

BioFire Defense, LLC David Rabiger Associate Director of Regulatory and Clinical Affairs 79 W 4500 S, Suite 14 Salt Lake City, Utah 84107 March 22, 2023

Re: K213362

Trade/Device Name: BioFire Global Fever Special Pathogens Panel; BIOFIRE SHIELD Control Kit

for the BioFire Global Fever Special Pathogens Panel

Regulation Number: 21 CFR 866.4000

Regulation Name: Device To Detect And Identify Biothreat Microbial Agents In Human Clinical

Specimens.

Regulatory Class: Class II

Product Code: QVR, QMV, PMN

Dated: October 8, 2021 Received: October 12, 2021

#### Dear David Rabiger:

We have reviewed your Section 510(k) premarket notification of intent to market the device referenced above and have determined the device is substantially equivalent (for the indications for use stated in the enclosure) to legally marketed predicate devices marketed in interstate commerce prior to May 28, 1976, the enactment date of the Medical Device Amendments, or to devices that have been reclassified in accordance with the provisions of the Federal Food, Drug, and Cosmetic Act (Act) that do not require approval of a premarket approval application (PMA). You may, therefore, market the device, subject to the general controls provisions of the Act. Although this letter refers to your product as a device, please be aware that some cleared products may instead be combination products. The 510(k) Premarket Notification Database located at <a href="https://www.accessdata.fda.gov/scripts/cdrh/cfdocs/cfpmn/pmn.cfm">https://www.accessdata.fda.gov/scripts/cdrh/cfdocs/cfpmn/pmn.cfm</a> identifies combination product submissions. The general controls provisions of the Act include requirements for annual registration, listing of devices, good manufacturing practice, labeling, and prohibitions against misbranding and adulteration. Please note: CDRH does not evaluate information related to contract liability warranties. We remind you, however, that device labeling must be truthful and not misleading.

If your device is classified (see above) into either class II (Special Controls) or class III (PMA), it may be subject to additional controls. Existing major regulations affecting your device can be found in the Code of Federal Regulations, Title 21, Parts 800 to 898. In addition, FDA may publish further announcements concerning your device in the <u>Federal Register</u>.

Please be advised that FDA's issuance of a substantial equivalence determination does not mean that FDA has made a determination that your device complies with other requirements of the Act or any Federal statutes and regulations administered by other Federal agencies. You must comply with all the Act's

requirements, including, but not limited to: registration and listing (21 CFR Part 807); labeling (21 CFR Part 801 and Part 809); medical device reporting (reporting of medical device-related adverse events) (21 CFR 803) for devices or postmarketing safety reporting (21 CFR 4, Subpart B) for combination products (see <a href="https://www.fda.gov/combination-products/guidance-regulatory-information/postmarketing-safety-reporting-combination-products">https://www.fda.gov/combination-products/guidance-regulatory-information/postmarketing-safety-reporting-combination-products</a>); good manufacturing practice requirements as set forth in the quality systems (QS) regulation (21 CFR Part 820) for devices or current good manufacturing practices (21 CFR 4, Subpart A) for combination products; and, if applicable, the electronic product radiation control provisions (Sections 531-542 of the Act); 21 CFR 1000-1050.

Also, please note the regulation entitled, "Misbranding by reference to premarket notification" (21 CFR Part 807.97). For questions regarding the reporting of adverse events under the MDR regulation (21 CFR Part 803), please go to <a href="https://www.fda.gov/medical-devices/medical-device-safety/medical-device-reporting-mdr-how-report-medical-device-problems">https://www.fda.gov/medical-device-problems</a>.

For comprehensive regulatory information about medical devices and radiation-emitting products, including information about labeling regulations, please see Device Advice (<a href="https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance">https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance</a>) and CDRH Learn (<a href="https://www.fda.gov/training-and-continuing-education/cdrh-learn">https://www.fda.gov/training-and-continuing-education/cdrh-learn</a>). Additionally, you may contact the Division of Industry and Consumer Education (DICE) to ask a question about a specific regulatory topic. See the DICE website (<a href="https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance/contact-us-division-industry-and-consumer-education-dice">https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance/contact-us-division-industry-and-consumer-education-dice">https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance/contact-us-division-industry-and-consumer-education-dice</a>) for more information or contact DICE by email (DICE@fda.hhs.gov) or phone (1-800-638-2041 or 301-796-7100).

Sincerely,

# Noel J. Gerald -S

Noel J. Gerald, Ph.D.
Branch Chief
Bacterial Respiratory and Medical Countermeasures Branch
Division of Microbiology Devices
OHT7: Office of In Vitro Diagnostics
Office of Product Evaluation and Quality
Center for Devices and Radiological Health

Enclosure

# DEPARTMENT OF HEALTH AND HUMAN SERVICES Food and Drug Administration

#### **Indications for Use**

Form Approved: OMB No. 0910-0120

Expiration Date: 06/30/2023
See PRA Statement below.

510(k) Number (if known)

K213362

**Device Name** 

BioFire Global Fever Special Pathogens Panel

Indications for Use (Describe)

The BioFire® Global Fever Special Pathogens Panel is a qualitative, multiplexed, nucleic acid-based test intended for use with BioFire® FilmArray® 2.0 and BioFire® FilmArray® Torch Systems. The BioFire Global Fever Special Pathogens Panel is for the simultaneous qualitative detection and identification of multiple bacterial, viral, and protozoan nucleic acids directly from EDTA whole blood collected from individuals with signs and/or symptoms of acute febrile illness or recent acute febrile illness and known or suspected exposure to the target pathogens described below.

#### Pathogens identified:

Chikungunya virus

Dengue virus (serotypes 1, 2, 3 and 4)

Leishmania spp. that cause visceral leishmaniasis (e.g., L. donovani and L. infantum)

Leptospira spp.

Plasmodium spp. (including species differentiation of Plasmodium falciparum and Plasmodium vivax/ovale)

West Nile virus

#### Pathogens presumptively identified:

Bacillus anthracis

Crimean-Congo hemorrhagic fever virus

Ebolavirus spp.

Francisella tularensis

Lassa virus

Marburgvirus

Yellow fever virus

Yersinia pestis

Pathogens for which the panel provides presumptive identification results require additional testing and confirmation procedures in consultation with the appropriate public health authorities for whom reports may be necessary.

Positive results do not rule out co-infections with pathogens not included on the BioFire Global Fever Special Pathogens Panel. Not all pathogens that cause acute febrile illness are detected by this test, and negative results do not preclude infection with the pathogens targeted by the device and should not be used as the sole basis for diagnosis, treatment, or other patient management decisions.

Evaluation for more common causes of acute febrile illness (e.g., infections of the upper and lower respiratory tract or gastroenteritis, as well as non-infectious causes) should be considered prior to evaluation with this panel. In the United States, patient travel history, exposure risk, and consultation of the CDC Yellow Book should be considered prior to use of the BioFire Global Fever Special Pathogens Panel as some pathogens are more common in certain geographical locations. Results are meant to be used in conjunction with other clinical, epidemiologic, and laboratory data, in accordance with the guidelines provided by the relevant public health authorities.

The BioFire Global Fever Special Pathogens Panel is indicated for use in laboratories having appropriate biosafety equipment, personal protective equipment (PPE), containment facilities and personnel trained in the safe handling of diagnostic clinical specimens potentially containing any of the pathogens detected by this panel.

The BioFire Global Fever Special Pathogens Panel is indicated for use in laboratories that follow public health guidelines that address appropriate biosafety conditions, interpretation of test results, and coordination of findings with public health authorities.

#### For In Vitro Diagnostic Use.

|   | ATE PAGE IF NEEDED.                         |
|---|---|
| Type of Use (Select one or both, as applicable)  Prescription Use (Part 21 CFR 801 Subpart D) | Over-The-Counter Use (21 CFR 801 Subpart C) |

This section applies only to requirements of the Paperwork Reduction Act of 1995.

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# 510(k) Summary

#### I. Submitter

BioFire Defense, LLC 79 West 4500 South, Suite 14 Salt Lake City, UT 84107 Phone: (801) 262-3592

Fax: (801) 447-6907

Contact Person: David Rabiger Date Prepared: 2022-Oct-07

#### II. Device

Name of Device: BioFire® Global Fever Special Pathogens Panel

Common or Usual Name: Same

**Product Code: QMV** 

**Regulation:** 21 CFR 866.3966

Classification Name: Device to detect and identify selected microbial agents that cause acute

febrile illness

**Regulatory Class:** Class II (Special Controls)

**Panel:** Microbiology – 83

**Additional Product Code:** QVR **Regulation:** 21 CFR 866.4000

Classification Name: A multiplex nucleic acid detection system for biothreat agents

**Regulatory Class:** Class II (Special Controls)

**Panel:** Microbiology – 83

#### **III.** Predicate Devices

BioFire<sup>®</sup> Global Fever Panel (BioFire Defense, LLC) [DEN200043] This predicate has not been subject to a design-related recall.

FilmArray® NGDS Warrior Panel (BioFire Defense, LLC) [K170883] This predicate has not been subject to a design-related recall.

# **IV.** Device Description

The BioFire® Global Fever Special Pathogens Panel is a multiplexed nucleic acid-based test designed to be used with BioFire® FilmArray® Systems (BioFire® FilmArray® 2.0 or BioFire® FilmArray® Torch). The BioFire Global Fever Special Pathogens Panel pouch contains freezedried reagents to perform nucleic acid purification and nested, multiplexed polymerase chain reaction (PCR) with DNA melt analysis. The BioFire Global Fever Special Pathogens Panel conducts tests for the identification of bacterial, viral, and protozoan pathogens from whole blood specimens collected in EDTA tubes (Table 1). Results from the BioFire Global Fever Special Pathogens Panel are available in about 1 hour.

Table 1. Pathogens Detected by the BioFire Global Fever Special Pathogens Panel

| Туре      | Disease                         | Pathogen  |  |  |
|-----------|---------------------------------|---|--|--|
|           | Anthrax                         | Bacillus anthracis <sup>1</sup>   |  |  |
| Bacterial | Leptospirosis                   | Leptospira spp.   |  |  |
| Dacteriai | Plague                          | Yersinia pestis <sup>1</sup>  |  |  |
| Tularemia |                                 | Francisella tularensis <sup>1</sup>   |  |  |
|           | Crimean-Congo hemorrhagic fever | Crimean-Congo hemorrhagic fever virus <sup>1</sup>                          |  |  |
|           | Chikungunya fever               | Chikungunya virus   |  |  |
|           | Dengue fever                    | Dengue virus (serotypes 1, 2, 3 and 4)                                      |  |  |
|           | Lassa fever                     | Lassa virus <sup>1</sup>  |  |  |
| Viral     | Ebola virus disease             | Ebolavirus spp. (Bundibugyo, Reston, Sudan, Taï Forest, Zaire) <sup>1</sup> |  |  |
|           | Marburg virus disease           | Marburgvirus <sup>1</sup>   |  |  |
|           | West Nile fever                 | West Nile virus   |  |  |
|           | Yellow fever                    | Yellow fever virus <sup>2</sup>   |  |  |
|           |                                 | Plasmodium spp.   |  |  |
|           | Malaria                         | Plasmodium falciparum   |  |  |
| Protozoan |                                 | Plasmodium vivax/ovale  |  |  |
|           | Visceral Leishmaniasis          | Leishmania spp. including L. donovani and L. infantum                       |  |  |

<sup>&</sup>lt;sup>1</sup> Select agents are subject to additional requirements. Definitive identification requires confirmatory testing.

A test is initiated by loading Hydration Solution into one port of the pouch and a whole blood specimen mixed with the provided Sample Buffer into the other port of the pouch. The pouch contains all the reagents required for specimen testing and analysis in a freeze-dried format; the addition of Hydration Solution and the Sample Buffer rehydrates the reagents. After the pouch is prepared, the BioFire® FilmArray® Software guides the user through the steps of placing the pouch into the instrument, scanning the pouch barcode, entering the sample identification, selecting the appropriate protocol, and initiating the run on the BioFire FilmArray system.

The BioFire FilmArray instruments contain a coordinated system of inflatable bladders and seal points, which act on the pouch to control the movement of liquid between the pouch blisters. When a bladder is inflated over a reagent blister, it forces liquid from the blister into connecting

<sup>&</sup>lt;sup>2</sup> Definitive identification requires confirmatory testing.

channels. Alternatively, when a seal is placed over a connecting channel it acts as a valve to open or close a channel. In addition, electronically controlled pneumatic pistons are positioned over multiple plungers to deliver the rehydrated reagents into the blisters at the appropriate times. Two Peltier devices control heating and cooling of the pouch to drive the PCR reactions and subsequent melt.

Nucleic acid extraction occurs within the BioFire FilmArray pouch using mechanical and chemical lysis followed by purification using standard magnetic bead technology. After extracting and purifying nucleic acids from the unprocessed sample, a nested multiplexed PCR is executed in two stages. During the first stage, a single, large volume, multiplexed reverse transcription PCR (rt-PCR) reaction is performed. The products from first stage PCR are then diluted and combined with a fresh, primer-free master mix and a fluorescent double stranded DNA binding dye (LC Green® Plus, BioFire Defense, LLC). The solution is then distributed to each well of the array. Array wells contain sets of primers designed specifically to amplify sequences internal to the PCR products generated during the first stage PCR reaction. The second stage PCR and melt, or nested singleplex PCR, is performed in each well of the array. At the end of the second stage PCR, the array is interrogated by melt curve analysis for the detection of signature amplicons denoting the presence of specific targets. A digital camera placed in front of the array captures fluorescent images of the PCR2 reactions and software interprets the data.

#### Materials provided in each BioFire Global Fever Special Pathogens Panel Kit:

- Individually packaged BioFire Global Fever Special Pathogens Panel Pouches
- Single-use (1.0 mL) BioFire® FilmArray® Sample Buffer Tubes
- Single-use pre-filled (1.5 mL) BioFire<sup>®</sup> FilmArray<sup>®</sup> Hydration Injection Vials
- Individually packaged BioFire® FilmArray® Sample Injection Vials
- Individually packaged Transfer Pipettes
- BioFire Global Fever Special Pathogens Panel Quick Guide
- Instructions available online at: www.biofiredefense.com
  - o BioFire Global Fever Special Pathogens Panel Instructions for Use

#### Materials required but not provided:

- BioFire FilmArray System including:
  - o BioFire® FilmArray® 2.0 or BioFire® FilmArray® Torch Instrument System including accompanying platform-specific core software
  - o BioFire® FilmArray® Pouch Loading Station
  - o BioFire<sup>®</sup> Global Fever Special Pathogens Pouch Module Software is required to run the BioFire Global Fever Special Pathogens Panel and is available by request at www.biofiredefense.com if not already installed on the instrument system.
- 10% bleach solution or a similar disinfectant

#### V. Intended Use

The BioFire® Global Fever Special Pathogens Panel is a qualitative, multiplexed, nucleic acid-based test intended for use with BioFire® FilmArray® 2.0 and BioFire® FilmArray® Torch Systems. The BioFire Global Fever Special Pathogens Panel is for the simultaneous qualitative detection and identification of multiple bacterial, viral, and protozoan nucleic acids directly from EDTA whole blood collected from individuals with signs and/or symptoms of acute febrile illness or recent acute febrile illness and known or suspected exposure to the target pathogens described below.

#### Pathogens identified:

Chikungunya virus

Dengue virus (serotypes 1, 2, 3 and 4)

*Leishmania* spp. that cause visceral leishmaniasis (e.g., *L. donovani* and *L. infantum*) *Leptospira* spp.

Plasmodium spp. (including species differentiation of Plasmodium falciparum and Plasmodium vivax/ovale)

West Nile virus

#### Pathogens presumptively identified:

Bacillus anthracis
Crimean-Congo hemorrhagic fever virus
Ebolavirus spp.
Francisella tularensis
Lassa virus
Marburgvirus
Yellow fever virus
Yersinia pestis

Pathogens for which the panel provides presumptive identification results require additional testing and confirmation procedures in consultation with the appropriate public health authorities for whom reports may be necessary.

Positive results do not rule out co-infections with pathogens not included on the BioFire Global Fever Special Pathogens Panel. Not all pathogens that cause acute febrile illness are detected by this test, and negative results do not preclude infection with the pathogens targeted by the device and should not be used as the sole basis for diagnosis, treatment, or other patient management decisions.

Evaluation for more common causes of acute febrile illness (e.g., infections of the upper and lower respiratory tract or gastroenteritis, as well as non-infectious causes) should be considered prior to evaluation with this panel. In the United States, patient travel history, exposure risk, and consultation of the CDC Yellow Book should be considered prior to use of the BioFire Global Fever Special Pathogens Panel as some pathogens are more common in certain geographical locations. Results are meant to be used in conjunction with other clinical, epidemiologic, and laboratory data, in accordance with the guidelines provided by the relevant public health authorities.

The BioFire Global Fever Special Pathogens Panel is indicated for use in laboratories having appropriate biosafety equipment, personal protective equipment (PPE), containment facilities and personnel trained in the safe handling of diagnostic clinical specimens potentially containing any of the pathogens detected by this panel.

The BioFire Global Fever Special Pathogens Panel is indicated for use in laboratories that follow public health guidelines that address appropriate biosafety conditions, interpretation of test results, and coordination of findings with public health authorities.

For In Vitro Diagnostic Use.

# VI. Substantial Equivalence

The BioFire Global Fever Special Pathogens Panel is substantially equivalent to the BioFire Global Fever Panel (previously known as the FilmArray Global Fever Panel) and the FilmArray NGDS Warrior Panel. The BioFire Global Fever Panel was granted De Novo classification on November 20, 2020 [DEN200043] and was categorized as a Class II device. The FilmArray NGDS Warrior Panel was cleared on June 22, 2017 and was also categorized as a Class II device [K170883].

In addition, the BioFire Global Fever Special Pathogens Panel is substantially equivalent to the FilmArray NGDS Warrior Panel in that they both have the ability to detect select agents in whole blood specimens. While there are differences between these panels regarding pouch chemistry, pouch protocols, and pouch module software, both panels were developed using the same FilmArray technology and design principles.

Table 2 outlines the similarities and differences between the BioFire Global Fever Special Pathogens Panel and the predicate devices.

Table 2. Comparison of the BioFire Global Fever Special Pathogens Panel and Predicate Devices

| Element                 | Subject Device:<br>BioFire Global Fever Special<br>Pathogens Panel   | Predicate:<br>BioFire Global Fever Panel<br>[DEN200043]    | Predicate:<br>FilmArray NGDS Warrior Panel<br>[K170883]   |
|-------------------------|--|--|---|
| Specimen Type           | Whole Blood (collected in EDTA tube)   | Same as BioFire Global<br>Fever Special Pathogens<br>Panel | Whole Blood (collected in EDTA tube), positive blood culture, or sputum   |
| Intended Use<br>Setting | Individuals with signs and/or symptoms of acute febrile illness or recent acute febrile illness and known or suspected exposure to pathogens on the panel. | Same as BioFire Global<br>Fever Special Pathogens<br>Panel | Individuals with signs and symptoms of infection from biothreat agents and/or individuals who are at risk for exposure or may have been exposed to pathogens tested by the panel. |

| Element                      | Subject Device:<br>BioFire Global Fever Special<br>Pathogens Panel   | Predicate:<br>BioFire Global Fever Panel<br>[DEN200043]  | Predicate:<br>FilmArray NGDS Warrior Panel<br>[K170883]   |
|------------------------------|--|--|---|
| Pathogens Detected           | Identification: Chikungunya virus, Dengue virus (serotypes 1, 2, 3 and 4), Leishmania spp., Leptospira spp., Plasmodium spp. (including species differentiation of Plasmodium falciparum and Plasmodium vivax/ovale), West Nile virus  Presumptive Identification: Bacillus anthracis, Crimean- Congo hemorrhagic fever virus, Ebolavirus spp., Francisella tularensis, Lassa virus, Marburgvirus, Yellow fever virus, Yersinia pestis | Identification: Chikungunya virus, Dengue virus (serotypes 1,2,3, and 4), Leptospira spp., Plasmodium spp. (including species differentiation of Plasmodium falciparum and Plasmodium vivax/ovale) | Presumptive Identification in whole blood EDTA: Bacillus anthracis, Coxiella burnetii, Ebolavirus spp., Francisella tularensis, Marburgvirus, Yersinia Pestis  Presumptive Identification in positive blood culture: Bacillus anthracis, Yersinia pestis  Presumptive Identification in sputum: Francisella tularensis, Yersinia pestis |
| Analyte                      | RNA/DNA Same as BioFire Global Fever Special Pathogens Panel   |  | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| Technological<br>Principles  | Highly multiplexed nested nucleic acid amplification test with melt analysis   | Same as BioFire Global<br>Fever Special Pathogens<br>Panel   | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| Instrumentation <sup>1</sup> | BioFire FilmArray 2.0 and<br>BioFire FilmArray Torch   | BioFire FilmArray 2.0  | BioFire FilmArray 2.0   |
| Time to result               | About 1 hour   | Same as BioFire Global<br>Fever Special Pathogens<br>Panel   | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| Reagent Storage              | Room temperature   | Same as BioFire Global<br>Fever Special Pathogens<br>Panel   | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| Test Interpretation          | Automated test interpretation and report generation. User cannot access raw data.  | Same as BioFire Global<br>Fever Special Pathogens<br>Panel   | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| Controls                     | Two controls are included in each reagent pouch to control for sample processing and both stages of PCR and melt analysis.   | Same as BioFire Global<br>Fever Special Pathogens<br>Panel   | Same as BioFire Global Fever<br>Special Pathogens Panel   |
| User Complexity              | High   | Moderate   | Moderate  |

<sup>&</sup>lt;sup>1</sup> The BioFire FilmArray 2.0 [K143178] and BioFire FilmArray Torch [K160068] instruments are based on the same technological principles.

# VII. Summary of Performance Data

# **Clinical Performance**

#### Prospective Clinical Study

The prospective clinical study was designed to evaluate the sensitivity/positive percent agreement (PPA) and specificity/negative percent agreement (NPA) for each analyte of the BioFire Global Fever Special Pathogens Panel when testing prospectively collected whole blood specimens in the intended use setting.

Between March 2018 and March 2021, 11 sites contributed 2139 prospectively collected whole blood specimens from individuals who had a recorded or self-reported fever within the past two days. A summary of demographic information for the 2139 specimens included in the prospective study is given in Table 3.

Table 3. Demographic Summary for Prospective BioFire Global Fever Special Pathogens Panel Clinical Evaluation

| Demographic |          | Overall      |
|-------------|----------|--------------|
| ×           | Female   | 1095 (51.2%) |
| Sex         | Male     | 1044 (48.8%) |
|             | <5       | 178 (8.3%)   |
| Age         | 5 to 21  | 822 (38.4%)  |
| ₹           | 22 to 50 | 779 (36.4%)  |
|             | >50      | 360 (16.8%)  |
| Total       |          | 2139         |

PPA for each analyte was calculated as  $100\% \times (TP / (TP + FN))$ . True positive (TP) indicates that both the BioFire Global Fever Special Pathogens Panel and the comparator method had a positive result for this specific analyte, and false negative (FN) indicates that the BioFire Global Fever Special Pathogens Panel result was negative while the comparator result was positive. NPA was calculated as  $100\% \times (TN / (TN + FP))$ . True negative (TN) indicates that both the BioFire Global Fever Special Pathogens Panel and the comparator method had negative results and a false positive (FP) indicates that the BioFire Global Fever Special Pathogens Panel result was positive, but the comparator result was negative. Results are summarized in Tables 4 (viruses), 5 (bacteria), and 6 (protozoa).

Table 4. BioFire Global Fever Special Pathogens Panel Clinical Performance Summary - Viruses

| BioFire Global Fever                        | Number            |                 | PPA   |                | NPA             |       |           |  |
|---|-------------------|-----------------|-------|----------------|-----------------|-------|-----------|--|
| Special Pathogens Panel Detected Result     | Tested            | TP/(TP +<br>FN) | %     | 95% CI         | TN/(TN +<br>FP) | %     | 95% CI    |  |
| Chikungunya virus <sup>a</sup>              | 1875 <sup>c</sup> | 25/25           | 100%  | 86.7-<br>100%  | 1848/1850       | 99.9% | 99.6-100% |  |
| Crimean-Congo<br>hemorrhagic<br>fever virus | 2139              | 1/1             | 100%  | 20.7-<br>100%  | 2138/2138       | 100%  | 99.8-100% |  |
| Dengue virus <sup>b</sup>                   | 1875°             | 266/283         | 94.0% | 90.6-<br>96.2% | 1592/1592       | 100%  | 99.8-100% |  |
| Ebola virus                                 | 2139              | 0/0             | -     | -              | 2139/2139       | 100%  | 99.8-100% |  |
| Lassa virus                                 | 2139              | 0/0             | -     | -              | 2139/2139       | 100%  | 99.8-100% |  |
| Marburg virus                               | 2139              | 0/0             | -     | -              | 2139/2139       | 100%  | 99.8-100% |  |
| West Nile virus                             | 2139              | 1/1             | 100%  | 20.7-<br>100%  | 2138/2138       | 100%  | 99.8-100% |  |
| Yellow fever virus                          | 2139              | 0/0             | -     | -              | 2139/2139       | 100%  | 99.8-100% |  |

<sup>&</sup>lt;sup>a</sup> Evidence of Chikungunya virus was found in 2/2 FP specimens by additional PCR.

Table 5. BioFire Global Fever Special Pathogens Panel Clinical Performance Summary – Bacteria

| <b>BioFire Global Fever Special</b> | Number PPA        |              |       |                | NPA             |       |            |  |
|-------------------------------------|-------------------|--------------|-------|----------------|-----------------|-------|------------|--|
| Pathogens Panel Detected<br>Result  | Tested            | TP/(TP + FN) | %     | 95% CI         | TN/(TN +<br>FP) | %     | 95% CI     |  |
| Bacillus anthracis                  | 2139              | 0/0          | -     | -              | 2139/2139       | 100%  | 99.8-100%  |  |
| Francisella tularensis              | 2139              | 0/0          | -     | -              | 2139/2139       | 100%  | 99.8-100%  |  |
| Leptospira spp.ª                    | 1875 <sup>b</sup> | 15/16        | 93.8% | 71.7-<br>98.9% | 1855/1859       | 99.8% | 99.4-99.9% |  |
| Yersinia pestis                     | 2139              | 0/0          | 1     | -              | 2139/2139       | 100%  | 99.8-100%  |  |

<sup>&</sup>lt;sup>a</sup> Evidence of *Leptospira* spp. was found in 1/1 FN specimens by BioFire Global Fever Special Pathogens Panel retest and by additional PCR, and in 3/4 FP specimens by additional PCR.

<sup>&</sup>lt;sup>b</sup> Evidence of Dengue virus was found in 15/17 FN specimens: five specimens were positive upon BioFire Global Fever Special Pathogens Panel retest and by additional PCR, two were positive only upon BioFire Global Fever Special Pathogens Panel retest, and eight were detected only by additional PCR.

<sup>&</sup>lt;sup>c</sup> Comparator analysis was not performed for Chikungunya virus or Dengue virus on specimens collected after September 2019.

<sup>&</sup>lt;sup>b</sup> Comparator analysis was not performed for *Leptospira* on specimens collected after September 2019.

Table 6. BioFire Global Fever Special Pathogens Panel Clinical Performance Summary – Protozoa

| <b>BioFire Global Fever Special</b> | Fever Special Number |              | PPA   |            |                 | NPA   |            |  |
|-------------------------------------|----------------------|--------------|-------|------------|-----------------|-------|------------|--|
| Pathogens Panel Detected<br>Result  | Tested               | TP/(TP + FN) | %     | 95% CI     | TN/(TN +<br>FP) | %     | 95% CI     |  |
| Leishmania spp.                     | 2139                 | 10/10        | 100%  | 72.2-100%  | 2129/2129       | 100%  | 99.8-100%  |  |
| Plasmodium spp.a,b                  | 1875 <sup>e</sup>    | 338/343      | 98.5% | 96.6-99.4% | 1519/1532       | 99.2% | 98.6-99.5% |  |
| Plasmodium falciparum <sup>c</sup>  | 1875 <sup>e</sup>    | 230/248      | 92.7% | 88.8-95.4% | 1624/1627       | 99.8% | 99.5-99.9% |  |
| Plasmodium vivax/ovale <sup>d</sup> | 1875 <sup>e</sup>    | 115/124      | 92.7% | 86.8-96.1% | 1751/1751       | 100%  | 99.8-100%  |  |

<sup>&</sup>lt;sup>a</sup> Four (4/5) *Plasmodium* FN specimens were also *P. falciparum* FN and one (1/5) was *P. vivax/ovale* FN. Three (3/13) *Plasmodium* FP specimens were also *P. falciparum* FP.

#### **Archived Specimen Study**

Many of the analytes detectable by the BioFire Global Fever Special Pathogens Panel were not observed or were not encountered in large enough numbers in the prospective clinical evaluation to adequately demonstrate system performance. In this study, archived specimens with known analyte content and/or archived specimens with a high likelihood of containing a given analyte were tested with the BioFire Global Fever Special Pathogens Panel to supplement the prospective clinical evaluation data. Wherever possible, archived whole blood specimens were tested. Where no whole blood specimens could be obtained, blood plasma and blood serum were tested instead. Although plasma and serum are not the intended specimen type for the BioFire Global Fever Special Pathogens Panel, these blood components provide similar results to whole blood specimens in a matrix equivalency study. Archived specimens were collected from a range of ages and sexes (Table 7).

<sup>&</sup>lt;sup>b</sup> Evidence of *Plasmodium* spp. was found in 2/5 FN specimens: one specimen was positive upon BioFire Global Fever Special Pathogens Panel retest and by additional PCR, and one was positive only upon BioFire Global Fever Special Pathogens Panel retest. Evidence of *Plasmodium* spp. was found in 11/13 FP specimens by additional PCR (10/13) or by species-level comparator assay (1/13).

<sup>&</sup>lt;sup>c</sup> Evidence of *P. falciparum* was found in 13/18 FN specimens: three specimens were positive upon BioFire Global Fever Special Pathogens Panel retest and by additional PCR, one was positive only upon BioFire Global Fever Special Pathogens Panel retest, and nine were positive only by additional PCR. Evidence of *P. falciparum* was found in 2/3 FP specimens by additional PCR.

<sup>&</sup>lt;sup>d</sup> Evidence of *P. vivax/ovale* was found in 7/9 FN specimens: two specimens were positive upon BioFire Global Fever Special Pathogens Panel retest and by additional PCR, two were positive only upon BioFire Global Fever Special Pathogens Panel retest, and three were positive only by additional PCR.

<sup>&</sup>lt;sup>e</sup> Comparator analysis was not performed for *Plasmodium, P. falciparum,* or *P. vivax/ovale* on specimens collected after September 2019.

Table 7. Overall and Per Site Archived Demographic Analysis

|     | Overall  |                         | Site 01 <sup>a</sup> | Site 02    | Site 03 <sup>a</sup> |
|-----|----------|-------------------------|----------------------|------------|----------------------|
|     | Female   | 160 (38.5%)             | 79 (39.7%)           | 49 (59.8%) | 32 (23.7%)           |
| Sex | Male     | 148 (35.6%)             | 97 (48.7%)           | 33 (40.2%) | 18 (13.3%)           |
|     | Unknown  | 108 (26.0%)             | 23 (11.6%)           | 0 (0%)     | 85 (63%)             |
|     | <5       | 14 (3.4%)               | 14 (7.0%)            | 0 (0%)     | 0 (0%)               |
|     | 5 to 21  | 37 (8.9%)               | 19 (9.5%)            | 14 (17.1%) | 4 (3%)               |
| Age | 22 to 50 | 200 (48.1%)             | 121 (60.8%)          | 56 (68.3%) | 23 (17%)             |
|     | >50      | 0 57 (13.7%) 22 (11.1%) |                      | 12 (14.6%) | 23 (17%)             |
|     | Unknown  | 108 (26.0%)             | 23 (11.6%)           | 0 (0%)     | 85 (63%)             |
|     | Total    | 416                     | 199                  | 82         | 135                  |

<sup>&</sup>lt;sup>a</sup> Demographic data was not available for 23 specimens from Site 01 and 85 specimens from Site 03.

Specimens were coded and either tested at the clinical site or shipped to BioFire Defense for testing on the BioFire Global Fever Special Pathogens Panel. Nucleic acid was extracted from each specimen and tested using plate-based PCR comparator methods. Samples for which false positive and/or false negative results (i.e., discrepant results) were obtained when comparing the BioFire Global Fever Special Pathogens Panel results to the comparator method results were further investigated. Results are summarized in Table 8.

Table 8. BioFire Global Fever Special Pathogens Panel Archived Performance Summary

| Pathogen <sup>a</sup>                 |              | PPA    |            | NPA          |         |            |  |  |  |
|---------------------------------------|--------------|--------|------------|--------------|---------|------------|--|--|--|
| Pathogen-                             | TP/(TP + FN) | %      | 95% CI     | TN/(TN + FP) | %       | 95% CI     |  |  |  |
| Viruses                               |              |        |            |              |         |            |  |  |  |
| Crimean-Congo hemorrhagic fever virus | 0/0          | -      | -          | 281/281      | 100%    | 98.7-100%  |  |  |  |
| Ebolavirus spp. b                     | 0/0          | -      | -          | 279/279      | 100%    | 98.6-100%  |  |  |  |
| Lassa virus <sup>b,c</sup>            | 10/12        | 83.3%  | 55.2-95.3  | 265/267      | 99.2%   | 97.3-99.8% |  |  |  |
| <i>Marburgvirus</i> sp. <sup>b</sup>  | 0/0          | -      | -          | 279/279      | 100%    | 98.6-100%  |  |  |  |
| West Nile virus <sup>b,d,e,f</sup>    | 59/65        | 90.8%  | 81.3-95.7% | 345/347      | 99.4%   | 97.9-99.8% |  |  |  |
| Yellow fever virus <sup>b</sup>       | 0/0          | -      | -          | 279/279      | 100%    | 98.6-100%  |  |  |  |
|                                       |              | Bacte  | ria        |              |         |            |  |  |  |
| Bacillus anthracis                    | 0/0          | -      | -          | 281/281      | 100%    | 98.7-100%  |  |  |  |
| Francisella tularensis                | 0/0          | -      | -          | 281/281      | 100%    | 98.7-100%  |  |  |  |
| Yersinia pestis                       | 0/0          | =      | -          | 281/281      | 100%    | 98.7-100%  |  |  |  |
|                                       |              | Protoz | oa         |              |         |            |  |  |  |
| Leishmania spp. <sup>g</sup>          | 0/0          | -      | -          | 283/283      | 283/283 | 98.7-100%  |  |  |  |

<sup>&</sup>lt;sup>a</sup> Results are not shown for assays previously granted in DEN200043 (i.e., chikungunya virus, dengue virus, *Leptospira* spp., and *Plasmodium* spp.)

<sup>&</sup>lt;sup>b</sup> Due to low specimen volume, comparator results for most pathogens were not obtained for two specimens. These specimens were only tested for a subset of pathogens including Crimean-Congo hemorrhagic fever virus, *Bacillus anthracis*, *Francisella tularensis*, *Yersinia pestis*, and *Leishmania* spp.

<sup>&</sup>lt;sup>c</sup> Evidence of Lassa virus was found in ½ FN specimens and ½ FP specimens by additional PCR.

<sup>&</sup>lt;sup>d</sup> A set of 133 specimens tested at Site 03 had been previously characterized and were expected to be negative for all panel targets or positive for West Nile virus. These specimens were only evaluated by comparator methods for West Nile virus.

<sup>&</sup>lt;sup>e</sup> Archived specimens included blood serum and blood plasma.

f Evidence of West Nile virus was found in 6/6 FN specimens: four specimens were positive upon Global Fever Special Pathogens Panel retesting and three of these were also positive by additional PCR. The other two FN specimens were positive only by additional PCR testing. Evidence of West Nile virus was detected in 1/2 FP specimens by additional PCR testing.

g Two specimens tested at Site 03 had been previously characterized and were only tested on comparator assays for *Leishmania* spp.

#### Contrived Specimen Study

For analytes for which no archived specimens were available, or for which there were an insufficient number of archived specimens, testing was performed using contrived specimens. This study evaluated BioFire Global Fever Special Pathogens Panel sensitivity and specificity when testing whole-blood specimens contrived with *Bacillus anthracis*, Crimean-Congo hemorrhagic fever virus, *Ebolavirus* spp., *Francisella tularensis*, Lassa fever virus, *Leishmania* spp., *Marburgvirus* sp., West Nile virus, Yellow fever virus, and *Yersinia pestis*.

Contrived specimens were prepared using residual human whole blood specimens from patients with signs/symptoms of acute febrile illness. For each analyte, fifty (50) replicates were contrived using quantified isolates at a range of concentrations relative to the limit of detection (LoD). If known, clinically relevant concentrations were used to adjust testing levels. The contrived specimens also served as a negative replicate for all other analytes evaluated. Specimens were prepared and randomized such that the analyte status of each contrived specimen was blinded to the users performing testing. The positive percent agreement (PPA) and negative percent agreement (NPA) were defined as agreement between the BioFire Global Fever Special Pathogens Panel result and the known composition of the contrived specimen. A summary of the results is shown in Table 9.

Table 9. Summary of BioFire Global Fever Special Pathogens Panel Contrived Specimen Performance Data

| Analista                |                     | РРА |           |                      | NPA  |          |  |  |
|-------------------------|---------------------|-----|-----------|----------------------|------|----------|--|--|
| Analyte                 | TP/(TP + FN)        | %   | 95% CI    | TN/(TN + FP)         | %    | 95% CI   |  |  |
| Bacillus anthracis      | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |
| CCHF virus <sup>a</sup> | 98/100 <sup>b</sup> | 98  | 93.0-99.5 | 282/282              | 100  | 98.7-100 |  |  |
| Ebolavirus spp.         | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |
| Francisella tularensis  | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |
| Lassa virus             | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |
| Leishmania spp.         | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |
| Marburgvirus sp.a       | 99/100°             | 99  | 94.6-99.8 | 282/282              | 100  | 98.7-100 |  |  |
| West Nile virus         | 50/50               | 100 | 92.9-100  | 331/332 <sup>d</sup> | 99.7 | 98.3-100 |  |  |
| Yellow fever virus      | 49/50 <sup>e</sup>  | 98  | 89.5-99.7 | 332/332              | 100  | 98.9-100 |  |  |
| Yersinia pestis         | 50/50               | 100 | 92.9-100  | 332/332              | 100  | 98.9-100 |  |  |

<sup>&</sup>lt;sup>a</sup> Tested at additional concentrations to better represent clinically relevant range.

<sup>&</sup>lt;sup>b</sup> Two false negative CCHF virus at 2× LoD.

<sup>&</sup>lt;sup>c</sup> Single false negative *Marburgvirus* sp. at 10× LoD.

<sup>&</sup>lt;sup>d</sup> Discrepancy testing showed near LoD levels of WNV in a single whole blood sample.

<sup>&</sup>lt;sup>e</sup> Single false negative yellow fever virus at 5× LoD.

### **Select Analytical Studies**

#### Limit of Detection

The Limit of Detection (LoD) for each analyte on the BioFire Global Fever Special Pathogens Panel was determined using quantified stocks within the BioFire Defense BioSafety Level 2 (BSL2) laboratory. For analytes that required BSL3/4 containment, inactivated strains were used. Contrived samples were prepared at known concentrations in a whole blood background. An estimated LoD concentration for each analyte was determined based on results of serial dilutions spanning at least four concentrations bracketing the anticipated LoD. Four replicates were tested at each dilution, with additional dilutions if needed, to reach a concentration at which loss of detection could be observed. The LoD concentration was confirmed by testing 20 replicates at the estimated LoD concentration and another 20 replicates at a ten-fold lower concentration. The required criteria for confirmation of LoD was a detection rate of at least 95% at LoD ((≥19/20) and a detection rate of less than 95% below LoD (<19/20).

The confirmed LoD concentrations for BioFire Global Fever Special Pathogens Panel analytes along with the detection rate at 1× LoD are shown in Table 10. The LoD concentration for each analyte is reported as target copies/mL determined using commercially available quantitative real-time PCR assay kits.

Table 10. Limit of Detection for BioFire Global Fever Special Pathogens Panel Test Results

| Global Fever Special                        |   |                  | LoD C                  | LoD Concentration              |  |  |  |
|---|---|------------------|------------------------|--------------------------------|--|--|--|
| Pathogens Panel<br>Analyte                  | Isolate Tested  | Live/Inactivated | Copies/mL <sup>1</sup> | Units/mL                       |  |  |  |
|   | BACTERIA  |                  |                        |                                |  |  |  |
| Bacillus anthracis                          | Ames35  | Live             | 4.2E+01                | N/A                            |  |  |  |
| Francisella tularensis                      | SCHU S4   | Inactivated      | 1.2E+03                | N/A                            |  |  |  |
| <i>Leptospira</i> spp.                      | interrogans: serovar icterohaemorrhagiae,<br>Serotype: Budapest | Live             | 3.4E+02                | N/A                            |  |  |  |
| Yersinia pestis                             | A1122   | Live             | 1.3E+02                | N/A                            |  |  |  |
|   | VIRUSES   |                  |                        |                                |  |  |  |
| Chikungunya virus                           | R80422  | Inactivated      | 5.5E+02                | 3.6E-01 units/mL               |  |  |  |
| Crimean-Congo<br>hemorrhagic fever<br>virus | Strain IbAr10200  | Inactivated      | 6.4E+00                | N/A                            |  |  |  |
|   | DENV-1: Hawaii  | Live             | 2.2E+02                | N/A                            |  |  |  |
|   | DENV-2-1: New Guinea C  | Live             | 3.4E+02                | N/A                            |  |  |  |
| Dengue virus                                | DENV-2-2: Dak AR A1247  | Live             | 2.7E+03                | 1.5E+02 TCID <sub>50</sub> /mL |  |  |  |
|   | DENV-3: H87   | Live             | 1.3E+02                | 3.7E+00 units/mL               |  |  |  |
|   | DENV-4: H241  | Live             | 6.4E+01                | 1.8E+02 units/mL               |  |  |  |
|   | Bundibugyo: 200706291 Uganda                                    | Inactivated      | 7.0E+04                | N/A                            |  |  |  |
|   | Taï Forest: Cote d'Ivoire 11/27/94                              | Inactivated      | 8.3E+03                | N/A                            |  |  |  |
| Ebolavirus spp.                             | Reston: 119810 RIID<br>(MKY 53) (prototype 1989)                | Inactivated      | 2.7E+04                | N/A                            |  |  |  |
|   | Sudan: Boniface   | Inactivated      | 1.1E+04                | N/A                            |  |  |  |
|   | Zaire:<br>Guéckédou/Guinea C07                                  | Inactivated      | 1.0E+02                | 1.5E+02<br>PFU/mL              |  |  |  |
| Lassa virus                                 | Josiah  | Inactivated      | 5.6E+04                | N/A                            |  |  |  |
| A 4 l                                       | Marburg virus: Musoke   | Inactivated      | 5.0E+02                | N/A                            |  |  |  |
| Marburgvirus                                | Ravn virus: Kenya Ravn  | Inactivated      | 2.6E+02                | N/A                            |  |  |  |
| 147 1 AI'I '                                | NY 2001-6263 (Lineage 1)  | Inactivated      | 1.1E+03                | 2.7E+01 units/mL               |  |  |  |
| West Nile virus                             | B-956 Uganda (Lineage 2)  | Inactivated      | 2.3E+04                | 6.2E+00 units/mL               |  |  |  |
| Yellow fever virus                          | Strain 17D  | Live attenuated  | 1.2E+02                | 1.2E+01 TCID <sub>50</sub> /mL |  |  |  |
|   | PROTOZOA  |                  |                        |                                |  |  |  |
| Leishmania spp.                             | donovani: 9515 (MHOM/IN/95/9515)                                | Live             | 1.0E+01                | 2.2E+01 cells/mL               |  |  |  |
|   | falciparum, IPC 4884 Pursat<br>Cambodia 2011                    | Live             | 1.8E+02                | N/A                            |  |  |  |
| _, ,,                                       | knowlesi, H strain  | gDNA             | 2.4E+01                | 20 pg/mL                       |  |  |  |
| Plasmodium spp.                             | malariae (Clinical Specimen)                                    | Live             | 1.9E+02                | 2.3E-01 cells/mL               |  |  |  |
|   | ovale, wallikeri (Clinical Specimen)                            | Live             | 2.4E+02                | N/A                            |  |  |  |
|   | vivax, Strain Chesson   | Live             | 1.5E+02                | N/A                            |  |  |  |
| Plasmodium falciparum                       | IPC 4884 Pursat<br>Cambodia 2011                                | Live             | 1.8E+02                | N/A                            |  |  |  |
| Plasmodium                                  | ovale, wallikeri (Clinical Specimen)                            | Live             | 2.4E+02                | N/A                            |  |  |  |
| vivax/ovale                                 | vivax, Strain Chesson   | Live             | 1.5E+02                | N/A                            |  |  |  |

<sup>&</sup>lt;sup>1</sup> Stock concentrations determined using commercially available quantitative real-time PCR assay kits.

Since decreased sensitivity may be observed due to nucleic acid damage when evaluating inactivated BSL3/4 pathogens, additional testing was performed using live strains within a subcontracted BSL3/4 laboratory.

The concentrations tested were based on the confirmed LoD for inactivated/attenuated material. In general, four test concentrations were tested for each analyte:  $10\times$ ,  $1\times$ ,  $0.1\times$ , and  $0.01\times$  the LoD for inactivated/attenuated stock. Some analytes required additional concentrations. For each concentration, four unique blood draws were individually spiked and tested on the BioFire Global Fever Special Pathogens Panel. The estimated LoD was identified as the lowest concentration at which 4/4 replicates were positive and fewer than four replicates were positive at the next lower concentration. The estimated LoD values for infectious material are provided in Table 11.

Table 11. BioFire Global Fever Special Pathogens Panel Estimated LoD in Whole Blood for Live BSL3/4

Agents

| BioFire Global Fever               |   | Estimated Lo | D Concentration |
|------------------------------------|---|--------------|-----------------|
| Special Pathogens Panel<br>Analyte | Species/Strain                              | Copies/mL    | Units/mL        |
|                                    | Bacteria                                    |              |                 |
| Bacillus anthracis                 | Ames  | 6.4E+01      | 3.5E+00 cfu/mL  |
| Francisella tularensis             | subsp. <i>tularensis</i> Schu               | 1.2E+01      | 2.1E+02 cfu/mL  |
| Yersinia pestis                    | CO92  | 1.5E+02      | 3.0E+01 cfu/mL  |
|                                    | Viruses                                     |              |                 |
| CCHF Virus                         | lbAr10200                                   | 6.4E+02      | 2.9E+03 pfu/mL  |
| Chilanaguana niana                 | B8635                                       | 5.5E+02      | 4.4E+01 pfu/mL  |
| Chikungunya virus                  | Indo23574                                   | 5.5E+02      | 9.0E+01 pfu/mL  |
|                                    | Bundibugyo virus / Uganda (811250)          | 7.0E+02      | 5.6E+00 pfu/mL  |
|                                    | Tai Forest virus / Taï Forest (Ivory Coast) | 1.8E+02      | N/A             |
| Ebolavirus spp.                    | Reston virus / H-28                         | 2.7E+03      | 4.7E+01 pfu/mL  |
|                                    | Sudan virus / Boniface                      | 1.1E+02      | 5.3E+00 pfu/mL  |
|                                    | Zaire ebolavirus / Makona                   | 1.1E+03      | 1.6E+01 pfu/mL  |
| Lassa Virus                        | Josiah                                      | 5.6E+03      | 2.6E+01 pfu/mL  |
|                                    | Ci67  | 5.0E+02      | 1.5E+03 pfu/mL  |
| Marburg marburgvirus               | Ravn  | 2.6E+02      | 4.7E+01 pfu/mL  |
| West Nile virus                    | Bz NY99 (lin. 1)                            | 1.6E+02      | 9.5E+00 pfu/mL  |
| Yellow fever virus                 | Asibi                                       | 1.2E+01      | 9.2E+00 pfu/mL  |

#### *Inclusivity (Reactivity)*

The analytical reactivity (inclusivity) of the BioFire Global Fever Special Pathogens Panel was evaluated by testing a diverse collection of strains/species/serotypes representing temporal, geographic, and genetic diversity. Available isolates were prepared as contrived whole blood samples with the isolate at a concentration near the LoD of the analyte. If the isolate was detected within  $3\times$  the LoD the BioFire Global Fever Special Pathogens Panel was considered inclusive for that isolate and those with similar sequences. Analytes that were detected at  $\ge 10\times$  the established LoD were considered to have reduced sensitivity.

Table 12 shows a summary of the BioFire Global Fever Special Pathogens Panel reactivity based on empirical data. Isolates used to determine LoD are bolded.

Table 12. BioFire Global Fever Special Pathogens Panel Analytical Reactivity (Inclusivity)

| Analyte         | # Isolates<br>Detected /<br>Tested | Isolates Tested          |                                   | Source / ID     | Concentration Detected Tested up to 100× LoD¹ | Limitations |
|-----------------|------------------------------------|--------------------------|-----------------------------------|-----------------|---|-------------|
|                 |                                    |                          |                                   |                 |   |             |
|                 |                                    | Ames35                   |                                   | BEI / NR-10355  | 4.2E+01 copies/mL                             |             |
| Bacillus        | 4/4                                | Sterne 34Fs              |                                   | BEI / NR-1400   |   | None        |
| anthracis       | 4/4                                | UM23                     |                                   | BEI / NR-10351  | 1.3E+02 copies/mL                             | None        |
|                 |                                    | Weybridge                |                                   | BEI / NR-10350  |   |             |
|                 |                                    | Subspecies               | Strain                            |                 |   |             |
|                 |                                    | tularensis               | SCHU S4                           | BEI / NR-15753  | 1.2E+03 copies/mL                             |             |
| Francisella     | 5/5                                | holarctica               | LVSR                              | BEI / NR-597    |   | None        |
| tularensis      | 3/3                                | Holarctica               | Type B LVS (CDC)                  | BEI / NR-646    | 3.6E+03 copies/mL                             | None        |
|                 |                                    | novicida                 | CG62                              | BEI / NR-580    | 3.0L+03 copies/file                           |             |
|                 |                                    | Hoviciaa                 | KM14S                             | BEI / NR-573    |   |             |
|                 |                                    | Species                  | Strain                            |                 |   |             |
|                 |                                    | interrogans              | Serovar (Budapest)                | ATCC / 23581    | 3.4E+02 copies/mL                             |             |
|                 |                                    |                          | HAI0156<br>(Copenhageni)          | BEI / NR-19891  | 1.2E+03 copies/mL                             |             |
|                 |                                    |                          | L495 (Manilae)                    | BEI / NR-19816  | -   |             |
|                 |                                    | alexanderi               | L60 (Manhao 3)                    | ATCC / 700520   | 1.0E+03 copies/mL                             |             |
|                 |                                    | alstonii                 | Sichuan 79601                     | ATCC / BAA-2439 |   |             |
|                 |                                    | h                        | Castellon 3<br>(Castellonis)      | ATCC / 23580    | 1.2E+03 copies/mL                             |             |
| Leptospira spp. | 19/19                              | borgpetersenii           | Veldrat Bataviae 46<br>(Javanica) | ATCC / 43292    | 1.0E+03 copies/mL                             | None        |
|                 |                                    | kirschneri               | 200701401<br>(Bogvere)            | BEI /NR-19942   | 1.2E+03 copies/mL                             |             |
|                 |                                    |                          | 3522 C (Cynopteri)                | ATCC / 49945    | 5.3E+02 copies/mL                             |             |
|                 | kmetyi                             | Bejo-Iso9T<br>(Malaysia) | BEI / NR-22254                    | , ,             |   |             |
|                 |                                    | mayottensis              | 200901116<br>(undesignated)       | KIT / 0254      | 1.2E+03 copies/mL                             |             |
|                 |                                    | noguchii                 | CZ 214T (Panama)                  | BEI / NR-22283  |   |             |
|                 |                                    | santarosai               | LT 821 (Shermani)                 | ATCC / 43286    | 8.7E+02 copies/mL                             |             |

| Analyte   | # Isolates<br>Detected /<br>Tested | Isola           | Isolates Tested       |                            | Concentration Detected Tested up to 100× LoD¹ | Limitations             |
|---|------------------------------------|-----------------|-----------------------|----------------------------|---|-------------------------|
|   |                                    |                 | 6712                  | KIT / 0220                 |   |                         |
|   |                                    |                 | 94-79970/3 (Topaz)    | KIT / 0237                 |   |                         |
|   |                                    | weilii          | A 102 (Mengrun)       | KIT / 0023                 | 1.2E+03 copies/mL                             |                         |
|   |                                    | weiiii          | Celledoni 20160426    | ATCC / 43285               | 1.2E+03 copies/iiiL                           |                         |
|   |                                    |                 | H 27 (Hekou)          | KIT / 0074                 |   |                         |
|   |                                    |                 | LT 89-68 (Vughia)     | KIT / 0127                 |   |                         |
|   |                                    | Biovar          | Strain                |                            |   |                         |
| Yersinia pestis                                 | 3/3                                | Orientalis      | A1122                 | BEI / NR-636               | 1.3E+02 copies/mL                             | None                    |
| Tersiniu pestis                                 | 3/3                                | Orientalis      | PY-013                | BEI / NR-51666             | 3.9E+02 copies/mL                             | None                    |
|   |                                    | Antiqua         | PH 80/63              | BEI / NR-51667             | 3.9L102 copies/iiiL                           |                         |
|   |                                    |                 | VIRUSES               |                            |   |                         |
|   |                                    | 1               |                       | ZeptoMetrix /              |   |                         |
|   |                                    | R80422          |                       | 0810105CFHI                | 5.5E+02 copies/mL                             |                         |
|   |                                    |                 |                       | BEI / NR-50884             |   |                         |
| Chikungunya                                     |                                    | DHS4263         |                       | (formerly NR-              | 1.7E+03 copies/mL                             | None                    |
| virus   | 3/3                                | 21131203        |                       | 50055)                     | 1.7L+03 copies/iiiL                           |                         |
|   |                                    |                 |                       | BEI / NR-50883             |   |                         |
|   | St. Martin 2013                    |                 | (formerly NR-         | 1.4E+03 copies/mL          |   |                         |
|   |                                    | 5t. Martin 2015 |                       |                            | 49901)  |                         |
| Crimean–<br>Congo<br>hemorrhagic<br>fever virus | 1/1                                |                 | Nigeria / IbAr10200   |                            | 6.4E+00 copies/mL                             | None                    |
|   |                                    | Serotype        | Strain                | ZeptoMetrix /              |   |                         |
|   |                                    |                 | Hawaii                | 0810088CF                  | 2.2E+02 copies/mL                             | _                       |
|   |                                    |                 | Strain 12150          | BEI / NR-3785              |   |                         |
|   |                                    |                 | 228690                | BEI / NR-3786              |   | None                    |
|   |                                    |                 | 276RK1                | BEI / NR-3782              |   |                         |
|   |                                    | Serotype 1      | BC89/94               | BEI / NR-3787              | 6.6E+02 copies/mL                             |                         |
|   |                                    |                 | SL-6-6-04             | BEI / NR-49744             |   |                         |
|   |                                    |                 | UIS 1162              | BEI / NR-49707             |   |                         |
|   |                                    |                 | VN/BID-<br>V1792/2007 | BEI / NR-44083             |   |                         |
|   |                                    |                 | New Guinea C          | ZeptoMetrix /              | 3.4E+02 copies/mL                             |                         |
| Dengue virus                                    | 27/28                              |                 | (DENV 2_1)            | 0810089CF                  | 3.42.02 copies/iiie                           |                         |
| 20640 11143                                     | _,,_0                              |                 | DakArA1247            | BEI / NR-12221             | 2.7E+03 copies/mL                             |                         |
|   |                                    |                 | (DENV 2_2)            |                            | 2.7.2.33 copies/iiii                          |                         |
|   |                                    |                 | 1349                  | BEI / NR-12219             | _   | Not inclusive           |
|   |                                    | Serotype 2      | 429557                | BEI / NR-12216             | 1.1E+03 copies/mL                             | of strain               |
|   |                                    |                 | ArA6894               | BEI / NR-12220             |   | DKA 811                 |
|   |                                    |                 | BC102/94              | BEI / NR-3789              | 9.4E+02 copies/mL                             |                         |
|   |                                    |                 | DKA 811               | BEI / NR-49747             | Not Detected                                  |                         |
|   |                                    |                 | VN/BID-               | BEI / NR-44085             | 1.1E+03 copies/mL                             |                         |
|   |                                    |                 | V1002/2006            | _                          |   |                         |
|   |                                    | Serotype 3      | Н87                   | ZeptoMetrix /<br>0810090CF | 1.3E+02 copies/mL                             | Reduced sensitivity for |
|   |                                    |                 | 271242                | BEI / NR-3802              | 3.9E+02 copies/mL                             | Schistivity 101         |

| Analyte                  | # Isolates<br>Detected /<br>Tested | Isolates Tested          |                    | Source / ID                 | Concentration  Detected  Tested up to  100× LoD¹ | Limitations     |
|--------------------------|------------------------------------|--------------------------|--------------------|-----------------------------|--|-----------------|
|                          |                                    |                          | BC188/97           | BEI / NR-3801               | 1.6E+04 copies/mL                                | strain          |
|                          |                                    |                          | C0360/94           | BEI / NR-48800              |  | BC188/97        |
|                          |                                    |                          | VN/BID-            | BEI / NR-44087              | 3.9E+02 copies/mL                                |                 |
|                          |                                    |                          | V1329/2006         |                             |  |                 |
|                          |                                    |                          | H241               | ZeptoMetrix /<br>0810091CF  | 6.4E+01 copies/mL                                |                 |
|                          |                                    |                          | 703                | BEI / NR-48801              |  | Reduced         |
|                          |                                    | Corotuno 1               | BC13/97            | BEI / NR-3805               | 1 0E+02 copies/ml                                | sensitivity for |
|                          |                                    | Serotype 4               | BC287/97           | BEI / NR-3806               | 1.9E+02 copies/mL                                | strain          |
|                          |                                    |                          | BC258/97           | BEI / NR-3807               | 7  | D85-019         |
|                          |                                    |                          | D85-019            | BEI / NR-3804               | 7.6E+03 copies/mL                                |                 |
|                          |                                    |                          | PR 06-65-740       | BEI / NR-49757              | 2.2E+02 copies/mL                                |                 |
|                          |                                    | Bundibugyo Uga           | anda               | BEI / NR-31813              | 7.0E+04 copies/mL                                | Reduced         |
|                          |                                    | Reston MKY 53            |                    | BEI / NR-44238              | 2.7E+04 copies/mL                                | sensitivity for |
| The lawings              | c Ic                               | Sudan                    |                    | BEI / NR-31810              | 1.1E+04 copies/mL                                | inactivated     |
| Ebolavirus               | 6/6                                | Taï Forest               |                    | BEI / NR-44241              | 8.3E+03 copies/mL                                | Zaire           |
|                          |                                    | Zoine                    | Guéckédou          | BEI / NR-49462              | 1.0E+02 copies/mL                                | ebolavirus      |
|                          |                                    | Zaire                    | Mayinga            | BEI / NR-31807              | 1.1E+04 copies/mL                                | Mayinga         |
| Lassa Virus              | 1/1                                | Josiah                   |                    | BEI / NR-31822              | 5.6E+04 copies/mL                                | None            |
|                          |                                    | Marburg virus Musoke     |                    | BEI / NR-48951              | 5.0E+02 copies/mL                                |                 |
| Marburgvirus             | 3/3                                | Marburg virus Voege      |                    | BEI / NR-31816              | 1.5E+03 copies/mL                                | None            |
| a.za.gas                 | 5,5                                | Ravn virus Kenya         |                    | BEI / NR-31819              | 2.6E+02 copies/mL                                | _               |
|                          |                                    | Ravii vii us keiry       | Ravii viius Reiiya |                             | Z.OZ.OZ COPICS/IIIZ                              |                 |
|                          |                                    | B-956 Uganda (           | lin. 2)            | Zeptometrix/<br>0810081cfhi | 2.3E+04 copies/mL                                |                 |
|                          |                                    | B-956 (lin. 2)           |                    | BEI / NR-50885              | 6.9E+04 copies/mL                                | <u> </u>        |
| West Nile virus          | 5/5                                | NY 2001-6263 (I          | in. 1)             | Zeptometrix/<br>0810033cfhi | 1.1E+03 copies/mL                                | None            |
|                          |                                    | 1986 (lin. 1)            |                    | Zeptometrix/<br>0810082cfhi | 2.2E+03 copies/mL                                |                 |
|                          |                                    | Bird 114 (lin. 1)        |                    | BEI / NR-50886              | 3.3E+03 copies/mL                                |                 |
| Yellow fever virus       | 1/1                                | 17D                      |                    | BEI / NR-116                | 1.2E+02 copies/mL                                | None            |
|                          |                                    |                          | PROTOZO            |                             |  | •               |
|                          |                                    | donovani 9515            |                    | BEI / NR-48822              | 1.0E+01 copies/mL                                |                 |
|                          |                                    | amazonensis              |                    | BEI / NR-49247              | 3.0E+01 copies/mL                                |                 |
|                          |                                    | <i>braziliensis</i> Vian | na                 | ATCC / 30879                | 1.0E+02 copies/mL                                | _               |
|                          |                                    | donovani, 1S             |                    | BEI / NR-48821              | 3.4E+01 copies/mL                                | Reduced         |
| Leishmania<br>spp. 12/12 |                                    | donovani chagasi         |                    | ATCC / 50133                | 3.0E+01 copies/mL                                | sensitivity for |
|                          | 12/12                              | infantum                 |                    | ATCC / 50134                | 1.0E+02 copies/mL                                | braziliensis    |
|                          | ,                                  | major IR173              |                    | BEI / NR-48816              | 3.0E+01 copies/mL                                | and infantum    |
|                          |                                    | mexicana MHOI            |                    | ATCC / 50157                | 2.9E+01 copies/mL                                | strains         |
|                          |                                    | panamensis PSC-1         |                    | BEI / NR-50162              | _  |                 |
|                          |                                    | tropica (MHOM,           | ·                  | BEI / NR-48820              | 3.0E+01 copies/mL                                |                 |
|                          |                                    | tropica (MHOM,           | /SU/58/strain-OD)  | ATCC / 50130                |  |                 |
|                          |                                    | venezuelensis M          | HOM-VE/80/H-16     | BEI / NR-29184              | 3.4E+01 copies/mL                                |                 |
| Plasmodium               | 10/10                              | falciparum               | IPC 4884           | BEI / MRA-1238              | 1.8E+02 copies/mL                                | None            |
| spp.                     | 10/10                              | , 4.0.04.4111            | SenTh021.09        | BEI / MRA-1182              | 3.0E+02 copies/mL                                | 140116          |

| Analyte                  | # Isolates<br>Detected /<br>Tested | Isolates Tested |                    | Source / ID            | Concentration Detected Tested up to 100× LoD¹ | Limitations     |
|--------------------------|------------------------------------|-----------------|--------------------|------------------------|---|-----------------|
| (Plasmodium              |                                    |                 | St. Lucia          | BEI / MRA-331          |   |                 |
| spp. assay)              |                                    |                 | Tanzania, 02000708 | BEI / MRA-1169         |   |                 |
|                          |                                    | vivax           | Chesson            | BEI / MRA-383          | 1.5E+02 copies/mL                             |                 |
|                          |                                    | VIVUX           | Panama             | ATCC / 30138           | 3.1E+02 copies/mL                             |                 |
|                          |                                    | ovale           | Wallikeri          | CDC / N8K9QKI9         | 2.4E+02 copies/mL                             |                 |
|                          |                                    | ovuie           | Curtisi            | CDC / N8K9QL0S         | 7.2E+02 copies/mL                             |                 |
|                          |                                    | knowlesi        | Strain H           | BEI / MRA-456G         | 2.4E+01 copies/mL                             |                 |
|                          |                                    | malariae        | Clinical Sample    | DLS / DLS17-<br>026015 | 1.9E+02 copies/mL                             |                 |
| Plasmodium<br>falciparum |                                    |                 | IPC 4884           | BEI / MRA-1238         | 1.8E+02 copies/mL                             | Reduced         |
| (Plasmodium              | 4/4                                | falciparum      | SenTh021.09        | BEI / MRA-1182         | 1.8E+03 copies/mL                             | sensitivity for |
| falciparum               |                                    |                 | St. Lucia          | BEI / MRA-331          | 2 0E 102 conios/ml                            | SenTh021.09     |
| assay)                   |                                    |                 | Tanzania, 02000708 | BEI / MRA-1169         | 3.0E+02 copies/mL                             |                 |
| Plasmodium               |                                    |                 | Chesson            | BEI / MRA-383          | 1.5E+02 copies/mL                             |                 |
| vivax/ovale              |                                    |                 | Panama             | ATCC / 30138           | 3.1E+02 copies/mL                             |                 |
| (Plasmodium              | 4/4                                |                 | Wallikeri          | CDC / N8K9QKI9         | 2.4E+02 copies/mL                             | None            |
| vivax/ovale<br>assay)    |                                    | ovale           | Curtisi            | CDC / N8K9QL0S         | 7.2E+02 copies/mL                             |                 |

<sup>&</sup>lt;sup>1</sup>Test concentrations were based on the LoD for the analyte in copies/mL with stock concentrations determined using a commercial qPCR assay.

Additionally, detection of BSL 3/4 infectious strains/isolates/serovars/species of *Bacillus* anthracis, Francisella tularensis, Yersinia pestis, West Nile virus, Zaire ebolavirus, Lassa virus, Marburgvirus, Yellow fever virus were evaluated by spiking whole blood at near infectious estimated LoD levels. Analytes detected only at ≥10-fold the estimated LoD were considered to have reduced sensitivity. Results are summarized in Table 13. Isolates used to evaluate the infectious estimated LoD are bolded.

Table 13. Summary of BioFire Global Fever Special Pathogens Panel Reactivity (Inclusivity) for Infectious BSL3/4 Analytes

| BSL3/4 An               | # Isolates Detected/Tested | Isolates 1            | ested                    | UCC <sup>1</sup> ID  | Concentration<br>Detected      | Limitations            |   |
|-------------------------|----------------------------|-----------------------|--------------------------|----------------------|--------------------------------|------------------------|---|
|                         |                            | Ames                  |                          | BACI008              | 6.4E+01 copies/mL              |                        |   |
|                         |                            | 108                   |                          | BACI226              | 1.9E+02 copies/mL              | †                      |   |
|                         | 2002013094                 |                       | BACI293                  | 1.9E+02 copies/mL    | 1                              |                        |   |
|                         |                            | Canadian Bison        |                          | BACI153              | 1.9E+02 copies/mL              | 1                      |   |
|                         |                            | K3                    |                          | BACI261              | 1.9E+02 copies/mL              | 1                      |   |
|                         |                            | Ohio ACB              |                          | BACI259              | 1.9E+02 copies/mL              | Reduced sensitivity    |   |
| Bacillus                | 10/12                      | PAK-1                 |                          | BACI309              | 1.9E+02 copies/mL              | for strains SK-102 and |   |
| anthracis               | 19,11                      | RA3                   |                          | BACI225              | 1.9E+02 copies/mL              | Vollum 1B              |   |
|                         |                            | SK-102 (Pakistan)     |                          | BACI126              | 6.4E+02 copies/mL              |                        |   |
|                         |                            | South Africa (BA 1035 | ·)                       | BACI207              | 1.9E+02 copies/mL              |                        |   |
|                         |                            | Sterne                | ,                        | BACI012              | 1.9E+02 copies/mL              |                        |   |
|                         |                            | Turkey #32            |                          | BACI260              | 1.9E+02 copies/mL              | =                      |   |
|                         |                            | Vollum 1B             |                          | BACI124              | 6.4E+02 copies/mL              |                        |   |
|                         |                            | Subspecies            | Strain                   | -                    | 2 22 23/                       |                        |   |
|                         |                            |                       | Schu4                    | FRAN016              | 1.2E+01 copies/mL              |                        |   |
|                         |                            | tularensis            | Scherm                   | FRAN031              | 3.6E+01 copies/mL              |                        |   |
|                         |                            |                       | WY96                     | FRAN072              | 3.6E+01 copies/mL              |                        |   |
|                         |                            |                       | Holarctica, LVS          | FRAN004              | 3.6E+01 copies/mL              |                        |   |
| E                       |                            |                       | Holarctica               | FRAN012              | 3.6E+01 copies/mL              |                        |   |
| Francisella             | 9/9                        | /9<br>holarctica      | Holarctica, 425          | FRAN029              | 3.6E+01 copies/mL              | None                   |   |
| tuiarensis              | tularensis                 |                       | HD,<br>DB082106G         | FRAN035              | 2.3E+01 copies/mL              |                        |   |
|                         |                            |                       | VT68                     | FRAN025              | 3.6E+01 copies/mL              | -                      |   |
|                         |                            | novicida              | F6168                    | FRAN134              | 3.6E+01 copies/mL              |                        |   |
|                         |                            |                       | U112,<br>GA993550        | FRAN003              | 3.6E+01 copies/mL              |                        |   |
|                         |                            | Biovar                | Strain                   |                      |                                |                        |   |
|                         |                            |                       | CO92                     | YERS023              | 1.5E+02 copies/mL              |                        |   |
|                         |                            |                       | A1122                    | YERS078              | 4.5E+02 copies/mL              |                        |   |
|                         |                            |                       | Dodson <sup>2</sup>      | YERS073              | 3.6E+02 CFU/mL                 |                        |   |
|                         |                            |                       | Orientalis               | Java 9               | YERS022                        | 4.5E+02 copies/mL      | - |
|                         |                            |                       | PBM19                    | YERS018              | 4.5E+02 copies/mL              |                        |   |
| Yersinia                |                            |                       | Shasta                   | YERS074              | 4.5E+02 copies/mL              |                        |   |
| Pestis                  | 12/12                      |                       | Angola                   | YERS080              | 4.5E+02 copies/mL              | None                   |   |
|                         |                            |                       | Antigua                  | YERS016              | 4.5E+02 copies/mL              | 1                      |   |
|                         |                            | Antiqua               | Nairobi                  | YERS017              | 4.5E+02 copies/mL              |                        |   |
|                         |                            |                       | Pestoides F <sup>2</sup> | YERS020              | 3.6E+02 CFU/mL                 |                        |   |
|                         |                            |                       | Harbin 35                | YERS021              | 5.3E+02 copies/mL <sup>3</sup> |                        |   |
|                         |                            | Madiavalia            | KIM5                     |                      | 6.4E+02 copies/mL <sup>3</sup> |                        |   |
|                         |                            | Medievalis            |                          | YERS082              |                                | -                      |   |
|                         |                            | Strain/Is             | Nicholisk 41             | YERS083              | 4.5E+02 copies/mL              |                        |   |
| Zaire<br>ebolavirus 4/4 | Makona – SL3864.1          | olate                 | Ebola027                 | 1.1E+03 copies/mL    | -                              |                        |   |
|                         |                            | Mayinga (Zaire 76)    |                          | Ebola001             | 2.1E+03 copies/mL              |                        |   |
|                         | 4/4                        | Gabon                 |                          | Ebola034             | 3.2E+03 copies/mL              | None                   |   |
|                         |                            | Kikwit '95            |                          | Ebola034<br>Ebola007 | · '                            | 1                      |   |
|                         |                            |                       |                          |                      | 3.1E+03 copies/mL              | -                      |   |
|                         |                            | Luebo                 |                          | Ebola035             | 3.1E+03 copies/mL              |                        |   |
|                         | 2 /2                       | Josiah                |                          | Arena002             | 2.4E+01 PFU/mL                 | <u>.</u> .             |   |
| Lassa virus             | 2/2                        | Macenta <sup>4</sup>  |                          | Arena009             | 7.2E+01 PFU/mL                 | None                   |   |
|                         | Pinneo <sup>4</sup>        |                       | Arena003                 | 7.2E+01 PFU/mL       |                                |                        |   |

| Analyte                | # Isolates<br>Detected/Tested | Isolates Tested |              | UCC <sup>1</sup> ID | Concentration<br>Detected        | Limitations        |
|------------------------|-------------------------------|-----------------|--------------|---------------------|----------------------------------|--------------------|
|                        |                               | Lineage         | Strain       |                     |                                  |                    |
| Marburg                |                               | Ravn virus      | RAVN         | Marb002             | 2.6E+02 copies/mL                |                    |
| Marburgvirus           | 2/2                           |                 | Ci67         | Marb003             | 5.0E+02 copies/mL                | None               |
| man sang mas           |                               | Marburg virus   | Angola       | Marb005             | 1.5E+03 copies/mL                |                    |
|                        |                               |                 | Musoke       | Marb001             | 7.8E+02 copies/mL                |                    |
| West Nile              | 1 /1                          | Bz NY99         |              | Flavi022            | 1.6E+02 copies/mL                | None for West Nile |
| virus<br>(Lineage 1)   | 1/1                           | Eg101           |              | Flavi016            | 4.8E+02 copies/mL                | Virus<br>Lineage 1 |
|                        |                               | Asibi           |              | FLAVI005            | 1.2E+01 copies/mL                | None               |
|                        |                               | SVM 3-18-09     |              | BEI NR-49799        | 1.2E+04 copies/mL                |                    |
| Yellow fever virus 4/4 | CAREC M2-09                   |                 | BEI NR-50062 | 1.2E+03 copies/mL   | Reduced sensitivity <sup>5</sup> |                    |
|                        | INHRR 7a-05                   |                 | BEI NR-50071 | 1.2E+03 copies/mL   |                                  |                    |
|                        |                               | INHRR 10a-10    |              | BEI NR-50063        | 1.2E+03 copies/mL                |                    |

<sup>&</sup>lt;sup>1</sup>U.S. Department of Defense Unified Culture Collection

#### Exclusivity (Specificity)

Analytical specificity (exclusivity) of the BioFire Global Fever Special Pathogens Panel assays was evaluated by challenging the system with a large collection of organisms and viruses prepared at high concentrations. On-panel organisms and viruses were tested to assess the potential for intra-panel cross-reactivity. Off-panel organisms and viruses (those not intended to be detected by the panel) were tested to assess the potential for non-specific amplification of other pathogens or potential contaminants. Organisms and viruses for off-panel testing were selected based on a combination of several factors including: relatedness to specific species detected by the BioFire Global Fever Special Pathogens Panel (near-neighbors); clinical relevance (cause illness or symptoms similar to the panel pathogens); likelihood of being present in blood as a co-infection based on a geographical region or specific population to which a panel pathogen is endemic; and genetic similarity to BioFire Global Fever Special Pathogens Panel assay primers, as determined by in silico analyses.

On-panel and off-panel testing included 217 isolates of bacteria, viruses, fungi, and protozoa tested at high concentration (typically >10<sup>6</sup> genomic copies/mL). Table 14 shows a complete list of the on- and off-panel bacteria, viruses, protozoa, and fungi that were tested and received the expected BioFire Global Fever Panel Special Pathogens test result (negative for all assays; no cross-reactivity) or for which in silico analysis does not predict cross-reactivity.

<sup>&</sup>lt;sup>2</sup>The *Y. pestis* bv. Orientalis Dodson and *Y. pestis* bv. Antiqua Pestoides F strains lack the pPCP plasmid targeted by the qPCR assay, and as a result the nucleic acid concentration could not be quantified. Inclusivity for these two strains was therefore based on concentrations obtained for these two strains, and the CO92 reference strain, by enumeration in Colony Forming Units (CFU)/mL.

<sup>&</sup>lt;sup>3</sup> After testing was completed, it was determined that *Yersinia pestis* bv. Medievalis strains Harbin35 and KIM5 were evaluated at higher than 3× the estimated LoD, at 3.5× and 4.2×, respectively. Based on amplification data it is expected that these two *Y. pestis* strains will be detected at 3×LoD.

<sup>&</sup>lt;sup>4</sup>The Lassa virus Macenta and Pinneo strains were not detected by the qPCR assay, only the Josiah strain evaluated in the Estimated LoD study was. Therefore, inclusivity testing for Lassa virus was performed based on concentrations obtained by enumeration in Plaque Forming Units (PFU)/mL.

<sup>&</sup>lt;sup>5</sup> YFV strains SVM 3-18-09, CAREC M2-09, NHRR 7a-05, and INHRR 10a-10 were isolated from neighboring regions (Trinidad and Venezuela). In silico analyses indicate these strains are closely related and do not represent a broad diversity of sequences.

Table 14. On and Off Panel Organisms and Viruses with No Cross-Reactivity with BioFire Global Fever Special Pathogens Panel Assays

| Special Pathogens Panel Assays        | ON-PANEL   |  |  |  |  |  |  |
|---------------------------------------|--|--|--|--|--|--|--|
|                                       | Bacteria   |  |  |  |  |  |  |
| Bacillus anthracis                    | Leptospira interrogans                           |  |  |  |  |  |  |
| Francisella tularensis subsp.         | Yersinia pestis (2 strains: A1122 and CO92)      |  |  |  |  |  |  |
| tularensis                            |  |  |  |  |  |  |  |
| Cuire and Canada have a unbania       | Viruses  | NASAH WAS AASAH WAS SING OF SING OF                    |  |  |  |  |  |
| Crimean-Congo hemorrhagic fever virus | Zaire ebolavirus                                 | Marburg Marburgvirus variant Musoke                    |  |  |  |  |  |
| Chikungunya virus                     | Sudan ebolavirus                                 | Marburg marburgvirus (RAVN)                            |  |  |  |  |  |
| Dengue virus Serotype 1               | Bundibugyo ebolavirus                            | West Nile virus Lineage 1                              |  |  |  |  |  |
| Dengue virus Serotype 2               | Taï Forest ebolavirus                            | West Nile virus Lineage 2                              |  |  |  |  |  |
| Dengue virus Serotype 3               | Reston ebolavirus                                | Yellow fever virus                                     |  |  |  |  |  |
| Dengue virus Serotype 4               | Lassa virus                                      |  |  |  |  |  |  |
|                                       | Protozoa   |  |  |  |  |  |  |
| Leishmania donovani                   |  |  |  |  |  |  |  |
| Plasmodium falciparum                 | Plasmodium vivax                                 | Plasmodium ovale                                       |  |  |  |  |  |
|                                       | OFF-PANEL  |  |  |  |  |  |  |
|                                       | Bacteria   |  |  |  |  |  |  |
|                                       | Tested   |  |  |  |  |  |  |
| Acinetobacter baumannii               | Enterococcus faecium                             | Salmonella enterica subs. enterica serovar Muenchen    |  |  |  |  |  |
| Bacillus brevis                       | Francisella persica (formerly Wolbachia persica) | Salmonella enterica subs. enterica serovar Newport     |  |  |  |  |  |
| Bacillus cereus                       | Francisella philomiragia (formerly Yersinia)     | Salmonella enterica subs. enterica<br>serovar Rubislaw |  |  |  |  |  |
| Bacillus circulans                    | Klebsiella oxytoca                               | Salmonella enterica susb. enterica                     |  |  |  |  |  |
|                                       |  | serovar Saintpaul Salmonella enterica subs. enterica   |  |  |  |  |  |
| Bacillus coagulans                    | Legionella pneumophila                           | serovar Tennessee                                      |  |  |  |  |  |
| Bacillus halodurans                   | Leptospira biflexa                               | Salmonella enterica subs. enterica serovar Thompson    |  |  |  |  |  |
| Bacillus licheniformis                | Leptospira meyeri                                | Salmonella enterica subs. enterica serovar Typhimurium |  |  |  |  |  |
| Bacillus megaterium                   | Leptospira terpstrae genomospecies 4             | Salmonella enterica subs. houtenae                     |  |  |  |  |  |
| Bacillus mycoides                     | Leptospira vanthielii genomospecies 3            | Salmonella enterica subs. indica                       |  |  |  |  |  |
| Bacillus pumilus                      | Leptospira wolbachii                             | Salmonella enterica subs. salamae                      |  |  |  |  |  |
| Bacillus subtilis                     | Leptospira yanagawae<br>genomospecies 5          | Salmonella enterica subsp. enterica<br>Paratyphi A     |  |  |  |  |  |
| Bacillus thuringiensis                | Listeria monocytogenes                           | Salmonella enterica subsp. enterica<br>Typhi           |  |  |  |  |  |
| Bacteroides fragilis                  | Mycobacterium tuberculosis                       | Serratia marcescens                                    |  |  |  |  |  |
| Bordetella bronchiseptica             | Mycoplasma pneumoniae                            | Staphylococcus aureus                                  |  |  |  |  |  |
| Borrelia burgdorferi                  | Neisseria meningitidis                           | Streptococcus agalactiae                               |  |  |  |  |  |
| Brucella melitensis                   | Orientia chuto (tsutsugamushi)                   | Streptococcus pneumoniae                               |  |  |  |  |  |
| Burkholderia cepacia                  | Proteus mirabilis                                | Streptococcus pyogenes                                 |  |  |  |  |  |

| Burkholderia mallei               | Pseudomonas aeruginosa                                 | Treponema pallidum pallidum               |
|-----------------------------------|--|---|
| Burkholderia pseudomallei         | Rickettsia prowazekii                                  | Vibrio cholerae                           |
| Chlamydophila pneumoniae          | Rickettsia ricketsii                                   | Yersinia aldovae                          |
| Chlamydophila psittaci            | Rickettsia typhi                                       | Yersinia bercovieri                       |
| Citrobacter koseri                | Salmonella enterica subs.bongori                       | Yersinia entericolitica                   |
| Clostridium bifermentans          | Salmonella enterica subs. arizonae                     | Yersinia fredericksenii                   |
| Clostridium                       |  | -   |
| botulinum/sporogenes              | Salmonella enterica subs. diarizoniae                  | Yersinia intermedia                       |
| Clostridium perfringens           | Salmonella enterica subs. enterica                     | Yersinia kristensenii                     |
| . , ,                             | serovar Enteritidis Salmonella enterica subs. enterica |   |
| Clostridium sordellii             | serovar Heidelberg                                     | Yersinia mollaretii                       |
| G : # 1 :::                       | Salmonella enterica subs. enterica                     | y   |
| Coxiella burnetii                 | serovar Javiana  | Yersinia pseudotuberculosis               |
| Enterobacter aerogenes            | Salmonella enterica subs. enterica                     | Yersinia rohdei                           |
| _                                 | serovar Montevideo                                     |   |
| Enterococcus faecalis             | Yersinia similis                                       | Yersinia ruckeri                          |
| 2 11 1 15                         | In silico Analysis Only                                |   |
| Bacillus luciferensis             | V2   |   |
|                                   | Viruses  |   |
| Adenovirus 1                      | Tested HPIV-1  | Omsk hemorrhagic fever                    |
| Adenovirus 3                      | HPIV-3   | Parvovirus                                |
| Adenovirus 5                      | Hughes virus   | Powassan virus                            |
| Aura virus                        | Human herpesvirus 6B                                   | Rabies virus                              |
|                                   | Human immunodeficiency virus, type                     |   |
| Barmah Forest virus               | 1  | Rift Valley fever virus                   |
| Bunyamwera virus                  | Human immunodeficiency virus, type 2                   | Zika virus                                |
| Coronavirus NL63                  | Human T-lymphotropic virus, type 1                     | Ross River virus                          |
| Cytomegalovirus                   | Human T-lymphotropic virus, type 2                     | Human respiratory syncytial virus         |
| Dugbe virus                       | Influenza A H1N1-2009                                  | Rubella virus                             |
| Eastern equine encephalitis virus | Influenza A H3N2                                       | SARS-CoV-2                                |
| Enterovirus, HEV-71               | Influenza B virus                                      | Saint Louis encephalitis virus            |
| Epstein Barr virus                | Japanese encephalitis virus                            | Semliki Forest virus                      |
| Flexal virus                      | Junin virus (2 strains: XJ and Candid)                 | Sindbis virus                             |
| Guanarito virus                   | Machupo virus  | Spondweni virus                           |
| Hantaan virus                     | Mayaro virus   | Tickborne encephalitis virus              |
| Hazara virus                      | Measles virus  | Tonate virus                              |
| Hendra virus                      | Metapneumovirus  | Una virus                                 |
| Hepatitis A virus                 | Middelburg virus                                       | Usutu virus                               |
| Hepatitis B virus                 | Mopeia virus   | Vaccinia virus                            |
| Hepatitis C virus                 | Mumps virus  | Varicella zoster virus                    |
| Herpes simplex virus type 2       | Murray Valley encephalitis virus                       | Venezuelan equine encephalomyelitis virus |

| In silico Analysis Only |                                     |   |  |  |  |
|-------------------------|-------------------------------------|---|--|--|--|
| Avalon virus            | Lymphocytic choriomeningitis virus  | Sabia virus (Brazilian hemorrhagic fever) |  |  |  |
| Bas-Congo virus         | Pirital virus                       | Variola major                             |  |  |  |
|                         | Protozoa                            |   |  |  |  |
|                         | Tested                              |   |  |  |  |
| Babesia microti         | Toxoplasma gondii                   | Trypanosoma cruzi                         |  |  |  |
| Cyclospora cayetanensis | Trypanosoma brucei                  | Trypanosoma cruzi                         |  |  |  |
|                         | Fungus                              |   |  |  |  |
|                         | Tested                              |   |  |  |  |
| Aspergillus fumigatus   | Cryptococcus neoformans var. grubii |   |  |  |  |
| Helminths               |                                     |   |  |  |  |
| Tested                  |                                     |   |  |  |  |
| Schistosoma mansoni     |                                     |   |  |  |  |

Table 15 lists the BioFire Global Fever Special Pathogens Panel assays and corresponding organisms and viruses for which some cross-reactivity was identified (either empirically observed in testing or predicted by in silico analyses).

Table 15. Observed or Predicted Cross-Reactivity of the BioFire Global Fever Special Pathogens Panel

| CROSS-REACTIVE ORGANISM  On-Pa                                    | BIOFIRE GLOBAL FEVER SPECIAL PATHOGENS PANEL TEST RESULT |  |
|---|--|--|
| Plasmodium knowlesi <sup>1</sup> Plasmodium malariae <sup>2</sup> | Plasmodium vivax/ovale Detected                          |  |
| Off-Pa  | anel   |  |
| Francisella hispaniensis <sup>3</sup>                             | - Francisella tularensis Detected                        |  |
| Francisella tularensis subsp. mediasiatica <sup>4</sup>           | Truncisena talarensis Detectea                           |  |
| O'nyong-nyong virus   | Chikungunya virus Detected                               |  |
| Crithidia fasciculata <sup>5</sup>                                | Laichmania can Dotacted                                  |  |
| Leptomonas seymouri <sup>6</sup>                                  | - <i>Leishmania</i> spp. Detected                        |  |
| Plasmodium berghei <sup>7</sup>                                   |  |  |
| Plasmodium brasilianum <sup>7</sup>                               |  |  |
| Plasmodium cynomolgi <sup>7</sup>                                 | Olympia diamana and                                      |  |
| Plasmodium fieldi <sup>7</sup>                                    | Plasmodium spp. and Plasmodium vivax/ovale Detected      |  |
| Plasmodium fragile <sup>7</sup>                                   | Plasificatum vivax/ovale Detected                        |  |
| Plasmodium inui <sup>7</sup>                                      |  |  |
| Plasmodium simiovale <sup>7</sup>                                 |  |  |

<sup>&</sup>lt;sup>1</sup>Cross-reactive with the Plasmodium vivax/ovale assay at concentrations ≥2.2E+04 copies/mL (~100×LoD).

<sup>&</sup>lt;sup>2</sup> In silico analysis predicts potential cross-reactivity with the *Plasmodium vivax* assay; cross-reactivity was not observed at the concentration evaluated, 1.9E+05 copies/mL.

<sup>&</sup>lt;sup>3</sup> Cross-reactivity was predicted by in silico analysis and observed during wet testing; *Francisella hispaniensis* is a pathogenic *Francisella* species.

<sup>&</sup>lt;sup>4</sup> In silico analysis predicts cross-reactivity; the potential for *Francisella tularensis* subsp. *mediasiatica* to cause disease in humans is unknown.

<sup>&</sup>lt;sup>5</sup> Crithidia fasciculata is a non-human infective trypanosomatid, however there have been very rare cases where a closely related Crithidia strain was found to be infective and act much like Leishmania spp.

<sup>&</sup>lt;sup>6</sup>Leptomonas seymouri is an opportunistic parasite in immunocompromised individuals, particularly those infected with visceral leishmaniasis.

<sup>&</sup>lt;sup>7</sup> Plasmodium spp. that typically infect non-human primates and rodents but are rarely found in humans.

#### Interference

Potentially interfering substances were selected for evaluation based upon whether the substance may normally be found in blood or may be introduced into blood specimens during collection, handling, or testing. These substances included endogenous substances (e.g., albumin, immunoglobulin, etc.), exogenous substances (e.g., antibiotic and antiviral drugs), microorganisms such as viral and bacterial species that may be present as co-infections, and technique-specific substances that may be introduced into a sample during routine laboratory handling including collection tubes.

The concentrations of endogenous and exogenous substances tested on the BioFire Global Fever Special Pathogens Panel were selected to represent a worst-case scenario based on a reference concentration of normal to high levels expected to be present in clinical specimens. Potentially competing microorganisms were tested at the highest levels possible based on stock concentrations. Technique-specific substances were tested at levels exceeding what would be expected to be found due to incidental contact, accidental addition, or during their use as solvents. Results are shown in Table 16.

Potential interference was only observed for heparin and TRIzol when testing analytes at near-LoD concentrations.

Table 16. Results for Potentially Interfering Substances Tested on the BioFire Global Fever Special Pathogens Panel

| Potentially Interfering Substance | Concentration Tested  | Results         |
|-----------------------------------|-----------------------|-----------------|
|                                   | Endogenous Substances |                 |
| Albumin                           | 60.0 mg/mL            | No interference |
| Bilirubin (Conjugated)            | 0.41 mg/mL            | No interference |
| Bilirubin (Unconjugated)          | 0.41 mg/mL            | No interference |
| Cholesterol (total)               | 4.2 mg/mL             | No interference |
| Glucose                           | 10.1 mg/mL            | No interference |
| Hemoglobin                        | 137.0 mg/mL           | No interference |
| Immunoglobulins                   | 60.0 mg/mL            | No interference |
| Triglycerides                     | 15.1 mg/mL            | No interference |
| White Blood Cells                 | 6.1E+06 cells/mL      | No interference |
|                                   | Exogenous Substances  |                 |
| Artemether-Lumefantrine           | 0.0004 mg/mL          | No interference |
| Atovaquone                        | 0.005 mg/mL           | No interference |
| Proguanil                         | 0.001 mg/mL           | No interference |
| Mefloquine                        | 0.0017 mg/mL          | No interference |
| Amphotericin B                    | 0.002 mg/mL           | No interference |
| Pentamidine                       | 0.0015 mg/mL          | No interference |
| Fluconazole                       | 0.026 mg/mL           | No interference |
| Amoxicillin                       | 0.062 mg/mL           | No interference |
| Azithromycin                      | 0.011 mg/mL           | No interference |
| Ceftriaxone                       | 1.0 mg/mL             | No interference |
| Ciprofloxacin                     | 0.012 mg/mL           | No interference |
| Clindamycin                       | 0.055 mg/mL           | No interference |
| Doxycycline                       | 0.02 mg/mL            | No interference |

| Potentially Interfering Substance                 | Concentration Tested          | Results                 |
|---|-------------------------------|-------------------------|
| Gentamicin  | 0.036 mg/mL                   | No interference         |
| Meropenem   | 0.39 mg/mL                    | No interference         |
| Sulfamethoxazole                                  | 0.38 mg/mL                    | No interference         |
| Vancomycin  | 0.12 mg/mL                    | No interference         |
| Cycloserine                                       | 75.0 mg/mL                    | No interference         |
| Isoniazid   | 0.06 mg/mL                    | No interference         |
| Oseltamivir                                       | 0.0005 mg/mL                  | No interference         |
| Ribavirin   | 0.011 mg/mL                   | No interference         |
| Tenofovir   | 0.001 mg/mL                   | No interference         |
| Acetaminophen                                     | 0.16 mg/mL                    | No interference         |
| Aspirin (Acetylsalicylic Acid)                    | 0.03 mg/mL                    | No interference         |
| Ibuprofen   | 0.22 mg/mL                    | No interference         |
| Prednisone  | 0.0001 mg/mL                  | No interference         |
| Prednisolone                                      | 1.2 mg/mL                     | No interference         |
| Cortisone   | 0.001 mg/mL                   | No interference         |
| Artesunate  | 0.1 mg/mL                     | No interference         |
|   | Competitive Microorganisms    |                         |
| Corynebacterium diphtheriae                       | 1:10 of Stock                 | No interference         |
| Staphylococcus epidermidis                        | 3.8E+06 CFU/mL                | No interference         |
| Escherichia coli                                  | 1:10 of Stock                 | No interference         |
| Klebsiella pneumoniae                             | 5.5E+04 CFU/mL                | No interference         |
| Haemophilus influenzae                            | 1.0E+08 CFU/mL                | No interference         |
| Herpes Simplex virus                              | 1.2E+05 PFU/mL                | No interference         |
| Epstein-Barr virus                                | 3.3E+07 copies/mL             | No interference         |
| Cytomegalovirus (CMV) AD-169                      | 1:10 of Stock                 | No interference         |
| Human Immunodeficiency virus<br>(HIV-1 and HIV-2) | 1:10 of Stocks                | No interference         |
| Plasmodium vivax                                  | 1.5E+06 copies/mL             | No interference         |
|   | Technique Specific Substances |                         |
| Bleach  | 1% v/v                        | No interference         |
| Povidone-iodine                                   | 1% v/v                        | No interference         |
| Ethanol   | 2% v/v                        | No interference         |
| TRIzol  | 2-3% v/v                      | Potentially Interfering |
| DMSO  | 2% v/v                        | No interference         |
| Methanol  | 2% v/v                        | No interference         |
| Saline  | 2% v/v                        | No interference         |
| Chloroform  | 2% v/v                        | No interference         |
| Acetone   | 2% v/v                        | No interference         |
| Hydrochloric Acid (HCl)                           | 0.0005N                       | No interference         |
|   | Blood Collection Tubes        |                         |
| Citrate (sodium)                                  | ~0.32%                        | No interference         |
| EDTA in excess (5x)                               | ~9.0 mg/mL                    | No interference         |
| Heparin   | ~19.0 USP/mL                  | Potentially Interfering |
| ·   | 2.2mg/mL (trisodium citrate)  |                         |
| Acid-citrate-dextrose (ACD)                       | 0.8 mg/mL (citric acid)       | No interference         |
| ` '   | 2.5 mg/mL (dextrose)          |                         |
| Sodium polyanethenole sulfonate (SPS)             | 0.72 mg/mL                    | No interference         |
| Serum Separation Tubes                            | N/A                           | No interference         |

#### Reproducibility

Assay reproducibility was evaluated using three contrived whole blood samples with a mixture of four representative panel analytes: one bacterium, one virus, and two protozoa. For each analyte, one sample was spiked at a moderate positive (3×LoD) level, another sample at a low positive (1×LoD) level, and the third sample was not spiked (negative). Six replicates of each sample were tested at three locations on five different days, providing a total of 90 replicate test results per sample. On each test day at each site, two different operators used three BioFire FilmArray 2.0 instruments; the BioFire Global Fever Special Pathogens Panel pouch lot was rotated daily. In total, 270 valid test results were obtained for the reproducibility evaluation of the BioFire Global Fever Special Pathogens Panel.

The primary assessment of reproducibility is based on a comparison of the observed test results (Detected/Not Detected) to the expected test results (Detected for spiked samples, Not Detected for un-spiked samples). The detection rate and percent agreement between observed and expected test results are shown in Table 17. The expected percent agreement was ≥95%. Based on the percent agreement between observed and expected test results, the reproducibility evaluation demonstrates that the BioFire Global Fever Special Pathogens Panel can provide accurate and highly reproducible test results in the context of multiple variables that may be expected in a clinical testing environment, including analyte concentration, location, test day, operator, instruments, and reagent lot.

Table 17. Reproducibility of the BioFire Global Fever Special Pathogens Panel

| rabie 17. Kepr                        | oducionity 01                                      | the BioFire Global F                                    | ever special rath | ogens ran   |                | Pato (n/N)                   |  |  |  |
|---------------------------------------|--|---|-------------------|---|----------------|------------------------------|--|--|--|
| Anal                                  | luto   | Concentration Tested                                    |                   | Detection Rate (n/N)  % Agreement with Expected Result  [95% Confidence Interval]  Site 1 Site 2 Site 3 All Sites |                |                              |  |  |  |
| (Source                               |  | (copies/mL)   | Expected Result   |   |                |                              |  |  |  |
| (35416)                               | c / 1.5 /  | (copies, init,  |                   |   |                |                              |  |  |  |
|                                       |  | Moderate Positive<br>3× LoD<br>(1.0E+03)                | Detected          | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | 90/90<br>100%<br>[95.9-100%]                     |  |  |
| Leptospira i<br>serovar icteroh       | aemorrhagiae                                       | Low Positive<br>1× LoD<br>(3.4E+02)                     | Detected          | 27/30<br>90.0%  | 28/30<br>93.3% | 26/30<br>86.7%               | <b>81/90</b><br><b>90.0%</b><br>[82.1-<br>94.6%] |  |  |
| (ATTC /                               | 23581)   | Negative<br>(No Analyte)                                | Not Detected      | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | <b>90/90</b><br><b>100%</b><br>[95.9-100%]       |  |  |
|                                       |  | Moderate Positive<br>3× LoD<br>(1.0E+03)                | Detected          | 29/30<br>96.7%  | 30/30<br>100%  | 30/30<br>100%                | <b>89/90 98.9%</b> [94.0- 99.8%]                 |  |  |
| <b>Dengue</b><br><b>DEN</b><br>New Gu | V-2  | Low Positive<br>1× LoD<br>(3.4E+02)                     | Detected          | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | <b>90/90 100%</b> [95.9-100%]                    |  |  |
| (Zeptometrix ,                        | / 0810089CF)                                       | Negative<br>(No Analyte)                                | Not Detected      | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | <b>90/90</b><br><b>100%</b><br>[95.9-100%]       |  |  |
|                                       |  | Moderate Positive<br>3.4× LoD <sup>1</sup><br>(3.4E+01) | Detected          | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | 90/90<br>100%<br>[95.9-100%]                     |  |  |
| 1S (MHOM/SD/62/1S)                    | Low Positive<br>1.1× LoD <sup>1</sup><br>(1.1E+01) | Detected  | 30/30<br>100%     | 30/30<br>100%   | 30/30<br>100%  | 90/90<br>100%<br>[95.9-100%] |  |  |  |
| (BEI / NR-48821)                      |  | Negative<br>(No Analyte)                                | Not Detected      | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | 90/90<br>100%<br>[95.9-100%]                     |  |  |
| Plasmodium<br>falciparum              | Plasmodium<br>spp.                                 | Moderate Positive<br>1.5× LoD <sup>2</sup><br>(2.7E+02) | Detected          | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                | <b>90/90</b><br><b>100%</b><br>[95.9-100%]       |  |  |

| Analyte<br>(Source / ID)                  |                                | Concentration Tested (copies/mL) Expected Result        |              | Detection Rate (n/N)  % Agreement with Expected Result  [95% Confidence Interval] |                |                                      |  |  |
|---|--------------------------------|---|--------------|---|----------------|--------------------------------------|--|--|
|   |                                |   |              | Site 1  | Site 2         | Site 3                               | All Sites  |  |
| IPC 4884<br>(BEI / MRA-<br>1238)          | Detection<br>Results           | Low Positive<br>0.5× LoD <sup>2</sup><br>(9.0E+01)      | Detected     | 28/30<br>93.3%  | 30/30<br>100%  | 29/30<br>96.7%                       | <b>87/90</b><br><b>96.7%</b><br>[90.7-<br>98.9%] |  |
|   |                                | Negative<br>(No Analyte)                                | Not Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                        | <b>90/90</b><br><b>100%</b><br>[95.9-100%]       |  |
|   | Plasmodium                     | Moderate Positive<br>1.5× LoD <sup>2</sup><br>(2.7E+02) | Detected     | 29/30<br>96.7%  | 30/30<br>100%  | 28/30<br>93.3%                       | <b>87/90</b><br><b>96.7%</b><br>[90.7-<br>98.9%] |  |
|   | falciparum  Detection  Results | Low Positive<br>0.5× LoD <sup>2</sup><br>(9.0E+01)      | Detected     | 18/30<br>60.0%  | 24/30<br>80.0% | 21/30<br>70.0%                       | <b>63/90</b><br><b>70.0%</b><br>[59.9-<br>78.5%] |  |
|   |                                | Negative<br>(No Analyte)                                | Not Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%                        | <b>90/90</b><br><b>100%</b><br>[95.9-100%]       |  |
| Overall Agreement with<br>Expected Result |                                | All Concentrations                                      | All Results  |   | 96             | <b>/1350</b><br><b>.8%</b><br>97.2%] |  |  |

<sup>&</sup>lt;sup>1</sup>Due to a correction in the stock concentration, *L. donovani* was evaluated at 3.4×LoD and 1.1×LoD.

Reproducibility Comparison Between BioFire FilmArray 2.0 and BioFire FilmArray Torch

Performance of the BioFire Global Fever Special Pathogens Panel was compared between the BioFire FilmArray 2.0 and BioFire FilmArray Torch systems by examining analyte detection of contrived whole blood samples. The testing incorporated a range of potential variation introduced by operator, instrument system, analyte concentration, and reagent lot for a total of 90 replicates for each analyte concentration distributed equally over three BioFire FilmArray 2.0 systems and three BioFire FilmArray Torch systems.

A summary of results (percent (%) agreement with the expected Detected or Not Detected result) for the analytes by BioFire FilmArray system is provided in Table 18. Overall agreement between observed and expected results was 99.5% for the BioFire FilmArray 2.0 system and 99.1% for the BioFire FilmArray Torch system demonstrating similar performance between both instrument systems.

<sup>&</sup>lt;sup>2</sup> Due to a correction in the stock concentration, *P. falciparum* was evaluated at 1.5×LoD and 0.5×LoD.

Table 18. Reproducibility of the BioFire Global Fever Special Pathogens Panel Test Results on BioFire FilmArray 2.0 and BioFire FilmArray Torch

| Finital                  | Luy 2.0 anu 1                  | Generation                              | TOTCH           | Detection Rate (n/N) % Agreement with Expected Result |                |                |  |                |                |                |  |
|--------------------------|--------------------------------|---|-----------------|---|----------------|----------------|--|----------------|----------------|----------------|--|
| Ana                      | -                              | Concentration<br>Tested                 | Expected        | Bio   | Fire Film      |                | BioFire FilmArray Torch Platform             |                |                |                |  |
| (Sourc                   | e / ID)                        | (copies/mL)                             | Result          | System<br>1   | System 2       | System 3       | All FA 2.0<br>Systems<br>[95% CI]            | System<br>1    | System 2       | System 3       | All FA Torch<br>Systems<br>[95% CI]          |
| Leptospira i             | interrogans                    | Moderate Positive<br>3×LoD<br>(1.0E+03) | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | 90/90<br>100%<br>[95.9-100]                  |
|                          | ovar<br>norrhagiae             | Low Positive<br>1×LoD<br>(3.4E+02)      | Detected        | 29/30<br>96.7%  | 29/30<br>96.7% | 28/30<br>93.3% | <b>86/90</b><br><b>95.6%</b><br>[89.1-98.3%] | 29/30<br>96.7% | 28/30<br>93.3% | 28/30<br>93.3% | <b>85/90</b><br><b>94.4%</b><br>[87.6-97.61] |
| (ATTC /                  | 23581)                         | Negative<br>(No Analyte)                | Not<br>Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100]    |
| Dengu                    |                                | Moderate Positive<br>3×LoD<br>(1.0E+03) | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| New G                    | IV-2<br>uinea C                | Low Positive<br>1×LoD<br>(3.4E+02)      | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 29/30<br>96.7% | 30/30<br>100%  | <b>89/90</b><br><b>98.9%</b><br>[94.0-99.8%] |
| (Zeptometrix             | / 0810089CF)                   | Negative<br>(No Analyte)                | Not<br>Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 29/30<br>96.7% | <b>89/90</b><br><b>98.9%</b><br>[94.0-99.8%] |
| Leishmanid               | a donovani                     | Moderate Positive<br>3×LoD<br>(3.0E+01) | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| 1S (MHOM                 | /SD/62/1S)                     | Low Positive<br>1×LoD<br>(1.0E+01)      | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| (BEI / NF                | R-48821)                       | Negative<br>(No Analyte)                | Not<br>Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
|                          | Plasmodium                     | Moderate Positive<br>3×LoD<br>(5.4E+02) | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| _,                       | spp.  Detection  Results       | Low Positive<br>1×LoD<br>(1.8E+02)      | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| Plasmodium<br>falciparum | nesuits                        | Negative<br>(No Analyte)                | Not<br>Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |
| (BEI / MRA-<br>1238)     | Plasmodium                     | Moderate Positive<br>3×LoD<br>(5.4E+02) | Detected        | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | 90/90<br>100%<br>[95.9-100%]                 | 30/30<br>100%  | 29/30<br>96.7% | 30/30<br>100%  | <b>89/90</b><br><b>98.9%</b><br>[94.0-99.8%] |
|                          | falciparum  Detection  Results | Low Positive<br>1×LoD<br>(1.8E+02)      | Detected        | 30/30<br>100%   | 28/30<br>93.3% | 29/30<br>96.7% | <b>87/90</b><br><b>96.7%</b><br>[90.7-98.9%] | 30/30<br>100%  | 28/30<br>93.3% | 28/30<br>93.3% | <b>86/90</b><br><b>95.6%</b><br>[89.1-98.3%] |
|                          |                                | Negative<br>(No Analyte)                | Not<br>Detected | 30/30<br>100%   | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   | 30/30<br>100%  | 30/30<br>100%  | 30/30<br>100%  | <b>90/90</b><br><b>100%</b><br>[95.9-100%]   |

|   | Concentration      |                     |             |                                | % <i>F</i>         | Detection I<br>Agreement with     |                                  | •        |             |                                     |
|---|--------------------|---------------------|-------------|--------------------------------|--------------------|-----------------------------------|----------------------------------|----------|-------------|-------------------------------------|
| Analyte                                   | Tested             | Expected<br>Result  | Expected    | BioFire FilmArray 2.0 Platform |                    |                                   | BioFire FilmArray Torch Platform |          |             |                                     |
| (Source / ID)                             | (copies/mL)        |                     | System<br>1 | System 2                       | System 3           | All FA 2.0<br>Systems<br>[95% CI] | System<br>1                      | System 2 | System<br>3 | All FA Torch<br>Systems<br>[95% CI] |
| Overall Agreement with<br>Expected Result | All Concentrations | 1343/1350 1338/1350 |             |                                | 1343/1350<br>99.5% |                                   |                                  |          |             |                                     |

Abbreviations: FA – FilmArray; 95% CI – 95% Confidence Interval

#### Specimen Storage

Stability of whole blood specimens was evaluated to support labeling recommendations for storage of samples at room temperature for up to 24 hours or at 2–8°C for up to seven days. These conditions were selected to be consistent with, or exceed, standard storage and transport conditions for most laboratory testing of clinical human whole blood specimens. Evaluation of frozen samples at ultra-low temperatures prior to testing was performed to support other analytical and clinical testing.

Testing was conducted using contrived samples composed of human whole blood and four representative BioFire Global Fever Special Pathogens Panel analytes in two mixes at a concentration of 3× their respective LoD or less. Organisms were selected to be representative of the analytes detected by the BioFire Global Fever Special Pathogens Panel, including bacteria, viruses, and protozoa. A time point zero was tested immediately after sample preparation as a no-storage control. Ten replicates were tested at each of the storage conditions. Two different room temperature conditions (18-21°C and 30°C) were also tested. The results are summarized in Table 19.

Table 19. Summary of Analyte Detections Over Total Number Tested by Storage Condition

|                        |            | Room Te                 | mperature                         | Re    | I Ilbus Issu |       |                       |
|------------------------|------------|-------------------------|-----------------------------------|-------|--------------|-------|-----------------------|
| Analyte                | No Storage | Ambient                 | Upper Limit                       | 2-8°C |              |       | Ultra-low<br>Freezing |
| Analyte                | (Control)  | 18-21°C<br>for 24 Hours | 30°C<br>for 24 Hours <sup>1</sup> | Day 1 | Day 3        | Day 7 | ≤ -70°C               |
| Leptospira interrogans | 10/10      | 10/10                   |                                   | 10/10 | 10/10        | 10/10 | 10/10                 |
| Leptospira interrogans | 10/10      |                         | 10/10                             |       |              |       |                       |
| Dongue virus           | 10/10      | 10/10                   |                                   | 10/10 | 10/10        | 10/10 | 10/10                 |
| Dengue virus           | 10/10      |                         | 10/10                             |       |              |       |                       |
| Leishmania donovani    | 10/10      | 10/10                   |                                   | 10/10 | 10/10        | 10/10 | 10/10                 |
| Leisiimumu uonovum     | 10/10      |                         | 10/10                             |       |              |       |                       |
| Plasmodium falciparum  | 10/10      | 10/10                   |                                   | 10/10 | 10/10        | 10/10 | 10/10                 |
|                        | 10/10      |                         | 10/10                             |       |              | _     |                       |

<sup>&</sup>lt;sup>1</sup>Testing of the upper limit of room temperature storage, 30°C, was performed separately and required a separate no storage control.

### VIII. Assayed External Controls

Name of Device: BIOFIRE® SHIELD<sup>TM</sup> Control Kit for the BioFire Global Fever Special

Pathogens Panel

Common or Usual Name: Same

**Product Code: PMN** 

**Regulation:** 21 CFR 866.3920

Classification Name: Assayed quality control material for clinical microbiology assays

**Regulatory Class:** Class II (Special Controls)

**Panel:** Microbiology – 83

**Predicate Device:** BIOFIRE® SHIELD<sup>TM</sup> Control Kit for the Global Fever Panel (BioFire

Defense, LLC) [K202382]

This predicate has not been subject to a design-related recall.

#### **Device Description**

The BIOFIRE SHIELD Control Kit for the BioFire Global Fever Special Pathogens (GF SP) Panel is a surrogate control to monitor performance of the BioFire GF SP Panel assays. The BIOFIRE SHIELD Control Kit for the BioFire GF SP Panel is designed to mitigate the risk of control contamination or misuse when evaluating clinical specimens on BioFire FilmArray Systems. Good laboratory practice recommends running positive and negative external controls regularly. Evaluation of external controls is recommended prior to using a new shipment or new lot of BioFire GF SP Panel kits, when there is a new operator, and following replacement or repair of a BioFire FilmArray System. It is the responsibility of each laboratory to determine the frequency of external control testing with the BioFire GF SP Panel as part of the laboratory's Quality Control program. Quality control materials should be used in accordance with local, state, federal regulations and accreditation requirements.

# Materials provided in each BIOFIRE SHIELD Control Kit for the BioFire Global Fever Special Pathogens Panel:

- Six individually packaged Positive External Control Injection Vials
- Six individually packaged Negative External Control Injection Vials
- Instructions available online at www.biofiredefense.com:
  - BIOFIRE® SHIELD™ Control Kit for the BioFire Global Fever Special Pathogens Panel Instructions for Use
  - BIOFIRE® SHIELD™ Control Kit for the BioFire Global Fever Special Pathogens Panel – Quick Guide

# Materials required but not provided:

- BioFire® FilmArray® System including:
  - o BioFire® FilmArray® 2.0 or BioFire® FilmArray® Torch Instrument System including accompanying platform-specific core software
  - o BioFire® FilmArray® Pouch Loading Station

- BioFire® Global Fever Special Pathogens Panel (Part No. DFA2-ASY-0018) and accompanying pouch module software
- 10% bleach solution or a similar disinfectant

#### **Intended Use**

The BIOFIRE® SHIELD™ Control Kit for the BioFire® Global Fever Special Pathogens Panel contains Positive and Negative External Controls intended for use as assayed quality controls to monitor the performance of *in vitro* diagnostic laboratory nucleic acid testing procedures for the qualitative detection of *Bacillus anthracis*, chikungunya virus, Crimean-Congo hemorrhagic fever virus, dengue virus (serotypes 1, 2, 3, and 4), *Ebolavirus* spp., *Francisella tularensis*, Lassa virus, *Leishmania* spp., Leptospira spp., *Marburgvirus*, *Plasmodium* spp. (including species differentiation of *Plasmodium falciparum* and *Plasmodium vivax/ovale*), West Nile virus, yellow fever virus, and *Yersinia pestis* when using the BioFire® Global Fever Special Pathogens Panel on BioFire® FilmArray® 2.0 and BioFire® FilmArray® Torch Systems. The BIOFIRE SHIELD Control Kit for the BioFire Global Fever Special Pathogens Panel is designed for and intended to be used solely with the BioFire Global Fever Special Pathogens Panel. This product does not replace manufacturer internal controls provided as part of the BioFire Global Fever Special Pathogens Panel.

Both the Positive and Negative External Controls are provided in a FilmArray Control Injection Vial format. The Positive Control Injection Vial contains dried synthetic DNA segments in buffer and stabilizer to assess the presence of each individual assay on the BioFire Global Fever Special Pathogens Panel. The Negative Control Injection Vial contains no DNA and is non-reactive with the BioFire Global Fever Special Pathogens Panel assays.

For In Vitro Diagnostic Use.

# **Substantial Equivalence**

| Element             | Subject Device: BIOFIRE SHIELD Control Kit for the BioFire Global Fever Special Pathogens Panel  | Predicate: BIOFIRE SHIELD Control Kit for the BioFire Global Fever Panel [K202382]   |
|---------------------|--|--|
| Device Description  | Positive and negative external assayed quality controls to monitor assay performance in the BioFire Global Fever Special Pathogens Panel   |  |
| Physical Format     | External control material dried on Control Injection Vial filter   | Same as subject device   |
| Composition         | Tm-shifted synthetic DNA (positive control only)   | Same as subject device   |
| Targets Monitored   | Bacillus anthracis, chikungunya virus, Crimean-Congo hemorrhagic fever virus, dengue virus (serotypes 1, 2, 3, and 4), Ebolavirus spp., Francisella tularensis, Lassa virus, Leishmania spp., Leptospira spp., Marburgvirus, Plasmodium spp. (including species differentiation of P. falciparum and P. vivax/ovale), West Nile virus, yellow fever virus, and Yersinia pestis | Chikungunya virus, dengue virus (serotypes 1, 2, 3, and 4), Leptospira spp., Plasmodium spp. (including species differentiation of P. falciparum and P. vivax/ovale) |
| Instrumentation     | BioFire Global Fever Special<br>Pathogens Panel run on BioFire<br>FilmArray 2.0 or BioFire FilmArray<br>Torch systems  | BioFire Global Fever Panel run on<br>BioFire FilmArray 2.0   |
| Test Interpretation | Automated test interpretation and report generation; user cannot access raw data   | Same as subject device   |
| Reagent Storage     | Room temperature   | Same as subject device   |

# **Summary of Performance Data**

#### Reproducibility

Reproducibility for the BIOFIRE SHIELD Control Kit for the BioFire Global Fever Special Pathogens Panel was evaluated both for the BioFire FilmArray 2.0 System and the BioFire FilmArray Torch System. Reproducibility on the BioFire FilmArray 2.0 System was evaluated using SHIELD lots from three different manufacturing events at three test sites by two operators and three instruments per test site using three different BioFire GF SP Panel reagent lots. Testing took place over 5 days at each site. Reproducibility on the BioFire FilmArray Torch was performed in a similar manner with the exception that sites were simulated using three different BioFire FilmArray Torch Systems. Reproducibility was evaluated by calculating the percent agreement between observed test results (Passed or Failed) and expected test results (Passed) for both negative and positive SHIELD controls. Table 20 and Table 21 show summarized results for BioFire FilmArray 2.0 and BioFire FilmArray Torch Systems, respectively.

Table 20. Multi-Site Reproducibility Test Results of the BIOFIRE SHIELD Control Kit for the BioFire GF

SP Panel on BioFire FilmArray 2.0 Systems

| SHIELD Control Type   | Expected Result     |                  | Observed/Expected<br>(Percent Agreement)<br>[95% Confidence Interval] |                               |  |  |  |
|-----------------------|---------------------|------------------|---|-------------------------------|--|--|--|
|                       |                     | Site 1           | Site 2  | Site 3                        | All Sites  |  |  |
| Positive              | Passed <sup>1</sup> | 42/45<br>(93.3%) | 45/45<br>(100%)   | 45/45<br>(100%)               | <b>132/135</b><br><b>(97.8%)</b><br>[93.7-99.2%] |  |  |
| Negative              | Passed <sup>2</sup> | 45/45<br>(100%)  | 44/45 <sup>3</sup> (97.8%)  | 44/45 <sup>4</sup><br>(97.8%) | <b>133/135</b><br><b>(98.5%)</b><br>[94.8-99.6%] |  |  |
| Overall Agreement wit | 265/270             |                  |   |                               |  |  |  |

<sup>&</sup>lt;sup>1</sup> All BioFire GF SP Panel assays have a positive amplicon melt in the SHIELD control melt range.

<sup>&</sup>lt;sup>2</sup> All BioFire GF SP Panel assays have no melt in both the SHIELD control melt range and in the pathogen melt range.

<sup>&</sup>lt;sup>3</sup> Unexpected detection of pathogen amplicon for *Leishmania* spp.

<sup>&</sup>lt;sup>4</sup> Unexpected detection of pathogen amplicon for dengue virus.

Table 21. Reproducibility Test Results of the BIOFIRE SHIELD Control Kit for the BioFire GF SP Panel on BioFire FilmArray Torch Systems

|  |                     | Observed/Expected (Percent Agreement) |                      |   |  |  |  |
|--|---------------------|---------------------------------------|----------------------|---|--|--|--|
| SHIELD Control Type                    | Expected Result     | FA Torch<br>System 1                  | FA Torch<br>System 2 | FA Torch<br>System 3                            | <b>Overall</b><br>[95% Confidence<br>Interval]   |  |  |
| Positive                               | Passed <sup>1</sup> | 43/45<br>(95.6%)                      | 43/45<br>(95.6%)     | 44/45<br>(97.8%)                                | <b>130/135</b><br><b>(96.3%)</b><br>[91.6-98.4%] |  |  |
| Negative                               | Passed <sup>2</sup> | 43/45<br>(95.6%)                      | 45/45<br>(100%)      | 45/45<br>(100%)                                 | <b>133/135</b> ( <b>98.5%)</b> [94.8-99.6%]      |  |  |
| Overall Agreement with Expected Result |                     |                                       | ]                    | <b>263/270</b><br><b>(97.4%)</b><br>94.7-98.7%] |  |  |  |

<sup>&</sup>lt;sup>1</sup> All BioFire GF SP Panel assays have a positive amplicon melt in the positive SHIELD control melt range.

#### Repeatability

Repeatability was performed by a single operator testing 45 Positive and 45 Negative External Controls from a single BIOFIRE SHIELD Control Kit, using a single BioFire GF SP Panel reagent lot on one BioFire FilmArray 2.0 instrument over a period of 14 days. The primary assessment is based on a comparison of the observed External Control test results (Passed/Failed) to the expected test results (Passed). Table 22 shows the results for the Repeatability evaluation.

Table 22. Summary of BIOFIRE SHIELD Control Kit for the BioFire GF SP Panel Repeatability Test Results

| a to baile b        |                                  |                                       |  |  |  |  |  |  |
|---------------------|----------------------------------|---------------------------------------|--|--|--|--|--|--|
| SHIELD Control Type | Expected Result                  | Observed/Expected (Percent Agreement) |  |  |  |  |  |  |
| Positive            | Passed <sup>1</sup>              | 45/45<br>(100%)                       |  |  |  |  |  |  |
| Negative            | Passed <sup>2</sup> 45/45 (100%) |                                       |  |  |  |  |  |  |

<sup>&</sup>lt;sup>1</sup> All BioFire GF SP Panel assays have a positive amplicon melt in the positive SHIELD control melt range.

<sup>&</sup>lt;sup>2</sup> All BioFire GF SP Panel assays have no melt in either the positive SHIELD control melt range or the pathogen melt range.

<sup>&</sup>lt;sup>2</sup> All BioFire GF SP Panel assays have no melt in either the positive SHIELD control melt range or the pathogen melt range.

#### Clinical Evaluation

Six clinical sites evaluated the BIOFIRE SHIELD Control Kit by testing a Positive or Negative External Control each day prior to testing clinical specimens. Results for the BioFire GF SP Panel on BioFire FilmArray 2.0 Systems are shown in Table 23.

Table 23. Summary of BIOFIRE SHIELD Control Kit for the BioFire GF SP Panel Clinical Evaluation Results

| SHIELD Control Type | Completed with Passed<br>Result | Total Completed | Percent Passed (%) |
|---------------------|---------------------------------|-----------------|--------------------|
| Positive            | 158°                            | 160             | 98.8%              |
| Negative            | 157 ª                           | 159             | 98.7%              |
| Overall             | 315                             | 319             | 98.7%              |

<sup>&</sup>lt;sup>a</sup> Site tested a Positive and a Negative External Control on the same day. Controls were most likely swapped as the Negative External Control failed because all External Control targets were Detected, and the Positive External Control failed because all External Control targets were Not Detected.