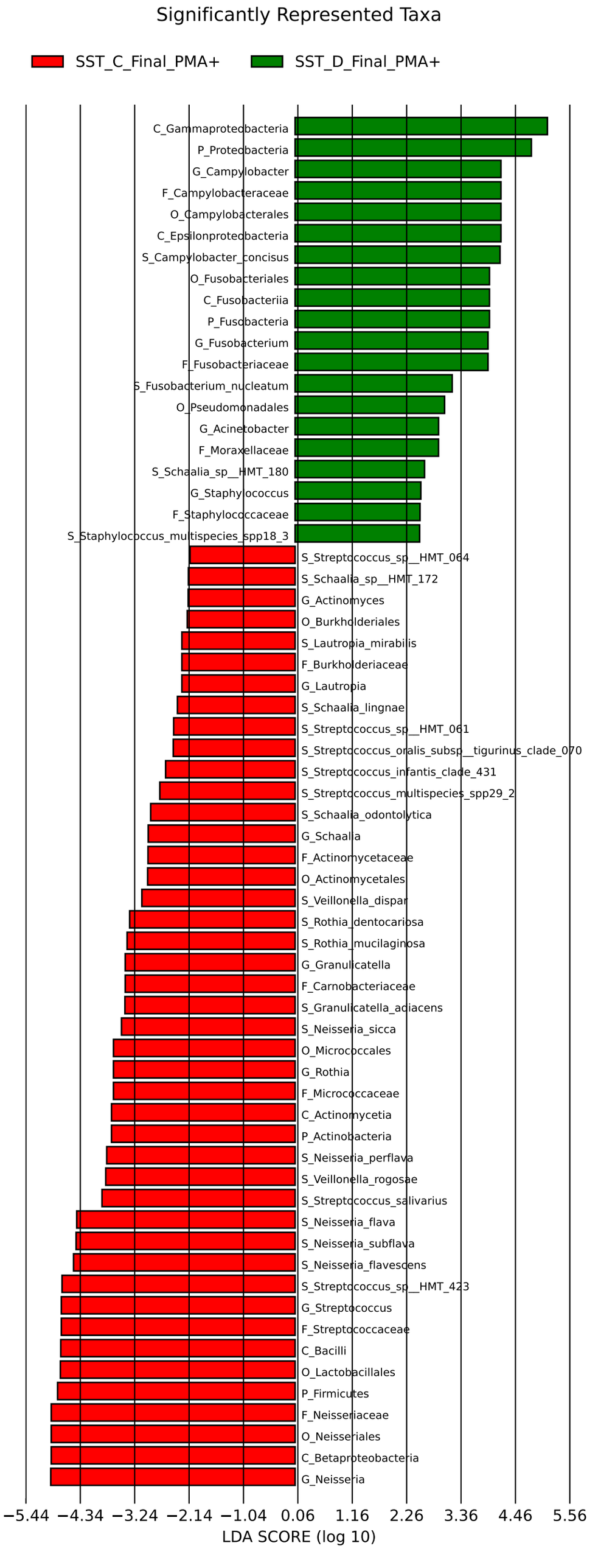
Significantly Represented Taxa
SST_C_Final_PMA+
SST_D_Final_PMA+
C_Gammaproteobacteria
P_Proteobacteria
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Campylobacter_concisus
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Fusobacterium_nucleatum
O_Pseudomonadales
G_Acinetobacter
F_Moraxellaceae
S_Schaalia_sp__HMT_180
G_Staphylococcus
F_Staphylococcaceae
S_Staphylococcus_multispecies_spp18_3
S_Streptococcus_sp__HMT_064
S_Schaalia_sp__HMT_172
G_Actinomyces
O_Burkholderiales
S_Lautropia_mirabilis
F_Burkholderiaceae
G_Lautropia
S_Schaalia_lingnae
S_Streptococcus_sp__HMT_061
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_infantis_clade_431
S_Streptococcus_multispecies_spp29_2
S_Schaalia_odontolytica
G_Schaalia
F_Actinomycetaceae
O_Actinomycetales
S_Veillonella_dispar
S_Rothia_dentocariosa
S_Rothia_mucilaginosa
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Neisseria_sicca
O_Micrococcales
G_Rothia
F_Micrococcaceae
C_Actinomycetia
P_Actinobacteria
S_Neisseria_perflava
S_Veillonella_rogosae
S_Streptococcus_salivarius
S_Neisseria_flava
S_Neisseria_subflava
S_Neisseria_flavescens
S_Streptococcus_sp__HMT_423
G_Streptococcus
F_Streptococcaceae
C_Bacilli
O_Lactobacillales
P_Firmicutes
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Neisseria
−5.44
−4.34
−3.24
−2.14
−1.04
0.06
1.16
2.26
3.36
4.46
5.56
LDA SCORE (log 10)