

**Population pharmacokinetic meta-analysis of individual data to design the first randomized efficacy trial of vancomycin in neonates and young Infants**

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Keywords:	neonate, pharmacometrics, infectious disease, therapeutic index, Nephrotoxicity

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Manuscripts

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2 **first randomized efficacy trial of vancomycin in neonates and young Infants**

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74 **Running title** : Population pharmacokinetic meta-analysis of vancomycin in neonates

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76 **Key words**: neonatal, population pharmacometrics, infectious disease, therapeutic  
77 index, toxicity

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**82 Synopsis****83 Objectives**

84 In the absence of consensus, the present meta-analysis was performed to determine an  
85 optimal dosing regimen of vancomycin for neonates.

**86 Methods**

87 A “meta-model” using NONMEM with 4894 concentrations from 1631 neonates was built  
88 and Monte Carlo simulations were performed to design an optimal intermittent infusion,  
89 aiming at reaching a target  $AUC_{0-24}$  of 400 mg\*h/L at steady state in at least 80% of  
90 neonates.

**91 Results**

92 A two-compartment model best fitted the data. Current weight, post-menstrual age (PMA)  
93 and serum creatinine were the significant covariates for clearance (CL). After model  
94 validation, simulations showed that a loading dose (25 mg/kg) and a maintenance dose  
95 (15 mg/kg twice daily if < 35 weeks PMA and 15 mg/kg three times daily if ≥ 35 weeks  
96 PMA) achieved the  $AUC_{0-24}$  target earlier than a standard “Blue Book” dosage regimen  
97 in more than 89% of the treated patients.

**98 Conclusions**

99 The results of a population meta-analysis of vancomycin data have been used to develop  
100 a new dosing regimen for neonatal use and assist in the design of the model-based,  
101 multinational European trial, NeoVanc.

102

## 103 **Introduction**

104 Vancomycin is one of the most widely used antibiotics in the world for the treatment of  
105 serious Gram-positive infections. It is a high molecular weight complex glycopeptide  
106 which inhibits the cell wall synthesis of Gram-positive bacteria by the formation of stable  
107 complex murein pentapeptides, thereby causing inhibition of further peptidoglycan  
108 formation. It became the treatment of choice for staphylococcal infections, when  
109 staphylococcal strains developed resistance to treatment with penicillin. It was then  
110 replaced by methicillin in the 1960s, but when the incidence of late onset neonatal sepsis  
111 increased due to coagulase negative and methicillin-resistant staphylococci, the use of  
112 vancomycin re-emerged and it is today the treatment of choice for many staphylococcal  
113 infections.<sup>1,2</sup>

114 According to recent surveys,<sup>3-7</sup> neonatal dosage recommendations for vancomycin are  
115 highly variable, and include a range of single or multiple clinical factors, such as  
116 gestational age (GA), post-natal age (PNA), postmenstrual age (PMA), weight and  
117 creatinine clearance. Even internationally recognised dosing guidelines gave different  
118 dosing recommendations, either as continuous (CVA) or intermittent intravenous (IVA)  
119 vancomycin administration. However, although vancomycin is one of the most studied  
120 antibiotics in neonates,<sup>2,8-10</sup> population pharmacokinetic (popPK) and pharmacokinetic–  
121 pharmacodynamic (popPKPD) approaches have had limited success in leading to a clear  
122 consensus on the optimal dosing regimen to use in routine clinical practice. This is partly  
123 because the models and results are dependent on study / centre-related factors, including  
124 differences in the covariates that were incorporated in the final analysis. The present  
125 study aimed to conduct a meta-analysis of published individual pharmacokinetic data and  
126 to build a popPKPD model that would take into account all available variables, as part of  
127 the programme of work to plan the NeoVanc trial.<sup>11</sup>

128

129

**130 Methods****131 Identification of individual patient data**

132 Published PK or PK/PD studies were identified through databases (PubMed, Embase) in  
133 2014. The investigator responsible for the publication was contacted, invited to participate  
134 in the present study and provided individual vancomycin dose and concentration data and  
135 associated covariates. Additional, non-published, routine Therapeutic Drug Monitoring  
136 (TDM) data were also used. All data were anonymised before transfer with a pre-defined  
137 data sharing agreement, according to Good Clinical and Laboratory Practices.

138

**139 Requested covariates and individual information**

140 The following dataset of mandatory variables was collected to ensure that individual  
141 patient data could be included in the pooled model.

142 *Vancomycin administration information:* dosing history for each infant (time of start of  
143 infusion, time of end of infusion and doses), continuous or intermittent infusion,  
144 vancomycin concentrations and exact sampling day and time.

145 *Demographic covariates:* gestational age, postnatal age, birth weight, current weight (at  
146 sampling), gender.

147 *Information on co-medications* was not collected and not analysed as it was available in  
148 only a limited number of neonates.

149 *Biological covariates:* serum creatinine concentrations

150 *Study-related covariates:* analytical method used to quantify vancomycin (FPIA EMIT,  
151 PENTINIA or CMIA CLIA), creatinine assay method (Jaffé or enzymatic) and  
152 corresponding units.



**153 Data analysis**

154 PK data were made available on a standard Microsoft Office Excel spreadsheet (CIC,  
155 1426, Hôpital Robert Debré) and formatted for subsequent modelling using NONMEM V  
156 7.2 (Icon Development Solutions, USA).

157 A first order conditional estimation (FOCE) method with interaction was used to estimate  
158 PK parameters and their variability. One and two compartment models with first order  
159 elimination were tested to estimate clearance (CL), central volume of distribution (V1),  
160 peripheral volume of distribution (V2) and intercompartmental clearance (Q) using the  
161 appropriate ADVAN subroutines. Inter-individual variability of the pharmacokinetic  
162 parameters was best described with an exponential model and was expressed as  $\theta_i =$   
163  $\theta_{\text{mean}} * e^{\eta_i}$ , where  $\theta_i$  represents the parameter value of the  $i^{\text{th}}$  subject,  $\theta_{\text{mean}}$  the typical value  
164 of the parameter in the population and  $\eta_i$  the variability between subjects, which is  
165 assumed to follow a normal distribution with a mean of zero and variance  $\omega^2$ .

166 Covariate analysis followed a forward and backward selection process. The likelihood  
167 ratio test was used to test the effect of each variable on model parameters. The effects  
168 of current weight, gestational age, postnatal age, postmenstrual age, serum creatinine  
169 concentration, analytical methods of vancomycin and creatinine, and ethnicity were  
170 investigated as potential covariates affecting PK parameters. During the first step of  
171 covariate model building, a covariate was included if a significant ( $p < 0.05$ ,  $\chi^2$  distribution  
172 with one degree of freedom) decrease (reduction  $> 3.84$ ) in the objective function value  
173 (OFV) from the basic model was obtained. All the significant covariates were then added  
174 simultaneously into a 'full' model. Subsequently, each covariate was independently  
175 removed from the full model. If the increase in the OFV was higher than 6.635 ( $p < 0.01$ ,  
176  $\chi^2$  distribution), the covariate was retained in the final model.

177 Model validation was based on graphical and statistical criteria. Goodness-of-fit plots,  
178 including observed (DV) *versus* population prediction (PRED); DV *versus* individual  
179 prediction (IPRED); conditional weighted residuals (CWRES) *versus* time and CWRES  
180 *versus* PRED were initially used for diagnostic purposes. The stability and performance  
181 of the final model was also assessed by means of a nonparametric bootstrap with re-  
182 sampling and replacement. Re-sampling was repeated 200 times and the values of  
183 estimated parameters from the bootstrap procedure were compared with those estimated  
184 from the original data set. The entire procedure was performed in an automated fashion,  
185 using Perl-speaks-Nonmem (PsN v2.30). The final model was also evaluated graphically  
186 and statistically by normalised prediction distribution errors (NPDE). One thousand  
187 datasets were simulated using the final population model parameters. NPDE results were  
188 summarized graphically by default as provided by the NPDE R package (v1.2): (i) QQ-  
189 plot of the NPDE; (ii) histogram of the NPDE. The NPDE is expected to follow the N (0,  
190 1) distribution.<sup>12</sup>

191 Monte Carlo simulations for dosage optimisation were performed to evaluate different  
192 weight adjusted (mg/kg) dosing regimens for three predefined neonatal groups:  
193 postmenstrual age (PMA) <29, 29-35 and >35 weeks. Drug exposure was simulated 100  
194 times for each set of patients including only the Caucasian patients. Area under the curve  
195 between 0 and 24h on the first treatment day ( $AUC_{0-24}$ ) and  $AUC_{0-24}$  at steady state  
196 ( $AUC_{SS0-24}$ ) were calculated for each simulated patient. The parameter estimates obtained  
197 from the final model were used to estimate the target attainment rate for an  $AUC_{0-24}$  of  
198  $\geq 400$  mg\*h/L with the standard dosage regimen recommended in the "Blue Book"<sup>13</sup> and  
199 to define the optimal dosing regimen able to attain this target in 80% of patients. The  
200 current dosage recommendations and a loading dose followed by a maintenance dose  
201 administered as an intermittent infusion were evaluated in the 3 PMA groups.

202

203 **Results**204 **Study population.**

205 A total of 1631 neonates and infants from 15 studies were included (Table 1). Their PMA  
206 and current weight (CW), expressed as mean (standard deviation) were 33.3 (5.7) weeks  
207 and 1785 (1127) grams, respectively. Patients' characteristics and vancomycin  
208 administration details are presented in Table 2. We refer to the original studies for  
209 additional factual information.<sup>14-26</sup>

210

211 **Population PK analysis**212 *Model building*

213 A total of 4894 concentrations from 1631 patients were included in the population  
214 analysis. A two-compartment model with first-order elimination best fitted the data; both  
215 the OFV and the residual variability were lower than with a one-compartment model.

216 *Covariate analysis*

217 Allowing separate estimates for each analytical method in the residual variability caused  
218 a significant drop in the OFV of 113.5 units. Body weight was the most important clinical  
219 covariate following a systematic covariate analysis, associated with a drop in the OFV of  
220 3367.1 units after incorporating it into the basic model using estimated allometric  
221 coefficients for CL,  $V_1$  and  $V_2$ . A further decrease in the OFV of 244.5 units was achieved  
222 by including PMA on CL and serum creatinine concentrations gave a further reduction  
223 ( $\Delta$ OFV 1087.9 units). The model was further improved ( $\Delta$ OFV 65.5 units) by introducing  
224 a conversion factor between the Jaffé and enzymatic assay methods for creatinine.  
225 Ethnicity (Malaysian patients) was identified as a sixth covariate ( $\Delta$ OFV 302.1 units) on  
226 CL. The final model had the following structure:

$$227 \quad CL = 0.0680 \times (CW/1350)^{0.863} \times RM \times RF \times F_{\text{Jaffé-Enzymatic}} \times F_{\text{race}}$$

228 where CW is current weight, RM reflects renal maturation and RF reflects renal function.

229 The population PK parameters of the final model are presented in Table 3.

### 230 Model evaluation

231 Model diagnostics showed acceptable goodness-of-fit for the final model. Predictions  
232 were unbiased and no trends were observed in the diagnostic plots of CWRES versus  
233 time. The NPDE distribution and histogram were consistent with the theoretical N (0, 1)  
234 distribution and density, indicating a good fit of the model to the individual data (Figure  
235 1). The mean and variance of the NPDEs were 0.09 and 0.98, respectively. Visual  
236 predictive checks (VPCs) of the final model for all neonates and in subgroups of neonates  
237 <29 weeks (L), 29-35 weeks (M) and >35 weeks (H) are shown in Fig. 2 (A and B). The  
238 plots confirm that the average predicted concentrations matched the observed  
239 concentrations and that the variability was well estimated in the 3 subgroups.

240 In addition, the median PK parameter estimates resulting from the bootstrap procedure  
241 closely agreed with the respective values from the final population model, indicating that  
242 the final model was stable (Table 3).

### 243 Dosing optimisation.

244 Dosing optimisation was conducted in the Caucasian population. Monte Carlo simulations  
245 were performed to evaluate different mg/kg dosing regimens for the three neonatal  
246 groups.

247 With the standard vancomycin dosing regimen at steady-state, the percentage of  
248 neonates reaching the target  $AUC_{\text{ss0-24}}$  of  $\geq 400$  mg\*h/ was 74.0% and the percentage  
249 exposed to an  $AUC_{\text{ss0-24}}$  above 700 mg\*h/L was 23.0% when considering all age groups.  
250 When considering only neonates <29 weeks PMA, the corresponding values fell to 27.7%  
251 and 1.1% respectively (Table 4).

252 With a loading dose of 25 mg/kg followed by the optimal maintenance dose of 15 mg/kg,  
253 either twice daily (BID) if  $\leq 35$  weeks PMA or three times daily (TID) if  $> 35$  weeks PMA,  
254 the percentage of neonates reaching the target  $AUC_{ss0-24}$  of 400 mg\*h/L was 89.3% while  
255 the percentage exposed to an  $AUC_{ss0-24}$  over 700 mg\*h/L was 33.3% when considering  
256 all age groups. When considering only neonates  $< 29$  weeks PMA, the corresponding  
257 values increased to 95.0% and 45.5% respectively (Table 5).

258 The target attainment rate on the first day of treatment increased from 42.6% with the  
259 standard regimen to 88.9% with the loading dose strategy.

260

## 261 Discussion

262 To the best of our knowledge, this is the largest meta-analysis that has assessed the  
263 population PK of vancomycin in neonates and young infants aged less than 3 months.

264 The analysis has combined vancomycin concentrations linked to key demographic and  
265 biological covariates from 15 pharmacokinetic studies conducted in 7 different countries.

266 Monte-Carlo simulations showed that the current dosage regimen was not suitable for the  
267 treatment of staphylococcal infection and that the optimal vancomycin dosing regimen  
268 should include a loading dose of 25 mg/kg for all neonates, irrespective of their PMA,  
269 followed by a maintenance dose adapted to their PMA.

270 Although widely used for many years, important questions remain on how to optimise  
271 vancomycin dosing in neonates.<sup>2,8,9</sup> In the absence of prospective evaluation, most

272 neonatal units have developed local dosing recommendations, resulting in variable  
273 exposures that may lead to poor efficacy, induction of resistance or toxicity<sup>3</sup>

274 Consequently, vancomycin dosage regimens adapted to neonates require harmonisation,  
275 taking into account the impact of developmental pharmacology on disposition and PK  
276 parameters from very preterm neonates through term neonates to older children.<sup>27</sup> This

277 issue is central and initiatives from both the FDA and EMA are currently being undertaken  
278 to revise vancomycin dosing.<sup>28,29</sup>

279 Drug pharmacokinetics and dynamics need to be linked to explicative individual  
280 characteristics either constitutional (age, weight, genetics, etc.) or environmental  
281 (pathology, drug interactions, etc.). In this context, population modelling allows  
282 assessment and quantification of sources of variability in drug exposure and response in  
283 the target population, even under sparse sampling conditions<sup>30-32</sup> The present study has  
284 confirmed the impact of serum creatinine and vancomycin assay methods as predictors  
285 of vancomycin concentrations in neonates.<sup>33,34</sup> Additional covariates, such as  
286 ventilation,<sup>35</sup> co-administered drugs (e.g. aminoglycosides or ibuprofen), ECMO,<sup>37</sup> whole  
287 body cooling,<sup>38</sup> as well as centre or country dependent effects linked to ethnic,  
288 environmental and nutritional differences, were not explored in the current study, as they  
289 were not available in all data sets. However, it is recognised that they may also contribute  
290 to PK variability in neonates.

291 Model-based approaches to characterise drug PK/PD have been recommended as  
292 powerful tools for overcoming the practical and ethical challenges associated with dose  
293 selection for neonatal indications.<sup>39,40</sup> For vancomycin, a model tailored dose had already  
294 been demonstrated to increase substantially the target attainment rate of vancomycin in  
295 treated neonates.<sup>10</sup> However, there were few neonates less than 29 weeks gestational  
296 age in that study and centre-effects could not be eliminated. These limitations were  
297 addressed in the present PK meta-analysis, which was conducted by pooling 4894  
298 vancomycin concentrations from 1631 neonates. Although robust parameter estimates  
299 were obtained with this strategy, different strategies may be necessary when data are  
300 heterogenous.<sup>41</sup>

301 For vancomycin, exposure, measured by  $AUC_{0-24}$ , is the PK/PD parameter influencing  
302 efficacy and emergence of resistance,<sup>42-44</sup> but also influencing toxicity. Nephrotoxicity is  
303 a multifactorial, well-identified risk of high vancomycin exposure and high trough  
304 concentrations.<sup>45</sup>  $AUC_{0-24}$  or trough levels can vary widely and independently, since the  
305 trough depends on both the daily dose and the frequency of administration, whereas  
306  $AUC_{0-24}$  only depends on the daily dose. Consequently, in the present study, simulations  
307 were performed to evaluate the current dosage regimen<sup>13</sup> and to optimize efficacy by  
308 determining the target attainment rate and exposure to vancomycin measured by the  
309  $AUC_{0-24}$ . A target  $AUC_{0-24}$  of at least 400 mg\*h/L was selected as an  $AUC_{0-24}/MIC$  ratio  
310 of 400 has been associated with favourable treatment outcomes in adults, assuming that  
311 bacterial strains have a vancomycin MIC  $\leq 1$  mg/L.<sup>43,46</sup> Simulations of the current dosage  
312 recommendation (see table 5) were performed after the first dose and at steady-state.  
313 Our results showed that the current daily dose was too low for all neonatal age groups  
314 but particularly for neonates <29 weeks, as less than 30% of neonates reached the  
315 steady-state target. As a loading dose strategy is recommended in adult settings in order  
316 to reduce the time needed to reach the target  $AUC_{0-24}$ <sup>47,48</sup> simulations were then  
317 performed with a loading dose and optimal maintenance doses in all age groups, based  
318 on weight and PMA. Increasing the maintenance dose to 15 mg/kg BID instead of OD  
319 was also tested in the group <29 PMA weeks to optimise dosage. These modifications  
320 led to an increase in the target attainment rate after the first dose and at steady-state in  
321 all age groups.

322 Nephrotoxicity is a recognised side-effect of vancomycin treatment, although its safety  
323 profile is considered favorable. The risk of nephrotoxicity primarily increases with high  
324 vancomycin exposure and duration of administration.<sup>45,49</sup> In studies in adults and  
325 children,<sup>45</sup> reported incidence varied widely, from 5% to 43%, occurrence increased with

326 longer durations of administration with a range of 4.3 to 17 days and nephrotoxicity was  
327 reversible in the majority of cases. In neonates, most studies were not sufficiently  
328 powered to detect nephrotoxicity and, when reported, renal impairment was frequently  
329 associated with concomitant administration of nephrotoxic drugs.<sup>50</sup>

330 Therefore, optimising exposure while reducing duration of administration would  
331 maximise clinical efficacy while minimising toxicity and selection of resistance.

332 The upper  $AUC_{0-24}$  limit remains a matter of debate. "Usual"  $AUC_{0-24}$  values of 700 or 800  
333 mg\*h/L have been used in both adults<sup>46,51,52</sup> and children,<sup>48</sup> however, more extreme  
334 values have also been reported, with breakpoints for nephrotoxicity of <600 or >1300  
335 mg\*h/L.<sup>51,53</sup> In the absence of specific neonatal data, a value of 700 mg\*h/L was used in  
336 the present study. With our simulated dosage regimen, 89% of neonates reached the  
337 predetermined  $AUC_{ss0-24}$  target and 21.0% had an  $AUC_{ss0-24}$ , over 800 mg\*h/L; this was  
338 slightly higher than the percentage expected with the dosing regimen that is currently  
339 used. Additional TDM is necessary to individualise therapy for patients at risk of high  
340 exposure rates.

341 Although the simulated dosage regimens increased the percentage of patients who would  
342 reach the target for efficacy, interpatient variability means that close monitoring is required  
343 with early analysis (ideally after the first dose) in patients for whom renal impairment is  
344 suspected. In the absence of pharmacokinetic interpretation software, it is easier to use  
345 trough concentrations than  $AUC_{0-24}$  to monitor vancomycin exposure but toxicity data in  
346 neonates remain sparse. A nephrotoxicity threshold of 15- 20 mg/L has been reported in  
347 both adult and paediatric studies,<sup>24,54,55</sup> although it is clear that a trough level is not a very  
348 good predictor of  $AUC_{0-24}$ .<sup>56,57</sup> Further data are also needed because additional factors  
349 specific to neonates may contribute to toxicity. These include hypovolemia, concurrent  
350 nephrotoxic drug use and duration of administration. This potential higher risk of



351 nephrotoxicity requires further evaluation. In our recent patient-tailored vancomycin dose  
352 study in 190 neonates, no patient developed nephrotoxicity after model-based TDM  
353 although the initial  $AUC_{0-24}$  reached 1200 mg·h/L in some patients.<sup>10</sup>

354 The drug regimen identified in the present, pre-clinical component of the NeoVanc  
355 programme is now being taken forward in a RCT of the optimised regimen in which the  
356 duration of vancomycin therapy is reduced to 5 days, compared to a standard dosing  
357 regimen and administration for 10 days. The aim of this change is to maximise clinical  
358 efficacy while minimising toxicity and selection of resistance. Due to multifactorial  
359 variability in vancomycin disposition, drug monitoring is being performed in the two arms  
360 to further guide dosing<sup>58</sup> and identify additional variability factors specific to neonates.

361

362

363

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373

374 **Authors' contributions**

375 EJA, WZ, SL, MS designed research

376 KA, EVC, VB, AT, NS, BM, YLL, RM, JEP, IL, JS, HN, JNA provided data and revised the  
377 manuscript

378 WZ and SL analysed data

379 EJA wrote the first version of the manuscript

380 EJA, WZ, AT, IL and MS revised it.

381

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387

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## Figure Legend

	Number	Mean (SD)	Median (Range)
<b>Patients</b>	1631		

558 **Figure 1:** Goodness-of-fit plots: **A)** Population predicted (PRED) versus observed  
559 concentrations (DV); **B)** Individual predicted (IPRED) versus DV; **C)** Conditional weighted  
560 residuals (CWRES) versus time; **D)** CWRES) versus PRED, Normalised prediction  
561 distribution errors: **E)** QQ-plot of the distribution of the Normalised Prediction Distribution  
562 Errors (NPDE) versus the theoretical N (0,1) distribution; **F)** Histogram of the distribution  
563 of the NPDE, with the density of the standard Gaussian distribution overlaid.

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565 **Figure 2:** Validation of the model by use of a visual predictive check (VPC) Visual  
566 predictive check after continuous (CVA: 2A) and intermittent (IVA) vancomycin  
567 administration. Comparison of the 5<sup>th</sup> (bottom dashed line), 50<sup>th</sup> (solid line), and 95<sup>th</sup> (top  
568 dashed line) percentiles obtained from 1,000 simulations and the observed data (circles)  
569 for vancomycin concentrations in premature neonates <29 weeks (L), 29-35 weeks (M)  
570 and >35 weeks (H). Open circles represents individual observed concentrations.

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	1463 Caucasian		
Ethnicity	116 Malaysian		
	52 Japanese		
GA (weeks)		31.2 (5.0)	30.0 (22.3 - 42.1)
PMA (weeks)		33.3 (5.7)	32.0 (23.3 - 52.4)
PNA (days)		16 (15)	11 (1 - 90)
Current weight (g)		1785 (1127)	1350 (415 - 11370)
Serum creatinine concentration ( $\mu\text{mol/L}$ )		59.2 (32.0)	53.9 (6.2 - 353.6)
<b>Vancomycin treatment</b>			
Continuous infusion	295		
Intermittent infusion	1336		

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**Table 1**

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**Demographic characteristics of the 1631 neonates and infants included**

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**Table 2**

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**Presentation of the studies included in the vancomycin meta-analysis**

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610 \*: Number of centers was given for multi centers study

611 **NP**: not published; **CVA**: Continuous intravenous Vancomycin Infusion; **IVA**: Intermittent612 intravenous Vancomycin infusion; **PETINIA**: particle enhanced turbidimetric inhibition

Study	N of patients N=1631	PK study	Adminis- tration	Location	Creatinine method	Vancomycin method	Ref
1	59	Single center	IVA	Glasgow, UK	Jaffe	FPIA	14
2	294	Multi centers (4)*	IVA	San Diego, US	Jaffe	EMIT and FPIA	15
3	35	Single center	IVA	Glasgow, UK	Jaffe	FPIA	16
4	210	Single center	IVA	Leuven, Belgium	Jaffe	PETINIA	17
5	116	Single center	IVA	Kuala Lumpur, Malaysia	Jaffe	FPIA	18
6	66	Single center	CVA	Marseille, Fran ce	Enzymatic	EMIT	19
7	61	Single center	CVA	Marseille, France	Enzymatic	EMIT	20
8	125	Single center	IVA	Memphis, US	Enzymatic	EMIT	21
9	55	Single center	CVA	Glasgow, UK	Enzymatic	CMIA	22
10	78	Single center	IVA	Paris, France	Enzymatic	EMIT and FPIA	23
11	113	Multi centers (3)*	CVA	Paris, France	Enzymatic	PETINIA and FPIA	24
12	199	Single center	IVA	Leuven, Belgium	Enzymatic	PETINIA and FPIA	25
15	68	Single center	IVA	Tartu, Estonia	Enzymatic	FPIA	26
13	52	Single center	IVA	Tokyo, Japan	Enzymatic	CLIA	NP
14	100	Single center	IVA	Valencia, Spain	Enzymatic	FPIA	NP

613 immunoassay **FPIA**: fluorescence polarization immunoassay method; **EMIT**: enzyme-614 multiplied immunoassay method; **CMIA**: chemiluminescent microparticle immunoassay;615 **CLIA**: chemiluminescent immunoassay; **CREA**: serum creatinine concentration in616  $\mu\text{mol/L}$ ; **PMA**: postmenstrual age in weeks.617 In our population, 1350 gram, 32 weeks and 52  $\mu\text{mol/L}$  are the median current weight

618 (day of the study), postmenstrual age, and serum creatinine concentration values,

619 respectively.

620 **Table 3: Population pharmacokinetic parameters of vancomycin and Bootstrap results**

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(n=500)

Parameters	Full dataset		Median	Bootstrap 2.5 <sup>th</sup> – 97.5 <sup>th</sup>
	Final	RSE(%)		
Central volume of distribution $V_1$ (L) $V_1 = \theta_1 \times (CW/1350)^{\theta_2}$				
$\theta_1$	0.728	1.5	0.714	0.414 – 0.742
$\theta_2$	1.13	3.0	1.12	0.596 – 1.200
Peripheral volume of distribution $V_2$ (L) $V_2 = \theta_3 \times (CW/1350)^{\theta_4}$				
$\theta_3$	0.358	11.1	0.335	0.185 – 0.474
$\theta_4$	1.15	14.9	1.25	0.75 – 1.93
Inter-compartment clearance $Q$ (L/h) $Q = \theta_5 \times (CW/1350)$				
$\theta_5$	0.0301	12.5	0.0361	0.0248 – 1.190
Clearance $CL$ (L/h) $CL = \theta_6 \times (CW/1350)^{\theta_7} \times RM \times RF \times F_{\text{Jaffé-Enzymatic}} \times F_{\text{race}}$				
$\theta_6$	0.0680	1.3	0.0686	0.0664 – 0.0717
$\theta_7$	0.863	5.3	0.895	0.787 – 0.968
$RM = (PMA/32)^{\theta_8}$				
$\theta_8$	0.544	30.3	0.544	0.143 – 0.816
$RF = 1/(F_{\text{Jaffé-Enzymatic}} \times CREA/54)^{\theta_9}$				
$\theta_9$	0.666	3.6	0.655	0.598 – 0.718
$F_{\text{Jaffé-Enzymatic}}$				
$\theta_{10}$	0.720	2.8	0.716	0.682 – 0.756
$F_{\text{race}}$				
$\theta_{11}$	0.724	2.8	0.710	0.646 – 0.757
Inter-individual variability (%)				
$V_1$	17.5	25.6	14.1	1.7 – 23.0
$V_2$	102.5	25.0	80.3	19.2 – 132.8
$CL$	18.2	21.6	15.2	2.2 – 21.0
Inter-occasion variability (%)				
$CL$	19.1	20.1	16.7	2.4 – 22.7
Residual proportional (%)				
FPIA	22.2	4.8	22.2	19.9 – 24.2
EMIT	20.9	7.3	21.1	18.0 – 24.0
PENTINIA	25.1	5.6	24.7	21.7 – 27.8
CMIA	10.7	21.2	10.6	5.7 – 14.2
CLIA	38.3	25.8	39.2	19.6 – 58.5
Residual additive (mg/L)				
FPIA	1.57	7.7	1.63	1.32 – 1.99
EMIT	1.53	16.7	1.54	0.94 – 2.08
PENTINIA	1.01	19.6	1.06	0.59 – 1.65
CMIA	2.02	26.1	2.07	0.71 – 2.78
CLIA	3.30	28.7	3.26	0.73 – 4.97

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626**Table 4: Monte Carlo simulations of vancomycin standard dose regimen\***

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<b>PMA (weeks)</b>	<b>&lt;29</b>	<b>29-35</b>	<b>&gt;35</b>	<b>Total</b>
Number of patients	335	618	510	1463#
Standard dose regimen (mg/kg)	15 OD	15 BID	15 TID	
<b>First day</b>				
AUC <sub>0-24h</sub> median (mg*h/L)	246	378	495	385
AUC <sub>0-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	163-356	264-523	332-725	203-638
Target attainment rate (%)	1.5	39.0	81.0	45.1
AUC <sub>0-24h</sub> 400-700mg*h/L (%)	1.5	38.9	74.0	42.6
AUC <sub>0-24h</sub> > 700mg*h/L (%)	0	0.1	7.0	2.5
AUC <sub>0-24h</sub> > 800mg*h/L (%)	0	0	2.0	0.7
C <sub>min24h</sub> median (mg/L)	3.8	9.0	14.3	9.0
C <sub>min24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	0.2-8.7	2.9-18.4	4.8-30.8	1.6-24.0
C <sub>min24h</sub> > 20mg/L (%)	0	3.4	24.2	9.9
<b>Steady-state</b>				
AUC <sub>ss-24h</sub> median (mg*h/L)	338	536	654	520
AUC <sub>ss-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	203-547	323-893	368-1276	259-1028
Target attainment rate (%)	27.7	84.3	91.9	74.0
AUC <sub>0-24h</sub> 400-700mg*h/L (%)	26.6	65.5	49.4	51.0
AUC <sub>ss-24h</sub> > 700mg*h/L (%)	1.1	18.7	42.5	23.0
AUC <sub>ss-24h</sub> > 800mg*h/L (%)	0.5	10.0	29.6	14.7
C <sub>minss-24h</sub> median (mg/L)	6.0	12.3	17.2	11.9
C <sub>minss-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	1.1-14.7	4.1-28.3	5.6-46.0	2.8-34.4
C <sub>minss-24h</sub> > 20mg/L (%)	12.8	17.6	40.0	21.7

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629 \* as indicated in the Blue Book (12),

630 # number of Caucasian patients,

631 AUC<sub>0-24h</sub>: 24h Area Under the Curve at the first day,632 C<sub>min24h</sub>: trough level at the first day633 AUC<sub>ss-24h</sub>: 24h Area Under the Curve at steady-state,634 C<sub>minss-24h</sub>: trough level at steady-state

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636**Table 5: Monte Carlo simulation of vancomycin dosage regimen with a loading**

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**dose 25 mg/kg following by optimal maintenance dose**

<b>PMA (weeks)</b>	<b>&lt;29</b>	<b>29-35</b>	<b>&gt;35</b>	<b>Total</b>
Number of patients	335	618	510	1463
Loading dose (mg/kg)	25	25	25	
Optimal maintenance dose (mg/kg)	15 BID	15 BID	15 TID	
<b>First day</b>				
AUC <sub>0-24h</sub> median (mg*h/L)	559	492	596	539
AUC <sub>0-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	384-787	336-692	426-820	358-812
Target attainment rate (%)	87.8	83.0	93.2	88.9
AUC <sub>0-24h</sub> 400-700mg*h/L (%)	74.0	78.6	67.5	74.7
AUC <sub>0-24h</sub> > 700mg*h/L (%)	13.7	4.4	25.7	14.2
AUC <sub>0-24h</sub> > 800mg*h/L (%)	4.1	1.0	12.0	5.6
C <sub>min24h</sub> median (mg/L)	14.4	10.7	15.5	13
C <sub>min24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	5.3-28.0	3.6-22.5	5.2-34.7	4.3-29.0
C <sub>min24h</sub> > 20mg/L (%)	21.2	8.9	31.3	19.8
<b>Steady-state</b>				
AUC <sub>ss-24h</sub> median (mg*h/L)	677	529	656	600
AUC <sub>ss-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	401-1102	325-883	368-1293	348-1093
Target attainment rate (%)	95.0	83.5	92.5	89.3
AUC <sub>0-24h</sub> 400-700mg*h/L (%)	49.5	65.0	49.4	56.0
AUC <sub>0-24h</sub> > 700mg*h/L (%)	45.5	17.6	43.1	33.3
AUC <sub>ss-24h</sub> > 800mg*h/L (%)	28.9	9.0	30.3	21.0
C <sub>minss-24h</sub> median (mg/L)	17.5	12.2	17.5	15
C <sub>minss-24h</sub> 5 <sup>th</sup> -95 <sup>th</sup> (mg*h/L)	6.5-38.0	4.0-28.5	5.6-46.1	4.9-37.6
C <sub>minss-24h</sub> > 20mg/L (%)	39.3	17.6	41.5	30.9

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AUC<sub>0-24h</sub>: 24h Area Under the Curve at the first day,

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C<sub>min24h</sub>: trough level at the first day

640

AUC<sub>ss-24h</sub>: 24h Area Under the Curve at steady-state,

641

C<sub>minss-24h</sub>: trough level at steady-state

Figure 1

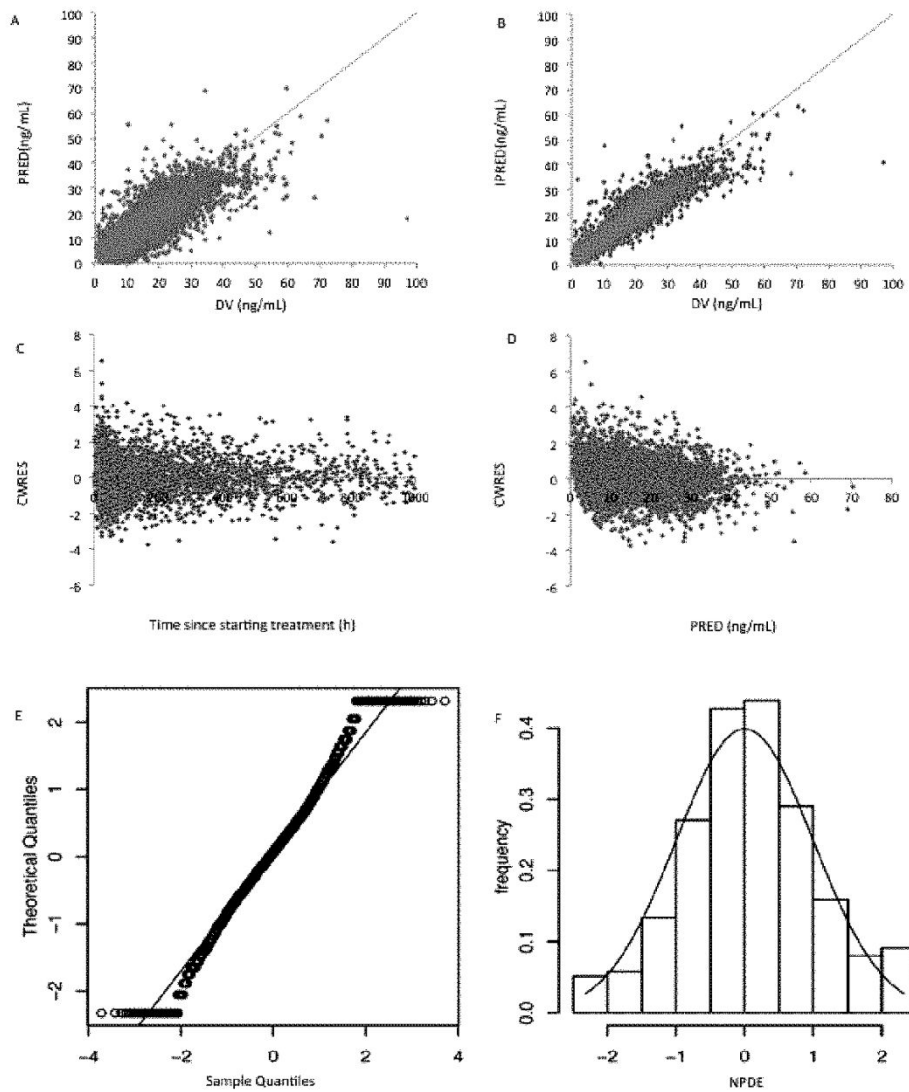




FIGURE 2

