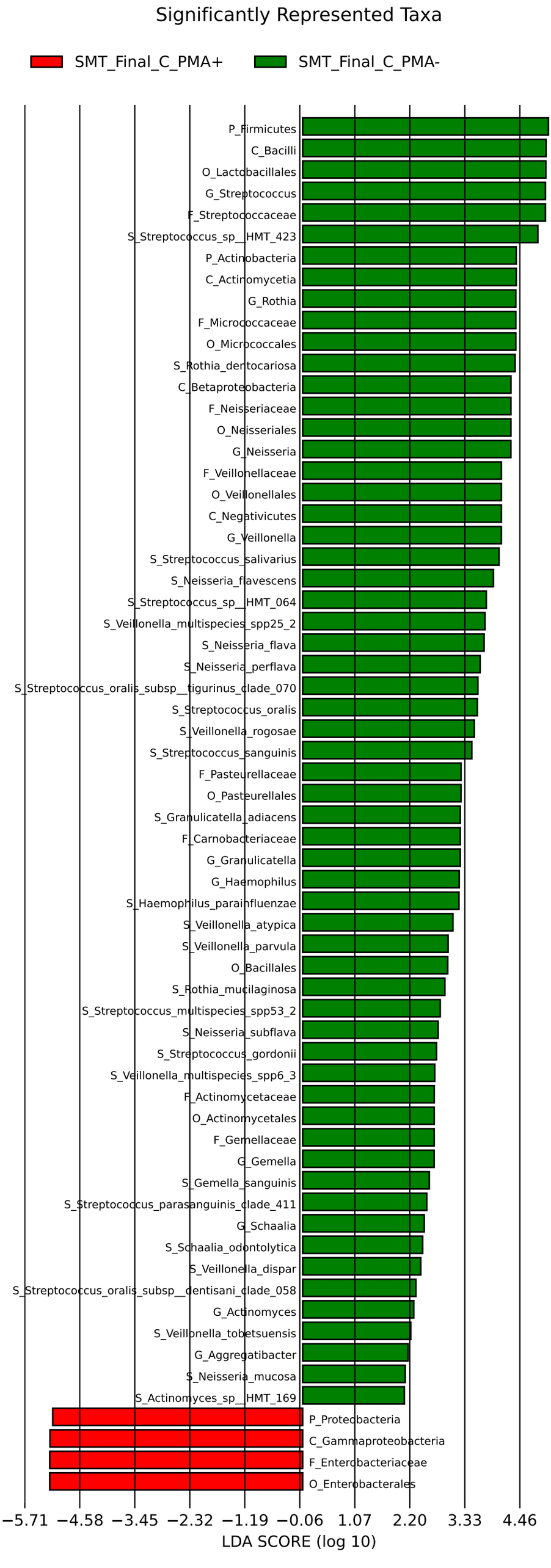
Significantly Represented Taxa
SMT_Final_C_PMA+
SMT_Final_C_PMA-
P_Firmicutes
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
P_Actinobacteria
C_Actinomycetia
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Rothia_dentocariosa
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
G_Neisseria
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
G_Veillonella
S_Streptococcus_salivarius
S_Neisseria_flavescens
S_Streptococcus_sp__HMT_064
S_Veillonella_multispecies_spp25_2
S_Neisseria_flava
S_Neisseria_perflava
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_oralis
S_Veillonella_rogosae
S_Streptococcus_sanguinis
F_Pasteurellaceae
O_Pasteurellales
S_Granulicatella_adiacens
F_Carnobacteriaceae
G_Granulicatella
G_Haemophilus
S_Haemophilus_parainfluenzae
S_Veillonella_atypica
S_Veillonella_parvula
O_Bacillales
S_Rothia_mucilaginosa
S_Streptococcus_multispecies_spp53_2
S_Neisseria_subflava
S_Streptococcus_gordonii
S_Veillonella_multispecies_spp6_3
F_Actinomycetaceae
O_Actinomycetales
F_Gemellaceae
G_Gemella
S_Gemella_sanguinis
S_Streptococcus_parasanguinis_clade_411
G_Schaalia
S_Schaalia_odontolytica
S_Veillonella_dispar
S_Streptococcus_oralis_subsp__dentisani_clade_058
G_Actinomyces
S_Veillonella_tobetsuensis
G_Aggregatibacter
S_Neisseria_mucosa
S_Actinomyces_sp__HMT_169
P_Proteobacteria
C_Gammaproteobacteria
F_Enterobacteriaceae
O_Enterobacterales
−5.71
−4.58
−3.45
−2.32
−1.19
−0.06
1.07
2.20
3.33
4.46
LDA SCORE (log 10)