

Impact of BioFire® FilmArray® Blood Culture Identification Panels on Time to Optimal Antimicrobial Therapy

Corresponding Author: Aiman Bandali aiman.bandali@atlantichealth.org

Shivam Vyas, PharmD¹; Aiman Bandali, PharmD, AAHIVP, BCPS, BCIDP²; Dimple Patel, PharmD, BCIDP¹; Pamela Giordano, PharmD, BCPS, BCIDP¹

¹Morristown Medical Center, Morristown, New Jersey; ²Overlook Medical Center, Summit, New Jersey

Background

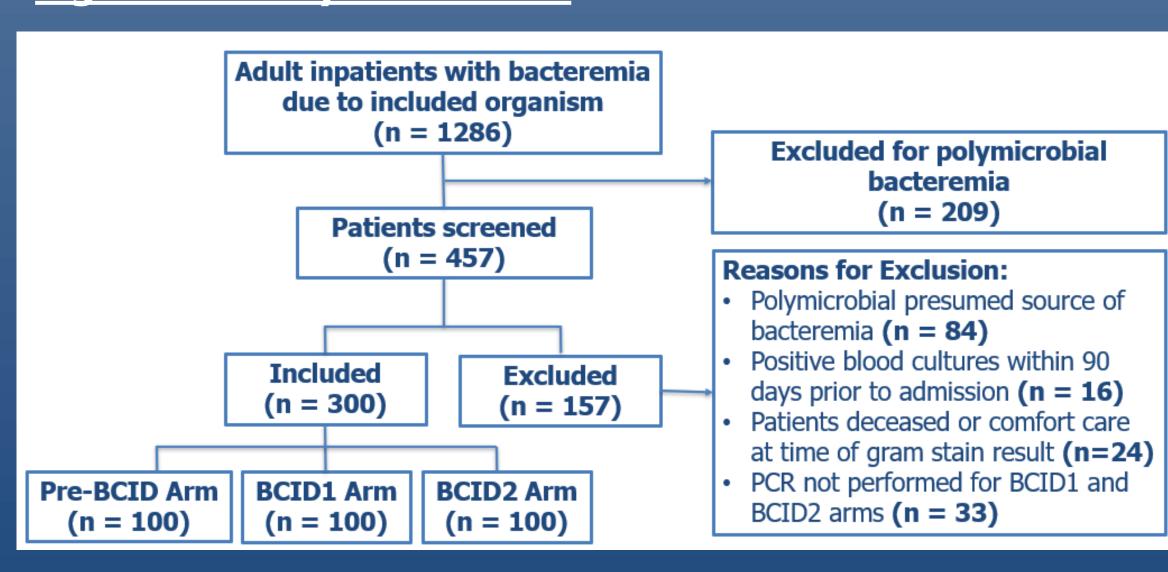
- Bloodstream infections are associated with significant morbidity, mortality, and costs
- Timely initiation of effective treatment is required for optimal outcomes
- Conventional microbiological techniques may require 72 hours for organism identification; however, rapid blood culture identification (BCID) polymerase chain reaction (PCR) panels have expedited this process
- At Atlantic Health System, the Biofire® FilmArray® was implemented June 2020 (BCID1), with subsequent upgrade to BCID2 March 2021
- Notable features of the BCID2 panel include species identification of *Enterococcus* and the addition of the CTX-M target
- The objective of this study is to assess time to optimal therapy with and without the BCID platforms for select organisms

Methods

- Retrospective chart review at two community-teaching hospitals in New Jersey
- **Study Arms:**
- Pre-BCID: May 2019 February 2020
- BCID1: June 2020 February 2021
- o BCID2: March 2021 December 2021
- Inclusion Criteria: Hospitalized patients ≥ 18 years old who have positive blood cultures for Enterococcus faecalis, E. faecium, Proteus spp, Escherichia coli, or non-aerogenes Klebsiella spp. in pre-BCID vs. BCID1 vs. BCID2 time periods
- **Exclusion Criteria**: Patients with polymicrobial or presumed polymicrobial source of bacteremia, recent positive blood cultures, PCR not performed in BCID arms, or who were deceased/comfort care at time of Gram stain
- **Statistics:** Continuous data analyzed using the student t-test and Wilcoxon rank sum test; nominal data analyzed using the chi-squared and Fisher's exact tests
- Primary Endpoint (Pre-BCID vs. BCID1/Pre-BCID vs. BCID2):
- Time to optimal therapy (TTOT) time from blood culture collection to start of pathogen-specific regimen as defined by local guidelines
- Secondary Endpoints (Pre-BCID vs. BCID1/Pre-BCID vs. BCID2):
- Time to effective therapy time from blood culture collection to initiation of antimicrobial therapy to which the organism was found to be susceptible
- Duration of vancomycin therapy
- Therapy modification 24 hours after Gram stain result
- Carbapenem use within 24 hours after Gram stain results for Gramnegative bacteremia
- Length of stay
- 30-day readmission
- Recurrence or reinfection within 30 days of discharge
- All-cause 30-day mortality

Results

Figure 1. Study Enrollment



Results

Table 1. Patient Characteristics

	Pre-BCID (n=100)	BCID1 (n=100)	BCID2 (n=100)
Age, years [median (range)]	75 (22 – 95)	74 (28 – 94)	67 (21 – 102)
Female	53	53	56
Pitt bacteremia score [median (range)]	1 (0 - 8)	1 (0 - 10)	1 (0 - 8)
ICU admission at time of Gram stain	9	13	11
Infectious diseases consult within 24 hours of Gram stain	76	84	81
Organisms E. faecalis E. faecium E. coli K. oxytoca K. pneumoniae Proteus spp.	13 2 66 0 15 4	7 2 69 1 16 5	13 3 60 1 1 14 9
Source of bacteremia Respiratory tract Endocarditis Catheter associated Urinary tract Intraabdominal Skin and soft tissue	8 2 5 65 18 2	2 2 3 69 23 1	4 2 4 60 26 4
Resistance mechanisms ESBL (CTX-M) CRE (KPC) VRE (VanA/B)	13 0 2	10 1 1	11 1 3

Data expressed as percentages unless otherwise specified

p>0.05 relative to pre-BCID, unless bolded

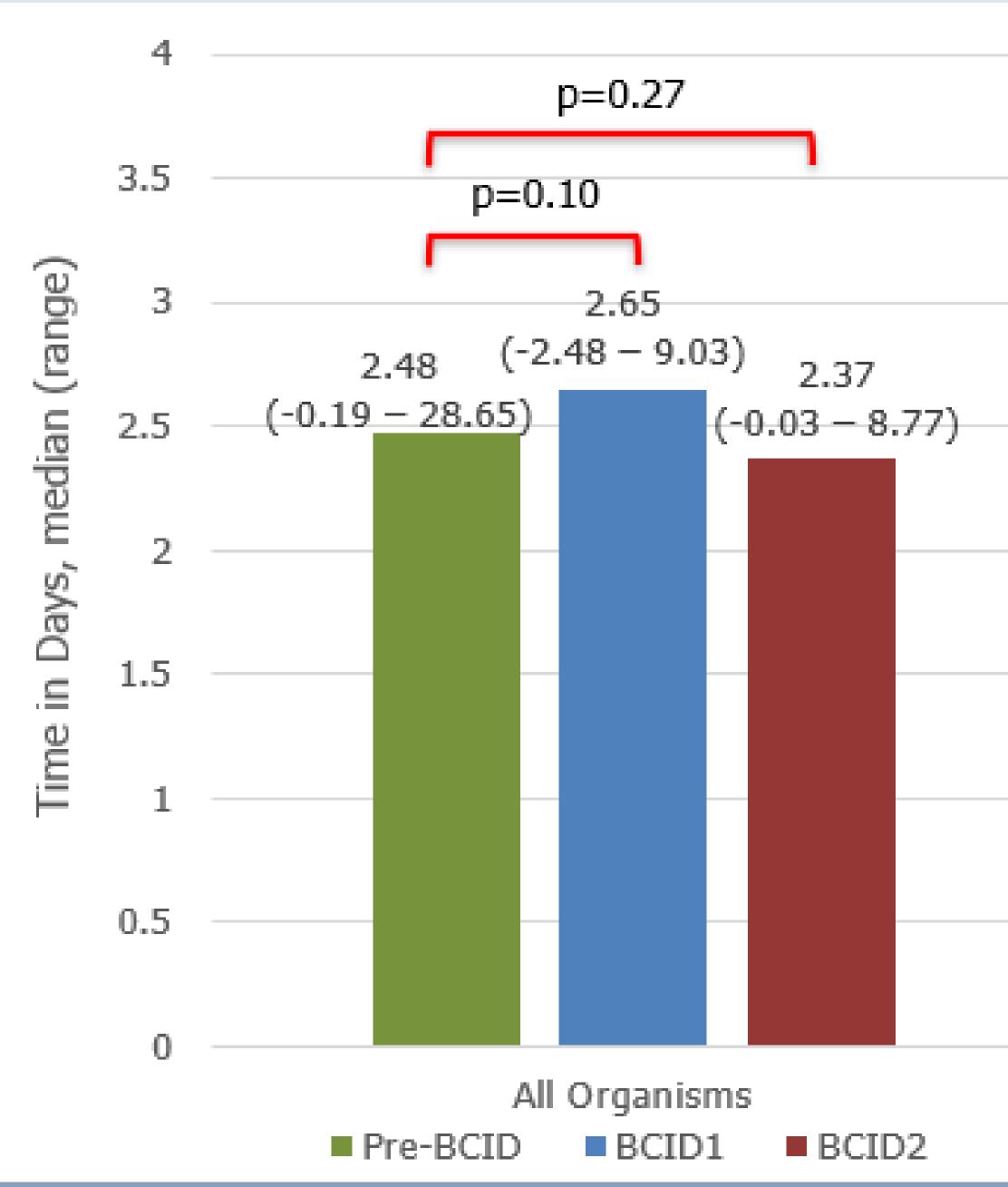
ESBL: Extended-spectrum beta-lactamases, **CRE**: Carbapenem-resistant *Enterobacterales*, **VRE**: Vancomycin-resistant *Enterococcus*

Table 2. Secondary Endpoints

Endpoint	Pre-BCID	BCID1	BCID2
	(n=100)	(n=100)	(n=100)
Time to optimal therapy, days Gram-negative Gram-positive	2.48 (-0.19 - 28.65)	2.65 (-2.48 - 9.03)	2.37 (-0.03 - 8.77)
	1.85 (-0.19 - 28.65); n=85	2.62 (-2.48 - 9.03); n= 91	2.23 (-0.03 - 7.85); n=84
	3.69 (1.17 - 6.99); n=15	2.93 (0.80 - 8.59); n=9	2.84 (0.82 - 8.77); n=16
Time to effective therapy, days Gram-negative Gram-positive	0.02 (-0.19 - 6.99)	0.05 (-2.48 - 6.06)	0.03 (-0.99 - 3.44)
	0.02 (-0.19 - 3.47); n=85	0.04 (-2.48 - 2.98); n=91	0.02 (-0.99 - 2.06); n=84
	0.01 (-0.12 - 6.99); n=15	0.84 (0.03 - 6.06); n=9	0.18 (-0.72 - 3.44) ; n=16
Duration of vancomycin, days Gram-negative Gram-positive	0 (0 - 28.59)	0 (0 - 2.65)	0.08 (0 - 4.78)
	0 (0 - 28.59); n=85	0 (0 - 2.65); n=91	0.07 (0 - 4.78); n=84
	2.46 (0 - 8.60); n=15	0.80 (0 - 2.64); n=9	0.84 (0 - 2.86); n=16
Therapy modification 24 hours after Gram stain result, %	23	30	44
Carbapenem use within 24 hours after Gram stain results for Gramnegative bacteremia, n/N (%) Non-ESBL ESBL	8/72 (11%)	18/81 (22%)	6/72 (8%)
	7/13 (54%)	2/10 (20%)	11/12 (92%)
Length of stay, days	7 (3 – 134)	7 (3 – 34)	7 (3 – 42)
30-day readmission, %	13	12	20
Recurrence or reinfection within 30 days of discharge, %	1	0	1
All-cause 30-day mortality, %	5	4	10

Data expressed as median (range), unless otherwise specified p>0.05 relative to pre-BCID, unless bolded

Figure 2. Primary Endpoint (TTOT)



Discussion/Conclusion

- There was no difference in time to optimal therapy with either the BCID1 or BICD2 panel
- In the BCID2 arm, the following findings were significant:
 - Reduction in the duration of vancomycin use
- More therapy changes within 24 hours of Gram stain result
- More appropriate carbapenem use within 24 hours of Gram stain result for gram-negative organisms
- Barriers to benefit
- Implementation at the emergence of COVID-19 pandemic limited prioritization of education efforts
- Lack of formal real-time antimicrobial stewardship team involvement
- Study Limitations:
- Retrospective design
- Generalizability
 - Local definitions of optimal antimicrobial therapy
 - Organism/source distribution (*E. coli* from urinary source was predominant)
- Future Directions
- Consideration of real-time antimicrobial stewardship team involvement

References

- 1. Nagel JL, et al. *J Clin Microbiol*. 2014;52(8):2849-54.
- 2. Messacar K, et al. *J Pediatr Infect Dis*. 2017;6(3):267-74.
- 3. Graff KE, et al. *Microbiol Spectr*. 2021;9(1):e0042921-e0042921

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