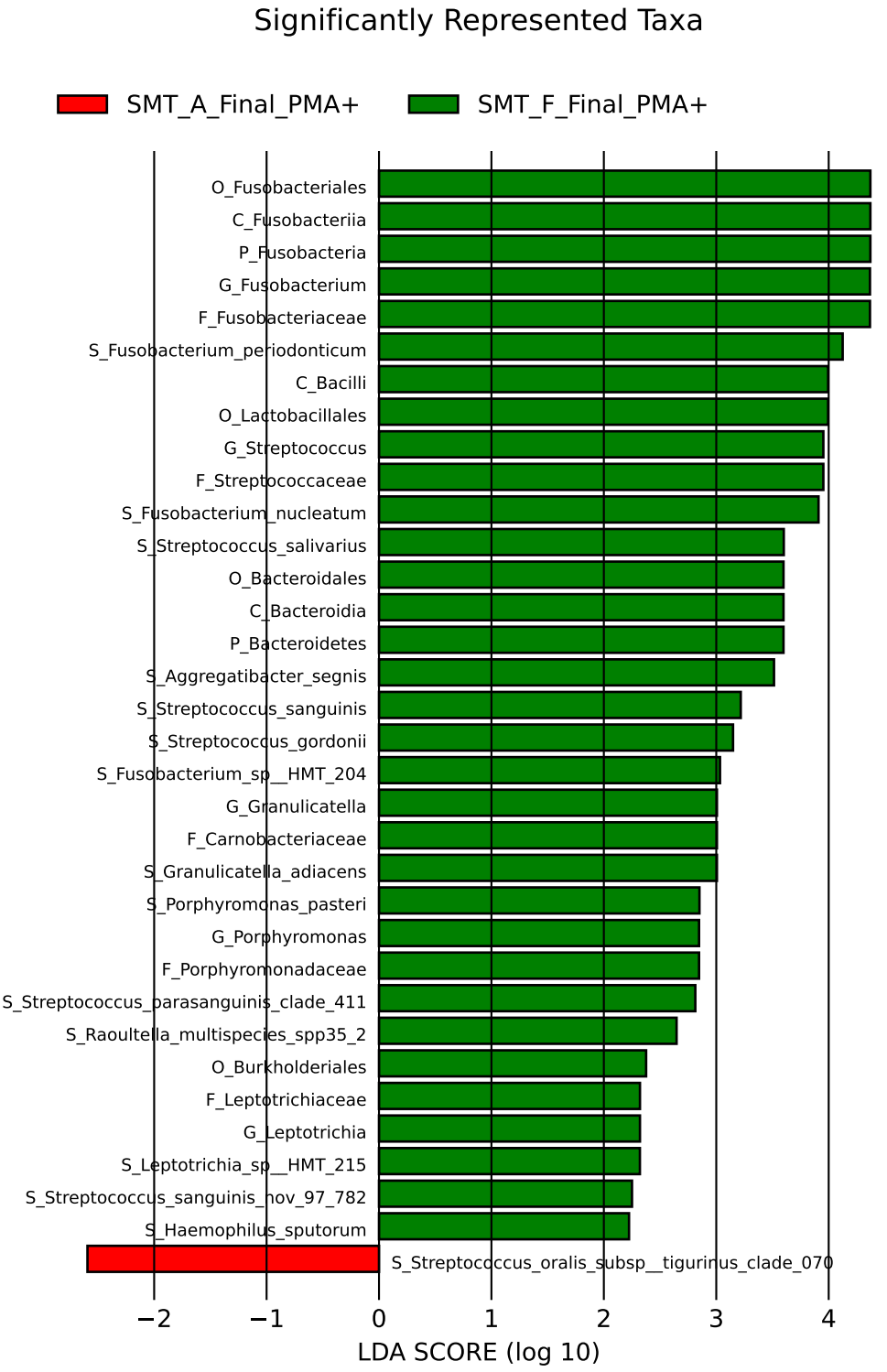
Significantly Represented Taxa
SMT_A_Final_PMA+
SMT_F_Final_PMA+
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
S_Fusobacterium_periodonticum
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
S_Fusobacterium_nucleatum
S_Streptococcus_salivarius
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Aggregatibacter_segnis
S_Streptococcus_sanguinis
S_Streptococcus_gordonii
S_Fusobacterium_sp__HMT_204
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
S_Porphyromonas_pasteri
G_Porphyromonas
F_Porphyromonadaceae
S_Streptococcus_parasanguinis_clade_411
S_Raoultella_multispecies_spp35_2
O_Burkholderiales
F_Leptotrichiaceae
G_Leptotrichia
S_Leptotrichia_sp__HMT_215
S_Streptococcus_sanguinis_nov_97_782
S_Haemophilus_sputorum
S_Streptococcus_oralis_subsp__tigurinus_clade_070
−2
−1
0
1
2
3
4
LDA SCORE (log 10)