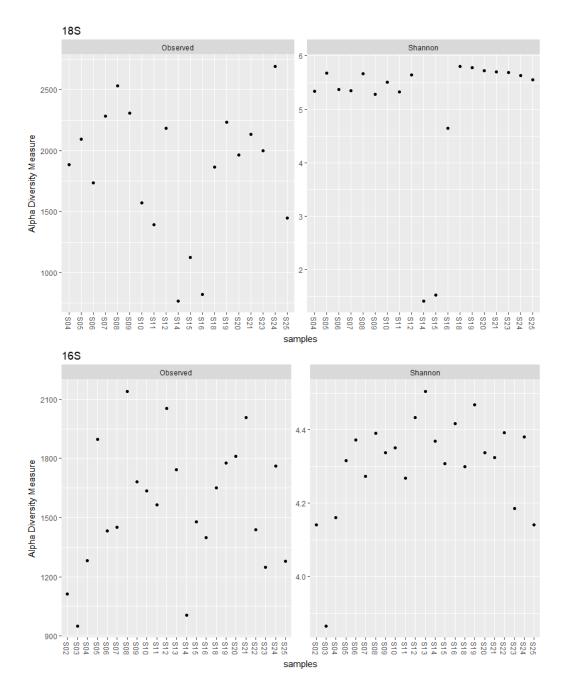


**Supplementary Figure 1:** Rarefaction plots showing OTUs observed versus sequencing depth for (**left**) 16S and (**right**) 18S rRNA amplicon samples. Samples were rarefied to the number of reads within the smallest included sample from the dataset.



**Supplementary Figure 2:** Observed OTUs per sample and Shannon diversity for **(top)** 18S and **(bottom)** 16S samples.