

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\_97.573%  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Bacteroidaceae;g\_Bacteroides;s\_fragilis  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Prevotellaceae;g\_Alloprevotella;s\_sp.\_HMT\_914  
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\_Flavobacteriia;o\_Flavobacteriales;f\_Flavobacteriaceae;g\_Capnocytophaga;s\_gingivalis\_HMT\_337  
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\_Fusobacteriaceae;g\_Fusobacterium;s\_periodonticum\_HMT\_201  
k\_Bacteria;p\_Fusobacteria;c\_Fusobacteriia;o\_Fusobacteriales;f\_Leptotrichiaceae;g\_Leptotrichia;s\_sp.\_HMT\_498  
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\_352