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General bacteria and archaea 16S-rRNA (515Fmod-806R) for Illumina amplicon sequencing

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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.
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KEYWORDS

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

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Jan 29, 2018  Eva Petrova Soil and Water Research Infrastructure

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9870

STEPS MATERIALS

NAME	CATALOG #	VENDOR
Agarose	A9539	Sigma Aldrich
GeneRuler DNA Ladder Mix	SM0331	Thermo Fisher Scientific
DNA Gel Loading Dye (6X)	R0611	Thermo Fisher Scientific
TAE buffer (50x), molecular biology grade	4254901	Serva, Germany
Fast Start PCR Master	196 10 126	Roche
PCR H2O	P040	Top Bio
Bovine Serum Albumin (BSA)	B14	Thermo Fisher Scientific
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
806mod_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

by Elisabeth Pharmacon

[View](#)



Primer: 806mod_CS2

by Elisabeth Pharmacon

[View](#)

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		9.3	930
Fast Start PCR Master (2x)	1x	12.5	1250
BSA (20 µg/µl)	0.6 µg/µl	0.7	70
515Fmod-CS1 (10 µM)	0.3 µM	0.75	75
806R-CS2 (10 µM)	0.3 µM	0.75	75
Final volume		24	2400



Fast Start PCR Master

by Roche

Catalog #: 196 10 126



PCR H2O

by Top Bio

Catalog #: P040





Bovine Serum Albumin (BSA)

by Thermo Fisher Scientific

Catalog #: B14

3 Vortex and spin down ⌚ 00:00:03

3s







4 Distribute  24 µl of the mixture to each tube and add  1 µl of template DNA or cDNA

PCR reaction

3s

5 Run the following PCR program:

17m 15s

1.  94 °C ⌚ 00:05:00
2. x 27 {
 - 2.1  94 °C ⌚ 00:00:45
 - 2.2  50 °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

by Sigma Aldrich

Catalog #: A9539



GeneRuler DNA Ladder Mix

by Thermo Fisher Scientific

Catalog #: SM0331



DNA Gel Loading Dye (6X)

by Thermo Fisher Scientific





Catalog #: R0611



TAE buffer (50x), molecular biology grade

by 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**
- 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.

40m