

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\_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\_Bacteroidia;o\_Bacteroidales;f\_Tannerellaceae;g\_Parabacteroides;s\_goldsteinii\_nov\_97.406%

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\_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\_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\_Pseudomonadales;f\_Pseudomonadaceae;g\_Pseudomonas;s\_resinovorans\_nov\_90.023%