

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_Archaea;D_1_Thaumarchaeota;D_2_Soil Crenarchaeotic Group(SCG);D_3_uncultured archaeon;D_4_uncultured archaeon
D_0_Bacteria;D_1_Acidobacteria;D_2_Acidobacteria;D_3_Acidobacteriales;D_4_Acidobacteriaceae (Subgroup 1)
D_0_Bacteria;D_1_Acidobacteria;D_2_Acidobacteria;D_3_Subgroup 3;D_4_Unknown Family
D_0_Bacteria;D_1_Acidobacteria;D_2_Acidobacteria;D_3_Subgroup 4;D_4_RB41
D_0_Bacteria;D_1_Acidobacteria;D_2_Acidobacteria;D_3_Subgroup 4;D_4_Unknown Family
D_0_Bacteria;D_1_Actinobacteria;D_2_Acidimicrobia;D_3_Acidimicrobiales;D_4_uncultured
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Actinomycetales;D_4_Actinomycetaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Corynebacteriales;D_4_Corynebacteriaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Corynebacteriales;D_4_Nocardiaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;D_4_Frankiaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;D_4_Geodermatophylaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;D_4_Nakamurellaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;D_4_Sporichthyaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;D_4_uncultured
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Frankiales;Other
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Kineosporiales;D_4_Kineosporiaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Micrococcales;D_4_Intrasporangiaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Micrococcales;D_4_Micrococcaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Propionibacteriales;D_4_Nocardioidaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Propionibacteriales;D_4_Propionibacteriaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Actinobacteria;D_3_Pseudonocardiales;D_4_Pseudonocardiaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Nitriliruptoria;D_3_Euzebyales;D_4_Euzebyaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Rubrobacteria;D_3_Rubrobacteriales;D_4_Rubrobacteriaceae
D_0_Bacteria;D_1_Actinobacteria;D_2_Thermoleophila;D_3_Solirubrobacteriales;D_4_0319-6M6
D_0_Bacteria;D_1_Actinobacteria;D_2_Thermoleophila;D_3_Solirubrobacteriales;D_4_480-2
D_0_Bacteria;D_1_Actinobacteria;D_2_Thermoleophila;D_3_Solirubrobacteriales;D_4_Patulibacteraceae
D_0_Bacteria;D_1_Armatimonadetes;D_2_Armatimonadia;D_3_Armatimonadales;D_4_uncultured bacterium
D_0_Bacteria;D_1_Armatimonadetes;D_2_uncultured bacterium;D_3_uncultured bacterium;D_4_uncultured bacterium
D_0_Bacteria;D_1_Bacteroidetes;D_2_Bacteroidia;D_3_Bacteroidales;D_4_Porphyromonadaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Bacteroidia;D_3_Bacteroidales;D_4_Prevotellaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia;D_3_Cytophagales;D_4_Cytophagaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia;D_3_Cytophagales;D_4_Flammeovirgaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia;D_3_Cytophagales;D_4_MWH-CFBk5
D_0_Bacteria;D_1_Bacteroidetes;D_2_Cytophagia;D_3_Order II;D_4_Rhodothermaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Flavobacteriia;D_3_Flavobacteriales;D_4_Cryomorphaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Flavobacteriia;D_3_Flavobacteriales;D_4_Flavobacteriaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Sphingobacteriia;D_3_Sphingobacteriales;D_4_Chitinophagaceae
D_0_Bacteria;D_1_Bacteroidetes;D_2_Sphingobacteriia;D_3_Sphingobacteriales;D_4_Sphingobacteriaceae
D_0_Bacteria;D_1_Chloroflexi;D_2_Ardenticatenia;D_3_Ardenticatenales;D_4_uncultured bacterium
D_0_Bacteria;D_1_Chloroflexi;D_2_Ktedonobacteria;D_3_C0119;Other
D_0_Bacteria;D_1_Chloroflexi;D_2_Thermomicrobia;D_3_JG30-KF-CM45;D_4_uncultured bacterium
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;D_3_uncultured bacterium;D_4_uncultured bacterium
D_0_Bacteria;D_1_Cyanobacteria;D_2_Chloroplast;D_3_uncultured eukaryote;D_4_uncultured eukaryote
D_0_Bacteria;D_1_Cyanobacteria;D_2_Chloroplast;D_3_uncultured organism;D_4_uncultured organism
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_Deinococcus-Thermus;D_2_Deinococci;D_3_Deinococcales;D_4_Deinococcaceae
D_0_Bacteria;D_1_Deinococcus-Thermus;D_2_Deinococci;D_3_Deinococcales;D_4_Trueperaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Bacillales;D_4_Bacillaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Bacillales;D_4_Family XI
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Bacillales;D_4_Staphylococcaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Lactobacillales;D_4_Aerococcaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Lactobacillales;D_4_Carnobacteriaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Bacilli;D_3_Lactobacillales;D_4_Streptococcaceae
D_0_Bacteria;D_1_Firmicutes;D_2_Clostridia;D_3_Clostridiales;D_4_Family XI
D_0_Bacteria;D_1_Fusobacteria;D_2_Fusobacteriia;D_3_Fusobacteriales;D_4_Fusobacteriaceae
D_0_Bacteria;D_1_Gemmatimonadetes;D_2_Gemmatimonadetes;D_3_AT425-EubC11 terrestrial group;D_4_uncultured bacterium
D_0_Bacteria;D_1_Plantcymycetes;D_2_Phycisphaerae;D_3_Phycisphaerales;D_4_Phycisphaeraceae
D_0_Bacteria;D_1_Plantcymycetes;D_2_Plantcymycetacia;D_3_Plantcymycetales;D_4_Plantcymycetaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Caulobacteriales;D_4_Caulobacteraceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhizobiales;D_4_1174-901-12
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhizobiales;D_4_D05-2
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhizobiales;D_4_JG34-KF-361
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_Rhodobacterales;D_4_Rhodobacteraceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Rhodospirillales;D_4_Acetobacteraceae
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_AKYG937
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Sphingomonadales;D_4_Ellin6055
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Sphingomonadales;D_4_JG34-KF-161
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Sphingomonadales;D_4_Sphingomonadaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Alphaproteobacteria;D_3_Sphingomonadales;Other
D_0_Bacteria;D_1_Proteobacteria;D_2_Betaproteobacteria;D_3_Burkholderiales;D_4_Comamonadaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Betaproteobacteria;D_3_Burkholderiales;D_4_Oxalobacteraceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Betaproteobacteria;D_3_Neisseriales;D_4_Neisseriaceae
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_Deltaproteobacteria;D_3_Myxococcales;D_4_KD3-10
D_0_Bacteria;D_1_Proteobacteria;D_2_Deltaproteobacteria;D_3_Myxococcales;D_4_Polyangiaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria;D_3_Alteromonadales;D_4_Shewanellaceae
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_Enterobacteriales;D_4_Enterobacteriaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria;D_3_Oceanospirillales;D_4_Halomonadaceae
D_0_Bacteria;D_1_Proteobacteria;D_2_Gammaproteobacteria;D_3_Pasteurellales;D_4_Pasteurellaceae
D_0_Bacteria;D_1_TM6;D_2_uncultured bacterium;D_3_uncultured bacterium;D_4_uncultured bacterium
D_0_Bacteria;D_1_Verrucomicrobia;D_2_Spartobacteria;D_3_Chthoniobacteriales;D_4_Chthoniobacteraceae
D_0_Bacteria;D_1_Verrucomicrobia;D_2_Spartobacteria;D_3_Chthoniobacteriales;D_4_DAI01 soil group
Unassigned;Other;Other;Other;Other