

k_Bacteria;p_Actinobacteria;g_Mycolicibacterium;s_Mycolicibacterium mucogenicum;x_unknown;y_unknown
k_Bacteria;p_Actinobacteria;g_unknown;s_Micrococcales bacterium UBA3913;x_unknown;y_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides caccae;x_unknown;y_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides gallinarum;x_unknown;y_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides ovatus;x_unknown;y_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides vulgatus;x_unknown;y_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides xylanisolvens;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Bacillus;s_Bacillus subtilis;x_Bacillus subtilis subsp. natto;y_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_Blautia obeum;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_[Ruminococcus] torques;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_uncultured Blautia sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Clostridium;s_uncultured Clostridium sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Clostridium;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus catus;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus eutactus;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus phoceensis;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Dialister;s_uncultured Dialister sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Dorea;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Drancourtella;s_Drancourtella massiliensis;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Eisenbergiella;s_Eisenbergiella massiliensis;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Eisenbergiella;s_Eisenbergiella tayi;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Erysipelatoclostridium;s_[Clostridium] innocuum;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Eubacterium;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Lachnoclostridium;s_[Clostridium] aldenense;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Lachnoclostridium;s_[Clostridium] bolteae;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Lachnoclostridium;s_[Clostridium] symbiosum;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Lactobacillus;s_Lactobacillus crispatus;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Marvinbryantia;s_uncultured Marvinbryantia sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Massilioclostridium;s_Massilioclostridium coli;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Oscillibacter;s_Oscillibacter sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Oscillibacter;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Roseburia;s_uncultured Roseburia sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Roseburia;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_Ruminococcus bromii;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_Ruminococcus lactaris;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_uncultured Ruminococcus sp.;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_Tyzzereella;s_Tyzzereella nexilis;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Firmicutes bacterium CAG:124;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Firmicutes bacterium CAG:194;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Firmicutes bacterium CAG:424;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Ruminococcaceae bacterium;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Clostridiales bacterium;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Lachnospiraceae bacterium;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Ruminococcaceae bacterium;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Acetobacter;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Cupriavidus;s_Cupriavidus metallidurans;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Delftia;s_Delftia tsuruhatensis;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Ralstonia;s_Ralstonia pickettii;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Tepidimonas;s_Tepidimonas fonticaldi;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Xanthomonas;s_Xanthomonas campestris;x_unknown;y_unknown