²⁰³² Clinical Impact of Rapid Identification and Phenotypic Antimicrobial Susceptibility Testing by Accelerate PhenoTM System for Gram-negative Bloodstream Infections



N. Matic^{1*}, B. Willey², B. Gascon², S. Lee², V. Koren², S. Surangiwala², P. Lo², T. Mazzulli^{1,2,3}, S. M. Poutanen^{1,2,3}

¹Department of Laboratory Medicine and Pathobiology, University of Toronto, ON, Canada ²Department of Microbiology, University Health Network/Sinai Health System, Toronto, ON, Canada ³Department of Medicine, University of Toronto – all in Toronto, Canada

Abstract

Background

- Laboratory turn-around-times (TATs) for identification (ID) and antimicrobial susceptibilities (AST) can delay prescription of adequate/optimal antimicrobial therapy (ABX) in septic patients, leading to poor outcomes.
- The Accelerate Pheno[™] system (Accelerate Diagnostics, USA) (AXDX) is a rapid ID and AST system with potential to improve TATs.

<u>Methods</u>

- 70 prospective non-duplicate blood cultures with Gram-negative bacilli were tested by AXDX.
- AXDX TATs were compared to TATs of current methods (MALDI-TOF with shortincubation subcultures & VITEK® 2), modified current methods (foregoing purity plate review or calling ID & AST results), and former methods (VITEK® 2).

<u>Results</u>

- Using current methods: Gram stain, ID and AST results led to tailoring of ABX in **88.6%** of patients, impacting **22.9%**, **31.4%**, and 64.3% of patients at 2.5h, 19.0h, and 62.1h respectively.
- AXDX generated the shortest ID and AST TATs with the potential to shorten time to ABX tailoring in response to ID and AST to **1.3h** and **6.7h**, respectively.

Conclusions

- Among the methods compared, AXDX has the greatest potential impact on time to appropriate ABX in Gram-negative bloodstream infections.
- Prospective studies evaluating the impact on patient outcomes are needed.

Introduction

- For every hour delay in the initiation of appropriate antimicrobial therapy (ABX) in septic patients, there is an average 7.6% decrease in survival.
- Laboratory methods for identification (ID) and antimicrobial susceptibility testing (AST) of organisms from positive blood cultures have traditionally relied on culture, which can have turn-around-times (TATs) up to >80 hours.
- Rapid ID and AST systems have the potential to improve laboratory TATs, with subsequent improvements in clinical outcomes such as mortality and hospital length of stay.
- Furthermore, rapid microbiology results have been shown to lead to earlier de-escalation of ABX in conjunction with an antimicrobial stewardship program (ASP).
- The Accelerate Pheno[™] system (Accelerate Diagnostics, USA) (AXDX) is a fully-automated, rapid diagnostic system that is used directly on positive blood cultures. It performs:
 - Gel electro-filtration and fluorescence in situ hybridization for ID
 - Automated microscopy for observation of bacterial growth and extrapolation of MICs for AST

Objective: Our laboratory evaluated the potential clinical impact of AXDX by reviewing ABX changes made by physicians as Gram stain, ID, and AST results are released.

Methods

- 70 prospective non-duplicate blood cultures with Gram-negative bacilli from 3 Toronto tertiary care hospitals from Jun-Sep 2016 were tested by AXDX as part of a concurrent study assessing AXDX accuracy (IDWeek Poster 2031).
- AXDX ID and AST TATs were compared to the TATs of: **Current Methods**
- ID by MALDI-TOF using VITEK® MS (bioMérieux) from short-incubation subcultures and AST by VITEK® 2 (bioMérieux) **Modified Methods**
- Releasing VITEK® 2 AST results prior to purity plate review
- Calling ID and AST results to ward

Former Methods

- ID and AST by VITEK® 2 from data review of 134 retrospective blood cultures from 2011
- Chart review was performed to assess time to tailoring of ABX by physicians following Gram stain, ID, and AST results.



Nancy.Matic@medportal.ca Susan.Poutanen@sinaihealthsystem.ca Toronto, ON, Canada M5G 1X5 Tel: 416-586-3139, Fax: 416-586-8746 IDWeek 2017, San Diego, USA

Sinai Health

System

Results

 Table 1: Blood culture isolates and patient
locations (n=70)

Blood Culture Isolates	
Escherichia coli	55.7%
Klebsiella pneumoniae	17.1%
Pseudomonas aeruginosa	8.6%
Enterobacter cloacae	5.7%
E. coli + K. pneumoniae	5.7%
Serratia marcescens	2.9%
Acinetobacter baumanii	1.4%
Klebsiella oxytoca	1.4%
S. marcescens + Enterococcus faecalis	1.4%
Patient Locations	
ER	54.3%
Medicine	11.4%
ICU	10.0%
Surgery	8.6%
Transplant Unit	5.7%
Other	10.0%

Conclusions

- AXDX generated the shortest ID and AST TATs with the greatest potential to significantly shorten the time to ABX tailoring in Gram-negative bacilli bloodstream infections thereby impacting patient outcomes.
- Calling ID and AST results directly to physicians or releasing AST results from VITEK® 2 prior to purity plate review would also have the potential to significantly improve time to ABX change compared to current methods.
- Prospective studies evaluating the impact of the reduced TATs associated with AXDX on patient outcomes are needed.

- Chemother 72: 299-304.

References

(1) Kumar A, Roberts D, et al. 2006. Duration of hypotension before initiation of effective antimicrobial therapy is the critical determinant of survival in human septic shock. Crit Care Med 34: 1589-96.

(2) Huang AM, Newton D, et al. 2013. Impact of Rapid Organism Identification via Matrix-Assisted Laser Desorption/Ionization Time-of-Flight Combined With Antimicrobial Stewardship Team Intervention in Adult Patients With Bacteremia and Candidemia. Clin Infect Dis 57: 1237-45.

(3) Bouza E, Sousa D, et al. 2004. Bloodstream infections: a trial of the impact of different methods of reporting positive blood culture results. Clin Infect Dis 39: 1161-69.

(4) Battle SE, Bookstaver PB, et al. 2017. Association between inappropriate empirical antimicrobial therapy and hospital length of stay in Gram-negative bloodstream infections: stratification by prognosis. J Antimicrob

(5) MacVane SH, Nolte FS. 2016. Benefits of adding a rapid PCR-based blood culture identification panel to an established antimicrobial stewardship program. J Clin Microbiol 54: 2455-63.