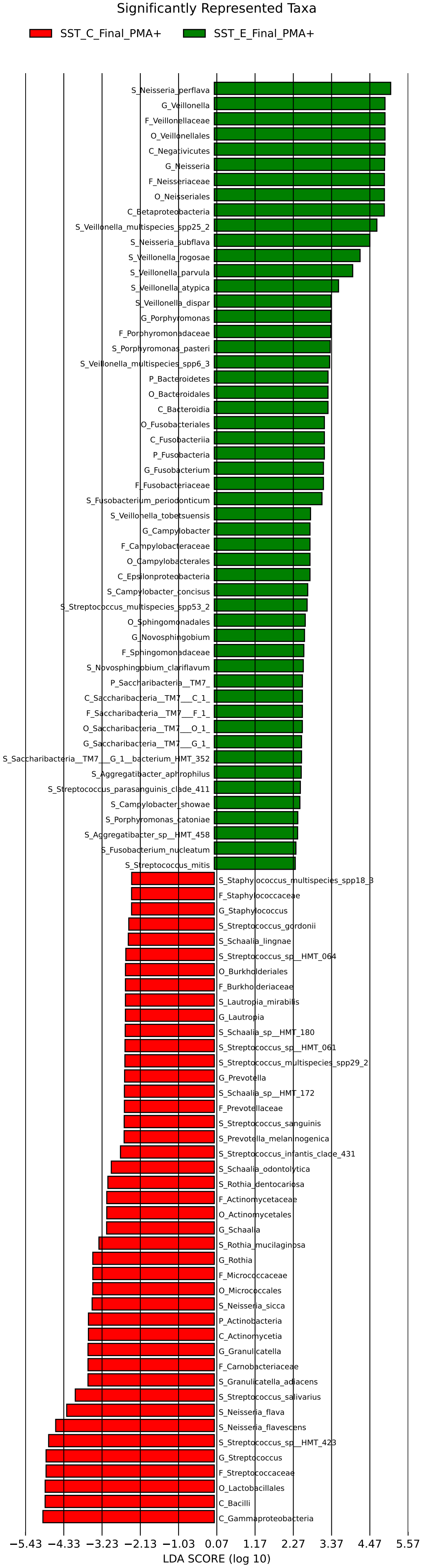
Significantly Represented Taxa
SST_C_Final_PMA+
SST_E_Final_PMA+
S_Neisseria_perflava
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
G_Neisseria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
S_Veillonella_multispecies_spp25_2
S_Neisseria_subflava
S_Veillonella_rogosae
S_Veillonella_parvula
S_Veillonella_atypica
S_Veillonella_dispar
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pasteri
S_Veillonella_multispecies_spp6_3
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Fusobacterium_periodonticum
S_Veillonella_tobetsuensis
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Campylobacter_concisus
S_Streptococcus_multispecies_spp53_2
O_Sphingomonadales
G_Novosphingobium
F_Sphingomonadaceae
S_Novosphingobium_clariflavum
P_Saccharibacteria__TM7_
C_Saccharibacteria__TM7___C_1_
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
S_Aggregatibacter_aphrophilus
S_Streptococcus_parasanguinis_clade_411
S_Campylobacter_showae
S_Porphyromonas_catoniae
S_Aggregatibacter_sp__HMT_458
S_Fusobacterium_nucleatum
S_Streptococcus_mitis
S_Staphylococcus_multispecies_spp18_3
F_Staphylococcaceae
G_Staphylococcus
S_Streptococcus_gordonii
S_Schaalia_lingnae
S_Streptococcus_sp__HMT_064
O_Burkholderiales
F_Burkholderiaceae
S_Lautropia_mirabilis
G_Lautropia
S_Schaalia_sp__HMT_180
S_Streptococcus_sp__HMT_061
S_Streptococcus_multispecies_spp29_2
G_Prevotella
S_Schaalia_sp__HMT_172
F_Prevotellaceae
S_Streptococcus_sanguinis
S_Prevotella_melaninogenica
S_Streptococcus_infantis_clade_431
S_Schaalia_odontolytica
S_Rothia_dentocariosa
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_Rothia_mucilaginosa
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Neisseria_sicca
P_Actinobacteria
C_Actinomycetia
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Streptococcus_salivarius
S_Neisseria_flava
S_Neisseria_flavescens
S_Streptococcus_sp__HMT_423
G_Streptococcus
F_Streptococcaceae
O_Lactobacillales
C_Bacilli
C_Gammaproteobacteria
−5.43
−4.33
−3.23
−2.13
−1.03
0.07
1.17
2.27
3.37
4.47
5.57
LDA SCORE (log 10)