

k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium GAM79;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium OF09-33XD;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium OM04-12BH;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium UBA7598;x_unknown;y_unknown

k_Bacteria;p_Firmicutes;g_unknown;s_Ruminococcaceae bacterium AM07-15;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Ruminococcaceae bacterium D16;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Ruminococcaceae bacterium D5;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_Ruminococcaceae bacterium;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_[Bacteroides] pectinophilus;x_unknown;y_unknown

k_Bacteria;p_Firmicutes;g_unknown;s_Lachnospiraceae bacterium;x_unknown;y_unknown

k_Bacteria;p_Firmicutes;g_unknown;s_[Clostridium] leptum;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_[Eubacterium] rectale;x_unknown;y_unknown
k_Bacteria;p_Firmicutes;g_unknown;s_[Eubacterium] siraeum;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 Erysipelotrichaceae 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_Lentisphaerae;g_unknown;s_Lentisphaeria bacterium;x_unknown;y_unknown
k_Bacteria;p_Lentisphaerae;g_unknown;s_Victivallales bacterium CCUG 44730;x_unknown;y_unknown

k_Bacteria;p_Proteobacteria;g_Acetobacter;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Acinetobacter;s_unknown;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Bilophila;s_Bilophila wadsworthia;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Desulfovibrio;s_Desulfovibrio desulfuricans;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Desulfovibrio;s_Desulfovibrio fairfieldensis;x_unknown;y_unknown
k_Bacteria;p_Proteobacteria;g_Desulfovibrio;s_uncultured Desulfovibrio sp.;x_unknown;y_unknown

k_Bacteria;p_Proteobacteria;g_Escherichia;s_Escherichia coli;x_unknown;y_unknown
 k_Bacteria;p_Proteobacteria;g_Klebsiella;s_Klebsiella oxytoca;x_unknown;y_unknown
 k_Bacteria;p_Proteobacteria;g_Oxalobacter;s_Oxalobacter formigenes;x_unknown;y_unknown

k_Bacteria;p_Tenericutes;g_Mycoplasma;s_unknown;x_unknown;y_unknown

 $k_Bacteria; p_unknown; g_unknown; s_bacterium \ LF-3; x_unknown; y_unknown$

k_Bacteria;p_unknown;g_unknown;s_uncultured bacterium;x_unknown;y_unknown

k Bacteria;p unknown;g unknown;s bacterium;x unknown;y unknown

 $k_Bacteria; p_Proteobacteria; g_Parasutterella; s_Parasutterella excrementihominis; x_unknown; y_unknown and protection of the protectio$

k_Bacteria;p_Proteobacteria;g_Sutterella;s_Sutterella wadsworthensis;x_unknown;y_unknown k_Bacteria;p_Proteobacteria;g_Sutterella;s_uncultured Sutterella sp.;x_unknown;y_unknown k_Bacteria;p_Proteobacteria;g_unknown;s_Proteobacteria bacterium CAG:495;x_unknown;y_unknown k_Bacteria;p_Proteobacteria;g_unknown;s_uncultured Rhodospirillaceae bacterium;x_unknown;y_unknown

k_Bacteria;p_Tenericutes;g_unknown;s_Acholeplasmatales bacterium;x_unknown;y_unknown
k_Bacteria;p_Tenericutes;g_unknown;s_uncultured Mollicutes bacterium;x_unknown;y_unknown
k_Bacteria;p_Tenericutes;g_unknown;s_uncultured Tenericutes bacterium;x_unknown;y_unknown
k_Bacteria;p_Verrucomicrobia;g_Akkermansia;s_Akkermansia muciniphila;x_unknown;y_unknown