

- D_0_Archaea;D_1_Euryarchaeota;D_2_Halobacteria;D_3_Halobacteriales
- D_0_Archaea;D_1_Nanohaloarchaeota;D_2_uncultured archaeon;D_3_uncultured archaeon
- D_0_Archaea;D_1_Nanohaloarchaeota;D_2_uncultured haloarchaeon;D_3_uncultured haloarchaeon
- D_0_Archaea;D_1_Nanohaloarchaeota;Other;Other
- D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Corynebacteriales
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia;D_3_Order II
- D_0_Bacteria;D_1_Bacteroidetes;D_2_Sphingobacteriia;D_3_Sphingobacteriales
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Chloroplast;D_3_uncultured Chlorophyta
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Chloroplast;Other
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;D_3_SubsectionI
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;D_3_SubsectionII
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;D_3_SubsectionIII
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;D_3_SubsectionIV
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;D_3_uncultured
- D_0_Bacteria;D_1_Cyanobacteria;D_2_Cyanobacteria;Other
- D_0_Bacteria;D_1_Planctomycetes;D_2_Phycisphaerae;D_3_Phycisphaerales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhizobiales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhodospirillales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Sphingomonadales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Deltaproteobacteria;D_3_Desulfuromonadales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria;D_3_Chromatiales
- D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria;D_3_Oceanospirillales
- D_0_Bacteria;D_1_TM6;D_2_uncultured bacterium;D_3_uncultured bacterium
- Unassigned;Other;Other;Other