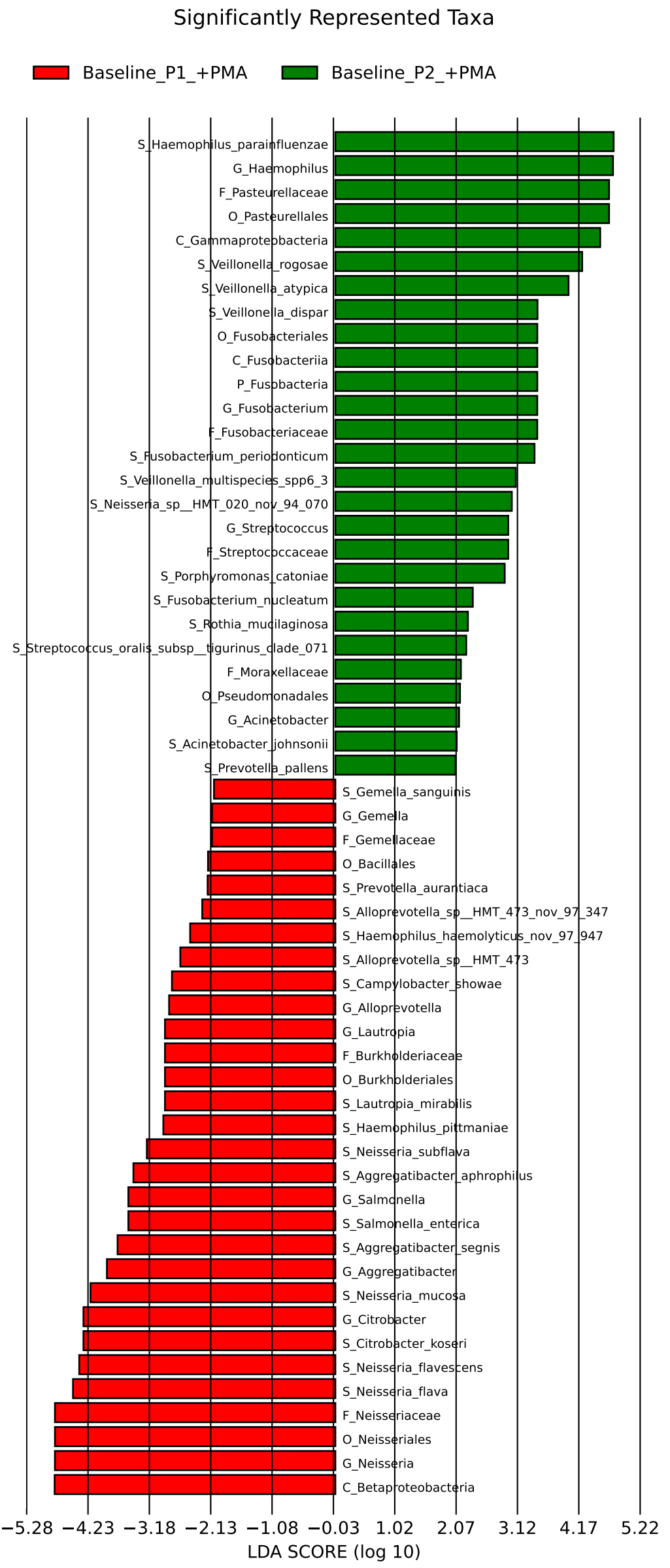
Significantly Represented Taxa
Baseline_P1_+PMA
Baseline_P2_+PMA
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
C_Gammaproteobacteria
S_Veillonella_rogosae
S_Veillonella_atypica
S_Veillonella_dispar
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Fusobacterium_periodonticum
S_Veillonella_multispecies_spp6_3
S_Neisseria_sp__HMT_020_nov_94_070
G_Streptococcus
F_Streptococcaceae
S_Porphyromonas_catoniae
S_Fusobacterium_nucleatum
S_Rothia_mucilaginosa
S_Streptococcus_oralis_subsp__tigurinus_clade_071
F_Moraxellaceae
O_Pseudomonadales
G_Acinetobacter
S_Acinetobacter_johnsonii
S_Prevotella_pallens
S_Gemella_sanguinis
G_Gemella
F_Gemellaceae
O_Bacillales
S_Prevotella_aurantiaca
S_Alloprevotella_sp__HMT_473_nov_97_347
S_Haemophilus_haemolyticus_nov_97_947
S_Alloprevotella_sp__HMT_473
S_Campylobacter_showae
G_Alloprevotella
G_Lautropia
F_Burkholderiaceae
O_Burkholderiales
S_Lautropia_mirabilis
S_Haemophilus_pittmaniae
S_Neisseria_subflava
S_Aggregatibacter_aphrophilus
G_Salmonella
S_Salmonella_enterica
S_Aggregatibacter_segnis
G_Aggregatibacter
S_Neisseria_mucosa
G_Citrobacter
S_Citrobacter_koseri
S_Neisseria_flavescens
S_Neisseria_flava
F_Neisseriaceae
O_Neisseriales
G_Neisseria
C_Betaproteobacteria
−5.28
−4.23
−3.18
−2.13
−1.08
−0.03
1.02
2.07
3.12
4.17
5.22
LDA SCORE (log 10)