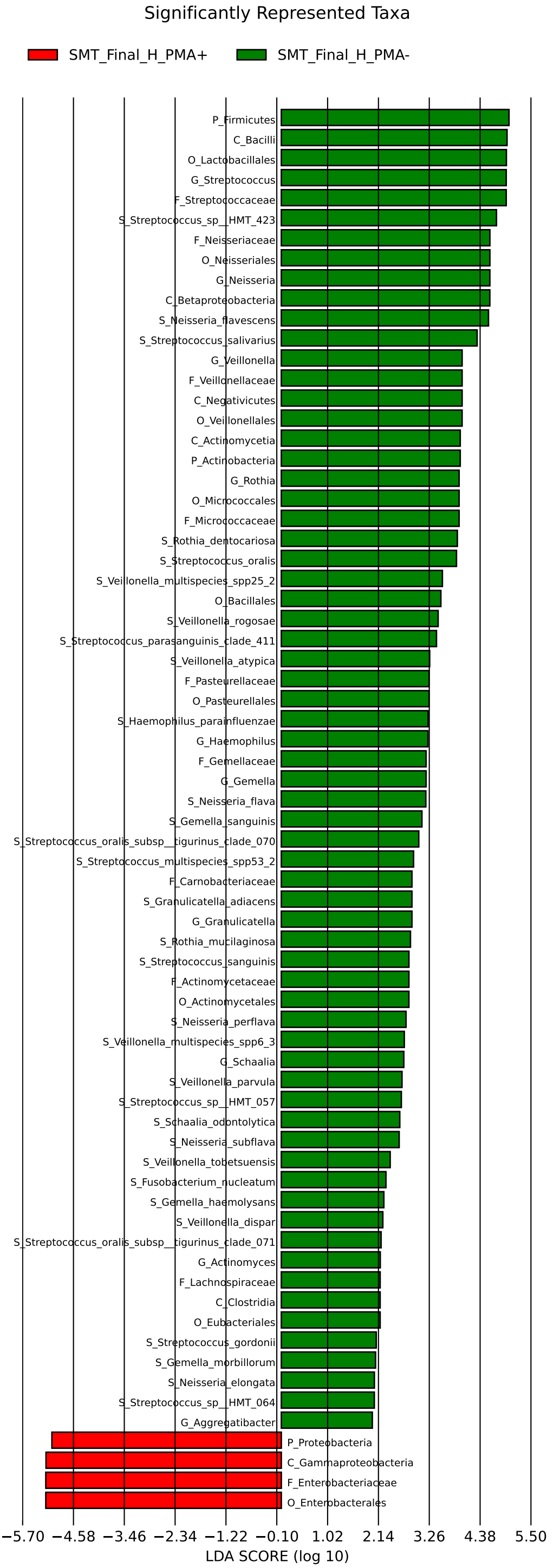
Significantly Represented Taxa
SMT_Final_H_PMA+
SMT_Final_H_PMA-
P_Firmicutes
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Streptococcus_sp__HMT_423
F_Neisseriaceae
O_Neisseriales
G_Neisseria
C_Betaproteobacteria
S_Neisseria_flavescens
S_Streptococcus_salivarius
G_Veillonella
F_Veillonellaceae
C_Negativicutes
O_Veillonellales
C_Actinomycetia
P_Actinobacteria
G_Rothia
O_Micrococcales
F_Micrococcaceae
S_Rothia_dentocariosa
S_Streptococcus_oralis
S_Veillonella_multispecies_spp25_2
O_Bacillales
S_Veillonella_rogosae
S_Streptococcus_parasanguinis_clade_411
S_Veillonella_atypica
F_Pasteurellaceae
O_Pasteurellales
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Gemellaceae
G_Gemella
S_Neisseria_flava
S_Gemella_sanguinis
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_multispecies_spp53_2
F_Carnobacteriaceae
S_Granulicatella_adiacens
G_Granulicatella
S_Rothia_mucilaginosa
S_Streptococcus_sanguinis
F_Actinomycetaceae
O_Actinomycetales
S_Neisseria_perflava
S_Veillonella_multispecies_spp6_3
G_Schaalia
S_Veillonella_parvula
S_Streptococcus_sp__HMT_057
S_Schaalia_odontolytica
S_Neisseria_subflava
S_Veillonella_tobetsuensis
S_Fusobacterium_nucleatum
S_Gemella_haemolysans
S_Veillonella_dispar
S_Streptococcus_oralis_subsp__tigurinus_clade_071
G_Actinomyces
F_Lachnospiraceae
C_Clostridia
O_Eubacteriales
S_Streptococcus_gordonii
S_Gemella_morbillorum
S_Neisseria_elongata
S_Streptococcus_sp__HMT_064
G_Aggregatibacter
P_Proteobacteria
C_Gammaproteobacteria
F_Enterobacteriaceae
O_Enterobacterales
−5.70
−4.58
−3.46
−2.34
−1.22
−0.10
1.02
2.14
3.26
4.38
5.50
LDA SCORE (log 10)