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-  Klebsiella pneumoniae
-  Candida orthopsilosis
-  Streptococcus prophage_u
-  Escherichia coli
-  Lactobacillus paracasei
-  Lactobacillus rhamnosus
-  Pyrenophora teres
-  Malassezia globosa
-  Lactobacillus prophage_u
-  Lactobacillus casei
-  Streptococcus pneumoniae
-  Streptococcus mitis
-  Corynebacterium matruchotii
-  Haemophilus parainfluenzae
-  Pyrenophora tritici-repentis
-  Citrobacter freundii
-  Setosphaeria turcica
-  Haemophilus prophage_u
-  Escherichia prophage_u
-  Pleosporaceae_u_u
-  Pyrenophora_u
-  Enterobacteriaceae prophage_u_u
-  Pleosporales_u_u_u
-  Haemophilus influenzae
-  Stereum hirsutum
-  Pseudomonas syringae group genomsp. 3
-  Pseudomonas brassicacearum
-  Lactobacillus zeae
-  Staphylococcus aureus
-  Aggregatibacter aphrophilus
-  Pseudomonas aeruginosa
-  Pseudomonas prophage_u
-  Streptococcus infantis
-  Pseudomonas fluorescens
-  Pseudomonas sp. CBZ-4
-  Zymoseptoria tritici
-  Ascomycota_u_u_u_u
-  Citrobacter rodentium
-  Pseudomonas chlororaphis
-  Pseudomonas sp. TJI-51
-  Bipolaris_u
-  Pseudomonas extremaustralis
-  Parastagonospora nodorum
-  Pasteurellaceae prophage_u_u
-  Escherichia sp. TW09231
-  Staphylococcus warneri
-  Staphylococcus epidermidis
-  Enterobacter mori
-  Acinetobacter baumannii