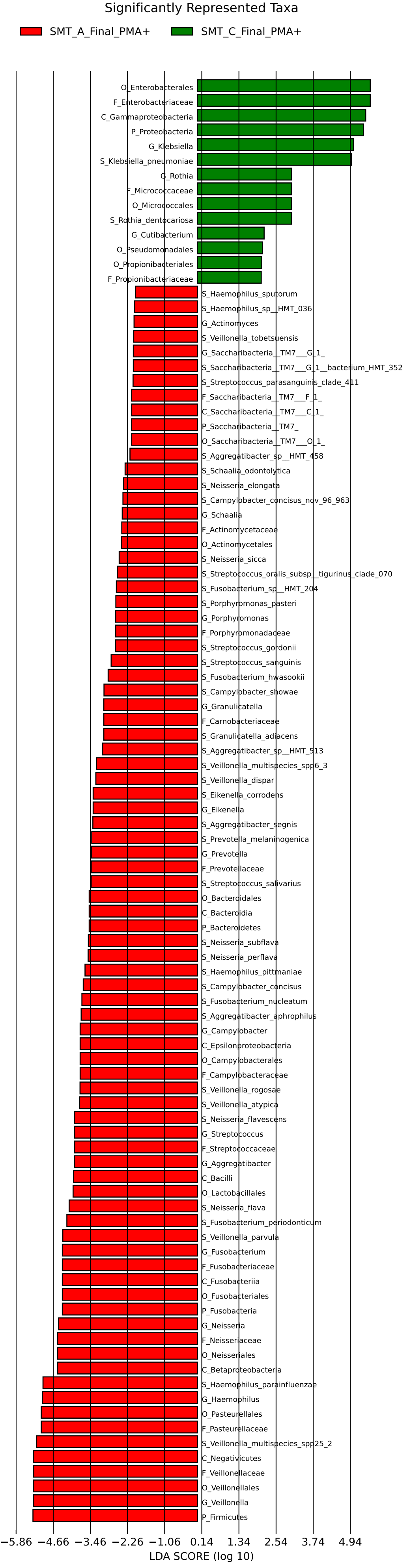
Significantly Represented Taxa
SMT_A_Final_PMA+
SMT_C_Final_PMA+
O_Enterobacterales
F_Enterobacteriaceae
C_Gammaproteobacteria
P_Proteobacteria
G_Klebsiella
S_Klebsiella_pneumoniae
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Rothia_dentocariosa
G_Cutibacterium
O_Pseudomonadales
O_Propionibacteriales
F_Propionibacteriaceae
S_Haemophilus_sputorum
S_Haemophilus_sp__HMT_036
G_Actinomyces
S_Veillonella_tobetsuensis
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
S_Streptococcus_parasanguinis_clade_411
F_Saccharibacteria__TM7___F_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
O_Saccharibacteria__TM7___O_1_
S_Aggregatibacter_sp__HMT_458
S_Schaalia_odontolytica
S_Neisseria_elongata
S_Campylobacter_concisus_nov_96_963
G_Schaalia
F_Actinomycetaceae
O_Actinomycetales
S_Neisseria_sicca
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Fusobacterium_sp__HMT_204
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Streptococcus_gordonii
S_Streptococcus_sanguinis
S_Fusobacterium_hwasookii
S_Campylobacter_showae
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Aggregatibacter_sp__HMT_513
S_Veillonella_multispecies_spp6_3
S_Veillonella_dispar
S_Eikenella_corrodens
G_Eikenella
S_Aggregatibacter_segnis
S_Prevotella_melaninogenica
G_Prevotella
F_Prevotellaceae
S_Streptococcus_salivarius
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Neisseria_subflava
S_Neisseria_perflava
S_Haemophilus_pittmaniae
S_Campylobacter_concisus
S_Fusobacterium_nucleatum
S_Aggregatibacter_aphrophilus
G_Campylobacter
C_Epsilonproteobacteria
O_Campylobacterales
F_Campylobacteraceae
S_Veillonella_rogosae
S_Veillonella_atypica
S_Neisseria_flavescens
G_Streptococcus
F_Streptococcaceae
G_Aggregatibacter
C_Bacilli
O_Lactobacillales
S_Neisseria_flava
S_Fusobacterium_periodonticum
S_Veillonella_parvula
G_Fusobacterium
F_Fusobacteriaceae
C_Fusobacteriia
O_Fusobacteriales
P_Fusobacteria
G_Neisseria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
S_Haemophilus_parainfluenzae
G_Haemophilus
O_Pasteurellales
F_Pasteurellaceae
S_Veillonella_multispecies_spp25_2
C_Negativicutes
F_Veillonellaceae
O_Veillonellales
G_Veillonella
P_Firmicutes
−5.86
−4.66
−3.46
−2.26
−1.06
0.14
1.34
2.54
3.74
4.94
LDA SCORE (log 10)