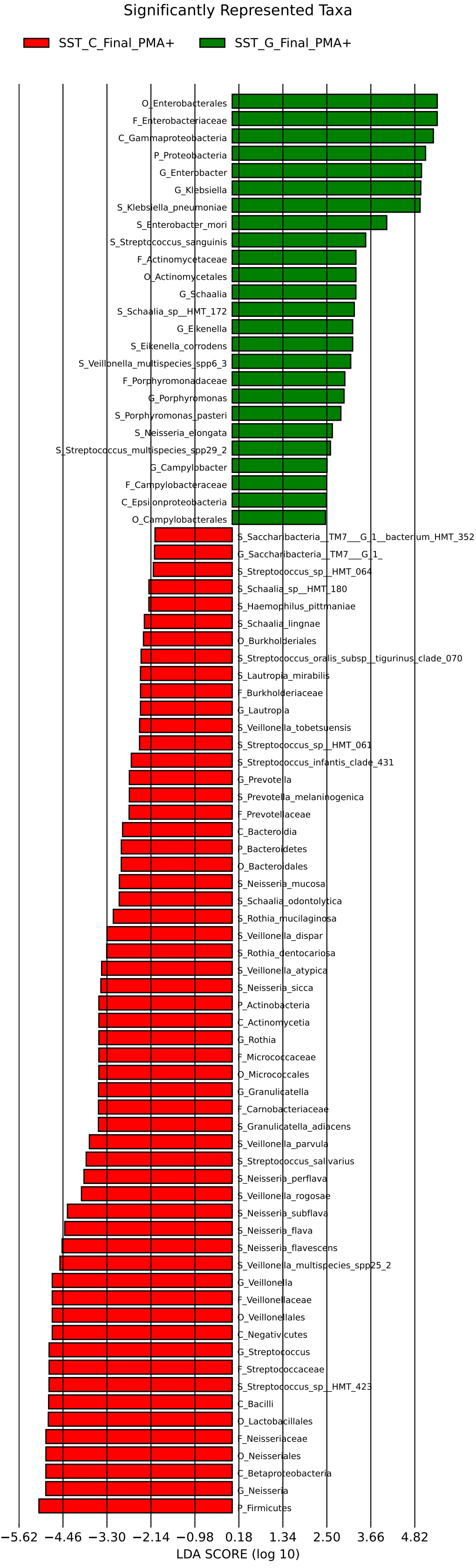
Significantly Represented Taxa
SST_C_Final_PMA+
SST_G_Final_PMA+
O_Enterobacterales
F_Enterobacteriaceae
C_Gammaproteobacteria
P_Proteobacteria
G_Enterobacter
G_Klebsiella
S_Klebsiella_pneumoniae
S_Enterobacter_mori
S_Streptococcus_sanguinis
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_Schaalia_sp__HMT_172
G_Eikenella
S_Eikenella_corrodens
S_Veillonella_multispecies_spp6_3
F_Porphyromonadaceae
G_Porphyromonas
S_Porphyromonas_pasteri
S_Neisseria_elongata
S_Streptococcus_multispecies_spp29_2
G_Campylobacter
F_Campylobacteraceae
C_Epsilonproteobacteria
O_Campylobacterales
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
G_Saccharibacteria__TM7___G_1_
S_Streptococcus_sp__HMT_064
S_Schaalia_sp__HMT_180
S_Haemophilus_pittmaniae
S_Schaalia_lingnae
O_Burkholderiales
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Lautropia_mirabilis
F_Burkholderiaceae
G_Lautropia
S_Veillonella_tobetsuensis
S_Streptococcus_sp__HMT_061
S_Streptococcus_infantis_clade_431
G_Prevotella
S_Prevotella_melaninogenica
F_Prevotellaceae
C_Bacteroidia
P_Bacteroidetes
O_Bacteroidales
S_Neisseria_mucosa
S_Schaalia_odontolytica
S_Rothia_mucilaginosa
S_Veillonella_dispar
S_Rothia_dentocariosa
S_Veillonella_atypica
S_Neisseria_sicca
P_Actinobacteria
C_Actinomycetia
G_Rothia
F_Micrococcaceae
O_Micrococcales
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Veillonella_parvula
S_Streptococcus_salivarius
S_Neisseria_perflava
S_Veillonella_rogosae
S_Neisseria_subflava
S_Neisseria_flava
S_Neisseria_flavescens
S_Veillonella_multispecies_spp25_2
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
C_Bacilli
O_Lactobacillales
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Neisseria
P_Firmicutes
−5.62
−4.46
−3.30
−2.14
−0.98
0.18
1.34
2.50
3.66
4.82
LDA SCORE (log 10)