



INSTRUCTION FOR USE

ABR-2 Panel PCR Kit

For Research Use Only



100



MBL ABR2017



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)	16
10.3.1.	In-Slico Analytical Reactivity	16
10.3.2.	Wet-Test Analytical Reactivity	18
10.4.	Analytical Specificity (Exclusivity)	19
10.4.1.	In-Slico Analytical Specificity	19
10.4.2.	Wet-Test Analytical Specificity	20
10.5.	Interferences	22
11.	TROUBLESHOOTING	23
12.	EXPLANATION of SYMBOLS	24

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 ABR-2 Panel PCR Kit** is a multiplexed qualitative Real-Time Polymerase Chain Reaction (qPCR) test intended for the simultaneous detection and identification of multiple antimicrobial resistance genes in research samples. The **MarinaBiolab ABR-2 Panel PCR Kit** allows to achieve qPCR result in less than 1 hour. The test is performed to detect gene sequences of the following antimicrobial resistance genes.

Targets	
AmpC-type beta-lactamase (class C) resistance gene (ampC)	Cefoxitin-hydrolyzing class C-beta-lactamase resistance gene (FOX)
Ambler class C beta-lactamase resistance gene (ACC)	Vietnam extended-spectrum beta-lactamase resistance gene (VEB)
Sulphydryl variable-beta-lactamase resistance gene (SHV)	Apramycin resistance gene (AAC(3)-IV)
Erythromycin resistance gene (ermA)	Guiana extended-spectrum beta-lactamase resistance gene (GES)
Erythromycin resistance gene (ermB)	Tetracycline efflux pump resistance genes (TET)
Erythromycin resistance gene (ermC)	Aminoglycoside modifying resistance gene (AAC(6')-Ie-APH(2'')-Ia)
DHA beta-lactamase resistance gene (DHA)	Aminoglycoside nucleotidyltransferase resistance gene (aadA)
Aminoglycoside-3'-phosphotransferase-Ia resistance gene (APH(3')-Ia)	2''-aminoglycoside nucleotidyltransferase resistance gene (ANT(2))
Controls	
Human RNase P (IC)	
<i>Bacillus atrophaeus</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 ABR-2 Panel PCR Kit** consists of four main components:

1. qPCR Enzyme and Buffer Mix (qPCR Master Mix)
2. Forward, Reverse and Probe Oligo Mix (ABR-2 Oligo Mix 1-5)
3. Mix of non-infectious DNA from artificial sample including targets in the table below (PC-ABR-2)
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 MBLABR2017
qPCR Master Mix	Ready-to-use mix for qPCR	2 x 1250 µL
ABR-2 Oligo Mix 1-5	Primers and probes complementary to specific regions of the targets in the table above	5 x 250 µL
PC-ABR-2	Mix of non-infectious DNA from artificial sample including targets in the table above	1 x 400 µL
NTC	DNase/RNase-Free Water	1 x 400 µL

Table 2. Oligo Mix target antimicrobial resistance genes and detection channels.

Vial Name	Target	Channel
ABR-2 Oligo Mix 1	ampC	FAM/Green
	ACC	HEX/VIC/JOE/Yellow
	SHV	ROX/Texas Red/Orange
	Human RNase P (IC)	CY5/Red
ABR-2 Oligo Mix 2	ermA	FAM/Green
	-	HEX/VIC/JOE/Yellow
	DHA	ROX/Texas Red/Orange
	<i>Bacillus atrophaeus</i> (EC)	CY5/Red
ABR-2 Oligo Mix 3	ermB	FAM/Green
	APH(3')-Ia	HEX/VIC/JOE/Yellow
	FOX	ROX/Texas Red/Orange
	VEB	CY5/Red

ABR-2 Oligo Mix 4	ermC	FAM/Green
	AAC(3)-IV	HEX/VIC/JOE/Yellow
	GES	ROX/Texas Red/Orange
	TET	CY5/Red
ABR-2 Oligo Mix 5	AAC(6')-Ie-APH(2'')-Ia	FAM/Green
	aadA	HEX/VIC/JOE/Yellow
	-	ROX/Texas Red/Orange
	ANT(2)	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 ABR-2 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 ABR-2 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
ABR-2 Oligo Mix 1-5	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 ABR-2 Oligo Mix 1-5 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 ABR-2 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 antimicrobial resistance genes.

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 ABR-2 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 ABR-2 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
ampC	ATCC BAA-3286	1.2E+02 copies/mL	20/20 100%
ACC	In-house	8.3E+01 copies/mL	20/20 100%
SHV	ATCC BAA-3261	1.1E+02 copies/mL	20/20 100%
ermA	ATCC BAA-1415	1.2E+02 copies/mL	20/20 100%
DHA	In-house	2.2E+02 copies/mL	20/20 100%
ermB	ATCC BAA-1415	1.8E+02 copies/mL	20/20 100%
APH(3')-Ia	ATCC BAA-3263	1.7E+02 copies/mL	20/20 100%
FOX	In-house	9.8E+01 copies/mL	20/20 100%
VEB	In-house	9.2E+01 copies/mL	20/20 100%
ermC	ATCC BAA-1415	1.0E+02 copies/mL	20/20 100%
AAC(3)-IV	ATCC BAA-3266	1.1E+02 copies/mL	20/20 100%
GES	ATCC BAA-3284	1.5E+02 copies/mL	20/20 100%
TET	ATCC BAA-3288	1.4E+02 copies/mL	20/20 100%

AAC(6')-Ile-APH(2'')-Ia	ATCC BAA-3288	1.2E+02 copies/mL	20/20 100%
aadA	ATCC BAA-3263	8.8E+01 copies/mL	20/20 100%
ANT(2)	ATCC BAA-3261	8.7E+01 copies/mL	20/20 100%

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 ampC, ACC, SHV, ermA, DHA, ermB, APH(3')-Ia, FOX, VEB, ermC, AAC(3)-IV, GES, TET, AAC(6')-Ile-APH(2'')-Ia, aadA, and ANT(2) 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
ampC	Sense Primer	5.911	100.00%	0.00%	0.00%	0.00%
ampC	Antisense Primer	5.911	99.13%	0.87	0.00%	0.00%
ampC	Hydrolysis Probe	5.952	99.78%	0.22%	0.00%	0.00%
ACC	Sense Primer	65	100.00%	0.00%	0.00%	0.00%
ACC	Antisense Primer	65	100.00%	0.00%	0.00%	0.00%
ACC	Hydrolysis Probe	71	100.00%	0.00%	0.00%	0.00%
SHV	Sense Primer	5.463	100.00%	0.00%	0.00%	0.00%
SHV	Antisense Primer	5.463	100.00%	0.00%	0.00%	0.00%
SHV	Hydrolysis Probe	5.470	100.00%	0.00%	0.00%	0.00%

ermA	Sense Primer	1.982	100.00%	0.00%	0.00%	0.00%
ermA	Antisense Primer	1.982	100.00%	0.00%	0.00%	0.00%
ermA	Hydrolysis Probe	1.987	100.00%	0.00%	0.00%	0.00%
DHA	Sense Primer	798	100.00%	0.00%	0.00%	0.00%
DHA	Antisense Primer	798	99.82%	0.18%	0.00%	0.00%
DHA	Hydrolysis Probe	792	100.00%	0.00%	0.00%	0.00%
ermB	Sense Primer	2.899	100.00%	0.00%	0.00%	0.00%
ermB	Antisense Primer	2.899	99.14%	0.86%	0.00%	0.00%
ermB	Hydrolysis Probe	2.890	100.00%	0.00%	0.00%	0.00%
APH(3')-Ia	Sense Primer	6.413	97.46%	2.54%	0.00%	0.00%
APH(3')-Ia	Antisense Primer	6.413	98.11%	1.89%	0.00%	0.00%
APH(3')-Ia	Hydrolysis Probe	6.420	98.20%	1.80%	0.00%	0.00%
FOX	Sense Primer	63	100.00%	0.00%	0.00%	0.00%
FOX	Antisense Primer	63	100.00%	0.00%	0.00%	0.00%
FOX	Hydrolysis Probe	65	100.00%	0.00%	0.00%	0.00%
VEB	Sense Primer	209	100.00%	0.00%	0.00%	0.00%
VEB	Antisense Primer	209	100.00%	0.00%	0.00%	0.00%
VEB	Hydrolysis Probe	215	100.00%	0.00%	0.00%	0.00%
ermC	Sense Primer	462	100.00%	0.00%	0.00%	0.00%
ermC	Antisense Primer	462	98.50%	1.50%	0.00%	0.00%
ermC	Hydrolysis Probe	465	99.00%	1.00%	0.00%	0.00%
AAC(3)-IV	Sense Primer	1.034	100.00%	0.00%	0.00%	0.00%
AAC(3)-IV	Antisense Primer	1.034	98.56%	1.44%	0.00%	0.00%
AAC(3)-IV	Hydrolysis Probe	1.030	100.00%	0.00%	0.00%	0.00%
GES	Sense Primer	408	100.00%	0.00%	0.00%	0.00%
GES	Antisense Primer	408	100.00%	0.00%	0.00%	0.00%
GES	Hydrolysis Probe	405	100.00%	0.00%	0.00%	0.00%
TET	Sense Primer	49.900	100.00%	0.00%	0.00%	0.00%
TET	Antisense Primer	49.900	97.45%	2.55%	0.00%	0.00%
TET	Hydrolysis Probe	49.915	99.26%	0.74%	0.00%	0.00%
AAC(6')-Ie-APH(2'')-Ia	Sense Primer	1.411	100.00%	0.00%	0.00%	0.00%
AAC(6')-Ie-APH(2'')-Ia	Antisense Primer	1.411	99.69%	0.31%	0.00%	0.00%

AAC(6')-Ie-APH(2'')-Ia	Hydrolysis Probe	1.410	100.00%	0.00%	0.00%	0.00%
aadA	Sense Primer	23.486	99.29%	0.71%	0.00%	0.00%
aadA	Antisense Primer	23.486	99.13%	0.87%	0.00%	0.00%
aadA	Hydrolysis Probe	23.480	99.74%	0.26%	0.00%	0.00%
ANT(2)	Sense Primer	1.090	100.00%	0.00%	0.00%	0.00%
ANT(2)	Antisense Primer	1.090	100.00%	0.00%	0.00%	0.00%
ANT(2)	Hydrolysis Probe	1.091	100.00%	0.00%	0.00%	0.00%

10.3.2. Wet-Test Analytical Reactivity

The analytical reactivity (inclusivity) of the **MarinaBiolab ABR-2 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 ABR-2 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
ampC	ATCC BAA-3286	1x
SHV	ATCC BAA-3261	1x
ermA	ATCC BAA-1415	1x
ermB	ATCC BAA-1415	1x
APH(3')-Ia	ATCC BAA-3263	1x
ermC	ATCC BAA-1415	1x
AAC(3)-IV	ATCC BAA-3266	1x
GES	ATCC BAA-3284	1x
TET	ATCC BAA-3288	1x
AAC(6')-Ie-APH(2'')-Ia	ATCC BAA-3288	1x
aadA	ATCC BAA-3263	1x
ANT(2)	ATCC BAA-3261	1x

10.4. Analytical Specificity (Exclusivity)

10.4.1. In-Slico 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	ampC	None	None	None
On-Panel	ACC	None	None	None
On-Panel	SHV	None	None	None
On-Panel	ermA	None	None	None
On-Panel	DHA	None	None	None
On-Panel	ermB	None	None	None
On-Panel	APH(3')-Ia	None	None	None
On-Panel	FOX	None	None	None
On-Panel	VEB	None	None	None
On-Panel	ermC	None	None	None
On-Panel	AAC(3)-IV	None	None	None
On-Panel	GES	None	None	None
On-Panel	TET	None	None	None
On-Panel	AAC(6')-Ie-APH(2'')-Ia	None	None	None
On-Panel	aadA	None	None	None
On-Panel	ANT(2)	None	None	None
Off-Panel	KPC	None	None	None
Off-Panel	VIM	None	None	None
Off-Panel	NDM	None	None	None
Off-Panel	IMP	None	None	None
Off-Panel	QNR	None	None	None
Off-Panel	OXA	None	None	None
Off-Panel	mecA	None	None	None

Off-Panel	mecC	None	None	None
Off-Panel	Van A	None	None	None
Off-Panel	Van B	None	None	None
Off-Panel	Sul1	None	None	None
Off-Panel	Sul2	None	None	None
Off-Panel	CTX	None	None	None
Off-Panel	DfrA	None	None	None
Off-Panel	mcr-3	None	None	None
Off-Panel	vanC	None	None	None
Off-Panel	vanD	None	None	None
Off-Panel	SME	None	None	None
Off-Panel	mcr-4	None	None	None
Off-Panel	blaRAHN	None	None	None
Off-Panel	CMY	None	None	None
Off-Panel	ompK36	None	None	None
Off-Panel	SPM	None	None	None
Off-Panel	vanM	None	None	None
Off-Panel	mcr-2	None	None	None
Off-Panel	OXA-24/65	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 ABR-2 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 ABR-2 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 ABR-2 Panel PCR Kit** analytical specificity.

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

ampC	ATCC BAA-3286	None
SHV	ATCC BAA-3261	None
ermA	ATCC BAA-1415	None
ermB	ATCC BAA-1415	None
APH(3')-Ia	ATCC BAA-3263	None
ermC	ATCC BAA-1415	None
AAC(3)-IV	ATCC BAA-3266	None
GES	ATCC BAA-3284	None
TET	ATCC BAA-3288	None
AAC(6')-Ie-APH(2'')-Ia	ATCC BAA-3288	None
aadA	ATCC BAA-3263	None
ANT(2)	ATCC BAA-3261	None

Table 14. Off-Panel organisms were tested for evaluation of *MarinaBiolab ABR-2 Panel PCR Kit* analytical specificity.

Organism	Isolate ID/Source	Cross Reactivity Detected
KPC	Zeptomatrix NATPPQ-BIO	None
VIM	Zeptomatrix NATPPQ-BIO	None
NDM	Zeptomatrix NATPPQ-BIO	None
IMP	Zeptomatrix NATPPQ-BIO	None
QNR	ATCC BAA-2728	None
OXA	Zeptomatrix NATPPQ-BIO	None
mecA	ATCC BAA-2094	None
mecC	ATCC BAA-2313	None
VanA	Zeptomatrix 0801892	None
VanB	Zeptomatrix 0801953	None
Sul1	ATCC BAA-3035	None
Sul2	ATCC BAA-2894	None
CTX	Zeptomatrix NATPPQ-BIO	None
DfrA	ATCC BAA-3041	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 ABR-2 Panel PCR Kit** and extrapolated from the interference evaluation of the **MarinaBiolab ABR-2 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 ABR-2 Panel PCR Kit**.

Table 15. Evaluation of potentially interfering substances on the **MarinaBiolab ABR-2 Panel PCR Kit**.

Substance Tested	Concentration Tested	Observed Interference
Endogenous Substances		
Human Genomic DNA	0.07 mg/mL	No Interference
Hemoglobin	10 mg/mL	No Interference
D-Glucose	10 mg/mL	No Interference
Cholesterol	4.0 mg/mL	No Interference
Exogenous Substances		
Acetaminophen	0.2 mg/mL	No Interference
Salicylic Acid	30 µg/mL	No Interference
Ibuprofen	0.2 mg/mL	No Interference
Specimen Collection Materials		
K ₂ EDTA	N/A	No Interference
Sodium Citrate	N/A	No Interference
Sodium Heparin	N/A	No Interference
Lithium Heparin	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.