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Working

## 16S Metagenomics in a Field Setting

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#### **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

CATALOG # **VENDOR** NAME A63880 Agencourt Ampure XP Beckman Coulter

SAFETY WARNINGS

### Amplification

- Remove samples and the following reagents and let thaw, once thawed keep on ice block
  - 10mM DNTPs
  - 25 mMqCl
  - 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



Electrophorogic

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Electrophoresis

### 2 Equipment

- BlueGel system
- MiniOne system

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

- Measure 1 g of agarose
- Mix agraose 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  $\frac{1}{2}$   $\mu$ l of each sample

Mix  $\frac{1}{2}$   $\mu$ I of loading dye to  $\frac{1}{2}$   $\mu$ I of each sample and load the geI. (If Green Taq buffer with built in loading dye was used, skip this step).

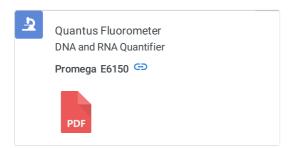
Load 5 µl of 100kp ladder into the agarose gel.

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

Quantification

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# **Equipment**



- dsDNA dye
- Qubit Assay Tubes
- Your DNA sample

Pulse vortex your sample and spin down.

Add  $200 \mu$  of dsDNA dye into the qubit assay tube.

Make sure to cover up the tube from light

Transfer 11 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 Fluormeter by standard and reference testing.

Follow the same process but use 
□1 μl ddH<sub>2</sub>O and Lambda DNA.

## Insert the qubit assay tube into the Quantus Flurometer

## Normalization and Pooling

- △ Dilute each sample to 50 nM
  - Mix 5 uL of the sample with calculated amound of ddH<sub>2</sub>O 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.



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