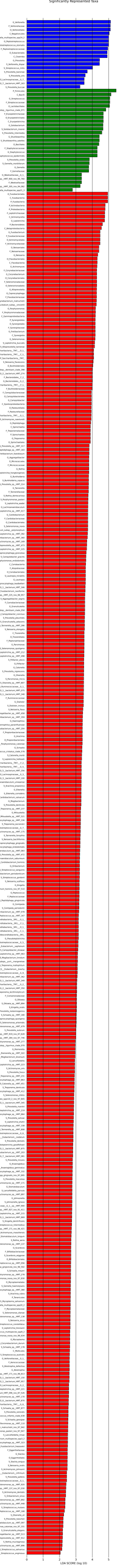
Significantly Represented Taxa
G_Verticinella
F_Methylococcaceae
F_Methylococcaceae
C_Megaphoto-phaga
alpha_multigenomic_taxa20_2
G_Mycobacteriaceae
Staphylococcus_pasteuri
F_Peptostreptococcaceae
O_Eubacteriales
C_Clostridia
G_Pleurofella
G_Lactococcus_piscium
G_Streptococcus_iniae
G_Streptococcus_parauberis
G_Pleurofella_parvum
G_Pleurofella_pleio
G_Lactococcaceae_G_2
C_Saccharomyces_cerevisiae
G_Pleurofella_bacculum
F_Clostridiaceae
C_Bacilli
G_Streptococcus
F_Streptococcaceae
F_Streptococcaceae
alpha_Proteobacteria_unclassified
F_Erysipelotrichaceae
C_Erysipelotrichia
C_Erysipelotrichia
G_Solobacterium
G_Solobacterium_moorei
G_Pleurofella_intermedia
G_Brachyspira
F_Streptococcaceae
F_Streptococcaceae
G_Bacillus
F_Staphylococcaceae
G_Staphylococcus
F_Staphylococcaceae
G_Streptococcus_sanguinis
G_Pleurofella_viridis
G_Gemella_morbillorum
G_Gemella
F_Carnobacteriaceae
G_Weissella_viridescens
F_Methylococcaceae
F_Methylococcaceae
alpha_multigenomic_taxa21_2
F_Fusobacteriaceae
F_Fusobacteria
F_Fusobacteria
F_Fusobacteria
F_Methylococcaceae
F_Methylococcaceae
F_Methylococcaceae
G_Neisseriaceae
G_Neisseria
G_Neisseria
O_Flavobacteriales
C_Flavobacteria
G_Aeromonas
F_Flavobacteriaceae
F_Flavobacteriaceae
F_Flavobacteriaceae
F_Flavobacteriaceae
G_Aeromonas
F_Flavobacteriaceae
F_Flavobacteriaceae
F_Flavobacteriaceae
F_Flavobacteriaceae