

Mar 13, 2020

Live/Dead qPCR of *B. pertussis* IS481

 PLOS One

DOI

[dx.doi.org/10.17504/protocols.io.bc5niy5e](https://doi.org/10.17504/protocols.io.bc5niy5e)

Iain MacArthur¹, Stacy Ramkissoon¹, Andrew Preston¹

¹[Milner Centre for Evolution, Department of Biology and Biochemistry, University of Bath, Bath, UK]



Stacy Ramkissoon

OPEN  ACCESS



DOI: [dx.doi.org/10.17504/protocols.io.bc5niy5e](https://doi.org/10.17504/protocols.io.bc5niy5e)

External link: <https://doi.org/10.1371/journal.pone.0232334>

Protocol Citation: Iain MacArthur, Stacy Ramkissoon, Andrew Preston 2020. Live/Dead qPCR of *B. pertussis* IS481. [protocols.io
https://doi.org/10.17504/protocols.io.bc5niy5e](https://doi.org/10.17504/protocols.io.bc5niy5e)

Manuscript citation:

Ramkissoon S, MacArthur I, Ibrahim M, Graaf Hd, Read RC, Preston A (2020) A qPCR assay for *Bordetella pertussis* cells that enumerates both live and dead bacteria. PLoS ONE 15(4): e0232334. doi: [10.1371/journal.pone.0232334](https://doi.org/10.1371/journal.pone.0232334)

License: This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: March 02, 2020

Last Modified: March 13, 2020

Protocol Integer ID: 33678

Keywords: enumeration, live, dead, qPCR, propidium monoazide, PMA,

Abstract

This protocol describes the use of qPCR for the quantitative determination of live *B. pertussis* bacteria in a sample. The method uses propidium monoazide, a compound that can cross the cell wall of dead bacteria and binds DNA following photo activation to inhibit amplification of DNA from dead bacteria. This method uses a TaqMan probe specific to IS481 to ensure maximum sensitivity due to the high copy number of the target. Although IS481 is not specific to *B. pertussis* for experimental purposes it provides acceptable specificity.

Attachments



[SOP_001.3_live_dead...](#)

30KB

Materials

MATERIALS

- ☒ Water
- ☒ Centrifuge
- ☒ Microcentrifuge
- ☒ Microcentrifuge tubes Denville Scientific Inc. Catalog #C2170
- ☒ StepOnePlus™ Real-Time PCR System Thermo Fisher Scientific Catalog #4376600
- ☒ MicroAmp® Fast Optical 96-Well Reaction Plate, 0.1 mL Thermo Fisher Catalog #4346907
- ☒ TaqMan® Gene Expression Master Mix Thermo Fisher Catalog #4369016
- ☒ Adhesive PCR Plate Seals Thermo Fisher Catalog #AB0558
- ☒ PMA-Lite™ LED Photolysis Device Biotium Catalog #E90002
- ☒ 1.5 ml Crystal Clear Microcentrifuge Tube StarLab Catalog #E1415-1500
- ☒ QIAamp DNA Mini Kit Qiagen Catalog #51304
- ☒ RNase A Merck MilliporeSigma (Sigma-Aldrich) Catalog #R4642
- ☒ PMA Dye 20 mM in H2O Biotium Catalog #40019
- ☒ 1000 µl Filter Tip (Sterile) Racked StarLab Catalog #S1126-7810
- ☒ 200 µl Graduated Filter Tip (Sterile) Racked StarLab Catalog #S1120-8810
- ☒ 20 µl Bevelled Filter Tip (Sterile) Racked StarLab Catalog #S1120-1810
- ☒ 10 µl Graduated Filter Tip (Sterile) Racked StarLab Catalog #S1121-3810
- ☒ Primers (IS481F/R)
- ☒ Probe (IS481)
- ☒ Control DNA for standard curve (Purified DNA from B1917 or strain being tested)

	Sequence (5'-3')
IS481 Forward Primer	ATCA AGCA CCG CTTT ACCC
IS481 Reverse Primer	TTGG GAGT TCTG GTAG GTGT G

IS481 Probe	FAM- AATG GCAA GGC CGAA CGCT TCA- BHQ1
----------------	-------------------------------------------------------------

Primers (IS481F/R) and Probe (IS481)

Safety warnings

- ! Good Laboratory Practice must be followed. Wear laboratory coats and gloves.

Before start

Principle

To use propidium monoazide (PMA) to inhibit the PCR amplification signal of DNA from dead bacteria. Purified DNA will be used as a template for the quantification of *IS481* using a TaqMan probe. A standard curve consisting of DNA from pure B1917 will be used to determine absolute quantity in unknowns.

PMA treatment

- 1 Pellet fresh samples by centrifuging at 2,000 x g for 10 minutes and resuspend in 1.2 ml of PBS. 10m
- 2 Transfer 200 μ l of resuspended sample into two clear microfuge tubes (continue with 1 sample to Step #3 and 1 sample skip to Step #4). 10m
- 3 Add 0.5 μ l of PMA to one of the 200 μ l of samples from Step #2. 10m
- 4 Incubate microfuge tubes in the dark for 10 min at room temperature. Cover samples with aluminium foil and incubate on a rocker. 10m
- 5 Expose samples to light using the PMA-Lite™ LED Photolysis Device for 5 min. 5m

DNA purification

- 6 Add 20 μ l of QIAGEN Protease and 4 μ l of RNase A.
- 7 Add 200 μ l of Buffer AL and vortex for 15 s. 15s
- 8 Incubate at 56°C for 10 min. 10m
- 9 Briefly spin tube to recover condensation. 10m
- 10 Add 200 μ l of ethanol (96-100%), vortex and spin briefly.
- 11 Add to spin column and centrifuge for 1 min at 6000xg. 1m
- 12 Transfer to a new collection tube.

13 Add 500µl AW1 buffer and spin for 1 min at 6000xg.

1m

14 Transfer to a new collection tube.

15 Add 500µl AW2 buffer and spin for 3 min at 17000xg

3m

16 Transfer to a 2ml microfuge tube.

17 Spin at max speed for 1 min.

1m

18 Transfer to a 1.5ml microfuge tube.

19 Add 200µl of Buffer AE and incubate at room temperature for 1 min.

1m

20 Elute by centrifuging for 1 min at 6000xg.

1m

qPCR

21 Dilute stock primers (50µM) 9 in 50µl, to give a reaction concentration of 900nm.

22 Dilute stock probe (50µM) 3 in 100µl, to give a reaction concentration of 150nm.

23

Prepare Mastermix as follows:

Taqm an MM	10µl	1x
Forw ard prime r	2µl	900n M

	Reverse primer	2µl	900nM
	Probe	2µl	150nM

Mastermix per reaction

- 24 Add 16µl of master mix per well.
- 25 Serial dilute 9 times, 5:50 positive control DNA (25ng/ul).
- 26 Use dilutions 3-9 for 1000, 100, 10, 1, 0.1, 0.01, 0.001pg.
- 27 Load in triplicate 4µl of DNA for each control dilution, unknown and water samples.

PCR and analysis

- 28 Cover plate with film and spin for 1 min at 1000xg

1m

- 29 Set up qPCR program as follows:

	50°C	2min
	95°C	10min
40 cycles:	95°C	15sec
	60°C	1min

qPCR Cycle

Results

- 30 Run analysis on data using StepOnePlus™ Software v2.3.
- 31 Outliers of triplicate samples should be ignored (omit).

- 32 Confirm standard curve has an $r^2 > 0.95$.
- 33 Convert pg of DNA to copy number: $B. pertussis/\text{sample} = (50 \times (\text{mass of template in pg}) \times 6.022 \times 10^{23}) / (\text{length of genome in bp} \times 1 \times 10^{12} \times 650)$.
- 34 Untreated sample (no PMA), represents total *B. pertussis* in the sample.
- 35 PMA treated sample gives number of live *B. pertussis* in the sample.
- 36 The number of dead can be determined from the difference between untreated and treated.