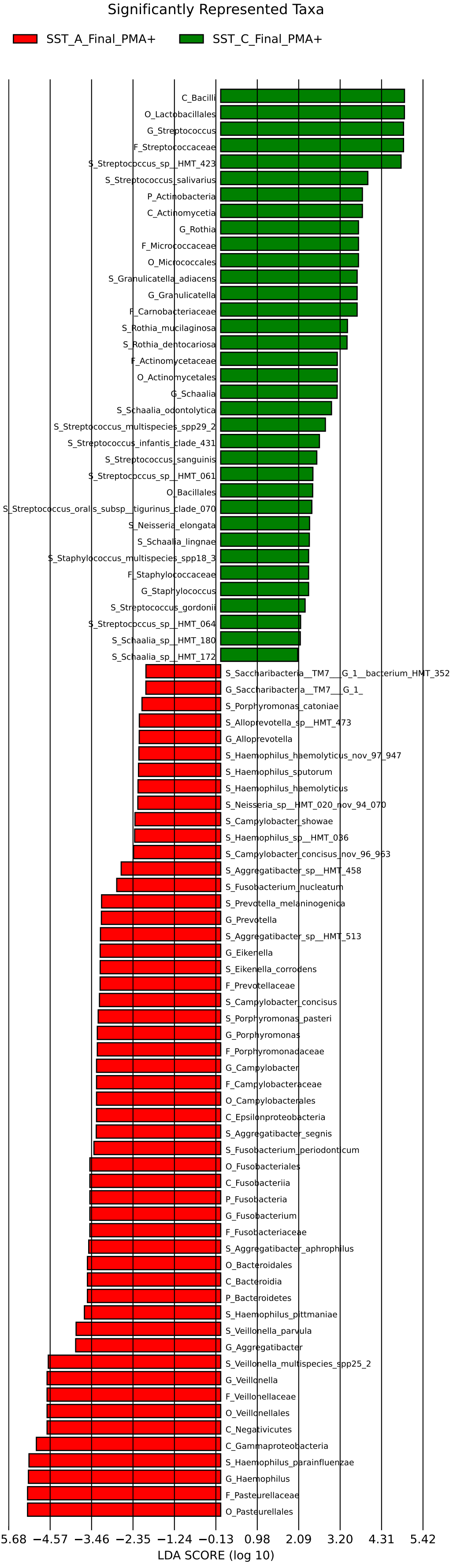
Significantly Represented Taxa
SST_A_Final_PMA+
SST_C_Final_PMA+
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
S_Streptococcus_salivarius
P_Actinobacteria
C_Actinomycetia
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Granulicatella_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_Rothia_mucilaginosa
S_Rothia_dentocariosa
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_Schaalia_odontolytica
S_Streptococcus_multispecies_spp29_2
S_Streptococcus_infantis_clade_431
S_Streptococcus_sanguinis
S_Streptococcus_sp__HMT_061
O_Bacillales
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Neisseria_elongata
S_Schaalia_lingnae
S_Staphylococcus_multispecies_spp18_3
F_Staphylococcaceae
G_Staphylococcus
S_Streptococcus_gordonii
S_Streptococcus_sp__HMT_064
S_Schaalia_sp__HMT_180
S_Schaalia_sp__HMT_172
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
G_Saccharibacteria__TM7___G_1_
S_Porphyromonas_catoniae
S_Alloprevotella_sp__HMT_473
G_Alloprevotella
S_Haemophilus_haemolyticus_nov_97_947
S_Haemophilus_sputorum
S_Haemophilus_haemolyticus
S_Neisseria_sp__HMT_020_nov_94_070
S_Campylobacter_showae
S_Haemophilus_sp__HMT_036
S_Campylobacter_concisus_nov_96_963
S_Aggregatibacter_sp__HMT_458
S_Fusobacterium_nucleatum
S_Prevotella_melaninogenica
G_Prevotella
S_Aggregatibacter_sp__HMT_513
G_Eikenella
S_Eikenella_corrodens
F_Prevotellaceae
S_Campylobacter_concisus
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Aggregatibacter_segnis
S_Fusobacterium_periodonticum
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Aggregatibacter_aphrophilus
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Haemophilus_pittmaniae
S_Veillonella_parvula
G_Aggregatibacter
S_Veillonella_multispecies_spp25_2
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
C_Gammaproteobacteria
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
−5.68
−4.57
−3.46
−2.35
−1.24
−0.13
0.98
2.09
3.20
4.31
5.42
LDA SCORE (log 10)