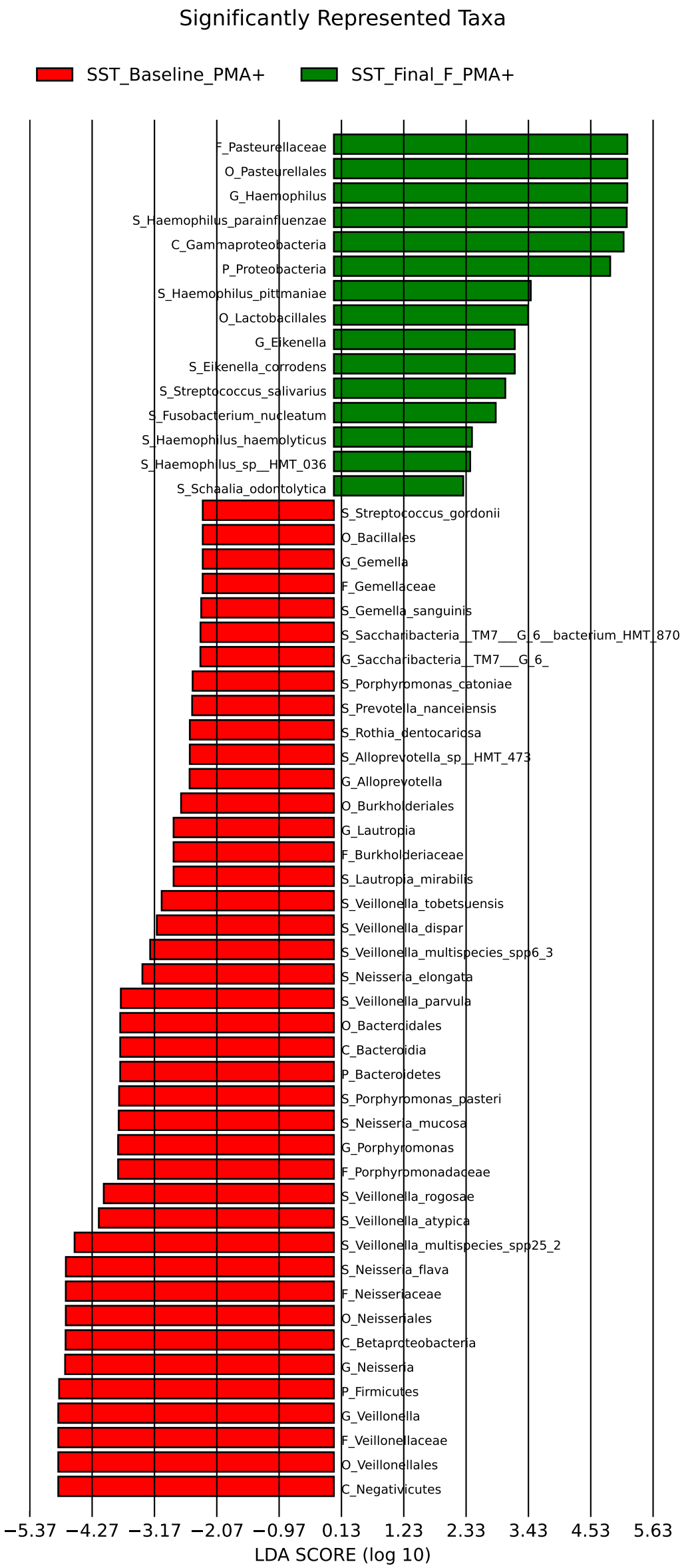
Significantly Represented Taxa
SST_Baseline_PMA+
SST_Final_F_PMA+
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_Haemophilus_parainfluenzae
C_Gammaproteobacteria
P_Proteobacteria
S_Haemophilus_pittmaniae
O_Lactobacillales
G_Eikenella
S_Eikenella_corrodens
S_Streptococcus_salivarius
S_Fusobacterium_nucleatum
S_Haemophilus_haemolyticus
S_Haemophilus_sp__HMT_036
S_Schaalia_odontolytica
S_Streptococcus_gordonii
O_Bacillales
G_Gemella
F_Gemellaceae
S_Gemella_sanguinis
S_Saccharibacteria__TM7___G_6__bacterium_HMT_870
G_Saccharibacteria__TM7___G_6_
S_Porphyromonas_catoniae
S_Prevotella_nanceiensis
S_Rothia_dentocariosa
S_Alloprevotella_sp__HMT_473
G_Alloprevotella
O_Burkholderiales
G_Lautropia
F_Burkholderiaceae
S_Lautropia_mirabilis
S_Veillonella_tobetsuensis
S_Veillonella_dispar
S_Veillonella_multispecies_spp6_3
S_Neisseria_elongata
S_Veillonella_parvula
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Porphyromonas_pasteri
S_Neisseria_mucosa
G_Porphyromonas
F_Porphyromonadaceae
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
−5.37
−4.27
−3.17
−2.07
−0.97
0.13
1.23
2.33
3.43
4.53
5.63
LDA SCORE (log 10)