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Version 3

## qPCR: Bacterial SSU rRNA 338F-516P-805R V.3

DOI

<https://dx.doi.org/10.17504/protocols.io.bi98kh9w>



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### Manuscript citation:

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**Protocol status:** Working

**We use this protocol and it's working**

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**Keywords:** qPCR, dual-labelled probe, 16S rRNA gene, bacteria, qpcr assay for bacteria, qpcr assay, rna probe, qpcr, 16s rna, rna, probe,

## Abstract

Universal 16S rRNA probe-based-qPCR assay for bacteria.

The primers and probe are taken from Yu et al. (2005).

### Citation

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>

LINK

## Materials

### MATERIALS

 iQ™ SYBR® Green Supermix **Bio-Rad Laboratories Catalog #1708880**

## Primers and probe

1

|  | Name                 | Type    | Sequence                   | Target region <sup>1</sup> |
|--|----------------------|---------|----------------------------|----------------------------|
|  | BAC338F              | Forward | ACT CCT ACG GGA GGC AG     | 338-354                    |
|  | BAC516P <sup>2</sup> | 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* SSU 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 $\mu$ l) | plate (20 $\mu$ l x 100) |
|--|-----------------------------------------|-----------------------------------|---------------------|--------------------------|
|  | PCR H <sub>2</sub> O                    |                                   | 4.6                 | 460                      |
|  | iQ <sup>TM</sup> Supermix               | 1x                                | 10                  | 1000                     |
|  | MgCl <sub>2</sub> (25 mM)               | 4.0 mM                            | 0.8 <sup>1</sup>    | 80                       |
|  | BSA (20 $\mu$ g $\mu$ l <sup>-1</sup> ) | 0.2 $\mu$ g $\mu$ l <sup>-1</sup> | 0.2                 | 20                       |
|  | <b>338F</b> (10 $\mu$ M)                | 0.5 $\mu$ M                       | 1.0                 | 100                      |
|  | <b>805R</b> (10 $\mu$ M)                | 0.5 $\mu$ M                       | 1.0                 | 100                      |
|  | <b>516P</b> (10 $\mu$ M)                | 0.2 $\mu$ M                       | 0.4                 | 40                       |
|  | Template                                |                                   | 2                   | 2 x 100                  |



1 Buffer contains  $\text{MgCl}_2$  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}

## Citations

Yu Y, Lee C, Kim J, Hwang S. Group-specific primer and probe sets to detect methanogenic communities using quantitative real-time polymerase chain reaction

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