

- D\_0\_\_Archaea;D\_1\_\_Euryarchaeota;D\_2\_\_Halobacteria;D\_3\_\_Halobacteriales;D\_4\_\_Halobacteriaceae
- D\_0\_\_Archaea;D\_1\_\_Nanohaloarchaeota;D\_2\_\_uncultured archaeon;D\_3\_\_uncultured archaeon;D\_4\_\_uncultured archaeon
- D\_0\_\_Archaea;D\_1\_\_Nanohaloarchaeota;D\_2\_\_uncultured haloarchaeon;D\_3\_\_uncultured haloarchaeon;D\_4\_\_uncultured haloarchaeon
- D\_0\_\_Archaea;D\_1\_\_Nanohaloarchaeota;Other;Other;Other
- D\_0\_\_Bacteria;D\_1\_\_Actinobacteria;D\_2\_\_Actinobacteria;D\_3\_\_Corynebacteriales;D\_4\_\_Nocardiaceae
- D\_0\_\_Bacteria;D\_1\_\_Bacteroidetes;D\_2\_\_Cytophagia;D\_3\_\_Order II;D\_4\_\_Rhodothermaceae
- D\_0\_\_Bacteria;D\_1\_\_Bacteroidetes;D\_2\_\_Sphingobacteriia;D\_3\_\_Sphingobacteriales;D\_4\_\_Chitinophagaceae
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Chloroplast;D\_3\_\_uncultured Chlorophyta;D\_4\_\_uncultured Chlorophyta
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Chloroplast;Other;Other
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_SubsectionI;D\_4\_\_FamilyI
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_SubsectionII;D\_4\_\_FamilyII
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_SubsectionIII;D\_4\_\_FamilyI
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_SubsectionIV;D\_4\_\_FamilyI
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_uncultured;D\_4\_\_uncultured bacterium
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;D\_3\_\_uncultured;Other
- D\_0\_\_Bacteria;D\_1\_\_Cyanobacteria;D\_2\_\_Cyanobacteria;Other;Other
- D\_0\_\_Bacteria;D\_1\_\_Planctomycetes;D\_2\_\_Phycisphaerae;D\_3\_\_Phycisphaerales;D\_4\_\_Phycisphaeraceae
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Alphaproteobacteria;D\_3\_\_Rhizobiales;D\_4\_\_Methylobacteriaceae
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Alphaproteobacteria;D\_3\_\_Rhodospirillales;D\_4\_\_Rhodospirillaceae
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Alphaproteobacteria;D\_3\_\_Sphingomonadales;D\_4\_\_Sphingomonadaceae
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Deltaproteobacteria;D\_3\_\_Desulfuromonadales;D\_4\_\_GR-WP33-58
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Gammaproteobacteria;D\_3\_\_Chromatiales;D\_4\_\_Ectothiorhodospiraceae
- D\_0\_\_Bacteria;D\_1\_\_Proteobacteria;D\_2\_\_Gammaproteobacteria;D\_3\_\_Oceanospirillales;D\_4\_\_Halomonadaceae
- D\_0\_\_Bacteria;D\_1\_\_TM6;D\_2\_\_uncultured bacterium;D\_3\_\_uncultured bacterium;D\_4\_\_uncultured bacterium
- Unassigned;Other;Other;Other;Other