

Taxonomy

< 15 % abund.

- p_Actinobacteriota;c_Actinobacteria;o_Bifidobacteriales;f_Bifidobacteriaceae;g_Alloscardovia
 p_Actinobacteriota;c_Actinobacteria;o_Bifidobacteriales;f_Bifidobacteriaceae;g_Bifidobacterium
 p_Actinobacteriota;c_Actinobacteria;o_Bifidobacteriales;f_Bifidobacteriaceae;g_Gardnerella
- p_Actinobacteriota;c_Actinobacteria;o_Corynebacteriales;f_Corynebacteriaceae;g_Corynebacterium
- p Actinobacteriota;c Coriobacteriia;o Coriobacteriales;f Atopobiaceae;g Atopobium
- p_Bacteroidota;c_Bacteroidia;o_Bacteroidales;f_Porphyromonadaceae;g_Porphyromonas
- p_Bacteroidota;c_Bacteroidia;o_Bacteroidales;f_Prevotellaceae;g_Prevotella
- p_Bacteroidota;c_Bacteroidia;o_Flavobacteriales;f_Flavobacteriaceae;g_Flavobacterium
- p_Firmicutes;c_Bacilli;o_Lactobacillales;f_Lactobacillaceae;g_Lactobacillus
- p_Firmicutes;c_Bacilli;o_Lactobacillales;f_Streptococcaceae;g_Streptococcus

- p_Firmicutes;c_Bacilli;o_Staphylococcales;f_Gemellaceae;g_Gemella
- p_Firmicutes;c_Bacilli;o_Staphylococcales;f_Staphylococcaceae;g_Staphylococcus
- p_Firmicutes;c_Clostridia;o_Peptostreptococcales-Tissierellales;f_Family XI;g_Anaerococcus
- p_Firmicutes;c_Clostridia;o_Peptostreptococcales-Tissierellales;f_Family XI;g_Finegoldia
- p_Firmicutes;c_Negativicutes;o_Veillonellales-Selenomonadales;f_Veillonellaceae;g_Dialister
- p Fusobacteriota;c Fusobacteriia;o Fusobacteriales;f Fusobacteriaceae;g Fusobacterium
- p_Fusobacteriota;c_Fusobacteriia;o_Fusobacteriales;f_Leptotrichiaceae;g_Sneathia

- p_Proteobacteria;c_Gammaproteobacteria;o_Burkholderiales;f_Neisseriaceae;g_Neisseria
 - $\verb|p_Proteobacteria|; \verb|c_Gamma| proteobacteria|; \verb|o_Enterobacteria|; \verb|f_Enterobacteria|; \verb|c_Gamma| proteobacteria|; \verb|o_Enterobacteria|; \verb|d_Enterobacteria|; \|d_Enterobacteria|; \|d_Enterobact$

p_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales;f_Beijerinckiaceae;g_Methylobacterium-Methylorubrum

taxonomic level: genus