



- k__Archaea; p__Euryarchaeota; c__Methanobacteria; o__Methanobacteriales
- k__Archaea; p__Euryarchaeota; c__Thermoplasmata; o__E2
- k__Bacteria; p__Acidobacteria; c__Acidobacteriia; o__Acidobacteriales
- k__Bacteria; p__Acidobacteria; c__DA052; o__Ellin6513
- k__Bacteria; p__Acidobacteria; c__Solibacteres; o__Solibacterales
- k__Bacteria; p__Actinobacteria; c__Acidimicrobiia; o__Acidimicrobiales
- k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
- k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
- k__Bacteria; p__Actinobacteria; c__Coriobacteriia; o__Coriobacteriales
- k__Bacteria; p__Actinobacteria; c__Nitriliruptoria; o__Euzebyales
- k__Bacteria; p__Actinobacteria; c__Thermoleophilia; o__Gaiellales
- k__Bacteria; p__Actinobacteria; c__Thermoleophilia; o__Solirubrobacterales
- k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
- k__Bacteria; p__Bacteroidetes; c__Cytophagia; o__Cytophagales
- k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales
- k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
- k__Bacteria; p__Bacteroidetes; c__[Rhodothermi]; o__[Rhodothermales]
- k__Bacteria; p__Bacteroidetes; c__[Saprospirae]; o__[Saprospirales]
- k__Bacteria; p__Chloroflexi; c__Anaerolineae; o__Anaerolineales
- k__Bacteria; p__Chloroflexi; c__Gitt-GS-136; o__
- k__Bacteria; p__Chloroflexi; c__TK10; o__AKYG885
- k__Bacteria; p__Cyanobacteria; c__4C0d-2; o__YS2
- k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
- k__Bacteria; p__Cyanobacteria; c__Oscillatoriophycideae; o__Oscillatoriales
- k__Bacteria; p__Elusimicrobia; c__Elusimicrobia; o__Elusimicrobiales
- k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
- k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
- k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
- k__Bacteria; p__Firmicutes; c__Bacilli; o__Turicibacterales
- k__Bacteria; p__Firmicutes; c__Clostridia; o__
- k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
- k__Bacteria; p__Firmicutes; c__Clostridia; o__SHA-98
- k__Bacteria; p__Firmicutes; c__Erysipelotrichi; o__Erysipelotrichales
- k__Bacteria; p__Fusobacteria; c__Fusobacteriia; o__Fusobacteriales
- k__Bacteria; p__Gemmatimonadetes; c__Gemmatimonadetes; o__Gemmatimonadales
- k__Bacteria; p__Lentisphaerae; c__[Lentisphaeria]; o__Victivallales
- k__Bacteria; p__Lentisphaerae; c__[Lentisphaeria]; o__Z20
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__BD7-3
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Caulobacterales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__RF32
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacteriales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodospirillales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales
- k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__
- k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
- k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Methylophilales
- k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
- k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Rhodocyclales
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__BPC076
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Bdellovibrionales
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Desulfobacteriales
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Desulfovibrionales
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Desulfuromonadales
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__GMD14H09
- k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Myxococcales
- k__Bacteria; p__Proteobacteria; c__Epsilonproteobacteria; o__Campylobacteriales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Alteromonadales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Cardiobacteriales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Chromatiales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Legionellales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Methylococcales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Oceanospirillales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Vibrionales
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
- k__Bacteria; p__SR1; c__; o__
- k__Bacteria; p__Spirochaetes; c__Spirochaetes; o__Sphaerochaetales
- k__Bacteria; p__Spirochaetes; c__Spirochaetes; o__Spirochaetales
- k__Bacteria; p__Spirochaetes; c__[Brachyspirae]; o__[Brachyspirales]
- k__Bacteria; p__Synergistetes; c__Synergistia; o__Synergistales
- k__Bacteria; p__TM7; c__TM7-3; o__
- k__Bacteria; p__TM7; c__TM7-3; o__CW040
- k__Bacteria; p__Tenericutes; c__Mollicutes; o__Anaeroplasmatales
- k__Bacteria; p__Tenericutes; c__Mollicutes; o__Mycoplasmatales
- k__Bacteria; p__Tenericutes; c__Mollicutes; o__RF39
- k__Bacteria; p__Tenericutes; c__RF3; o__ML615J-28
- k__Bacteria; p__Verrucomicrobia; c__Opitutae; o__[Cerasicoccales]
- k__Bacteria; p__Verrucomicrobia; c__Verruco-5; o__WCHB1-41
- k__Bacteria; p__Verrucomicrobia; c__Verrucomicrobiae; o__Verrucomicrobiales
- k__Bacteria; p__Verrucomicrobia; c__[Spartobacteria]; o__[Chthoniobacteriales]
- k__Bacteria; p__[Thermi]; c__Deinococci; o__Deinococcales