

- D_0__Archaea;D_1__Euryarchaeota;D_2__Halobacteria;D_3__Halobacteriales
- 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__Actinobacteria;D_2__Actinobacteria;D_3__Propionibacteriales
- D_0__Bacteria;D_1__Actinobacteria;D_2__Nitriliruptoria;D_3__Euzebyales
- D_0__Bacteria;D_1__Bacteroidetes;D_2__Cytophagia;D_3__Order II
- D_0__Bacteria;D_1__Bacteroidetes;D_2__Cytophagia;D_3__Order III
- D_0__Bacteria;D_1__Bacteroidetes;D_2__Sphingobacteriia;D_3__Sphingobacteriales
- D_0__Bacteria;D_1__Cyanobacteria;D_2__Chloroplast;Other
- 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__Planctomycetes;D_2__Phycisphaerae;D_3__Phycisphaerales
- D_0__Bacteria;D_1__Proteobacteria;D_2__Betaproteobacteria;D_3__Burkholderiales
- D_0__Bacteria;D_1__Proteobacteria;D_2__Betaproteobacteria;D_3__Neisseriales
- 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__Proteobacteria;D_2__Gammaproteobacteria;D_3__Pseudomonadales
- D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Salinisphaerales
- D_0__Bacteria;D_1__TM6;D_2__uncultured bacterium;D_3__uncultured bacterium
- Unassigned;Other;Other;Other