

- k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Actinomycetales;f\_Propionibacteriaceae;g\_Propionibacterium;s\_acnes
- k\_Bacteria;p\_Cyanobacteria;c\_NA;o\_Chroococcales;f\_NA;g\_Gloeothecae;s\_membranacea\_nov\_86.56%
- k\_Bacteria;p\_Cyanobacteria;c\_NA;o\_Nostocales;f\_Nostocaceae;g\_Trichormus;s\_azollae\_nov\_88.37%
- k\_Bacteria;p\_Fusobacteria;c\_Fusobacteriia;o\_Fusobacteriales;f\_Leptotrichiaceae;g\_Leptotrichia;s\_sp\_oral\_taxon\_212
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_NA;f\_NA;g\_Candidatus\_Puniceispirillum;s\_Puniceispirillum\_marinum\_nov\_90.85%
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rhizobiales;f\_Bradyrhizobiaceae;g\_Bradyrhizobium;s\_sp\_str\_LMG10689
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rhizobiales;f\_Methylobacteriaceae;g\_Methylobacterium;s\_zatmanii
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rhodobacterales;f\_Rhodobacteraceae;g\_Paracoccus;s\_kocurii
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rhodospirillales;f\_Rhodospirillaceae;g\_Nisaea;s\_multispecies\_sppn8\_2\_nov\_87.50%
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rickettsiales;f\_Pelagibacteraceae;g\_Pelagibacter;s\_ubique
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Sphingomonadales;f\_Sphingomonadaceae;g\_Sphingomonas;s\_changbaiensis
- k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Sphingomonadales;f\_Sphingomonadaceae;g\_Sphingomonas;s\_paucimobilis
- k\_Bacteria;p\_Proteobacteria;c\_Betaproteobacteria;o\_Burkholderiales;f\_NA;g\_Tepidicella;s\_xavieri
- k\_Bacteria;p\_Proteobacteria;c\_Betaproteobacteria;o\_Burkholderiales;f\_Oxalobacteraceae;g\_Massilia;s\_timonae
- k\_Bacteria;p\_Proteobacteria;c\_Betaproteobacteria;o\_Nitrosomonadales;f\_Nitrosomonadaceae;g\_Nitrosomonas;s\_sp\_AJ224410.1
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pseudomonadales;f\_Moraxellaceae;g\_Acinetobacter;s\_johnsonii
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pseudomonadales;f\_Moraxellaceae;g\_Moraxella;s\_osloensis
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pseudomonadales;f\_Pseudomonadaceae;g\_Pseudomonas;s\_aeruginosa
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pseudomonadales;f\_Pseudomonadaceae;g\_Rugamonas;s\_rubra\_nov\_97.93%