



Version 3

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# General bacteria and archaea 16S-rRNA (515Fmod-806Rmod) for Illumina amplicon sequencing V.3

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SoWa RI Anaerobic and Molecular Microbiology (public)

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SUBMIT TO PLOS ONE

## ABSTRACT

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

The primers target the V4 region of the 16S rRNA gene and were specifically designed for Illumina amplicon sequencing. The original primers were designed by Caporaso *et al.* (2012) and modified by Walters *et al.* (2015). For barcoding, we use the [Fludigm Access Array](#) for barcoding the sample and therefore the primers are synthesized with the CS1 and CS2 regions.

Caporaso JG, Lauber CL, Walters WA, Berg-Lyons D, Huntley J, Fierer N, Owens SM, Betley J, Fraser L, Bauer M, Gormley N, Gilbert JA, Smith G, Knight R (2012). Ultra-high-throughput microbial community analysis on the Illumina HiSeq and MiSeq platforms.. The ISME journal.  
<https://doi.org/10.1038/ismej.2012.8>

Walters W, Hyde ER, Berg-Lyons D, Ackermann G, Humphrey G, Parada A, Gilbert JA, Jansson JK, Caporaso JG, Fuhrman JA, Apprill A, Knight R (2015). Improved Bacterial 16S rRNA Gene (V4 and V4-5) and Fungal Internal Transcribed Spacer Marker Gene Primers for Microbial Community Surveys.. mSystems.  
<http://10.1128/msystems.00009-15>

## DOI

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## PROTOCOL CITATION

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## KEYWORDS

PCR, 16S rRNA, SSU rRNA, Amplicon sequencing, Illumina sequencing, Barcoded sequencing, Targeted metagenomics, Microbiome

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47193

## MATERIALS TEXT

### STEP MATERIALS

- [Agarose Sigma](#)
- Aldrich Catalog #A9539** **Step 6**
- [GeneRuler DNA Ladder Mix Thermo Fisher](#)
- Scientific Catalog #SM0331** **Step 6**
- [DNA Gel Loading Dye \(6X\) Thermo Fisher](#)
- Scientific Catalog #R0611** **Step 6**
- [TAE buffer \(50x\), molecular biology grade Serva,](#)
- Germany Catalog #4254901** **Step 6**
- [Primer: 515Fmod\\_CS1 Elisabeth Pharmacon](#) **Step 1**
- [Primer: 806mod\\_CS2 Elisabeth Pharmacon](#) **Step 1**
- [DreamTaq Green DNA Polymerase \(5 U/μL\) Thermo Fisher](#)
- Scientific Catalog #EP0712** **Step 2**
- [dNTP Set \(100 mM each\) Contributed by](#)
- users Catalog #BR0600601** **Step 2**
- [PCR H2O Top](#)
- Bio Catalog #P040** **Step 2**
- [Bovine Serum Albumin \(BSA\) Thermo Fisher](#)
- Scientific Catalog #B14** **Step 2**

## Primers

1

Name	Direction	Sequence <sup>1</sup>	Target region <sup>2</sup>
515Fmod_CS1	Forward	ACA CTG ACG ACA TGG TTC TAC AGT <b>GYC AGC MGC CGC CGT AA</b>	515-533
806Rmod_CS2	Reverse	TAC GGT AGC AGA GAC TTG GTC <b>TGG ACT ACN VGG GTW TCT AAT</b>	787-806

1. CS + primer sequence (in bold)
2. Relative to *E. coli* SSU rRNA gene

[🔗 Primer: 515Fmod\\_CS1 Elisabeth Pharmacon](#)

[🔗 Primer: 806mod\\_CS2 Elisabeth Pharmacon](#)

#### PCR reaction

- 2 Prepare the following master mixture [🔗 On ice](#).

Don't forget to prepare an additional mixture for the negative (NTC) and positive controls, and to account for pipetting errors.

Reagent	Final. conc.	1 tube (25 $\mu$ l)	100 reactions (96-well plate; $\mu$ l)
PCR H <sub>2</sub> O		17.525	1752.5
10X DreamTaq Green Buffer	1X	2.5	250
dNTP (2 mM each)	0.2 mM	2.5	250
BSA (20 $\mu$ g/ $\mu$ l)	80 ng $\mu$ l <sup>-1</sup>	0.1	10
<b>515Fmod-CS1 (10 <math>\mu</math>M)</b>	0.2 $\mu$ M	0.625	62.5
<b>806Rmod-CS2 (10 <math>\mu</math>M)</b>	0.2 $\mu$ M	0.625	<b>62.5</b>
DreamTaq Green DNA Polymerase	0.625 U	0.125	12.5
<b>Final volume</b>		<b>24</b>	<b>2400</b>

[🔗 DreamTaq Green DNA Polymerase \(5 U/ \$\mu\$ L\) Thermo Fisher](#)

**Scientific Catalog #EP0712**

[🔗 dNTP Set \(100 mM each\) Contributed by](#)

**users Catalog #BR0600601**

[🔗 PCR H<sub>2</sub>O Top](#)

**Bio Catalog #P040**

[🔗 Bovine Serum Albumin \(BSA\) Thermo Fisher](#)

**Scientific Catalog #B14**

- 3 Vortex and spin down [🕒 00:00:03](#)

3s

- 4 Distribute [📄 24  \$\mu\$ l](#) of the mixture to each tube and add [📄 1  \$\mu\$ l](#) of template DNA or cDNA

PCR reaction 3s

17m 15s

5 Run the following PCR program:

1. 94 °C 00:05:00
2. x 28 {
  - 2.1 94 °C 00:00:45
  - 2.2 52 °C 00:00:45
  - 2.3 72 °C 00:00:45
3. 72 °C 00:10:00
4. 4 °C hold

Evaluate PCR products on an agarose gel 40m

6 Prepare a 1.5% agarose gel by mixing:

100 mL TAE

1.5 g agarose

Heat in the microwave until dissolved and pour into a gel frame.  
Place solid gel into an electrophoresis bath filled with TAE buffer.

Agarose Sigma

Aldrich Catalog #A9539

GeneRuler DNA Ladder Mix Thermo Fisher

Scientific Catalog #SM0331

DNA Gel Loading Dye (6X) Thermo Fisher

Scientific Catalog #R0611

TAE buffer (50x), molecular biology grade Serva,

Germany Catalog #4254901

7 Mix up to 5 µl of the PCR reaction sample with 1 µl of loading dye and load the sample into a well.  
In addition load 5 µl of DNA ladder mix (80-10,000 bp) into an empty well, as a marker.

40m

8 Run the gel at 110V, 265mA for approx. 00:40:00

9 Stain gel for at least 40min in an Ethidium bromide TAE bath (or any other DNA stain).

10 Visualise the gel using a gel documentation system.