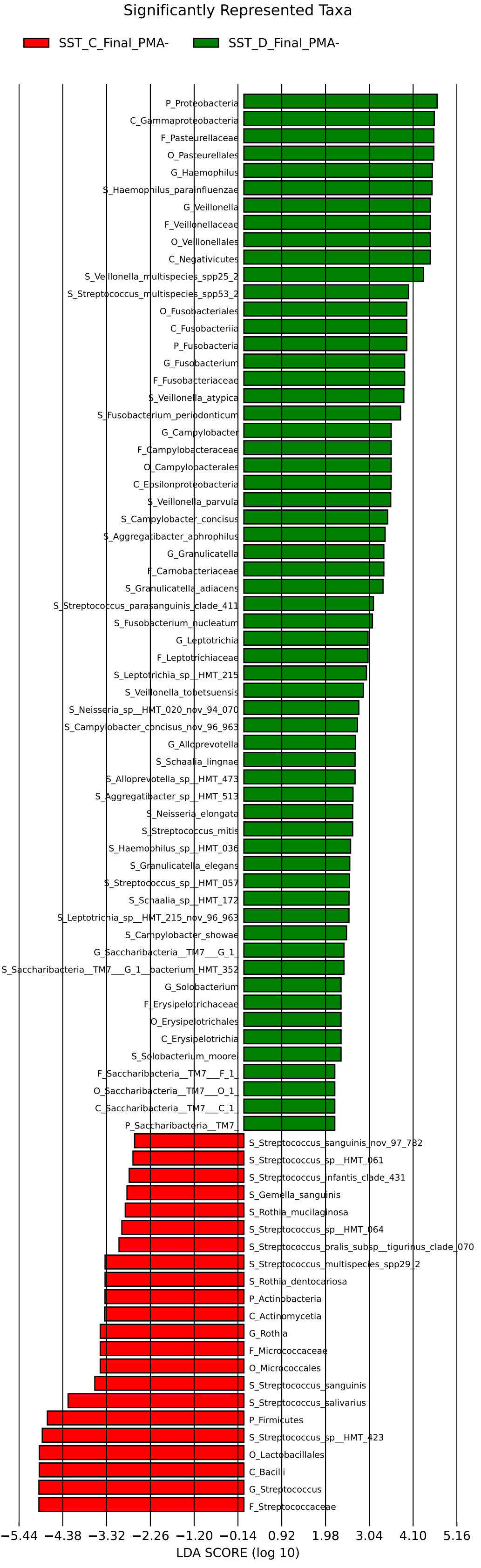
Significantly Represented Taxa
SST_C_Final_PMA-
SST_D_Final_PMA-
P_Proteobacteria
C_Gammaproteobacteria
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_Haemophilus_parainfluenzae
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
S_Veillonella_multispecies_spp25_2
S_Streptococcus_multispecies_spp53_2
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Veillonella_atypica
S_Fusobacterium_periodonticum
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Veillonella_parvula
S_Campylobacter_concisus
S_Aggregatibacter_aphrophilus
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Streptococcus_parasanguinis_clade_411
S_Fusobacterium_nucleatum
G_Leptotrichia
F_Leptotrichiaceae
S_Leptotrichia_sp__HMT_215
S_Veillonella_tobetsuensis
S_Neisseria_sp__HMT_020_nov_94_070
S_Campylobacter_concisus_nov_96_963
G_Alloprevotella
S_Schaalia_lingnae
S_Alloprevotella_sp__HMT_473
S_Aggregatibacter_sp__HMT_513
S_Neisseria_elongata
S_Streptococcus_mitis
S_Haemophilus_sp__HMT_036
S_Granulicatella_elegans
S_Streptococcus_sp__HMT_057
S_Schaalia_sp__HMT_172
S_Leptotrichia_sp__HMT_215_nov_96_963
S_Campylobacter_showae
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_Solobacterium_moorei
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
S_Streptococcus_sanguinis_nov_97_782
S_Streptococcus_sp__HMT_061
S_Streptococcus_infantis_clade_431
S_Gemella_sanguinis
S_Rothia_mucilaginosa
S_Streptococcus_sp__HMT_064
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_multispecies_spp29_2
S_Rothia_dentocariosa
P_Actinobacteria
C_Actinomycetia
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Streptococcus_sanguinis
S_Streptococcus_salivarius
P_Firmicutes
S_Streptococcus_sp__HMT_423
O_Lactobacillales
C_Bacilli
G_Streptococcus
F_Streptococcaceae
−5.44
−4.38
−3.32
−2.26
−1.20
−0.14
0.92
1.98
3.04
4.10
5.16
LDA SCORE (log 10)