

16S rRNA Amplicon PCR (Adapted from EMP 16S amplification protocol*)

PCR reagent Master Mix

| Reagent | Volume |
|---|----------------|
| PCR Grade H ₂ O (ddH ₂ O) | 13.0 µL |
| 5 Primer Hot MM | 10.0 µL |
| Forward primer (10µM) | 0.5 µL |
| Reverse primer (10µM) | 0.5 µL |
| Template DNA | 1.0 µL |
| Total reaction volume | 25.0 µL |

Thermocycler Conditions for 96 well thermocyclers:

| Step | Temp | Time | # cycles |
|-----------------|------|------|----------|
| Hot Start | 94°C | 3m | 1 |
| Denature | 94°C | 45s | 35 |
| Anneal | 50°C | 60s | |
| Extend | 72°C | 90s | |
| Final Extension | 72°C | 10m | 1 |
| Hold | 4°C | Hold | Hold |

Note: Samples can be prepared in strip tubes or plates provided that the plastic is compatible with the thermocycler.

1. Prepare master mix in the order above for each sample in triplicate (*i.e.*, each sample is amplified in 3 individual replicate 25µL reactions).
Note: Each individual sample must have a unique barcoded primer, but all replicate PCR reactions of that sample will share that barcoded primer.
2. Place tubes into the thermocycler and start the program above.
3. At the end of the PCR program remove the tubes and place the samples on ice for 2m.
4. Spin down the samples briefly (~3-5sec).
5. Combine the triplicate PCR reactions for each sample into a single tube. Combination will result in a total of 75 µL of amplicon for each sample. **Do NOT combine amplicons from different samples at this point!**
6. Proceed to amplicon visualization and quantification protocols.

*Earth microbiome project(EMP; <http://www.earthmicrobiome.org>)