

Background

- Current guidelines endorse vancomycin area under the concentration-time curve (AUC)-based monitoring over trough-only monitoring for serious methicillin-resistant Staphylococcus aureus (MRSA) infections¹
- Vancomycin AUC can be estimated using either Bayesian modeling software or first-order pharmacokinetic (PK) calculations
- Both methods have pros and cons; Bayesian software is generally more expensive, but can estimate an AUC using potentially just one vancomycin level
- **Purpose**: to compare a Bayesian modeling program and first order PK equations in terms of efficiency for calculating the estimated vancomycin AUC



Methods

- Single-center, prospective, crossover study
- 4 medical/surgical units at Brigham & Women's Hospital over a 2-month time period (11/11/19 – 1/13/20)
- Intervention
 - Vancomycin concentration monitoring by Bayesian modeling (InsightRX[®]) versus PK calculations (EPIC[®] Kinetics Navigator)
- Extensive education to clinicians, pharmacists, and nursing on the changes to the protocol for vancomycin monitoring (see specific examples in QR code)
- Primary endpoint
- Time taken to estimate a vancomycin AUC and determine dose adjustments with PK calculations compared to Bayesian modeling, with and without electronic health record (EHR) integration (see full protocol in QR code for details)
- Secondary endpoint Number of vancomycin levels drawn & percentage that were usable for AUC calculations
- Partners Healthcare Institutional Review Board deemed project exempt as it was a quality improvement initiative

- Inclusion Criteria:

124 patients evaluated for inclusion

Vancomycin Area Under the Concentration-Time Curve Estimation Using a Bayesian **Approach Versus First-Order Pharmacokinetic Equations: a Pilot Study**

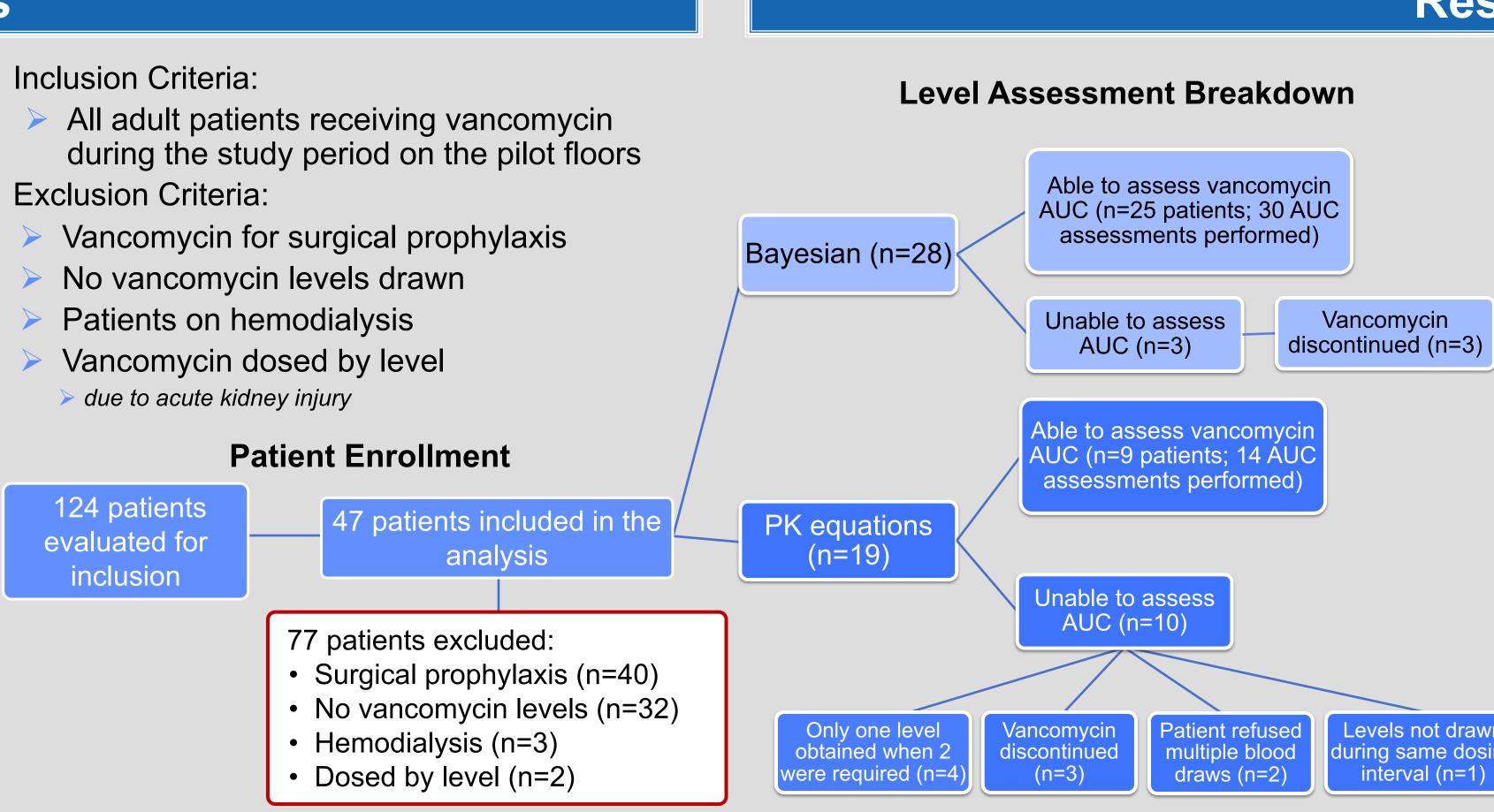
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> Compared to first-order PK equations, estimating a vancomycin AUC with Bayesian software (if integrated in the EHR):

Required less time to calculate vancomycin AUC

Allowed for greater flexibility with regards to timing levels, which led to fewer unusable vancomycin levels

Required fewer overall vancomycin levels to be drawn



Results

Primary Endpoint: Time to Estimate AUC

Variable (n=level assessments)	Bayesian (n=30)	PK equations (n=14)	p-value
Total time: Bayesian versus PK equations	9.3 (7.8-12.4)	6.8 (4.8-8)	0.004
Adjusted time if Bayesian software is EHR-integrated versus PK equations	3.8 (2.3-6.9)	6.8 (4.8-8)	0.019

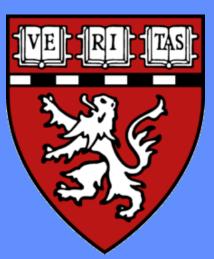
*All data presented in minutes as median (IQR)

Secondary Endpoint: Vancomycin Levels

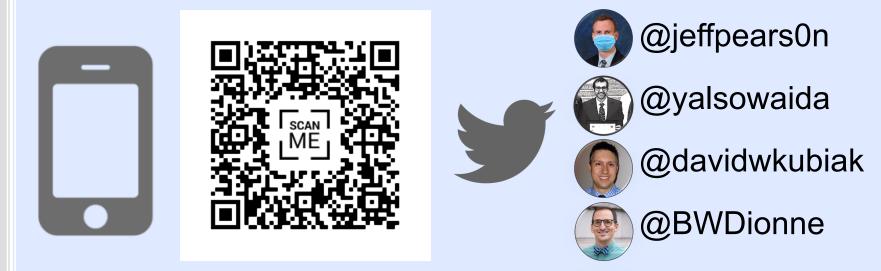
Variable (n=levels)	Bayesian (n=34)	PK equations (n=58)	p-value
Usable vancomycin levels to calculate AUC, n (%)	30/34 (88.2%)	28/58 (48.3%)	0.001



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Take a picture of the QR code for more info (protocol, education examples, methods poster), and/or reach out on Twitter to further discuss!

Limitations

- Single-center study with a small sample size
- Primary endpoint only applicable to Bayesian modeling using InsightRX®
- Access to InsightRX[®] was through a web-based program, so time was adjusted to infer pharmacists' time if integrated into EHR

Conclusion

- The Bayesian approach using InsightRX[®] takes less time to calculate vancomycin AUC and recommend dose changes if integrated into the EHR
- Individual institutions should weigh the pros and cons of different vancomycin monitoring solutions to determine the best methods for their needs

Disclosures

- Bayesian modeling software was donated by InsightRx[®] for the purposes of this project
- All authors have nothing else to disclose

Reference

1. Rybak M, Le J, Lodise, TP, et al. Therapeutic monitoring of vancomycin: A revised consensus guideline and review of the American Society of Health-System Pharmacists, the Infectious Diseases Society of America, the Pediatric Infectious Diseases Society and the Society of Infectious Diseases Pharmacists. Am J Health Syst Pharm. 2020;77(11):835-864. doi:10.1093/ajhp/zxaa036.