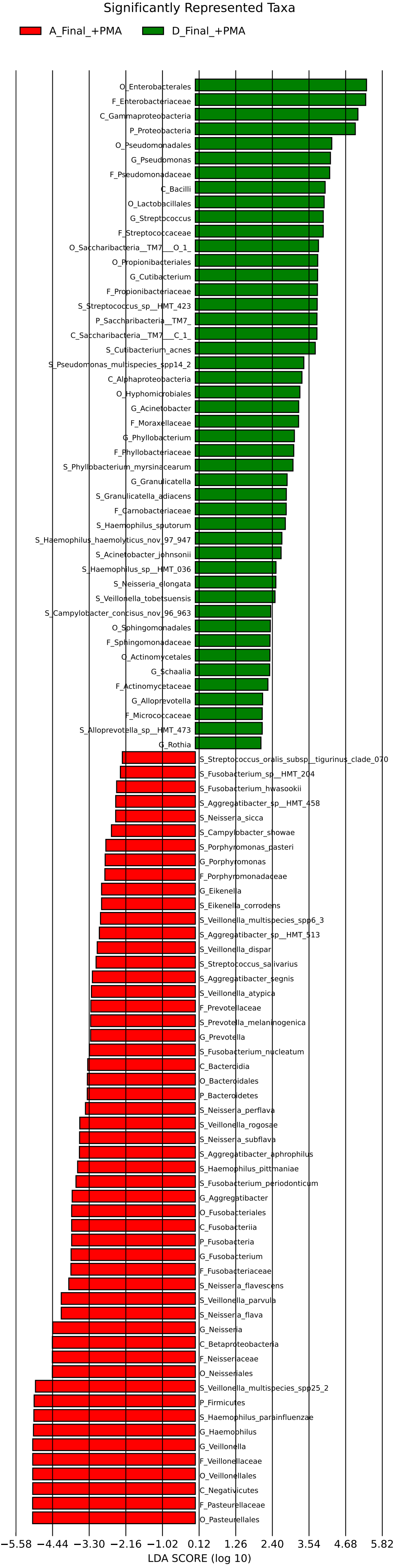
Significantly Represented Taxa
A_Final_+PMA
D_Final_+PMA
O_Enterobacterales
F_Enterobacteriaceae
C_Gammaproteobacteria
P_Proteobacteria
O_Pseudomonadales
G_Pseudomonas
F_Pseudomonadaceae
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
O_Saccharibacteria__TM7___O_1_
O_Propionibacteriales
G_Cutibacterium
F_Propionibacteriaceae
S_Streptococcus_sp__HMT_423
P_Saccharibacteria__TM7_
C_Saccharibacteria__TM7___C_1_
S_Cutibacterium_acnes
S_Pseudomonas_multispecies_spp14_2
C_Alphaproteobacteria
O_Hyphomicrobiales
G_Acinetobacter
F_Moraxellaceae
G_Phyllobacterium
F_Phyllobacteriaceae
S_Phyllobacterium_myrsinacearum
G_Granulicatella
S_Granulicatella_adiacens
F_Carnobacteriaceae
S_Haemophilus_sputorum
S_Haemophilus_haemolyticus_nov_97_947
S_Acinetobacter_johnsonii
S_Haemophilus_sp__HMT_036
S_Neisseria_elongata
S_Veillonella_tobetsuensis
S_Campylobacter_concisus_nov_96_963
O_Sphingomonadales
F_Sphingomonadaceae
O_Actinomycetales
G_Schaalia
F_Actinomycetaceae
G_Alloprevotella
F_Micrococcaceae
S_Alloprevotella_sp__HMT_473
G_Rothia
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Fusobacterium_sp__HMT_204
S_Fusobacterium_hwasookii
S_Aggregatibacter_sp__HMT_458
S_Neisseria_sicca
S_Campylobacter_showae
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
G_Eikenella
S_Eikenella_corrodens
S_Veillonella_multispecies_spp6_3
S_Aggregatibacter_sp__HMT_513
S_Veillonella_dispar
S_Streptococcus_salivarius
S_Aggregatibacter_segnis
S_Veillonella_atypica
F_Prevotellaceae
S_Prevotella_melaninogenica
G_Prevotella
S_Fusobacterium_nucleatum
C_Bacteroidia
O_Bacteroidales
P_Bacteroidetes
S_Neisseria_perflava
S_Veillonella_rogosae
S_Neisseria_subflava
S_Aggregatibacter_aphrophilus
S_Haemophilus_pittmaniae
S_Fusobacterium_periodonticum
G_Aggregatibacter
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Neisseria_flavescens
S_Veillonella_parvula
S_Neisseria_flava
G_Neisseria
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
S_Veillonella_multispecies_spp25_2
P_Firmicutes
S_Haemophilus_parainfluenzae
G_Haemophilus
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
F_Pasteurellaceae
O_Pasteurellales
−5.58
−4.44
−3.30
−2.16
−1.02
0.12
1.26
2.40
3.54
4.68
5.82
LDA SCORE (log 10)