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QlAstat-Dx[®] Gastrointestinal Panel 2 Instructions for Use



Version 1



For In Vitro Diagnostic Use

For use with QIAstat-Dx $^{\mathbb{R}}$ Analyzer 1.0, QIAstat-Dx $^{\mathbb{R}}$ Analyzer 2.0, and QIAstat-Dx $^{\mathbb{R}}$ Rise





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Intended Use

The QIAstat-Dx Gastrointestinal Panel 2 is a multiplexed nucleic acid test intended for use with the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise for the simultaneous qualitative detection and identification of nucleic acids from multiple viruses, bacteria, and parasites directly from stool samples in Cary-Blair transport media obtained from individuals with signs and/or symptoms of gastrointestinal infection. The following viruses, bacteria (including several diarrheagenic *E. coli/ Shigella* pathotypes), and parasites are identified with the QIAstat-Dx Gastrointestinal Panel 2:

- Adenovirus F40/F41
- Astrovirus
- Norovirus (GI/GII)
- Rotavirus A
- Sapovirus (GI, GII, GIV, GV)
- Campylobacter (C. jejuni, C. coli and C. upsaliensis)
- Clostridium difficile (toxin A/B)
- Enteroaggregative Escherichia coli (EAEC)
- Shigella/Enteroinvasive Escherichia coli (EIEC)
- Enteropathogenic Escherichia coli (EPEC)
- Enterotoxigenic Escherichia coli (ETEC) lt/st
- Plesiomonas shigelloides
- Salmonella spp.

- Shiga-like toxin-producing Escherichia coli (STEC) stx1/stx2* (including specific identification of E. coli O157 serogroup within STEC)
- Vibrio vulnificus
- Vibrio parahaemolyticus
- · Vibrio cholerae
- Yersinia enterocolitica
- Cryptosporidium
- Cyclospora cayetanensis
- Entamoeba histolytica
- Giardia lamblia

Concomitant culture is necessary for organism recovery and further typing of bacterial agents.

^{*}Shiga-like toxin-producing E. coli (STEC) genes (stx1 and stx2) are differentiated by QIAstat-Dx Gastrointestinal Panel 2

The QIAstat- Dx Gastrointestinal Panel 2 is indicated as an aid in the diagnosis of specific agents of gastrointestinal illness in conjunction with other clinical, laboratory, and epidemiological data. Confirmed positive results do not rule-out co-infection with organisms not detected by the QIAstat-Dx Gastrointestinal Panel 2. The organisms detected may not be the sole or definitive cause of the disease.

QlAstat- Dx Gastrointestinal Panel 2 is not intended to monitor or guide treatment for *C. difficile* infections.

Negative QIAstat-Dx Gastrointestinal Panel 2 results in the setting of clinical illness compatible with gastroenteritis may be due to infection by pathogens that are not detected by this assay test or non-infectious causes such as ulcerative colitis, irritable bowel syndrome, or Crohn's disease.

The QIAstat-Dx Gastrointestinal Panel 2 also aids in the detection and identification of acute gastroenteritis in the context of outbreaks. The QIAstat-Dx Gastrointestinal Panel 2 is intended for professional use only and is not intended for self-testing. The QIAstat-Dx Gastrointestinal Panel 2 is intended for in vitro diagnostic use.

Intended User

This kit is intended for professional use.

The product is to be used only by personnel specifically instructed and trained in molecular biology techniques and familiar with this technology.

Summary and Explanation

QIAstat-Dx Gastrointestinal Panel 2 cartridge description

The QIAstat-Dx Gastrointestinal Panel 2 Cartridge (Figure 1) is a disposable plastic device that allows performance of fully automated molecular assays for the detection of gastrointestinal pathogens. The main features of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge include compatibility with a liquid sample type, hermetical containment of the pre-loaded reagents necessary for testing, and true walk-away operation. All sample preparation and assay testing steps are performed within the cartridge.

All reagents required for the complete execution of a test run are pre-loaded and self-contained in the QIAstat-Dx Gastrointestinal Panel 2 Cartridge. The user does not need to come in contact with and/or manipulate any reagents. The QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise house air filters for both incoming and outgoing air, further safeguarding the environment. After testing, the cartridge stays hermetically closed at all times, greatly enhancing its safe disposal.

Within the cartridge, multiple steps are automatically performed in sequence using pneumatic pressure to transfer samples and fluids via the transfer chamber to their intended destinations.

After the sample is manually loaded, the diagnostic tests with the QIAstat-Dx Gastrointestinal Panel 2 are performed on the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise. All of the sample preparation and analysis steps are performed automatically by the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise.

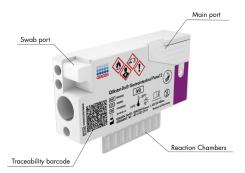


Figure 1. Layout of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge and its features.

Pathogen Information

Acute gastrointestinal infections can be caused by a variety of pathogens, including parasites, bacteria and viruses, and generally present with nearly indistinguishable clinical signs and symptoms. The rapid and accurate determination of the presence or absence of potential causative agent (s) helps make timely decisions regarding treatment, hospital admission, infection control and return of the patient to work and family. It may also greatly support improved antimicrobial stewardship and other important public health initiatives.

The QIAstat-Dx Gastrointestinal Panel 2 Cartridge allows detection and differentiation of 22 parasitic, viral and bacterial pathogens that cause gastrointestinal symptoms, which includes specific identification of *E. coli* O157 serogroup within STEC, resulting in 23 targets in total. Testing requires a small sample volume and minimal hands-on time, and the results are available in approximately 78 minutes.

Pathogens that can be detected and identified with the QIAstat-Dx Gastrointestinal Panel 2 are listed in Pathogens detected by the QIAstat-Dx Gastrointestinal Panel 2.

Sample collection and cartridge loading

The collection of samples and their subsequent loading into the QIAstat-Dx Gastrointestinal Panel 2 Cartridge should be performed by personnel trained in safe handling of biological samples.

The following steps are performed:

Fresh unpreserved stool specimen is collected and resuspended into Cary-Blair transport
medium as soon as possible after collection following the manufacturer's instructions.

Attention should be given not to exceed the maximum fill line of the Cary-Blair container.

2. The sample information is manually written on or a sample label is affixed to the top of a QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

Table 1. Pathogens detected by the QIAstat-Dx Gastrointestinal Panel 2

| Pathogen | Classification (genome type) |
|---|---------------------------------|
| Adenovirus F40/F41 | Adenovirus (DNA) |
| Astrovirus | Astrovirus (RNA) |
| Norovirus GI/GII | Calicivirus (RNA) |
| Rotavirus A | Reovirus (RNA) |
| Sapovirus (GI, GII, GIV, GV) | Calicivirus (RNA) |
| Campylobacter (C. jejuni, C. upsaliensis, C. colí) | Bacterium (DNA) |
| Clostridium difficile (toxin A/B) | Bacterium (DNA) |
| Enteroaggregative E. coli (EAEC) | Bacterium (DNA) |
| Enteroinvasive <i>E. coli</i> (EIEC)/Shigella | Bacterium (DNA) |
| Enteropathogenic E. coli (EPEC) | Bacterium (DNA) |
| Enterotoxigenic E. coli (ETEC) lt/st | Bacterium (DNA) |
| Plesiomonas shigelloides | Bacterium (DNA) |
| Salmonella spp. | Bacterium (DNA) |
| Shiga-like toxin-producing <i>E. coli</i> (STEC) stx1/stx2 (including specific identification of <i>E. coli</i> O157 serogroup within STEC) | Bacterium (DNA) |
| Vibrio vulnificus | Bacterium (DNA) |
| Vibrio parahaemolyticus | Bacterium (DNA) |
| Vibrio cholerae | Bacterium (DNA) |
| Yersinia enterocolitica | Bacterium (DNA) |
| | |

Table 1. Pathogens detected by the QIAstat-Dx Gastrointestinal Panel 2 (continued)

| Pathogen | Classification (genome type) |
|-------------------------|---------------------------------|
| Cryptosporidium | Parasite (DNA) |
| Cyclospora cayetanensis | Parasite (DNA) |
| Entamoeba histolytica | Parasite (DNA) |
| Giardia lamblia | Parasite (DNA) |

3. Liquid sample (stool resuspended in Cary-Blair transport medium) is loaded manually into the QlAstat-Dx Gastrointestinal Panel 2 Cartridge.

Note: Cary-Blair preserved stool specimens should present a homogenous suspension (easily vortexed).

Note: The user must perform a visual check of the sample inspection window to confirm that the liquid sample has been loaded.

- 4. The sample bar code (if available) and the QIAstat-Dx Gastrointestinal Panel 2 Cartridge bar code are scanned by the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, or QIAstat-Dx Rise. If sample bar code is not available, the sample ID is manually written using the virtual keyboard of the touchscreen.
- 5. The QIAstat-Dx Gastrointestinal Panel 2 Cartridge is introduced into the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, or QIAstat-Dx Rise.
- 6. The test is started on the QIAstat-Dx Analyzer 1.0, QIAstat-Dx Analyzer 2.0, or QIAstat-Dx Rise.

Sample preparation, nucleic acid amplification, and detection

The extraction, amplification, and detection of nucleic acids in the sample are performed automatically by the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0.

- The liquid sample is homogenized, and cells are lysed in the lysis chamber of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge, which includes a rotor that turns at high speed and silica beads that provide effective cell disruption.
- 2. Nucleic acids are purified from the lysed sample via binding to a silica membrane in the purification chamber of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge in the presence of chaotropic salts and alcohol.
- The purified nucleic acids are eluted from the membrane in the purification chamber and are mixed with the lyophilized PCR chemistry in the dried-chemistry chamber of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge.
- 4. The mixture of sample and PCR reagents is dispensed into the QIAstat-Dx Gastrointestinal Panel 2 Cartridge PCR chambers, which contain air- dried assay- specific primers and probes.
- The QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Rise creates the optimal temperature profiles
 to carry out effective multiplex real- time RT- PCR and performs real- time fluorescence
 measurements to generate amplification curves.
- 6. The QIAstat-Dx Analyzer 1.0 and QIAstat-Dx Rise Software interprets the resulting data and process controls and delivers a test report.

Materials Provided

Kit contents

| QIAstat-Dx Gastrointestinal Panel 2 Cartridge* Catalog number Number of tests | 691412 6 | |
|---|-------------|--|
| QIAstat-Dx Gastrointestinal Panel 2 Cartridges* | 6 | |
| Transfer pipettes† | 6 | |

^{* 6} individually packaged cartridges containing all reagents needed for sample preparation and multiplex real-time RT-PCR, plus Internal Control.

¹⁶ individually packaged transfer pipettes for dispensing liquid sample into the QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

Materials Required but Not Provided

Equipment

The QIAstat-Dx Gastrointestinal Panel 2 is designed for use with the QIAstat-Dx Analyzer 1.0, QIAstat- Dx Analyzer 2.0, and QIAstat- Dx Rise. Before beginning a test, make sure the following are available:

 QIAstat-Dx Analyzer 1.0 (at least one Operational Module and one Analytical Module) with software version 1.4 or later OR a QIAstat-Dx Rise (at least two Analytical Modules must be inside for the machine to work) with software version 2.2 or higher OR QIAstat-Dx Analyzer 2.0 (at least one Operational Module PRO and one Analytical Module) with software version 1.6 or higher.

Note: Application software version 1.6 or higher cannot be installed on QIAstat-Dx Analyzer 1.0.

- QIAstat-Dx Analyzer 1.0 and QIAstat-Dx Analyzer 2.0 User Manual (for use with software version from 1.4 to 1.5) OR QIAstat-Dx Rise User Manual (for use with software version 2.2 or later) OR QIAstat-Dx Analyzer 2.0 User Manual (for use with software version 1.6 or higher)
- QIAstat-Dx-specific Assay Definition File software for Gastrointestinal Panel 2 installed on the Operational Module or the Operational Module PRO.

^{*}Prior to use, ensure that instruments have been checked and calibrated according to the manufacturer's recommendations.

Warnings and Precautions

For in vitro diagnostic use.

The QIAstat-Dx Gastrointestinal Panel 2 is to be used by laboratory professionals trained in the use of QIAstat-Dx Analyzer 1.0, the QIAstat-Dx Analyzer 2.0, and the QIAstat-Dx Rise.

Safety information

When working with chemicals, always wear a suitable lab coat, disposable gloves, and protective goggles. For more information, please consult the appropriate safety data sheets (SDSs). These are available online in convenient and compact PDF format at www.qiagen.com/safety, where you can find, view and print the SDS for each QIAGEN kit and kit component.

Always wear appropriate personal protective equipment, including but not limited to disposable powder-free gloves, a lab coat, and protective eyewear. Protect skin, eyes and mucus membranes. Change gloves often when handling samples.

Handle all samples, used cartridges, and transfer pipettes as if they are capable of transmitting infectious agents. Always observe safety precautions as outlined in relevant guidelines, such as the Clinical and Laboratory Standards Institute [®] (CLSI) Protection of Laboratory Workers from Occupationally Acquired Infections; Approved Guideline (M29), or other appropriate documents provided by:

- OSHA®: Occupational Safety and Health Administration (United States of America)
- ACGIH®: American Conference of Government Industrial Hygienists (USA)
- COSHH: Control of Substances Hazardous to Health (United Kingdom)

Follow your institution's safety procedures for handling biological samples. Dispose of samples, QIAstat-Dx Gastrointestinal Panel 2 Cartridges, and transfer pipettes according to the appropriate regulations.

The QIAstat-Dx Gastrointestinal Panel 2 Cartridge is a closed, single-use device that contains all reagents needed for sample preparation and multiplex real-time RT-PCR within the QIAstat-Dx Analyzer 1.0, the QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise. Do not use a QIAstat-Dx Gastrointestinal Panel 2 Cartridge that is past its expiration date, appears damaged, or leaks fluid. Dispose of used or damaged cartridges in accordance with all national, state, and local health and safety regulations and laws.

Observe standard laboratory procedures for keeping the working area clean and contamination- free. Guidelines are outlined in publications such as the Biosafety in Microbiological and Biomedical Laboratories from the Centers for Disease Control and Prevention and the National Institutes of Health (www.cdc.gov/od/ohs/biosfty/biosfty.htm).

Precautions

The following hazard and precautionary statements apply to components of the QIAstat-Dx Gastrointestinal Panel 2.



Contains: ethanol; quanidine hydrochloride; quanidine thiocyanate; isopropanol; proteinase K; t-Octylphenoxypolyethoxyethanol. Danger! Highly flammable liquid and vapor. Harmful if swallowed or if inhaled. May be harmful in contact with skin. Causes severe skin burns and eye damage. May cause allergy or asthma symptoms or breathing difficulties if inhaled. May cause drowsiness or dizziness. Harmful to aquatic life with long lasting effects. Contact with acids liberates very toxic aas. Corrosive to the respiratory tract. Keep away from heat/sparks/open flames/hot surfaces. No smoking. Avoid breathing dust/fume/gas/mist/vapors/spray. Wear protective gloves/protective clothing/eye protection/face protection. Wear respiratory protection. IF IN EYES: Rinse cautiously with water for several minutes. Remove contact lenses, if present and easy to do. Continue rinsing. IF exposed or concerned: Immediately call a POISON CENTER or doctor/ physician. Remove person to fresh air and keep comfortable for breathing.

To reduce the risk of contamination when handling stool samples, it is recommended that the below guidelines are applied:

- When handling the stool sample, a biosafety cabinet, dead air box, splash shield, or face shield should be used.
- The work area used for cartridge loading should be separate from the work area used for stool pathogen testing (i.e., stool culture, EIA).
- Prior to sample handling, the work area should be thoroughly cleaned using 10% bleach or similar disinfectant.
- QIAstat-Dx Gastrointestinal Panel 2 Cartridges and samples should be processed one at a time.
- Change gloves prior to removing cartridges from shipping boxes.
- Change gloves and clean the work area between processing each sample.
- Dispose of used cartridges in a biohazard container immediately after the run is completed and avoid excessive handling.

Cartridge Storage and Handling

Store the QIAstat-Dx Gastrointestinal Panel 2 Cartridges in a dry, clean storage space at room temperature (15–25°C). Do not remove the QIAstat-Dx Gastrointestinal Panel 2 Cartridges or the transfer pipettes from their individual packaging until actual use. Under these conditions, QIAstat-Dx Gastrointestinal Panel 2 Cartridges can be stored until the expiration date printed on the individual packaging. The expiration date is also included in the QIAstat-Dx Gastrointestinal Panel 2 Cartridge bar code and is read by the QIAstat-Dx Analyzer 1.0, the QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise when the cartridge is inserted into the instrument to run a test. Once the cartridge is removed from the pouch it should be protected from sunlight.

Attention should be paid to expiration dates and storage conditions printed on the box and labels of all components. Do not use expired or incorrectly stored components.

Specimen Handling, Storage and Preparation

The QIAstat-Dx Gastrointestinal Panel 2 is for use with QIAstat-Dx Analyzer 1.0, the QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise. All samples should be treated as potentially hazardous.

Specimen collection

Stool samples should be collected and handled according to the Cary-Blair transport medium manufacturer's recommended procedures.

Recommended storage conditions for stool resuspended in Cary-Blair transport medium specimens are listed below:

- Room temperature up to 4 days at 15-25°C
- Refrigerated up to 4 days at 2-8°C

Protocol: Processing Raw Stool Samples in Cary-Blair transport medium

Sample collection, transport, and storage

Collect and resuspend the stool sample in Cary-Blair transport medium according to the manufacturer's recommended procedures.

Loading a sample into the QIAstat-Dx Gastrointestinal Panel 2 cartridge

Note: applicable for both the QIAstat-Dx 1.0, QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise

1. Open the package of a QlAstat-Dx Gastrointestinal Panel 2 Cartridge using the tear notches on the sides of the packaging (Figure 2).

IMPORTANT: After the package is opened, sample should be introduced into the QIAstat-Dx Gastrointestinal Panel 2 Cartridge within 30 minutes. Sample-loaded cartridges should be loaded into the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer

2.0 within 90 minutes or immediately into QIAstat-Dx Rise.



Figure 2. Opening the QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

- 2. Remove the QIAstat- Dx Gastrointestinal Panel 2 Cartridge from the packaging and position it so that the bar code on the label faces you.
- 3. Manually write the sample information or place a sample information label on the top of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge. Make sure that the label is properly positioned and does not block the lid opening (Figure 3). See QIAstat-Dx Rise workflow section for proper cartridge labelling.



Figure 3. Sample information placement on top of QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

4. Place the QIAstat-Dx Gastrointestinal Panel 2 Cartridge flat on the clean work surface so that the bar code on the label faces upwards. Open the sample lid of the main port on the front of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge (Figure 4).

IMPORTANT: Do not flip the QIAstat-Dx Gastrointestinal Panel 2 Cartridge or agitate it while the main port lid is open. The main port contains silica beads used in the sample disruption. The silica beads could fall out of the QIAstat- Dx Gastrointestinal Panel 2 Cartridge if it is agitated while the lid is open.

Note: The swab port is not used for the QIAstat-Dx Gastrointestinal Panel 2 assay.



Figure 4. Opening the sample lid of main port.

5. Thoroughly mix the stool in the Cary-Blair transport medium, for example, by vigorously agitating the tube 3 times (Figure 5).



Figure 5. Mixing stool sample in Cary-Blair transport medium.

6. Open the tube with the sample to be tested. Use the supplied transfer pipette to draw up fluid. Draw the sample to the second fill line on the pipette (i.e., 200 µl) (Figure 6).

IMPORTANT: Do not draw air, mucus, or particles into the pipette. If air, mucus, or particles are drawn into the pipette, carefully expel the sample fluid in the pipette back into the sample tube and draw up fluid again. In the event that the supplied transfer pipette is lost please use another one from the package or any other commercially available pipette with a minimum volume of 200 µl.

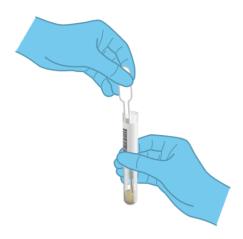


Figure 6. Drawing up sample into the supplied transfer pipette.

7. Carefully transfer the sample into the main port of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge using the supplied single-use transfer pipette (Figure 7).

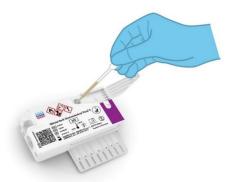


Figure 7. Transferring sample to main port of QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

8. Firmly close the lid of the main port until it clicks (Figure 8).



Figure 8. Closing the lid of the main port.

9. Visually confirm that the sample has been loaded by checking the sample inspection window of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge (Figure 9). A mixture of sample and silica beads should be observed.

IMPORTANT: After the sample is placed inside the QIAstat-Dx Gastrointestinal Panel 2 Cartridge, the cartridge must be loaded into the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0 within 90 minutes or immediately placed on the QIAstat-Dx Rise tray once all samples are loaded into the cartridges. The maximum waiting time for a cartridge that is already loaded into the QIAstat-Dx Rise (on-board stability) is about 145 minutes. The QIAstat-Dx Rise will automatically detect and warn the user if the cartridge has been placed into the instrument for a longer time than permitted.

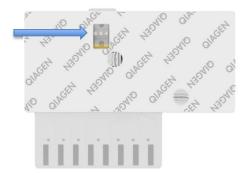


Figure 9. Sample inspection window (blue arrow).

Running a test with the QIAstat-Dx Analyzer 1.0

1. Power ON the QIAstat- Dx Analyzer 1.0 using the On/Off button on the front of the instrument

Note: The power switch on the back of the Analytical Module must be set in the "I" position. The QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 status indicators will turn blue.

 Wait until the Main screen appears and the QIAstat- Dx Analyzer 1.0 or QIAstat- Dx Analyzer 2.0 status indicators turn green and stop blinking. 3. Log in to the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 by entering the user name and password.

Note: The Login screen will appear if User Access Control is activated. If the User Access Control is disabled, no user name/password will be required, and the Main screen will appear.

- 4. If the Assay Definition File software has not been installed on the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0, follow the installation instructions prior to running the test (see "Appendix A: Installing the Assay Definition File", for additional information).
- 5. Press the Run Test button in the top right corner of the touchscreen of the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0.
- 6. When prompted, scan the sample ID bar code on the Cary-Blair sample or scan the specimen information bar code located on the top of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge (see step 3) using the integrated front bar code reader of the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 (Figure 10).

Note: It is also possible to enter the sample ID using the virtual keyboard of the touchscreen by selecting the Sample ID field.

Note: Depending on the chosen system configuration, entering the patient ID may also be required at this point.

Note: Instructions from the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 appear in the Instructions Bar at the bottom of the touchscreen.



Figure 10. Scanning sample ID bar code.

7. When prompted, scan the bar code of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge to be used (Figure 11). The QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 will automatically recognize the assay to be run based on the cartridge bar code.

Note: The QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 will not accept QIAstat-Dx Gastrointestinal Panel 2 Cartridges with lapsed expiration dates, previously used cartridges or cartridges for assays that have not been installed on the unit. An error message will be shown in these cases and the QIAstat- Dx Gastrointestinal Panel 2 Cartridge will be rejected. Refer to the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 User Manual or Appendix A for further details on how to install assays.



Figure 11. Scanning QIAstat-Dx Gastrointestinal Panel 2 Cartridge bar code.

- 8. The Confirm screen will appear. Review the entered data and make any necessary changes by selecting the relevant fields on the touchscreen and editing the information.
- 9. Press Confirm when all the displayed data are correct. If needed, select the appropriate field to edit its content, or press Cancel to cancel the test (Figure 12).



Figure 12. Confirming data entry.

- Ensure that both sample lids of the swab port and main port of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge are firmly closed.
- When the cartridge entrance port on the top of the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx
 Analyzer 2.0 automatically opens, insert the QIAstat-Dx Gastrointestinal Panel 2

Cartridge with the bar code facing to the left and the reaction chambers facing down (Figure 13).

Note: Depending on the system configuration, the operator may be required to re-enter their user password to start the test run.

Note: Up to this point, it is possible to cancel the test run by pressing the Cancel button in the bottom right corner of the touchscreen.

12. Upon detecting the QIAstat- Dx Gastrointestinal Panel 2 Cartridge, the QIAstat- Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0 will automatically close the lid of the cartridge entrance port and start the test run. No further action from the operator is required to start the run.

Note: There is no need to push the QIAstat-Dx Gastrointestinal Panel 2 Cartridge into the QIAstat-Dx Analyzer 1.0 or QIAstat-Dx Analyzer 2.0.

Note: The QIAstat-Dx Analyzer 1.0 and the QIAstat-Dx Analyzer 2.0 will not accept a QIAstat-Dx Gastrointestinal Panel 2 Cartridge other than the one used and scanned during the test setup. If a cartridge other than the one scanned is inserted, an error will be generated, and the cartridge will be automatically ejected.

Note: The lid of the cartridge entrance port will close automatically after 30 seconds if a QIAstat-Dx Gastrointestinal Panel 2 Cartridge is not positioned in the port. If this occurs, repeat the procedure starting from step 5.



Figure 13. Inserting QIAstat-Dx Gastrointestinal Panel 2 Cartridge into QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0.

- 13. While the test is running, the remaining run time is displayed on the touchscreen.
- 14. After the test run is completed, the Eject screen will appear (Figure 14) and the Module status bar will display the test result as one of the following options:
 - TEST COMPLETED: The test was completed successfully
 - TEST FAILED: An error occurred during the test
 - TEST CANCELED: The user canceled the test

IMPORTANT: If the test fails, refer to the 'Troubleshooting' section in the QIAstat- Dx Analyzer 1.0 or the QIAstat- Dx Analyzer 2.0 User Manual for possible reasons and instructions on how to proceed. For additional information about specific QIAstat- Dx Gastrointestinal Panel 2 error codes and messages, refer to the 'Troubleshooting' section of this document.



Figure 14. Eject screen display.

- 15. Press Eject on the touchscreen to remove the QIAstat- Dx Gastrointestinal Panel 2 Cartridge and dispose of it as biohazardous waste in accordance with all national, state and local health and safety regulations and laws. The QIAstat-Dx Gastrointestinal Panel 2 Cartridge should be removed when the cartridge entrance port opens and ejects the cartridge. If the cartridge is not removed after 30 seconds, it will automatically move back into the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0 and the cartridge entrance port lid will close. If this occurs, press Eject to open the lid of the cartridge entrance port again and then remove the cartridge.
 - IMPORTANT: Used QIAstat-Dx Gastrointestinal Panel 2 Cartridges must be discarded. It is not possible to re-use cartridges for tests for which the execution was started but then subsequently canceled by the operator, or for which an error was detected.
- 16. After the QIAstat- Dx Gastrointestinal Panel 2 Cartridge has been ejected, the results Summary screen will appear. Refer to "Interpretation of Results", page 52 for further details. To begin the process for running another test, press Run Test.
 - Note: For further information on the use of the QlAstat-Dx Analyzer 1.0 or QlAstat-Dx Analyzer 2.0, refer to the QlAstat-Dx Analyzer 1.0 or QlAstat-Dx Analyzer 2.0 User Manual.

Running a test on the QIAstat-Dx Rise

Starting the QIAstat-Dx Rise

1. Press the ON/OFF button on the front panel of the QIAstat-Dx Rise to start the unit.

Note: The power switch at the rear-left connection box must be set to the "I" position.

- 2. Wait until the Login screen appears and the LED status indicators turn green.
- 3. Log in to the system once the login screen appears (Figure 15).

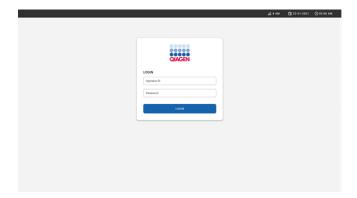


Figure 15. Log in screen.

Note: After successful initial installation of the QIAstat-Dx Rise, the system administrator needs to log in for the initial configuration of the software.

Preparing the QIAstat-Dx Gastrointestinal Panel 2 cartridge

Remove the QIAstat-Dx Gastrointestinal Panel 2 cartridge from its packaging. For details about adding the sample to the QIAstat-Dx Gastrointestinal Panel 2 cartridge and for

information specific to the assay to be run, refer to "Loading a sample into the QIAstat-Dx Gastrointestinal Panel 2 cartridge".

Always make sure that both sample lids are firmly closed after adding a sample to the QIAstat-Dx Gastrointestinal Panel 2 cartridge.

Adding a sample barcode to the QIAstat-Dx Gastrointestinal Panel 2 cartridge

Place a barcode on the top-right side of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge (indicated by the arrow) (Figure 16).



Figure 16. Placing sample ID barcode.

The maximum barcode size is: 22 mm x 35 mm. The barcode must always be on the right side of the cartridge (as it is shown above with red marked area), as the left side of the cartridge is critical for sample autodetection (Figure 17).

Note: To process samples on the QIAstat-Dx Rise, it is required to provide a machine-readable sample ID barcode on the QIAstat-Dx Gastrointestinal Panel 2 Cartridge.



Figure 17. Positioning sample ID barcode.

1D and 2D barcodes can be used. Usable 1D barcodes are the following: EAN-13 and EAN-8, UPC-A and UPC-E, Code128, Code39, Code 93, and Codabar. Usable 2D barcodes are Aztec Code, Data Matrix, and QR code.

Make sure that the barcode quality is sufficient. The system is capable of reading a printing quality of grade C or better, as defined in ISO/IEC 15416 (linear) or ISO/IEC 15415 (2D).

Procedure to run a test

Note: All operators should wear appropriate personal protective equipment, such as gloves, lab coat, and protective glasses when handling the QIAstat- Dx Rise touchscreen and cartridges.

- Press the OPEN WASTE DRAWER button at the lower-right corner of the main test screen (Figure 18).
- Open the waste drawer and remove used cartridges from previous runs. Check the waste drawer for spilled liquids. If necessary, clean the waste drawer as described in the Maintenance section of the QIAstat-Dx Rise User Manual.

- 3. Close the waste drawer after removal of the cartridges. The system will scan the tray and return to the main screen (Figure 18). If the tray was removed for maintenance purposes, make sure it is correctly inserted before closing the drawer.
- 4. Press the OPEN INPUT DRAWER button on the lower-right corner of the screen (Figure 18).

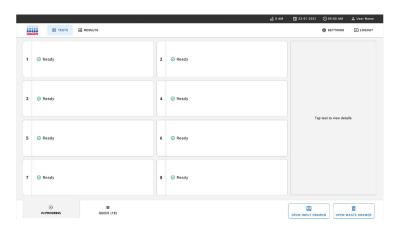


Figure 18. Main test screen.

5. Wait until the input drawer is unlocked (Figure 19).

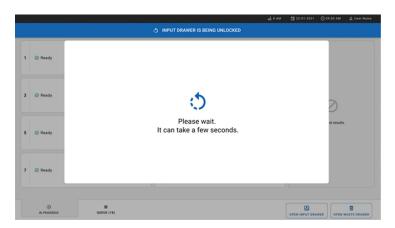


Figure 19. Input drawer waiting dialog box.

6. When prompted, pull the input drawer to open (Figure 20).

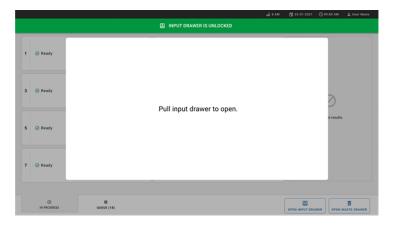


Figure 20. Input drawer open dialog box.

7. The Add Cartridge dialog appears and the scanner in front of the instrument will be activated. Scan the sample ID barcode on top of the QIAstat-Dx Gastrointestinal 2

cartridge in front of the instrument (position indicated by the arrow (Figure 21).

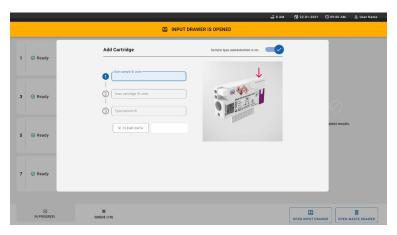


Figure 21. Scan sample ID screen.

8. After entering the sample ID barcode, scan the bar code of the QIAstat-Dx Gastrointestinal Panel 2 cartridge to be used (position indicated by the arrow). The QIAstat-Dx Rise will automatically recognize the assay to be run, based on the QIAstat-Dx Gastrointestinal Panel 2 cartridge barcode (Figure 22).

Note: Make sure that Sample type autodetection is set to on. The system will automatically recognize the used sample type (if applicable for the assay used).

If Sample type autodetection is set to off, you might need select the appropriate sample type manually (if applicable for the assay used).

Note: The QIAstat-Dx Rise will not accept QIAstat-Dx Gastrointestinal Panel 2 cartridges that have lapsed expiration dates, were previously used, or if the QIAstat-Dx Gastrointestinal Panel 2 assay definition file is not installed on the unit. An error message will be shown in this case.

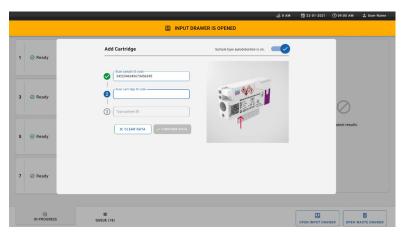


Figure 22. Scanning the QIAstat-Dx Gastrointestinal Panel 2 cartridge ID screen.

9. Input the patient ID (Patient ID has to be set to on) then confirm the data (Figure 23 and 24).

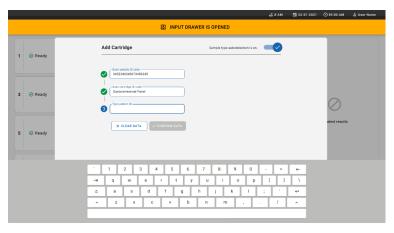


Figure 23. Typing the patient ID.

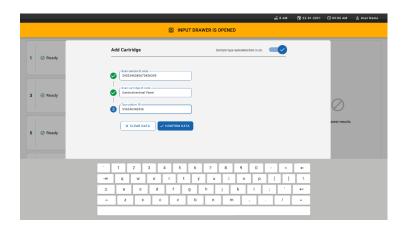


Figure 24. Type patient ID then confirm the data screen.

10. After a successful scan, the following dialog box appears briefly on top of the screen (Figure 25).



Figure 25. Cartridge saved screen.

- 11. Place the cartridge into the input drawer. Make sure that the cartridge is inserted properly into the tray (Figure 26).
- 12. Continue scanning and inserting cartridges by following the previous steps.

IMPORTANT: Please be aware that QIAstat-Dx Rise can handle up to 16 QIAstat-Dx Gastrointestinal Panel 2 cartridges at the same time, within the input drawer.

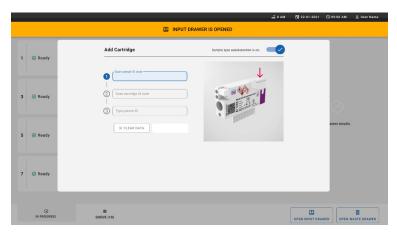


Figure 26. Add cartridge screen.

13. Close the input drawer when all cartridges have been scanned and inserted. The system will scan the cartridges and prepare a queue (Figure 27).

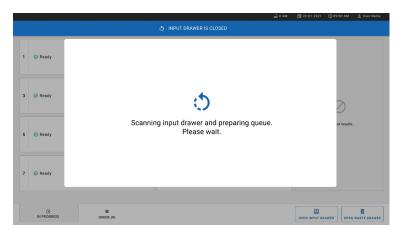


Figure 27. Preparing queue screen.

14. After successful scanning, the queue will be shown (Figure 28). Review the data and in case of an error, press the OPEN INPUT DRAWER button remove and re-scan the respective cartridge, following steps 10-13.

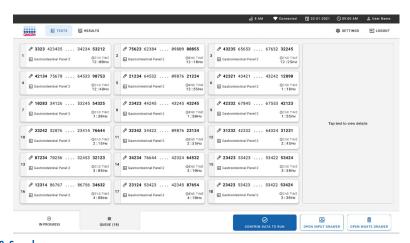


Figure 28. Sample queue screen.

Note: The sample order on the screen may not match the cartridge order in the input drawer (it only matches when all the cartridges are queued together) and cannot be changed without opening the input tray and removing cartridges.

The sample queue/processing order is generated by QIAstat-Dx Rise based on the following rules:

- Stability time. QIAstat-Dx Gastrointestinal Panel 2 cartridges with the shortest on-board stability time will be prioritized irrespective of the position in the loading tray.
- Within the same assay type, the position in the loading tray determines the order in queue.

If you select a test on the touchscreen, additional information is displayed in the TEST DETAILS section of the screen (Figure 29).

Note: The system will reject cartridges that exceed the maximum on-board stability time within the input drawer (about 145 minutes)

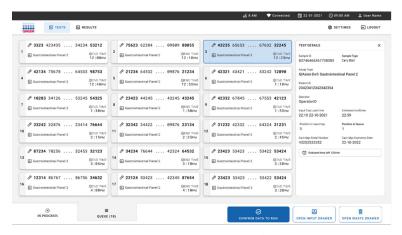


Figure 29. Sample queue screen with selected assay showing additional information.

The following information is shown in the Test Details section (Figure 30):

- Sample ID
- · Sample Type (depending on the assay)
- Assay Type (QIAstat-Dx Gastrointestinal Assay Panel 2)
- Patient ID
- Operator
- Input Tray Load Time
- Estimated end time
- Position in input drawer
- Position in Queue (Note: the position may differ based on sample stability time)

- · Cartridge Serial Number
- Cartridge Expiration Date
- · Onboard time left

Note: The on-board time is defined in the respective assay and triggers the order of samples in the queue.

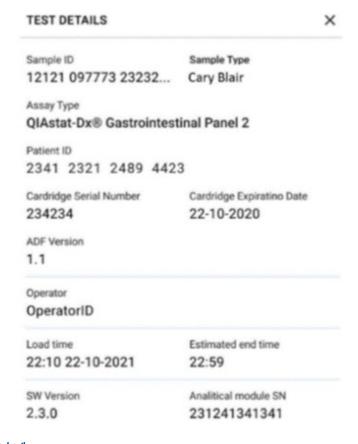


Figure 30. Test details.

15. Press the CONFIRM DATA TO RUN button at the bottom of the screen when all the displayed data are correct (Figure 29). thereafter, a final confirmation is required from the operator to run the tests (Figure 31).

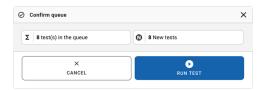


Figure 31. Final confirmation to run rest.

While the tests are running, the remaining run time and other information for all queued tests are displayed on the touchscreen (Figure 32).

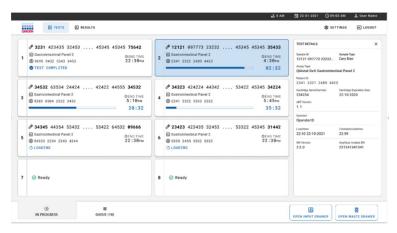


Figure 32. Test execution information on queue screen.

If the cartridge is being loaded into an Analytical Module, a TEST LOADING message and the estimated end time are displayed (Figure 33).



Figure 33. Test loading message and end time.

If the test is running, the elapsed run time and the approximate end time are displayed (Figure 34).

```
₱ 23423 423435 32453 .... 45345 45345 80855
   ■ Gastrointestinal Panel 2
3

② 9383 6904 4836 3855

                                                   5:10<sub>PM</sub>
                                                    20:32
```

Figure 34. Elapsed run time and approximate end time view.

If the test is completed, a TEST COMPLETED message and the run end time are displayed (Figure 35).

```
№ 23423 423435 32453 .... 45345 45345 80855
III Gastrointestinal Panel 2
                                         OEND TIME
                                         22:30рм
2 9695 9432 5543 3453
TEST COMPLETED
```

Figure 35. Test completed view.

Prioritizing samples

If a sample needs to be run urgently, it is possible to select this sample on the sample queue screen and run as a first sample (Figure 36). Please note that it is not possible to prioritize a sample after confirmation of the queue

Prioritizing sample before starting run

The urgent sample is selected on the queue screen and marked URGENT from right hand side of the sample queue screen before confirm data to run. (Figure 36). Following this, the sample is moved to the first position of the queue (Figure 37). Note that only one sample can be prioritized.

Note: It is required to open and close the input drawer otherwise it is not possible to prioritize a cartridge that has already been confirmed. At this point, if the Urgent button is not active, the operator needs to switch between QUEUE and IN PROGRESS tabs on the GUI to see the active Urgent button.

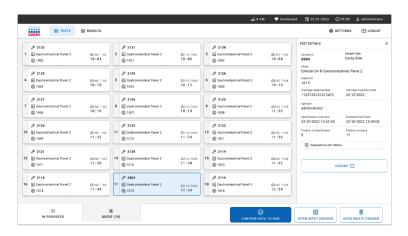


Figure 36. Sample queue screen while selecting sample to be prioritized.

Some other samples may run out of stability time due to prioritization of a sample. This warning can be seen on the right corner of the screen (Figure 37).

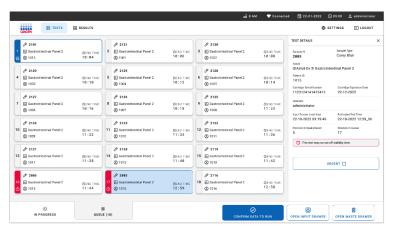


Figure 37. Sample queue screen after a sample is prioritized.

After confirmation of the gueue the run can be started (Figure 38).

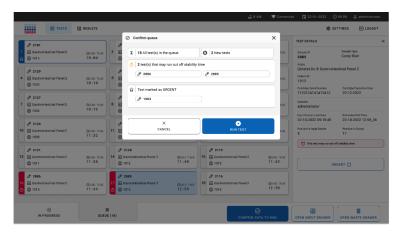


Figure 38. Confirmation of the run screen.

Prioritizing sample during run

A sample can be also prioritized for any reason during the run. In this case, if there is no available AM, any other ongoing sample needs to be aborted to perform prioritization (Figure 39).

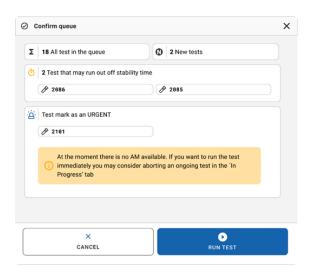


Figure 39. Confirmation dialog during run.

Abortion of running sample

A sample can be aborted during scanning, loading and running. Please note that the sample cannot be used again once it is aborted. This is also true for the sample that is aborted during scanning and loading.

To abort a sample, go to "in progress" tab of the screen and select the sample and push "abort" option on the right corner of the screen (Figure 40).

It is not possible to abort a run while a sample is about to load into AM or about to complete to run and the system is retrieving result data or/and technical logs from the respective AM.

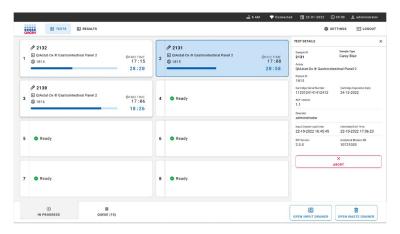


Figure 40. Abortion of a running sample.

The system needs a confirmation to abort the sample (Figure 41).



Figure 41. Confirmation dialog to abort running sample.

After a while, the sample can be seen as "aborted" on the screen (Figure 42 and Figure 43).

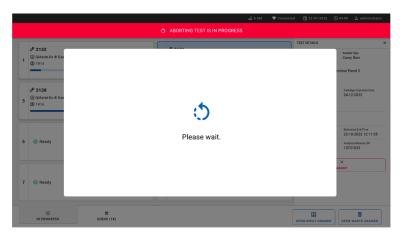


Figure 42. Sample abortion waiting dialog.

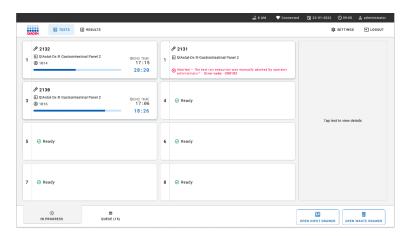


Figure 43. Aborted sample after confirmation of the abortion.

Interpretation of results

Viewing results with the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0

The QIAstat- Dx Analyzer 1.0 or the QIAstat- Dx Analyzer 2.0 automatically interprets and saves test results. After ejecting the QIAstat-Dx Gastrointestinal Panel 2 Cartridge, the results Summary screen is automatically displayed. Figure 44 shows the screen for the QIAstat-Dx Analyzer 1.0.

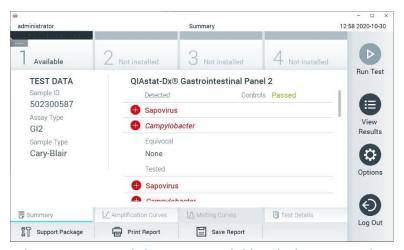


Figure 44. Results Summary screen example showing Test Data on the left panel and Test Summary in the main panel in the QIAstat-Dx Analyzer 1.0.

From this screen, other tabs with more information, which will be explained in the following chapters, are available:

- **Amplification Curves**
- Melting Curves. This tab is disabled for the QIAstat-Dx Respiratory SARS-CoV-2 Panel.

· Test Details.

Figure 45 shows the screen for the QIAstat-Dx Analyzer 2.0.

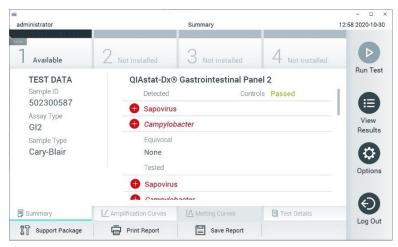


Figure 45. Results Summary screen example showing Test Data on the left panel and Test Summary in the main panel in QIAstat-Dx Analyzer 2.0.

QIAstat-Dx Analyzer 2.0 includes an additional tab:

AMR Genes, It is disabled for the QIAstat-Dx Gastrointestinal Panel 2.

Note: From this point forward, example screen shots will be used when referring to the QIAstat-Dx Analyzer 1.0 and/or QIAstat-Dx Analyzer 2.0 where the functions being explained are the same.

The main part of the screen provides the following lists and uses color-coding and symbols to indicate the results:

- The first list, under the heading "Detected", includes all pathogens detected and identified in the sample, which are preceded by a 👽 sign and are colored red.
- The second list, under the heading "Equivocal" is not used. "Equivocal" results are not applicable for the QIAstat-Dx Gastrointestinal Panel 2. Therefore, the "Equivocal" list will always be empty.
- The third list, under the heading "Tested", includes all pathogens tested in the sample. Pathogens detected and identified in the sample are preceded by a 🕀 sign and are colored red. Pathogens that were tested but not detected are preceded by a lacktriangle sian and are colored green. Invalid and not applicable pathogens are also displayed in this list.

Note: Pathogens detected and identified in the sample are shown in both the "Detected" and "Tested" lists.

If the test failed to complete successfully, a message will indicate "Failed" followed by the specific Error Code.

The following Test Data is shown on the left side of the screen:

- · Sample ID
- · Patient ID (if available)
- Assay Type
- · Sample Type

Further data about the assay is available, depending on the operator's access rights, through the tabs at the bottom of the screen (e.g., amplification plots and test details).

A report with the assay data can be exported to an external USB storage device. Insert the USB storage device into one of the USB ports of the QIAstat-Dx Analyzer 1.0 and press Save Report in the bottom bar of the screen. This report can be exported later at any time by selecting the test from the View Result List.

The report can also be sent to the printer by pressing Print Report in the bottom bar of the screen.

Viewing amplification curves

To view test amplification curves of pathogens detected, press the L Amplification Curves tab (Figure 46).



Figure 46. Amplification Curves screen (PATHOGENS tab).

Details about the tested pathogens and controls are shown on the left and the amplification curves are shown in the center.

Note: If User Access Control is enabled on the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0, the Amplification Curves screen is only available for operators with access rights.

Press the **PATHOGENS** tab on the left side to display the plots corresponding to the tested pathogens. Press the pathogen name to select which pathogens are shown in the amplification plot. It is possible to select single, multiple, or no pathogens. Each pathogen in the selected list will be assigned a color corresponding to the amplification curve associated with the pathogen. Unselected pathogens will be shown in gray. The corresponding C_T and endpoint fluorescence (EP) values are shown below each pathogen name.

Press the **CONTROLS** tab on the left side to view the controls in the amplification plot. Press the circle next to the control name to select or deselect it (Figure 47).



Figure 47. Amplification Curves screen (CONTROLS tab).

The amplification plot displays the data curve for the selected pathogens or controls. To alternate between logarithmic or linear scale for the Y-axis, press the Lin or Log button at the bottom left corner of the plot.

The scale of the X-axis and Y-axis can be adjusted using the blue pickers on each axis. Press and hold a blue picker and then move it to the desired location on the axis. Move a blue picker to the axis origin to return to the default values.

Viewing test details

Press Test Details in the Tab Menu bar at the bottom of the touchscreen to review the results in more detail. Scroll down to see the complete report. The following Test Details are shown in the center of the screen (Figure 48):

- User ID
- Cartridge SN (serial number)
- Cartridge Expiration Date
- Module SN (serial number)
- Test Status (Completed, Failed or Canceled by operator)
- · Error Code (if applicable)
- · Test Start Date and Time
- Test Execution Time
- · Assay Name
- Test ID
- · Test Result:
 - · Positive (if at least one gastrointestinal pathogen is detected/identified)
 - Positive with warning (if at least one pathogen is detected, but the Internal Control failed)

- Negative (if no gastrointestinal pathogen is detected)
- Failed (an error occurred, or the test was canceled by the user)
- · List of analytes tested in the assay, with C_T and endpoint fluorescence in the event of a positive signal
- Internal Control, with C_T and endpoint fluorescence



Figure 48. Example screen showing Test Data on the left panel and Test Details in the main panel.

Browsing results from previous tests

To view results from previous tests that are stored in the results repository, press Results on the Main Menu bar (Figure 49).

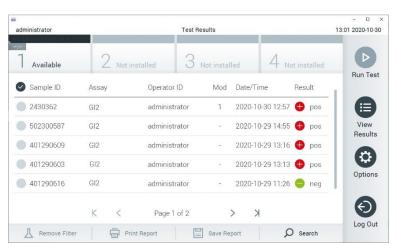


Figure 49. View Results screen.

The following information is available for every executed test (Figure 48):

- Sample ID
- Assay (name of test assay, which is "GI2" for Gastrointestinal Panel 2)
- Operator ID
- Mod (Analytical Module on which the test was executed)
- Date/Time (date and time when the test was finished)
- Result (outcome of the test: positive [pos], positive with warning [pos*], negative [neg], failed [fail] or successful [suc])

Note: If User Access Control is enabled on the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0, the data for which the user has no access rights will be hidden with asterisks.

Select one or more test results by pressing the gray circle to the left of the sample ID. A checkmark will appear next to selected results. Unselect test results by pressing this checkmark.

The entire list of results can be selected by pressing the checkmark circle in the top row (Figure 50).

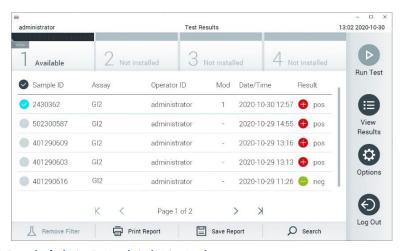


Figure 50. Example of selecting Test Results in the View Results screen.

Press anywhere in the test row to view the result for a particular test.

Press a column headline (e.g., Sample ID) to sort the list in ascending or descending order according to that parameter. The list can be sorted according to only one column at a time.

The Result column shows the outcome of each test (Descriptions of test results displayed on View Results screen):

Table 2. Descriptions of test results displayed on View Results screen

| Outcome | Result | Description | Action |
|-----------------------|---------------|---|---|
| Positive | pos | At least one pathogen is positive | Refer to the Summary Result Screen or Result Printout for pathogen specific results. Description of pathogen results can be found in Description of Pathogen results as displayed on Summary Result Screen and the Result Printout. |
| Positive with warning | pos* | At least one pathogen is positive, but the Internal Control failed | Refer to the Summary Result Screen or Result Printout for pathogen specific results. Description of pathogen results can be found in Description of Pathogen results as displayed on Summary Result Screen and the Result Printout. |
| Negative | neg | No pathogen were detected | Refer to the Summary Result Screen or Result Printout for pathogen specific results. Description of pathogen results can be found in Description of Pathogen results as displayed on Summary Result Screen and the Result Printout. |
| Failed | ⊗ fail | The test failed because either an error occurred, the test was canceled by the user, or no pathogens were detected and the internal control failed. | Repeat the test using a new cartridge. Accept the results of the repeat testing. If the error persists, contact QIAGEN Technical Services for further instructions. |
| Successful | Suc | The test is either positive or negative, but the user does not have the access rights to view the test results | Login from a user profile with rights to view the results. |

Make sure a printer is connected to the QIAstat-Dx Analyzer 1.0 and the proper driver is installed. Press Print Report to print the report(s) for the selected result(s).

Press **Save Report** to save the report(s) for the selected result(s) in PDF format to an external USB storage device.

Select the report type: List of Tests or Test Reports.

Press **Search** to search the test results by Sample ID, Assay and Operator ID. Enter the search string using the virtual keyboard and press Enter to start the search. Only the records containing the search text will be displayed in the search results.

If the results list has been filtered, the search will only apply to the filtered list. Press and hold a column headline to apply a filter based on that parameter. For some parameters, such as Sample ID, the virtual keyboard will appear so the search string for the filter can be entered.

For other parameters, such as Assay, a dialog box will open with a list of assays stored in the repository. Select one or more assays to filter only the tests that were performed with the selected assays.

The r symbol to the left of a column headline indicates that the column's filter is active.

A filter can be removed by pressing **Remove Filter** in the Submenu bar.

Exporting results to a USB drive

From any tab of the View Results screen, select Save Report to export and save a copy of the test results in PDF format to a USB drive. The USB port is located on the front of the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0.

Printing results

Make sure a printer is connected to the QIAstat-Dx Analyzer 1.0 or the QIAstat-Dx Analyzer 2.0 and the proper driver is installed. Press Print Report to send a copy of the test results to the printer.

Sample result interpretation

A result for a gastrointestinal organism is interpreted as "Positive" when the corresponding PCR assay is positive, except for EPEC, STEC, and E. coli O157. The result interpretation for EPEC, STEC, and E. coli O157 follows the rationale explained in Interpretation of results, below.

Table 3. Interpretation of EPEC, STEC, and E. coli O157 results

| | STEC stx 1/stx 2 Result* | | | E. coli 0157 | |
|-------------|--------------------------|-------|--------------|--------------|---|
| EPEC RESULT | stx1 | stx 2 | stx1 + stx 2 | Result | Description |
| Negative | | | Negative | N/A | Enteropathogenic <i>E. coli</i> (EPEC) was not detected and Shiga-like toxin-producing <i>E. coli</i> (STEC) stx1/stx2 is negative as both stx1 and stx2 have not been detected. |
| | | | | | E. coli O157 result is not applicable (N/A) when Shiga-like toxin-producing E. coli (STEC) stx1/stx2 is not detected due to E.coli O157 being a specific serotype of STEC |
| | | | | | Enteropathogenic <i>E. coli</i> (EPEC) was detected and Shiga-like toxin-producing <i>E. coli</i> (STEC) stx1/stx2 is negative as both stx1 and stx2 have not been detected. |
| Positive | | | Negative | N/A | E. coli O157 result is not applicable (N/A) when Shiga-like toxin-producing <i>E. coli</i> (STEC) stx1/stx2 is not detected due to E.coli O157 being a specific serotype of STEC. |
| N/A | Positive | | | Negative | EPEC result is not applicable because EPEC detection cannot be differentiated when STEC stx1 or stx2 is detected. |
| | | | | | E. coli O157 was not detected. |

Table 3. Interpretation of EPEC, STEC, and E. coli O157 results (continued)

| | STEC stx 1/stx 2 Result* | | | E. coli 0157 | |
|-------------|--------------------------|----------|--------------|--|--|
| EPEC RESULT | stx1 | stx 2 | stx1 + stx 2 | Result | Description |
| N/A | | Positive | | Negative | EPEC result is not applicable because EPEC detection cannot be differentiated when STEC stx1 or stx2 is detected. E. coli O157 was not detected. |
| N/A | | Positive | Negative | EPEC result is not applicable because EPEC detection cannot be differentiated when both STEC stx1 and stx2 are detected. | |
| | | | | | E. coli 0157 was not detected. |
| N/A | Positive | | | Positive | EPEC result is not applicable because EPEC detection cannot be differentiated when STEC stx1 or stx2 is detected. |
| | | | | | E. coli 0157 was detected. |
| N/A | | Positive | | Positive | EPEC result is not applicable because EPEC detection cannot be differentiated when STEC stx1 or stx2 is detected. |
| | | | | | E. coli 0157 was detected. |
| N/A | | | Positive | Positive | EPEC result is not applicable because EPEC detection cannot be differentiated when both STEC stx1 and stx2 are detected. E. coli O157 was detected. |

Internal control results are to be interpreted according to Interpretation of Internal Control results.

Table 4. Interpretation of Internal Control results

| Control Result | Explanation | Action |
|-------------------|---|---|
| Passed | The Internal Control amplified successfully | The run was completed with success. All results are validated and can be reported. Detected pathogens are reported as "positive" and undetected pathogens are reported as "negative". |
| Failed | The Internal Control failed | Positively detected pathogen(s) are reported, but all negative results (tested but not detected pathogen[s]) are invalid. Repeat the testing using a new Cartridge. Accept the results of the repeat testing. If the invalid result persists, contact QIAGEN Technical Services for further instruction |

The software provides an overall test result (Descriptions of test results displayed on View Results screen) as well as a result for individual pathogens. Possible results for each organism include Detected/Positive, Not Detected/Negative, N/A, and Invalid (Description of Pathogen results as displayed on Summary Result Screen and the Result Printout). If the internal control has failed and no positive signal was detected or if there is an instrument error, there will be no pathogen results provided.

Table 5. Description of Pathogen results as displayed on Summary Result Screen and the Result Printout

| Result | Symbol | Explanation | Action |
|---------------------------------------|--------|--|---|
| Positive/ Detected | 0 | A positive signal was detected for this pathogen. Result of the Internal Control is passed. | None. Report results. |
| Positive/ Detected with Warning | pos* | A positive signal was detected for this pathogen, but the result of the internal control has failed. | Report positive analyte. Repeat the test using a new cartridge. Accept the results of the repeat testing. If the invalid result persists, contact QIAGEN Technical Services for further instructions. |
| Negative/ Not Detected | | No signal was detected for this pathogen. The Internal Control passed. | None. Report results. |

Table 5. Description of Pathogen results as displayed on Summary Result Screen and the Result Printout (continued)

| Result | Symbol | Explanation | Action |
|---|--------|--|--|
| N/A (applies to E. coli O157 and EPEC only) | 8 | The run was successfully completed and the Internal Control passed. For <i>E. coli</i> O157 N/A: Shiga-like toxin-producing <i>E. coli</i> (STEC) was not detected. For EPEC N/A: Shiga-like toxin producing <i>E. coli</i> (STEC) was detected. | None. Report results. |
| Invalid | 8 | No signal was detected for this pathogen and the Internal Control failed (but other pathogens have been detected). | Repeat the test using a new cartridge. Accept the results of the repeat testing. If the invalid result persists, contact QIAGEN Technical Services for further instructions. |

Interpretation of results with QIAstat-Dx Rise

Viewing results with QIAstat-Dx Rise

The QIAstat-Dx Rise automatically interprets and saves test results. After the run completed, the results can be seen in the Results summary screen (Figure 51).

Note: Visible information will be dependent on the operator's access rights.

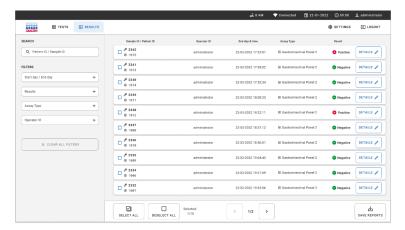


Figure 51. The results summary screen.

- The main part of the screen provides an overview of the completed runs and uses colorcoding and symbols to indicate the results:
 - If at least one pathogen is detected in the sample, the word Positive is shown in the result column, preceded by a \bigoplus sign.
- If no pathogen is detected, and the internal control is valid, the word Negative is shown in the result column, preceded by a sign.
- If at least one pathogen is detected in the sample, and the internal control was invalid, the term Positive with warning is shown in the result column, preceded by a •! sign.
- If the test failed to complete successfully, a message will indicate Failed followed by the specific Error Code.

The following Test Data are on the screen (Figure 50):

- Sample ID/Patient ID
- Operator ID

- · End day and time
- · Assay Type

Viewing test details

Further data about the assay is available, depending on the operator's access rights, through the Details button at the right side of the screen (e.g., amplification plots, and test details (Figure 52).

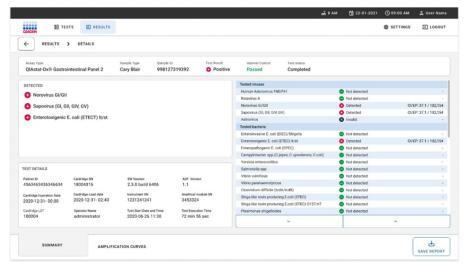


Figure 52. The test details screen.

The upper part of the screen shows general information about the test. It includes assay and sample type, Sample ID, overall test result, status of the internal control, and the test status.

On the left side of the screen, all detected pathogens are shown, the middle part of the screen shows all pathogens that the assay can detect.

Note: Categories and type of pathogens displayed depend on the assay used.

On the right side of the screen, the following test details are shown: Sample ID, operator ID, cartridge lot number, cartridge serial number, cartridge expiration date, cartridge load date and time, test execution date and time, test execution duration, Software and ADF version, and the analytical Module serial number.

Viewing amplification curves

To view the test amplification curves, press the Amplification Curves tab at the bottom of the screen (Figure 53).

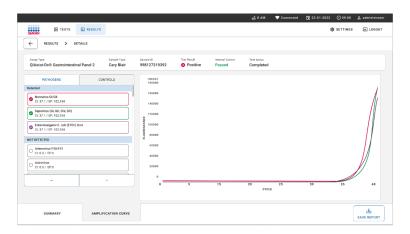


Figure 53. The amplification curves screen.

Press the PATHOGENS tab on the left side to display the plots corresponding to the tested pathogens. Press the pathogen name to select which pathogens are shown in the amplification plot. You can select single, multiple, or no pathogens. Each pathogen in the selected list will be assigned a color corresponding to the amplification curve associated with the pathogen. Unselected pathogens will not be shown.

The corresponding C_T and endpoint fluorescence values are shown below each pathogen name. Pathogens are grouped into detected and not detected.

"Equivocal" results are not applicable for the QIAstat-Dx Gastrointestinal Panel 2. Therefore, the "Equivocal" list will always be empty.

Press the CONTROLS tab on the left side to view the controls and select which controls are shown in the amplification plot.

Browsing results from previous tests

To view results from previous tests that are stored in the results repository, use the search functionality in the main results screen (Figure 53).

Note: The functionality may be restricted or disabled due to user profile settings.

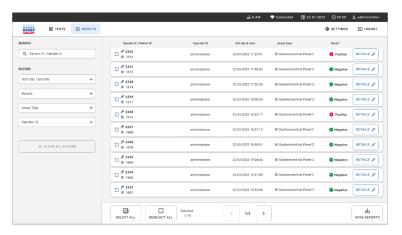


Figure 54. Search functionality in the results screen.

Exporting results to a USB storage device

From the Results screen, select individually or all with Select All button to export and save a copy of the test reports in PDF format to a USB storage device (Figure 53). The USB port is located in front and on the rear of the instrument.

Note: It is recommended to use the USB storage device for short-term data saving and transfer only. The use of a USB storage device is subject to restrictions (e.g. the memory capacity or the risk of overwriting, which should be considered before usage).

Quality Control

Internal control interpretation

The QIAstat-Dx Gastrointestinal Panel Cartridge includes a full process Internal Control, which is titered *Schizosaccharomyces pombe*. *Schizosaccharomyces pombe* is a yeast (fungi) that is included in the cartridge in dried form and is rehydrated upon sample loading. This Internal Control material verifies all steps of the analysis process, including sample homogenization, lysis of viral, and cellular structures (by means of chemical and mechanical disruption), nucleic acid purification, reverse transcription, and real-time PCR.

A passed result for the Internal Control indicates that all processing steps performed by the QIAstat-Dx Gastrointestinal Panel Cartridge were successful.

A failed result of the Internal Control does not negate any positive results for detected and identified targets, but it does invalidate all negative results in the analysis. Therefore, the test should be repeated if the Internal Control signal is negative.

External control information

All external quality control requirements and testing should be performed in accordance with local, state, and federal regulations or accreditation organizations and should follow the user's laboratory standard quality control procedures.

Limitations

- Results from the QIAstat-Dx Gastrointestinal Panel 2 are not intended to be used as the sole basis for diagnosis, treatment, or other patient management decisions.
- · For prescription use only.
- The performance of this test has only been validated with human stool collected in Cary-Blair transport medium, according to the media manufacturers' instructions. It has not been validated for use with other stool transport media, rectal swabs, raw stool, vomitus, or endoscopy stool aspirates.
- The QIAstat-Dx Gastrointestinal Panel 2 should not be used to test Cary-Blair vials from collection devices that have been overfilled with stool. Only stool resuspended following the collection device manufacturer's instructions should be used.
- The performance of this test has not been determined for patients without signs and symptoms of gastrointestinal illness.
- Results from this test must be correlated with the clinical history, epidemiological data, and
 other data available to the clinician evaluating the patient. Due to high rates of
 asymptomatic carriage of Clostridium difficile, especially in very young children and
 hospitalized patients, the detection of toxigenic C. difficile should be interpreted within the
 context of guidelines developed by the testing facility or other experts.
- Positive results do not rule out co-infection with organisms not included in the QIAstat-Dx Gastrointestinal Panel 2. The agent detected may not be the definitive cause of the disease.
- Negative results do not preclude infection of the gastrointestinal tract. Not all agents of
 acute gastrointestinal infection are detected by this assay and sensitivity in some clinical
 settings may differ from that described in the Instructions for Use.
- A negative result with the QIAstat- Dx Gastrointestinal Panel 2 does not exclude the infectious nature of the syndrome. Negative assay results may originate from several

factors and their combinations, including sample handling mistakes, variation in the nucleic acid sequences targeted by the assay, infection by organisms not included in the assay, organism levels of included organisms that are below the limit of detection for the assay and use of certain medications (e.g., calcium carbonate).

- The QIAstat-Dx Gastrointestinal Panel 2 is not intended for testing of samples other than
 those described in this Instructions for Use. Test performance characteristics have been
 established only with unpreserved stool samples resuspended in Cary- Blair transport
 medium.
- The QIAstat-Dx Gastrointestinal Panel 2 is intended to be used in conjunction with standard of care culture for organism recovery, serotyping and/or antimicrobial susceptibility testing where applicable.
- The results from the QIAstat-Dx Gastrointestinal Panel 2 must be interpreted by a trained healthcare professional within the context of all relevant clinical, laboratory, and epidemiological findings.
- The QIAstat-Dx Gastrointestinal Panel 2 can be used only with the QIAstat-Dx Analyzer 1.0,
 QIAstat-Dx Analyzer 2.0, and QIAstat-Dx Rise.
- The identification of multiple diarrheagenic E. coli pathotypes has historically relied upon phenotypic characteristics, such as adherence patterns or toxigenicity in certain tissue culture cell lines. The QIAstat- Dx Gastrointestinal Panel 2 targets genetic determinants characteristic of most pathogenic strains of these organisms but may not detect all strains having phenotypic characteristics of a pathotype. In particular, the QIAstat- Dx Gastrointestinal Panel 2 will only detect Enteroaggregative E. coli (EAEC) strains carrying the aggR and/or aatA markers on the pAA (aggregative adherence) plasmid; it will not detect all strains exhibiting an aggregative adherence pattern.
- Genetic virulence markers associated with diarrheagenic *E.coli/Shigella* pathotypes are often carried on mobile genetic elements (MGEs) that can be transferred horizontally

between different strains, therefore "Detected" results for multiple diarrheagenic *E. coli/Shigella* may be due to co-infection with multiple pathotypes or, less frequently, may be due to the presence of a single organism containing genes characteristic of multiple pathotypes. An example of the latter is the 2019 *E. coli* hybrid ETEC/STEC strains found in Sweden*.

*Bai X, Zhang J, Ambikan A, et al. Molecular Characterization and Comparative Genomics of Clinical Hybrid Shiga Toxin-Producing and Enterotoxigenic *Escherichia coli* (STEC/ETEC) Strains in Sweden. Sci Rep. 2019;9(1):5619. Published 2019 Apr 4. doi:10.1038/s41598-019-42122-z

- The QIAstat-Dx Gastrointestinal Panel 2 detects heat-stable toxin variants (ST1a and ST1b) and the heat-labile toxin (LT) of Enterotoxigenic E. coli (ETEC), which are associated with human disease. The variant LT- II toxin (structurally similar to LT) and the STB/ST2 toxin (structurally dissimilar to ST1) are not targeted by the ETEC oligonucleotide designs and have not been established as important in human disease.
- The QIAstat-Dx Gastrointestinal Panel 2 detects Enteropathogenic E. coli (EPEC) through targeting of the eae gene, which encodes the adhesin intimin. As some Shiga-like toxin-producing E. coli (STEC) also carry eae (in particular, strains identified as enterohemorrhagic E. coli; EHEC), the QIAstat-Dx Gastrointestinal Panel 2 cannot distinguish between STEC containing eae and a co-infection of EPEC and STEC. Therefore, the EPEC result is not applicable (N/A) and not reported for specimens in which STEC has also been detected. In rare cases, STEC may be reported as EPEC when a STEC carrying eae (EHEC) is present in a specimen below the LoD of the STEC oligonucleotide design(s) (stx1/stx2). Rare instances of other organisms carrying eae have been documented; e.g., Escherichia albertii, and Shigella boydii.
- Shigella dysenteriae serotype 1 possess a shiga toxin gene (stx) that is identical to the stx1 gene of STEC. Stx genes have been more recently found in other Shigella species (e.g., S. sonnei and S. flexneri). The detection of both Shigella/Enteroinvasive E. coli (EIEC) and STEC stx1/stx2 analytes in the same specimen may indicate the presence of Shigella species such as S. dysenteriae. Rare instances of the detection of Shiga-like toxin genes in

- other genera/species have been reported; e.g., Acinetobacter haemolyticus, Enterobacter cloacae and Citrobacter freundii.
- The presence of Shigella species carrying the stx1 gene, such as S. dysenteriae in the specimen will be reported as STEC stx1 + Shigella. Being the EPEC result not applicable (N/A) due to the reporting of STEC. Therefore, the QIAstat-Dx Gastrointestinal Panel will not report EPEC in the event of a co-infection with Shigella species carrying stx1 gene.
- E. coli O157 result is only reported as specific serogroup identification in association with STEC stx1/stx2. While non-STEC O157 strains have been detected in human stool, their role in disease has not been established. Serotype O157 EPEC has been identified and will be detected by the QIAstat-Dx Gastrointestinal Panel 2 (by the EPEC oligonucleotides design) due to their carriage of the eae gene. The E. coli O157 result will be not applicable (N/A) due to the absence of STEC.
- The QIAstat-Dx Gastrointestinal Panel 2 cannot distinguish between infections with a single toxigenic STEC O157 or rare co-infections of STEC (non-O157) with a stx-negative E. coli O157, that will also be detected as STEC O157.
- This test only detects Campylobacter jejuni, C. coli and C. upsaliensis and does not differentiate between these three species of Campylobacter. Additional testing is required to differentiate between these species and to detect other Campylobacter species that may be present in stool specimens. In particular the Campylobacter upsaliensis oligonucleotides design may cross-react with the Campylobacter species C. lari and C. helveticus organisms.
- A negative QIAstat-Dx Gastrointestinal Panel 2 result does not exclude the possibility of gastrointestinal infection. Negative test results may occur from sequence variants in the region targeted by the assay, the presence of inhibitors, technical error, sample mix-up, or an infection caused by an organism not detected by the panel. Test results may also be affected by concurrent antimicrobial therapy or levels of organism in the sample that are below the limit of detection for the test. Negative results should not be used as the sole

basis for diagnosis, treatment, or other management decisions.

- Organism and amplicon contamination may produce erroneous results for this test.
 Particular attention should be given to the Laboratory Precautions noted under the Laboratory Precautions section.
- The performance of the QIAstat-Dx Gastrointestinal Panel 2 has not been established in individuals who received Rotavirus A vaccine. Recent oral administration of a Rotavirus A vaccine may cause positive results for Rotavirus A if the virus is passed in the stool.
- Based on the available sequences, a few Cryptosporidium species, or certain variants of species, including C. wrari, may not be efficiently detected by the Cryptosporidium design.
 These species are rarely detected in human samples.
- There is a risk of false negative results due to the presence of strains with sequence variability in the target regions of the oligonucleotides design. Refer to the inclusivity testing section of this document for additional information.
- Not all Salmonella serotypes were tested in validation studies; however, representatives of
 the 20 most prevalent serotypes recently circulating in the US (CDC National Salmonella
 Surveillance Annual Summary 2016) were evaluated during analytical reactivity studies. In
 silico sequence analysis supports detection of all subspecies and serotypes of Salmonella.
- The performance of this test has not been evaluated for immunocompromised individuals.
- State and local public health authorities have published guidelines for notification of reportable diseases in their jurisdictions including Salmonella, Shigella, V. cholerae, E. coli O157, Enterotoxigenic E. coli (ETEC) It/st, and Shiga-like toxin-producing E. coli (STEC) stx1/stx2 to determine necessary measures for verification of results to identify and trace outbreaks. Laboratories are responsible for following their state or local regulations for submission of clinical material or isolates on positive specimens to their state public health laboratories.

- There is a risk of false-positive values resulting from cross-contamination by target organisms, their nucleic acids, or the amplified product.
- All assay results should be used and interpreted in the context of a full clinical evaluation as an aid in the diagnosis of gastrointestinal infection.
- There is a risk of false-positive values resulting from non-specific signals in the assay.
- Analyte targets (virus, bacteria, or parasite nucleic acid sequences) may persist in vivo, independent of virus, bacteria, or parasite viability. Detection of analyte target(s) does not guarantee that the corresponding live organism(s) is present, or that the corresponding organism(s) is the causative agent for clinical symptoms.
- The detection of viral, bacterial, or parasitic sequences is dependent upon proper specimen collection, handling, transportation, storage, and preparation (including extraction). Failure to observe proper procedures in any one of these steps can lead to incorrect results.
- Underlying polymorphisms in primer-binding regions can affect the targets being detected and subsequently the test results returned.
- There is a risk of false negative values resulting from improperly collected, transported, or handled specimens.
- There is a risk of false negative values due to the presence of strain/species sequence variability in the targets of the assay, procedural errors, amplification inhibitors in specimens, or inadequate numbers of organisms for amplification.
- The performance of this test has not been established for monitoring treatment of infection with any of the targeted microorganisms.
- Positive and negative predictive values are highly dependent on prevalence. False negative
 test results are more likely when prevalence of disease is high. False positive test results are
 more likely when prevalence is low.

- The effect of interfering substances has only been evaluated for those listed in the labeling at its indicated amount or concentration. Interference by substances other than those described in the "Interfering Substances" section of the Instruction for Use can lead to erroneous results.
- Cross- reactivity with gastrointestinal tract organisms other than those listed in the "Analytical Specificity" section of the package insert may lead to erroneous results.
- This test is a qualitative test and does not provide the quantitative value of detected organism present.
- The assay sensitivity to detect Cyclospora cayetanensis, Adenovirus F41, Entamoeba histolytica and the Shiga-like toxin- producing Escherichia coli (STEC) might be reduced up to 3.16-fold when using half-input sample volume (100 μL) workflow detailed in Appendix C.

Performance Characteristics

Analytical performance

The analytical performance shown below was demonstrated using QIAstat-Dx Analyzer 1.0. The QIAstat-Dx Analyzer 2.0 uses the same Analytical Module as QIAstat-Dx Analyzer 1.0 therefore the performance is not impacted by QIAstat-Dx Analyzer 2.0.

With regards to QIAstat- Dx Rise, specific studies to demonstrate the carryover and the repeatability were executed. The rest of analytical performance parameters shown below were demonstrated using QIAstat-Dx Analyzer 1.0. The QIAstat-Dx Rise uses the same Analytical Module as QIAstat-Dx Analyzer 1.0 therefore the performance is not impacted by QIAstat-Dx Rise.

Sensitivity (Limit of Detection)

The Analytical Sensitivity, or Limit of Detection (LoD), is defined as the lowest concentration at which \geq 95% of the tested samples generate a positive call.

The LoD for each of the QIAstat-Dx Gastrointestinal Panel 2 target pathogenic organisms was assessed, using in total 48 pathogen strains, by analyzing serial dilutions of analytical samples prepared from culture isolates from commercial suppliers (e.g., ZeptoMetrix® and ATCC®), confirmed clinical isolates, or artificial samples for target analytes commercially unavailable. Each sample tested was prepared in human stool matrix, which consists of a pool of previously tested negative clinical stool specimens resuspended in Cary-Blair transport medium.

Each of the 48 strains was tested in human stool matrix prepared following the manufacturer's instructions for the Para-Pak C&S® collection device.

Individual LoD values for each QIAstat- Dx Gastrointestinal Panel 2 target is shown in LoD values obtained for the different gastrointestinal target strains tested with the QIAstat- Dx Gastrointestinal Panel 2.

Table 6. LoD values obtained for the different gastrointestinal target strains tested with the QIAstat-Dx Gastrointestinal Panel 2

| Pathogen | Strain | Source | Concentration (molecular units: copies/ml) | Concentration (microbiological units) | Detection rate |
|-----------------------------|---|------------------------|---|---|----------------|
| | Campylobacter coli 76-GA2 [LMG 21266] | ATCC 43478 | 5802 | 1.2 CFU/ml | 20/20 |
| | Campylobacter coli CIP 7080 | ATCC 33559 | 8941 | 0.6 CFU/ml | 20/20 |
| | Campylobacter jejuni Z086 | ZeptoMetrix 801650 | 14491 | 1660 CFU/ml | 20/20 |
| Campylobacter | Campylobacter jejuni subsp. Jejuni RM3193 | ATCC BAA- 1234 | 7210 | 110 CFU/ml | 19/20 |
| | Campylobacter upsaliensis NCTC 11541 | ZeptoMetrix 0801999 | 56165 | 2259.4 CFU/ml | 20/20 |
| | Campylobacter upsaliensis RM3195 | ATCC BAA- 1059 | 7631 | 35 CFU/vial | 19/20 |
| Clostridium difficile | (NAP1A) Toxinotype III A+ B+ | ZeptoMetrix 801619 | 11083 | 515 CFU/ml | 19/20 |
| toxin A/B | Toxinotype 0 A+ B+ | ATCC 9689 | 101843 | 853.2 CFU/ml | 20/20 |
| Plesiomonas shigelloides | Z130 | ZeptoMetrix 801899 | 481 | 2291 CFU/ml | 20/20 |
| singenoides | Bader | ATCC 14029 | 116 | 2.7 CFU/vial | 19/20 |

| Pathogen | Strain | Source | Concentration (molecular units: copies/ml) | Concentration (microbiological units) | Detection rate |
|----------------------------|---|------------------------|---|---|----------------|
| Salmonella | Salmonella enterica Serovar choleraseus | ATCC 13312 | 647 | 91.6 CFU/ml | 20/20 |
| Salmonella | Salmonella enterica Serovar Typhimurium Z005 | ZeptoMetrix 801437 | 1441 | 4518.8 CFU/ml | 20/20 |
| Vibrio cholerae | Z132; toxigenic | ZeptoMetrix 801901 | 28298 | 13600 CFU/ml | 20/20 |
| VIDIIO CNOIErae | Z133; non-toxigenic | ZeptoMetrix 801902 | 79749 | 54668 CFU/ml | 20/20 |
| Vibrio | EB 101 | ATCC 17802 | 12862 | 1600 CFU/ml | 20/20 |
| parahaemolyticus | Z134 | ZeptoMetrix 801903 | 8904 | 143 CFU/ml | 20/20 |
| Vibrio vulnificus | 329 [CDC B3547] | ATCC 33817 | 109131 | 260 CFU/ml | 20/20 |
| VIDITO VUITITICUS | 324 [CDC B629] | ATCC 27562 | 2983 | 1305.1 CFU/ml | 20/20 |
| | Z036 | ZeptoMetrix 0801734 | 719 | 2070 CFU/ml | 20/20 |
| Yersinia enterocolitica | subsp. enterocolitica NTCC 11175, Biotype 4, serotype 3 | ATCC 700822 | 2496 | 120.1 CFU/ml | 20/20 |
| Enteroaggregative | Escherichia coli 92.0147, 077:HN | ZeptoMetrix 0801919 | 1075 | 634 CFU/ml | 20/20 |
| E. coli (EAEC) | Escherichia coli CDC3250-76, O111a, 111b: K58:H21 | ATCC 29552 | 842 | 87 CFU/ml | 19/20 |

| Pathogen | Strain | Source | Concentration (molecular units: copies/ml) | Concentration (microbiological units) | Detection rate |
|-----------------------------|--|-------------------------------------|---|---|----------------|
| Enteroinvasive | Shigella sonnei Z004 | ZeptoMetrix 25931 | 488 | 0.2 CFU/ml | 20/20 |
| E. coli (EIEC)/ Shigella | Escherichia coli CDC EDL 1282, O29:NM | ATCC 43892 | 1431 | 41.3 CFU/ml | 20/20 |
| Enteropathogenic E. | Escherichia coli O111:NM (EPEC) | ZeptoMetrix 0801747 | 1817 | 2581.7 CFU/ml | 20/20 |
| coli (EPEC) | Escherichia coli 7.1493; EPEC; O84:H28 | Zeptometrix 801938 | 29021 | 1190 CFU/ml | 20/20 |
| Enterotoxigenic | Escherichia coli H10407, O78:H11 | ATCC 35401 | 367 | 10.1 CFU/ml | 19/20 |
| E. coli (ETEC) lt/st | Escherichia coli ETEC; ST+, LT+ | ZeptoMetrix 801624 | 855 | 567 CFU/ml | 20/20 |
| | Cryptosporidium hominis | Public Health Wales UKM 84 | 357 | N/A | 20/20 |
| Cryptosporidium | Cryptosporidium parvum – lowa isolate | Waterborne® P102C | 661 | N/A | 20/20 |
| Cyclospora | N/A | LACNY-Clinical sample LAC2825 | 53 | N/A | 19/20 |
| cayetanensis | N/A | LACNY Clinical sample LAC2827 | 137 | N/A | 20/20 |
| Entamoeba histolytica | HM-1:IMSS (Mexico City 1967) | ATCC 30459 | 7 | 0.2 cells/ml | 20/20 |
| | HK-9 (Korea) | ATCC 30015 | 1 | 0.01 cells/ml | 19/20 |

| Pathogen | Strain | Source | Concentration (molecular units: copies/ml) | Concentration (microbiological units) | Detection rate |
|-----------------|----------------------------|---|---|---|----------------|
| Giardia lamblia | WB (Bethesda) | ATCC 30957 | 11850 | 632 cells/ml | 19/20 |
| | Portland-1 | ATCC 30888 | 14500 | 635 cells/ml | 20/20 |
| Adenovirus | Type 40 (Dugan) | ZeptoMetrix 0810084CF | 11726 | 0.1 TCID50/ml | 20/20 |
| F40/F41 | Type 41 (Tak) | ZeptoMetrix 0810085CF | 979 | 0.5 TCID50/ml | 19/20 |
| Astrovirus | ERE IID 2371 (type 8) | Zeptometrix 0810277CF | 11586371 | 11.7 TCID50/ml | 20/20 |
| Asilovilus | ERE IID 2868 (type 4) | Zeptometrix 0810276CF | 52184 | 1.3 TCID50/ml | 19/20 |
| Norovirus Gl | GI.1 (recombinant) | ZeptoMetrix 0810086CF | 24629 | 891.1 TCID50/ml | 19/20 |
| Norovirus GII | GII.4 (recombinant) | ZeptoMetrix 0810087CF | 8998 | 1.1 TCID50/ml | 20/20 |
| Rotavirus A | 69M | ZeptoMetrix 0810280CF | 5787 | 436.1 TCID50/ml | 19/20 |
| KOIQVII US A | Wa | ZeptoMetrix 0810041CF | 5201 | 14.1 TCID50/ml | 19/20 |
| Sapovirus | Genogroup I, genotype 1 | QIAGEN Barcelona - Clinical sample GI-88 | 187506 | N/A | 20/20 |
| | Genogroup V | Universitat de Barcelona 160523351 | 3007 | N/A | 20/20 |

Exclusivity (Analytical Specificity)

The analytical specificity study was carried out by in vitro testing and in silico analysis (9) to assess the potential cross-reactivity and exclusivity of the QIAstat-Dx Gastrointestinal Panel 2. On-panel organisms were tested to assess the potential for intra-panel cross-reactivity and Offpanel organisms were tested to evaluate cross-reactivity with organisms not covered by the panel content. The On-panel and Off-panel organisms tested are shown in Table 7 and Table 8, respectively.

Samples were prepared by single spiking organisms into negative stool resuspended in Cary-Blair at the highest concentration possible based on the organism stock, preferably at 10^5 TCID₅₀/ml for viral, 10^5 cells/ml for parasite targets and 10^6 CFU/ml for bacterial targets. The pathogens were tested in 3 replicates. There was no intra-panel or Off-panel cross-reactivity for all pathogens tested in vitro, except for two non-targeted *Campylobacter* species (*C. helveticus* and *C. lari*) that cross-reacted with the *Campylobacter* assay oligonucleotides included in the QIAstat-Dx Gastrointestinal Panel 2.

Table 7. List of Analytical Specificity on-panel pathogens tested

| Туре | Pathogen | | |
|-----------|---------------------------|--------------------------|--|
| | Campylobacterx coli | Plesiomonas shigelloides | |
| | Campylobacter jejuni | Salmonella enterica | |
| | Campylobacter upsaliensis | Shigella sonnei | |
| | Clostridium difficile | Vibrio cholerae | |
| Bacteria | Escherichia coli (EAEC) | Vibrio parahaemolyticus | |
| | Escherichia coli (EPEC) | Vibrio vulnificus | |
| | Escherichia coli (ETEC) | Yersinia enterocolitica | |
| | Escherichia coli (STEC) | | |
| | | | |
| Parasites | Cryptosporidium parvum | Entamoeba histolytica | |
| | Cyclospora cayetanensis | Giardia lamblia | |

Table 7. List of Analytical Specificity on-panel pathogens tested (continued)

| Туре | Pathogen | | |
|---------|----------------|---------------|--|
| Viruses | Adenovirus F41 | Norovirus GII | |
| | Astrovirus | Rotavirus A | |
| | Norovirus Gl | Sapovirus | |
| | | | |

| Туре | Pathogen (potential cross-reactant) | |
|----------|-------------------------------------|-------------------------------------|
| Bacteria | Abiotrophia defectiva | Enterobacter cloacae |
| | Acinetobacter baumannii | Enterococcus faecalis |
| | Aeromonas hydrophila | Enterococcus faecium |
| | Arcobacter cryaerophilus | Escherichia fergusonii |
| | Bacillus subtilis | Escherichia hermannii |
| | Bifidobacterium bifidum | Escherichia vulneris |
| | Campylobacter fetus | Faecalibacterium prausnitzii |
| | Campylobacter gracilis | Gardnerella vaginalis |
| | Campylobacter helveticus | Haemophilus influenzae |
| | Campylobacter hominis | Helicobacter pylori |
| | Campylobacter lari | Klebsiella pneumoniae |
| | Campylobacter mucosalis | Lactobacillus casei |
| | Campylobacter rectus | Listeria monocytogenes |
| | Chamydia trachomatis | Proteus mirabilis |
| | Citrobacter freundii | Proteus vulgaris |
| | Clostridium difficile non-toxigenic | Pseudomonas aeruginosa |
| | Clostridium perfringens | Staphylococcus aureus |
| | Clostridium septicum | Staphylococcus aureus subsp. Aureus |
| | Clostridium tetani | Staphylococcus epidermidis |
| | Corynebacterium genitalium | Streptococcus agalactiae |
| | Enterobacter aerogenes | Streptococcus pyogenes |
| | | |
| Fungi | Aspergillus fumigatus | Saccharomyces boulardii |
| | Candida albicans | Sacabara myoo aarayisiaa |
| | | Saccharomyces cerevisiae |

Table 8. List of Analytical Specificity off-panel pathogens tested (continued)

| Туре | Pathogen (potential cross-reactant) | |
|-----------|-------------------------------------|-----------------------------|
| Parasites | Babesia microti | Toxoplasma gondii |
| | Blastocystis hominis | Trichomonas tenax |
| | Giardia muris | |
| Viruses | Adenovirus C:2 | Coronavirus 229E |
| | Adenovirus B:34 | Coxsackievirus B3 |
| | Adenovirus B3 | Cytomegalovirus |
| | Adenovirus E:4a | Enterovirus 6 (Echovirus) |
| | Adenovirus serotype 1 | Enterovirus 68 |
| | Adenovirus serotype 5 | Herpes Simplex Virus Type 2 |
| | Adenovirus serotype 8 | Rhinovirus 1A |
| | Bocavirus Type 1 | |

In silico predictions of potential cross-reactions showed that the following cross-reactions may occur when testing stool samples with the QIAstat-Dx Gastrointestinal Panel 2 (Potential cross-reactions based on in silico analysis) (5, 15–17).

Table 9. Potential cross-reactions based on in silico analysis

| QIAstat-Dx Gastrointestinal Panel 2 Target | Potential cross-reactive organisms |
|---|---|
| Enteropathogenic <i>E. coli</i> (EPEC) | Shigella boydii *†‡, Escherichia albertii *† |
| Campylobacter spp. | Campylobacter lari §, Campylobacter helveticus § |
| Shiga-like toxin-producing <i>E. coli</i> (STEC) stx1 | Shigella sonnei *‡, Shigella dysenteriae*‡ |
| Shiga-like toxin-producing <i>E. coli</i> (STEC) stx2 | Acinetobacter haemolyticus*¶, Citrobacter freundi*¶, Enterobacter cloacae*¶, Aeromonas caviae*¶ Escherichia albertii *¶ |

Table 9. Potential cross-reactions based on in silico analysis (continued)

QIAstat-Dx Gastrointestinal Panel 2 Target

Potential cross-reactive organisms

E. coli O157

Non-STEC E. coli O157 strains**

[†]Rare or less common eae intimin carrier organisms.

‡On-panel target.

§In vitro testing of Campylobacter lari and Campylobacter helveticus strains at high concentration confirmed potential cross-reaction of these Campylobacter species with the QIAstat-Gastrointestinal Panel 2 assay.

¶Rare or less common Stx toxins producers.

**E. coli 0157 will only be called when there is a positive amplification for the E. coli (STEC) design according to the calling algorithm. An infrequent case of an E. coli (STEC) and an E. coli O157 co-infection will not be differentiated from a single infection caused by an STEC O157:H7 strain.

Inclusivity (Analytical Reactivity)

Analytical Reactivity (Inclusivity) was evaluated with gastrointestinal pathogen isolates/strains that were selected based on clinical relevance and genetic, temporal and geographical diversity. Based on in vitro (wet) testing and in silico analysis, the QIAstat-Dx Gastrointestinal Panel 2 primers and probes are specific and inclusive for clinically prevalent and relevant strains for each pathogen tested.

In vitro (Wet) testing

QIAstat-Dx Gastrointestinal Panel 2 is inclusive for 100% (143 out of 143) of the pathogen strains tested in vitro. Most pathogen strains evaluated in wet testing (133/143) were detected at \leq 3-fold of the corresponding LoD reference strain. (Table 10. Inclusivity test results for all the pathogens tested with the QIAstat-Dx Gastrointestinal Panel 2 Assay. LoD reference strain for every pathogen is written in bold.).

^{*}Note that these potential cross-reactions affect designs with target genes responsible of the pathogenicity of the corresponding QIAstat-Dx Gastrointestinal Panel 2 target pathogens which can be acquired within species in a known biological process in bacteria called horizontal gene transfer.

Table 10. Inclusivity test results for all the pathogens tested with the QIAstat- Dx Gastrointestinal

Panel 2 Assay.

LoD reference strain for every pathogen is written in bold.

Table 10. a. Inclusivity test results for Campylobacter strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------|---------------------------------------|-------------------------------|-------------|------------|--------------|
| Campylobacter | Campylobacter coli | 76-GA2 [LMG 21266] | ATCC | 43478* | 1x LoD |
| | Campylobacter coli | Z293 | ZeptoMetrix | 0804272 | 1x LoD |
| | Campylobacter coli | CIP 7080 [1407, CIP 70.80] | ATCC | 33559* | 3x LoD |
| | Campylobacter jejuni | Z086 | ZeptoMetrix | 0801650* | 1x LoD |
| | Campylobacter jejuni | subsp. jejuni RM3193 | ATCC | BAA-1234* | 0.1x LoD |
| | Campylobacter jejuni subsp. jejuni | O:19 HL7; D3180 | ATCC | BAA-218 | 0.1x LoD |
| | Campylobacter jejuni subsp. jejuni | AS-83-79 | ATCC | 33291 | 0.1x LoD |
| | Campylobacter jejuni subsp. doylei | NCTC 11951 | ATCC | 49349 | 0.1x LoD |
| | Campylobacter upsaliensis | NCTC 11541 | ZeptoMetrix | 0801999* | 1x LoD |
| | Campylobacter upsaliensis | RM 3195 (1994) | ATCC | BAA-1059* | 0.3x LoD |
| | Campylobacter upsaliensis | NCTC 11541 [C231] | ATCC | 43954 | 1x LoD |

^{*} Strain tested during LoD verification study.

Table 10b. Inclusivity test results for Clostridium difficile strains.

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|------------------------------------|--------------------------|-----------------------------------|-------------|------------|--------------|
| Clostridium difficile toxin A/B | Clostridium difficile | (90556-M6S) Toxinotype 0 A+ B+ | ATCC | 9689* | 1x LoD |
| | Clostridium difficile | NAP1, toxinotype IIIb A+B+ | ATCC | BAA-1805 | 1x LoD |
| | Clostridium difficile | 5325, toxinotype V A+B+ | ATCC | BAA-1875 | 1x LoD |
| | Clostridium difficile | 1470, toxinotype VIII A-B+ | ATCC | 43598 | 1x LoD |
| | Clostridium difficile | toxinotype XII A+B+ | ATCC | BAA-1812 | 1x LoD |
| | Clostridium difficile | toxinotype XXII A+B (unknown) | ATCC | BAA-1814 | 1x LoD |
| | Clostridium difficile | NAP1A, toxinotype III A+B+ | ATCC | 0801619* | 0.1x LoD |
| | Clostridium difficile | NAP1, toxinotype III A+B+ | ZeptoMetrix | 0801620 | 3x LoD |

^{*}Strain tested during LoD verification study.

Table 10c. Inclusivity test results for Plesiomonas shigelloides strains.

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-----------------------------|-----------------------------|---|-------------|------------|--------------|
| Plesiomonas shigelloides | Plesiomonas shigelloides | Z130 | ZeptoMetrix | 0801899* | 1x LoD |
| | Plesiomonas shigelloides | GNI 14 | ATCC | 51903 | 1x LoD |
| | Plesiomonas shigelloides | CDC 3085-55 [Bader M51, NCIB 9242, NCTC 10360, RH 798] | ATCC | 14029* | 0.3x LoD |

^{*}Strain tested during LoD verification study.

Table 10d. Inclusivity test results for Salmonella strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|------------------------|--|-------------|------------|--------------|
| Salmonella | Salmonella enterica | Serovar Typhimurium Z005 | ZeptoMetrix | 0801437* | 1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Bareilly | NCTC | NC05745 | 1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar typhi, Z152 | ZeptoMetrix | 0801933 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Enteridis, CDC K-1891 [ATCC 25928] | ATCC | 13076 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Infantis, MZ1479 [SARB27] | ATCC | BAA-1675 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Montevideo, G4639 | ATCC | BAA-710 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Javiana | NCTC | NC06495 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Thompson | NCTC | NC08496 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Saintpaul | ATCC | 9712 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Berta | NCTC | NC05770 | 0.1x LoD |
| | Salmonella enterica | Subps. Salame, II NCTC 10310 [JT945, SS140/61] | ATCC | 700151 | 0.1x LoD |
| | Salmonella enterica | Subps. diarizonae IIIb, 62 | ATCC | 29934 | 0.1x LoD |
| | Salmonella enterica | Subps. houtenae IV, CIP 82.32 [264.66] | ATCC | 43974 | 0.1x LoD |
| | Salmonella enterica | Subps. Indica VI, CIP 102501 [F. Kauffmann 1240] | ATCC | 43976 | 0.1x LoD |
| | | | | | |

Table 10d. Inclusivity test results for Salmonella strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|------------------------|--|----------|------------|--------------|
| | Salmonella enterica | Subsp. Enterica, serovar Agona, CDC 873 [CDC 1111-61] | ATCC | 51957 | 0.1x LoD |
| Salmonella | Salmonella enterica | Subsp. Enterica, serovar Muenchen, 54 | ATCC | 8388 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Oranienburg, E1093 | ATCC | 9239 | 0.1x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Paratyphi B var. Java, CDC 5 | ATCC | 51962 | 0.1x LoD |
| | Salmonella bongori | CIP 82.33 [1224.72] | ATCC | 43975 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Choleraesius, NCTC 5735 [1348, K.34] | ATCC | 13312* | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Newport, C487-69 | ATCC | 27869 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, 4, 5, 12:7:-, serovar Typhimurium | NCTC | NC13952 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Braenderup | ATCC | 700136 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Anatum | NCTC | NC05779 | 0.3x LoD |
| | Salmonella enterica | Subps. arizonae Illa, NCTC 7311 [CDAI 426] | ATCC | 700156 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Heidelberg, [16] | ATCC | 8326 | 0.3x LoD |
| | Salmonella enterica | Subsp. Enterica, serovar Mississippi, CDC 2012K-0487 | ATCC | BAA-2739 | 0.3x LoD |

^{*} Strain tested during LoD verification study.

Table 10e. Inclusivity test results for Vibrio cholerae strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|--------------------|--------------------------------------|-------------|------------|--------------|
| Vibrio cholerae | Vibrio cholerae | Z133; non-toxigenic | ZeptoMetrix | 801902* | 1x LoD |
| | Vibrio cholerae | Pacini 1854; NCTC 8021, O:1 Ogawa | CECT | 514 | 1x LoD |
| | Vibrio cholerae | Z132; toxigenic | ZeptoMetrix | 0801901* | 0.3x LoD |

^{*} Strain tested during LoD verification study.

Table 10f. Inclusivity test results for Vibrio parahaemolyticus strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------------|----------------------------|-----------------------------------|-------------|------------|--------------|
| Vibrio parahaemolyticus | Vibrio parahaemolyticus | EB101 [P. Baumann 113] (Japan) | ATCC | 17802* | 1x LoD |
| | Vibrio parahaemolyticus | VP250,O1:KUT | ATCC | BAA-242 | 1x LoD |
| | Vibrio parahaemolyticus | 205 [9302] | ATCC | 33846 | 3x LoD |
| | Vibrio parahaemolyticus | Z134 | ZeptoMetrix | 0801903* | 0.3x LoD |

^{*} Strain tested during LoD verification study.

Table 10g. Inclusivity test results for Vibrio vulnificus strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------|-------------------|---------------------------|-------------|------------|-----------|
| Vibrio vulnificus | Vibrio vulnificus | 324 [CDC B9629] | ATCC | 27562* | 1x LoD |
| | Vibrio vulnificus | 329 [CDC B3547],Biotype 2 | ATCC | 33817* | 1x LoD |
| | Vibrio vulnificus | Z473 | ZeptoMetrix | 0804349 | 3x LoD |

^{*} Strain tested during LoD verification study.

Table 10h. Inclusivity test results for Yersinia enterocolitica strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------------|----------------------------|--|-------------|------------|--------------|
| Yersinia enterocolitica | Yersinia enterocolitica | Z036 | ZeptoMetrix | 801734* | 1x LoD |
| | Yersinia enterocolitica | NTCC 11175, Biotype 4, serotype 3 (O:3) | ATCC | 700822* | 1x LoD |
| | Yersinia enterocolitica | 33114 [CCUG 11291, CCUG 12369, CIP 80.27, DSM 4780, LMG 7899, NCTC 12982], Biovar 1, O:8 | ATCC | 9610 | 1x LoD |
| | Yersinia enterocolitica | O:9 | ATCC | 55075 | 3x LoD |

^{*} Strain tested during LoD verification study.

Table 10i. Inclusivity test results for Enteroaggregative E. coli (EAEC) strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------------------------|-------------------------------------|--|---------------|--|--------------|
| Enteroaggregative E. coli (EAEC) | Enteroaggregative E. coli (EAEC) | 92.0147 | ZeptoMetrix | 0801919* | 1x LoD |
| | Enteroaggregative E. coli (EAEC) | CDC3250-76, O111a, 111b: K58:H21, CVD432+, aggR+, stx1-, stx2-, eae- | ATCC | 29552* | 1x LoD |
| | Enteroaggregative E. coli (EAEC) | - | Vall d'Hebrón | Clinical sample; VH 529140369015 | 3x LoD |

^{*} Strain tested during LoD verification study.

Table 10j. Inclusivity test results for Enteropathogenic E. coli (EPEC) strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|---------------------------------|--|-----------------------------|-------------|------------|--------------|
| Enteropathogenic E. coli (EPEC) | Enteropathogenic <i>E.</i> coli (EPEC) | O111:NM | ZeptoMetrix | 0801747* | 1x LoD |
| | Enteropathogenic <i>E.</i> coli (EPEC) | 7.1493,O84:H28 | ZeptoMetrix | 0801938* | 1x LoD |
| | Enteropathogenic E. coli (EPEC) | Stoke W,O111:K58 (B4):H- | ATCC | 33780 | 1x LoD |

^{*} Strain tested during LoD verification study.

Table 10k. Inclusivity test results for Enterotoxigenic E. coli (ETEC) strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|--------------------------------------|---|-------------------------------------|--------------------|------------|--------------|
| Enterotoxigenic E. coli (ETEC) It/st | Enterotoxigenic <i>E.</i> coli (ETEC) lt/st | ST+, LT+ | ZeptoMetrix | 0801624* | 1x LoD |
| | Enterotoxigenic E. coli (ETEC) lt/st | H10407,O78:H11,LT (+)/ctx A11(+) | ATCC | 35401* | 0.3x LoD |
| | Enterotoxigenic <i>E.</i> coli (ETEC) lt/st | O27:H7,ST (+)/ LT (-) | SSI Diagnostica | 82173 | 0.1x LoD |
| | Enterotoxigenic E. coli (ETEC) lt/st | O115:H15,ST (+)/LT (-) | SSI Diagnostica | 82174 | 3x LoD |
| | Enterotoxigenic <i>E.</i> coli (ETEC) lt/st | O169:H-,ST (-)/LT (+) | SSI Diagnostica | 82172 | 10x LoD |

^{*} Strain tested during LoD verification study.

Table 101. Inclusivity test results for Enteroinvasive E. coli (EIEC)/Shigella strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|---|--------------------------------------|---------------------------------|--------------------|------------|--------------|
| Enteroinvasive <i>E. coli</i> (EIEC)/ Shigella | Enteroinvasive <i>E. coli</i> (EIEC) | CDC EDL 1282, O29:NM | ATCC | 43892* | 1x LoD |
| | Enteroinvasive <i>E. coli</i> (EIEC) | O172:H- | SSI Diagnostica | 82171 | 3x LoD |
| | Shigella boydii | Z004 | ATCC | 25931* | 1x LoD |
| | Shigella boydii (Serogroup C) | Z131 | ZeptoMetrix | 0801900 | 1x LoD |
| | Shigella flexneri (Serogroup B) | AMC 43-G-68 [EVL 82, M134] | ATCC | 9199 | 1x LoD |
| | Shigella flexneri (Serogroup B) | Z046 | ZeptoMetrix | 0801757 | 1x LoD |
| | Shigella sonnei (Serogroup D) | WRAIR I virulent | ATCC | 29930 | 1x LoD |
| | Shigella sonnei (Serogroup D) | Z004 | ZeptoMetrix | 801627 | 3x LoD |
| | Shigella boydii (Serogroup C) | AMC 43-G-58 [M44 (Type 170)] | ATCC | 9207 | 10x LoD |

^{*} Strain tested during LoD verification study

Table 10m. Inclusivity test results for Shiga-like toxin-producing E. coli (STEC)(stx1-carrier strains)

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|---|---|--|-----------------|------------|--------------|
| Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O1 <i>57</i> :H7; EDL933 | ZeptoMetrix | 0801622* | 1 x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O26:H4,stx1 (+) | ZeptoMetrix | 0801748* | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O22:H8,stx1c (+), stx2b (+) | SSI Diagnostica | 91350 | 1 x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O8,stx1d (+) | SSI Diagnostica | 91349 | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | Reference ATCC 35150 (EDL 931), O157:H7,stx1 (+), stx2 (+) | Microbiologics | 617 | 1x LoD |
| Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | Reference CDC 00-3039, O45:H2,unknown | Microbiologics | 1098 | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O103:H2,stx1 (+) | SSI Diagnostica | 82170 | 3x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx1 | O128ac:H-,stx2f (+) | SSI Diagnostica | 91355 | 10x LoD |

^{*} Strain tested during LoD verification study

Table 10n. Inclusivity test results for Shiga-like toxin-producing E. coli (STEC) (stx2-carrier strains)

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|---|---|---|-----------------|------------|--------------|
| Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O157:H7; EDL933 | ZeptoMetrix | 0801622* | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O22:H8,stx1c (+), stx2b (+) | SSI Diagnostica | 91350 | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O26:H11,stx2a (+) | SSI Diagnostica | 95211 | 1x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O101:K32:H-,stx2e (+) | SSI Diagnostica | 91354 | 0.3x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | Reference ATCC 35150 (EDL 931),O157:H7,stx1 (+), stx2 (+) | Microbiologics | 617 | 3x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O92,O107:K+:H48, stx2d (+) | SSI Diagnostica | 91352 | 10x LoD |
| | Shiga-like toxin producing <i>E. coli</i> (STEC) - stx2 | O128ac:H-,stx2f (+) | SSI Diagnostica | 91355 | 10x LoD |

^{*} Strain tested during LoD verification study

Table 10o. Inclusivity test results for Shiga-like toxin producing E. coli (STEC) stx1/stx2 O157 strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalogue ID | Times LoD |
|---|--|--|-----------------|--------------|--------------|
| Shiga-like toxin producing <i>E. coli</i> (STEC) O157 | Shiga-like toxin producing <i>E.</i> coli (STEC) - O157 | O157:H7; EDL933 | ZeptoMetrix | 0801622* | 1x LoD |
| | Shiga-like toxin producing <i>E.</i> coli (STEC) O157 | O128ac:H-,stx2f (+) | SSI Diagnostica | 91355† | 1x LoD |
| | Shiga-like toxin producing <i>E.</i> <i>coli</i> (STEC) O1 <i>57</i> | Reference ATCC 35150 (EDL 931), O157:H7, stx1 (+), stx2 (+) | Microbiologics | 617 | 1x LoD |

^{*} Strain tested during LoD verification study.

[†] The *E. coli* strain 91355 from SSI Diagnostica is reported as following in its catalog: vtx2f+, eae+. However, it was found to amplify for *E. coli* O157 in both QIAstat-Dx and FilmArray devices

Table 10p. Inclusivity test results for Cryptosporidium strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------|--------------------------------|-----------------|------------------------|---------------------------------|--------------|
| Cryptosporidium | Cryptosporidium parvum | lowa isolate | Waterborne | P102C* | 1x LoD |
| | Cryptosporidium hominis | n/a | Public Health Wales | Clinical sample; UKM 84* | 0.01x LoD |
| | Cryptosporidium parvum | - | ATCC | PRA-67DQ (isolated genomic DNA) | <0.01 LoD |
| | Cryptosporidium meleagridis | - | Public Health Wales | Clinical sample; UKMEL 14 | <0.01 LoD |
| | Cryptosporidium meleagridis | - | Public Health Wales | Clinical sample; UKMEL 14 | <0.01 LoD |

^{*} Strain tested during LoD verification study

Table 10q. Inclusivity test results for Cyclospora cayetanensis strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalogue ID | Times LoD |
|-------------------------|-------------------------|--------|-----------------|--------------|-----------|
| Cyclospora cayetanensis | Cyclospora cayetanensis | n/a | Clinical sample | LAC2825* | 1 x LoD |
| | Cyclospora cayetanensis | n/a | Clinical sample | LAC2827* | 1x LoD |
| | Cyclospora cayetanensis | - | ATCC | PRA-3000SD | 1x LoD |

^{*} Strain tested during LoD verification study

Table 10r. Inclusivity test results for Entamoeba histolytica strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalogue ID | Times LoD |
|--------------------------|--------------------------|---------------------------------|------------------|------------------|--------------|
| Entamoeba histolytica | Entamoeba histolytica | HM-1:IMSS (Mexico City 1967) | ATCC | 30459* | 1x LoD |
| | Entamoeba histolytica | HK-9 (Korea) | ATCC | 30015* | 1x LoD |
| | Entamoeba histolytica | - | Vall d'Hebrón | Clinical sample; | 1x LoD |

^{*} Strain tested during LoD verification study

Table 10s. Inclusivity test results for Giardia lamblia strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------|----------------------|----------------------------------|------------|------------|-----------|
| Giardia lamblia | Giardia lamblia | Portland -1 (Portland, OR, 1971) | ATCC | 30888* | 1 x LoD |
| | Giardia lamblia | WB (Bethesda, MD, 1979) | ATCC | 30957* | 1x LoD |
| | Giardia intestinalis | H3 isolate | Waterborne | P101 | 1 x LoD |

^{*} Strain tested during LoD verification study.

Table 10t. Inclusivity test results for Adenovirus F40/F41 targets

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-----------------------|-----------------------------|----------------------|-------------|------------|--------------|
| Adenovirus F40/F41 | Human Adenovirus F41 | Tak | ZeptoMetrix | 0810085CF* | 1x LoD |
| | Human Adenovirus F41 | Tak (73-3544) | ATCC | VR-930 | 10x LoD |
| | Human Adenovirus F40 | Dugan [79- 18025] | ATCC | VR-931 | 10x LoD |
| | Human Adenovirus Type 40 | Dugan | ZeptoMetrix | 0810084CF* | 3x LoD |

^{*} Strain tested during LoD verification study

Table 10u. Inclusivity test results for Astrovirus strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|---------------------|-----------------------|-----------------------------|-------------------------------|--------------|
| Astrovirus | Human Astrovirus | ERE IID 2371 (type 8) | ZeptoMetrix | 0810277CF* | 1x LoD |
| | Human Astrovirus | HAstV-1 | Universitat de Barcelona | Clinical sample; 160521599 | 1x LoD |
| | Human Astrovirus | ERE IID 2868 (type 4) | Zepto/Metrix | 0810276CF* | 1x LoD |
| | Human Astrovirus | HAstV-3 | Universitat de Barcelona | Clinical sample; 151601306 | 1x LoD |

^{*} Strain tested during LoD verification study.

Table 10v. Inclusivity test results for Norovirus GI/GII strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|--------------------------------|----------------------|-----------------------------------|-------------------------------|--------------|
| Norovirus GI/GII | Human Norovirus Genogroup 1 | Recombinant Gl. 1 | ZeptoMetrix | 0810086CF* | 1x LoD |
| | Human Norovirus Genogroup 1 | - | Indiana University Health | Clinical sample; IU3156 | 1x LoD |
| | Human Norovirus Genogroup 1 | - | Indiana University Health | Clinical sample; IU3220 | 1x LoD |
| | Human Norovirus Genogroup 1 | - | TriCore Reference Laboratories | Clinical sample; TC4274 | 3x LoD |
| | Human Norovirus Genogroup 2 | Recombinant GII.4 | ZeptoMetrix | 0810087CF* | 1x LoD |
| | Human Norovirus Genogroup 2 | GII.2 | Vall d'Hebrón | Clinical sample; 198058327 | 1x LoD |
| | Human Norovirus Genogroup 2 | GII.4 | Universitat de Barcelona | Clinical sample; N26.2TA | 1x LoD |
| | Human Norovirus Genogroup 2 | - | Lacny Hospital | Clinical sample; LAC2019 | 1x LoD |
| | Human Norovirus Genogroup 2 | - | Nationwide Children's Hospital | Clinical sample; NWC6063 | 1x LoD |
| | Human Norovirus Genogroup 2 | GII.6 | QIAGEN Barcelona (STAT-Dx) | Clinical sample; GI 12 | 3x LoD |
| | Human Norovirus Genogroup 2 | - | Lacny Hospital | Clinical sample; LAC2133 | 10x LoD |
| | Human Norovirus Genogroup 2 | - | Lacny Hospital | Clinical sample; LAC2074 | 10x LoD |

^{*} Strain tested during LoD verification study.

Table 10w. Inclusivity test results for Rotavirus A strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|-------------------|-------------------|-----------------|-------------|------------|-----------|
| Rotavirus A | Human Rotavirus A | 69M | ZeptoMetrix | 0810280CF* | 1x LoD |
| | Human Rotavirus A | Wa, G1P1A[8] | ZeptoMetrix | 0810041CF* | 1x LoD |
| | Human Rotavirus A | DS-1, G2P1B[4]) | ATCC | VR-2550 | 1x LoD |
| | Human Rotavirus A | Va70 | ZeptoMetrix | 0810281CF | 1x LoD |
| | Human Rotavirus A | RRV | ZeptoMetrix | 0810530CF | 10x LoD |

^{*} Strain tested during LoD verification study

Table 10x. Inclusivity test results for Sapovirus strains

| QIAstat-Dx target | Pathogen | Strain | Supplier | Catalog ID | Times LoD |
|----------------------|---------------------------------|--------|-----------------------------|--------------------------------|--------------|
| Sapovirus | Human Sapovirus Genogroup I | - | QIAGEN Barcelona | Clinical sample; GI-88* | 1x LoD |
| | Human Sapovirus Genogroup V | n/a | Universitat Barcelona | Clinical Sample; 160523351* | 1x LoD |
| | Human Sapovirus Genogroup I | Gl.1 | Universitat de Barcelona | Clinical sample; 171016324 | 1x LoD |
| | Human Sapovirus Genogroup II | GII.3 | Universitat de Barcelona | Clinical sample; 215512 | 1x LoD |

^{*} Strain tested during LoD verification study.

In silico analysis

In silico analysis of potential reactivity showed that the following organisms (including species, subspecies, subtypes, serotypes or serovars) are predicted to be detected with the QIAstat-Dx Gastrointestinal Panel 2 (Organisms with predicted reactivity based on in silico analysis).

Table 11. Organisms with predicted reactivity based on in silico analysis

QIAstat-Dx GI Panel

| 2 Target | Organisms with predicted reactivity (species, subspecies, subtypes, serotypes or serovars) |
|--|--|
| | Bacteria |
| | |
| | |
| Clostridium difficile | Clostridium difficile (including ribotypes 01 and 17 and strains BI1, BI9, NAP1, SD1, SD2, M68, M120) |
| | |
| | |
| | |
| | |
| | |
| Plesiomonas shigelloides | Plesiomonas shigelloides (e.g. strains NCTC10360, ATCC 14029T, R4605035) |
| | |
| Vibrio cholerae | Vibrio cholerae (including serotypes O:1 and non-O:1 (O:37) and biovars El Tor, Bengal) |
| Vibrio | Vibrio parahaemolyticus |
| parahaemolyticus | |
| | |
| Vibrio vulnificus | Vibrio vulnificus |
| Yersinia enterocolitica | Yersinia enterocolitica, Yersinia enterocolitica subsp. palearctica, Yersinia enterocolitica subsp. enterocolitica |
| | |
| Enteroaggregative <i>E.</i> | |
| coli (EAEC) | O25:H4, O86:H2, O86:HND, OUT:H4, OUT:HND) |
| Enteroinvasive E. coli (EIEC)/Shigella | Enteroinvasive <i>E. coli</i> (EIEC), Escherichia coli sp., Shigella flexneri, Shigella dysenteriae, Shigella boydii, Shigella sonnei |
| con (LiLC)/ onigelia | onigona soyan, onigona sonner |
| Enteropathogenic <i>E.</i> | Enteropathogenic E. coli (EPEC) (e.g. including serotypes OUT: HND, OUT:H6, OUT:H34, |
| coli (EPEC) | OUT:H21, O55:H7, O119:HNM, O117) |
| | Other eae-carriers bacteria: some Shiga-like toxin-producing <i>E. coli</i> (STEC), STEC 0157:H7 |

and few Shigella boydii strains

Table 11. Organisms with predicted reactivity based on in silico analysis (continued)

| QIAstat-Dx GI Panel 2 Target | Organisms with predicted reactivity (species, subspecies, subtypes, serotypes or serovars) |
|--|---|
| Enterotoxigenic <i>E.</i> coli (ETEC) | Enterotoxigenic <i>E. coli</i> (ETEC) (including H10407 and E24377A strains and serotypes O169:H41, O25:H42, O148:H28, O6:H16) |
| Shiga-like toxin- producing <i>E. coli</i> (STEC) - stx1 | Shiga-like toxin-producing <i>E. coli</i> (STEC) (including non-O157 serotypes O111:NM, O111:H-, O26:H11, O145:NM, O145:H28, O45:H2, O26:H11, ONT:NM, and including STEC O157 serotypes O157:H7) |
| | Stx1 toxin subtypes predicted to be detected include stx1a, stx1c and stx1d |
| | Other stx-carriers bacteria: Shigella sonnei, Shigella dysenteriae |
| Shiga-like toxin- producing <i>E. coli</i> (STEC) - stx2 | Shiga-like toxin-producing <i>E. coli</i> (STEC) (including non-0157 serotypes O111:NM, O104:H4, O111:H-, O26:H11, O121:H19, O145:H34, O113:H21, ONT:H-, O128:H2, OUT:HNM, O124:HNM and including STEC O157 serotypes O157:H7, O157:NM) Stx2 toxin subtypes predicted to be detected include stx2a, stx2b, stx2c, stx2d, stx2e, stx2f, stx2g and stx2h. |
| Shiga-like toxin- producing <i>E. coli</i> (STEC) O1 <i>57</i> | Escherichia coli O157 including: STEC O157:H7 strains (e.g. EDL933) and <i>E. coli</i> O157: non-H7 groups including Non-Shiga-toxigenic <i>E. coli</i> O157 bacteria (e.g. serotype O157:H45) Other bacteria with O157 O-antigen: Escherichia fergusonii O157 |

Parasites

| Cyclospora cayetanensis | Cyclospora cayetanensis (including strains LG, CY9, NP20 and NP21) | |
|----------------------------|--|--|
| Entamoeba | Entamoeba histolytica (e.g. strains HM-1: IMSS, EHMfas1, HK-9) | |
| | | |
| Giardia lamblia | Giardia lamblia (aka Giardia duodenalis, Giardia intestinalis)f | |
| Viruses | | |
| Adenovirus | Human Adenovirus F 40/41 | |
| Astrovirus | Human Astrovirus (including types 1, 2, 3, 4, 5, 6, 7, 8) | |

Table 11. Organisms with predicted reactivity based on in silico analysis (continued)

| QIAstat-Dx GI Panel 2 Target | Organisms with predicted reactivity (species, subspecies, subtypes, serotypes or serovars) |
|---------------------------------|---|
| Norovirus GI/GII | Norovirus genogroup II genotypes: GII.1, GII.2, GII.3, GII.4, GII.4_Sydney 2012, GII.P4_ New Orleans 2009, GII.4_DenHaag, GII.4_Hong Kong, GII.5, GII.6, GII.7, GII.8, GII.10, GII.12, GII.13, GII.17, GII.21. |
| | Norovirus genogroup I genotypes: Gl.1, Gl.3, Gl.4, Gl.5, Gl.6, Gl.7, Gl.8, Gl.9. |
| Rotavirus | Rotavirus A (including strains Wa, ST3, 69M, DS-1, RVA and serotypes G1P[8], G12P[6], G2P[4], G3P[6], G4P[6], G6P[6], G8P[8], G9P[19]) |
| Sapovirus | Genogroups GI (including genotypes GI.1, GI.2, GI.3, GI.4, GI.6), GII (including genotypes GII.1, GII.2, GII.3, GII.4, GII.5, GII.6), GIV (including genotype GIV.1) and GV (including genotypes GV.1). |

Interfering Substances

The effect of potentially interfering substances on the detectability of the QIAstat-Dx Gastrointestinal Panel 2 organisms was evaluated. Forty-three (43) potentially interfering substances were spiked into the sample mixes at a level predicted to be above the concentration of the substance likely to be found in stool specimens. Each organism was tested at 3x LoD and testing was performed in triplicates. Endogenous substances such as human whole blood, human genomic DNA and several pathogens were tested alongside exogenous substances like antibiotics, other gastrointestinal-related medications and different technique-specific substances.

For the vast majority of substances tested, no inhibition was observed, with the exceptions of mucin from bovine submaxillary, Human genomic DNA, bisacodyl, calcium carbonate, nonoxynol-9 and Rotavirus reassortants, that may cause inhibition at high concentration.

Mucin from bovine submaxillary was found to interfere with the detection of *Vibrio cholerae*, EAEC and *Entamoeba* at concentrations above 2.5% w/v.

Human genomic DNA was found to interfere with the detection of *E. coli* O157 and *Entamoeba* at concentrations above $5 \, \mu g/mL$.

Bisacodyl was found to interfere with the detection of EAEC at concentrations above 0.15% w/v.

Calcium carbonate was found to interfere with the detection of all the QIAstat- Dx Gastrointestinal Panel 2 targets at concentrations above 0.5% w/v.

Nonoxynol-9 was found to interfere with the detection of Entamoeba at concentrations above 0.02% v/v.

Rotavirus reassortants WC3:2-5, R574(9) and WI79-4,9 used in Rotavirus A vaccines were predicted to be reactive with Rotavirus A in the QIAstat- Dx Gastrointestinal Panel 2. Final concentrations without observable interfering effects on the detection of targets at 3x LoD concentration for WC3:2-5, R574(9) and WI79-4,9 were 8.89x10⁻⁵ TCID₅₀/ml and 1.10 PFU/ml, respectively (see Table 12) for other concentrations tested.

Competitive interference was tested in a subset of pathogens. No interference was observed when evaluating competitive interference by target pathogens when two QIAstat-Dx Gastrointestinal Panel target pathogens were tested by spiking samples with one pathogen target at 3x LoD and one at 50x LoD. Results from the pathogen targets tested are provided in Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration.

Results from the 43 interfering substances that could be present or introduced in a stool specimen are provided in Table 12.

Table 12. Final highest concentration without observable inhibitory effect

| Substance tested | Concentration tested | Result |
|---|------------------------------|-----------------|
| Endogenous substances | | |
| Bovine and ovine bile | 12% w/v | No Interference |
| Cholesterol | 1.5% w/v | No Interference |
| Fatty acids (palmitic acid) | 0.2% w/v | No Interference |
| Fatty acids (stearic acid) | 0.4% w/v | No Interference |
| Human genomic DNA | 20 μg/ml | Interference |
| | 10 μg/ml | Interference |
| | 5 μg/ml | No Interference |
| Human stool (overfill of Cary Blair vial) | 300 mg/ml | No Interference |
| Human urine | 50% v/v | No Interference |
| Human whole blood with Na Citrate | 40% v/v | No Interference |
| Mucin from bovine submaxillary | 5% w/v | Interference |
| | 2.5% w/v | No Interference |
| Triglycerides | 5% w/v | No Interference |
| Non-target microorganisms | | |
| Aeromonas hydrophila | 1 x 106 units/ml | No Interference |
| Bacteroides vulgatus | 1 x 106 units/ml | No Interference |
| Bifidobacterium bifidum | 1 x 106 units/ml | No Interference |
| Enterovirus Species D, Serotype EV-D68 | 1 x 10 ⁵ units/ml | No Interference |
| Non-pathogenic E. coli | 1 x 106 units/ml | No Interference |
| Helicobacter pylori | 1 x 106 units/ml | No Interference |

Table 12. Final highest concentration without observable inhibitory effect (continued)

| Substance tested | Concentration tested | Result |
|--|------------------------------|-----------------|
| Saccharomyces cerevisiae (deposited as S. boulardii) | 1 x 10 ⁵ units/ml | No Interference |
| Exogenous substances | | |
| Bacitracin | 250U/ml | No Interference |
| Bisacodyl | 0.3% w/v | Interference |
| | 0.15% w/v | No Interference |
| Bismuth subsalicylate | 0.35% w/v | No Interference |
| Calcium carbonate (TUMS® Extra Strength 750) | 5%w/v | Interference |
| | 0.5% w/v | No Interference |
| Docusate sodium | 2.5% w/v | No Interference |
| Doxycycline hydrochloride | 0.05% w/v | No Interference |
| Glycerin | 50% v/v | No Interference |
| Hydrocortisone | 0.5% w/v | No Interference |
| Loperamide hydrochloride | 0.078% w/v | No Interference |
| Magnesium hydroxide | 0.1% w/v | No Interference |
| Metronidazole | 1.5% w/v | No Interference |
| Mineral oil | 50% v/v | No Interference |
| Naproxen sodium | 0.7% w/v | No Interference |

Table 12. Final highest concentration without observable inhibitory effect (continued)

| Substance tested | Concentration tested | Result |
|--|-------------------------------|-----------------|
| Nonoxynol-9 | 1.2% v/v | Interference |
| | 0.6% v/v | Interference |
| | 0.3% v/v | Interference |
| | 0.15% v/v | Interference |
| | 0.075% v/v | Interference |
| | 0.02% v/v | No Interference |
| Nystatin | 10000 USP units/ml | No Interference |
| Phenylephrine hydrochloride | 0.075% w/v | No Interference |
| Sodium phosphate | 5% w/v | No Interference |
| Vaccine components | | |
| Rotavirus reassortant WC3:2-5, R574(9) - VR 2195 | 8.89 x 10-3 TCID50/ml | Interference |
| | 8.89 x 10-4 TCID50/ml | Interference |
| | 8.89 x 10-5 TCID50/ml | No Interference |
| Rotavirus reassortant WI79-4,9 - VR 2415 | 1.10 x 102 pfu/ml | Interference |
| | 1.10 x 10 ¹ pfu/ml | Interference |
| | 1.10 pfu/ml | No Interference |
| Technique-specific Substances | | |
| Bleach | 0.5% v/v | No Interference |
| | | |
| Ethanol | 0.2% v/v | No Interference |

Table 12. Final highest concentration without observable inhibitory effect (continued)

| Substance tested | Concentration tested | Result |
|-----------------------------------|---------------------------|-----------------|
| Fecal Opti-Swab Cary-Blair Medium | 100% | No Interference |
| PurSafe® DNA/RNA Preservative | 100% | No Interference |
| Para-Pak C&S spoon | 1 spoon/2ml Cary Blair | No Interference |
| Sigma transwab | 1 swab/2ml Cary Blair | No Interference |

Table 13. QIAstat-Dx Gastrointestinal Panel 2 results for competitive interference

| Sample Mix | Target | Final concentration tested x LoD | Co-infection detected | |
|------------------------------|---------------------------------|----------------------------------|-----------------------|--|
| Norovirus 50x - Rotavirus 3x | Norovirus GI/GII | 50x | Yes | |
| | Rotavirus A | 3x | | |
| Norovirus 3x - Rotavirus 50x | Norovirus GI/GII | 3x | Yes | |
| | Rotavirus A | 50x | | |
| Giardia 50x - Adenovirus 3x | Giardia lamblia | 50x | Yes | |
| | Adenovirus F40/F41 | 3x | | |
| Adenovirus 50x - Giardia 3x | Giardia lamblia | 3x | Yes | |
| | Adenovirus F40/F41 | 50x | | |
| Norovirus 50x -C.diff 3x | Norovirus GII | 50x | Yes | |
| | Clostridium difficile toxin A/B | 3x | | |
| Norovirus 3x - C.diff 50x | Norovirus GII | 3x | Yes | |
| | Clostridium difficile toxin A/B | 50x | | |
| EPEC 50x - EAEC 3x | EPEC | 50x | Yes | |
| | EAEC | 3x | | |

Table 13. QIAstat-Dx Gastrointestinal Panel 2 results for competitive interference (continued)

| Sample Mix | Target | Final concentration tested x LoD | Co-infection detected |
|----------------------|---------------------------------|----------------------------------|-----------------------|
| EPEC 3x - EAEC 50x | EPEC | 3x | Yes |
| | EAEC | 50x | |
| EPEC 50x - C.diff 3x | EPEC | 50x | Yes |
| | Clostridium difficile toxin A/B | 3x | |
| EPEC 3x - C.diff 50x | EPEC | 3x | Yes |
| | Clostridium difficile toxin A/B | 50x | |
| EPEC 50x - ETEC 3x | EPEC | 50x | Yes |
| | ETEC | 3x | |
| EPEC 3x - ETEC 50x | EPEC | 3x | Yes |
| | ETEC | 50x | |
| ETEC 50x - EIEC 3x | ETEC | 50x | Yes |
| | EIEC/ Shigella | 3x | |
| ETEC 3x - EIEC 50x | ETEC | 3x | Yes |
| | EIEC/ Shigella | 50x | |
| | | | |

Carryover

A carryover study was performed to evaluate the potential occurrence of cross-contamination between consecutive runs when using the QIAstat-Dx Gastrointestinal Panel 2 on the QIAstat-Dx Analyzer 1.0.

Pathogen samples of stool sample matrix, with alternating high-positive (10 5 - 10 6 organism/ml) and negative samples, were conducted on two QIAstat- Dx Analyzer 1.0 instruments.

No carryover between samples was observed in the QlAstat-Dx Gastrointestinal Panel 2, demonstrating that the system design and recommended sample handling and testing practices are effective in preventing false-positive results due to carryover or cross- contamination between samples.

Reproducibility

Reproducibility testing of contrived samples was performed at three test sites including one internal site (Site A) and two external sites (Site B and Site C). The study incorporated a range of potential variation introduced by sites, days, replicates, cartridge lots, operators, and QIAstat-Dx analyzers. For each site, testing was performed across 5 non-consecutive days with 6 replicates per day (leading to a total of 30 replicates per target, concentration and site), 4 QIAstat-Dx Analyzers (2 analyzers per operator and per site), and at least 2 operators on each testing day. A total of 5 sample mixes (two combined samples at 1x LoD and 3x LoD plus one negative sample) were prepared. For each mix, 6 replicates were tested and evaluated.

Table 14 shows the detection rate per target and concentration for each site of the Reproducibility study. In addition, data obtained at all three sites have been compiled to calculate the exact 2-sided 95% Confidence Interval by target and concentration.

Table 14. Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration

% Agreement with Expected Result

| Pathogen tested | Concentration tested | Expected result | Site A | Site B | Site C | All Sites (95% Confidence Interval) |
|---|----------------------|-----------------|---------------|---------------|---------------|---|
| Adenovirus F41 ZeptoMetrix 0810085CF | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Clostridium difficile ZeptoMetrix 0801619 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Campylobacter ZeptoMetrix 0801650 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |

Table 14. Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration (continued)

% Agreement with Expected Result

| | | _ | | | | |
|---|----------------------|-----------------|---------------|-----------------|-----------------|---|
| Pathogen tested | Concentration tested | Expected result | Site A | Site B | Site C | All Sites (95% Confidence Interval) |
| Escherichia coli EPEC ZeptoMetrix 0801747 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 29/30 96.67% | 30/30 100% | 89/90 98.89% (93.96 – 99.97%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Entamoeba histolytica ATCC 30459 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 29/30 96.67% | 89/90 98.89% (93.96 – 99.97%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Giardia lamblia ATCC 30888 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | | | | | | |

Table 14. Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration (continued)

% Agreement with Expected Result

| Pathogen tested | Concentration tested | Expected result | Site A | Site B | Site C | All Sites (95% Confidence Interval) |
|---|----------------------|-----------------|-----------------|-----------------|-----------------|---|
| Norovirus GII ZeptoMetrix 0810087CF | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 29/30 96.67% | 30/30 100% | 30/30 100% | 89/90 100% (93.96 – 99.97%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Rotavirus A ZeptoMetrix 0810280CF | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 29/30 96.67% | 30/30 100% | 89/90 98.89% (93.96 – 99.97%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Escherichia coli (STEC) O157:H7 ZeptoMetrix 0801622 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 29/30 96.67% | 89/90 98.89% (93.96 – 99.97%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |

Table 14. Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration (continued)

| | | | % Agreement with Expected Result | | | |
|--|----------------------|-----------------|----------------------------------|-----------------|-----------------|---|
| Pathogen tested | Concentration tested | Expected result | Site A | Site B | Site C | All Sites (95% Confidence Interval) |
| Escherichia coli (STEC) stx1 ZeptoMetrix 0801622 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Escherichia coli (STEC) stx2 ZeptoMetrix 801622 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| Salmonella enterica ZeptoMetrix 801437 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 29/30 96.67% | 29/30 96.67% | 88/90 97.78% (92.20 - 99.73%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 – 99.73%) |

Table 14. Detection rate per target and concentration for each site of the Reproducibility study and exact 2-sided 95% Confidence Interval by target and concentration (continued)

| % Agreemen | t with Fx | nected | Result |
|------------|-----------|--------|--------|
| | | | |

| Pathogen tested | Concentration tested | Expected result | Site A | Site B | Site C | All Sites (95% Confidence Interval) |
|---|----------------------|-----------------|---------------|---------------|---------------|---|
| | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 – 99.73%) |
| Yersinia enterocolitica Zeptometrix 801734 | 3x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | 1x LoD | Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 - 100.00%) |
| | None | Not Detected | 30/30 100% | 30/30 100% | 30/30 100% | 90/90 100% (95.98 – 99.73%) |

Repeatability

A repeatability study was conducted on the QIAstat-Dx Analyzer 1.0 instruments using a set of samples composed of low-concentrated analytes spiked into stool matrix (3x LoD and 1x LoD) and negative stool samples. Pathogens included in the positive samples were Adenovirus, Clostridium difficile, Campylobacter, Enteropathogenic *E. coli* (EPEC), Entamoeba histolytica, Giardia lamblia, Norovirus GII, Rotavirus, *E. coli* O157, STEC stx1, STEC stx2, Salmonella enterica, Vibrio parahaemolyticus and Yersinia enterocolitica. Each sample was tested with the same instrument over 12 days. In total, 60 replicates of 1x LoD and 60 replicates of 3x

LoD per each of the tested targets and 60 replicates of negative samples were run. Overall results showed a 93.33-100.00-% and 95.00-100.00% detection rate for 1x LoD and 3x LoD samples, respectively. Negative samples showed 100% of negative calls for all panel analytes.

Repeatability in the QIAstat-Dx Rise instrument was also evaluated in comparison with QIAstat-Dx Analyzers. A study was conducted on two QIAstat-Dx Rise instruments using a representative set of samples composed of low-concentrated analytes (3x LoD and 1x LoD) spiked into stool matrix and negative stool samples. Pathogens included in the positive samples were *Norovirus GII*, *Entamoeba histolytica*, *Clostridium difficile*, *Yersinia enterocolitica*, *Salmonella enterica*, Adenovirus F 40 and Rotavirus A. Samples were tested in replicates using two lots of cartridges. In total, 128 replicates of 1x LoD positive samples, 128 replicates of 3x LoD positive samples, and 64 replicates of negative samples were run on the QIAstat-Dx Rise instrument. Overall results showed a 99.22-100.00-% detection rate for both 1x LoD and 3x LoD samples. Negative samples showed 100% of negative calls for all panel analytes. Testing with two QIAstat-Dx Analyzers (each with four Analytical Modules) was included in the study for results comparison. QIAstat-Dx Rise performance was shown to be equivalent to QIAstat-Dx Analyzer 1.0.

Clinical Performance

The clinical performance shown below was demonstrated using QIAstat-Dx Analyzer 1.0. The QIAstat-Dx Rise and the QIAstat-Dx Analyzer 2.0 use the same Analytical Modules as QIAstat-Dx Analyzer 1.0; therefore the performance is not impacted by QIAstat-Dx Rise or the QIAstat-Dx Analyzer 2.0. A multi-center international observational clinical study was conducted using prospectively and retrospectively collected samples to evaluate the performance of QIAstat-Dx Gastrointestinal Panel 2 during normal conditions of use. The study was conducted in 13 clinical sites across 5 countries (4 sites in Europe and 9 sites in USA) from May 2021 to July 2021

The final data set consisted of a total of 2,085 leftover de-identified specimens, which were prospectively collected from patients who underwent stool specimen collection for clinical indications of diarrhea caused by gastrointestinal infection at the 13 investigational sites. In addition, testing was performed on archived known positives and contrived specimens to further augment the positive specimen numbers (Table 15). Samples used in the study were all stool samples in Cary-Blair transport media collected using either Para-Pak C&S (Meridian Bioscience), FecalSwab® (COPAN), Fecal Transwab® (Medical Wire & Equipment Co. (Bath) Ltd), or C & S Medium (Medical Chemical).

Table 15. Prospective and Archived specimens summary across each clinical study site

| Site/Country | Specimen Type | Total | |
|--------------|---------------------|-------------------|------|
| | Prospective (Fresh) | Archived (Frozen) | |
| Germany | 339 | 21 | 360 |
| Denmark | 293 | 37 | 330 |
| Spain | 247 | 60 | 307 |
| France | 63 | 7 | 70 |
| USA site 1 | 186 | 6 | 192 |
| USA site 2 | 43 | 9 | 52 |
| USA site 3 | 282 | 84 | 366 |
| USA site 4 | 177 | 0 | 177 |
| USA site 5 | 44 | 0 | 44 |
| USA site 6 | 39 | 0 | 39 |
| USA site 7 | 148 | 0 | 148 |
| USA site 8 | 131 | 0 | 131 |
| USA site 9 | 95 | 0 | 95 |
| Total | 2087 | 224 | 2311 |

All prospectively collected specimens that had age, sex, and patient population status were collected by the site. Subject demographics (evaluable specimens) are summarized in Table 16 below.

Table 16. Demographic data for enrolled prospective specimens

| Demographic data | N | % |
|---|------------------|------|
| Gender | | |
| Female | 1158 | 55.5 |
| Male | 927 | 44.5 |
| Age group | | |
| 0-6 years | 221 | 10.6 |
| 6-21 years | 167 | 8.0 |
| 22-49 years | 540 | 25.9 |
| 50+ years | 1150 | 55.2 |
| Not Reported | 7 | 0.3 |
| Patient population | | |
| Emergency room | 114 | 5.5 |
| Hospitalized | 500 | 24.0 |
| Immunocompromised | 3 | 0.1 |
| Outpatient | 908 | 43.5 |
| Not Reported | 560 | 26.9 |
| No. of days between symptom onset and QIA | Astat-Dx testing | |
| >7 days | 152 | 7.3 |
| ≤7 days | 222 | 10.6 |
| Not Reported | 1711 | 82.1 |

Performance of the QIAstat- Dx Gastrointestinal Panel 2 was compared to the reference method: BioFire [®] FilmArray [®] GI Panel for all targets. For the majority of targets, direct comparison of the two results could be made as a binary result (positive or negative). However, for certain targets the QIAstat-Dx GI Assay provides additional differentiation, so further comparators were required to determine agreement., The appropriate comparator/reference method used for each member of the panel are detailed in the QIAstat- Dx Gastrointestinal Panel 2 Clinical studies reference method below.

Table 17. QIAstat-Dx Gastrointestinal Panel 2 Clinical studies reference method

QIAstat-Dx GI Panel 2 target

Comparator method

Adenovirus F40/F41

Astrovirus

Norovirus GI/GII

Rotavirus A

Sapovirus (GI, GII, GIV, GV)

Campylobacter (C. jejuni, C. coli and C. upsaliensis)

Clostridium difficile (toxin A/B)

Enteroaggregative Escherichia coli (EAEC)

Shigella/Enteroinvasive Escherichia coli (EIEC)

Enteropathogenic Escherichia coli (EPEC)

Enterotoxigenic Escherichia coli (ETEC) lt/st

Shiga-like toxin-producing Escherichia coli (STEC)

stx1/stx2

E. coli O157 serogroup

Salmonella

Plesiomonas shigelloides

Vibrio cholerae

Yersinia enterocolitica

Cryptosporidium

Cyclospora cayetanensis

Entamoeba histolytica

BioFire FilmArray Gastrointestinal (GI) Panel

Table 17. QIAstat-Dx Gastrointestinal Panel 2 Clinical studies reference method (continued)

QIAstat-Dx GI Panel 2 target Comparator method

| Vibrio parahaemolyticus | BioFire FilmArray GI Panel Vibrio + PCR-BDS* assay to identify V. parahaemolyticus |
|-------------------------|--|
| Vibrio vulnificus | BioFire FilmArray GI Panel Vibrio + PCR-BDS* assay to identify V. vulnificus |

^{*}This is a targeted Polymerase Chain Reaction (PCR) assay which was developed and validated for the performance evaluation, when amplification is observed in the PCR, the amplicon was verified by Bi-directional Sequencing (BDS).

Discordant Results Resolution

Upon discordance with the comparator method, resolution testing was performed to determine the presence/absence of specific targets. Table 18 below details the methods used for discordance resolution.

Table 18. Discordant Specimen Testing

| QIAstat-Dx GI Panel 2 target | Discordant testing method |
|---|---|
| Adenovirus F40/F41 | BD-MAX Enteric Viral Panel |
| Astrovirus | |
| Norovirus GI/GII | |
| Rotavirus A | |
| Sapovirus (GI, GII, GIV, GV) | |
| Campylobacter (C. jejuni, C. coli and C. upsaliensis) Shigella/Enteroinvasive E. coli (EIEC) Salmonella | BD-MAX Enteric Bacterial Panel |
| Enterotoxigenic E. coli (ETEC) lt/st Plesiomonas shigelloides Yersinia enterocolitica | BD-MAX Extended Enteric Bacterial Panel |

Table 18. Discordant Specimen Testing (continued)

QIAstat-Dx GI Panel 2 target

Discordant testing method

Clostridium difficile (toxin A/B)

Enteroaggregative E. coli (EAEC)

Enteropathogenic E. coli (EPEC)

Shiga-like toxin- E. coli (STEC) stx1

Shiga-like toxin- E. coli (STEC) stx2

Vibrio cholerae

Vibrio parahaemolyticus

Vibrio vulnificus

Cryptosporidium

Giardia lamblia

Clinical Performance – PPA and NPA

A total of 2,309 prospective and archived clinical samples were evaluated to determine the Clinical Performance characteristics of the QIAstat-Dx Gastrointestinal Panel 2. The Positive Percentage Agreement (PPA) and Negative Percent Agreement (NPA) was calculated for each target after discordance resolution for all clinical samples (prospective and archived).

Additionally, to supplement the prospective and archived clinical samples data, an evaluation of contrived specimens was performed for several pathogens (Adenovirus F40/F41, Astrovirus, Rotavirus, Sapovirus, Campylobacter, ETEC, EIEC/Shigella, STEC stx1/stx2, E. coli O157, Plesiomonas shigelloides, Salmonella, Vibrio cholerae, Vibrio parahaemolyticus, Vibrio vulnificus, Yersinia enterocolitica, Cryptosporidium, Cyclospora cayetanensis, Entamoeba histolytica, and Giardia lamblia), due to low number of prospective and archived clinical samples found during the study. Surrogate specimens were prepared using residual clinical specimens that had previously tested negative for all GI panel analytes targeted by QIAstat-Dx Gastrointestinal Panel 2 and comparator methods. Specimens were spiked around

^{*} All Polymerase Chain Reaction (PCR)- Bidirectional Sequencing (BDS) assays represent a validated nucleic acid amplification test (NAAT) followed by bi-directional sequencing. For Vibrio parahaemolyticus and Vibrio vulnificus, the same PCR-BDS method was used for both discordant testing and differentiation testing.

the assay LoD and at clinically relevant levels using different quantified strains for each organism. The analyte status of each contrived specimen was blinded to the users analyzing the specimens. A total of 1,254 cartridge test runs were performed for the contrived samples providing additional data on the rarer pathogens measured by QIAstat-Dx Gastrointestinal Panel 2. PPA was established for the mentioned targets on contrived samples.

The total combined PPA and NPA per pathogen and overall was calculated alongside with the corresponding exact binomial two-sided 95% confidence interval. The results are summarized in Table 19 below.

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI

| | | Sensitivity (P | PA) | | | Specificity (NP) | A) | | |
|------------|---------------|----------------|--------|---------|--------|------------------|------------|--------|--------|
| | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| Target | type | TP/(TP+FN) | % | Lower | Upper | TP/(TP+FN) | % | Lower | Upper |
| | | | | Viruses | 5 | | | | |
| Adenovirus | Clinical | 9/9 | 100.00 | 66.37 | 100.00 | 2287/2288 | 99.96 | 99.76 | 100.00 |
| F40/F41 | Contrive d | 68/70 | 97.14 | 90.06 | 99.65 | N/A | N/A | N/A | N/A |
| | Total | 77/79 | 97.47 | 91.15 | 99.69 | 2287/2288 | 99.96 | 99.76 | 100.00 |
| | | | | | | | | | |
| Astrovirus | Clinical | 13/14 | 92.86 | 66.13 | 99.82 | 2282/2282 | 100.00 | 99.84 | 100.00 |
| | Contrive d | 67/68 | 98.53 | 92.08 | 99.96 | N/A | N/A | N/A | N/A |
| | Total | 80/82 | 97.56 | 91.47 | 99.70 | 2282/2282 | 100.00 | 99.84 | 100.00 |
| Norovirus | Clinical | 69/73 | 94.52 | 86.56 | 98.49 | 2221/2222 | 99.95 | 99.75 | 100.00 |
| | Contrive d | 0/0 | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| | Total | 69/73 | 94.52 | 86.56 | 98.49 | 2221/2222 | 99.95 | 99.75 | 100.00 |

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI (continued)

| | | Sensitivity (P | PA) | | | Specificity (NP | A) | | |
|---------------------------------------|---------------|----------------|--------|----------|--------|-----------------|-----------|--------|--------|
| | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| Target | type | TP/(TP+FN) | % | Lower | Upper | TN/(TN+FP) | % | Lower | Upper |
| Rotavirus | Clinical | 34/36 | 94.44 | 81.34 | 99.32 | 2257/2260 | 99.87 | 99.61 | 99.97 |
| | Contrive d | 69/70 | 98.57 | 92.30 | 99.96 | N/A | N/A | N/A | N/A |
| | Total | 103/106 | 97.17 | 91.95 | 99.41 | 2256/2259 | 99.87 | 99.61 | 99.97 |
| Sapovirus | Clinical | 16/16 | 100.00 | 79.41 | 100.00 | 2280/2281 | 99.96 | 99.76 | 100.00 |
| | Contrive d | 69/69 | 100.00 | 94.79 | 100.00 | N/A | N/A | N/A | N/A |
| | Totals | 85/85 | 100.00 | 95.75 | 100.00 | 2280/2281 | 99.96 | 99.76 | 100.00 |
| | | | | Bacterio | 1 | | | | |
| Campylobacter | Clinical | 146/146 | 100.00 | 97.51 | 100.00 | 2148/2152 | 99.81 | 99.52 | 99.95 |
| | Contrive d | 45/46 | 97.83 | 88.47 | 99.94 | N/A | N/A | N/A | N/A |
| | Total | 191/192 | 99.48 | 97.13 | 99.99 | 2148/2152 | 99.81 | 99.52 | 99.95 |
| Clostridium difficile toxin A/B | Clinical | 234/245 | 95.51 | 92.11 | 97.74 | 2053/2056 | 99.85 | 99.57 | 99.97 |
| | Contrive d | 0/0 | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| | Total | 234/245 | 95.51 | 92.11 | 97.74 | 2053/2056 | 99.85 | 99.57 | 99.97 |
| Enteroaggre- | Clinical | 83/96 | 86.46 | 77.96 | 92.59 | 2196/2201 | 99.77 | 99.47 | 99.93 |
| gative <i>E. coli</i> (EAEC) | Contrive d | 0/0 | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| | Total | 83/96 | 86.46 | 77.96 | 92.59 | 2196/2201 | 99.77 | 99.47 | 99.93 |

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI (continued)

| | | Sensitivity (P | PA) | | | Specificity (NP) | A) | | |
|---|---------------|----------------|--------|--------|--------|------------------|------------|--------|--------|
| | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| Target | type | TP/(TP+FN) | % | Lower | Upper | TN/(TN+FP) | % | Lower | Upper |
| Enteropathoge | Clinical | 236/256 | 92.19 | 88.19 | 95.16 | 1980/1984 | 99.80 | 99.48 | 99.95 |
| nic <i>E. coli</i> (EPEC) | Contrive d | 0/0 | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| | Total | 236/256 | 92.19 | 88.19 | 95.16 | 1980/1984 | 99.80 | 99.48 | 99.95 |
| Enterotoxigenic | Clinical | 59/62 | 95.16 | 86.50 | 98.99 | 2235/2236 | 99.96 | 99.75 | 100.00 |
| E. coli (ETEC) lt/st | Contrive d | 43/43 | 100.00 | 91.78 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 102/105 | 97.14 | 91.88 | 99.41 | 2235/2236 | 99.96 | 99.75 | 100.00 |
| Shigella/ Enteroinvasive E. coli (EIEC) | Clinical | 37/38 | 97.37 | 86.19 | 99.93 | 2259/2259 | 100.00 | 99.84 | 100.00 |
| , , | Contrive d | 69/69 | 100.00 | 94.79 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 106/107 | 99.07 | 94.90 | 99.98 | 2259/2259 | 100.00 | 99.84 | 100.00 |
| Shiga-like toxin | Clinical | 43/50 | 86.00 | 73.26 | 94.18 | 2244/2246 | 99.91 | 99.68 | 99.99 |
| E.coli (STEC) stx1/stx2* | Contrive d | 200/200 | 100.00 | 98.17 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 243/250 | 97.20 | 94.32 | 98.87 | 2244/2246 | 99.91 | 99.68 | 99.99 |
| E. coli 0157 | Clinical | 2/2 | 100.00 | 15.81 | 100.00 | 38/38 | 100.00 | 90.75 | 100.00 |
| | Contrive d | 67/69 | 97.10 | 89.92 | 99.65 | N/A | N/A | N/A | N/A |
| | Total | 69/71 | 97.18 | 90.19 | 99.66 | 38/38 | 100.00 | 90.75 | 100.00 |

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI (continued)

| | | Sensitivity (PPA) | | Specificity (NPA) | | | | | |
|---------------------------------|---------------|-------------------|--------|-------------------|--------|------------|--------|--------|--------|
| | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| Target | type | TP/(TP+FN) | % | Lower | Upper | TN/(TN+FP) | % | Lower | Upper |
| Plesiomonas | Clinical | 8/8 | 100.00 | 63.06 | 100.00 | 2283/2288 | 99.78 | 99.49 | 99.93 |
| shigelloides | Contrive d | 67/68 | 98.53 | 92.08 | 99.96 | N/A | N/A | N/A | N/A |
| | Total | 75/76 | 98.68 | 92.89 | 99.97 | 2283/2288 | 99.78 | 99.49 | 99.93 |
| Salmonella | Clinical | 71/71 | 100.00 | 94.94 | 100.00 | 2225/2227 | 99.91 | 99.68 | 99.99 |
| | Contrive d | 33/33 | 100.00 | 89.42 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 104/104 | 100.00 | 96.52 | 100.00 | 2225/2227 | 99.91 | 99.68 | 99.99 |
| Vibrio cholerae | Clinical | 2/2 | 100.00 | 15.81 | 100.00 | 2294/2294 | 100.00 | 99.84 | 100.00 |
| | Contrive d | 67/70 | 95.71 | 87.98 | 99.11 | N/A | N/A | N/A | N/A |
| | Total | 69/72 | 95.83 | 88.30 | 99.13 | 2294/2294 | 100.00 | 99.84 | 100.00 |
| Vibrio parahaemoly- ticus | Clinical | 3/4 | 75.00 | 19.41 | 99.37 | 2291/2292 | 99.96 | 99.76 | 100.00 |
| | Contrive d | 70/70 | 100.00 | 94.87 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 73/74 | 98.65 | 92.70 | 99.97 | 2291/2292 | 99.96 | 99.76 | 100.00 |
| Vibrio | Clinical | 0/0 | N/A | N/A | N/A | 2296/2296 | 100.00 | 99.84 | 100.00 |
| vulnificus | Contrive d | 69/69 | 100.00 | 94.79 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 69/69 | 100.00 | 94.79 | 100.00 | 2296/2296 | 100.00 | 99.84 | 100.00 |

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI (continued)

| | | Sensitivity (P | PA) | | | Specificity (NP | A) | | |
|----------------------------|---------------|----------------|--------|----------|--------|-----------------|---------------|--------|--------|
| | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| Target | type | TP/(TP+FN) | % | Lower | Upper | TN/(TN+FP) | - % | Lower | Upper |
| Yersinia enterocolitica | Clinical | 51/51 | 100.00 | 93.02 | 100.00 | 2232/2246 | 99.38 | 98.96 | 99.66 |
| enieroconnica | | | | | | | | | |
| | Total | 119/120 | 99.17 | 95.44 | 99.98 | 2232/2246 | 99.38 | 98.96 | 99.66 |
| | | | | Parasite | Э | | | | |
| Cryptosporidiu | Clinical | 19/21 | 90.48 | 69.62 | 98.83 | 2272/2275 | 99.87 | 99.62 | 99.97 |
| m spp. | Contrive d | 58/58 | 100.00 | 93.84 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 77/79 | 97.47 | 91.15 | 99.69 | 2272/2275 | 99.87 | 99.62 | 99.97 |
| Cyclospora | Clinical | 25/26 | 96.15 | 80.36 | 99.90 | 2270/2270 | 100.00 | 99.84 | 100.00 |
| cayetanensis | Contrive d | 56/56 | 100.00 | 93.62 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 81/82 | 98.78 | 93.39 | 99.97 | 2270/2270 | 100.00 | 99.84 | 100.00 |
| Entamoeba | Clinical | 0/0 | N/A | N/A | N/A | 2296/2296 | 100.00 | 99.84 | 100.00 |
| histolytica | Contrive d | 69/70 | 98.57 | 92.30 | 99.96 | N/A | N/A | N/A | N/A |
| | Total | 69/70 | 98.57 | 92.30 | 99.96 | 2296/2296 | 100.00 | 99.84 | 100.00 |
| Giardia | Clinical | 36/36 | 100.00 | 90.26 | 100.00 | 2255/2260 | 99.78 | 99.48 | 99.93 |
| lamblia | Contrive d | 56/56 | 100.00 | 93.62 | 100.00 | N/A | N/A | N/A | N/A |
| | Total | 92 /92 | 100.00 | 96.07 | 100.00 | 2255/2260 | 99.78 | 99.48 | 99.93 |
| Overall clinical samples | | 1196/126 2 | 94.77 | 93.39 | 95.93 | 49195/492 50 | 99.89 | 99.85 | 99.92 |

Table 19. Summary of Clinical Study Results for all clinical specimens (prospective and retrospective), contrived samples and total combined, including the exact binomial two-sided 95% CI (continued)

| | | Sensitivity (PPA) | | | Specificity (NPA) | | | | |
|---------------------------------------|--------|-------------------|-------|-------------|-------------------|-----------------|-------|--------|-------|
| _ | Sample | Fraction | | 95% CI | | Fraction | | 95% CI | |
| | type | TP/(TP+FN) | % | Lower Upper | TN/(TN+FP) | % | Lower | Upper | |
| Overall con- trived spe- cimens | | 1310/132 3 | 99.02 | 98.33 | 99.48 | N/A | N/A | N/A | N/A |
| | | | | | | | | | |
| Overall total combined | | 2506/258 5 | 96.94 | 96.21 | 97.57 | 49195/492 50 | 99.89 | 99.85 | 99.92 |

Troubleshooting Guide

This troubleshooting guide may be helpful in solving any problems that may arise. For more information, see also the Frequently Asked Questions page at our Technical Support Center: www.qiagen.com/FAQ/FAQList.aspx . The scientists in QIAGEN Technical Services are always happy to answer any questions you may have about either the information and/or protocols in this Instructions for Use or sample and assay technologies (for contact information, visit www.qiagen.com).

Additional information about specific QIAstat-Dx Gastrointestinal Panel 2 error codes and messages can be found in Information about specific QIAstat-Dx Gastrointestinal Panel 2 error codes and messages:

Table 20. Information about specific QIAstat-Dx Gastrointestinal Panel 2 error codes and messages

| Error Code | Error message displayed |
|------------|---|
| 0x02C9 | Cartridge execution failure: Sample concentration too high. |
| 0x032D | Please repeat by loading 100 microliters of the sample in a new cartridge (per IFU explanation) |
| 0x0459 | |
| 0x045A | |
| 0x04BF | |
| 0x0524 | |
| 0x058B | |
| 0x05E9 | |
| 0x0778 | |
| 0x077D | |
| 0x14023 | |

When the sample concentration is too high and the test must be repeated by loading 100 µl, follow the workflow detailed in the Appendix C of this document.

Symbols

The following table describes the symbols that may appear on the labeling or in this document.

| Symbols | Description |
|--------------------|---|
| <u>\\\</u> <n></n> | Contains reagents sufficient for <n> reactions</n> |
| | Use by |
| IVD | For in vitro diagnostic use |
| *** | Manufacturer |
| REF | Catalog number |
| LOT | Lot number |
| MAT | Material number (i.e., component labeling) |
| | Gastrointestinal application |
| Rn | R is for revision of the Instructions for Use and n is the revision |
| | Temperature limitation |
| | Consult instructions for use |
| <u>^</u> | Caution |

| Symbols | Description |
|-------------|---|
| SN | Serial number |
| 3 | Do not reuse |
| 类 | Keep away from sunlight |
| ® | Do not use if package is damaged |
| GTIN | Global Trade Item Number |
| ® | Flammable, risk of fire |
| | Corrosive, risk of chemical burn |
| ③ | Health hazard, risk of sensitization, carcinogenicity |
| (!) | Risk of harm |

Contact Information

For technical assistance and more information, please see our Technical Support Center at www.qiagen.com/Support, call 00800- 22- 44- 6000, or contact one of the QIAGEN Technical Service Departments or local distributors (see back cover or visit www.qiagen.com).

Appendices

Appendix A: Installing the Assay Definition File

The Assay Definition File (ADF 1.1) of the QIAstat-Dx Gastrointestinal Panel 2 must be installed on the QIAstat- Dx Analyzer 1.0 and QIAstat- Dx Rise prior to testing with QIAstat- Dx Gastrointestinal Panel 2 Cartridges.

Note: For QlAstat-Dx Rise, please contact Technical Service or your sales representative to upload new assay definition files.

Note: Whenever a new version of the QIAstat-Dx Gastrointestinal Panel 2 assay is released, the new QIAstat-Dx Gastrointestinal Panel 2 Assay Definition File must be installed prior to testing.

The Assay Definition File (.asy file type) is available at **www.qiagen.com**. The Assay Definition file (.asy file type) must be saved onto a USB Drive prior to installation on the QIAstat-Dx Analyzer 1.0. This USB Drive must be formatted with a FAT32 file system.

To import an ADF from the USB to the QIAstat-Dx Analyzer 1.0, proceed with the following steps:

- Insert the USB stick containing the Assay Definition File into one of the USB ports on the QIAstat-Dx Analyzer 1.0.
- 2. Press the Options button and then select Assay Management. The Assay Management screen appears in the Content area of the display (Figure 55).

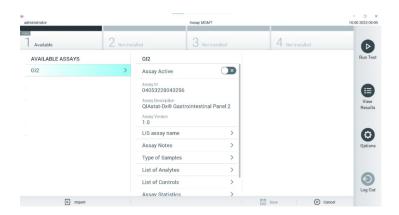


Figure 55. Assay Management screen.

- 3. Press the Import icon in the bottom left of the screen (Figure 55).
- 4. Select the file corresponding to the assay to be imported from the USB drive.
- 5. A dialog box will appear to confirm upload of the file.
- 6. A dialog box may appear to override the current version by a new one. Press Yes to override (Figure 56).



Figure 56. Dialog that appears when upgrading the ADF version.

7. The assay becomes active by selecting Assay Active (Figure 57).

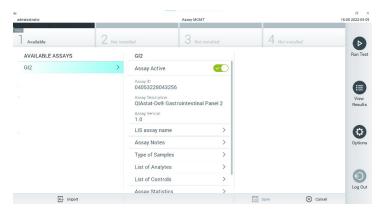


Figure 57. Activating the assay.

8. Assign the active assay to the user by pressing the Options button and then the User Management button. Select the user who should be allowed to run the assay. If it is needed, this action can be repeated for every user created in the system. Next, select Assign Assays from the "User Options". Enable the assay and press the Save button (Figure 58).

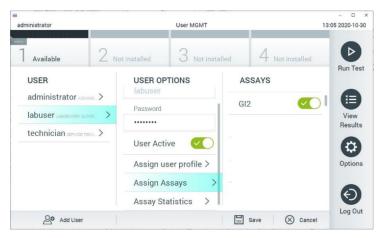


Figure 58. Assigning the active assay.

Appendix B: Glossary

- Amplification curve: Graphical representation of the multiplex real-time RT- PCR amplification data.
- Analytical Module (AM): The main QlAstat-Dx Analyzer 1.0 hardware module, in charge of executing tests on QlAstat-Dx Gastrointestinal Panel 2 Cartridges. It is controlled by the Operational Module. Several Analytical Modules can be connected to one Operational Module.
- QIAstat-Dx Analyzer 1.0: The QIAstat-Dx Analyzer 1.0 consists of an Operational Module and an Analytical Module. The Operational Module includes elements that provide connectivity to the Analytical Module and enables user interaction with the QIAstat-Dx

- Analyzer 1.0. The Analytical Module contains the hardware and software for sample testing and analysis.
- QlAstat- Dx Rise: The QlAstat- Dx Rise Base is an in-vitro diagnostic device for use with QlAstat-Dx assays and QlAstat-Dx 1.0 Analytical Modules, which provides full automation from sample preparation to real-time PCR detection for molecular applications. The system can be operated either in random access and batch testing, and the system throughput can be escalated up to 160 test/day by including up to 8 Analytical Modules. The system also includes a multi-test front drawer that can accommodate up to 16 tests at the same time, and a waste drawer to automatically discard the performed tests, enhancing the walk-away efficiency of the system.
- QlAstat-Dx Gastrointestinal Panel 2 Cartridge: A self-contained disposable plastic device
 with all pre-loaded reagents required for the complete execution of fully automated
 molecular assays for the detection of gastrointestinal pathogens.
- IFU: Instructions For Use.
- Main port: In the QIAstat-Dx Gastrointestinal Panel 2 Cartridge, inlet for transport medium liquid samples.
- Nucleic acids: Biopolymers, or small biomolecules composed of nucleotides, which are monomers made of three components: a 5- carbon sugar, a phosphate group and a nitrogenous base.
- Operational Module (OM): The dedicated QIAstat- Dx Analyzer 1.0 hardware that provides the user interface for 1–4 Analytical Modules (AM).
- PCR: Polymerase Chain Reaction.
- IUO: For investigational use only
- RT: Reverse Transcription.

- Swab port: In the QIAstat-Dx Gastrointestinal Panel 2 Cartridge, inlet for dry swabs. The swab port is not used for the QIAstat-Dx Gastrointestinal Panel 2 assay.
- User: A person who operates the QIAstat-Dx Analyzer 1.0/ QIAstat-Dx Rise/QIAstat-Dx Gastrointestinal Panel 2 Cartridge in the intended way.

Appendix C: Additional Instructions for use

In case of cartridge execution failures, corresponding to error codes (0x02C9, 0x032D, 0x0459, 0x045A, 0x04BF, 0x0524, 0x058B, 0x05E9, 0x077B, 0x077D, 0x14023) that occur during the testing, the following error message will be displayed in the QIAstat-Dx Analyzer 1.0 screen after the run has finalized:

Cartridge execution failure: Sample concentration too high. Please repeat by loading 100 microliters of the sample in a new cartridge (as per IFU explanation)'.

In this case, the test should be repeated using $100~\mu L$ of the same sample following equivalent testing procedures detailed in the "Procedure" Section in the Instructions for Use adapted to $100~\mu l$ sample input volume:

- 1. Open the package of a new QIAstat-Dx Gastrointestinal Panel 2 Cartridge using the tear notches on the sides of the packaging.
- 2. Remove the QIAstat-Dx Gastrointestinal Panel 2 Cartridge from the packaging.
- Manually write the sample information, or place a sample information label on the top of the QIAstat- Dx Gastrointestinal Panel 2 Cartridge. Ensure that the label is properly positioned and does not block the lid opening.
- 4. Place the QIAstat-Dx Gastrointestinal Panel 2 Cartridge flat on the clean work surface so that the bar code on the label faces upwards. Open the sample lid of the main port on the front of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge.

- 5. Thoroughly mix the stool in the Cary-Blair transport medium, for example, by vigorously agitating the tube 3 times.
- 6. Open the tube with the sample to be tested. Use the supplied transfer pipette to draw up fluid. Draw the sample to the first fill line on the pipette (i.e., $100 \, \mu$ l)
- 7. IMPORTANT: Do not draw air, mucus, or particles into the pipette. If air, mucus, or particles are drawn into the pipette, carefully expel the sample fluid in the pipette back into the sample tube and draw up fluid again.
- 8. Carefully transfer the sample into the main port of the QIAstat-Dx Gastrointestinal Panel 2 Cartridge using the supplied single-use transfer pipette (Figures 6 and 7).
- 9. Firmly close the lid of the main port until it clicks (Figure 8).

From this point, proceed following the instructions described in the IFU.

Ordering Information

| Product | Contents | Cat. no. |
|---|---|----------|
| QlAstat-Dx Gastrointestinal Panel 2 | For 6 tests: 6 individually packaged QIAstat-Dx Gastrointestinal Panel 2 Cartridges and 6 individually packaged transfer pipettes | 691412 |
| Related Products | | |
| QlAstat-Dx Analyzer 1.0 | 1 QIAstat-Dx Analytical Module, 1 QIAstat-Dx Operational Module and related hardware and software to run molecular diagnostic QIAstat-Dx assay cartridges | 9002824 |
| QIAstat-Dx Rise | 1 QIAstat-Dx Rise Base Module and related hardware and software to run molecular diagnostics on QIAstat-Dx assay cartridges | 9003163 |

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Document Revision History

| Date | Changes |
|-------------------------|--|
| R1, December 2024 | Initial Release for Singapore. Edited Table 19 to correct table headers. Replaced instances of "handbook" with "Instructions for Use". |

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