




Jul 30, 2020

16S Bacteria 338F-516P-805R BSA


Roey Angel¹, Eva Petrova¹¹Soil and Water Research Infrastructure**1** Works for me dx.doi.org/10.17504/protocols.io.qckdsuw

SoWa RI Anaerobic and Molecular Microbiology (public)

Tech. support email: eva.petrova@bc.cas.cz Eva Petrova
Soil and Water Research Infrastructure

ABSTRACT

Universal 16S rRNA probe-based-qPCR assay for bacteria.
The primers and probe are taken from [Yu et al. \(2005\)](#).

 Yu Y, Lee C, Kim J, Hwang S (2005). Group-specific primer and probe sets to detect methanogenic communities using quantitative real-time polymerase chain reaction. Biotechnology and bioengineering. <http://dx.doi.org/10.1002/bit.20347>

THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Yu, Y., Lee, C., Kim, J., and Hwang, S. (2005). Group-specific primer and probe sets to detect methanogenic communities using quantitative real-time polymerase chain reaction. Biotechnol Bioeng 89, 670–679. doi:10.1002/bit.20347.

ATTACHMENTS

[Introduction_QPCR_Strata](#) [AB_rt-QPCRguide.pdf](#)
[gene.pdf](#)

DOI

dx.doi.org/10.17504/protocols.io.qckdsuw

PROTOCOL CITATION

Roey Angel, Eva Petrova 2020. 16S Bacteria 338F-516P-805R BSA. **protocols.io**
dx.doi.org/10.17504/protocols.io.qckdsuw

MANUSCRIPT CITATION please remember to cite the following publication along with this protocol

Yu, Y., Lee, C., Kim, J., and Hwang, S. (2005). Group-specific primer and probe sets to detect methanogenic communities using quantitative real-time polymerase chain reaction. Biotechnol Bioeng 89, 670–679. doi:10.1002/bit.20347.

KEYWORDS

qPCR, dual-labelled probe, 16S rRNA gene, bacteria

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

CREATED

May 24, 2018

LAST MODIFIED

Jul 30, 2020

PROTOCOL INTEGER ID

12396

MATERIALS

NAME	CATALOG #	VENDOR
iQ™ SYBR® Green Supermix	1708880	BioRad Sciences

Primers and probe

1

Name	Type	Sequence	Target region ¹
BAC338F	Forward	ACT CCT ACG GGA GGC AG	338-354
BAC516P ²	Probe	TGC CAG CAG CCG CGG TAA TA	516-536
BAC805R	Reverse	GAC TAC CAG GGT ATC TAA TC	785-805

1. Relative to *E. coli* 16S rRNA gene.

2. The probe must be dual-labelled either with 5'-6-FAM, 3'-BHQ1 or any other valid combination.

qPCR mixture

2

Reagent	Final concentration	1 tube (20 µl)	plate (20 µl x 100)
PCR H ₂ O		4.6	460
iQ™ Supermix	1x	10	1000
MgCl ₂ (25 mM)	4.0 mM	0.8 ¹	80
BSA (20 µg µl ⁻¹)	0.2 µg µl ⁻¹	0.2	20
338F (10 µM)	0.5 µM	1.0	100
805R (10 µM)	0.5 µM	1.0	100
516P (10 µM)	0.2 µM	0.4	40
Template		2	2 x 100

1 Buffer contains MgCl₂ at final conc. of 3.0 mM

Thermocycler programme

3

1. **95 °C** for **00:05:00**
2. x 40 {
 - 2.1 **95 °C** for **00:00:30**
 - 2.2 **62 °C** for **00:00:30** take snapshot}