

The BioFire® FilmArray® Blood Culture Identification 2 (BCID2) Panel is an FDA-approved multiplex PCR assay that rapidly detects a limited number of commonly-identified bloodstream pathogens. While detection of organisms and resistance markers can serve as a guide for empiric antimicrobial selection, culture-based identification and susceptibility testing is required for directed antimicrobial therapy.

**Note:** All management decisions should be made with consideration of the patient’s clinical status, medication allergies, and prior recent culture/susceptibility results.

- **Considerations to warrant broader or alternative therapy recommendations:**
  - **Critical illness and/or recent history of multi-drug resistant organisms**
  - **Source of infection that may include other pathogens**
  - **Discordant BCID2 and Gram stain results – await final culture and susceptibility data**
  - **Suspected bloodstream infection based on clinical presentation, but negative BCID2**
- **^ For cases with suspected or confirmed intra-abdominal infections (IAI), consider addition of Gram-negative or anaerobic coverage (add metronidazole, piperacillin-tazobactam)**
- Please contact ADSP/ID pharmacist or consider ID consultation for other questions related to interpretation and management. For non-urgent questions email [nmhaspconsult@nm.org](mailto:nmhaspconsult@nm.org).
- See [Appendix A](#) for additional clinical considerations & *pharmacist interpretation workflow*

#### Gram-positive Bacteria:

- *Enterococcus faecalis*
- *Enterococcus faecium*
- *Listeria monocytogenes*
- *Staphylococcus aureus*
- *Staphylococcus epidermidis*
- *Staphylococcus lugdunensis*
- *Staphylococcus* species
- *Streptococcus agalactiae*
- *Streptococcus pneumoniae*
- *Streptococcus pyogenes*
- *Streptococcus* species

#### Gram-negative Bacteria:

- *Acinetobacter calcoaceticus-baumannii complex*
- *Bacteroides fragilis*
- Enterobacterales species
- *Enterobacter cloacae complex*
- *Escherichia coli*
- *Haemophilus influenzae*
- *Klebsiella aerogenes*
- *Klebsiella oxytoca*
- *Klebsiella pneumoniae* group
- *Neisseria meningitidis*
- *Proteus* species
- *Pseudomonas aeruginosa*
- *Salmonella*
- *Serratia marcescens*
- *Stenotrophomonas maltophilia*

Gram-negative Resistance Genes: CTX-M, KPC, IMP, OXA-48-like, NDM, VIM, *mcr-1*

#### Yeast:

- *Candida albicans*
- *Candida auris*
- *Candida glabrata*
- *Candida krusei*
- *Candida parapsilosis*
- *Candida tropicalis*
- *Cryptococcus neoformans/gattii*

**Table 1. Gram-positive Bacteria**

Organism	Resistance Gene	Antibiotic Recommendation	Comments
<i>Enterococcus faecalis</i>	---	Ampicillin ^	<u>Severe PCN allergy</u> : Vancomycin ^Consider piperacillin-tazobactam if concern for intra-abdominal source of infection caused by other GNRs
<i>Enterococcus faecium</i>	---	Vancomycin ^	
	<i>vanA/B</i>	Linezolid or Daptomycin ^	Daptomycin 8-12 mg/kg IV q24h
<i>Listeria monocytogenes</i>	---	Ampicillin	<u>Severe PCN allergy</u> : TMP/SMX. If dual allergy of PCN and TMP/SMX, consult ID or ADSP/ID pharmacist.
<i>Staphylococcus aureus</i>	---	Cefazolin	<u>CNS infections</u> : Use Oxacillin <i>ID consult recommended</i>
	<i>mecA/C</i> and MREJ or <i>mecA/C</i>	Vancomycin	<i>ID consult recommended</i>
<i>Staphylococcus epidermidis</i>	---	Vancomycin*	*Consider discontinuing antibiotics if only 1 of 2 cultures are positive, the patient lacks systemic signs/symptoms of infection (fever, hypotension, leukocytosis), & the patient does not have an implanted device such as hemodialysis line, PICC line, cardiac devices or prosthetic valves. This suggests contamination.
<i>Staphylococcus species</i>	---	Vancomycin	
<i>Staphylococcus lugdunensis</i>	---	Cefazolin	
	<i>mecA/C</i>	Vancomycin	
<i>Streptococcus agalactiae</i>	---	Ceftriaxone	
<i>Streptococcus pneumoniae</i>	---	Ceftriaxone	Consider addition of Azithromycin or Levofloxacin <u>CNS infection</u> : add Vancomycin & increase Ceftriaxone dosing to 2 g IV q12h
<i>Streptococcus pyogenes</i>	---	Ceftriaxone	<u>Toxic shock syndrome</u> : add Linezolid or Clindamycin IV 900 mg q8h for anti-toxin effects
<i>Streptococcus species</i>	---	Ceftriaxone	For patients with febrile neutropenia, Cefepime first line

**Table 2. Gram-negative Bacteria**

Organism	Resistance Gene	Antibiotic Recommendation	Comments
<i>Acinetobacter calcoaceticus-baumannii</i> complex	---	Ampicillin-sulbactam + Minocycline	Ampicillin-sulbactam 9 g IV q8h Minocycline 200 mg IV BID  Severe PCN allergy: Meropenem 2 g IV q8h + Minocycline 200 mg IV BID  If critically ill: <i>ID or ASP/ID pharmacist consultation is strongly recommended.</i>  Consider adding Meropenem, Cefiderocol, or Polymyxin B
<i>Bacteroides fragilis</i>	---	Piperacillin-tazobactam	Alternative for low-risk, hemodynamically-stable patients: Metronidazole ^
Enterobacterales species	---*	Cefepime ^ (Please read comments to prevent misinterpretation)	This is a broad target for all bacteria within the Enterobacterales (formerly Enterobacteriaceae) group. <b>If another target within the group is also positive (<i>E. cloacae</i>, <i>E. coli</i>, <i>Klebsiella spp.</i>, <i>Proteus spp.</i>, or <i>Salmonella</i>), follow the antibiotic recommendation for that specific organism instead as this likely represents a monomicrobial bacteremia</b>  *See below if resistance genes detected
<i>Enterobacter cloacae</i> complex	---*	Cefepime ^	*See below if resistance genes detected
<i>Escherichia coli</i>	---*	Ceftriaxone ^	*See below if resistance genes detected
<i>Haemophilus influenzae</i>	---	Ceftriaxone	CNS infection: dose Ceftriaxone 2 g IV q12h
<i>Klebsiella aerogenes</i>	---*	Cefepime ^	*See below if resistance genes detected
<i>Klebsiella oxytoca</i>	---*	Ceftriaxone ^	*See below if resistance genes detected
<i>Klebsiella pneumoniae</i> group	---*	Ceftriaxone ^	*See below if resistance genes detected
<i>Neisseria meningitidis</i>	---	Ceftriaxone	CNS infection: dose Ceftriaxone 2 g IV q12h
<i>Proteus</i>	---*	Ceftriaxone ^	*See below if resistance genes detected
<i>Pseudomonas aeruginosa</i>	---*	NMH: Cefepime  Sites other than NMH: Piperacillin-tazobactam	Alternatives: Piperacillin-tazobactam, Ceftazidime, Meropenem  If concern for MDR <i>P. aeruginosa</i> based on culture history, consider

			Ceftolozane-tazobactam with ID consultation  Septic shock: Add Tobramycin 7-10 mg/kg x 1  *See below if resistance genes detected
<i>Salmonella</i>	---*	Ceftriaxone	*See below if resistance genes detected
<i>Serratia marcescens</i>	---*	Ceftriaxone^	*See below if resistance genes detected
<i>Stenotrophomonas maltophilia</i>	---	TMP/SMX	TMP/SMX 8-12 mg/kg/day based on AdjBW  Allergy: Levofloxacin
*Resistance Genes			
Any	CTX-M	Meropenem	<i>ID consult required</i>
Any	IMP	Ceftazidime-avibactam + Aztreonam	
Any	KPC	Meropenem-vaborbactam OR Ceftazidime-avibactam	
Any	OXA-48-like	Ceftazidime-avibactam	
Any	NDM	Ceftazidime-avibactam + Aztreonam	
Any	VIM	Ceftazidime-avibactam + Aztreonam	
Any	<i>mcr-1</i>	<u>Avoid</u> Colistin/Polymyxin B	

**Table 3. Yeast**

Organism	Resistance Gene	Antifungal Recommendation	Comments
<i>Candida albicans</i>	---	Fluconazole	<i>ID consult recommended</i> Consider Micafungin if previous azole exposure or critically ill
<i>Candida auris</i>	---	Micafungin	<i>ID consult recommended</i> Micafungin 200 mg IV daily
<i>Candida glabrata</i>	---	Micafungin	<i>ID consult recommended</i>
<i>Candida krusei</i>	---	Micafungin	<i>ID consult recommended</i>
<i>Candida parapsilosis</i>	---	Fluconazole	<i>ID consult recommended</i> Consider Micafungin if previous azole exposure or critically ill
<i>Candida tropicalis</i>	---	Fluconazole	<i>ID consult recommended</i> Consider Micafungin if previous azole exposure or critically ill
<i>Cryptococcus neoformans/gattii</i>	---	Liposomal Amphotericin B + Flucytosine	<i>ID consult recommended</i> Liposomal Amphotericin B IV 3-4 mg/kg/day Flucytosine 25 mg/kg PO q6h

## Appendix A. Pharmacist Interpretation Workflow for BCID2 Results

Purpose of this section: interpretation and management for results that occur in off-peak hours (8am – 4pm) and clinical pharmacist outreach, when appropriate, to primary teams.

**Note:** All management decisions should be made conjunctively with primary teams and should be guided by the patient’s clinical status, medication allergies, and prior recent culture/susceptibility results.

Please complete the following steps while covering Blood Biofire Results:

1. Review Biofire Result inbox 3 times per shift (every 2-3 hours).
2. Assess new results and follow the below actions based the diagram below (Traffic light).
3. Contact the primary team as indicated.
4. Complete the Antimicrobial Stewardship Navigator: under intervention, choose “Rapid diagnostic” → Blood Biofire → Choose active abx → Indication→ Overall recommendations
5. Mark the inbox message as “read” but do not delete the message.

### Actions Based on Biofire Result

Must reach out to provider:	Consider reaching out to provider:	No need to reach out to provider:
<ul style="list-style-type: none"> <li>Uncovered pathogen/Drug Bug mismatch</li> <li>Complex patient or restricted agent requiring ID consult</li> <li>Unnecessary antimicrobial causing increased risk of adverse event</li> </ul> <p>High priority pathogens to review carefully and likely intervene:</p> <ul style="list-style-type: none"> <li><i>Staphylococcus aureus</i> <ul style="list-style-type: none"> <li>Discontinue vancomycin in setting of <i>Staph aureus</i> when no resistance is detected (i.e. MSSA)</li> </ul> </li> <li><i>Pseudomonas</i></li> <li><i>Candida</i></li> <li><i>Enterococcus</i></li> <li>Gram-negative resistance genes</li> </ul>	<ul style="list-style-type: none"> <li>De-escalation opportunity</li> <li>IV to PO recommendation</li> </ul> <p>Straightforward, high-value stewardship interventions:</p> <ul style="list-style-type: none"> <li>D/c vancomycin for single positive for species outlined above with high suspicion for contamination in patients without guidance listed risk factors</li> <li>Cefepime or pip-tazo de-escalation to ceftriaxone for urinary source Gram-negatives as recommended in interpretation tables above</li> </ul>	<ul style="list-style-type: none"> <li>Receiving recommended therapy</li> <li>Complex patient actively followed by ID Consult</li> </ul> <p>Examples:</p> <ul style="list-style-type: none"> <li>De-escalation opportunity: <i>E. coli</i> bacteremia from an intra-abdominal source on pip-tazo</li> <li>MRSA bacteremia on vancomycin</li> <li>Gram-negative resistance gene detected and receiving appropriate empiric coverage per ID consult</li> </ul>