

- D_0_Archaea;D_1_Bathyarchaeota
- D_0_Archaea;D_1_Diapherotrites
- D_0_Archaea;D_1_Euryarchaeota
- D_0_Archaea;D_1_Miscellaneous Euryarchaeotic Group(MEG)
- D_0_Archaea;D_1_Parvarchaeota
- D_0_Archaea;D_1_Thaumarchaeota
- D_0_Archaea;D_1_Woesearchaeota (DHVEG-6)
- D_0_Bacteria;D_1_Acidobacteria
- D_0_Bacteria;D_1_Actinobacteria
- D_0_Bacteria;D_1_Aminicenantes
- D_0_Bacteria;D_1_Armatimonadetes
- D_0_Bacteria;D_1_BJ-169
- D_0_Bacteria;D_1_BRC1
- D_0_Bacteria;D_1_Bacteroidetes
- D_0_Bacteria;D_1_Candidatus Berkelbacteria
- D_0_Bacteria;D_1_Chlamydiae
- D_0_Bacteria;D_1_Chlorobi
- D_0_Bacteria;D_1_Chloroflexi
- D_0_Bacteria;D_1_Cyanobacteria
- D_0_Bacteria;D_1_Deferribacteres
- D_0_Bacteria;D_1_Deinococcus-Thermus
- D_0_Bacteria;D_1_Elusimicrobia
- D_0_Bacteria;D_1_FBP
- D_0_Bacteria;D_1_FCPU426
- D_0_Bacteria;D_1_Fibrobacteres
- D_0_Bacteria;D_1_Firmicutes
- D_0_Bacteria;D_1_Fusobacteria
- D_0_Bacteria;D_1_GAL15
- D_0_Bacteria;D_1_Gemmatimonadetes
- D_0_Bacteria;D_1_Gracilibacteria
- D_0_Bacteria;D_1_Hydrogenedentes
- D_0_Bacteria;D_1_Ignavibacteriae
- D_0_Bacteria;D_1_Latescibacteria
- D_0_Bacteria;D_1_Lentisphaerae
- D_0_Bacteria;D_1_Microgenomates
- D_0_Bacteria;D_1_Nitrospirae
- D_0_Bacteria;D_1_Omnitrophica
- D_0_Bacteria;D_1_PAUC34f
- D_0_Bacteria;D_1_Parcubacteria
- D_0_Bacteria;D_1_Peregrinibacteria
- D_0_Bacteria;D_1_Planctomycetes
- D_0_Bacteria;D_1_Proteobacteria
- D_0_Bacteria;D_1_RBG-1 (Zixibacteria)
- D_0_Bacteria;D_1_RsaHf231
- D_0_Bacteria;D_1_SBR1093
- D_0_Bacteria;D_1_SR1 (Absconditabacteria)
- D_0_Bacteria;D_1_Saccharibacteria
- D_0_Bacteria;D_1_Spirochaetae
- D_0_Bacteria;D_1_Synergistetes
- D_0_Bacteria;D_1_TM6 (Dependentiae)
- D_0_Bacteria;D_1_Tectomicrobia
- D_0_Bacteria;D_1_Tenericutes
- D_0_Bacteria;D_1_Verrucomicrobia
- D_0_Bacteria;D_1_WS1
- D_0_Bacteria;D_1_WS2
- D_0_Bacteria;D_1_WWE3
- D_0_Bacteria;Other
- Unassigned;Other