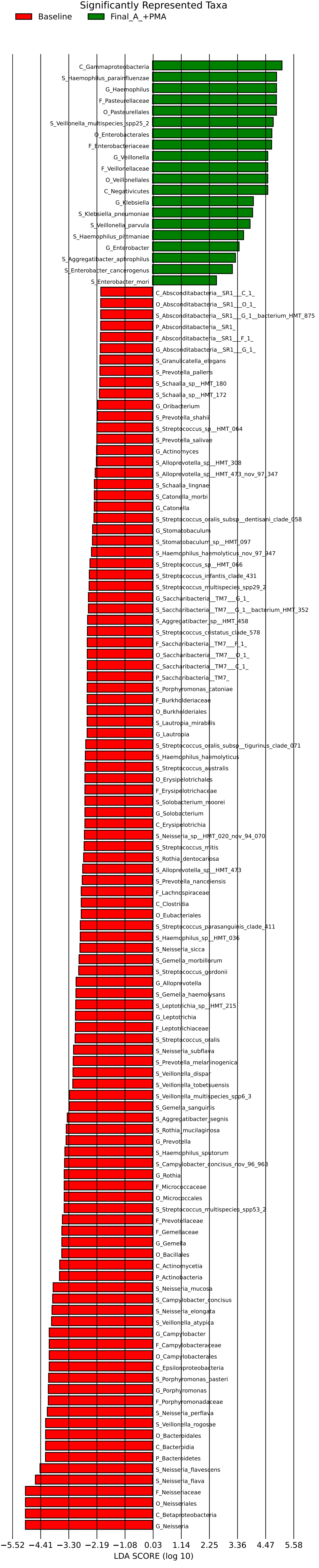
Significantly Represented Taxa
Baseline
Final_A_+PMA
C_Gammaproteobacteria
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
S_Veillonella_multispecies_spp25_2
O_Enterobacterales
F_Enterobacteriaceae
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
G_Klebsiella
S_Klebsiella_pneumoniae
S_Veillonella_parvula
S_Haemophilus_pittmaniae
G_Enterobacter
S_Aggregatibacter_aphrophilus
S_Enterobacter_cancerogenus
S_Enterobacter_mori
C_Absconditabacteria__SR1___C_1_
O_Absconditabacteria__SR1___O_1_
S_Absconditabacteria__SR1___G_1__bacterium_HMT_875
P_Absconditabacteria__SR1_
F_Absconditabacteria__SR1___F_1_
G_Absconditabacteria__SR1___G_1_
S_Granulicatella_elegans
S_Prevotella_pallens
S_Schaalia_sp__HMT_180
S_Schaalia_sp__HMT_172
G_Oribacterium
S_Prevotella_shahii
S_Streptococcus_sp__HMT_064
S_Prevotella_salivae
G_Actinomyces
S_Alloprevotella_sp__HMT_308
S_Alloprevotella_sp__HMT_473_nov_97_347
S_Schaalia_lingnae
S_Catonella_morbi
G_Catonella
S_Streptococcus_oralis_subsp__dentisani_clade_058
G_Stomatobaculum
S_Stomatobaculum_sp__HMT_097
S_Haemophilus_haemolyticus_nov_97_947
S_Streptococcus_sp__HMT_066
S_Streptococcus_infantis_clade_431
S_Streptococcus_multispecies_spp29_2
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
S_Aggregatibacter_sp__HMT_458
S_Streptococcus_cristatus_clade_578
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
S_Porphyromonas_catoniae
F_Burkholderiaceae
O_Burkholderiales
S_Lautropia_mirabilis
G_Lautropia
S_Streptococcus_oralis_subsp__tigurinus_clade_071
S_Haemophilus_haemolyticus
S_Streptococcus_australis
O_Erysipelotrichales
F_Erysipelotrichaceae
S_Solobacterium_moorei
G_Solobacterium
C_Erysipelotrichia
S_Neisseria_sp__HMT_020_nov_94_070
S_Streptococcus_mitis
S_Rothia_dentocariosa
S_Alloprevotella_sp__HMT_473
S_Prevotella_nanceiensis
F_Lachnospiraceae
C_Clostridia
O_Eubacteriales
S_Streptococcus_parasanguinis_clade_411
S_Haemophilus_sp__HMT_036
S_Neisseria_sicca
S_Gemella_morbillorum
S_Streptococcus_gordonii
G_Alloprevotella
S_Gemella_haemolysans
S_Leptotrichia_sp__HMT_215
G_Leptotrichia
F_Leptotrichiaceae
S_Streptococcus_oralis
S_Neisseria_subflava
S_Prevotella_melaninogenica
S_Veillonella_dispar
S_Veillonella_tobetsuensis
S_Veillonella_multispecies_spp6_3
S_Gemella_sanguinis
S_Aggregatibacter_segnis
S_Rothia_mucilaginosa
G_Prevotella
S_Haemophilus_sputorum
S_Campylobacter_concisus_nov_96_963
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Streptococcus_multispecies_spp53_2
F_Prevotellaceae
F_Gemellaceae
G_Gemella
O_Bacillales
C_Actinomycetia
P_Actinobacteria
S_Neisseria_mucosa
S_Campylobacter_concisus
S_Neisseria_elongata
S_Veillonella_atypica
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Neisseria_perflava
S_Veillonella_rogosae
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Neisseria_flavescens
S_Neisseria_flava
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
G_Neisseria
−5.52
−4.41
−3.30
−2.19
−1.08
0.03
1.14
2.25
3.36
4.47
5.58
LDA SCORE (log 10)