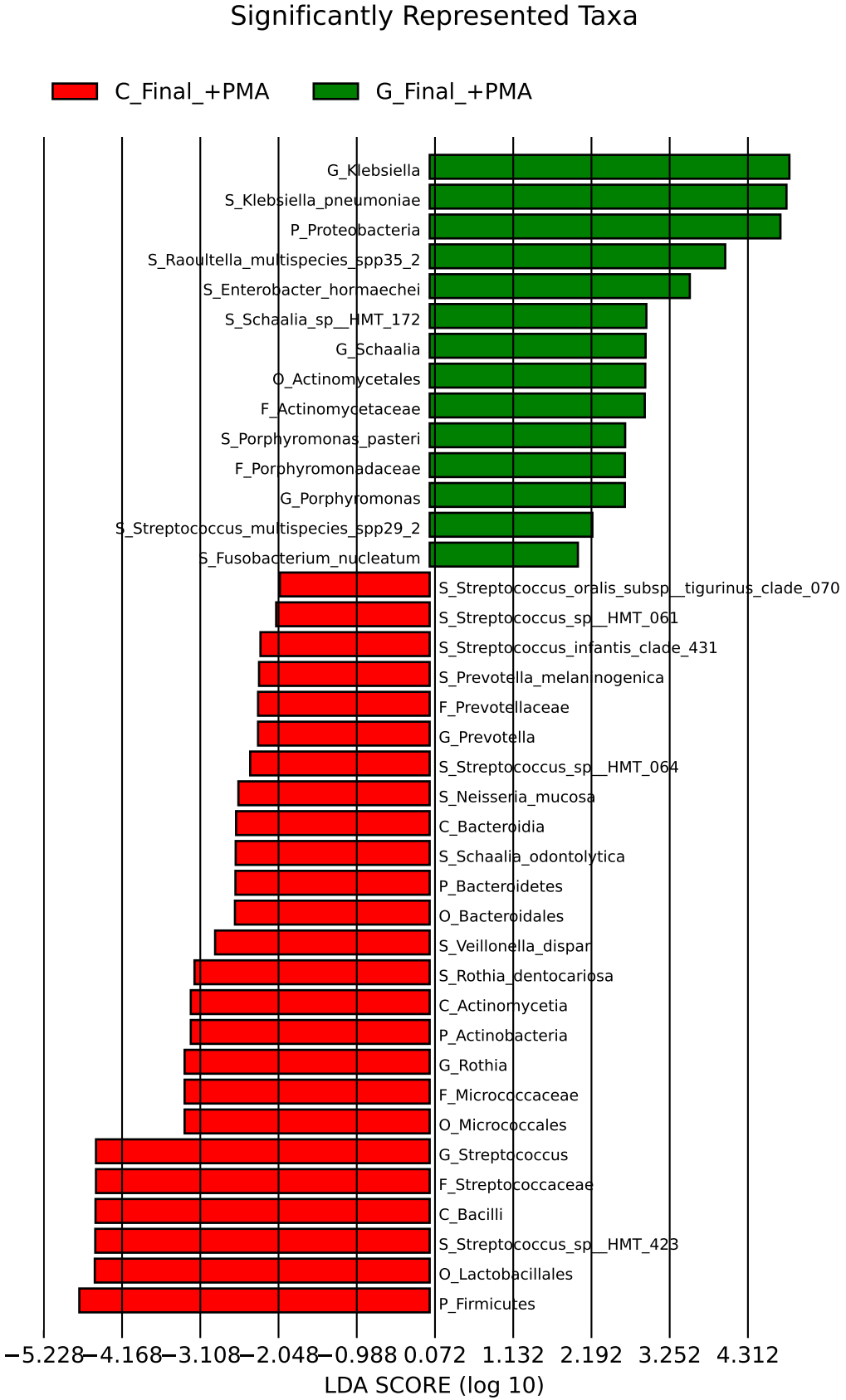
Significantly Represented Taxa
C_Final_+PMA
G_Final_+PMA
G_Klebsiella
S_Klebsiella_pneumoniae
P_Proteobacteria
S_Raoultella_multispecies_spp35_2
S_Enterobacter_hormaechei
S_Schaalia_sp__HMT_172
G_Schaalia
O_Actinomycetales
F_Actinomycetaceae
S_Porphyromonas_pasteri
F_Porphyromonadaceae
G_Porphyromonas
S_Streptococcus_multispecies_spp29_2
S_Fusobacterium_nucleatum
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Streptococcus_sp__HMT_061
S_Streptococcus_infantis_clade_431
S_Prevotella_melaninogenica
F_Prevotellaceae
G_Prevotella
S_Streptococcus_sp__HMT_064
S_Neisseria_mucosa
C_Bacteroidia
S_Schaalia_odontolytica
P_Bacteroidetes
O_Bacteroidales
S_Veillonella_dispar
S_Rothia_dentocariosa
C_Actinomycetia
P_Actinobacteria
G_Rothia
F_Micrococcaceae
O_Micrococcales
G_Streptococcus
F_Streptococcaceae
C_Bacilli
S_Streptococcus_sp__HMT_423
O_Lactobacillales
P_Firmicutes
−5.228
−4.168
−3.108
−2.048
−0.988
0.072
1.132
2.192
3.252
4.312
LDA SCORE (log 10)