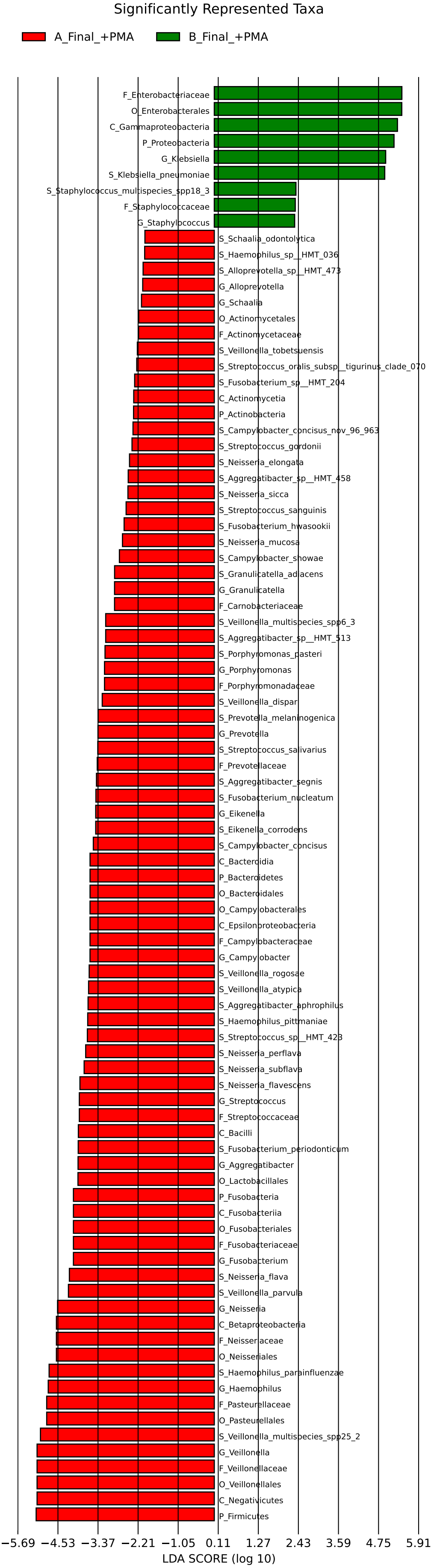
Significantly Represented Taxa
A_Final_+PMA
B_Final_+PMA
F_Enterobacteriaceae
O_Enterobacterales
C_Gammaproteobacteria
P_Proteobacteria
G_Klebsiella
S_Klebsiella_pneumoniae
S_Staphylococcus_multispecies_spp18_3
F_Staphylococcaceae
G_Staphylococcus
S_Schaalia_odontolytica
S_Haemophilus_sp__HMT_036
S_Alloprevotella_sp__HMT_473
G_Alloprevotella
G_Schaalia
O_Actinomycetales
F_Actinomycetaceae
S_Veillonella_tobetsuensis
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Fusobacterium_sp__HMT_204
C_Actinomycetia
P_Actinobacteria
S_Campylobacter_concisus_nov_96_963
S_Streptococcus_gordonii
S_Neisseria_elongata
S_Aggregatibacter_sp__HMT_458
S_Neisseria_sicca
S_Streptococcus_sanguinis
S_Fusobacterium_hwasookii
S_Neisseria_mucosa
S_Campylobacter_showae
S_Granulicatella_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_Veillonella_multispecies_spp6_3
S_Aggregatibacter_sp__HMT_513
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Veillonella_dispar
S_Prevotella_melaninogenica
G_Prevotella
S_Streptococcus_salivarius
F_Prevotellaceae
S_Aggregatibacter_segnis
S_Fusobacterium_nucleatum
G_Eikenella
S_Eikenella_corrodens
S_Campylobacter_concisus
C_Bacteroidia
P_Bacteroidetes
O_Bacteroidales
O_Campylobacterales
C_Epsilonproteobacteria
F_Campylobacteraceae
G_Campylobacter
S_Veillonella_rogosae
S_Veillonella_atypica
S_Aggregatibacter_aphrophilus
S_Haemophilus_pittmaniae
S_Streptococcus_sp__HMT_423
S_Neisseria_perflava
S_Neisseria_subflava
S_Neisseria_flavescens
G_Streptococcus
F_Streptococcaceae
C_Bacilli
S_Fusobacterium_periodonticum
G_Aggregatibacter
O_Lactobacillales
P_Fusobacteria
C_Fusobacteriia
O_Fusobacteriales
F_Fusobacteriaceae
G_Fusobacterium
S_Neisseria_flava
S_Veillonella_parvula
G_Neisseria
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
S_Veillonella_multispecies_spp25_2
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
P_Firmicutes
−5.69
−4.53
−3.37
−2.21
−1.05
0.11
1.27
2.43
3.59
4.75
5.91
LDA SCORE (log 10)