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General bacteria and archaea 16S-rRNA (515Fmod-806Rmod) for Illumina amplicon sequencing V.2

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SOWA

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Protocol status: Working

We use this protocol and it's working

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



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.

CITATION

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.

LINK

<https://doi.org/10.1038/ismej.2012.8>

CITATION

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.

LINK

<https://doi.org/pii:e00009-15>



Materials

STEP MATERIALS

- ✕ 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**
- ✕ Primer: 515Fmod_CS1 **Elisabeth Pharmacon**
- ✕ Primer: 806mod_CS2 **Elisabeth Pharmacon**
- ✕ DreamTaq Green DNA Polymerase (5 U/ μ L) **Thermo Fisher Scientific Catalog #EP0712**
- ✕ dNTP Set (100 mM each) **Catalog #BR0600601**
- ✕ PCR H₂O **Top Bio Catalog #P040**
- ✕ Bovine Serum Albumin (BSA) **Thermo Fisher Scientific Catalog #B14**



Protocol materials

☒ DNA Gel Loading Dye (6X) **Thermo Fisher Scientific Catalog #R0611**

☒ TAE buffer (50x), molecular biology grade **Serva, Germany Catalog #4254901**

☒ dNTP Set (100 mM each) **Catalog #BR0600601**

☒ Agarose **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A9539**

☒ GeneRuler DNA Ladder Mix **Thermo Fisher Scientific Catalog #SM0331**

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☒ Primer: 515Fmod_CS1 **Elisabeth Pharmacon**

☒ Primer: 806mod_CS2 **Elisabeth Pharmacon**

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Primers


1

Name	Direction	Sequence ¹	Target region ²
515Fmod_CS1	Forward	ACA CTG ACG ACA TGG TTC TAC AGT GYC AGC MGC CGC CGT AA	515-533
806Rmod_CS2	Reverse	TAC GGT AGC AGA GAC TTG GTC TGG ACT ACN VGG GTW TCT AAT	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 µl)	100 reactions (96-well plate; µl)
PCR H ₂ 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 µg/µl)	80 ng µl ⁻¹	0.1	10
515Fmod-CS1 (10 µM)	0.2 µM	0.625	62.5
806Rmod-CS2 (10 µM)	0.2 µM	0.625	62.5
DreamTaq Green DNA Polymerase	0.625 U	0.125	12.5
Final volume		24	2400



DreamTaq Green DNA Polymerase (5 U/ μ L) **Thermo Fisher Scientific Catalog #EP0712**



dNTP Set (100 mM each) **Catalog #BR0600601**





PCR H₂O **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 μ L of the mixture to each tube and add  1 μ L of template DNA or cDNA

PCR reaction

3s

5 Run the following PCR program:

17m 15s

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 **Merck MilliporeSigma (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.

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

40m

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.

Citations

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. Ultra-high-throughput microbial community analysis on the Illumina HiSeq and MiSeq platforms.

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