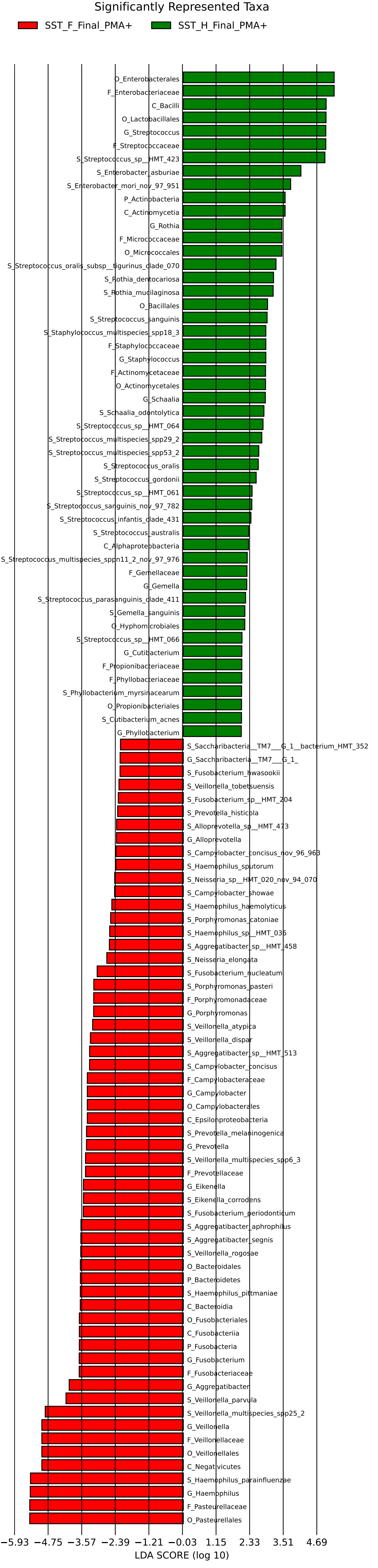
Significantly Represented Taxa
SST_F_Final_PMA+
SST_H_Final_PMA+
O_Enterobacterales
F_Enterobacteriaceae
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
S_Enterobacter_asburiae
S_Enterobacter_mori_nov_97_951
P_Actinobacteria
C_Actinomycetia
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Rothia_dentocariosa
S_Rothia_mucilaginosa
O_Bacillales
S_Streptococcus_sanguinis
S_Staphylococcus_multispecies_spp18_3
F_Staphylococcaceae
G_Staphylococcus
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_Schaalia_odontolytica
S_Streptococcus_sp__HMT_064
S_Streptococcus_multispecies_spp29_2
S_Streptococcus_multispecies_spp53_2
S_Streptococcus_oralis
S_Streptococcus_gordonii
S_Streptococcus_sp__HMT_061
S_Streptococcus_sanguinis_nov_97_782
S_Streptococcus_infantis_clade_431
S_Streptococcus_australis
C_Alphaproteobacteria
S_Streptococcus_multispecies_sppn11_2_nov_97_976
F_Gemellaceae
G_Gemella
S_Streptococcus_parasanguinis_clade_411
S_Gemella_sanguinis
O_Hyphomicrobiales
S_Streptococcus_sp__HMT_066
G_Cutibacterium
F_Propionibacteriaceae
F_Phyllobacteriaceae
S_Phyllobacterium_myrsinacearum
O_Propionibacteriales
S_Cutibacterium_acnes
G_Phyllobacterium
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
G_Saccharibacteria__TM7___G_1_
S_Fusobacterium_hwasookii
S_Veillonella_tobetsuensis
S_Fusobacterium_sp__HMT_204
S_Prevotella_histicola
S_Alloprevotella_sp__HMT_473
G_Alloprevotella
S_Campylobacter_concisus_nov_96_963
S_Haemophilus_sputorum
S_Neisseria_sp__HMT_020_nov_94_070
S_Campylobacter_showae
S_Haemophilus_haemolyticus
S_Porphyromonas_catoniae
S_Haemophilus_sp__HMT_036
S_Aggregatibacter_sp__HMT_458
S_Neisseria_elongata
S_Fusobacterium_nucleatum
S_Porphyromonas_pasteri
F_Porphyromonadaceae
G_Porphyromonas
S_Veillonella_atypica
S_Veillonella_dispar
S_Aggregatibacter_sp__HMT_513
S_Campylobacter_concisus
F_Campylobacteraceae
G_Campylobacter
O_Campylobacterales
C_Epsilonproteobacteria
S_Prevotella_melaninogenica
G_Prevotella
S_Veillonella_multispecies_spp6_3
F_Prevotellaceae
G_Eikenella
S_Eikenella_corrodens
S_Fusobacterium_periodonticum
S_Aggregatibacter_aphrophilus
S_Aggregatibacter_segnis
S_Veillonella_rogosae
O_Bacteroidales
P_Bacteroidetes
S_Haemophilus_pittmaniae
C_Bacteroidia
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
G_Aggregatibacter
S_Veillonella_parvula
S_Veillonella_multispecies_spp25_2
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
−5.93
−4.75
−3.57
−2.39
−1.21
−0.03
1.15
2.33
3.51
4.69
LDA SCORE (log 10)