



Mar 11,
2019

Working

16S Metagenomics in a Field Setting

Version 1

Mrinalini Watsa¹, Gideon Erkenwick², Aaron Pomerantz³, Stefan Prost⁴

¹University of Missouri - Saint Louis; Washington University in Saint Louis; Field Projects International, ²University of Missouri - Saint Louis; Field Projects International, ³University of California, Berkeley, CA, USA, Department of Integrative Biology; Marine Biological Laboratory, Woods Hole, MA - USA, ⁴LOEWE-Center for Translational Biodiversity Genomics, Senckenberg Museum, 60325 Frankfurt, Germany; South African National Biodiversity Institute, National Zoological Garden, Pretoria 0184, South Africa

[dx.doi.org/10.17504/protocols.io.y2ffbybn](https://doi.org/10.17504/protocols.io.y2ffbybn)

Mrinalini Watsa

ABSTRACT

This protocol is was used to conduct DNA 16S metagenomics on FPI's Genomics in the Jungle - 2018 field course at the Green Lab, located and Inkaterra Guides Field Station, Madre de Dios, Peru.

PROTOCOL STATUS

Working

We use this protocol in our group and it is working

GUIDELINES

This protocol starts from already extracted DNA

STEPS MATERIALS

NAME

Agencourt Ampure XP

CATALOG

A63880

VENDOR

Beckman Coulter

SAFETY WARNINGS

Amplification

1 Remove samples and the following reagents and let thaw, once thawed keep on ice block

- 10mM DNTPs
- 25 mMgCl
- 5x Go Taq Buffer
- Forward primer 16S-27F (10uM)
- Reverse primer 16S-1429R (10uM)
- GoTaq Hotstart Polymerase 5u/ul

Make PCR cocktail for # of samples * 1.1 (10% extra). Don't forget to include 1 PCR negative control for each separate PCR

Run PCR according to the following cycle conditions:

- Initial denaturation **95 °C** for 120s
- 25 cycles of **95 °C** for 30s, **51 °C** for 30s, **72 °C** for 30s
- Final extension **72 °C** for 420s

Upon completion remove, label, and store at **4 °C** , or take directly to electrophoresis

2 Equipment



- BlueGel system
- MiniOne system

Create .8 - 1.0% agarose 1 gel with 13 combs

- Measure 1 g of agarose
- Mix agarose with 100 mL of 1xTBE
- Microwave the mixture until agarose is completely dissolved (1-3 min)
- Pour the agarose gel into the tray with the comb in place.
- Allow the agarose gel to harden (20-30 min)

Insert the agarose gel into electrophoresis equipment and add 1xTBE buffer until the agarose gel is submerged

Spot check with  2 µl of each sample


Mix  1 µl of loading dye to  2 µl of each sample and load the gel. (If Green Taq buffer with built in loading dye was used, skip this step).

Load  5 µl of 100kp ladder into the agarose gel.


Turn on the electrode and let the DNA run until the band is identifiable (


Quantification

3 Equipment



Quantus Fluorometer
DNA and RNA Quantifier

Promega E6150 



- dsDNA dye
- Qubit Assay Tubes
- Your DNA sample


Pulse vortex your sample and spin down.

Add  200 µl of dsDNA dye into the qubit assay tube.

- Make sure to cover up the tube from light

Transfer  1 µl of DNA sample into qubit assay tube

Pulse vortex and spin down

Let it sit for  00:05:00 in room temperature

- Covered from the light

Gently vortex and spindown the sample

Calibrate the Quantus Fluorometer by standard and reference testing.

- Follow the same process but use  1 µl ddH₂O and Lambda DNA.

Insert the qubit assay tube into the Quantus Flurometer

Normalization and Pooling

4 Dilute each sample to 50 nM

- Mix 5 uL of the sample with calculated amount of ddH₂O to make each sample the same concentration of 50 nM
- Then pool 5 uL of each dilution into a single tube.
- This is now the library.

SPRI Clean-up

- 5 Run a SPRI cleanup of the library using your choice of bead purification systems/kits in a 1:1 ratio. Resuspend in the same volume.



Agencourt Ampure XP

by Beckman Coulter

Catalog #: [A63880](#)



This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited