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|----------------|----------------------|-----------------------|---------------|
| SR1 | OP10 | Gemmatimonadetes | Nitrospira |
| OD1 | Chlamydiae | OP11 | Synergistetes |
| Lentisphaerae | Chlorobi | TM7 | Thermotogae |
| BRC1 | Deferribacteres | Deinococcus-Thermus | WS3 |
| Fibrobacteres | unclassified_Archaea | Tenericutes | Fusobacteria |
| Crenarchaeota | Acidobacteria | Verrucomicrobia | Cyanobacteria |
| Planctomycetes | Chloroflexi | Euryarchaeota | Spirochaetes |
| Actinobacteria | Firmicutes | unclassified_Bacteria | Bacteroidetes |
| Proteobacteria | | | |

Supplemental Figure 2. SSU rRNA-based microbial community composition from datasets obtained from non-ribodepleted RNA (left) vs. ribodepleted RNA (right).