

k__Archaea; p__Euryarchaeota; c__Methanobacteria; o__Methanobacteriales
k__Bacteria; p__; c__; o__
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
k__Bacteria; p__Aquificae; c__Aquificae; o__Aquificales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Cyanobacteria; c__4C0d-2; o__YS2
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__
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__Turicibacteriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Coriobacteriales
k__Bacteria; p__Firmicutes; c__Erysipelotrichi; o__Erysipelotrichales
k__Bacteria; p__Firmicutes; c__RF3; o__
k__Bacteria; p__Firmicutes; c__RF3; o__ML615J-28
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Caulobacterales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__RF32
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales
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__Neisseriales
k__Bacteria; p__Proteobacteria; c__Deltaproteobacteria; o__Desulfovibrionales
k__Bacteria; p__Proteobacteria; c__Epsilonproteobacteria; o__Campylobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Cardiobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
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__Xanthomonadales
k__Bacteria; p__Synergistetes; c__Synergistia; o__Synergistales
k__Bacteria; p__TM7; c__TM7-3; o__
k__Bacteria; p__Tenericutes; c__Mollicutes; o__RF39
k__Bacteria; p__Verrucomicrobia; c__Verrucomicrobiae; o__Verrucomicrobiales