SHENTEK

Mycobacteria DNA Detection Kit

User Guide

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Version: A/1 For Research Use Only Product No.:1503602 Reagents for 50 Reactions

Huzhou Shenke Biotechnology Co., Ltd

(IMPORTANT: Please read this document carefully before experiment.)

1. Product information

Product description

MycoSHENTEK[®] Mycobacteria DNA Detection Kit is used, together with MycoSHENTEK[®] Mycobacteria DNA Extraction Kit, to qualitatively determine whether there is Mycobacteria contamination in cells, cell culture-derived biologicals, etc., using real-time qPCR technology.

The integrated system has been fully validated at a detection limit of 10-100 CFU/mL to detect more than 100 different Mycobacteria species. The system contains dUTP, and if used with the UNG enzyme to prevent contamination and

eliminate carry-over contamination.

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Extraction Kit User Guide (Product No.: 1503601)

Kit contents and storage

WARNING: Please read the Material Safety Data Sheets (MSDSs) and follow the handling instructions. Wear appropriate protective eyewear, clothing and gloves.

	-	-		
Reagent	Part No.	Quantity	Storage	
MB qPCR Reaction Buffer	NNB011	400 μ L × 1 tube	-20°C,	
MB Primer & Probe MIX	NNC035	75 μ L × 1 tube	protect from light	
MB Internal Control (IC)	NNA037	600 μ L × 1 tube		
MB Positive Control (PC)*	NNA041	500 μ L × 1 tube	-20°C	
DNA Dilution Buffer(DDB)	NND001	$1.5 \text{ mL} \times 1 \text{ tube}$		

Table 1. Kit components and storage

The kit components can be stored at appropriate conditions for up to 24 months. Please check the expiration date on the labels.

Applied instruments, including but not limited to the following

- SHENTEK-96S Real-time PCR System
- CFX96 Real-Time PCR System
- > 7500 Real-Time PCR System

Required materials not included in the kit

- Nonstick, RNase-free microfuge tubes, 1.5mL, 2.0mL
- > PCR 8-well strip tubes with caps or 96-well plates with seals
- Low retention filter tips 1000µL, 100µL, 10µL
- 75% Ethanol

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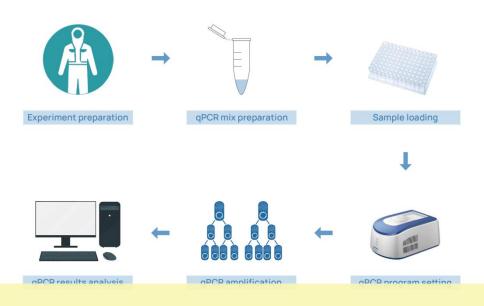
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Microplate and micro test tube shaker

- > Vortex mixer
- Real-time PCR System
- ➢ Pipettes, 1000µL, 100µL, 10µL

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Workflow



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with 75% ethanol

3. Thaw the kit completely at 2-8°C or melt on ice, vortex and spin briefly.

qPCR MIX preparation

1. Determine the number of reaction wells according to the number of test samples and control samples with 2 replicates for each sample generally.

Number of reaction wells = (1 Positive control (PC) + 1 No Template control)

(NTC) + 1 Negative control sample (NCS) + N Unknown samples)×2

2. Thaw the kit completely at 2-8°C or melt on ice, then prepare qPCR MIX according to the following table.

Reagents	Volume/reaction	Volume for 30 reaction (includes 10% overage)		
MB qPCR Reaction Buffer	8 μL	264 µL		
MB Primer&Probe MIX	1.5 μL	49.5 μL		
IC*	0.5 μL	16.5 μL		
Total volume	10 µL	330 µL		
UNG enzyme (optional)	0.1 U	3.3 U		

* If IC has been added during sample extraction, an equal volume of DNA dilution buffer should be added instead of IC to prepare qPCR MIX.

qPCR Reaction MIX preparation

. Vortex each solution and mix well. Then add the solution according to Table 3

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able 3. CR Reaction MIX Prepar

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(Nagetive control sample) 10μ L qPCR MIX + 20 μ L of extracted NCS

 $\frac{10\mu L \text{ qPCR MIX} + 20\mu L \text{ of extracted unknown}}{\text{sample}}$

	_		Tau	ne 4.E	kample	01 90-	wen p	Tale lay	youi		-	
PC	PC				S1	S1						А
					S2	S2						В
					S3	S3						С
					S4	S4						D
					S5	S5						Е
					S6	S6						F
					S7	S7						G
NTC	NTC				S8	S8				NCS	NCS	Н
1	2	3	4	5	6	7	8	9	10	11	12	

Table 4.Example of 96-well plate layout

• This example shows one Positive control (PC), one No-template control (NTC),

one Negative control sample (NCS), and eight unknown samples, with duplicate wells for each sample in qPCR analysis.

• *Adjust the layout according to the number of test samples to be run.*

2. Close PCR 8-well strip tubes with caps, or seal the 96-well plate with sealing film. Mix well in microplate or micro test tube shaker, then spin down the reagents for 10 seconds in centrifuge and place it in the qPCR instrument.

■ qPCR program setting

Please refer to the program setting as follows:

- 1. Run a new method program and select the quantitative PCR assay template.
- 2. Run a new Probe template, and type the name "Mycobacteria". Select FAM in the Reporter Dye drop-down list and select (none) in the Quencher Dye drop-down list. Select VIC in the Reporter Dye drop-down list and select (none).

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erence fluorescence as ROX (optional

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Step	Temp.	Time(mm:sec)	Cycles
UNG treatment	25°C	10:00	1
Activation	95°C	10:00	1
Denaturation	95°C	00:15	
Annealing	60°C	00:30	45
Extension	72°C*	01:30	

able 5. qPCR running temperature and time

*Instrument will read the fluorescence signal during this step.

Result analysis

The following instructions apply only to the 7500 instrument with SDS v1.4.

1. Select the Results tab, then click Amplification Plot. In the Analysis Settings

window, enter the following settings:

- a. Select Manual Ct.
- b. In the **Threshold** field, enter 0.02.
- c. Click Analyze, and check whether it is a normal amplification curve.
- 2. Select the **Results** tab, and click **Plate** tab, then set tasks for each sample type by clicking on the Task Column drop-down list and then click **>**:
 - a. NTC wells: target DNA detector task = NTC.
 - b. NCS, unknown samples wells: target DNA detector task = Unknown.
- 3. Select File >> Export >> Results. In the Save as type drop-down list, select

Results Export Files, then click Save.

NOTE: If you use a different instrument or software, refer to the applicable

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Quality control		
samples	FAM	VIC
NTC	Ct≥40.00 or absence of specific	Ct<35.00 and normal
NIC	amplification for duplicate runs	amplification for duplicate runs
NCC	Ct≥40.00 or absence of specific	Ct<35.00 and normal
NCS	amplification for duplicate runs	amplification for duplicate runs
DC	Ct<35.00 and normal	Ct<35.00 and normal
PC	amplification for duplicate runs	amplification for duplicate runs

The QC sample analysis shall be considered in relationship to LOD (Limit of Detection) based on method validation data.

2. Guidance for unknown samples as in the following table.

FAM	VIC	Conclusion
Ct<40.00 (at least	Ct<40.00 and normal amplification for duplicate runs	Positive
one well) and normal amplification	Ct≥40.00 or absence of specific amplification for duplicate runs	Presumptive Positive and presence of PCR inhibitors
Ct≥40.00 or	Ct<40.00 and normal amplification for duplicate runs	Negative
non-specific amplification	Ct≥40.00 or absence of specific amplification for duplicate runs	Not conclusive, and presence of PCR inhibitors

Table 7.Test sample results analysis

* If the VIC signal is inhibited, it is necessary to repeat the sample preparation or

appropriately remove the inhibitors during the sample preparation and repeat the

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