

- 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;Other;Other
- D_0__Bacteria;D_1__Actinobacteria;D_2__Actinobacteria;D_3__Propionibacteriales
- D_0__Bacteria;D_1__Bacteroidetes;D_2__Cytophagia;D_3__Order II
- D_0__Bacteria;D_1__Cyanobacteria;D_2__Chloroplast;Other
- D_0__Bacteria;D_1__Cyanobacteria;D_2__Cyanobacteria;D_3__SubsectionIII
- D_0__Bacteria;D_1__Cyanobacteria;D_2__Cyanobacteria;D_3__uncultured
- D_0__Bacteria;D_1__Planctomycetes;D_2__Phycisphaerae;D_3__Phycisphaerales
- 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__Betaproteobacteria;D_3__Burkholderiales
- D_0__Bacteria;D_1__Proteobacteria;D_2__Deltaproteobacteria;D_3__Desulfuromonadales
- D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Alteromonadales
- 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__Proteobacteria;D_2__Gammaproteobacteria;D_3__Xanthomonadales
- Unassigned;Other;Other;Other