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-  Streptococcaceae prophage\_u\_u
-  Malassezia globosa
-  Clostridioides difficile
-  Staphylococcus aureus
-  Bacilli prophage\_u\_u\_u\_u
-  Lactobacillales prophage\_u\_u\_u
-  Bacillales prophage\_u\_u\_u
-  Basidiomycota\_u\_u\_u\_u\_u
-  Firmicutes prophage\_u\_u\_u\_u\_u
-  Pseudomonas syringae group genomsp. 3
-  Bacillus thuringiensis
-  Enterobacteriaceae prophage\_u\_u
-  Ascomycota\_u\_u\_u\_u\_u
-  Pediococcus acidilactici
-  Candida orthopsilosis
-  Gammaproteobacteria prophage\_u\_u\_u\_u\_u
-  Lactococcus garvieae
-  undetermined Eukaryota
-  Yarrowia lipolytica
-  Azospirillum lipoferum
-  Penicillium rubens
-  Paenibacillus larvae
-  Escherichia coli
-  [Haemophilus] parasuis
-  Staphylococcus epidermidis
-  Lactobacillus casei
-  undetermined Prophage
-  Coccidioides immitis
-  Pseudomonas aeruginosa
-  Wallemia mellicola
-  Schizosaccharomyces octosporus
-  Coniosporium apollinis
-  Haemophilus influenzae
-  Clavispora lusitaniae
-  Spizellomyces punctatus
-  Penicillium digitatum
-  Pneumocystis murina
-  Lactobacillaceae prophage\_u\_u
-  Histoplasma capsulatum
-  Penicillium\_u
-  Talaromyces\_u
-  Propionibacterium acnes
-  Streptococcus mitis
-  Streptococcus prophage\_u
-  Salmonella enterica
-  Ustilaginaceae\_u\_u
-  Pyrenophora tritici-repentis
-  Saccharomyces cerevisiae
-  Beauveria bassiana