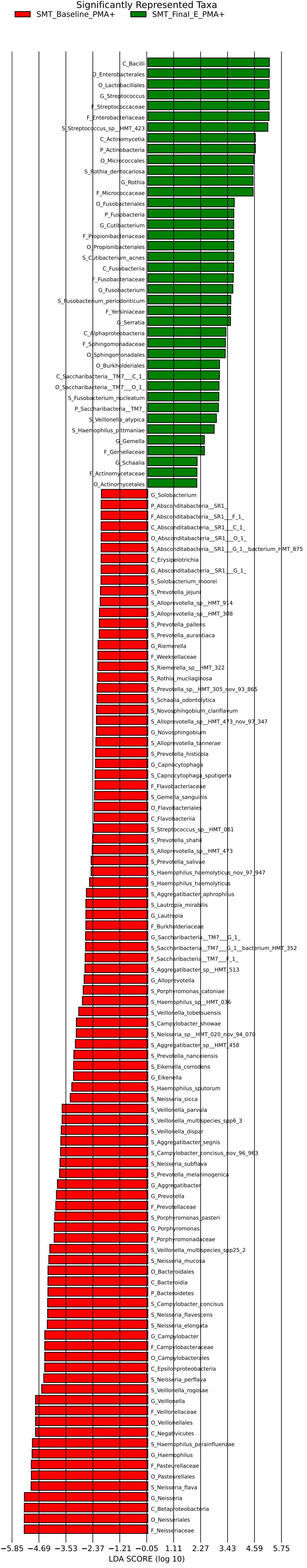
Significantly Represented Taxa
SMT_Baseline_PMA+
SMT_Final_E_PMA+
C_Bacilli
O_Enterobacterales
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
F_Enterobacteriaceae
S_Streptococcus_sp__HMT_423
C_Actinomycetia
P_Actinobacteria
O_Micrococcales
S_Rothia_dentocariosa
G_Rothia
F_Micrococcaceae
O_Fusobacteriales
P_Fusobacteria
G_Cutibacterium
F_Propionibacteriaceae
O_Propionibacteriales
S_Cutibacterium_acnes
C_Fusobacteriia
F_Fusobacteriaceae
G_Fusobacterium
S_Fusobacterium_periodonticum
F_Yersiniaceae
G_Serratia
C_Alphaproteobacteria
F_Sphingomonadaceae
O_Sphingomonadales
O_Burkholderiales
C_Saccharibacteria__TM7___C_1_
O_Saccharibacteria__TM7___O_1_
S_Fusobacterium_nucleatum
P_Saccharibacteria__TM7_
S_Veillonella_atypica
S_Haemophilus_parainfluenzae
G_Gemella
F_Gemellaceae
G_Schaalia
F_Actinomycetaceae
O_Actinomycetales
G_Solobacterium
P_Absconditabacteria__SR1_
F_Absconditabacteria__SR1___F_1_
C_Absconditabacteria__SR1___C_1_
O_Absconditabacteria__SR1___O_1_
S_Absconditabacteria__SR1___G_1__bacterium_HMT_875
C_Erysipelotrichia
G_Absconditabacteria__SR1___G_1_
S_Solobacterium_moorei
S_Prevotella_jejuni
S_Alloprevotella_sp__HMT_914
S_Alloprevotella_sp__HMT_308
S_Prevotella_pallens
S_Prevotella_aurantiaca
G_Riemerella
F_Weeksellaceae
S_Riemerella_sp__HMT_322
S_Rothia_mucilaginosa
S_Prevotella_sp__HMT_305_nov_93_865
S_Schaalia_odontolytica
S_Novosphingobium_clariflavum
S_Alloprevotella_sp__HMT_473_nov_97_347
G_Novosphingobium
S_Alloprevotella_tannerae
S_Prevotella_histicola
G_Capnocytophaga
S_Capnocytophaga_sputigena
F_Flavobacteriaceae
S_Gemella_sanguinis
O_Flavobacteriales
C_Flavobacteriia
S_Streptococcus_sp__HMT_061
S_Prevotella_shahii
S_Alloprevotella_sp__HMT_473
S_Prevotella_salivae
S_Haemophilus_haemolyticus_nov_97_947
S_Haemophilus_haemolyticus
S_Aggregatibacter_aphrophilus
S_Lautropia_mirabilis
G_Lautropia
F_Burkholderiaceae
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
F_Saccharibacteria__TM7___F_1_
S_Aggregatibacter_sp__HMT_513
G_Alloprevotella
S_Porphyromonas_catoniae
S_Haemophilus_sp__HMT_036
S_Veillonella_tobetsuensis
S_Campylobacter_showae
S_Neisseria_sp__HMT_020_nov_94_070
S_Aggregatibacter_sp__HMT_458
S_Prevotella_nanceiensis
S_Eikenella_corrodens
G_Eikenella
S_Haemophilus_sputorum
S_Neisseria_sicca
S_Veillonella_parvula
S_Veillonella_multispecies_spp6_3
S_Veillonella_dispar
S_Aggregatibacter_segnis
S_Campylobacter_concisus_nov_96_963
S_Neisseria_subflava
S_Prevotella_melaninogenica
G_Aggregatibacter
G_Prevotella
F_Prevotellaceae
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Veillonella_multispecies_spp25_2
S_Neisseria_mucosa
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Campylobacter_concisus
S_Neisseria_flavescens
S_Neisseria_elongata
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Neisseria_perflava
S_Veillonella_rogosae
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
S_Neisseria_flava
G_Neisseria
C_Betaproteobacteria
O_Neisseriales
F_Neisseriaceae
−5.85
−4.69
−3.53
−2.37
−1.21
−0.05
1.11
2.27
3.43
4.59
5.75
LDA SCORE (log 10)