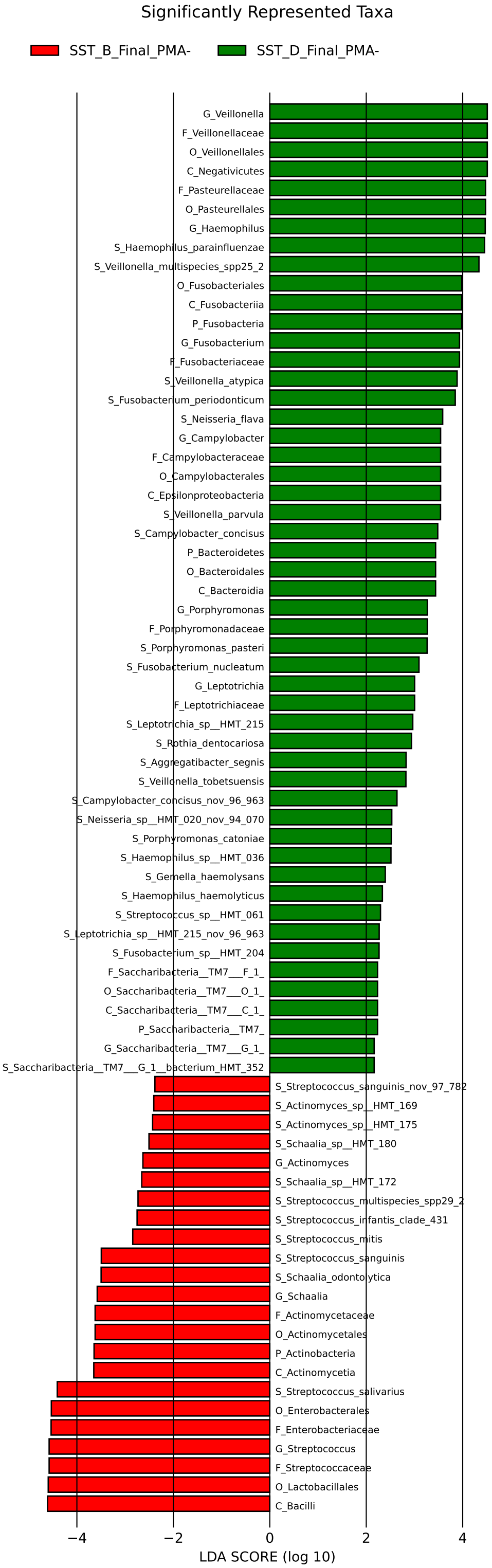
Significantly Represented Taxa
SST_B_Final_PMA-
SST_D_Final_PMA-
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
F_Pasteurellaceae
O_Pasteurellales
G_Haemophilus
S_Haemophilus_parainfluenzae
S_Veillonella_multispecies_spp25_2
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Veillonella_atypica
S_Fusobacterium_periodonticum
S_Neisseria_flava
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Veillonella_parvula
S_Campylobacter_concisus
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
G_Porphyromonas
F_Porphyromonadaceae
S_Porphyromonas_pasteri
S_Fusobacterium_nucleatum
G_Leptotrichia
F_Leptotrichiaceae
S_Leptotrichia_sp__HMT_215
S_Rothia_dentocariosa
S_Aggregatibacter_segnis
S_Veillonella_tobetsuensis
S_Campylobacter_concisus_nov_96_963
S_Neisseria_sp__HMT_020_nov_94_070
S_Porphyromonas_catoniae
S_Haemophilus_sp__HMT_036
S_Gemella_haemolysans
S_Haemophilus_haemolyticus
S_Streptococcus_sp__HMT_061
S_Leptotrichia_sp__HMT_215_nov_96_963
S_Fusobacterium_sp__HMT_204
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
S_Streptococcus_sanguinis_nov_97_782
S_Actinomyces_sp__HMT_169
S_Actinomyces_sp__HMT_175
S_Schaalia_sp__HMT_180
G_Actinomyces
S_Schaalia_sp__HMT_172
S_Streptococcus_multispecies_spp29_2
S_Streptococcus_infantis_clade_431
S_Streptococcus_mitis
S_Streptococcus_sanguinis
S_Schaalia_odontolytica
G_Schaalia
F_Actinomycetaceae
O_Actinomycetales
P_Actinobacteria
C_Actinomycetia
S_Streptococcus_salivarius
O_Enterobacterales
F_Enterobacteriaceae
G_Streptococcus
F_Streptococcaceae
O_Lactobacillales
C_Bacilli
−4
−2
0
2
4
LDA SCORE (log 10)