Bacterial Taxonomic Composition by Site and Layer 1.00 -Phylum D 1 Acidobacteria D 1 Fusobacteria D\_1\_\_Actinobacteria D\_1\_\_Gemmatimonadetes D\_1\_\_Armatimonadetes D\_1\_Nanoarchaeaeota 0.75 -D\_1\_Bacteroidetes D\_1\_Nitrospirae D\_1\_BRC1 D\_1\_\_Omnitrophicaeota D\_1\_Chlamydiae D\_1\_Patescibacteria Abundance D\_1\_Chloroflexi D\_1\_Planctomycetes 0.50 -D\_1\_Cyanobacteria D\_1\_Proteobacteria D\_1\_Deinococcus-Thermus D\_1\_Rokubacteria D\_1\_ Dependentiae D\_1\_SAR D\_1\_Elusimicrobia D\_1\_Thaumarchaeota D\_1\_Entotheonellaeota D\_1\_Verrucomicrobia 0.25 -D\_1\_\_Euryarchaeota D\_1\_WPS-2 D\_1\_FBP D\_1\_\_WS2 D\_1\_Fibrobacteres NA D\_1\_\_Firmicutes 0.00 -JTNP\_Sub CIMA\_SF GMT\_SF KELSO\_SF KELSO\_Sub

Site\_layer