

- 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\_mastitidis\_HMT\_328
- k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_mastitidis\_HMT\_328\_nov\_84.541%
- k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_mastitidis\_HMT\_328\_nov\_97.101%
- k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Bacteroidaceae;g\_Bacteroides;s\_fragilis
- k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Tannerellaceae;g\_Parabacteroides;s\_goldsteinii
- k\_Bacteria;p\_Bacteroidetes;c\_Cytophagia;o\_Cytophagales;f\_Cytophagaceae;g\_Cytophaga;s\_xylanolytica\_nov\_83.803%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Acetatifactor;s\_muris\_nov\_94.307%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Lachnoclostridium;s\_pacaense\_nov\_96.782%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Lachnoclostridium;s\_populeti\_nov\_94.059%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_multigenus;s\_multispecies\_sppn2\_2\_nov\_95.297%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae\_[XIVa];g\_Lachnoclostridium;s\_hathewayi\_nov\_95.320%
- k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Oscillospiraceae;g\_Oscillibacter;s\_ruminantium\_nov\_95.567%
- k\_Bacteria;p\_Fusobacteria;c\_Fusobacteriia;o\_Fusobacteriales;f\_Leptotrichiaceae;g\_Leptotrichia;s\_shahii\_HMT\_214
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pasteurellales;f\_Pasteurellaceae;g\_Bisgaardia;s\_hudsonensis\_nov\_86.343%
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pasteurellales;f\_Pasteurellaceae;g\_Haemophilus;s\_pittmaniae\_HMT\_946\_nov\_90.443%
- k\_Bacteria;p\_Proteobacteria;c\_Gammaproteobacteria;o\_Pasteurellales;f\_Pasteurellaceae;g\_Mannheimia;s\_caviae\_nov\_90.443%
- k\_Bacteria;p\_Saccharibacteria\_(TM7);c\_Saccharibacteria\_(TM7)\_[C-1];o\_Saccharibacteria\_(TM7)\_[O-1];f\_Saccharibacteria\_(TM7)\_[F-1];g\_Saccharibacteria\_(TM7)\_[G-1];s\_bacterium\_HMT\_952
- All Other Categories