariates. Additionally, we were limited with the covariates included in our dataset. Only two covariates were tested in the popPK model building (gestational age and maternal bodyweight) and both were not retained. In future research, it would be of interest to test other covariates, such as estimated glomerular filtration rate and albumin, on the cefazolin PK in this clinical setting. Another limitation is that our dataset only included total cefazolin concentrations. The unbound concentrations were not measured. This limitation hindered our ability to fully explore the cefazolin maternal protection. Instead, we postulated a
cefazolin protein-bound fraction of 75% in the maternal blood as previously published. To conclude, we showed that the standard pre-delivery maternal cefazolin dosing provided adequate maternal and neonatal antimicrobial protection against GBS in our patient cohort up to the time of delivery. In addition, we developed a popPK model to describe the cefazolin PK in women during labour with subsequent vaginal delivery. This popPK model was used for simulations that showed that standard pre-delivery cefazolin dosing of a 2 g infusion at the start and a 1 g infusion at the eighth hour provides adequate unbound cefazolin concentrations during the entire 12 h time window of the simulations.

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**Transparency declarations**

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**Supplementary data**

Supplementary data, including Figures S1 to S3 and Tables S1 and S2, are available at JAC Online.

**References**


19 MIC EUCAST. https://mic.eucast.org/search/show-registration/12248.
