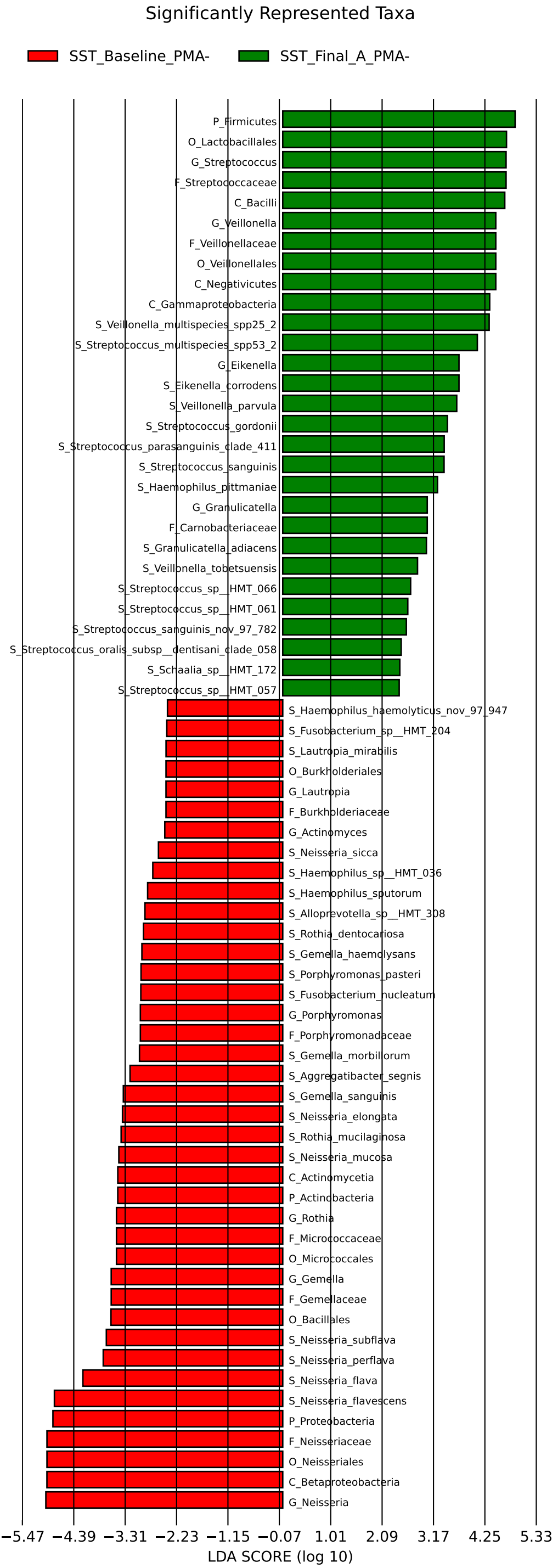
Significantly Represented Taxa
SST_Baseline_PMA-
SST_Final_A_PMA-
P_Firmicutes
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
C_Bacilli
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
C_Gammaproteobacteria
S_Veillonella_multispecies_spp25_2
S_Streptococcus_multispecies_spp53_2
G_Eikenella
S_Eikenella_corrodens
S_Veillonella_parvula
S_Streptococcus_gordonii
S_Streptococcus_parasanguinis_clade_411
S_Streptococcus_sanguinis
S_Haemophilus_pittmaniae
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Veillonella_tobetsuensis
S_Streptococcus_sp__HMT_066
S_Streptococcus_sp__HMT_061
S_Streptococcus_sanguinis_nov_97_782
S_Streptococcus_oralis_subsp__dentisani_clade_058
S_Schaalia_sp__HMT_172
S_Streptococcus_sp__HMT_057
S_Haemophilus_haemolyticus_nov_97_947
S_Fusobacterium_sp__HMT_204
S_Lautropia_mirabilis
O_Burkholderiales
G_Lautropia
F_Burkholderiaceae
G_Actinomyces
S_Neisseria_sicca
S_Haemophilus_sp__HMT_036
S_Haemophilus_sputorum
S_Alloprevotella_sp__HMT_308
S_Rothia_dentocariosa
S_Gemella_haemolysans
S_Porphyromonas_pasteri
S_Fusobacterium_nucleatum
G_Porphyromonas
F_Porphyromonadaceae
S_Gemella_morbillorum
S_Aggregatibacter_segnis
S_Gemella_sanguinis
S_Neisseria_elongata
S_Rothia_mucilaginosa
S_Neisseria_mucosa
C_Actinomycetia
P_Actinobacteria
G_Rothia
F_Micrococcaceae
O_Micrococcales
G_Gemella
F_Gemellaceae
O_Bacillales
S_Neisseria_subflava
S_Neisseria_perflava
S_Neisseria_flava
S_Neisseria_flavescens
P_Proteobacteria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Neisseria
−5.47
−4.39
−3.31
−2.23
−1.15
−0.07
1.01
2.09
3.17
4.25
5.33
LDA SCORE (log 10)