

k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Actinomycetales;f\_Micrococcaceae;g\_Rothia;s\_mucilaginoso\_HMT\_681  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Bifidobacteriales;f\_Bifidobacteriaceae;g\_Bifidobacterium;s\_pseudolongum  
k\_Bacteria;p\_Actinobacteria;c\_Actinobacteria;o\_Corynebacteriales;f\_Corynebacteriaceae;g\_Corynebacterium;s\_macginleyi\_HMT\_050  
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\_Prevotella;s\_melaninogenica\_HMT\_469  
k\_Bacteria;p\_Bacteroidetes;c\_Bacteroidia;o\_Bacteroidales;f\_Prevotellaceae;g\_Prevotella;s\_multispecies\_spp106\_2  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Clostridiaceae;g\_Clostridium;s\_paraputrificum  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Clostridiaceae;g\_Clostridium;s\_perfringens  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Dorea;s\_formicigenerans\_nov\_97.030%  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Faecalicatena;s\_orotica  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Lachnoclostridium;s\_multispecies\_sppn93\_2\_nov\_94.802%  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_Muricomes;s\_intestini  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_multigenus;s\_multispecies\_sppn26\_2\_nov\_97.778%  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae;g\_multigenus;s\_multispecies\_sppn87\_2\_nov\_95.556%  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_Lachnospiraceae\_[XIVa];g\_Lacnoclostridium;s\_hathewayi  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_NA;g\_Flavonifractor;s\_plautii  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_multifamily;g\_Lacnoclostridium;s\_scindens  
k\_Bacteria;p\_Firmicutes;c\_Clostridia;o\_Clostridiales;f\_multifamily;g\_Pseudoflavonifractor;s\_capillosus  
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