# A NEW APPROACH

TO THERAPEUTIC DRUG MONITORING OF TOBRAMYCIN IN CYSTIC FIBROSIS

THE DOSING INSTITUTE CLINICAL REVIEW SERIES: TOBRAMYCIN IN CYSTIC FIBROSIS MAY 2017



# THE DOSING INSTITUTE

The Dosing Institute (TDI) is an Australian led initiative translating the latest research on therapeutic drug monitoring (TDM) and precision dosing into education.

The primary aim is to upskill healthcare practitioners and students to integrate TDM and precision dosing into clinical practice, to enhance patient care and improve clinical outcomes.

Teaching hospitals and universities worldwide are increasingly realising the benefits of integrating effective educational programs on TDM and precision dosing into their curriculums. TDI was founded in collaboration with world leading universities and DoseMe to facilitate and advocate for high-quality education in this space.

## **EXECUTIVE SUMMARY**

### INTRODUCTION

Intravenous (IV) tobramycin, an aminoglycoside antibiotic, is a standard treatment for *Pseudomonas* infection in patients with cystic fibrosis (CF). However, drug exposure must be maintained within a narrow therapeutic range. If the dose is too high, there is an increased risk of adverse events, such as nephrotoxicity and ototoxicity.<sup>1,2</sup> If the dose is too low, there is an increased risk of therapeutic failure, antibiotic resistance and poor patient outcomes.<sup>3,4</sup> As a result, therapeutic drug monitoring (TDM) - measurement of drug concentrations in the blood at known times - of tobramycin is essential to enable appropriate dose individualization.

# KEY CHALLENGES OF CONVENTIONAL METHODS ESTIMATING AUC FOR TOBRAMYCIN

Dose individualization of tobramycin is usually performed by taking two blood samples at fixed time points (relative to dose administration) and interpolating these into an estimation formula. This conventional method for estimating tobramycin area under the curve (AUC) – the actual body exposure to a drug after administration of a dose - is associated with a number of practical challenges, including difficulties surrounding drawing two blood samples, where patients often fail to present for their second blood test at the proposed time.<sup>5</sup>

### CONCLUSION

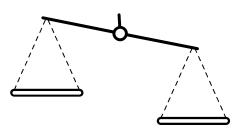
In this study, DoseMe, a Bayesian dose individualization platform, is shown to be more accurate at estimating AUC than the conventional two-sample method while requiring only one blood sample and allowing the collection of this sample at almost any patientpreferred time.



# CURRENT CHALLENGES ASSOCIATED WITH THERAPEUTIC DRUG MONITORING OF TOBRAMYCIN

### THE CHALLENGE

To be effective and well tolerated, tobramycin drug exposure must be maintained within a narrow therapeutic range. Achieving this is further complicated by a high degree of inter- and intra-patient variability in the PK parameters of tobramycin. As a result, therapeutic drug monitoring (TDM) of tobramycin is essential to enable appropriate dose individualization beyond covariate-based dosing (i.e. using age and/or mg/kg).



**TOBRAMYCIN DOSE TOO HIGH** Increased risk of nephrotoxicity and ototoxicity, of which the latter may be irreversible<sup>1,2</sup> **TOBRAMYCIN DOSE TOO LOW** Increased risk of therapeutic failure, antibiotic resistance and poor patient outcomes<sup>3,4</sup>

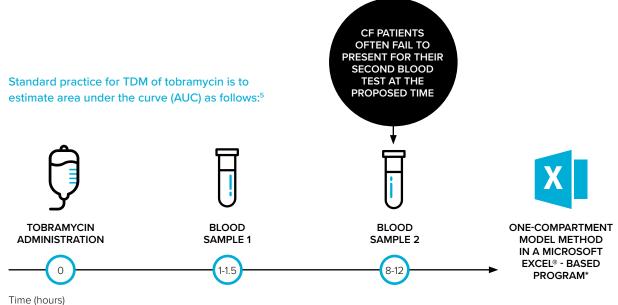
### WHAT IS THERAPEUTIC DRUG MONITORING?

TDM is the measurement of drugs at specific intervals to maintain a constant concentration in a patient's bloodstream - to optimize individual dosage regimens. Performing TDM requires a multidisciplinary approach that is integral to the clinical success of dose individualization for drugs with narrow therapeutic ranges and pharmacokinetic variability.<sup>6</sup>

### CLINICAL RELEVANCE

Accurate and clinically meaningful drug concentrations are attainable only by collaboration across multiple departments, with most methods relying on multiple blood samples to be taken at specific times.<sup>7</sup>

In a busy and demanding clinical environment, the optimal collection period for TDM samples is short, and unsurprisingly, inaccurately timed blood collections are common. In one study, only 4.3% of blood samples were deemed to have been taken at the correct time.<sup>8</sup> This increases the potential for misleading clinical decision making on these results, having implications for patient safety, including sub- or supratherapeutic dosing.



\*Dose individualization tools are regulated under law in many jurisdictions. The Dosing Institute does not endorse the use of unathorized tools.

# **TOBRAMYCIN TDM STUDY OVERVIEW**

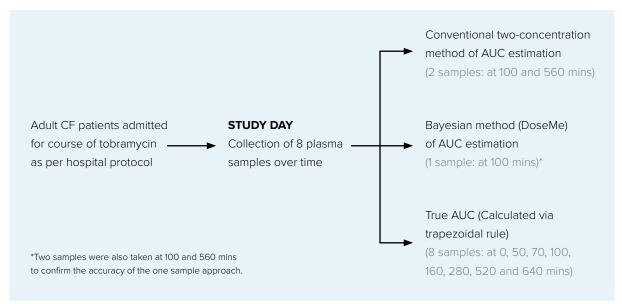
### AIM

To compare the conventional two-sample AUC estimation method with a one-concentration Bayesian method\* using DoseMe, and compare these with the true AUC, calculated from sampling across eight timepoints.

#### STUDY DESIGN

Data were collected from adult CF patients prescribed once-daily intravenous tobramycin at the Mater Health Service (MHS) Respiratory Unit, Brisbane, Australia.

#### Figure 1. Flow chart for data collection



#### RESULTS

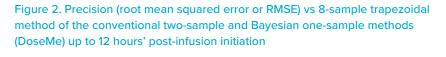
Data were collected from 12 patients with CF (Table 1).

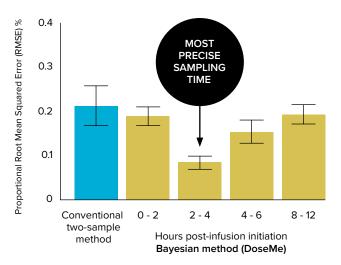
#### Table 1. Patient demographics

Demographics (n=12)	No. or median (range)
Age, year	25 (18–36)
Weight, kg	66.5 (51–76)
Height, cm	170 (154–185)
Gender: Male	7
Gender: Female	5
Serum creatinine, µmol/L	73 (55–92)
Dose at study day, mg	400 (320–640)
Dose, mg/kg	6.3 (5.2–8.4)
Day of therapy for sampling	6 (4–7)

\*Bayesian dose individualization decision support systems, such as DoseMe, provide pharmacokinetic model-based clinical decision support. These methods integrate patient data and laboratory results with a prior population pharmacokinetic and pharmacodynamic to estimate a patient's ability to absorb, process and clear a drug from their system.

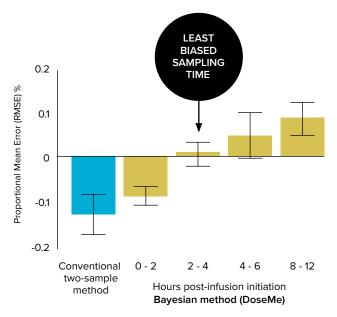
Bayesian dosing tools adjust the parameters so that a patient-specific, individualized drug model is created. This individual model is then used to provide a patient-specific dosing recommendation to reach a specific target exposure. With one sample, the Bayesian method using DoseMe provided similar estimated AUC values to the true, 8-sample trapezoidal AUC. Furthermore, the Bayesian method was both more precise and less biased than the two-sample conventional method using any one sample taken between 2-4 hours after initiation of tobramycin infusion.





The Bayesian method (2–4 hours) using DoseMe was significantly more precise than the conventional method (RMSE 8% vs 21%, respectively)

Figure 3. Bias (Mean Error) vs 8-sample trapezoidal method of the conventional two-sample and Bayesian one-sample method using DoseMe up to 12 hours' post-infusion initiation



The Bayesian method (2–4 hours) using DoseMe was significantly less biased (ME 0% vs – 14%, respectively)

# THE BENEFITS OF A ONE-SAMPLE BAYESIAN METHOD FOR TDM OF TOBRAMYCIN

### CONCLUSION

This study demonstrates that tobramycin TDM in CF patients can be simplified safely with a one-sample Bayesian method. The reduced requirement for blood collection samples not only lessens the burden placed on CF patients by alleviating overall time spent in hospital, but additionally reduces cost by removing unnecessary assays. Overall, this study shows that using a one-sample Bayesian method such as DoseMe, reduces the restriction on the sampling times, which is known to be problematic in the hospital environment. By allowing the use of any sampling time, Bayesian methods reduce the likelihood of inappropriate use of assays, assisting healthcare practitioners to select the most appropriate dose to achieve an optimal response and reduce potential toxicity.



DoseMe using a one-sample Bayesian dosing method achieves higher accuracy and precision in calculating AUC using half as many laboratory results as the conventional clinical method.



Using the DoseMe Bayesian dosing method, only one assay result is required for accurate AUC estimation.



DoseMe using a one-sample Bayesian method allows collection to be at a patient or providerpreferred time. While ideally 2-4 hours after initiation of infusion any single time up to 12 hours is as accurate as conventional methods.

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