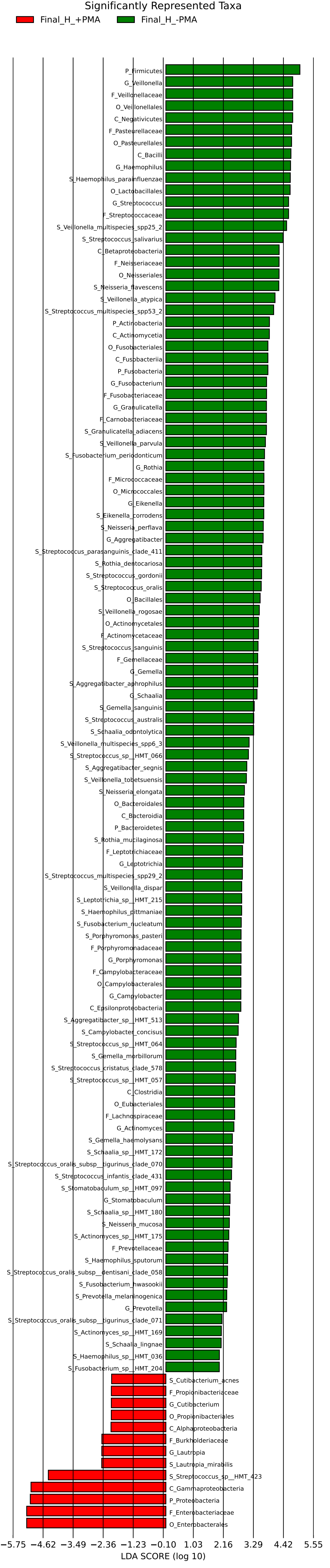
Significantly Represented Taxa
Final_H_+PMA
Final_H_-PMA
P_Firmicutes
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
F_Pasteurellaceae
O_Pasteurellales
C_Bacilli
G_Haemophilus
S_Haemophilus_parainfluenzae
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Veillonella_multispecies_spp25_2
S_Streptococcus_salivarius
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
S_Neisseria_flavescens
S_Veillonella_atypica
S_Streptococcus_multispecies_spp53_2
P_Actinobacteria
C_Actinomycetia
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Veillonella_parvula
S_Fusobacterium_periodonticum
G_Rothia
F_Micrococcaceae
O_Micrococcales
G_Eikenella
S_Eikenella_corrodens
S_Neisseria_perflava
G_Aggregatibacter
S_Streptococcus_parasanguinis_clade_411
S_Rothia_dentocariosa
S_Streptococcus_gordonii
S_Streptococcus_oralis
O_Bacillales
S_Veillonella_rogosae
O_Actinomycetales
F_Actinomycetaceae
S_Streptococcus_sanguinis
F_Gemellaceae
G_Gemella
S_Aggregatibacter_aphrophilus
G_Schaalia
S_Gemella_sanguinis
S_Streptococcus_australis
S_Schaalia_odontolytica
S_Veillonella_multispecies_spp6_3
S_Streptococcus_sp__HMT_066
S_Aggregatibacter_segnis
S_Veillonella_tobetsuensis
S_Neisseria_elongata
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Rothia_mucilaginosa
F_Leptotrichiaceae
G_Leptotrichia
S_Streptococcus_multispecies_spp29_2
S_Veillonella_dispar
S_Leptotrichia_sp__HMT_215
S_Haemophilus_pittmaniae
S_Fusobacterium_nucleatum
S_Porphyromonas_pasteri
F_Porphyromonadaceae
G_Porphyromonas
F_Campylobacteraceae
O_Campylobacterales
G_Campylobacter
C_Epsilonproteobacteria
S_Aggregatibacter_sp__HMT_513
S_Campylobacter_concisus
S_Streptococcus_sp__HMT_064
S_Gemella_morbillorum
S_Streptococcus_cristatus_clade_578
S_Streptococcus_sp__HMT_057
C_Clostridia
O_Eubacteriales
F_Lachnospiraceae
G_Actinomyces
S_Gemella_haemolysans
S_Schaalia_sp__HMT_172
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_infantis_clade_431
S_Stomatobaculum_sp__HMT_097
G_Stomatobaculum
S_Schaalia_sp__HMT_180
S_Neisseria_mucosa
S_Actinomyces_sp__HMT_175
F_Prevotellaceae
S_Haemophilus_sputorum
S_Streptococcus_oralis_subsp__dentisani_clade_058
S_Fusobacterium_hwasookii
S_Prevotella_melaninogenica
G_Prevotella
S_Streptococcus_oralis_subsp__tigurinus_clade_071
S_Actinomyces_sp__HMT_169
S_Schaalia_lingnae
S_Haemophilus_sp__HMT_036
S_Fusobacterium_sp__HMT_204
S_Cutibacterium_acnes
F_Propionibacteriaceae
G_Cutibacterium
O_Propionibacteriales
C_Alphaproteobacteria
F_Burkholderiaceae
G_Lautropia
S_Lautropia_mirabilis
S_Streptococcus_sp__HMT_423
C_Gammaproteobacteria
P_Proteobacteria
F_Enterobacteriaceae
O_Enterobacterales
−5.75
−4.62
−3.49
−2.36
−1.23
−0.10
1.03
2.16
3.29
4.42
5.55
LDA SCORE (log 10)