Protocol for Laboratory Verification of Performance of the **BioFire® Blood Culture** Identification 2 (BCID2) Panel

A Laboratory Protocol for Use with Live **Organisms**

Purpose

The Clinical Laboratory Improvement Amendments (CLIA), passed in 1988, establishes quality standards for all laboratory testing to ensure the accuracy and reliability of patient test results, regardless of where the test is performed. The CLIA regulations include a requirement for verifying the performance specifications of unmodified, moderate complexity tests cleared or approved by the FDA.

This document provides an example of a verification procedure to assist your laboratory in developing a protocol for the verification of the BioFire BCID2 Panel performance on BioFire® FilmArray® Systems as required by CLIA. A verification scheme, compatible with the BioFire BCID2 Panel, has been designed using nonclinical specimens. This scheme provides positive and negative tests for each organism detected by the BioFire BCID2 Panel and may be easily modified or expanded to meet specific criteria.

Day-to-day variation is evaluated by testing each sample on two separate days. To evaluate user-to-user variation, multiple laboratory technicians may test the same sample. In addition, testing patient samples for verification of the performance of the BioFire BCID2 Panel should be done under the guidance of the Laboratory Director, but is not described here.

As per the CLIA regulation, the Laboratory Director is ultimately responsible for ensuring that verification procedures meet the appropriate standards for CLIA and applicable laboratory accrediting agencies.

BioFire Intended Use

The BioFire BCID2 Panel is a multiplexed nucleic acid test intended for use with BioFire® FilmArray® 2.0 or BioFire® FilmArray® Torch Systems for the simultaneous qualitative detection and identification of multiple bacterial and yeast nucleic acids and select genetic determinants

Technical Note BioFire Diagnostics, LLC www.biofiredx.com BFR0000-6592-01 QS-339B-02

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associated with antimicrobial resistance. The BioFire BCID2 Panel test is performed directly on blood culture samples identified as positive by a continuous monitoring blood culture system. Results are intended to be interpreted in conjunction with Gram stain results. The following organism types and subtypes are identified using the BioFire BCID2 Panel:

| <u> </u> | <u>v</u> | | | | | | | |
|---|----------------------------|------------------------------------|--|--|--|--|--|--|
| Gram Positive Bacteria | | | | | | | | |
| Enterococcus faecalis | Staphylococcus spp. | Streptococcus spp. | | | | | | |
| Enterococcus faecium | Staphylococcus aureus | Streptococcus agalactiae (Group B) | | | | | | |
| Listeria monocytogenes | Staphylococcus epidermidis | Streptococcus pneumoniae | | | | | | |
| | Staphylococcus lugdunensis | Streptococcus pyogenes (Group A) | | | | | | |
| | Gram Negative Bacteria | | | | | | | |
| Acinetobacter calcoaceticus-baumannii complex | | Enterobacterales | | | | | | |
| Bacteroides fragilis | | Enterobacter cloacae complex | | | | | | |
| Haemophilus influenzae | | Escherichia coli | | | | | | |
| Neisseria meningitidis (encapsulated) | | Klebsiella aerogenes | | | | | | |
| Pseudomonas aeruginosa | | Klebsiella oxytoca | | | | | | |
| Stenotrophomonas maltophilia | | Klebsiella pneumoniae group | | | | | | |
| | | Proteus spp. | | | | | | |
| | | Salmonella spp. | | | | | | |
| | | Serratia marcescens | | | | | | |
| Yeast | | | | | | | | |
| Candida albicans | Candida krusei | Cryptococcus neoformans/gattii | | | | | | |
| Candida auris | Candida parapsilosis | | | | | | | |
| Candida glabrata | Candida tropicalis | | | | | | | |

The BioFire BCID2 Panel contains assays for the detection of genetic determinants associated with resistance to methicillin (*mecA/C* and *mecA/C* in conjunction with MREJ), vancomycin (*vanA* and *vanB*), ß-lactams including penicillins, cephalosporins, monobactams, and carbapenems (*bla*_{CTX-M}, *bla*_{IMP}, *bla*_{KPC}, *bla*_{NDM}, *bla*_{OXA48-like}, *bla*_{VIM}) to aid in the identification of potentially antimicrobial-resistant organisms in positive blood culture samples. In addition, the panel includes an assay for the detection of the mobilized genetic determinant *mcr-1*, an emerging marker of public health importance. The antimicrobial resistance gene or marker detected may or may not be associated with the agent responsible for disease. Negative results for these select antimicrobial resistance gene and marker assays do not indicate susceptibility, as multiple mechanisms of resistance to methicillin, vancomycin, ß-lactams, and colistin exist.

| Antimicrobial Resistance Genes | | | | | | | | |
|--------------------------------|-------|------------------------|-------------|--------|--|--|--|--|
| CTX-M | KPC | mecA/C | NDM | vanA/B | | | | |
| IMP | mcr-1 | mecA/C and MREJ (MRSA) | OXA-48-like | VIM | | | | |

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The complete intended use statement and additional information about the use of the BioFire System can be found in the *BioFire[®] Blood Culture* Identification 2 (BCID2) Panel Instructions for Use.

Performance Verification: Overview

The verification procedure described here may be used to evaluate the performance of each assay on the BioFire BCID2 Panel. The performance verification protocol has been designed to take advantage of the multiplex nature of the BioFire BCID2 Panel. Verification testing efficiency is maximized by evaluating multiple target organisms in a single test run. The procedure described below will generate multiple positive and negative detections for each of the BioFire BCID2 assays. The procedures were developed using organism strains available from Microbiologics[®], Saint Cloud, MN. Table 3 provides a list of part numbers.

A BioFire[®] System is defined as all BioFire[®] FilmArray[®] Instruments or Modules that are connected to and controlled by a single computer system. If the laboratory director chooses not to perform the entire verification protocol on each individual instrument, it is advised that test replicates are evenly distributed among the instruments or modules. An example of a performance verification workflow using 2, 4, or 6 modules is provided in Figure 2.

Clinical/patient samples may be used in place of, or in addition to the verification scheme described here in order to assess clinical sensitivity/specificity and sample matrix effects as part of the performance verification of the BioFire BCID2 Panel.

| Table 1. Ove | | | | | | | |
|---------------------------------|------------------------|---------------------------------|-------------------------------------|---------------------|--|---------------------------------|--|
| Verification Protocol | Organisms per Poolª | Number of Sample Pools | Replicates per Sample Pool | Pouches Required | Expected Positive Results ^b | Expected Negative Results | Approximate Days of Testing ^c |
| Simple Protocol ^d | 1, 8, or 9 | 5 | 4 | 20 | ≥4 per organism | 4-16 per organism | 4 |

Table 1 Overview of Verification Protocol

^a Depending on the material used for verification, pooling of organisms may not be appropriate and the values in the table may need to be modified.

^b The expected number of positives and negatives per organism is dependent upon the number strains of a particular organism used to complete the verification. The proposed verification procedure recommends two E. coli strains; therefore the number of expected E. coli positives would be 8 and the number of expected negatives would be 12.

^c The approximate number of days for testing assumes a system configured with one instrument/module and does not include time to grow microbial cultures.

^dThis simple protocol may be easily expanded to increase the number of pouches tested on one instrument/module or for the verification of multiple instruments/modules.

Note: The BioFire BCID2 Panel contains assays for the detection of genetic determinants of resistance, including colistin (mcr-1). mcr-1, the mobilized colistin resistance gene, confers plasmid-borne resistance to colistin, a last-resort drug for some

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multidrug-resistant infections. Laboratories conducting verification of performance of the BioFire BCID2 Panel may choose to not verify the *mcr-1* colistin resistance interpretation on the BioFire BCID2 Panel. In this case, verification can be performed by preparing Pools 1 - 4 as described in Table 4. If a lab chooses to validate the *mcr-1* colistin resistance detection, the organism conferring the resistance (*Escherichia coli* CDC AR-0346) <u>should not</u> be pooled with any other organisms to reduce the risk of interspecies transfer of antibiotic resistance genes. **Proper decontamination and disposal procedures must be strictly followed**. More information can be found in the *BioFire*[®] *Blood Culture Identification 2 (BCID2) Panel Instructions for Use* and on the Center for Disease Control and Preventions (CDC) website:

https://www.cdc.gov/hicpac/2007IP/2007ip_part3.htm Infection control guidelines:

https://www.cdc.gov/infectioncontrol/guidelines/isolation/index.html#a4

Performance Verification: Materials

The following materials may be used to perform the verification procedure:

| Material | Part Number |
|---|--|
| BioFire [®] Blood Culture Identification 2 (BCID2) Panel (30-test kit) | BioFire Diagnostics, LLC RFIT-ASY-0147 |
| BioFire [®] Blood Culture Identification 2 (BCID2) Panel Instructions for Use | BioFire Diagnostics, LLC RFIT-PRT-0841 |
| BioFire [®] Blood Culture Identification 2 (BCID2) Panel Quick Guide | BioFire Diagnostics, LLC RFIT-PRT-0867 |
| Blood culture media | BACT/ALERT® FA PLUS , 410851, 442192ª (or equivalent) |
| Human Whole Blood | BioIVT, HUMANWBACDAUZN (or equivalent, with anticoagulant) |
| McFarland Turbidity Standard | Fisher Scientific, R20411 (or equivalent) |
| Phosphate Buffered Saline, pH 7.4 | Hardy Diagnostics, R201 (or equivalent) |
| Polystyrene tubes with cap (15 mL or 50 mL conical) | VWR, 82050-278 or 82050-350 (or equivalent) |
| 10-mL syringe and 18 gauge needle | VWR, 75846-756 and BD-305196 (or equivalent) |
| Serological pipette, 5 mL | VWR, 82050-478 (or equivalent) |
| Disposable Transfer pipets, graduated | VWR, 414004-024 (or equivalent) |

| Table 2. Recommended | d materials for the | verification protocol: |
|----------------------|---------------------|------------------------|
|----------------------|---------------------|------------------------|

^a SeeTable 84 in the BioFire[®] Blood Culture Identification 2 (BCID2) Panel Instructions for Use for other compatible blood culture bottle types.

Note: It is important to use pathogen free human whole blood for the verification procedure. The blood may be screened on the BioFire BCID2 Panel prior to starting the verification procedure. The optimal human whole blood specimen will be negative for all analytes tested on the BioFire BCID2 Panel.

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| Table 3. Recommended organism strain and source for verification protocol | Table 3. R | Recommended | organism | strain ar | nd source fo | or verification | protocol |
|---|------------|-------------|----------|-----------|--------------|-----------------|----------|
|---|------------|-------------|----------|-----------|--------------|-----------------|----------|

| Table 3. Recommended organism strain and source for ve | |
|---|------------------------------------|
| Organism | Microbiologics® Catalog Numberª |
| Acinetobacter baumannii ATCC® 19606™ KWIK-STIK | 0357P |
| Bacteroides fragilis ATCC® 25285™ KWIK-STIK | 0320P |
| Candida albicans ATCC® 10231™ KWIK-STIK | 0443P |
| Candida auris CDC B11903 KWIK-STIK | 01256P |
| Candida glabrata ATCC® 15126™ KWIK-STIK | 0737P |
| Candida krusei ATCC® 14243™ KWIK-STIK | 0809P |
| Candida parapsilosis ATCC® 22019™ KWIK-STIK | 0726P |
| Candida tropicalis ATCC® 1369™ KWIK-STIK | 01036P |
| Cryptococcus gattii ATCC® MYA-4560™ KWIK-STIK | 01051P |
| Cryptococus neoformans serotype A ATCC® 66031™ KWIK-STIK | 0985P |
| Enterobacter cloacae subsp.cloacae ATCC® 13047™ KWIK-STIK | 0323P |
| Enterococcus faecalis ATCC® 51299™ KWIK-STIK | 0959P ^b |
| Enterococcus faecium NCTC 12204 KWIK-STIK | 01143P° |
| Escherichia coli NCTC 13476 KWIK-STIK | 01136P ^d |
| Escherichia coli CDC AR-0346 KWIK-STIK | 01259P ^e |
| Haemophilus influenzae ATCC® 10211™ KWIK-STIK | 0441P |
| Klebsiella aerogenes ATCC® 35029™ KWIK-STIK | 0399P |
| Klebsiella oxytoca ATCC® 13182™ KWIK-STIK | 0530P |
| Klebsiella pneumoniae NCTC 13443 KWIK-STIK | 01145P ^f |
| Klebsiella pneumoniae ATCC® BAA-1705™ KWIK-STIK | 01005P ^g |
| Klebsiella pneumoniae CDC AR-0039 KWIK-STIK | 01241P ^h |
| Klebsiella pneumoniae NCTC 13440 KWIK-STIK | 01112P ⁱ |
| Listeria monocytogenes ATCC® 19111™ KWIK-STIK | 0277P |
| Neisseria meningitidis ATCC® 13077™ KWIK-STIK | 0453P |
| Proteus mirabilis ATCC® 35659™ KWIK-STIK | 0944P |
| Pseudomonas aeruginosa ATCC® 27853™ KWIK-STIK | 0353P |
| Salmonella enterica subsp. arizonae ATCC® 13314™ KWIK-STIK | 0901P |
| Serratia marcescens ATCC® 13880™ KWIK-STIK | 0247P |
| Staphylococcus aureus subsp. aureus ATCC® 33591™ KWIK-STIK | 0496P ^j |
| Staphylococcus epidermidis ATCC® 12228™ KWIK-STIK | 0371P |
| Staphylococcus lugdunensis ATCC® 49576™ KWIK-STIK | 0217P |
| Stenotrophomonas maltophilia ATCC® 17666™ KWIK-STIK | 0759P |
| Streptococcus agalactiae ATCC® 12386™ KWIK-STIK | 0439P |
| Streptococcus pneumoniae ATCC® 10015™ KWIK-STIK | 0865P |
| Streptococcus pyogenes ATCC® 19615™ KWIK-STIK | 0385P |
| Blood Culture Identification 2 (BCID2) Verification Panel (Live Culture) | 5258P ^k |
| Any appropriate source of organism may be used for verification of any or all of th | |

^a Any appropriate source of organism may be used for verification of any or all of the assays in the BioFire BCID2 Panel. However, when alternate organism sources are used, the sample volumes or pooling schemes suggested in the examples below may need to be adjusted. Alternate organism strains may not provide the same results for antimicrobial resistance genes as those suggested here.

^b This strain of *E. faecalis* (ATCC® 51299) carries the *vanB* gene (vancomycin resistance) .

^c This strain of *E. faecium*(NCTC 12204) carries the *vanA* gene (vancomycin resistance).

^d This strain of *E. coli* (NCTC 13476) carries the IMP gene (ß-lactam resistance).

^e This strain of *E. coli* (CDC AR-0346) carries the *mcr-1* gene (colistin resistance) and the CTX-M gene (ß-lactam resistance).

^f This strain of *K. pneumoniae* (NCTC 13443) carries the CTX-M and NDM-1 genes (ß-lactam resistance)

⁹ This strain of K. pneumoniae (ATCC® BAA-1705) carries the KPC gene (ß-lactam resistance).

^h This strain of K. pneumoniae (CDC AR-0039) carries the CTX-M and OXA-48 like genes (ß-lactam resistance).

ⁱ This strain of *K. pneumoniae* (NCTC 13440) carries the VIM-1 gene (ß-lactam resistance).

^j This strain of *S. aureus* subsp. *aureus* (ATCC® 33591) carries the *mecA* and MREJ (MRSA) genes (methicillin resistance).

^k This is a bundled part number containing all the organisms listed above.

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Note: Extreme care should be used in properly handling and disposing of organisms containing antibiotic resistance genes. Follow your institution's guidelines for proper handling and disposal of pathogens or refer to the CDC Infection control guidelines: https://www.cdc.gov/infectioncontrol/guidelines/isolation/index.html#a4

Performance Verification Protocol

The protocol described below evaluates the BioFire BCID2 Panel performance by pooling together up to 9 different organism suspensions in a simulated blood culture matrix. The pooling scheme described in Table 4 should be followed to obtain the expected number of positive and negative results for each organism in a time and resource-efficient manner.

Note: Dilution of organisms beyond levels proposed in these guidelines may lead to inconsistent results and is not recommended.

The protocol and workflow schemes (Figures 1 and 2) illustrate testing 4 replicates per pool for 5 pools over multiple days. This produces a total of 20 verification sample test runs and provides at least 4 positive results and as many as 16 negative results per assay. Some organisms, such as *Klebsiella pneumoniae*, are represented multiple times. This is done to ensure all antimicrobial resistance genes are represented in the verification protocol.

The number of samples tested per day should be determined by the individual laboratory. This testing scheme can be modified to run more samples per day based on the number of modules in the BioFire[®] System. The pooling scheme provides sufficient volume for testing more replicates if desired.

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| Fable 4 . Proposed Organism | | | Blood | Humon | |
|---|----------------------------|--------------------|-----------------------------|-----------------------------------|----------------------------|
| Organism and Resistance Genes | Microbiologics Part No. | Organism Volume | Culture Medium Volume | Human Whole Blood Volume | Approximate Pool Volume |
| Pool 1 | | | | | |
| Candida albicans | 0443P | 0.1 mL | | | |
| Candida krusei | 0809P | 0.1 mL | | | |
| Klebsiella pneumoniae (CTX-M, NDM) | 01145P | 0.1 mL | | | |
| Listeria monocytogenes | 0277P | 0.1 mL | | | |
| Neisseria meningitidis | 0453P | 0.1 mL | 8 mL | 3 mL | 12 mL |
| Pseudomonas aeruginosa | 0353P | 0.1 mL | | | |
| Staphylococcus aureus (mecA and MREJ) | 0496P | 0.1 mL | | | |
| Streptococcus agalactiae | 0439P | 0.1 mL | | | |
| Streptococcus pyogenes | 0385P | 0.1 mL | | | |
| Pool 2 | | | | | |
| Acinetobacter baumannii | 0357P | 0.1 mL | | | |
| Candida glabrata | 0737P | 0.1 mL | | | |
| Candida parapsilosis | 0726P | 0.2 mL | | | |
| Candida tropicalis | 01036P | 0.4 mL | 0 | 3 mL | 10 |
| Escherichia coli (IMP) | 01136P | 0.2 mL | 8 mL | | 12 mL |
| Klebsiella oxytoca | 0530P | 0.1 mL | | | |
| Klebsiella pneumoniae (KPC) | 1005P | 0.1 mL | | | |
| Staphylococcus epidermidis | 0371P | 0.4 mL | | | |
| Pool 3 | | | | | |
| Candida auris | 01256P | 0.2 mL | | | |
| Cryptococcus gattii | 01051P | 0.1 mL | | | |
| Enterococcus faecalis (vanB) | 0959P | 0.2 mL | | | |
| Haemophilus influenzae | 0411P | 0.1 mL | | | |
| Klebsiella pneumoniae (VIM) | 01112P | 0.1 mL | 8 mL | 3 mL | 12 mL |
| Proteus mirabilis | 0944P | 0.2 mL | | | |
| Serratia marcescens | 0247P | 0.1 mL | | | |
| Staphylococcus lugdunensis | 0217P | 0.1 mL | | | |
| Streptococcus pneumoniae | 0865P | 0.1 mL | | | |
| Pool 4 | | | | | |
| Klebsiella pneumoniae (CTX-M, OXA-48 like) | 01241P | 0.1 mL | | | |
| Bacteroides fragilis | 0320P | 0.1 mL | | | |
| Cryptococcus neoformans | 0985P | 0.1 mL | | | |
| Enterobacter cloacae | 0323P | 0.1 mL | 8 mL | 3 mL | 12 mL |
| Enterococcus faecium (vanA) | 01143P | 0.1 mL | | | |
| Klebsiella aerogenes | 0399P | 0.1 mL | | | |
| Salmonella enterica typhimurium | 0901P | 0.1 mL | | | |
| Stenotrophomonas maltophilia | 0759P | 0.1 mL | | | |
| Pool 5 | | | | | |
| Escherichia coli (CTX-M, mcr-1) | 01259P | 0.2 mL | 8 mL | 3 mL | 11 mL |

Table 4. Proposed Organism Pooling Scheme

Note: *E.coli* (CTX-M. *mcr-1*) <u>should not</u> be pooled and should be properly disposed of to avoid possible transfer of the resistance gene to other bacteria. Follow your institution's guidelines for proper handling and disposal of pathogens or refer to the CDC Infection control guidelines:

https://www.cdc.gov/infectioncontrol/guidelines/isolation/index.html#a4

Protocol Example

The estimated total time for completion for this verification example is 4 days for a BioFire System configured with 1 module, not including the time required to grow microbial cultures.

A proposed organism pooling scheme is presented above in Table 4. Figure 1 illustrates a simplified workflow schematic. The number of samples tested per day should be determined by the individual laboratory. The protocol can be modified to run more samples per day (or fewer) based upon the number of modules in the BioFire System. The proposed organism pooling scheme in Table 4 provides sufficient volume for testing more replicates, if desired. Figure 2 provides an examples of user-to-user, day-to-day, and module-to-module testing for labs with multiple BioFire Modules.

Note: Dilution of organisms beyond levels proposed in these guidelines may lead to inconsistent results and is not recommended.

Pooled samples may be stored overnight (or up to 2 days) at refrigeration temperature (2–8°C) for subsequent testing to evaluate day-to-day variation. To evaluate user-to-user variation, multiple laboratory technicians may perform testing.

Note: It is important to prepare only the number of sample pools that will be tested within 3 days of preparation. The suggestion to prepare 3 sample pools is based on testing up to 6 pouches per day. The number of samples prepared may be increased or decreased based on the laboratory's work schedule and number of modules connected within a BioFire System.

<u>Day 1</u>

- 1. Organize materials needed (Table 2). Human whole blood should be screened in the BioFire BCID2 Panel in order to characterize the sample prior to preparing pools. The optimal human whole blood specimen will be negative for all organisms tested on the BioFire BCID2 Panel.
- 2. Obtain a pure culture of each organism that has been streaked for isolation on agar media appropriate for the organism. It is recommended to use agar plate cultures that are less than 1 week old. See Microbiologics product insert for use of KWIK-STIK cultures.

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- Prepare a suspension of each organism equivalent to McFarland turbidity standard 1.0 using approximately 3 mL of phosphate buffered saline (PBS), pH 7.4.
- 4. Prepare three sample pools according to the organism pooling scheme presented in Table 4. The sample pool preparation worksheet in the Appendix may be useful to ensure all components are added to each pool.
 - a. Use a serological pipette to add 3 mL of human whole blood to a 15 mL conical bottom tube. Alternatively, a 50 mL tube may be used.
 - b. Use a 10-mL syringe and an 18 gauge needle to remove 8 mL of blood culture medium from a blood culture bottle and add it to the conical tube containing whole blood. Care should be taken to minimize transferring resin beads into the sample.
 - c. Use a disposable transfer pipette to add the appropriate volume (Table 4) of organism suspension to the blood-medium mixture.
 - d. Repeat for the remaining organisms in the pool to combine the appropriate organisms into a single tube.
 - e. Ensure the pooled sample is effectively mixed prior to removing a sample for testing.
- Prepare and test two replicates from a single sample pool (i.e. Figure 1: Pool #1 replicates A and B). The replicate samples should be tested in a single day by different users.

Note: For each sample, follow instructions in the *BioFire® Blood Culture Identification 2 (BCID2) Panel Instruction Booklet* and *BioFire® Blood Culture Identification 2 (BCID2) Panel Quick Guide* for pouch preparation, pouch hydration, sample loading, and sample testing.

- 6. Repeat Step 5 for the remaining sample pool replicates to be tested that day (i.e. Pool 2 replicates A and B)
- 7. Refrigerate samples (2–8°C) for up to 2 days for the evaluation of dayto-day variation.

<u>Day 2</u>

To evaluate day-to-day variation, test replicates from the same sample pools prepared on Day 1 by repeating Step 5 above (i.e. replicates C and D).

<u>Day 3</u>

Prepare 2 new sample pools (i.e. pools #4 and #5) as described in Steps 3 to 4. Test replicates as described in Step 5 above.

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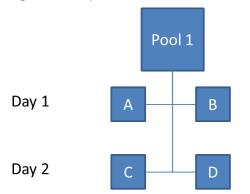
<u>Day 4</u>

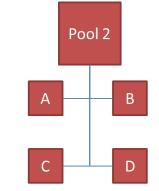
To evaluate day-to-day variation, test replicates from the same sample pools prepared on Day 3 by repeating Step 5 above (i.e. replicates C and D).

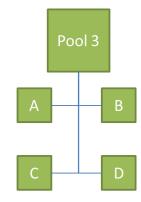
Note: Extreme care should be used in properly handling and disposing of organisms containing antibiotic resistance genes. Follow your institution's guidelines for proper handling and disposal of pathogens or refer to the CDC Infection control guidelines: https://www.cdc.gov/infectioncontrol/guidelines/isolation/index.html#a4

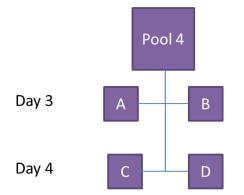


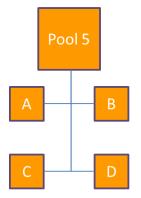












| Modules | | | | | | | |
|--------------|--------------------|--------------------|--------------------|--------------------|--------------------|----------------|--|
| 2 modules | Module 1 | | | Module 2 | | | |
| Day 1 | Pool 1/ User 1 | Pool 2/ User 2 | Pool 3 / User 1 | Pool 1/ User 2 | Pool 2/ User 1 | Pool 3 User | |
| Day 2 | Pool 1/ User 2 | Pool 2/ User 1 | Pool 3/ User 2 | Pool 1/ User 1 | Pool 2/ User 2 | Pool 3 User | |
| Day 3 | Pool 4 / User 1 | Pool 5 / User 2 | | Pool 4 / User 2 | Pool 5 / User 1 | | |
| Day 4 | Pool 4 / User 2 | Pool 5 / User 1 | | Pool 4 / User 1 | Pool 5 / User 2 | | |

Figure 2. Example of a Verification workflow for use with multiple BioFire Modules

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| 4 modules | Module 1 | | Module 2 | | Module 3 | | Module 4 | |
|--------------|--------------------|-------------------|--------------------|-------------------|--------------------|-------------------|-------------------|-------------------|
| Day 1 | Pool 1/ User 1 | Pool 2/ User 2 | Pool 1/ User 2 | Pool 2/ User 1 | Pool 3 / User 1 | | Pool 3/ User 2 | |
| Day 2 | Pool 3 / User1 | | Pool 3/ User 2 | | Pool 1/ User 1 | Pool 2/ User 2 | Pool 1/ User 2 | Pool 2/ User 1 |
| | Pool 4 / | | Pool 4 / | | Pool 5 / | | Pool 5 / User | |
| Day 3 | User 1 | | User 2 | | User 2 | | 1 Pool 4 | |
| Day 4 | Pool 5 / User 1 | | Pool 5 / User 2 | | Pool 4 / User 1 | | / User 2 | |

| 6 | Module | Module | Module | Module | Module | Module |
|---------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
| modules | 1 | 2 | 3 | 4 | 5 | 6 |
| Day 1 | Pool 1/ | Pool 1/ | Pool 2/ | Pool 2/ | Pool 3 / | Pool 3/ |
| | User 1 | User 2 | User 1 | User 2 | User 1 | User 2 |
| Day 2 | Pool 3/ | Pool 3 / | Pool 1/ | Pool 1/ | Pool 2/ | Pool 2/ |
| | User 2 | User1 | User 2 | User 1 | User 2 | User 1 |
| Day 3 | | | Pool 4 / User 1 | Pool 4 / User 2 | Pool 5 / User 1 | Pool 5 / User 2 |
| Day 4 | Pool 5 / User 1 | Pool 4 / User 2 | Pool 5 / User 2 | | Pool 4 / User 1 | |

Expanding the protocols

The protocol described above can be expanded by increasing the number of tests from each of the organism pools. Each organism pool contains

approximately 12 mL, which is enough material to complete many replicates for each pool.

Verification of Loaner, Repaired, and Permanent Replacement Instruments

If it becomes necessary to verify the performance of a loaner, repaired, or permanent replacement instrument, the following protocol <u>may</u> serve as a guideline but should be verified by the Laboratory Director.

- Select a few specimens and/or proficiency samples (any combination of positives and negatives) previously tested on the BioFire BCID2 Panel. The Laboratory Director should determine the appropriate number of samples to test. Proficiency samples should not be pooled or diluted.
- 2. Select a set of controls that verify detection of all targets on the BioFire BCID2 Panel.
- 3. Test the selected samples on the loaner, repaired, or permanent replacement instrument and document the results.

Technical Support Contact Information

BioFire is dedicated to providing the best customer support available. If you have any questions or concerns about this process, please contact the BioFire Technical Support team for assistance.

BioFire Technical Support

Email: <u>support@biofiredx.com</u> Phone: +1-801-736-6354, select Option 5



* All product names, trademarks and registered trademarks are property of their respective owners.

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Appendix

Sample Pool Preparation Worksheet

| Organism and Resistance Genes | Or | ganism olume | Blood Culture Medium Volume | Human Whole Blood Volume | Approximate Pool Volume |
|--|----|-----------------|-----------------------------------|-----------------------------|----------------------------|
| | | 0.4 ml | | | |
| Candida albicans | | 0.1 mL | - | | |
| Candida krusei | | 0.1 mL | - | | |
| Klebsiella pneumoniae (CTX-M, NDM) | | 0.1 mL | - | | |
| Listeria monocytogenes | | 0.1 mL | - | | |
| Neisseria meningitidis | | 0.1 mL | 🗌 8 mL | 🗌 3 mL | ∼12 mL |
| Pseudomonas aeruginosa | | 0.1 mL | - | | |
| Staphylococcus aureus (mecA and MREJ) | | 0.1 mL | - | | |
| Streptococcus agalactiae | | 0.1 mL | - | | |
| Streptococcus pyogenes | | 0.1 mL | | | |
| Pool 2 | | | 1 | 1 | |
| Acinetobacter baumannii | | 0.1 mL | - | | |
| Candida glabrata | | 0.1 mL | | | |
| Candida parapsilosis | | 0.2 mL | | | |
| Candida tropicalis | | 0.4 mL | | | |
| Escherichia coli (IMP) | | 0.2 mL | 🗌 8 mL | ☐ 3 mL | ~ 12 mL |
| Klebsiella oxytoca | | 0.1 mL | | | |
| Klebsiella pneumoniae (KPC) | | 0.1 mL | | | |
| Staphylococcus epidermidis | | 0.4 mL | | | |
| Pool 3 | 3 | - | | | |
| Candida auris | | 0.2 mL | | | |
| Cryptococcus gattii | | 0.1 mL | - | | |
| Enterococcus faecalis (vanB) | | 0.2 mL | | | |
| Haemophilus influenzae | | 0.1 mL | | | |
| Klebsiella pneumoniae (VIM) | | 0.1 mL | □ 8 mL | □ 3 mL | ~12 mL |
| Proteus mirabilis | | 0.2 mL | | | |
| Serratia marcescens | | 0.1 mL | | | |
| Staphylococcus lugdunensis | | 0.1 mL | - | | |
| Streptococcus pneumoniae | | 0.1 mL | - | | |
| Pool 4 | | | | | |
| <i>Klebsiella pneumoniae</i> (CTX-M, OXA-48 like) | | 0.1 mL | | | |
| Bacteroides fragilis | | 0.1 mL | | | |
| Cryptococcus neoformans | | 0.1 mL | | | |
| Enterobacter cloacae | | 0.1 mL | | _ | 12 mL |
| Enterococcus faecium (vanA) | | 0.1 mL | □ 8 mL | 🗌 3 mL | 12 111 |
| Klebsiella aerogenes | | 0.1 mL | 1 | | |
| Salmonella enterica typhimurium | | 0.1 mL | | | |
| Stenotrophomonas maltophilia | | 0.1 mL | | | |
| Pool 5 | | | | | |
| Escherichia coli (CTX-M, mcr-1) | | 0.2 mL | 🗆 8 mL | 🗌 3 mL | 11 mL |

TECHNICAL ::: NOTE



BioFire® Blood Culture Identification 2 (BCID2) Panel Verification Record-Page 1

TECHNICAL

| BioFire® Blood Culture Identification 2 (BCID2) Panel Module Serial # Kit Part # Module Serial # | | | | | | | | | | Module Serial # Module Serial # | | | | | | | | | | | | | | | | | |
|---|---------------------------------------|-----------------|-----|--------|-----|-----------------|-----|-----|-----|------------------------------------|-----|-------|-----|-----------------|-----|-----|-----|-----|-----|-----|-----|-------------|-------------|---------|--------|-----------|---------------------|
| Lot # | | | | | | Module Serial # | | | | | | | | Module Serial # | | | | | | | | | | | | | |
| | | Replicate Testi | | | | | | | | | | sting | | | | | | | | | | | Summary | | | | |
| | Organism and Resistance Genes | 1-A | 1-B | - Ļ | 1-D | 2-A | 2-B | 2-C | 2-D | 3-A | 3-B | 3.C | 3-D | 4-A | 4-B | 4-C | 4-D | 5-A | 5-B | 5-C | 5-D | # Positives | # Negatives | # Users | # Days | # Modules | Patient Samples? |
| Pool 1 | Candida albicans | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Candida krusei | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Klebsiella pneumoniae (CTX-M, NDM) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Listeria monocytogenes | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Neisseria meningitidis (encapsulated) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Pseudomonas aeruginosa | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Staphylococcus aureus (mecA and MREJ) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Streptococcus agalactiae | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Streptococcus pyogenes | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | CTX-M | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | mecA/C and MREJ (MRSA) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | NDM | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Acinetobacter baumannii | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Candida glabrata | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Candida parapsilosis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Candida tropicalis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| 12 | Escherichia coli (IMP) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Pool 2 | Klebsiella oxytoca | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Klebsiella pneumoniae (KPC) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | Staphylococcus epidermidis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | KPC | | | | | | | | | | | | | | | | | | | | | | | | | | |
| | IMP | | | | | | | 1 | | | | | | | | | | | | | | | | | | | |



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| | TEALINIAN | | | | | | | | | | | | | | | | | | | | | | | | | |
|--|-----------|-------------------|-------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|---------|-----|-----|-----|-------------|-------------|---------|--------|-----------|------------------|
| | | Replicate Testing | | | | | | | | | | | | | | | Summary | | | | | | | | | |
| Organism and Resistance Genes | 1-A | 1-B | - | 1-D | 2-A | 2-B | 2-C | 2-D | 3-A | 3-B | 3-C | 3-D | 4-A | 4-B | 4-C | 4-D | 5-A | 5-B | 5-C | 5-D | # Positives | # Negatives | # Users | # Days | # Modules | Patient Samples? |
| Candida auris | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Cryptococcus gattii | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Enterococcus faecalis (vanB) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Haemophilus influenzae | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Klebsiella pneumoniae (VIM) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Proteus mirabilis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Serratia marcescens | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Staphylococcus lugdunensis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Streptococcus pneumoniae | | | | | | | | | | | | | | | | | | | | | | | | | | |
| vanA/B | | | | | | | | | | | | | | | | | | | | | | | | | | |
| VIM | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Klebsiella pneumoniae (CTX-M, OXA-48 like) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Bacteroides fragilis | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Cryptococcus neoformans | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Enterobacter cloacae | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Enterococcus faecium (vanA) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Klebsiella aerogenes | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Salmonella sp. | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Stenotrophomonas maltophilia | | | | | | | | | | | | | | | | | | | | | | | | | | |
| CTX-M | | | | | | | | | | | | | | | | | | | | | | | | | | |
| OXA-48 like | | | | | | | | | | | | | | | | | | | | | | | | | | |
| vanA/B | | | | | | | | | | | | | | | | | | | | | | | | | | |
| Escherichia coli (CTX-M, mcr-1) | | | | | | | | | | | | | | | | | | | | | | | | | | |
| CTX-M | | | | | | | | | | | | | | | | | | | | | | | | | | |
| mcr-1 | | | | | | | | | | | | | | | | | | | | | | | | | | |

Reviewed by:

Signature

Date