



INSTRUCTION FOR USE

Urinary Tract Infections Panel PCR Kit

For Research Use Only



100



MBLUTI004



06 2024



Hürriyet Mah. Erler Sok. No: 4/1 B Blok Nexus No: 13, Kartal/Istanbul-TURKEY

Document Revision History

Rev.No_Date	Revision Description
Rev.00_June 20, 2024	First Release

CONTENTS

1.	INTENDED USE	3
2.	PRINCIPLE of the PROCEDURE	4
3.	KIT COMPONENTS	5
4.	EQUIPMENT and MATERIALS REQUIRED but NOT PROVIDED	7
5.	WARNING and PRECAUTIONS	8
6.	HANDLING, STORAGE, and STABILITY	9
7.	TEST PROCEDURE	10
7.1.	Sample Preparation and Nucleic Acid Extraction	10
7.2.	PCR Reaction Preparation and Processing	10
8.	INTERPRETATION OF RESULTS	12
8.1.	Calculation of Cq Values and Instrument-Specific Requirements	12
8.2.	Overall Validity of Detection	12
8.3.	Interpretation of Unknown Specimen Results	13
9.	ASSAY LIMITATIONS	14
10.	PERFORMANCE CHARACTERISTICS	15
10.1.	Analytical Sensitivity (Limit of Detection, LoD)	15
10.2.	Device Equivalence Study	16
10.3.	Analytical Reactivity (Inclusivity)	17
10.3.1.	In-Slico Analytical Reactivity	17
10.3.2.	Wet-Test Analytical Reactivity	20
10.4.	Analytical Specificity (Exclusivity)	21
10.4.1.	In-Slico Analytical Specificity	21
10.4.2.	Wet-Test Analytical Specificity	23
10.5.	Interferences	25
11.	TROUBLESHOOTING	28
12.	EXPLANATION of SYMBOLS	29

1. INTENDED USE

For Research Use Only (RUO). Not for use in diagnostic procedures. No claim or representation is intended to provide information for the diagnosis, prevention, or treatment of disease. Furthermore, the test kit is not intended to diagnose infectious animal diseases.

The **MarinaBiolab Urinary Tract Infections Panel PCR Kit** is a multiplexed qualitative Real-Time Polymerase Chain Reaction (qPCR) test intended for the simultaneous detection and identification of multiple pathogenic nucleic acids in research samples. The **MarinaBiolab Urinary Tract Infections Panel PCR Kit** allows to achieve qPCR result in less than 1 hour. The test is performed to detect gene sequences of the following organisms.

Targets	
<i>Escherichia coli</i>	<i>Corynebacterium urealyticum</i>
<i>Streptococcus agalactiae</i>	<i>Enterococcus faecium</i>
<i>Klebsiella oxytoca</i>	<i>Enterococcus faecalis</i>
<i>Staphylococcus saprophyticus</i>	<i>Acinetobacter baumannii</i>
<i>Serratia marcescens</i>	<i>Proteus vulgaris</i>
<i>Proteus mirabilis</i>	<i>Staphylococcus aureus</i>
<i>Aerococcus urinae</i>	<i>Ureplasma (Ureaplasma urealyticum/parvum)</i>
<i>Treponema pallidum</i>	<i>Providencia stuartii</i>
<i>Enterobacter cloacae</i>	<i>Candida albicans</i>
<i>Pseudomonas aeruginosa</i>	<i>Candida glabrata</i>
<i>Citrobacter freundii</i>	<i>Candida parapsilosis</i>
<i>Klebsiella aerogenes</i>	<i>Candida tropicalis</i>
<i>Klebsiella pneumoniae</i>	<i>Candida krusei</i>
<i>Morganella morganii</i>	<i>Candida auris</i>
Controls	
Human RNase P (IC)	
<i>Bacillus atropheus</i> (EC)	

2. PRINCIPLE of the PROCEDURE

DNA target regions are amplified via real-time PCR instruments using the primer and probe sets in the kit. In the process, the probe anneals a specific target sequence located between the forward and reverse primers. During the extension phase of the PCR cycle, the 5' nuclease activity of Taq polymerase degrades the probe, causing the reporter dye to separate from the quencher, generating a fluorescent signal. With each cycle, additional reporter dye molecules are cleaved from their respective probes, increasing the fluorescence intensity. Fluorescence intensity is monitored at each PCR cycle by the real-time PCR instruments. Probes labeled with different fluorophores are used to detect specific amplicons originating from targets and Internal Control.

PCR instruments measure these signals at the end of each amplification cycle in real time and interpret the data to provide a qualitative result for each of the above targets. A positive result for the detection of target DNA is indicated by the presence of a real-time PCR growth curve and an associated C_q (Quantification Cycle) value.

3. KIT COMPONENTS

The **MarinaBiolab Urinary Tract Infections Panel PCR Kit** consists of four main components:

1. qPCR Enzyme and Buffer Mix (qPCR Master Mix)
2. Forward, Reverse and Probe Oligo Mix (UTIP Oligo Mix 1-8)
3. Mix of non-infectious DNA from artificial sample including targets in the table below (PC-UTIP)
4. DNase/RNase-Free Water (NTC)

The kit components are provided in Table 1-2.

Table 1. Kit components.

Component	Description	Quantity x Volume
		100 rxn MBLUT1004
qPCR Master Mix	Ready-to-use mix for qPCR	4 x 1000 µL
UTIP Oligo Mix 1-8	Primers and probes complementary to specific regions of the targets in the table above	8 x 250 µL
PC-UTIP	Mix of non-infectious DNA from artificial sample including targets in the table above	2 x 400 µL
NTC	DNase/RNase-Free Water	2 x 400 µL

Table 2. Oligo Mix target organisms and detection channels.

Vial Name	Target	Channel
UTIP Oligo Mix 1	<i>Escherichia coli</i>	FAM/Green
	<i>Streptococcus agalactiae</i>	HEX/VIC/JOE/Yellow
	<i>Klebsiella oxytoca</i>	ROX/Texas Red/Orange
	Human RNase P (IC)	CY5/Red
UTIP Oligo Mix 2	<i>Staphylococcus saprophyticus</i>	FAM/Green
	<i>Serratia marcescens</i>	HEX/VIC/JOE/Yellow
	<i>Proteus mirabilis</i>	ROX/Texas Red/Orange
	<i>Aerococcus urinae</i>	CY5/Red
UTIP Oligo Mix 3	<i>Treponema pallidum</i>	FAM/Green
	<i>Enterobacter cloacae</i>	HEX/VIC/JOE/Yellow
	<i>Pseudomonas aeruginosa</i>	ROX/Texas Red/Orange
	<i>Citrobacter freundii</i>	CY5/Red

UTIP Oligo Mix 4	<i>Klebsiella aerogenes</i>	FAM/Green
	<i>Klebsiella pneumoniae</i>	HEX/VIC/JOE/Yellow
	<i>Morganella morganii</i>	ROX/Texas Red/Orange
	<i>Corynebacterium urealyticum</i>	CY5/Red
UTIP Oligo Mix 5	<i>Enterococcus faecium</i>	FAM/Green
	<i>Enterococcus faecalis</i>	HEX/VIC/JOE/Yellow
	<i>Acinetobacter baumannii</i>	ROX/Texas Red/Orange
	<i>Proteus vulgaris</i>	CY5/Red
UTIP Oligo Mix 6	<i>Staphylococcus aureus</i>	FAM/Green
	-	HEX/VIC/JOE/Yellow
	<i>Ureaplasma (Ureaplasma urealyticum/parvum)</i>	ROX/Texas Red/Orange
	-	CY5/Red
UTIP Oligo Mix 7	<i>Candida albicans</i>	FAM/Green
	<i>Candida glabrata</i>	HEX/VIC/JOE/Yellow
	<i>Candida parapsilosis</i>	ROX/Texas Red/Orange
	<i>Providencia stuartii</i>	CY5/Red
UTIP Oligo Mix 8	<i>Candida tropicalis</i>	FAM/Green
	<i>Candida krusei</i>	HEX/VIC/JOE/Yellow
	<i>Candida auris</i>	ROX/Texas Red/Orange
	<i>Bacillus atrophaeus</i> (EC)	CY5/Red

The oligonucleotide set targeting the human *RNase P* mRNA (Internal Control: IC) and *Bacillus atrophaeus* (External Control: EC) are used to monitor sampling, nucleic acid extraction, and inhibition of qPCR. The kit also contains negative and positive control templates for evaluating the contamination and the qPCR reagent stability, respectively.

4. EQUIPMENT and MATERIALS REQUIRED but NOT PROVIDED

- 2-8°C Refrigerator
- ≤ -20°C Freezer
- ≤ -70°C Freezer (Optional)
- Vortex mixer
- Benchtop centrifuge with rotor for 1.5 mL tubes
- Benchtop mini centrifuge with rotor for PCR strips
- Benchtop plate centrifuge
- Biological Safety Cabinet (BSC)
- PCR cabinet for PCR Setup
- Adjustable Micropipettes: 1-10, 10-100, 100-1000 µL
- Sterile DNase/RNase free micropipettes tips - Compatible with the micropipettes
- Cold tube rack for microfuge tubes (1.5/2 mL) and for PCR tubes (0.1/0.2 mL)
- Disposable, powder-free, nitrile gloves
- Disposable (preferably) laboratory coat
- Surface decontaminants - Freshly diluted 10% bleach solution (0.5% NaClO)
- Applied Biosystems QuantStudio 5, 7, and 12K with Design & Analysis software and consumables
- Bio-Rad CFX96 Touch™/CFX96™ Dx/CFX Opus 96™/CFX Opus 96™ Dx/CFX384 Touch™/CFX Opus 384™ with Maestro software v1.1 and consumables
- Qiagen Rotor-Gene Q 5plex Platform with Rotor-Gene Q series software v2.1.0.9 and consumables
- Roche LightCycler 480 with software and consumables

5. WARNING and PRECAUTIONS

- The **MarinaBiolab Urinary Tract Infections Panel PCR Kit** is designed for research use only and should be used by professionally trained, qualified staff only. All work should be performed using Good Laboratory Practices.
- Biological material used for extraction of nucleic acid should be handled as potentially infectious. When handling biological material appropriate safety precautions are recommended (do not pipet by mouth; wear disposable gloves while handling biological material and performing the test; disinfect hands when finished the test).
- Biological material should be inactivated before disposal (e.g., in an autoclave). Disposables should be autoclaved or incinerated after use.
- Spillage of potentially infectious materials should be removed immediately with absorbent paper tissue and the contaminated areas swabbed with a suitable standard disinfectant or 70% alcohol. Material used to clean spills, including gloves, should be inactivated before disposal (e.g., in an autoclave).
- Disposal of all samples, unused reagents and waste should be in accordance with country, federal, state, and local regulations.
- Microbial contamination of the reagents while taking aliquots should be avoided. It is recommended to use sterile one-way pipettes and tips. Reagents that look cloudy or show any signs of microbial contamination must not be used.
- The kit should be stored away from nucleic acid sources and qPCR amplicons.
- Always check the expiration date on the kit. Do not use expired or incorrectly stored kit.
- The components in the kit should not be mixed with components with different lot numbers or chemicals of the same name but from different manufacturers.
- Kit components should be mixed by gently shaking before use.
- A common concern with PCR-based assays is false positive results caused by contamination of the work area with PCR amplicon. To prevent amplicon contamination:
 - It shall be ensured that separate work areas with their own apparatus are available.
 - Do not open the reaction tubes/plates post amplification to avoid contamination with amplicons.
 - Discard used tubes/plates in a biohazard container immediately after the run has completed.
 - Avoid excessive handling of tubes/plates after test runs.
 - Change gloves after handling a used tubes/plate.

6. HANDLING, STORAGE, and STABILITY

- The **MarinaBiolab Urinary Tract Infections Panel PCR Kit** is shipped on dry ice. If any component except qPCR Master Mix of the kit is not frozen on arrival, or if the outer packaging has been compromised during shipment, please contact **MarinaBiolab** or the local distributors as soon as possible.
- All components should be stored between -25°C and -15°C upon arrival.
- Repeated freezing and thawing of the kit components may result in lower detection quality. The kit can undergo up to 15 freeze/thaw cycles without affecting performance.
- When stored under the specified storage conditions, the kit is stable until the stated expiration date printed on the package. The expiration date of the kit is 12 months from date of manufacture.
- All components must be thawed at ambient temperature for a minimum of 30 minutes before use.
- It is recommended that all components should be kept on ice when setting up the assay mixes.
- The primer and probe mixes contain fluorophore labeled probes and should be protected from direct sunlight or long-term ambient light.
- Do not use expired or incorrectly stored components.

7. TEST PROCEDURE

7.1. Sample Preparation and Nucleic Acid Extraction

The sample material for the isolation of nucleic acid must be sent in appropriate cell collection systems. The performance of the kit strongly depends on the amount and quality of the extracted nucleic acid. It must be ensured that the system used for nucleic acid extraction is compatible with real-time PCR technology.

If the established standard method of the lab is used for nucleic acid isolation, it must be validated by the user.

For frozen samples or frozen extracted nucleic acid, only thaw the number of specimen extracts that will be tested in a single day.

Do not freeze/thaw extracted nucleic acid more than once before testing as each freeze/thaw cycle can decrease the nucleic acid quality. For optimal results, use it directly.

7.2. PCR Reaction Preparation and Processing

- Completely thaw the components at room temperature for a minimum of 30 minutes before each use.
- Place all components on ice once thawed during the whole test procedure.
- Determine the number of reactions and create the PCR plate plan.
- Include the following reactions to the plan:
 - Reactions for each test sample and extraction negative control.
 - PCR control reactions:
 - Positive Control (included in the kit)
 - Negative (No Template) Control (NTC) (included in the kit)
 - No Template Addition Control (NRC)
- Vortex and spin down briefly the components before each use.
- Combine the following components for the number of reactions required plus 10% overage to compensate for pipetting errors:

Table 3. Reaction set-up.

Reaction Mix Component	1X Reaction (µL) per well
qPCR Master Mix	5 µL
UTIP Oligo Mix 1-8	2.5 µL
Template Nucleic Acid	2.5 µL
Total Reaction Volume	10 µL

- Add 5 µL of qPCR Master Mix and 2.5 µL of UTIP Oligo Mix 1-8 into PCR tubes.
- Add 2.5 µL of the isolated sample into the individual tubes.
- The final reaction mix volume is 10 µL.
- Close the tubes, centrifuge briefly, insert tubes into the real-time PCR instrument and amplify according to the following PCR profile.

Table 4. Amplification profile.

Step	Number of Cycles	Temperature	Time	Data Collection
Initial Denaturation	1	95 °C	10 sec	FAM/Green, HEX/VIC/JOE/Yellow, ROX/Texas Red/Orange, CY5/Red
Denaturation	40	95 °C	5 sec	
Annealing/Extension		55 °C	15 sec	

8. INTERPRETATION OF RESULTS

MarinaBiolab Urinary Tract Infections Panel PCR Kit provides a qualitative result for the presence (Detected) or absence (Not Detected) of the target genes.

8.1. Calculation of Cq Values and Instrument-Specific Requirements

Perform the following instrument settings before evaluating the results.

Table 5. Instrument-specific requirements before evaluating the results.

Instrument	Threshold Level	Other Settings
CFX96 Touch™/CFX96™ Dx/CFX Opus 96™/CFX Opus 96™ Dx/CFX384 Touch™/CFX Opus 384™ (Bio-Rad)	500 RFU	-
Rotor-Gene Q Splex Platform (QIAGEN)	0.02 RFU	Dynamic Tube: Active Slope Correct: Active Outlier Removal: 0
QuantStudio™ 5, 7 and 12K (Applied Biosystems™)	Auto	-
Roche LightCycler 480 (Roche)	Auto	-

The shape of the amplification curves should be examined. If a Cq value is assigned to a sample by the instruments' software and the curve is sigmoidal, the Cq value can be used in the final evaluation. *Non-sigmoidal curves should be recorded as negative.*

The result is recorded as positive if $Cq \leq 38$ or as established by your lab.

8.2. Overall Validity of Detection

Table 6. Expected performance of controls.

Control Type	Used to Monitor	Signal	
		Target Channel	Internal/External Control Channel
Negative Control	Cross-contamination during extraction and reaction setup	-	-
No template addition	Reagent and/or environmental contamination	-	-
Positive Control	qPCR reaction setup and reagent integrity	+	+
Internal/External Control	To monitor the integrity of nucleic acid extraction and qPCR from each specimen	Not applicable	+

Before analyzing samples results, we recommend verifying if the real-time PCR test is valid. Thus, for each run, please confirm if the results for Positive and Negative controls performed as expected, according to the following criteria:

Table 7. Run validity/positive and negative control pass criteria.

Positive Control		Negative Control		Results	Recommendation
Target Channel	Internal/External Control Channel	Target Channel	Internal/External Control Channel		
+	+	-	-	VALID	Continue to result interpretation of samples.
Any of them is Negative		Not considered		INVALID	Contact the manufacturer, renew the reagents, and repeat the reaction.
Not considered		Any of them is Positive		INVALID	Repeat analysis, paying attention to "Warnings and Precautions" in IFU.

If any control does not perform as described above, the run is considered invalid, and the test is repeated. If the problem persists, contact the manufacturer.

If all the controls are valid, proceed to the interpretation of the results.

8.3. Interpretation of Unknown Specimen Results

The data produced by the instruments can manually be evaluated and reported using their software.

Table 8. Interpretation of unknown specimen results for DNA pathogens.

DNA Pathogens	Internal Control (<i>RNase P</i>)	External Control (<i>Bacillus atrophaeus</i>)	Results	Interpretation
Positive (+) (Cq<38)	Positive (+) (Cq<38)	Positive (+) (Cq<38)	Positive for Target	Target DNA is detected
Positive (+) (Cq<38)	Negative (-) (Cq≥38 or N/A)	Positive (+) (Cq<38)	Positive for Target	Target DNA is detected
Positive (+) (Cq<38)	Negative (-) (Cq≥38 or N/A)	Negative (-) (Cq≥38 or N/A)	Invalid	Repeat test by re-extracting the sample. If the repeat result remains invalid, consider collecting a new sample.
Positive (+) (Cq<38)	Positive (+) (Cq<38)	Negative (-) (Cq≥38 or N/A)	Invalid	Repeat test by re-extracting the sample. If the repeat result remains invalid, consider collecting a new sample.
Negative (-) (Cq≥38 or N/A)	Positive (+) (Cq<38)	Positive (+) (Cq<38)	Negative for Target	Target DNA is not detected
Negative (-) (Cq≥38 or N/A)	Negative (-) (Cq≥38 or N/A)	Positive (+) (Cq<38)	Negative for Target	Target DNA is not detected
Negative (-) (Cq≥38 or N/A)	Negative (-) (Cq≥38 or N/A)	Negative (-) (Cq≥38 or N/A)	Invalid	Repeat test by re-extracting the sample. If the repeat result remains invalid, consider collecting a new sample.
Negative (-) (Cq≥38 or N/A)	Positive (+) (Cq<38)	Negative (-) (Cq≥38 or N/A)	Invalid	Repeat test by re-extracting the sample. If the repeat result remains invalid, consider collecting a new sample.

9. ASSAY LIMITATIONS

- The ***MarinaBiolab Urinary Tract Infections Panel PCR Kit*** is intended for use by professionally trained, qualified staff only.
- A false negative result may occur if a specimen is improperly collected, transported, or handled. False negative results may also occur if amplification inhibitors are present in the specimen or if inadequate numbers of organisms are present in the specimen.
- Spontaneous mutations within the target sequences may result in failure to detect the target sequence. While this risk is mitigated in the test's design, if failure to detect the target is expected it is recommended to test the specimen with a different test that detects different target sequences from the target's genome.
- There is a risk of false positive results due to cross-contamination by target viruses and/or bacteria, their nucleic acids or amplified product, or from non-specific signals in the assay. Attention should be given to the handling of consumables under the Warnings and Precautions section to help minimize this risk.
- This assay is a qualitative test and does not provide a quantitative assessment of the concentration of the detected organism.
- All instruments (e.g., pipettes, real-time cyclers) must be calibrated according to the manufacturer's instructions.

10. PERFORMANCE CHARACTERISTICS

10.1. Analytical Sensitivity (Limit of Detection, LoD)

The LoD was defined as the concentration at which the test produces a positive result >95% of the time. Serial dilutions of the strains were tested and the initial tentative LoD confirmed with twenty (20) replicates. To ensure the accuracy of the LoD determination, if the initial detection rate was 100%, a further twenty (20) replicates were performed at the next lower concentration until ≤95% was achieved. For nucleic acid extraction, simulated research matrix was spiked with strains and loaded onto the Automatic Nucleic Acids Extraction Instrument. The tests were carried out using the CFX96 Touch™ (Bio-Rad) Real-Time PCR system. The confirmed LoDs for the strains tested and the corresponding LoDs for the **MarinaBiolab Urinary Tract Infections Panel PCR Kit** reportable targets are shown in Table 9 below.

Table 9. Summary of LoD study results.

Analyte	Isolate ID/Source	LoD Concentration (copies/mL)	Detected/Total
<i>Escherichia coli</i>	ATCC 25922	3.5E+01 copies/mL	20/20 100%
<i>Streptococcus agalactiae</i>	ATCC 12386	6.7E+01 copies/mL	19/20 95%
<i>Klebsiella oxytoca</i>	ATCC 700324	2.6E+01 copies/mL	20/20 100%
<i>Staphylococcus saprophyticus</i>	Zeptomatrix 0804014	5.7E+01 copies/mL	20/20 100%
<i>Serratia marcescens</i>	ATCC 29021	2.1E+02 copies/mL	20/20 100%
<i>Proteus mirabilis</i>	Zeptomatrix 0801544	2.1E+02 copies/mL	20/20 100%
<i>Aerococcus urinae</i>	ATCC 51268	4.5E+01 copies/mL	20/20 100%
<i>Treponema pallidum</i>	ATCC BAA-2642SD	1.8E+07 copies/mL	20/20 100%
<i>Enterobacter cloacae</i>	Zeptomatrix 0801830	7.4E+01 copies/mL	19/20 95%
<i>Pseudomonas aeruginosa</i>	ATCC 27853	6.7E+02 copies/mL	20/20 100%
<i>Citrobacter freundii</i>	Zeptomatrix 0801563	4.2E+01 copies/mL	20/20 100%
<i>Klebsiella aerogenes</i>	ATCC 13048	2.4E+02 copies/mL	20/20 100%
<i>Klebsiella pneumoniae</i>	NCTC 13465	3.0E+01 copies/mL	20/20 100%

<i>Morganella morganii</i>	Zeptomatrix 0804010	4.8E+01 copies/mL	20/20 100%
<i>Corynebacterium urealyticum</i>	ATCC 43044	4.1E+01 copies/mL	20/20 100%
<i>Enterococcus faecium</i>	ATCC BAA-2127	4.5E+01 copies/mL	20/20 100%
<i>Enterococcus faecalis</i>	Zeptomatrix 0804216	3.6E+02 copies/mL	20/20 100%
<i>Acinetobacter baumannii</i>	ATCC 19606	1.7E+02 copies/mL	20/20 100%
<i>Proteus vulgaris</i>	Zeptomatrix 0810290CF	1.5E+02 copies/mL	20/20 100%
<i>Staphylococcus aureus</i>	ATCC 10832	5.5E+01 copies/mL	20/20 100%
<i>Ureaplasma urealyticum</i>	ATCC 27618	5.0E+01 copies/mL	20/20 100%
<i>Ureaplasma parvum</i>	ATCC 27815	4.0E+01 copies/mL	20/20 100%
<i>Providencia stuartii</i>	Zeptomatrix 0810452CF	4.5E+01 copies/mL	20/20 100%
<i>Candida albicans</i>	ATCC 10231	3.4E+02 copies/mL	20/20 100%
<i>Candida glabrata</i>	ATCC 90030	4.4E+01 copies/mL	20/20 100%
<i>Candida parapsilosis</i>	ATCC 22019	5.8E+01 copies/mL	20/20 100%
<i>Candida tropicalis</i>	ATCC 750	5.7E+01 copies/mL	20/20 100%
<i>Candida krusei</i>	ATCC 2159	6.8E+01 copies/mL	20/20 100%
<i>Candida auris</i>	ATCC MYA-5003	7.2E+01 copies/mL	19/20 95%

10.2. Device Equivalence Study

Device equivalence study was carried out to observe the differences between the results to be obtained using the kit in different instruments. For this purpose, the same LoD determination study was performed again with the Bio-Rad CFX96™ Dx/CFX Opus 96™/CFX Opus 96™ Dx/CFX384 Touch™/CFX Opus 384™, Applied Biosystems QuantStudio 5, 7, and 12K, Qiagen Rotor-Gene Q

5plex Platform and Roche LightCycler 480. Similar test results were obtained with the 1x LoD concentration level of the targets in the “device equivalence study” performed with the other instruments.

10.3. Analytical Reactivity (Inclusivity)

10.3.1. In-Silico Analytical Reactivity

BLAST search of the oligonucleotides was performed on the *Escherichia coli*, *Streptococcus agalactiae*, *Klebsiella oxytoca*, *Staphylococcus saprophyticus*, *Serratia marcescens*, *Proteus mirabilis*, *Aerococcus urinae*, *Treponema pallidum*, *Enterobacter cloacae*, *Pseudomonas aeruginosa*, *Citrobacter freundii*, *Klebsiella aerogenes*, *Klebsiella pneumoniae*, *Morganella morganii*, *Corynebacterium urealyticum*, *Enterococcus faecium*, *Enterococcus faecalis*, *Acinetobacter baumannii*, *Proteus vulgaris*, *Staphylococcus aureus*, *Ureplasma (Ureaplasma urealyticum/parvum)*, *Providencia stuartii*, *Candida albicans*, *Candida glabrata*, *Candida parapsilosis*, *Candida tropicalis*, *Candida krusei*, and *Candida auris* genome sequences available in the NCBI database, using the Primer-BLAST tool of NCBI.

The aggregated result of all in-silico analyzes performed in NCBI database is provided in Table below. The melting temperatures (T_m) of the oligonucleotide sequences with 1-base mismatch, are still higher than the annealing temperature specified in the PCR cycle parameters of the kit. Hence, the single mismatches in the sequences are not expected to affect the inclusivity of the test.

Table 10. In-silico analysis results performed in the NCBI database.

Target	Primer	Total number of target sequences	Ratio of the sequences without mismatch	Ratio of the sequences with 1 base mismatch	Ratio of the sequences with 2 base mismatches	Ratio of the sequences with 3 base mismatches
Escherichia coli	Sense Primer	5.547	99.25%	0.75%	0.00%	0.00%
Escherichia coli	Antisense Primer	5.579	99.65%	0.35%	0.00%	0.00%
Escherichia coli	Hydrolysis Probe	5.579	99.78%	0.22%	0.00%	0.00%
Streptococcus agalactiae	Sense Primer	226	99.95%	0.05%	0.00%	0.00%
Streptococcus agalactiae	Antisense Primer	236	100.00%	0.00%	0.00%	0.00%
Streptococcus agalactiae	Hydrolysis Probe	236	100.00%	0.00%	0.00%	0.00%
Klebsiella oxytoca	Sense Primer	150	99.74%	0.26%	0.00%	0.00%
Klebsiella oxytoca	Antisense Primer	158	99.56%	0.44%	0.00%	0.00%
Klebsiella oxytoca	Hydrolysis Probe	158	99.83%	0.27%	0.00%	0.00%
Staphylococcus saprophyticus	Sense Primer	26	99.52%	0.48%	0.00%	0.00%
Staphylococcus saprophyticus	Antisense Primer	26	97.52%	2.48%	0.00%	0.00%
Staphylococcus saprophyticus	Hydrolysis Probe	26	99.74%	0.26%	0.00%	0.00%
Serratia marcescens	Sense Primer	296	99.80%	0.20%	0.00%	0.00%
Serratia marcescens	Antisense Primer	296	99.80%	0.20%	0.00%	0.00%
Serratia marcescens	Hydrolysis Probe	292	99.82%	0.18%	0.00%	0.00%
Proteus mirabilis	Sense Primer	164	100.00%	0.00%	0.00%	0.00%

Proteus mirabilis	Antisense Primer	164	100.00%	0.00%	0.00%	0.00%
Proteus mirabilis	Hydrolysis Probe	160	99.80%	0.20%	0.00%	0.00%
Aerococcus urinae	Sense Primer	80	100.00%	0.00%	0.00%	0.00%
Aerococcus urinae	Antisense Primer	80	100.00%	0.00%	0.00%	0.00%
Aerococcus urinae	Hydrolysis Probe	76	100.00%	0.00%	0.00%	0.00%
Treponema pallidum	Sense Primer	538	99.64%	0.36%	0.00%	0.00%
Treponema pallidum	Antisense Primer	538	99.64%	0.36%	0.00%	0.00%
Treponema pallidum	Hydrolysis Probe	538	99.50%	0.50%	0.00%	0.00%
Enterobacter cloacae	Sense Primer	683	99.63%	0.37%	0.00%	0.00%
Enterobacter cloacae	Antisense Primer	669	99.12%	0.88%	0.00%	0.00%
Enterobacter cloacae	Hydrolysis Probe	669	99.82%	0.18%	0.00%	0.00%
Pseudomonas aeruginosa	Sense Primer	1.162	99.75%	0.25%	0.00%	0.00%
Pseudomonas aeruginosa	Antisense Primer	1.167	99.79%	0.21%	0.00%	0.00%
Pseudomonas aeruginosa	Hydrolysis Probe	1.167	99.84%	0.16%	0.00%	0.00%
Citrobacter freundii	Sense Primer	175	99.70%	0.30%	0.00%	0.00%
Citrobacter freundii	Antisense Primer	175	99.70%	0.30%	0.00%	0.00%
Citrobacter freundii	Hydrolysis Probe	168	99.90%	0.10%	0.00%	0.00%
Klebsiella aerogenes	Sense Primer	83	98.52%	1.48%	0.00%	0.00%
Klebsiella aerogenes	Antisense Primer	82	97.11%	2.89%	0.00%	0.00%
Klebsiella aerogenes	Hydrolysis Probe	82	96.85%	3.15%	0.00%	0.00%
Klebsiella pneumoniae	Sense Primer	2.816	100.00%	0.00%	0.00%	0.00%
Klebsiella pneumoniae	Antisense Primer	2.711	100.00%	0.00%	0.00%	0.00%
Klebsiella pneumoniae	Hydrolysis Probe	2.711	99.66%	0.34%	0.00%	0.00%
Morganella morganii	Sense Primer	81	99.84%	0.16%	0.00%	0.00%
Morganella morganii	Antisense Primer	81	99.84%	0.16%	0.00%	0.00%
Morganella morganii	Hydrolysis Probe	80	99.52%	0.48%	0.00%	0.00%
Corynebacterium urealyticum	Sense Primer	30	100.00%	0.00%	0.00%	0.00%
Corynebacterium urealyticum	Antisense Primer	30	100.00%	0.00%	0.00%	0.00%
Corynebacterium urealyticum	Hydrolysis Probe	30	100.00%	0.00%	0.00%	0.00%
Enterococcus faecium	Sense Primer	552	98.68%	1.32%	0.00%	0.00%
Enterococcus faecium	Antisense Primer	555	98.68%	1.32%	0.00%	0.00%
Enterococcus faecium	Hydrolysis Probe	555	98.46%	1.54%	0.00%	0.00%

Enterococcus faecalis	Sense Primer	575	100.00%	0.00%	0.00%	0.00%
Enterococcus faecalis	Antisense Primer	578	100.00%	0.00%	0.00%	0.00%
Enterococcus faecalis	Hydrolysis Probe	578	99.89%	0.11%	0.00%	0.00%
Acinetobacter baumannii	Sense Primer	1.703	99.35%	0.65%	0.00%	0.00%
Acinetobacter baumannii	Antisense Primer	1.701	99.89%	0.21%	0.00%	0.00%
Acinetobacter baumannii	Hydrolysis Probe	1.701	99.47%	0.53%	0.00%	0.00%
Proteus vulgaris	Sense Primer	155	99.83%	0.17%	0.00%	0.00%
Proteus vulgaris	Antisense Primer	155	99.83%	0.17%	0.00%	0.00%
Proteus vulgaris	Hydrolysis Probe	157	99.85%	0.15%	0.00%	0.00%
Staphylococcus aureus	Sense Primer	2.491	99.65%	0.35%	0.00%	0.00%
Staphylococcus aureus	Antisense Primer	2.703	99.74%	0.26%	0.00%	0.00%
Staphylococcus aureus	Hydrolysis Probe	2.703	99.62%	0.38%	0.00%	0.00%
Ureplasma	Sense Primer	90	99.90%	0.10%	0.00%	0.00%
Ureplasma	Antisense Primer	90	99.90%	0.10%	0.00%	0.00%
Ureplasma	Hydrolysis Probe	88	99.90%	0.10%	0.00%	0.00%
Providencia stuartii	Sense Primer	23	100.00%	0.00%	0.00%	0.00%
Providencia stuartii	Antisense Primer	23	100.00%	0.00%	0.00%	0.00%
Providencia stuartii	Hydrolysis Probe	22	100.00%	0.00%	0.00%	0.00%
Candida auris	Sense Primer	501	100,00%	0,00%	0.00%	0.00%
Candida auris	Antisense Primer	501	100,00%	0,00%	0.00%	0.00%
Candida auris	Hydrolysis Probe	499	100,00%	0,00%	0.00%	0.00%
Candida krusei	Sense Primer	1.415	100%	0.00%	0.00%	0.00%
Candida krusei	Antisense Primer	1.415	100%	0.00%	0.00%	0.00%
Candida krusei	Hydrolysis Probe	1.415	100%	0.00%	0.00%	0.00%
Candida albicans	Sense Primer	3.629	99.69%	0.31%	0.00%	0.00%
Candida albicans	Antisense Primer	3.728	98.85%	2.25%	0.00%	0.00%
Candida albicans	Hydrolysis Probe	3.728	98.52%	2.48%	0.00%	0.00%
Candida parapsilosis	Sense Primer	2.559	99.74%	0.26%	0.00%	0.00%
Candida parapsilosis	Antisense Primer	2.463	100%	0.00%	0.00%	0.00%
Candida parapsilosis	Hydrolysis Probe	2.463	100%	0.00%	0.00%	0.00%
Candida tropicalis	Sense Primer	1.164	98.40%	2.60%	0.00%	0.00%
Candida tropicalis	Antisense Primer	1.906	97.83%	2.17%	0.00%	0.00%

Candida tropicalis	Hydrolysis Probe	1.906	97.12%	2.88%	0.00%	0.00%
Candida glabrata	Sense Primer	763	100%	0.00%	0.00%	0.00%
Candida glabrata	Antisense Primer	1.111	99.20%	0.80%	0.00%	0.00%
Candida glabrata	Hydrolysis Probe	1.111	99.64%	0.36%	0.00%	0.00%

10.3.2. Wet-Test Analytical Reactivity

The analytical reactivity (inclusivity) of the **MarinaBiolab Urinary Tract Infections Panel PCR Kit** was demonstrated with a comprehensive panel representing temporal, evolutionary, and geographic diversity for each of the target organisms.

Each sample was tested with the **MarinaBiolab Urinary Tract Infections Panel PCR Kit** in triplicate at an initial concentration 3-fold higher than the LoD determined for each analyte. In cases where the expected targets were not detected in one or more replicates, concentrations at a 3-fold higher level were evaluated.

The individual strains and concentrations at which positive test results were obtained for all three (3) replicates are presented by target organism in Table 11 below.

Table 11. Results of the wet inclusivity test.

Variant/Type/Subtype/Lineage/Genotype/Species	Isolate ID/Source	xLoD Detected
<i>Escherichia coli</i>	ATCC 25922	1x
<i>Streptococcus agalactiae</i>	ATCC 12386	1x
<i>Klebsiella oxytoca</i>	ATCC 700324	1x
<i>Staphylococcus saprophyticus</i>	Zeptomatrix 0804014	1x
<i>Serratia marcescens</i>	ATCC 29021	1x
<i>Proteus mirabilis</i>	Zeptomatrix 0801544	1x
<i>Aerococcus urinae</i>	ATCC 51268	1x
<i>Treponema pallidum</i>	ATCC BAA-2642SD	1x
<i>Enterobacter cloacae</i>	Zeptomatrix 0801830	1x
<i>Pseudomonas aeruginosa</i>	ATCC 27853	1x
<i>Citrobacter freundii</i>	Zeptomatrix 0801563	1x
<i>Klebsiella aerogenes</i>	ATCC 13048	1x
<i>Klebsiella pneumoniae</i>	NCTC 13465	1x
<i>Morganella morganii</i>	Zeptomatrix 0804010	1x
<i>Corynebacterium urealyticum</i>	ATCC 43044	1x
<i>Enterococcus faecium</i>	ATCC BAA-2127	1x
<i>Enterococcus faecalis</i>	Zeptomatrix 0804216	1x

<i>Acinetobacter baumannii</i>	ATCC 19606	1x
<i>Proteus vulgaris</i>	Zeptomatrix 0810290CF	1x
<i>Staphylococcus aureus</i>	ATCC 10832	1x
<i>Ureaplasma urealyticum</i>	ATCC 27618	1x
<i>Ureaplasma parvum</i>	ATCC 27815	1x
<i>Providencia stuartii</i>	Zeptomatrix 0810452CF	1x
<i>Candida albicans</i>	ATCC 10231	1x
<i>Candida glabrata</i>	ATCC 90030	1x
<i>Candida parapsilosis</i>	ATCC 22019	1x
<i>Candida tropicalis</i>	ATCC 750	1x
<i>Candida krusei</i>	ATCC 2159	1x
<i>Candida auris</i>	ATCC MYA-5003	1x

10.4. Analytical Specificity (Exclusivity)

10.4.1. In-Silico Analytical Specificity

Primers and probes intended for a target sequence may also attach to similar sequences if they closely match or differ by only a few base pairs from the non-targeted sequence. To ensure specificity to the target amplicon sequence, it's essential to screen the primers and probe against the reference database transcript or genome database for the intended templates, as well as any databases containing potential contaminating templates.

Table 12. The results of On-Panel and Off-Panel organisms tested for cross-reactivity.

On-Panel/Off-Panel	Name of the organism	Cross Reactivity*		
		Forward	Probe	Reverse
On-Panel	<i>Escherichia coli</i>	None	None	None
On-Panel	<i>Streptococcus agalactiae</i>	None	None	None
On-Panel	<i>Klebsiella oxytoca</i>	None	None	None
On-Panel	<i>Staphylococcus saprophyticus</i>	None	None	None
On-Panel	<i>Serratia marcescens</i>	None	None	None
On-Panel	<i>Proteus mirabilis</i>	None	None	None
On-Panel	<i>Aerococcus urinae</i>	None	None	None
On-Panel	<i>Treponema pallidum</i>	None	None	None
On-Panel	<i>Enterobacter cloacae</i>	None	None	None
On-Panel	<i>Pseudomonas aeruginosa</i>	None	None	None

On-Panel	<i>Citrobacter freundii</i>	None	None	None
On-Panel	<i>Klebsiella aerogenes</i>	None	None	None
On-Panel	<i>Klebsiella pneumoniae</i>	None	None	None
On-Panel	<i>Morganella morganii</i>	None	None	None
On-Panel	<i>Corynebacterium urealyticum</i>	None	None	None
On-Panel	<i>Enterococcus faecium</i>	None	None	None
On-Panel	<i>Enterococcus faecalis</i>	None	None	None
On-Panel	<i>Acinetobacter baumannii</i>	None	None	None
On-Panel	<i>Proteus vulgaris</i>	None	None	None
On-Panel	<i>Staphylococcus aureus</i>	None	None	None
On-Panel	<i>Ureaplasma urealyticum</i>	None	None	None
On-Panel	<i>Ureaplasma parvum</i>	None	None	None
On-Panel	<i>Providencia stuartii</i>	None	None	None
On-Panel	<i>Candida albicans</i>	None	None	None
On-Panel	<i>Candida glabrata</i>	None	None	None
On-Panel	<i>Candida parapsilosis</i>	None	None	None
On-Panel	<i>Candida tropicalis</i>	None	None	None
On-Panel	<i>Candida krusei</i>	None	None	None
On-Panel	<i>Candida auris</i>	None	None	None
Off-Panel	<i>Staphylococcus epidermidis</i>	None	None	None
Off-Panel	<i>Staphylococcus haemolyticus</i>	None	None	None
Off-Panel	<i>Staphylococcus lugdunensis</i>	None	None	None
Off-Panel	<i>Streptococcus dysgalactiae</i>	None	None	None
Off-Panel	<i>Streptococcus pyogenes</i>	None	None	None
Off-Panel	<i>Fusarium solani</i>	None	None	None
Off-Panel	<i>Microsporum spp.</i>	None	None	None
Off-Panel	<i>Trichophyton spp.</i>	None	None	None
Off-Panel	<i>Acinetobacter iwoffi</i>	None	None	None
Off-Panel	<i>Acinetobacter nosocomialis</i>	None	None	None
Off-Panel	<i>Stenotrophomonas maltophilia</i>	None	None	None
Off-Panel	<i>Moraxella catarrhalis</i>	None	None	None
Off-Panel	<i>Pasteurella stomatis</i>	None	None	None

Off-Panel	<i>Epidermophyton floccosum</i>	None	None	None
Off-Panel	<i>Finegoldia magna</i>	None	None	None
Off-Panel	<i>Bartonella henselae</i>	None	None	None
Off-Panel	<i>Haemophilus influenzae</i>	None	None	None
Off-Panel	<i>Candida sojae</i>	None	None	None
Off-Panel	<i>Candida oregonensis</i>	None	None	None
Off-Panel	<i>Malessezia restricta</i>	None	None	None
Off-Panel	<i>Peptoniphilus harei</i>	None	None	None
Off-Panel	<i>Peptoniphilus ivorii</i>	None	None	None
Off-Panel	<i>Peptostreptococcus prevotii</i>	None	None	None
Off-Panel	<i>Peptostreptococcus anaerobius</i>	None	None	None
Off-Panel	<i>Listeria monocytogenes</i>	None	None	None
Off-Panel	<i>Candida lusitanae</i>	None	None	None
Off-Panel	<i>Kingella kingae</i>	None	None	None
Off-Panel	<i>Chlamydia trachomatis</i>	None	None	None
Off-Panel	<i>Legionella dumoffii</i>	None	None	None
Off-Panel	<i>Corynebacterium diphtheriae</i>	None	None	None
Off-Panel	<i>Neisseria meningitidis</i>	None	None	None

* Homology should be <80% between the cross-reactivity microorganisms and the test primers/ probe(s).

** In silico sequence analysis indicates the potential for cross-reactivity of Bordetella pertussis with certain strains of Bordetella bronchiseptica.

10.4.2. Wet-Test Analytical Specificity

The potential for non-specific amplification by assays for detection of analytes was evaluated by testing high concentrations of organisms or nucleic acids with the **MarinaBiolab Urinary Tract Infections Panel PCR Kit**. On-panel organisms were tested to assess the potential for intra-panel cross-reactivity, and off-panel organisms were tested to evaluate panel specificity. Off-panel organisms included normal flora and pathogens that may be present in specimens as well as near-neighbors or species genetically related to the organisms detected by the **MarinaBiolab Urinary Tract Infections Panel PCR Kit**. The concentration of organism tested (in triplicate) was at least 1.0E+06 CFU/mL for bacteria, fungi and parasite, and at least 1.0E+05 unit/mL for viruses. For the few organisms of interest that were not available for laboratory testing, results of in silico analysis of the organism's whole genome sequences are indicated. The on-panel and off-panel organisms tested are shown in Table 13 and Table 14.

Table 13. On-Panel organisms tested for evaluation of **MarinaBiolab Urinary Tract Infections Panel PCR Kit** analytical specificity.

Organism	Isolate ID/Source	Cross Reactivity Detected
<i>Escherichia coli</i>	ATCC 25922	None

<i>Streptococcus agalactiae</i>	ATCC 12386	None
<i>Klebsiella oxytoca</i>	ATCC 700324	None
<i>Staphylococcus saprophyticus</i>	Zeptomatrix 0804014	None
<i>Serratia marcescens</i>	ATCC 29021	None
<i>Proteus mirabilis</i>	Zeptomatrix 0801544	None
<i>Aerococcus urinae</i>	ATCC 51268	None
<i>Treponema pallidum</i>	ATCC BAA-2642SD	None
<i>Enterobacter cloacae</i>	Zeptomatrix 0801830	None
<i>Pseudomonas aeruginosa</i>	ATCC 27853	None
<i>Citrobacter freundii</i>	Zeptomatrix 0801563	None
<i>Klebsiella aerogenes</i>	ATCC 13048	None
<i>Klebsiella pneumoniae</i>	NCTC 13465	None
<i>Morganella morganii</i>	Zeptomatrix 0804010	None
<i>Corynebacterium urealyticum</i>	ATCC 43044	None
<i>Enterococcus faecium</i>	ATCC BAA-2127	None
<i>Enterococcus faecalis</i>	Zeptomatrix 0804216	None
<i>Acinetobacter baumannii</i>	ATCC 19606	None
<i>Proteus vulgaris</i>	Zeptomatrix 0810290CF	None
<i>Staphylococcus aureus</i>	ATCC 10832	None
<i>Ureaplasma urealyticum</i>	ATCC 27618	None
<i>Ureaplasma parvum</i>	ATCC 27815	None
<i>Providencia stuartii</i>	Zeptomatrix 0810452CF	None
<i>Candida albicans</i>	ATCC 10231	None
<i>Candida glabrata</i>	ATCC 90030	None
<i>Candida parapsilosis</i>	ATCC 22019	None
<i>Candida tropicalis</i>	ATCC 750	None
<i>Candida krusei</i>	ATCC 2159	None
<i>Candida auris</i>	ATCC MYA-5003	None

Table 14. Off-Panel organisms were tested for evaluation of *MarinaBiolab Urinary Tract Infections Panel PCR Kit* analytical specificity.

Organism	Isolate ID/Source	Cross Reactivity Detected
----------	-------------------	---------------------------

<i>Acinetobacter calcoaceticus</i>	ATCC 23055	None
<i>Staphylococcus epidermidis</i>	Zeptomatrix 0804281	None
<i>Staphylococcus haemolyticus</i>	Zeptomatrix 0801591	None
<i>Staphylococcus lugdunensis</i>	Zeptomatrix 0801555	None
<i>Streptococcus dysgalactiae</i>	Zeptomatrix 0801516	None
<i>Streptococcus pyogenes</i>	Zeptomatrix 0801512	None
<i>Fusarium solani</i>	Zeptomatrix 0801806	None
<i>Acinetobacter iwoffii</i>	Zeptomatrix 0801909	None
<i>Stenotrophomonas maltophilia</i>	Zeptomatrix 0801569	None
<i>Moraxella catarrhalis</i>	Zeptomatrix 0801509	None
<i>Candida sojae</i>	Zeptomatrix 0801825	None
<i>Listeria monocytogenes</i>	Zeptomatrix 0804339	None
<i>Chlamydia trachomatis</i>	Zeptomatrix 0801775	None
<i>Acinetobacter baumannii</i>	ATCC 19606	None
<i>Legionella pneumophila</i>	Zeptomatrix 0801530	None
<i>Neisseria gonorrhoeae</i>	ATCC 19424	None
<i>Neisseria meningitidis</i>	ATCC 13090	None
<i>Aspergillus flavus</i>	Zeptomatrix 0801598	None

10.5. Interferences

The ability of endogenous or exogenous substances that could be present in research samples (or introduced during sample collection and handling) to interfere with accurate detection of analytes was evaluated with select direct testing on the **MarinaBiolab Urinary Tract Infections Panel PCR Kit** and extrapolated from the interference evaluation of the **MarinaBiolab Urinary Tract Infections Panel PCR Kit**.

Potentially interfering substances were evaluated using contrived samples spiked with substance. Results from samples containing a substance were compared to results from control samples without substance. The substances tested included endogenous substances that may be found in sample at normal or elevated levels (e.g., blood, mucus/mucin, human genomic DNA), various commensal or infectious microorganisms, medications, washes or topical applications, various swabs and transport media for sample collection, and substances used to clean, decontaminate, or disinfect work areas. Each substance was added to contrived samples containing representative organisms at concentrations near (3x) LoD. The concentration of substance added to the samples was equal to or greater than the highest level expected to be in research sample and each sample was tested in triplicate.

None of the substances were shown to interfere with the **MarinaBiolab Urinary Tract Infections Panel PCR Kit**.

Table 15. Evaluation of potentially interfering substances on the **MarinaBiolab Urinary Tract Infections Panel PCR Kit**.

Substance Tested	Concentration Tested	Observed Interference
Endogenous Substances		
Whole Blood	10% v/v	No Interference
Human serum	5% v/v	No Interference
Human Urine	-	No Interference
Competitive Microorganisms		
<i>Escherichia coli</i>	1.0E+06 CFU/mL	No Interference
<i>Streptococcus agalactiae</i>	1.0E+06 CFU/mL	No Interference
<i>Klebsiella oxytoca</i>	1.0E+06 CFU/mL	No Interference
<i>Staphylococcus saprophyticus</i>	1.0E+06 CFU/mL	No Interference
<i>Serratia marcescens</i>	1.0E+06 CFU/mL	No Interference
<i>Proteus mirabilis</i>	1.0E+06 CFU/mL	No Interference
<i>Aerococcus urinae</i>	1.0E+06 CFU/mL	No Interference
<i>Treponema pallidum</i>	1.0E+06 CFU/mL	No Interference
<i>Enterobacter cloacae</i>	1.0E+06 CFU/mL	No Interference
<i>Pseudomonas aeruginosa</i>	1.0E+06 CFU/mL	No Interference
<i>Citrobacter freundii</i>	1.0E+06 CFU/mL	No Interference
<i>Klebsiella aerogenes</i>	1.0E+06 CFU/mL	No Interference
<i>Klebsiella pneumoniae</i>	1.0E+06 CFU/mL	No Interference
<i>Morganella morganii</i>	1.0E+06 CFU/mL	No Interference
<i>Corynebacterium urealyticum</i>	1.0E+06 CFU/mL	No Interference
<i>Enterococcus faecium</i>	1.0E+06 CFU/mL	No Interference
<i>Enterococcus faecalis</i>	1.0E+06 CFU/mL	No Interference
<i>Acinetobacter baumannii</i>	1.0E+06 CFU/mL	No Interference
<i>Proteus vulgaris</i>	1.0E+06 CFU/mL	No Interference
<i>Staphylococcus aureus</i>	1.0E+06 CFU/mL	No Interference
<i>Ureaplasma urealyticum</i>	1.0E+06 CFU/mL	No Interference
<i>Ureaplasma parvum</i>	1.0E+06 CFU/mL	No Interference
<i>Providencia stuartii</i>	1.0E+06 CFU/mL	No Interference
<i>Candida albicans</i>	1.0E+06 CFU/mL	No Interference
<i>Candida glabrata</i>	1.0E+06 CFU/mL	No Interference
<i>Candida parapsilosis</i>	1.0E+06 CFU/mL	No Interference

<i>Candida tropicalis</i>	1.0E+06 CFU/mL	No Interference
<i>Candida krusei</i>	1.0E+06 CFU/mL	No Interference
<i>Candida auris</i>	1.0E+06 CFU/mL	No Interference
Exogenous Substances		
Feminine Spray/talcum powder	5% v/v	No Interference
Phenazopyridine Hydrochloride (Pyridium)	10 µg/mL	No Interference
Ascorbic acid	0.6 mmol/L	No Interference
High pH	pH = 8.0	No Interference
Low pH	pH = 4.0	No Interference
<u>Antibiotic Pool</u> Amoxicillin trihydrate Metronidazole Tetracycline Hydrochloride Sodium Cefotaxime	1 mg/mL (Each)	No Interference
Specimen Collection Materials		
Urine Tubes (BD Vacutainer® 364992)	N/A	No Interference
Starplex™ Scientific Urine Preservative Tube (22046414)	N/A	No Interference

11. TROUBLESHOOTING

Problem	Cause	Solution
Target-specific and/or IC signals are detected in the Negative Control well.	Contamination from the environment, contamination of extraction and/or qPCR reagents, or well-to-well cross contamination. The signal is not true target amplification, but background curves generated by the software of the qPCR instrument.	Repeat the qPCR with new reagents. Follow the general rules of GLP in a PCR lab (e.g., Decontaminate all surfaces and instruments with sodium hypochlorite or ethanol. Ensure that filters tips are used during the procedure and changed between samples.). It is recommended to set up the qPCR reactions in a separate area, where no RNA/DNA is handled and with equipment designated for pre-PCR activities. Ignore the Cq value of NTC if the amplification curve looks not real but background noise. If the problem persists, contact Technical Support.
No IC signal is detected, but target-specific signal is detected in sample wells.	A high copy number of target nucleic acid exists in samples, resulting in preferential amplification of the target-specific nucleic acid.	No action is required. The result is considered positive.
The Positive Control did not meet the criteria set for acceptable values of the kit. The assay is invalid.	Positive Control was not stored at the recommended conditions. Kit shelf-life expired.	Check the kit label for storage conditions and expiration date. Replace the Positive Control. Use a new kit if necessary.
High Cq values observed for repeated samples.	Frozen samples were not mixed properly after thawing. Degraded nucleic acids.	Make sure, thaw frozen samples with mild agitation to ensure thorough mixing. Ensure that samples are stored correctly and not subjected to multiple freeze-thaw cycles
Target-specific and/or IC signal detected after 38 Cycles in Positive Control.	Incorrect qPCR set-up or the kit reagents may have been compromised (e.g., improper storage or more than 15 freeze-thaw cycles).	Replace the control. If the problem persists, contact Technical Support.
No target-specific and IC signal is detected in sample wells.	Sampling, extraction, or inhibition problem.	Dilute the nucleic acid isolate 1/10 and repeat the qPCR. If the diluted sample does not give a positive result in the IC channel, request for a new sample and repeat the NA extraction. Repeat the NA extraction and the qPCR. Request for a new sample, repeat the NA extraction and the qPCR. If the problem persists, contact Technical Support.

12. EXPLANATION of SYMBOLS

Symbol	Title of Symbol	Symbol	Title of Symbol
	Research Use Only		Use-by date
	Manufacturer		Batch code
	Negative control		Non-sterile
	Positive control		Consult instructions for use or consult electronic instructions for use
	Control		Caution
	Temperature limit		Catalogue number
	Keep away from sunlight		Do not use if package is damaged and consult instructions for use
	Keep dry		Keep upright
	Contains sufficient for <n> tests		Protect from heat and radioactive sources

Custom care and technical support

Tel: +1 510 579-5802

e-mail customer care: accounting@marinabiolab.com

e-mail Technical Support: rd@marinabiolab.com



MarinaBiolab LTD.

Address: Hürriyet Mah. Erler So. No: 4/1A Nexus No: 13 Kartal/Istanbul/Turkey

For research use only (RUO)! Not for use in diagnostic procedures.

© 2024 MarinaBiolab LTD.; all rights reserved.