

- D_0_Archaea;D_1_Euryarchaeota;D_2_Halobacteria
- D_0_Archaea;D_1_Nanohaloarchaeota;D_2_uncultured archaeon
- D_0_Archaea;D_1_Nanohaloarchaeota;D_2_uncultured haloarchaeon
- D_0_Archaea;D_1_Nanohaloarchaeota;Other
- D_0_Archaea;D_1_Thaumarchaeota;D_2_Soil Crenarchaeotic Group(SCG)
- D_0_Bacteria;D_1_Acidobacteria;D_2_Acidobacteria
- D_0_Bacteria;D_1_Actinobacteria;D_2_Acidimicrobiia
- D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria
- D_0_Bacteria;D_1_Actinobacteria;D_2_Nitriliruptoria
- D_0_Bacteria;D_1_Actinobacteria;D_2_Rubrobacteria
- D_0_Bacteria;D_1_Actinobacteria;D_2_Thermoleophilia
- D_0_Bacteria;D_1_Armatimonadetes;D_2_Armatimonadia
- D_0_Bacteria;D_1_Armatimonadetes;D_2_uncultured bacterium
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Bacteroidia
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Flavobacteriia
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Sphingobacteriia
- D_0_Bacteria;D_1_Chloroflexi;D_2_Ardenticatenia
- D_0_Bacteria;D_1_Chloroflexi;D_2_Ktedonobacteria
- D_0_Bacteria;D_1_Chloroflexi;D_2_Thermomicrobia
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Chloroplast
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria
- D_0_Bacteria;D_1_Deinococcus-Thermus;D_2_Deinococci
- D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli
- D_0_Bacteria;D_1_Firmicutes;D_2_Clostridia
- D_0_Bacteria;D_1_Fusobacteria;D_2_Fusobacteriia
- D_0_Bacteria;D_1_Gemmatimonadetes;D_2_Gemmatimonadetes
- D_0_Bacteria;D_1_Planctomycetes;D_2_Phycisphaerae
- D_0_Bacteria;D_1_Planctomycetes;D_2_Planctomycetacia
- D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria
- D_0_Bacteria;D_1_Proteobacteria;D_2_Betaproteobacteria
- D_0_Bacteria;D_1_Proteobacteria;D_2_Deltaproteobacteria
- D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria
- D_0_Bacteria;D_1_TM6;D_2_uncultured bacterium
- D_0_Bacteria;D_1_Verrucomicrobia;D_2_Spartobacteria
- Unassigned;Other;Other