

k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Actinomycetales;f\_Actinomycetaceae;g\_Actinomyces;s\_lingnae\_[Not\_Validly\_Published]\_HMT\_181  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Actinomycetales;f\_Micrococcaceae;g\_Rothia;s\_mucilaginoso\_HMT\_681  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_aurimucosum\_HMT\_034  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_coyleae\_HMT\_341  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_vitaeruminis  
k\_Bacteria;p\_Actinobacteria;c\_Coriobacteriia;o\_Coriobacteriales;f\_Coriobacteriaceae;g\_Atopobium;s\_parvulum\_HMT\_723  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Bacteroidaceae;g\_Bacteroides;s\_fragilis  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Dysgonamonadaceae;g\_Proteiniphilum;s\_acetatigenes\_nov\_91.822%  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Prevotellaceae;g\_Alloprevotella;s\_sp.\_HMT\_308  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Prevotellaceae;g\_Prevotella;s\_melaninogenica\_HMT\_469  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Prevotellaceae;g\_Prevotella;s\_sp.\_HMT\_313  
k\_Bacteria;p\_Bacteroidetes;c\_Chitinophagia;o\_Chitinophagales;f\_Chitinophagaceae;g\_Vibrionimonas;s\_magnilacihabitans  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Clostridiaceae;g\_Clostridium;s\_multispecies\_spp252\_2  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Clostridiaceae;g\_Clostridium;s\_perfringens  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae\_[XIV];g\_Oribacterium;s\_sinus\_HMT\_457  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae\_[XIVa];g\_Lacnoclostridium;s\_hathewayi  
k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Rhizobiales;f\_Bradyrhizobiaceae;g\_Bradyrhizobium;s\_multispecies\_spp135\_14  
k\_Bacteria;p\_Proteobacteria;c\_Alphaproteobacteria;o\_Sphingomonadales;f\_Sphingomonadaceae;g\_Sphingomonas;s\_changbaiensis  
k\_Bacteria;p\_Synergistetes;c\_Synergistetes\_[C];o\_Synergistetes\_[O];f\_Synergistetes\_[F];g\_Synergistetes\_[G];s\_sp.\_HMT\_B52