

ASSAY NAME: FUN3A-BR
(Fungal Panel 3A for Bio-Rad 384 or 96)

Quantity: 100 x 20µL PCR reactions

4-plex assay: *Malassezia* genus, *Nannizzia* genus, *Trichosporon* genus, and human RPP30 DNA

SKU: BUN-FUN3A-D-BR-100 (Bio-Rad)

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

SCOPE OF THIS PRODUCT INFORMATION SHEET (PIS):

The oligonucleotide recipes are optimized for each instrument (Bio-Rad, QuantStudio). The verification data presented in this PIS were performed using kit BUN-FUN3A-D-BR-100 on a Bio-Rad CFX96 instrument using a white-bottomed plate. This kit is compatible with both CFX-96 and CFX-384 instruments. The performance of the other SKUs on their corresponding instrument should be similar. Contact PCRassays.com if you are planning to use a different qPCR instrument.

CONTENTS

The primers and probes in the FUN3A assay are provided in Tube 1 as a 5X concentrated working solution that detects the 3 pathogens and a human extraction control. The species detected by the *Malassezia* genus primers include: *M. furfur*, *M. globosa*, *M. pachydermatis*, *M. restricta*, and *M. sympodialis*. The species detected by the *Nannizzia* genus primers include: *N. fulva*, *N. incurvata*, *N. gypsea*, and *N. nana*. The species detected by the *Trichosporon* genus primers include: *Cutaneotrichosporon dermatitis*, *Cutaneotrichosporon oleaginosum*, *T. asahii*, *T. coremiiforme*, *T. faecale*, *T. inkin*, and *T. ovoides*.

Table of Dyes used in this assay:

Pathogen/Target	Dyes	Quencher	Refs.
<i>Malassezia</i> genus	FAM	BHQ-1	1, 2
RPP30-DNA control	HEX	BHQ-1	3
<i>Nannizzia</i> genus	TEX615	BHQ-2	4, 5
<i>Trichosporon</i> genus	Cy5	BHQ-2	6

The probes are designed as TaqMan⁷ cleavage mechanism and thus the reaction requires a DNA polymerase with 5'-exonuclease activity.

ASSAY HANDLING AND CONTAMINATION

The FUN3A kit is shipped on ice, and should be stored at -20 °C. The tubes should be kept on ice once thawed. Do not subject the enzyme to multiple freeze-thaw cycles.

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 *Malassezia* genus, *Nannizzia* genus, *Trichosporon* genus, and hRPP30DNA.

Tube 2: (Do NOT add to specimen unknowns) Positive control: Synthetic 500 bp DNA fragments of *Malassezia* genus, *Nannizzia* genus, *Trichosporon* genus, and human RPP30DNA.

Tube 3: InhibiTaq Standard 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).

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

Component	Volume (µL)
InhibiTaq qPCR enzyme mastermix (2X)	10
Primer/Probe mix (5X)	4
Sample or Positive Control	2
Water (Molecular Biology Grade)	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.

For 10 µL reactions, divide all of the amounts above by a factor of 2.

A PCR protocol was used for verification on a Bio-Rad CFX96 system using white plates, 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 20 seconds*
4	Plate Read
5	Go to Step 2, repeat 44x more

* A 15 second extension time may be used for Bio-Rad CFX96 instruments.

RESULT INTERPRETATION

After running the qPCR reaction, use the instrument software to determine the quantification cycle, C_q (or use C_T if your instrument does not have the capability to compute a C_q). PCR reactions are considered “positive” or “+” in the table below if the RFU > “threshold” and the C_q <38 cycles. “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 Bio-Rad CFX-96 with 200 µL wells (with white-bottomed plate) and 20 µL reaction volume, the average RFU was approximately 3,300, we used a threshold of 330 for calling positives or “+” in the Table below.

<i>Malassezia</i> genus FAM TM	<i>Nannizzia</i> genus TEX615 TM	<i>Trichosporon</i> genus Cy5 TM	RPP30 HEX TM	Recommended Interpretation
—	—	—	—	The PCR reaction failed. Please repeat the experiment.
—	—	—	+	The sample contains human RPP30 DNA. The sample doesn't contain pathogen DNA.
+	—	—	—	The sample contains <i>Malassezia</i> genus DNA. The sample may not contain human RPP30 DNA.
+	—	—	+	The sample contains <i>Malassezia</i> genus DNA and human RPP30 DNA.
—	+	—	—	The sample contains <i>Nannizzia</i> genus DNA. The sample may not contain human RPP30 DNA.
—	+	—	+	The sample contains <i>Nannizzia</i> genus DNA and human RPP30 DNA.
—	—	+	—	The sample contains <i>Trichosporon</i> genus DNA. The sample may not contain human RPP30 DNA.
—	—	+	+	The sample contains <i>Trichosporon</i> genus DNA and human RPP30 DNA.
+	+	+	—	The sample contains <i>Malassezia</i> genus, <i>Nannizzia</i> genus, and <i>Trichosporon</i> genus DNA. The sample may not contain human RPP30 DNA.
+	+	+	+	The sample contains <i>Malassezia</i> genus, <i>Nannizzia</i> genus, <i>Trichosporon</i> genus DNA, and human RPP30 DNA.

VERIFICATION EXPERIMENTS

The FUN3A assay verification was carried out as a 4-plex assay, which simultaneously detects DNA from *Malassezia* genus, *Nannizzia* genus, *Trichosporon* genus, 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 1×10^4 copies/reaction of synthetic 500 bp DNA constructs (from Twist Biosciences) harboring the regions of interest from the pathogen genomes, human RPP30 DNA gene, and human genomic DNA. **Figure 1** shows the results of these experiments, which indicate that the 4-plex specifically detects the different pathogens in the human genomic DNA matrix.

The limit of detection (LOD) was estimated by performing serial dilution experiments in triplicate (**Figure 2**). For each dilution series only one target construct was added. The results show a limit of detection (LOD) <10 copies/reaction.

Conclusion: The data in **Figure 1** indicate that the 4-plex primers and probes specifically detect and differentiate the pathogens and are also compatible with RPP30 DNA positive control primers. Human genomic DNA doesn't interfere with the detection of the pathogens.

NOTES

¹ FAMTM (Carboxyfluorescein), a trademark of Life Technologies Corp.

² BHQ-1TM (Black Hole Quencher) is a trademark of Biosearch Tech, Inc.

³ HEXTM (Hexachloro-fluorescein), a trademark of Thermo Fisher Scientific.

⁴ TEX615TM is a trademark of Thermo Fisher Scientific.

⁵ BHQ-2TM (Black Hole Quencher) is a trademark of Biosearch Tech, Inc.

⁶ Cy5TM is a trademark of GE Healthcare.

⁷ TaqManTM is a trademark of Roche Diagnostics, Inc.

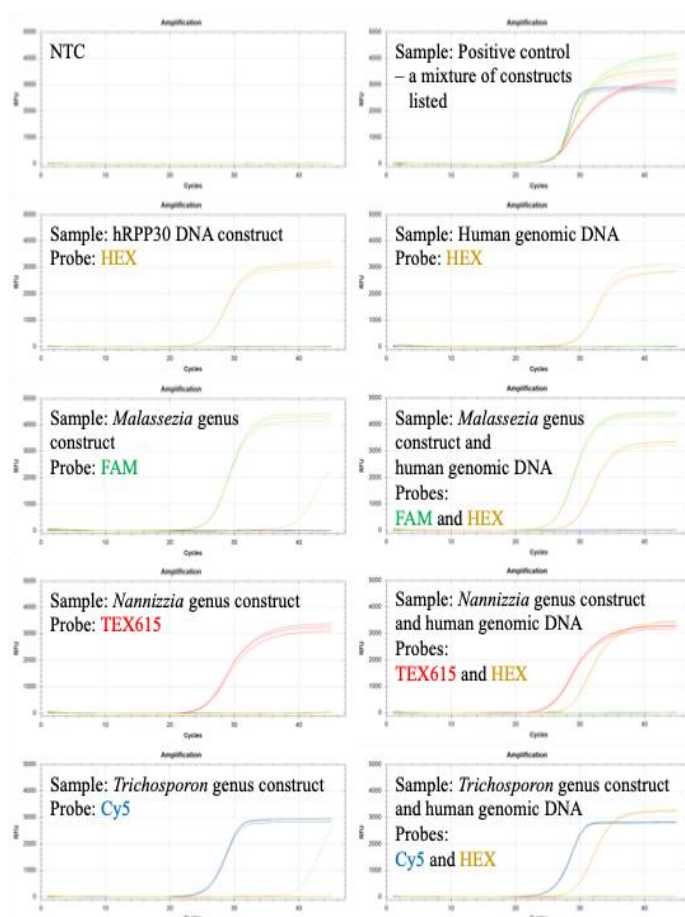


Figure 1: Verification experiments with single and multiple targets (given in text boxes for each panel). All sets of probes and primers are present in every reaction, but positive signal is only observed when the target(s) is present, indicating that the amplification is specific.

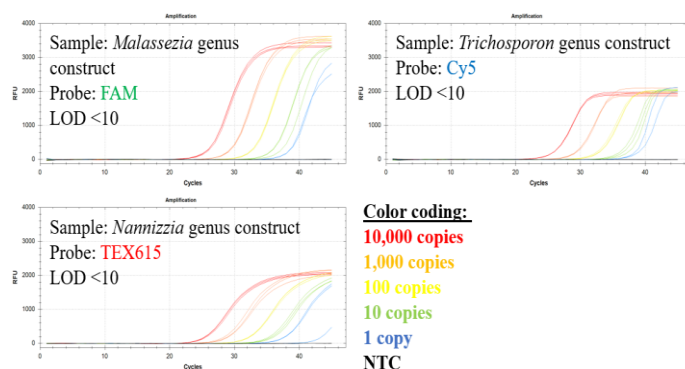


Figure 2: Serial dilution experiments show LOD <10 molecules for the synthetic DNA construct of each target.

CONTACT US

For assistance, please contact DNA Software using the link:

<https://www.pcrassays.com/contact/>

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