

- 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\_Porphyromonadaceae;g\_Porphyromonas;s\_pasteri\_HMT\_279
- 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\_Tannerellaceae;g\_Parabacteroides;s\_goldsteinii
- k\_Bacteria;p\_Bacteroidetes;c\_Chitinophagia;o\_Chitinophagales;f\_Chitinophagaceae;g\_Vibrionimonas;s\_magnilacihabitans
- 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\_Lacnoclostridium;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\_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\_Proteobacteria;c\_Gammaproteobacteria;o\_Pasteurellales;f\_Pasteurellaceae;g\_multigenus;s\_multispecies\_sppn96\_3\_nov\_91.375%
- All Other Categories