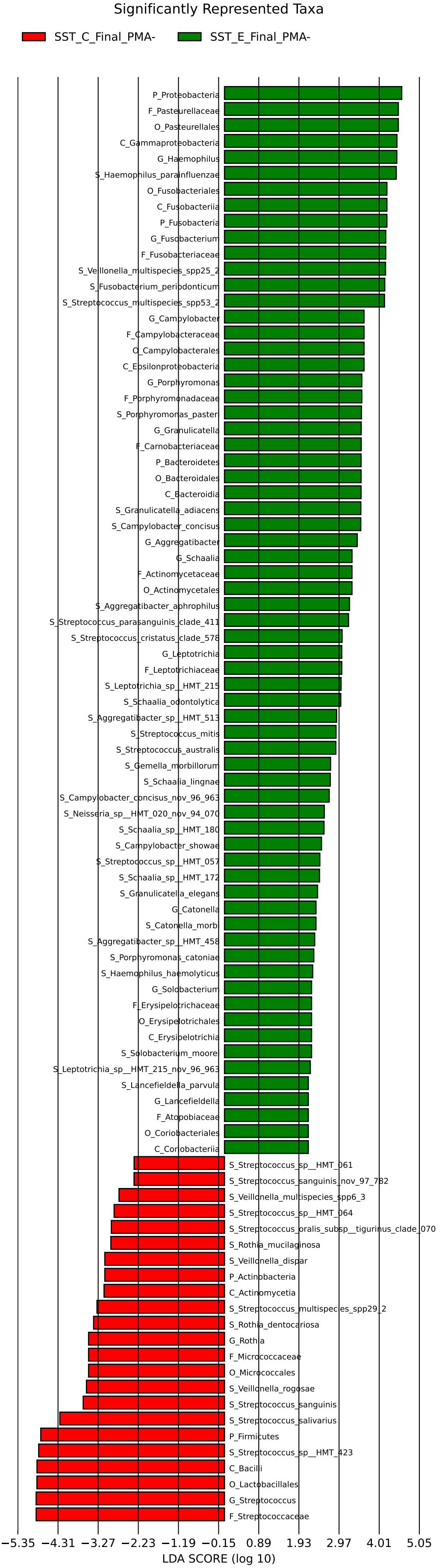
Significantly Represented Taxa
SST_C_Final_PMA-
SST_E_Final_PMA-
P_Proteobacteria
F_Pasteurellaceae
O_Pasteurellales
C_Gammaproteobacteria
G_Haemophilus
S_Haemophilus_parainfluenzae
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Veillonella_multispecies_spp25_2
S_Fusobacterium_periodonticum
S_Streptococcus_multispecies_spp53_2
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pasteri
G_Granulicatella
F_Carnobacteriaceae
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
S_Granulicatella_adiacens
S_Campylobacter_concisus
G_Aggregatibacter
G_Schaalia
F_Actinomycetaceae
O_Actinomycetales
S_Aggregatibacter_aphrophilus
S_Streptococcus_parasanguinis_clade_411
S_Streptococcus_cristatus_clade_578
G_Leptotrichia
F_Leptotrichiaceae
S_Leptotrichia_sp__HMT_215
S_Schaalia_odontolytica
S_Aggregatibacter_sp__HMT_513
S_Streptococcus_mitis
S_Streptococcus_australis
S_Gemella_morbillorum
S_Schaalia_lingnae
S_Campylobacter_concisus_nov_96_963
S_Neisseria_sp__HMT_020_nov_94_070
S_Schaalia_sp__HMT_180
S_Campylobacter_showae
S_Streptococcus_sp__HMT_057
S_Schaalia_sp__HMT_172
S_Granulicatella_elegans
G_Catonella
S_Catonella_morbi
S_Aggregatibacter_sp__HMT_458
S_Porphyromonas_catoniae
S_Haemophilus_haemolyticus
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_Solobacterium_moorei
S_Leptotrichia_sp__HMT_215_nov_96_963
S_Lancefieldella_parvula
G_Lancefieldella
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteriia
S_Streptococcus_sp__HMT_061
S_Streptococcus_sanguinis_nov_97_782
S_Veillonella_multispecies_spp6_3
S_Streptococcus_sp__HMT_064
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Rothia_mucilaginosa
S_Veillonella_dispar
P_Actinobacteria
C_Actinomycetia
S_Streptococcus_multispecies_spp29_2
S_Rothia_dentocariosa
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Veillonella_rogosae
S_Streptococcus_sanguinis
S_Streptococcus_salivarius
P_Firmicutes
S_Streptococcus_sp__HMT_423
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
−5.35
−4.31
−3.27
−2.23
−1.19
−0.15
0.89
1.93
2.97
4.01
5.05
LDA SCORE (log 10)