


















































-  undetermined Bacteria
-  Propionibacterium_u
-  Actinobacteria_u_u_u_u
-  Corynebacterium_u
-  Streptococcus pneumoniae
-  Pasteurellaceae_u_u
-  Proteobacteria_u_u_u_u_u_u
-  Bacteroides_u
-  Pseudomonas_u
-  Streptococcaceae prophage_u_u
-  Firmicutes_u_u_u_u_u_u
-  Malassezia globosa
-  Clostridiales_u_u_u_u
-  Actinomyces_u
-  Negativicutes_u_u_u_u_u_u
-  Bacteroidales_u_u_u_u
-  Clostridioides difficile
-  Bacteroidetes_u_u_u_u_u_u_u_u
-  Staphylococcus aureus
-  Bacilli prophage_u_u_u_u_u_u
-  Lactobacillales prophage_u_u_u_u
-  Micrococcales_u_u_u_u_u_u
-  Veillonella_u
-  Propionibacteriaceae_u_u_u_u
-  Porphyromonadaceae_u_u_u_u
-  Bacillales prophage_u_u_u_u_u_u
-  Mycobacterium_u
-  Basidiomycota_u_u_u_u_u_u_u_u
-  Staphylococcus_u
-  Vibrio parahaemolyticus
-  Firmicutes prophage_u_u_u_u_u_u_u_u
-  Pseudomonas syringae group genomsp. 3
-  Streptococcus_u
-  Bacillales_u_u_u_u_u_u_u_u
-  Prevotella_u
-  Escherichia coli
-  Rothia mucilaginosa
-  Porphyromonas_u
-  Corynebacteriales_u_u_u_u_u_u_u_u
-  Bacillus thuringiensis
-  Gammaproteobacteria_u_u_u_u_u_u_u_u
-  Lawsonella clevelandensis
-  Ruminococcaceae_u_u_u_u_u_u_u_u
-  Corynebacterium afermentans
-  Enterobacteriaceae prophage_u_u_u_u_u_u_u_u
-  Deinococcus_u
-  Betaproteobacteria_u_u_u_u_u_u_u_u
-  Clostridia_u_u_u_u_u_u_u_u_u_u
-  Brvibacterium_u
-  Brachybacterium_u