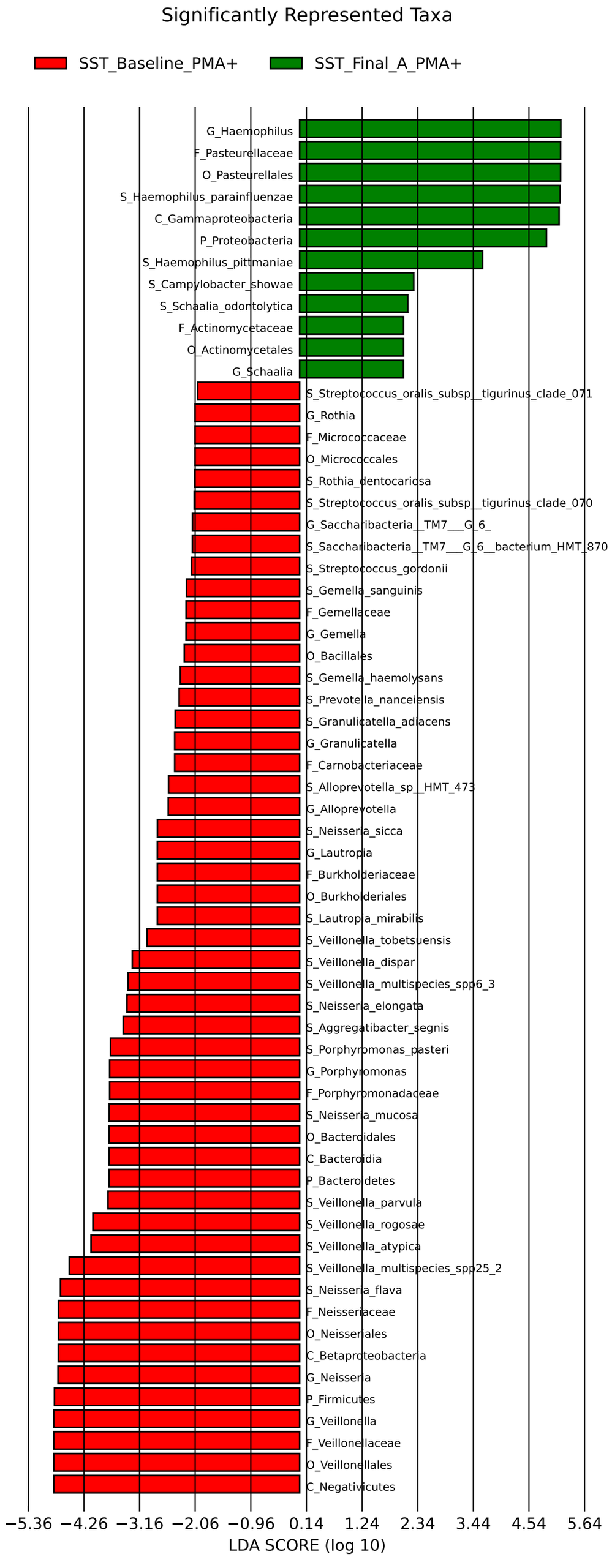
Significantly Represented Taxa
SST_Baseline_PMA+
SST_Final_A_PMA+
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
S_Haemophilus_parainfluenzae
C_Gammaproteobacteria
P_Proteobacteria
S_Haemophilus_pittmaniae
S_Campylobacter_showae
S_Schaalia_odontolytica
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_Streptococcus_oralis_subsp__tigurinus_clade_071
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Rothia_dentocariosa
S_Streptococcus_oralis_subsp__tigurinus_clade_070
G_Saccharibacteria__TM7___G_6_
S_Saccharibacteria__TM7___G_6__bacterium_HMT_870
S_Streptococcus_gordonii
S_Gemella_sanguinis
F_Gemellaceae
G_Gemella
O_Bacillales
S_Gemella_haemolysans
S_Prevotella_nanceiensis
S_Granulicatella_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_Alloprevotella_sp__HMT_473
G_Alloprevotella
S_Neisseria_sicca
G_Lautropia
F_Burkholderiaceae
O_Burkholderiales
S_Lautropia_mirabilis
S_Veillonella_tobetsuensis
S_Veillonella_dispar
S_Veillonella_multispecies_spp6_3
S_Neisseria_elongata
S_Aggregatibacter_segnis
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Neisseria_mucosa
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Veillonella_parvula
S_Veillonella_rogosae
S_Veillonella_atypica
S_Veillonella_multispecies_spp25_2
S_Neisseria_flava
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Neisseria
P_Firmicutes
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
−5.36
−4.26
−3.16
−2.06
−0.96
0.14
1.24
2.34
3.44
4.54
5.64
LDA SCORE (log 10)