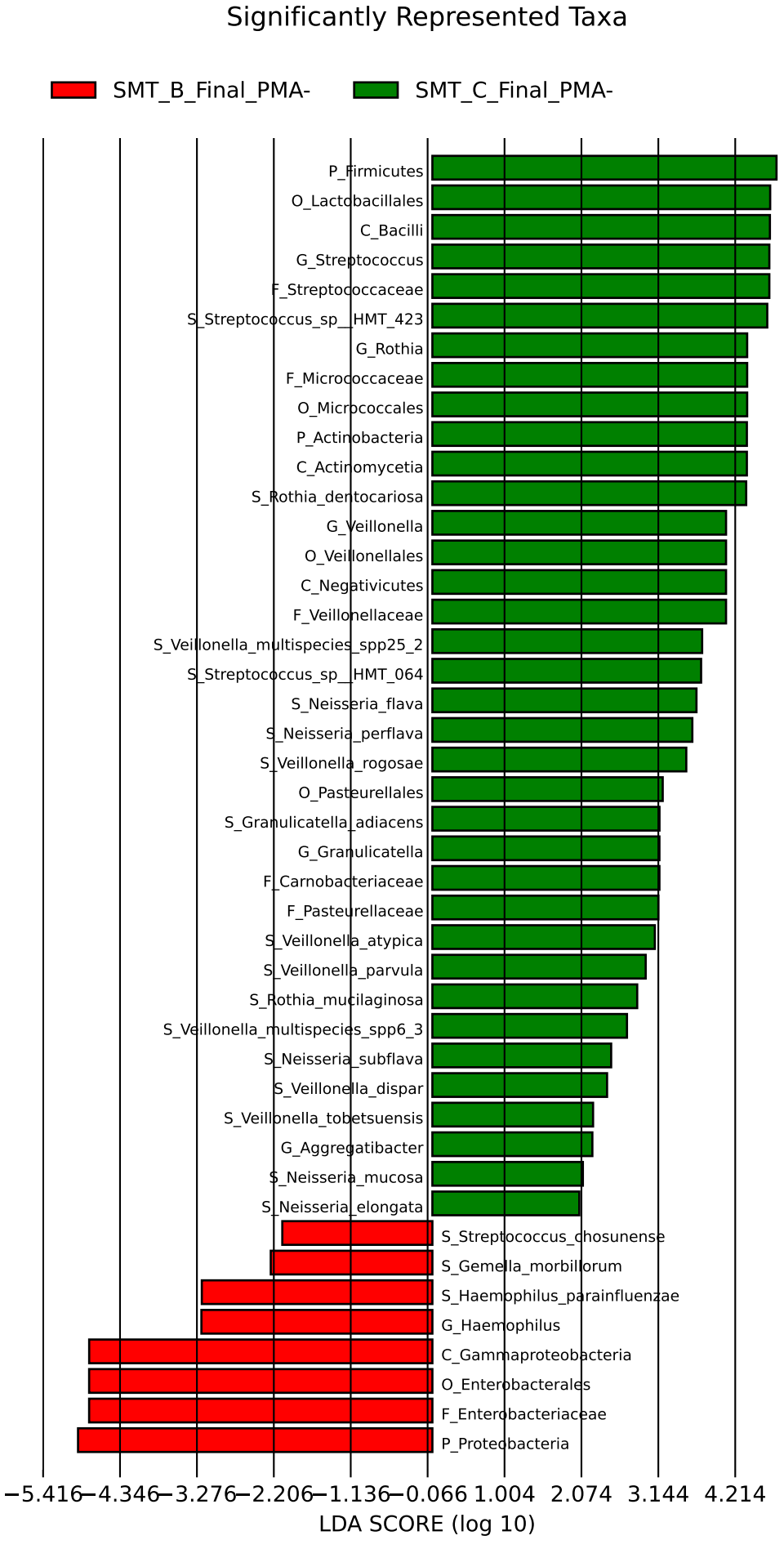
Significantly Represented Taxa
SMT_B_Final_PMA-
SMT_C_Final_PMA-
P_Firmicutes
O_Lactobacillales
C_Bacilli
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
G_Rothia
F_Micrococcaceae
O_Micrococcales
P_Actinobacteria
C_Actinomycetia
S_Rothia_dentocariosa
G_Veillonella
O_Veillonellales
C_Negativicutes
F_Veillonellaceae
S_Veillonella_multispecies_spp25_2
S_Streptococcus_sp__HMT_064
S_Neisseria_flava
S_Neisseria_perflava
S_Veillonella_rogosae
O_Pasteurellales
S_Granulicatella_adiacens
G_Granulicatella
F_Carnobacteriaceae
F_Pasteurellaceae
S_Veillonella_atypica
S_Veillonella_parvula
S_Rothia_mucilaginosa
S_Veillonella_multispecies_spp6_3
S_Neisseria_subflava
S_Veillonella_dispar
S_Veillonella_tobetsuensis
G_Aggregatibacter
S_Neisseria_mucosa
S_Neisseria_elongata
S_Streptococcus_chosunense
S_Gemella_morbillorum
S_Haemophilus_parainfluenzae
G_Haemophilus
C_Gammaproteobacteria
O_Enterobacterales
F_Enterobacteriaceae
P_Proteobacteria
−5.416
−4.346
−3.276
−2.206
−1.136
−0.066
1.004
2.074
3.144
4.214
LDA SCORE (log 10)