

k_Bacteria;p_Actinobacteria;g_Mycolicibacterium;s_Mycolicibacterium mucogenicum;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Actinobacteria;g_unknown;s_Micrococcales bacterium UBA3913;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides caccae;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides gallinarum;x_unknown;y_unknown;z_Bacteroides gallinarum DSM 18171 = JCM 13658
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides ovatus;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides vulgatus;x_unknown;y_unknown;z_Bacteroides vulgatus ATCC 8482
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides vulgatus;x_unknown;y_unknown;z_Bacteroides vulgatus CL09T03C04
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides vulgatus;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides xylinisolvans;x_unknown;y_unknown;z_Bacteroides xylinisolvans XB1A
k_Bacteria;p_Bacteroidetes;g_Bacteroides;s_Bacteroides xylinisolvans;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Bacillus;s_Bacillus subtilis;x_Bacillus subtilis subsp. natto;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_Blautia obicum;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_[Ruminococcus] torques;x_unknown;y_unknown;z_[Ruminococcus] torques ATCC 27756
k_Bacteria;p_Firmicutes;g_Blautia;s_uncultured Blautia sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Blautia;s_unknown;x_unknown;y_unknown;z_AF19-10LB
k_Bacteria;p_Firmicutes;g_Blautia;s_unknown;x_unknown;y_unknown;z_KGMB0111
k_Bacteria;p_Firmicutes;g_Blautia;s_unknown;x_unknown;y_unknown;z_Marseille-P3087
k_Bacteria;p_Firmicutes;g_Blautia;s_unknown;x_unknown;y_unknown;z_N6H1-15
k_Bacteria;p_Firmicutes;g_Clostridium;s_uncultured Clostridium sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Clostridium;s_unknown;x_unknown;y_unknown;z_AT4
k_Bacteria;p_Firmicutes;g_Clostridium;s_unknown;x_unknown;y_unknown;z_CAG:167
k_Bacteria;p_Firmicutes;g_Clostridium;s_unknown;x_unknown;y_unknown;z_CAG:221
k_Bacteria;p_Firmicutes;g_Clostridium;s_unknown;x_unknown;y_unknown;z_CAG:253
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus catus;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus eutactus;x_unknown;y_unknown;z_Coprococcus eutactus ATCC 27759
k_Bacteria;p_Firmicutes;g_Coprococcus;s_Coprococcus phoenensis;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Dialister;s_uncultured Dialister sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Dorea;s_unknown;x_unknown;y_unknown;z_AM10-31
k_Bacteria;p_Firmicutes;g_Drancourtella;s_Drancourtella massiliensis;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Eisenbergiella;s_Eisenbergiella massiliensis;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Eisenbergiella;s_Eisenbergiella tayi;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Erysipelatoclostridium;s_[Clostridium] innocuum;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Eubacterium;s_unknown;x_unknown;y_unknown;z_CAG:38
k_Bacteria;p_Firmicutes;g_Lachnosporaceae;s_[Clostridium] aldenense;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Lachnosporaceae;s_[Clostridium] bolteae;x_unknown;y_unknown;z_[Clostridium] bolteae 90A9
k_Bacteria;p_Firmicutes;g_Lachnosporaceae;s_[Clostridium] bolteae;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Lachnosporaceae;s_[Clostridium] symbiosum;x_unknown;y_unknown;z_[Clostridium] symbiosum WAL-14163
k_Bacteria;p_Firmicutes;g_Lactobacillus;s_Lactobacillus crispatus;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Marvinbryantia;s_uncultured Marvinbryantia sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Massilioclostridium;s_Massilioclostridium colii;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Oscillibacter;s_Oscillibacter sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Oscillibacter;s_Oscillibacter;s_unknown;x_unknown;y_unknown;z_57_20
k_Bacteria;p_Firmicutes;g_Oscillibacter;s_unknown;x_unknown;y_unknown;z_CAG:241_62_21
k_Bacteria;p_Firmicutes;g_Roseburia;s_uncultured Roseburia sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Roseburia;s_unknown;x_unknown;y_unknown;z_CAG:10041_57
k_Bacteria;p_Firmicutes;g_Roseburia;s_unknown;x_unknown;y_unknown;z_CAG:182
k_Bacteria;p_Firmicutes;g_Roseburia;s_unknown;x_unknown;y_unknown;z_CAG:197
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_Ruminococcus bromii;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_Ruminococcus lactaris;x_unknown;y_unknown;z_Ruminococcus lactaris CC59_002D
k_Bacteria;p_Firmicutes;g_Ruminococcus;s_uncultured Ruminococcus sp.;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Tyzzerella;s_Tyzzerella nexilis;x_unknown;y_unknown;z_Tyzzerella nexilis DSM 1787
k_Bacteria;p_Firmicutes;g_unknown;s_Firmicutes bacterium CAG:124;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Firmicutes bacterium CAG:424;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Lachnospiraceae;bacterium;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_Ruminococcaceae;bacterium;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Clostridiales bacterium;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Lachnospiraceae bacterium;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_uncultured Ruminococcaceae bacterium;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Proteobacteria;g_Aacetobacter;s_unknown;x_unknown;y_unknown;z_46_36
k_Bacteria;p_Proteobacteria;g_Cupriavidus;s_Cupriavidus metallidurans;x_unknown;y_unknown;z_Cupriavidus metallidurans CH34
k_Bacteria;p_Proteobacteria;g_Cupriavidus;s_Cupriavidus metallidurans;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Proteobacteria;g_Delftia;s_Delftia tsuruhatensis;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Proteobacteria;g_Ralstonia;s_Ralstonia picketti;x_unknown;y_unknown;z_Ralstonia picketti 12D
k_Bacteria;p_Proteobacteria;g_Ralstonia;s_Ralstonia picketti;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Proteobacteria;g_Tepidimonas;s_Tepidimonas fonticaldi;x_unknown;y_unknown;z_unknown
k_Bacteria;p_Proteobacteria;g_Xanthomonas;s_Xanthomonas campestris;x_unknown;y_unknown;z_ATCC 33913