



Becton, Dickinson and Company  
Joseph Basore  
Staff Regulatory Specialist  
7 Loveton Circle  
Sparks, Maryland 21152

July 28, 2022

Re: K214122

Trade/Device Name: BD MAX Enteric Bacterial Panel, BD MAX Extended Enteric Bacterial Panel  
Regulation Number: 21 CFR 866.3990  
Regulation Name: Gastrointestinal Microorganism Multiplex Nucleic Acid-Based Assay  
Regulatory Class: Class II  
Product Code: PCI, PCH, OOI  
Dated: December 22, 2021  
Received: December 30, 2021

Dear Joseph Basore:

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 <https://www.accessdata.fda.gov/scripts/cdrh/cfdocs/cfpmn/pmnmn.cfm> 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 Federal Register.

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 <https://www.fda.gov/combination-products/guidance-regulatory-information/postmarketing-safety-reporting-combination-products>); 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 <https://www.fda.gov/medical-devices/medical-device-safety/medical-device-reporting-mdr-how-report-medical-device-problems>.

For comprehensive regulatory information about medical devices and radiation-emitting products, including information about labeling regulations, please see Device Advice (<https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance>) and CDRH Learn (<https://www.fda.gov/training-and-continuing-education/cdrh-learn>). 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 (<https://www.fda.gov/medical-devices/device-advice-comprehensive-regulatory-assistance/contact-us-division-industry-and-consumer-education-dice>) for more information or contact DICE by email ([DICE@fda.hhs.gov](mailto:DICE@fda.hhs.gov)) or phone (1-800-638-2041 or 301-796-7100).

Sincerely,

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

## Indications for Use

510(k) Number (if known)  
K214122

Device Name  
BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel

Indications for Use (Describe)  
BD MAX™ Enteric Bacterial Panel

The BD MAX™ Enteric Bacterial Panel performed on the BD MAX™ System is an automated in vitro diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. The BD MAX™ Enteric Bacterial Panel detects nucleic acids from:

- *Salmonella* spp.
- *Campylobacter* spp. (*jejuni* and *coli*)
- *Shigella* spp. / Enteroinvasive *E. coli* (EIEC)
- Shiga toxin 1 (*stx1*) / Shiga toxin 2 (*stx2*) genes (found in Shiga toxin-producing *E. coli* [STEC]) as well as *Shigella dysenteriae*, which can possess a Shiga toxin gene (*stx*) that is identical to the *stx1* gene of STEC.

Testing is performed on unpreserved soft to diarrheal stool specimens or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of *SpaO*, a *Campylobacter* specific *tuf* gene sequence, *ipaH* and *stx1/stx2*. The test utilizes fluorogenic sequence-specific hybridization probes for detection of the amplified DNA.

This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of *Salmonella*, *Shigella*/EIEC, *Campylobacter* and Shiga toxin-producing *E. coli* (STEC) infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.

BD MAX™ Extended Enteric Bacterial Panel

The BD MAX™ Extended Enteric Bacterial Panel performed on the BD MAX™ System, is an automated in vitro diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. It is used in conjunction with the BD MAX™ Enteric Bacterial Panel as an optional Master Mix. The BD MAX™ Extended Enteric Bacterial Panel detects nucleic acids from:

- *Plesiomonas shigelloides*
- *Vibrio* (*V. vulnificus*, *V. parahaemolyticus*, and *V. cholerae*)
- Enterotoxigenic *Escherichia coli* (ETEC) heat-labile enterotoxin (LT)/ heat-stable enterotoxin (ST) genes
- *Yersinia enterocolitica*

Testing is performed on unpreserved soft to diarrheal or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of relevant gene target DNA. The test utilizes fluorogenic gene-specific hybridization probes for the detection of the amplified DNA.

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This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of *Plesiomonas shigelloides*, *Vibrio* (*V. vulnificus*, *V. parahaemolyticus*, and *V. cholerae*) Enterotoxigenic *Escherichia coli* (EPEC) LT/ST and *Yersinia enterocolitica* infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.

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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)

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**CONTINUE ON A SEPARATE PAGE IF NEEDED.**

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## **510(k) Summary**

BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel

### **Summary Preparation Date:**

12/22/2021

### **Submitted by:**

Becton, Dickinson and Company  
7 Loveton Circle  
Sparks, MD 21152

### **Contact:**

Joseph Basore, Ph.D., RAC  
Staff Regulatory Affairs Specialist  
Tel: 616-301-4068  
Email: Joseph.Basore@bd.com

### **Proprietary Names:**

*For the instrument:*

BD MAX™ System

*For the assay:*

BD MAX™ Enteric Bacterial Panel  
BD MAX™ Extended Enteric Bacterial Panel

### **Common Names:**

*For the instrument:*

Bench-top molecular diagnostics workstation

*For the assay:*

Gastrointestinal Bacterial Panel Multiplex Nucleic Acid-Based Assay System  
Enteric Bacterial Panel  
Enteric Bacterial Nucleic Acid Test  
Enteric Bacterial identification and differentiation system  
Enteric assay  
Enteric test

## **Regulatory Information**

*Regulation section:* 21 CFR 866.3990 – Gastrointestinal Bacterial Panel Multiplex Nucleic Acid-Based Assay System

*Classification:* Class II (Special Controls)

*Panel:* Microbiology (83)

*Product Code(s):*

PCI Gastrointestinal Bacterial Panel Multiplex Nucleic Acid-Based Assay System  
PCH Gastrointestinal Pathogen Panel Multiplex Nucleic Acid-Based Assay System  
OOI Real Time Nucleic Acid Amplification System

## **Predicate Device**

BD MAX™ Enteric Bacterial Panel (K140111)  
BD MAX™ Extended Enteric Bacterial Panel (K170308)

## **Device Establishment**

Becton, Dickinson and Company  
7 Loveton Circle  
Sparks, MD 21152  
Registration Number: 1119779

## **Performance Standards**

Class II Special Controls Guideline: Gastrointestinal Microorganism Multiplex Nucleic Acid-Based Assays for Detection and Identification of Microorganisms and Toxin Genes from Human Stool Specimens, November 2, 2015.

## **Intended Use**

### **BD MAX™ Enteric Bacterial Panel**

The BD MAX™ Enteric Bacterial Panel performed on the BD MAX™ System is an automated *in vitro* diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. The BD MAX™ Enteric Bacterial Panel detects nucleic acids from:

- *Salmonella* spp.
- *Campylobacter* spp. (*jejuni* and *coli*)
- *Shigella* spp. / Enteroinvasive *E. coli* (EIEC)
- Shiga toxin 1 (*stx1*) / Shiga toxin 2 (*stx2*) genes (found in Shiga toxin-producing *E. coli* [STEC]) as well as *Shigella dysenteriae*, which can possess a Shiga toxin gene (*stx*) that is identical to the *stx1* gene of STEC.

Testing is performed on unpreserved soft to diarrheal stool specimens or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of *SpaO*, a *Campylobacter* specific *tuf* gene sequence, *ipaH* and *stx1/stx2*. The test utilizes fluorogenic sequence-specific hybridization probes for detection of the amplified DNA.

This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of *Salmonella*, *Shigella*/EIEC, *Campylobacter* and Shiga toxin-producing *E. coli* (STEC) infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.

### **BD MAX™ Extended Enteric Bacterial Panel**

The BD MAX™ Extended Enteric Bacterial Panel performed on the BD MAX™ System, is an automated in vitro diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. It is used in conjunction with the BD MAX™ Enteric Bacterial Panel as an optional Master Mix. The BD MAX™ Extended Enteric Bacterial Panel detects nucleic acids from

- *Plesiomonas shigelloides*
- *Vibrio* (*V. vulnificus*, *V. parahaemolyticus*, and *V. cholerae*)
- Enterotoxigenic *Escherichia coli* (ETEC) heat-labile enterotoxin (LT)/ heat-stable enterotoxin (ST) genes
- *Yersinia enterocolitica*

Testing is performed on unpreserved soft to diarrheal or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of relevant gene target DNA. The test utilizes fluorogenic gene-specific hybridization probes for the detection of the amplified DNA.

This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of *Plesiomonas shigelloides*, *Vibrio* (*V. vulnificus*, *V. parahaemolyticus*, and *V. cholerae*) Enterotoxigenic *Escherichia coli* (ETEC) LT/ST and *Yersinia enterocolitica* infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this

test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.

**Special Conditions for Use Statement:** For Prescription Use Only

**Special Instrument Requirements:** BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel are performed on the BD MAX™ System

### **Device Description**

The BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays along with the BD MAX System are comprised of an instrument with associated hardware and accessories, disposable microfluidic cartridges, master mixes, unitized reagent strips, and extraction reagents. The instrument automates sample preparation including target lysis, DNA extraction and concentration, reagent rehydration, target nucleic acid amplification and detection using real-time PCR. The assay includes a Sample Processing Control (SPC) that is present in the Extraction Tube. The SPC monitors DNA extraction steps, thermal cycling steps, reagent integrity and the presence of inhibitory substances. The BD MAX™ System software automatically interprets test results. For the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel, a test result may be called as POS, NEG or UNR (Unresolved) based on the amplification status of the targets and of the Sample Processing Control. IND (Indeterminate) or INC (Incomplete) results are due to BD MAX™ System failure.

### **Test Principle**

The BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays are designed for use with unpreserved or Cary-Blair preserved stool samples. Unpreserved samples are placed in a BD MAX sample buffer tube (SBT) with a 10 µL transfer loop for analysis on the BD Max System. The current Cary-Blair preserved specimen claim utilizes a plastic paddle (scoop) to place a stool sample into 15 ml of Cary-Blair media for transport before being placed into a SBT with a 10 µL transfer loop prior to analysis on the BD Max System.

To use the FecalSwab Collection, Transport, and Preservation System, the operator transfers fecal material from an unpreserved stool specimen to the vial of FecalSwab transport medium using the nylon flocked specimen collection swab. The FecalSwab transport medium tube is filled with 2 ml of a semi-solid modified Cary-Blair medium that is designed to maintain the viability of enteric pathogenic bacteria during transit to the testing laboratory. Last, before analysis on the BD MAX system, samples collected/stored with the FecalSwab system are vortexed and then pipetted (50 µl) into a BD MAX™ sample buffer tube (SBT).

Once specimens (Unpreserved, Cary-Blair, or FecalSwab Cary-Blair) are placed into a BD MAX SBT, the test principles are as described in K140111 and K170308. For all specimen types the SBTs are vortexed and then loaded into the BD MAX system along with the Unitized Reagent Strips, Master Mix, Extraction Tubes, and PCR Cartridges. No further operator intervention is necessary, and the following automated procedures occur. The microbial cells are lysed and DNA



is extracted using a combination of lytic and extraction reagents at elevated temperatures. Nucleic acids released from the target organisms are captured on magnetic affinity beads. The beads, together with the bound nucleic acids, are washed and the nucleic acids are eluted by a combination of heat and pH. Eluted DNA is neutralized and transferred to the Master Mix Tube to rehydrate the PCR reagents. After reconstitution, the BD MAX System dispenses a fixed volume of PCR-ready solution containing the extracted nucleic acids into the PCR Cartridge. Microvalves in the cartridge are sealed by the system prior to initiating PCR in order to contain the amplification mixture and thus prevent evaporation and contamination.

The amplified DNA targets are detected using hydrolysis (TaqMan®) probes, labeled at one end with a fluorescent reporter dye (fluorophore), and at the other end, with a quencher moiety. Probes labeled with different fluorophores are used to detect the target analytes in different optical channels of the BD MAX System. The probes are used to detect amplicons for enteric bacterial targets and the Sample Processing Control in five different optical channels of the BD MAX System. When the probes are in their native state, the fluorescence of the fluorophore is quenched due to its proximity to the quencher. However, in the presence of target DNA, the probes hybridize to their complementary sequences and are hydrolyzed by the 5'-3' exonuclease activity of the DNA polymerase as it synthesizes the nascent strand along the DNA template. As a result, the fluorophores are separated from the quencher molecules and fluorescence is emitted. The amount of fluorescence detected in the optical channels used for the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays are directly proportional to the quantity of the corresponding probe that is hydrolyzed. The BD MAX System monitors these signals at each cycle of the PCR and interprets the data at the end of the reaction to provide qualitative test results for each analyte (i.e., positive or negative). The assay includes a Sample Processing Control, which monitors the integrity of the reagents as well as the process steps involved in DNA extraction, amplification and detection, and checks for the presence of potential assay inhibitor.

### **Substantial Equivalence<sup>1</sup>**

Table 1 and Table 2 provides the similarities and differences between the submitted device and the legally marketed predicate device.

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<sup>1</sup> The term “substantial equivalence” as used in this 510(k) notification is limited to the definition of substantial equivalence as found in the Federal Food, Drug and Cosmetic Act, as amended and as applied under 21 CFR 807, Subpart E under which a device can be marketed without pre-market approval or reclassification. A determination of substantial equivalency under this notification is not intended to have any bearing whatsoever on the resolution of patent infringement suits or any other patent matters. No statements related to, or in support of substantial equivalence herein shall be construed as an admission against interest under the US Patent Laws or their application by the courts.

**Table 1: Comparison of BD MAX™ Enteric Bacterial Panel to Predicate Device**

| Item         | Predicate - BD MAX™ Enteric Bacterial Panel (K140111)  | Submitted Device - BD MAX™ Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System |
|--------------|--|--|
| Intended Use | <p>The BD MAX™ Enteric Bacterial Panel performed on the BD MAX™ System is an automated <i>in vitro</i> diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. The BD MAX™ Enteric Bacterial Panel detects nucleic acids from:</p> <ul style="list-style-type: none"> <li>• <i>Salmonella</i> spp.</li> <li>• <i>Campylobacter</i> spp. (<i>jejuni</i> and <i>coli</i>)</li> <li>• <i>Shigella</i> spp. / Enteroinvasive <i>E. coli</i> (EIEC)</li> <li>• Shiga toxin 1 (<i>stx1</i>) / Shiga toxin 2 (<i>stx2</i>) genes (found in Shiga toxin-producing <i>E. coli</i> [STEC]) as well as <i>Shigella dysenteriae</i>, which can possess a Shiga toxin gene (<i>stx</i>) that is identical to the <i>stx1</i> gene of STEC.</li> </ul> <p>Testing is performed on unpreserved soft to diarrheal stool specimens or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of <i>SpaO</i>, a <i>Campylobacter</i> specific <i>tuf</i> gene sequence, <i>ipaH</i> and <i>stx1/stx2</i>. The test utilizes fluorogenic sequence-specific hybridization probes for detection of the amplified DNA.</p> <p>This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of <i>Salmonella</i>, <i>Shigella</i>/EIEC, <i>Campylobacter</i> and Shiga toxin-producing <i>E. coli</i> (STEC) infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test, and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn’s disease.</p> | Same   |

| Item                           | Predicate - BD MAX™ Enteric Bacterial Panel (K140111)   | Submitted Device - BD MAX™ Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System  |
|--------------------------------|---|---|
| Organisms Detected             | <ul style="list-style-type: none"> <li>• <i>Salmonella</i> spp.</li> <li>• <i>Campylobacter</i> spp. (<i>jejuni</i> and <i>coli</i>)</li> <li>• <i>Shigella</i> spp. / Enteroinvasive <i>E. coli</i> (EIEC)</li> <li>• Shiga toxin 1 (<i>stx1</i>) / Shiga toxin 2 (<i>stx2</i>) genes (found in Shiga toxin-producing <i>E. coli</i> [STEC]) as well as <i>Shigella dysenteriae</i>, which can possess a Shiga toxin gene (<i>stx</i>) that is identical to the <i>stx1</i> gene of STEC.</li> </ul> | Same  |
| Specimen Type                  | Unpreserved stool or Cary-Blair preserved stool   | Same  |
| Assay Format                   | Amplification: PCR<br>Detection: Fluorogenic target-specific hybridization  | Same  |
| Mode of Detection              | Presence of <ul style="list-style-type: none"> <li>• <i>tuf</i> gene specific for <i>Campylobacter</i></li> <li>• <i>SpaO</i> gene specific for <i>Salmonella</i></li> <li>• <i>ipaH</i> gene specific for <i>Shigella</i></li> <li>• <i>stx1a</i> and <i>stx2a</i> genes specific to Shiga-toxin producing organisms</li> </ul>  | Same  |
| Interpretation of Test Results | Automated (BD MAX™ System diagnostic software)  | Same  |
| Analysis Platform              | BD MAX™ System  | Same  |
| PCR Sample Preparation         | Automated by the BD MAX™ System   | Same  |
| Detection Probes               | TaqMan® Probe   | Same  |
| Assay Controls                 | Sample Processing Control (SPC)   | Same  |
| Cary-Blair Buffer Formulation  | <ul style="list-style-type: none"> <li>-Sodium Chloride</li> <li>-Calcium Chloride</li> <li>-Phosphate Buffer</li> <li>-Thioglycolic Acid Sodium Salt</li> <li>-Phenol Red</li> <li>-Agar</li> <li>-Water</li> </ul>  | <ul style="list-style-type: none"> <li>-Chloride salts</li> <li>-Sodium salts</li> <li>-Phosphate buffer</li> <li>-L-Cysteine</li> <li>-Agar</li> <li>-Water</li> </ul> |
| Cary-Blair Buffer Container    | Plastic Container w/Lid prefilled 15 ml of media  | Plastic Container w/Lid prefilled 2 ml of media   |

| <b>Item</b>  | <b>Predicate - BD MAX™ Enteric Bacterial Panel (K140111)</b> | <b>Submitted Device - BD MAX™ Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System</b> |
|--|--|---|
| Specimen Transfer Tool (unpreserved to Cary-Blair) | Plastic Paddle   | Flocked Swab  |
| Transport Method to SBT Tube                       | 10 µL Transport Loop   | 50 µL Pipette   |
| Sterility of FecalSwab                             | Not Applicable   | Yes, Irradiation (FecalSwab)  |

**Table 2: Comparison of BD MAX™ Extended Enteric Bacterial Panel to Predicate Device**

| Item         | Predicate - BD MAX Extended Enteric Bacterial Panel (K170308)   | Proposed - BD MAX Extended Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System |
|--------------|---|--|
| Intended Use | <p>The BD MAX™ Extended Enteric Bacterial Panel performed on the BD MAX™ System, is an automated <i>in vitro</i> diagnostic test for the direct qualitative detection and differentiation of enteric bacterial pathogens. It is used in conjunction with the BD MAX™ Enteric Bacterial Panel as an optional Master Mix. The BD MAX™ Extended Enteric Bacterial Panel detects nucleic acids from</p> <ul style="list-style-type: none"> <li>• <i>Plesiomonas shigelloides</i></li> <li>• <i>Vibrio</i> (<i>V. vulnificus</i>, <i>V. parahaemolyticus</i>, and <i>V. cholerae</i>)</li> <li>• Enterotoxigenic <i>Escherichia coli</i> (ETEC) heat-labile enterotoxin (LT)/ heat-stable enterotoxin (ST) genes</li> <li>• <i>Yersinia enterocolitica</i></li> </ul> <p>Testing is performed on unpreserved soft to diarrheal or Cary-Blair preserved stool specimens from symptomatic patients with suspected acute gastroenteritis, enteritis or colitis. The test is performed directly on the specimen, utilizing real-time polymerase chain reaction (PCR) for the amplification of relevant gene target DNA. The test utilizes fluorogenic gene-specific hybridization probes for the detection of the amplified DNA.</p> <p>This test is intended for use, in conjunction with clinical presentation, laboratory findings, and epidemiological information, as an aid in the differential diagnosis of <i>Plesiomonas shigelloides</i>, <i>Vibrio</i> (<i>V. vulnificus</i>, <i>V. parahaemolyticus</i>, and <i>V. cholerae</i>) Enterotoxigenic <i>Escherichia coli</i> (ETEC) LT/ST and <i>Yersinia enterocolitica</i> infections. Results of this test should not be used as the sole basis for diagnosis, treatment, or other patient management decisions. Positive results do not rule out co-infection with other organisms that are not detected by this test, and may not be the sole or definitive cause of patient illness. Negative results in the setting of clinical illness compatible with gastroenteritis may be due to infection</p> | Same   |

| Item                           | Predicate - BD MAX Extended Enteric Bacterial Panel (K170308)   | Proposed - BD MAX Extended Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System |
|--------------------------------|---|--|
|                                | by pathogens that are not detected by this test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.  |  |
| Organisms Detected             | <ul style="list-style-type: none"> <li>• <i>Plesiomonas shigelloides</i></li> <li>• <i>Vibrio</i> (<i>V. vulnificus</i>, <i>V. parahaemolyticus</i>, and <i>V. cholerae</i>)</li> <li>• Enterotoxigenic <i>Escherichia coli</i> (ETEC) heat-labile enterotoxin (LT)/ heat-stable enterotoxin (ST) genes</li> <li>• <i>Yersinia enterocolitica</i></li> </ul>                      | Same   |
| Specimen Type                  | Unpreserved stool or Cary-Blair preserved stool   | Same   |
| Assay Format                   | Amplification: PCR<br>Detection: Fluorogenic target-specific hybridization  | Same   |
| Mode of Detection              | Presence of <ul style="list-style-type: none"> <li>• Undefined gene suspected to be implicated in Fe<sup>3+</sup> transport for <i>Plesiomonas shigelloides</i></li> <li>• <i>atpA</i> gene specific for <i>Vibrio</i></li> <li>• <i>eltA</i> gene specific for <i>Enterotoxigenic Escherichia coli</i></li> <li>• <i>invA</i> gene for <i>Yersinia enterocolitica</i></li> </ul> | Same   |
| Interpretation of Test Results | Automated (BD MAX™ System diagnostic software)  | Same   |
| Analysis Platform              | BD MAX™ System  | Same   |
| PCR Sample Preparation         | Automated by the BD MAX™ System   | Same   |
| Detection Probes               | TaqMan® Probe   | Same   |
| Assay Controls                 | Sample Processing Control (SPC)   | Same   |
| Cary-Blair Buffer Formulation  | Sodium Chloride<br>Calcium Chloride<br>Phosphate Buffer<br>Thioglycolic Acid Sodium Salt<br>Phenol Red<br>Agar<br>Water   | -Chloride salts<br>-Sodium salts<br>-Phosphate buffer<br>-L-Cysteine<br>-Agar<br>-Water                          |

| <b>Item</b>                  | <b>Predicate - BD MAX Extended Enteric Bacterial Panel (K170308)</b> | <b>Proposed - BD MAX Extended Enteric Bacterial Panel with FecalSwab Collection, Preservation, and Transport System</b> |
|------------------------------|--|---|
| Cary-Blair Buffer Container  | Plastic Container w/Lid prefilled 15 ml of media                     | Plastic Container w/Lid prefilled 2 ml of media   |
| Transfer Tool                | Plastic Paddle   | Nylon Flocked Swab  |
| Transport Method to SBT Tube | Transport Loop   | Pipette   |
| Sterility of FecalSwab       | Not Applicable   | Yes, Irradiation (FecalSwab)  |

## **Performance Evaluation**

Four studies were conducted to demonstrate the substantial equivalence between the current predicate specimen collection (Cary-Blair) and the additional specimen collection (FecalSwab) for use in the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays:

- Confirmation of equivalent analytical sensitivity with the Copan FecalSwab™ preserved stool specimen (FecalSwab) compared to the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel from a Cary-Blair Para-Pak® stool sample was performed by the limiting dilution LoD model. Acceptable performance was demonstrated when the detection break points between the FecalSwab and Cary-Blair Para-Pak® specimen types were within one five-fold dilution of each other. Break point is defined as the highest concentration where the positivity rate is <95% (<23/24). To achieve this comparison, a negative stool pool was prepared and divided into five aliquots, to which serially diluted multiplex organism mix was added. The organism mix contained a representative strain for each of the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel targets. The organism mixes contain one species from all claimed genera.

For the BD MAX™ Enteric Bacterial Panel the organism mix was prepared by combining the following targets in PBS: *Salmonella typhimurium* (ATCC 14028), *Shigella sonnei* (ATCC 9290), *Campylobacter jejuni* (ATCC 43429), and *Escherichia coli stx1* (ATCC 43890). For the BD MAX™ Extended Enteric Bacterial Panel the organism mix was prepared by combining the following targets in PBS: *Plesiomonas shigelloides* (ATCC 14029), *V. parahaemolyticus* (ATCC 178020), *Y. enterocolitica* (ATCC 9610), and *Escherichia coli* ETEC (ATCC 35401). The diluted BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel mixtures were spiked into aliquots of the stool pool. Three separate lots of FecalSwabs and one lot of scaled volume Cary-Blair were inoculated with each of the serially diluted stool samples. A 0.5 McFarland turbidity suspension was prepared for each organism. This suspension was diluted in PBS to create concentration 1 (Conc 1). Five additional concentrations (Conc 2-6) were prepared by performing 5-fold serial dilutions from Conc 1.

Limiting dilutions of specimens for each organism prepared using the FecalSwab™ and Para-Pak® exhibited drop-out rates at similar levels when tested with the BD MAX™ Enteric Bacterial Panel and the BD MAX™ Extended Enteric Bacterial Panel Assays on the BD MAX™ System. All FecalSwab™ break points were within one five-fold concentration when compared to Para-Pak® (Table 3). There was no indication that the FecalSwab™ collection device negatively impacted the analytical sensitivity of the BD MAX™ Enteric Bacterial Panel or BD MAX™ Extended Enteric Bacterial Panel.



**Table 3: Number of Positive Samples for the BD MAX™ Enteric Bacterial Panel**

| Enteric Bacterial Panel          |                        |           |                          |           |                            |           |                     |           |
|----------------------------------|------------------------|-----------|--------------------------|-----------|----------------------------|-----------|---------------------|-----------|
| Organism                         | <i>S. typhimurium</i>  |           | <i>E. coli stx1</i>      |           | <i>C. jejuni</i>           |           | <i>S. sonnei</i>    |           |
| Collection Type                  | FecalSwab™             | Para-Pak® | FecalSwab™               | Para-Pak® | FecalSwab™                 | Para-Pak® | FecalSwab™          | Para-Pak® |
| Conc 1                           | 24/24                  | 24/24     | 24/24                    | 24/24     | 24/24                      | 24/24     | 24/24               | 24/24     |
| Conc 2                           | 24/24                  | 24/24     | 24/24                    | 24/24     | 24/24                      | 24/24     | 24/24               | 24/24     |
| Conc 3                           | 22/24                  | 16/24     | 24/24                    | 23/24     | 24/24                      | 24/24     | 24/24               | 23/24     |
| Conc 4                           | 8/24                   | 6/24      | 16/24                    | 9/24      | 23/24                      | 22/24     | 20/24               | 21/24     |
| Conc 5                           | 3/24                   | 7/24      | 3/24                     | 6/24      | 14/24                      | 13/24     | 13/24               | 15/24     |
| Conc 6                           | 0/24                   | 0/24      | 0/24                     | 0/24      | 3/24                       | 3/24      | 3/24                | 4/24      |
| Extended Enteric Bacterial Panel |                        |           |                          |           |                            |           |                     |           |
| Organism                         | <i>P. shigelloides</i> |           | <i>Y. enterocolitica</i> |           | <i>V. parahaemolyticus</i> |           | <i>E. coli ETEC</i> |           |
| Collection Type                  | FecalSwab™             | Para-Pak® | FecalSwab™               | Para-Pak® | FecalSwab™                 | Para-Pak® | FecalSwab™          | Para-Pak® |
| Conc 1                           | 24/24                  | 23/23*    | 24/24                    | 23/23*    | 24/24                      | 23/23*    | 24/24               | 23/23*    |
| Conc 2                           | 24/24                  | 24/24     | 24/24                    | 24/24     | 24/24                      | 24/24     | 24/24               | 24/24     |
| Conc 3                           | 24/24                  | 17/24     | 24/24                    | 17/24     | 24/24                      | 18/24     | 24/24               | 18/24     |
| Conc 4                           | 14/24                  | 9/24      | 10/24                    | 10/24     | 7/24                       | 3/24      | 13/24               | 7/24      |
| Conc 5                           | 3/24                   | 6/24      | 2/24                     | 1/24      | 1/24                       | 0/24      | 6/24                | 4/24      |
| Conc 6                           | 0/24                   | 0/24      | 1/24                     | 2/24      | 0/24                       | 0/24      | 1/24                | 1/24      |

\*One non-reportable sample was not retested, and 23 replicates were accepted for the highest concentration.

- Specimen Stability of stool specimen collected with the FecalSwab was tested against all target organisms. The results showed that specimen stability of FecalSwab meets the current BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assay stability claims. For each organism tested across both the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays, a detection  $\geq 95\%$  occurred at all the target stability time points claimed in the package insert. Therefore, stool preserved with FecalSwab can be stored for 24 hours (1 days) at  $25 \pm 2$  °C and 120 hours (5 days) at 2 - 8 °C, and sample buffer tube inoculated with FecalSwab specimen can be stored for 48 hours (2 days) at  $25 \pm 2$  °C and 120 hours (5 days) at 2 - 8 °C.
- A user variability study was performed using the FecalSwab since there are differences in workflow (unpreserved sample to preservation media to SBT) between the FecalSwab and Cary-Blair Para-Pak® specimen collection. The data demonstrate that expected assay results are obtained when FecalSwab stool specimens were prepared by multiple users and shows that the difference in workflow between Cary-Blair Para-Pak® and FecalSwab specimen collection has no effect on the ability of the user to place the sample into the SBT for the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays.

The user variability study was performed to confirm that the preparation of the FecalSwab™ by different users does not induce variability in the expected results for the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel assays. Six (6) different users prepared two (2) different FecalSwab™ specimens from each of the five (5) panel members provided (one (1) negative panel member, three (3) low-positive panel members, and one (1)

moderate-positive panel member;). Once the FecalSwab™ specimens were prepared by various users, all subsequent steps, including the transfer to SBTs from each FecalSwab™, were performed by a single experienced BD MAX™ user. *Campylobacter jejuni* was used for BD MAX™ Enteric Bacterial Panel assay because it is the most prevalent Enteric Bacterial Panel target and has the lowest LoD from Enteric Bacterial Panel targets. ETEC ST/LT was used for BD MAX™ Extended Enteric Bacterial Panel assay because it is the most prevalent Extended Enteric Bacterial target and has the second lowest LoD from Extended Enteric Bacterial targets.

Acceptance criteria were: 100% negative results for the twelve (12) negative samples, ≥95% positive results for the thirty-six (36) low-positive samples, and 100% positive for the twelve (12) moderate-positive samples. All conditions met acceptance criteria (Table 4)

**Table 4. Acceptance criteria for user variability**

| Target                               | Panel Member      | Acceptance Criteria | Assay Results | Results |
|--------------------------------------|-------------------|---------------------|---------------|---------|
| <i>Campylobacter jejuni</i> and ETEC | Moderate-Positive | 100% POS            | 100% POS      | Pass    |
|                                      | Low-Positive      | ≥95% POS            | 100% POS      |         |
| Negative samples                     | Negative          | 100% NEG            | 100% NEG      |         |

Results met all acceptance criteria. The data demonstrate that expected assay results are obtained when the FecalSwab™ fecal specimens were prepared by multiple users.

- The performance of the BD FecalSwab™ Collection, Transport and Preservation System when tested with the BD MAX™ Enteric Bacterial Panel and BD MAX™ Extended Enteric Bacterial Panel was evaluated in a comparison study by comparing the results obtained for specimens using Cary-Blair Para-Pak® preserved stool samples to those using the BD FecalSwab™ Collection, Transport and Preservation System. Both the BD FecalSwab™ and Copan FecalSwab™ are identical other than branding and were incorporated into the performance evaluation. Unpreserved stool samples were collected from pediatric and adult patients suspected of acute bacterial gastroenteritis, enteritis, or colitis from eight (8) geographically diverse clinical centers where specimens were collected as part of routine patient care. At these locations, the fresh, unpreserved stool samples were transferred into both Cary-Blair Para-Pak® collection vials and BD FecalSwab™ devices. All samples were subsequently shipped to a centralized testing laboratory and tested with the BD MAX™ Enteric Bacterial panel. A total

of 621 prospective specimens and 295 retrospective specimens were enrolled in the clinical evaluation where three (3) prospective samples were excluded from the data analysis due to subject exclusion criteria. Table 5 describes the 913 (618 prospective and 295 retrospective) compliant specimens enrolled by patient age, sex, and specimen type. Sixteen (16) additional prospective samples were excluded from the data analysis due to specimen or instrument level exclusion criteria. The final data analysis included 897 compliant subjects for *Campylobacter*, *Salmonella*, *Shigella* spp. / Enteroinvasive *E. coli* (EIEC), Shiga toxin producing *E. coli* (STEC), *Plesiomonas shigelloides*, *Vibrio* (*V. vulnificus*, *V. parahaemolyticus*, *V. cholerae*), Enterotoxigenic *E. coli* (ETEC) LT/ST, and *Yersinia enterocolitica* targets.

**Table 5: Compliant Clinical Trial Enrollment Summary by Age, Sex, and Specimen Type**

| Specimen Type                  | Mean Age in years (SD) | Median Age in years | Min Age in years | Max Age in years | Sex of Total N |
|--------------------------------|------------------------|---------------------|------------------|------------------|----------------|
| Prospective<br>Total N = 618   | 47.1 (22.4)            | 49.0                | <1               | 95               | Male: 44.8%    |
| Unknown Age: 0                 |                        |                     |                  |                  | Female: 55.2%  |
| Known Age: 618                 |                        |                     |                  |                  | Unknown: 0.0%  |
| Retrospective<br>Total N = 295 | 37.2 (20.8)            | 33.5                | <1               | 86               | Male: 30.5%    |
| Unknown Age: 149               |                        |                     |                  |                  | Female: 24.1%  |
| Known Age: 146                 |                        |                     |                  |                  | Unknown: 45.4% |
| Overall<br>Total N = 913       | 45.2 (22.4)            | 47.0                | <1               | 95               | Male: 40.2%    |
| Unknown Age: 149               |                        |                     |                  |                  | Female: 45.1%  |
| Known Age: 764                 |                        |                     |                  |                  | Unknown: 14.7% |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% and 99.8% of the *Campylobacter* spp. prospective positive and negative specimens, respectively, and 100.0% and 97.1% of the retrospective positive and negative specimens, respectively (Table 6).

**Table 6: *Campylobacter* spp. PPA and NPA of the BD MAX™ Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Campylobacter</i> spp.                                 |           | Cary-Blair |          | Total |
|---|-----------|------------|----------|-------|
| Specimen Origin   | FecalSwab | Positive   | Negative |       |
| Prospective   | Positive  | 9          | 1        | 10    |
|   | Negative  | 0          | 585      | 585   |
|   | Total     | 9          | 586      | 595   |
| PPA: 100.0% (70.1%, 100.0%)<br>NPA: 99.8% (99.0%, 100.0%) |           |            |          |       |
| Retrospective   | Positive  | 88         | 6        | 94    |
|   | Negative  | 0          | 200      | 200   |
|   | Total     | 88         | 206      | 294   |
| PPA: 100.0% (95.8%, 100.0%)<br>NPA: 97.1% (93.8%, 98.7%)  |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% of the *Salmonella* spp. prospective positive and negative specimens, and 93.3% and 95.9% of the retrospective positive and negative specimens, respectively (Table 7).

**Table 7: *Salmonella* spp. PPA and NPA of the BD MAX™ Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Salmonella</i> spp.                                     |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 4          | 0        | 4     |
|  | Negative  | 0          | 591      | 591   |
|  | Total     | 4          | 591      | 595   |
| PPA: 100.0% (51.0%, 100.0%)<br>NPA: 100.0% (99.4%, 100.0%) |           |            |          |       |
| Retrospective  | Positive  | 70         | 9        | 79    |
|  | Negative  | 5          | 210      | 215   |
|  | Total     | 75         | 219      | 294   |
| PPA: 93.3% (85.3%, 97.1%)<br>NPA: 95.9% (92.4%, 97.8%)     |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100% of the *Shigella* spp. prospective positive and negative specimens, and 98.1% and 99.6% of the retrospective positive and negative specimens, respectively (Table 8).

**Table 8: *Shigella* spp. PPA and NPA of the BD MAX™ Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Shigella</i> spp.                                       |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 7          | 0        | 7     |
|  | Negative  | 0          | 588      | 588   |
|  | Total     | 7          | 588      | 595   |
| PPA: 100.0% (64.6%, 100.0%)<br>NPA: 100.0% (99.4%, 100.0%) |           |            |          |       |
| Retrospective  | Positive  | 52         | 1        | 53    |
|  | Negative  | 1          | 240      | 241   |
|  | Total     | 53         | 241      | 294   |
| PPA: 98.1% (90.1%, 99.7%)<br>NPA: 99.6% (97.7%, 99.9%)     |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% and 99.5% of the *stx1/stx2* (STX) prospective positive and negative specimens, respectively, and 92.9% and 100.0% of the retrospective positive and negative specimens, respectively (Table 9).

**Table 9: *stx1/stx2* (STX) PPA and NPA of the BD MAX™ Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| STX  |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 1          | 3        | 4     |
|  | Negative  | 0          | 591      | 591   |
|  | Total     | 1          | 594      | 595   |
| PPA: 100.0% (20.7%, 100.0%)<br>NPA: 99.5% (98.5%, 99.8%) |           |            |          |       |
| Retrospective  | Positive  | 13         | 0        | 13    |
|  | Negative  | 1          | 281      | 282   |
|  | Total     | 14         | 281      | 295   |
| PPA: 92.9% (68.5%, 98.7%)<br>NPA: 100.0% (98.7%, 100.0%) |           |            |          |       |

In addition, due to the small number of *stx1/stx2* (STX) positive specimens in the study, contrived specimens were evaluated. The BD FecalSwab™ Collection, Transport and Preservation System on the BD MAX™ Enteric Bacterial Panel identified 100% of the *stx1/stx2* (STX) contrived positive and negative specimens, when compared to expected results (Table 10).

**Table 10: STX Contrived FecalSwab™ Specimen Results**

| STX  | Expected Results |          |    |     |
|--|------------------|----------|----|-----|
|  | Contrived        | Positive | 53 | 0   |
| Negative   |                  | 0        | 53 | 53  |
| Total  |                  | 53       | 53 | 106 |
| PPA: 100.0% (93.2%, 100.0%)<br>NPA: 100.0% (93.2%, 100.0%) |                  |          |    |     |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% and 99.0% of the *Plesiomonas shigelloides* prospective positive and negative specimens, respectively, and 33.3% and 100.0% of the retrospective positive and negative specimens, respectively (Table 11).

**Table 11: *Plesiomonas shigelloides* PPA and NPA of the BD MAX™ Extended Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Plesiomonas shigelloides</i>                          |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 2          | 6        | 8     |
|  | Negative  | 0          | 586      | 586   |
|  | Total     | 2          | 592      | 594   |
| PPA: 100.0% (34.2%, 100.0%)<br>NPA: 99.0% (97.8%, 99.5%) |           |            |          |       |
| Retrospective  | Positive  | 1          | 0        | 1     |
|  | Negative  | 2          | 291      | 293   |
|  | Total     | 3          | 291      | 294   |
| PPA: 33.3% (6.1%, 79.2%)<br>NPA: 100.0% (98.7%, 100.0%)  |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 99.7% of the *Vibrio* spp. prospective negative specimens (zero prospective positive specimens were analyzed), and 100.0% of the retrospective positive and negative specimens (Table 12).

**Table 12: *Vibrio* spp. PPA and NPA of the BD MAX™ Extended Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Vibrio</i>  |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 0          | 2        | 2     |
|  | Negative  | 0          | 592      | 592   |
|  | Total     | 0          | 594      | 594   |
| PPA: Not Available<br>NPA: 99.7% (98.8%, 99.9%)            |           |            |          |       |
| Retrospective  | Positive  | 4          | 0        | 4     |
|  | Negative  | 0          | 290      | 290   |
|  | Total     | 4          | 290      | 294   |
| PPA: 100.0% (51.0%, 100.0%)<br>NPA: 100.0% (98.7%, 100.0%) |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% of the ETEC prospective positive and negative specimens, and 100.0% and 99.6%-of the retrospective positive and negative specimens, respectively (Table 13).

**Table 13: ETEC PPA and NPA of the BD MAX™ Extended Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| ETEC   |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 2          | 0        | 2     |
|  | Negative  | 0          | 592      | 592   |
|  | Total     | 2          | 592      | 594   |
| PPA: 100.0% (34.2%, 100.0%)<br>NPA: 100.0% (99.4%, 100.0%) |           |            |          |       |
| Retrospective  | Positive  | 14         | 1        | 15    |
|  | Negative  | 0          | 279      | 279   |
|  | Total     | 14         | 280      | 294   |
| PPA: 100.0% (78.5%, 100.0%)<br>NPA: 99.6% (98.0%, 99.9%)   |           |            |          |       |

For the BD FecalSwab™ Collection, Transport and Preservation System, the BD MAX™ Enteric Bacterial Panel identified 100.0% of the *Yersinia enterocolitica* prospective negative specimens (zero prospective positive specimens were analyzed), and 100.0% and 99.0% of the retrospective positive and negative specimens, respectively (Table 14).



**Table 14: *Yersinia enterocolitica* PPA and NPA of the BD MAX™ Extended Enteric Bacterial Panel - FecalSwab™ Compared to Cary-Blair**

| <i>Yersinia enterocolitica</i>                           |           | Cary-Blair |          | Total |
|--|-----------|------------|----------|-------|
| Specimen Origin  | FecalSwab | Positive   | Negative |       |
| Prospective  | Positive  | 0          | 0        | 0     |
|  | Negative  | 1          | 593      | 594   |
|  | Total     | 1          | 593      | 594   |
| PPA: 0.0% (0.0%, 79.3%)<br>NPA: 100.0% (99.4%, 100.0%)   |           |            |          |       |
| Retrospective  | Positive  | 4          | 3        | 7     |
|  | Negative  | 0          | 287      | 287   |
|  | Total     | 4          | 290      | 294   |
| PPA: 100.0% (51.0%, 100.0%)<br>NPA: 99.0% (97.0%, 99.6%) |           |            |          |       |

Additional contrived specimens were evaluated due to low prevalence of the targets in the study. The BD FecalSwab™ Collection, Transport and Preservation System on the BD MAX™ Extended Enteric Bacterial Panel identified 100.0% of the *Plesiomonas shigelloides* contrived positive and negative specimens, when compared to expected results (Table 15).

**Table 15: *Plesiomonas shigelloides* Contrived FecalSwab™ Specimen Results**

| <i>Plesiomonas shigelloides</i>                            | Expected Results |    |    |     |
|--|------------------|----|----|-----|
| Contrived  | Positive         | 53 | 0  | 53  |
|  | Negative         | 0  | 53 | 53  |
|  | Total            | 53 | 53 | 106 |
| PPA: 100.0% (93.2%, 100.0%)<br>NPA: 100.0% (93.2%, 100.0%) |                  |    |    |     |

The BD FecalSwab™ Collection, Transport and Preservation System on the BD MAX™ Extended Enteric Bacterial Panel identified 98.1% and 100.0% of the positive and negative *Vibrio* contrived positive and negative specimens, respectively, when compared to expected results (Table 16).

**Table 16: *Vibrio* Contrived FecalSwab™ Specimen Results**

| <i>Vibrio</i> spp.                                       | Expected Results |    |    |     |
|--|------------------|----|----|-----|
| Contrived  | Positive         | 52 | 0  | 52  |
|  | Negative         | 1  | 53 | 54  |
|  | Total            | 53 | 53 | 106 |
| PPA: 98.1% (90.1%, 99.7%)<br>NPA: 100.0% (93.2%, 100.0%) |                  |    |    |     |

The BD FecalSwab™ Collection, Transport and Preservation System on the BD MAX™ Extended Enteric Bacterial Panel identified 100.0% of the ETEC contrived positive and negative specimens, when compared to expected results (Table 17).

**Table 17: ETEC Contrived FecalSwab™ Specimen Results**

| ETEC   | Expected Results |    |    |     |
|--|------------------|----|----|-----|
| Contrived  | Positive         | 53 | 0  | 53  |
|  | Negative         | 0  | 53 | 53  |
|  | Total            | 53 | 53 | 106 |
| PPA: 100.0% (93.2%, 100.0%)<br>NPA: 100.0% (93.2%, 100.0%) |                  |    |    |     |

The BD FecalSwab™ Collection, Transport and Preservation System on the BD MAX™ Extended Enteric Bacterial Panel identified 98.1% and 100.0% of the positive and negative *Yersinia enterocolitica* contrived positive and negative specimens, respectively, when compared to expected results (Table 18).

**Table 18: *Yersinia enterocolitica* Contrived FecalSwab™ Specimen Results**

| <i>Yersinia enterocolitica</i>                           | Expected Results |    |    |     |
|--|------------------|----|----|-----|
| Contrived  | Positive         | 52 | 0  | 52  |
|  | Negative         | 1  | 53 | 54  |
|  | Total            | 53 | 53 | 106 |
| PPA: 98.1% (90.1%, 99.7%)<br>NPA: 100.0% (93.2%, 100.0%) |                  |    |    |     |