- 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