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# Development and Evaluation of Bayesian Software for Improving Therapeutic Drug Monitoring of Gentamicin in Neonates

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## **Overview**

- Introduction
- Aims
- Methods
- Results
- Conclusions

#### TDM = therapeutic drug monitoring

# Introduction

- Neonatal infection → morbidity, mortality
- Gentamicin: antibiotic used for treating newborns
- Narrow therapeutic index
- Small blood volumes
- Renal function not fully developed
- $\rightarrow$  Need for TDM





## Introduction



#### **Background:**

• NPSA alert



#### **Patient Safety Alert**

NPSA/2010/PSA001 09 February 2010 National Patient Safety Agency

National Reporting and Learning Service

# Safer use of intravenous gentamicin for neonates

Patient safety incidents have been reported involving administration of gentamicin at the incorrect time, prescribing errors and issues relating to blood level monitoring.

#### **Action for the NHS**

NHS organisations, clinical directors and those responsible for the provision of neonatal services should ensure that by **9 February 2011:** 

 a local neonatal gentamicin protocol is available that clarifies the initial dose and frequency of administration, blood level monitoring requirements, and arrangements

## Introduction



## **UK neonatal units:**

• No single dosing & monitoring regimen

J Antimicrob Chemother 2011; **66**: 2647–2650 doi:10.1093/jac/dkr351 Advance Access publication 22 August 2011

#### Variation in gentamicin and vancomycin dosage and monitoring in UK neonatal units

S. Kadambari<sup>1\*</sup>, P. T. Heath<sup>1</sup>, M. Sharland<sup>1</sup>, S. Lewis<sup>2</sup>, A. Nichols<sup>3</sup> and M. A. Turner<sup>4</sup>

Forty-three units across the ENN responded to the gentamicin questionnaire, revealing 24 different combinations of dose, timing of dose and timing of monitoring. Figure 1 demonstrates

# Introduction



# Usually:

- Trough levels
  - taken at pre-set intervals (pre-dose)
  - require separate blood test
  - at inconvenient times

## **Bayesian methods:**

- Combine TDM with routine blood sampling
- $\rightarrow \downarrow$  invasive



#### **Published PK models:**

- Model: 1-, 2-, 3-compartment
- Covariates: weight, age, creatinine, APGAR score, sepsis, gender, co-medication
- $\rightarrow$  a more mechanistic approach is required



## Aim

- 1. Develop a software neoGent  $\rightarrow$  improve TDM
  - predict safe trough levels from routine blood samples
- 2. Evaluate the developed software
  - with new prospectively collected data



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#### Data

- Retrospective: published studies  $\rightarrow$  model building
  - Literature search: data from two studies<sup>a,b</sup>
  - N=174, samples=1163
  - GA=23.3-42.1 weeks, PNA=0-65 days
- Prospective: neoGent study → model evaluation
  - St George's, Liverpool, Oxford, Portsmouth, Coventry
  - Collected: June 2012 November 2013
  - N=163, samples=483
  - GA=23.9-42.3 weeks, PNA=0-77 days

GA = gestational age PNA = postnatal age

## **Methods**



## **Model development**

- PK meta-analysis performed on pooled data from published studies
- NONMEM VII; FOCE with interaction
- Mechanistic covariates:
  - Allometric scaling and a function, describing maturation of the glomerular filtration rate included *a priori*
  - PNA and serum creatinine tested

## **Methods**



## **Model predictions evaluation**

- Simulate from final model
  - Internal and external VPC
- Use prospectively collected data:
  - Take the 1<sup>st</sup> level = opportunistic, study sample
  - Predict the 2<sup>nd</sup> level = trough sample
  - Compare measured with model predicted trough level

## **Methods**



#### neoGent software (pilot version)

• Individual patient's data put into a file:

ID	GA	SEX	DATE	TIME	PNA	WT	CREAT	RATE	AMT	DV
1	226	1	03/09/2013	12:00	3	1710	59	240	8	0
1	226	1	03/09/2013	15:15	3	1710	59	0	0	6.8

- Read into R; changed to appropriate format
- Predictions: R calls NONMEM
- NONMEM results read back into R
- Prediction of the time when concentration <2 mg/L → safe to give the next dose</li>



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#### Final model

- 3-compartment model
- Residual error model: proportional + additive
- Inter-individual and inter-occasion variability: exponential model
- PNA and serum creatinine standardized for PMA: significant

$$CL = \theta_{CL} \cdot \left(\frac{WT}{70}\right)^{0.632} \cdot \frac{PMA^{3.33}}{55.4^{3.33} + PMA^{3.33}} \cdot \left(\frac{MSCr}{TSCr}\right)^{\theta_{SCr}} \cdot \frac{PNA}{\theta_{P_{50}} + PNA} \cdot e^{(\eta_{CL} + \kappa_{CL})}$$
$$V = \theta_V \cdot \left(\frac{WT}{70}\right) \cdot e^{\eta_V} \qquad Q = \theta_Q \cdot \left(\frac{WT}{70}\right)^{0.75} \cdot e^{\eta_Q}$$

 $TSCr = -2.8488 \cdot PMA [weeks] + 166.48$ 

Cuzzolin L, et al., Pediatr Nephrol 2006; **21**: 931-8. Rudd PT, et al., Arch Dis Child 1983; **58**: 212-5.



#### **Internal evaluation**





#### Visual predictive check

Internal VPC

External VPC





#### An example of the neoGent output





#### Predictions

#### PE = observed - predicted

dataset	Limit = 1 mg/L			Limit = 2 mg/L			Median	95% CI	
	n correct (%)	ΟΡ	UP	n correct (%)	ОР	UP	PE (mg/L)	2.5%ile	97.5%ile
paired + unpaired	215/254 (84.6)	17	22	246/254 (96.9)	6	2	-0.0016	-0.87	0.85
paired: study≥3mg/L	18/20 (90.0)	0	2	20/20 (100)	0	0	-0.061	-0.53	0.84
XV: paired: study≥3mg/L	428/456 (93.9)	13	15	421/456 (92.3)	20	15	-0.062	-1.55	1.04

OP is overprediction, UP is underprediction; PE is prediction error, CI is confidence interval for the prediction errors, XV is cross-validation



#### Paired samples with study level ≥3 mg/L





#### **Comparison with other published models**



Prediction error (mg/L)



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## Conclusions

- Final model good descriptive & predictive properties
- Provisional version of neoGent software developed

#### **Future work**

- Develop user-friendly interface
- Further clinical trial



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# **Questions?**