

ASSAY NAME: UTI18_QS

Quantity: 100 x 20µL PCR reactions

6-plex assay: *Enterobacter cloacae*, *Acinetobacter baumannii*, *Klebsiella aerogenes*, *Citrobacter freundii*, *Citrobacter koseri*, and human RPP30 DNA

SKU #: PNP-UTI18-D-QS-100 (QuantStudio)

(RUO). Research Use Only. Not for use in Diagnostic Procedures.

SCOPE OF THIS PRODUCT INFORMATION SHEET:

The oligonucleotide recipes are optimized for each instrument (QuantStudio, BioRad). The verification data presented in this IFU were performed using PNP-UTI18-D-QS-100 on a QuantStudio™ 7 Flex Real-Time System. The performance of the other SKUs on their corresponding instrument should be similar. Contact PCRassays.com if you are planning to use a different instrument.

CONTENTS

The primers and probes in the UTI18_QS assay are provided in Tube 1 as a 5X concentrated working solution that detects 4 pathogens and a human control.

Table of Dyes used in this assay:

| Pathogen/Target | Dyes | Quencher | Refs. |
|---------------------------------------|---------------|----------|-------|
| <i>E. cloacae</i> | FAM | BHQ-1 | 1,2 |
| <i>A. baumannii</i> | HEX | BHQ-1 | 3 |
| RPP30-DNA control | TAMRA | BHQ-2 | 4, 5 |
| <i>K. aerogenes</i> | TEX615 | BHQ-2 | 6 |
| <i>C. freundii</i> / <i>C. koseri</i> | Cy5 | BHQ-2 | 7 |

The probes are designed as TaqMan⁸ cleavage mechanism and thus the reaction requires a DNA polymerase with 5'-exonuclease activity.

ASSAY HANDLING

The UTI18_QS assay is shipped at ambient temperature, and should be stored at -20 °C. The assay should be kept on ice once thawed. Do not subject the enzyme to multiple freeze-thaw cycles.

Any contamination should be avoided by using appropriate personal protective equipment (PPE), powder free gloves, aerosol barrier pipette tips, and a clean hood.

ASSAY CONTENTS:

Tube 1: 5X Primer/Probe mix for *E. cloacae*, *A. baumannii*, *K. aerogenes*, *C. freundii*, *C. koseri*, and hRPP30DNA.

Tube 2: (Do NOT add to specimen unknowns) Positive control: 5000 copies/µl of synthetic 500 bp DNA fragments for *E. cloacae*, *A. baumannii*, *K. aerogenes*, *C. freundii*, *C. koseri*, and hRPP30DNA.

Tube 3: InhibiTaq qPCR enzyme Mastermix (enough for 100 rxns. with 20 µL total volume). This is a custom formulation from Fortis Life Sciences to the specifications of PCRassays.com.



EXPERIMENTAL

Perform nucleic acid extraction/purification (recommended). Since some of the bacteria in this assay are Gram+, it is important to use an extraction procedure with an appropriate cell wall lysis agent.

Note: molecular biology grade water (NOT included) should be used to prepare the PCR reactions.

Set up your PCR reaction (20 µL) as follows on ice:

| Component | Volume (µL) |
|---------------------------------|-------------|
| InhibiTaq enzyme mastermix (2X) | 10 |
| Primer/Probe mix (5X) | 4 |
| Sample | 2 |
| Water | 4 |

Notes: To improve assay sensitivity, up to 6 µL of sample can be added (water volume adjusted accordingly) for a total reaction volume of 20 µL. For positive control rxns., add 2 µL of the solution from Tube 2 (i.e., the "sample").

A PCR protocol was used for verification on a QuantStudio™ 7 Flex Real-Time System, with the following program:

| Step | Thermocycling Protocol: |
|------|---------------------------------|
| 1 | Incubate @ 95 °C for 2 minutes |
| 2 | Incubate @ 95 °C for 3 seconds |
| 3 | Incubate @ 55 °C for 22 seconds |
| 4 | Plate Read |
| 5 | Go to Step 2, repeat 44× more |

RESULT INTERPRETATION

After running the qPCR reaction, use the instrument software to determine the quantification cycle, Cq (or use CT if your instrument does not have the capability to compute a Cq). Fluorescence channels with a Cq < 38 cycles, and final RFU >Threshold is considered "positive" or "+" in the Table below. The "Threshold" value for calling a PCR positive is dependent on the instrument model, well size, and sample volume; thus the user must determine the threshold that is appropriate for their method. For our QuantStudio 7 Flex with 96 well plate with 100 µL wells and 20 µL reaction volume, the average RFU was approximately 1,000,000 and we used a threshold of 200,000 for calling positives or "+" in the Table below.

| <i>E. cloacae</i> FAM™ | <i>A. baumannii</i> HEX™ | RPP30 TAMRA [†] M | <i>K. aerogenes</i> TEX615™ | <i>C. freundii</i> / <i>C. koseri</i> Cy5™ | Recommended Interpretation |
|---------------------------|-----------------------------|----------------------------------|--------------------------------|--------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| - | - | - | - | - | The PCR reaction failed. Please repeat the experiment. |
| - | - | + | - | - | The sample contains human RPP30 DNA. The sample doesn't contain bacterial DNA. |
| + | - | - | - | - | The sample contains <i>E. cloacae</i> DNA. The sample may not contain human RPP30 DNA. |
| + | - | + | - | - | The sample contains <i>E. cloacae</i> DNA and human RPP30 DNA. |
| - | + | - | - | - | The sample contains <i>A. baumannii</i> DNA. The sample may not contain human RPP30 DNA. |
| - | + | + | - | - | The sample contains <i>A. baumannii</i> DNA and human RPP30 DNA. |
| - | - | - | + | - | The sample contains <i>K. aerogenes</i> DNA. The sample may not contain human RPP30 DNA. |
| - | - | + | + | - | The sample contains <i>K. aerogenes</i> DNA and human RPP30 DNA. |
| - | - | - | - | + | The sample contains <i>C. freundii</i> / <i>C. koseri</i> DNA. The sample may not contain human RPP30 DNA. |
| - | - | + | - | + | The sample contains <i>C. freundii</i> / <i>C. koseri</i> DNA and human RPP30 DNA. |
| + | + | - | + | + | The sample contains <i>E. cloacae</i> , <i>A. baumannii</i> , <i>K. aerogenes</i> , and <i>C. freundii</i> / <i>C. koseri</i> DNA. The sample may not contain human RPP30 DNA. |
| + | + | + | + | + | The sample contains <i>E. cloacae</i> , <i>A. baumannii</i> , <i>K. aerogenes</i> , <i>C. freundii</i> / <i>C. koseri</i> DNA, and human RPP30 DNA. |

VERIFICATION EXPERIMENTS

The UTI18_QS assay verification was carried out as a 6-plex assay, which simultaneously detects DNA from *E. cloacae*, *A. baumannii*, *K. aerogenes*, *C. freundii*, *C. koseri*, and human RPP30 DNA, which serves as a positive extraction-control assay.

Experiments were performed in triplicate using the experimental procedure given above, but with different samples added to each reaction. The samples used for the verification experiments contained 1'10⁴ copies/reaction of 500 bp synthetic DNA constructs (from Twist Biosciences) harboring the regions of interest from the pathogen genomes, human RPP30 DNA gene, and human genomic DNA. DNA extracts were provided by ATCC (*E. cloacae* #13047DQ, *A. baumannii* #17978DQ, *K. aerogenes* #13048DQ, *C. freundii* #8090 bacteria, *C. koseri* #27027 bacteria). The results of these experiments are shown in **Figure 1** and indicate that the 6-plex specifically detects the different bacterial species in the human genomic DNA matrix.

The limit of detection (LOD) was estimated by performing serial dilution experiments in triplicate (Figure 2). For dilution series only target construct was added. The results show a limit of detection (LOD)

Conclusion: The data in Figure 1 indicate that the 6-plex primers and probes specifically detect and differentiate the bacterial types and are also compatible with RPP30_DNA positive control primers.

NOTES

- ¹ FAM™ (Carboxyfluorescein) is a trademark of Life Technologies, Inc
- ² BHQ-1™ (Black Hole Quencher) is a trademark of Biosearch Technologies, Inc.
- ³ HEX™ (Hexachloro-fluorescein) is a trademark of Applera Corp
- ⁴ TAMRA (Carboxyltetramethylrhodamine) is a trademark of Applera Cor.
- ⁵ BHQ-2™ (Black Hole Quencher) is a trademark of Biosearch Technologies, Inc.
- ⁶ TEX615™ is a trademark of Thermo Fisher Scientific
- ⁷ Cy5™ is a trademark of GE Healthcare
- ⁸ "TaqMan" is a trademark of Roche Molecular Systems, Inc.

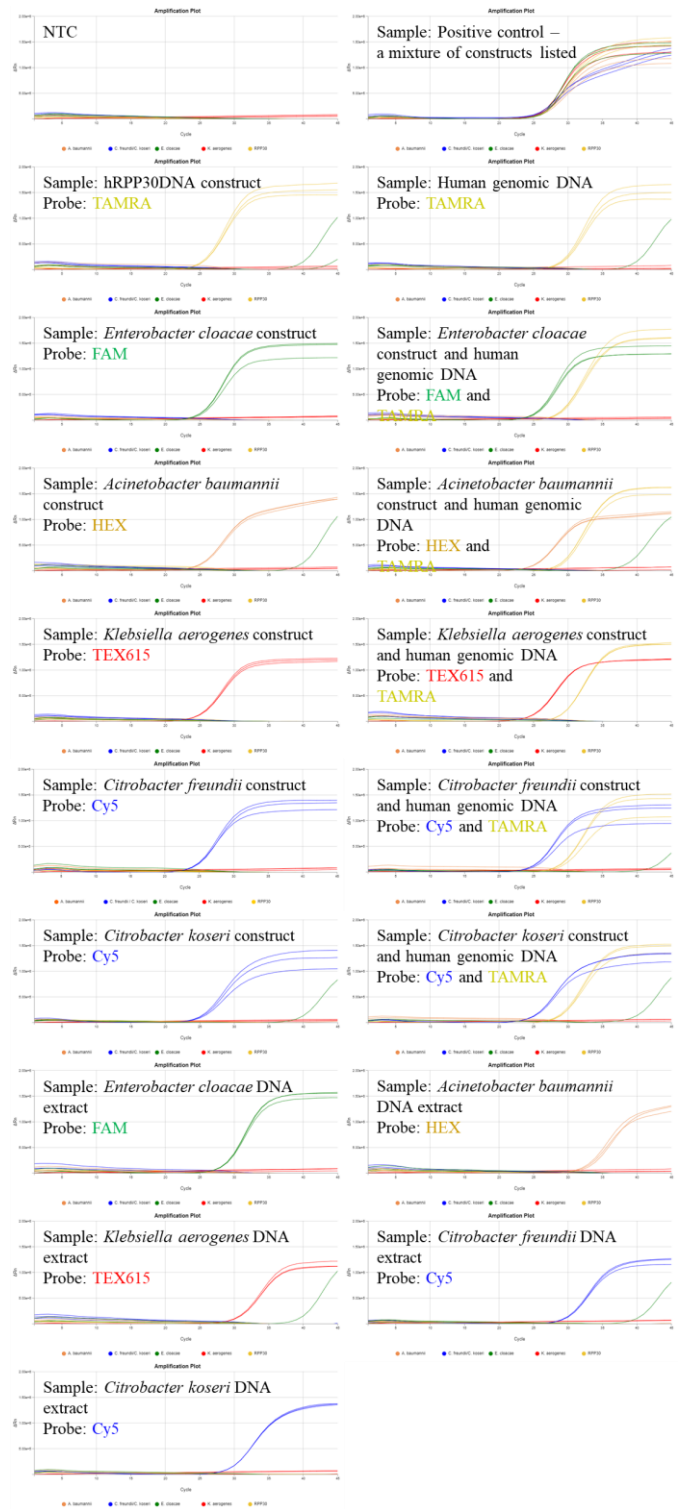


Figure 1: Verification experiments with single targets (given in text boxes for each panel). All sets of probes and primers are present in every reaction, but positive signal is only observed when the target(s) is present, indicating that the amplification is specific. Signal from single molecule events show Cq >38 and are considered negative.

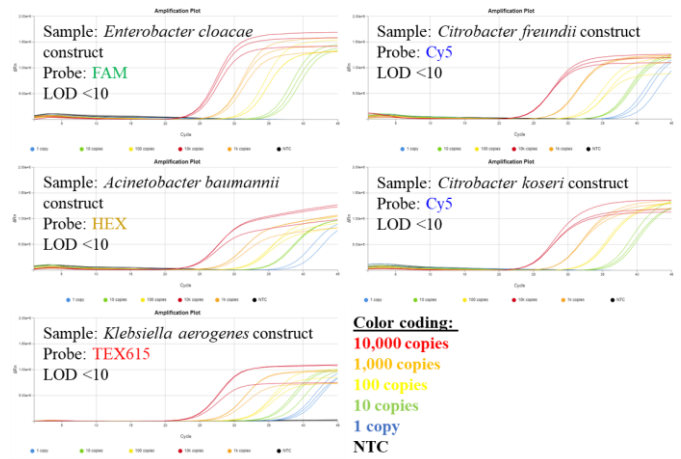


Figure 2: Serial dilution experiments show LOD <math><10</math> molecules for the synthetic DNA construct of each target.

CONTACT US

For assistance, please contact DNA Software using the link:
<https://www.pcrassays.com/contact/>

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