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Protocol for quantification of BdDV-1 mycovirus of Batrachochytrium dendrobatidis by qPCR

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Protocol status: Working We use this protocol and it's working

ABSTRACT

This protocol is used for quantifying the amount of the DNA mycovirus BdDV-1 in DNA extracts or cDNA libraries. This virus is associated with the fungus Batrachochytrium dendrobatidis (Bd), and it is known to be endogenized in the genome of a large number of Bd strains. The assay uses real time quantitative PCR with Taqman probes. Setup of the reaction is typically done in a biosafety cabinet. This protocol has been optimized on a QuantStudio 3 (Applied Biosystems).

ATTACHMENTS

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80313

Keywords: BdDV-1 qPCR

GUIDELINES

Notes:

BdV F: CCTGAGTACCCTGATCACAATGTBdV R: GGGTCATTGGTCGTATCTTCA

■ BdV Probe: MBGNFQ-CCATGGTGGCGTTCT-NED

Standards for viral copy number were created by cloning PCR products from BdV F and BdV R primers into pCR 2.1-TOPO vector using the TOPO TA cloning kit. Plasmids from transformed E. coli colonies with successful insert were extracted using the Zyppy Plasmid Miniprep kit. DNA of plasmid extracts was quantified with a Qubit. Plasmid copy numbers were estimated using the formula:

Number of copies per uL = [(g/uL) / plasmid length in bp * 650)] * (6.022 * 10²³)

Extracts were then diluted to a final concentration of 10⁶ viral copies per total 35uL volume for qPCR standards.

MATERIALS

Materials:

- 96 well optical qPCR plate (Applied Biosystems 4306737)
- p1000, p200, p20, p10 pipettes
- Barrier tips for all pipettes (recommend using a new unopened box of 10uL tips)
- Optical adhesive sealing film (Applied Biosystems 4311971)
- Tabletop centrifuge (for spinning down samples and reagents)
- Large centrifuge (for spinning down 96 well plate)
- PCR water
- Radiant Probe Lo-ROX qPCR Kit (Alkali Scientific QP9005)
- BdV F and BdV R Primers ([M] 18 micromolar (µM))
- BdV Probe ([M] 5 micromolar (µM))—**light sensitive**, tubes should be wrapped in aluminum foil when not in use
- BSA (<u>A</u> 400 ng/µl)
- 10⁶ BdDV-1 standard
- MicroAmp™ Optical 96-Well Reaction Plate with Barcode **Thermo Fisher Catalog** #4306737
- ABI prism optical adhesive covers **Thermo Fisher Scientific Catalog** #4311971
- Radiant™ Probe Lo-ROX qPCR Kits **Alkali Scientific Catalog**#OP9005

Protocol:

15m 41s

Wipe down pipettes with 70% ethanol and place in the biosafety hood with tips and 96 well plate.

UV for at least (5) 00:15:00 (longer is fine).

15m

After setting up hood to UV, prepare standard dilutions on the bench using barrier tips. Fill 6



- 2.1 Add \angle 10 μ L of the 10⁶ virus standard to the 10⁵ tube, pipetting up and down several times to mix.
- Repeat this, adding \square 10 μ L from the 10⁵ tube to the 10⁴ tube and so on. You should have 7 standards total, 10^6 - 10^0 .
- Prepare the master mix in the biosafety hood. All reagents can be thawed quickly on the bench, flicked to mix, and briefly spun down on a tabletop centrifuge before preparation. All the reagents and master mix should be kept on ice during the preparation. Volumes per well:

A	В
Radiant Probe Lo-ROX qPCR Kit	12.5 uL
PCR water	2.75 uL
BdV F primer (18 uM)	1.25 uL
BdV R primer (18 uM)	1.25 uL
BdV probe (5 uM) –protect from light when not in use	1.25 uL
BSA (400 ng/uL)	1 uL

- Add master mix to the 96 well plate. Put the plate on a 96 well rack and place it On ice. Add

 20 µL of master mix to each well on the plate that will be used. A single barrier tip can be used for all the wells.
- Add standards and samples to the 96 well plate. Standard tubes (prepped in step 2) and samples should be thawed, flicked to mix, and briefly spun down on a tabletop centrifuge. Add 5 µL of each standard or sample to the appropriate well, using a new 10uL tip for each well. If you are using a new box of tips, you can use the placement of tips in the box to help keep track of which

wells have been loaded. Negative control wells should be loaded with PCR water. Example plate setup below (where RAC### are experimental sample numbers):

Α	В	С	D	E	F	G	Н	I	J	K	L	М
	1	2	3	4	5	6	7	8	9	10	11	12
A	10^6	10^6	10^6	RAC 70	RAC 70	RAC8	RAC84	RAC1 08	RAC1 08	RAC1 23	RAC1 23	RAC1 36
В	10^5	10^5	10^5	RAC 71	RAC 71	RAC8	RAC85	RAC1	RAC1	RAC1 26	RAC1 26	RAC1 36
С	10^4	10^4	10^4	RAC 72	RAC 72	RAC8	RAC86	RAC1	RAC1	RAC1 27	RAC1 27	RAC1 37
D	10^3	10^3	10^3	RAC 73	RAC 73	RAC8	RAC87	RAC1	RAC1	RAC1	RAC1	RAC1 37
Ε	10^2	10^2	10^2	RAC 80	RAC 80	RAC8	RAC88	RAC1	RAC1	RAC1	RAC1	RAC1 38
F	10^1	10^1	10^1	RAC 81	RAC 81	RAC8	RAC89	RAC1 20	RAC1 20	RAC1	RAC1	RAC1 38
G	10^0	10^0	10^0	RAC 82	RAC 82	RAC1 01	RAC10	RAC1 21	RAC1 21	RAC1 34	RAC1 34	RAC1
Н	negati ve	negati ve	negati ve	RAC 83	RAC 83	RAC1 06	RAC10	RAC1 22	RAC1 22	RAC1 35	RAC1 35	RAC1 39

- Once all samples have been added to the plate, seal the plate with optical adhesive sealing film. Make sure that the edges and corners are well sealed. Using the corner of a plastic sealing tool to press down the edges and corners is helpful.
- Spin down the entire plate in a centrifuge briefly to make sure the liquid is at the bottom of the wells. Try to remove any bubbles be gently tapping the plate against the bench and spinning again.
- Set up the qPCR machine for the run. Make sure that the plate is On ice while you are doing this step (it can also be done ahead of time).
- 8.1 Open up the QuantStudio software. Name the experiment. Set Run Mode to "Fast".

- 8.2 Make sure total well volume is set to $\mathbb{Z}_{25 \, \mu L}$. Run conditions are:
 - 1. § 95 °C for 00:00:20 (set ramp up to 3.96)
- 8.3 Input the names for your wells into the software. This can be done manually or by copying and pasting the sample names in a list format from excel, where wells are labeled 1-96 and sample names are listed in order of A1-A12, then B1-12, etc.
- 8.4 Mark all of the standard wells as standards in the software. Input the correct quantities for each of the standards (e.g., the quantity for the 10⁶ sample should be input as 1000000).
- 8.5 Label the negative control wells as negatives and the rest of the occupied wells as sample wells (or unknowns).
- **9** Place the 96 well plate into the machine (making sure it is in the correct orientation) and start the run!
- At the end of the run, check the standards to make sure that the replicates reasonably overlap with one another and that there is no amplification in the negative controls. It is not unusual to have outliers or no amplification in some of the lower dilutions. Check the standard curve. Efficiency should be between 90-110%, R2 should be as close to 1 as possible, and error should be as close to 0 as possible. If necessary, outlier standard replicates can be removed in the software to change these values and the samples can be reanalyzed.
- 11 Export your data as an .xls file. This will contain the quantities for each sample, including the mean and SD of the replicates.