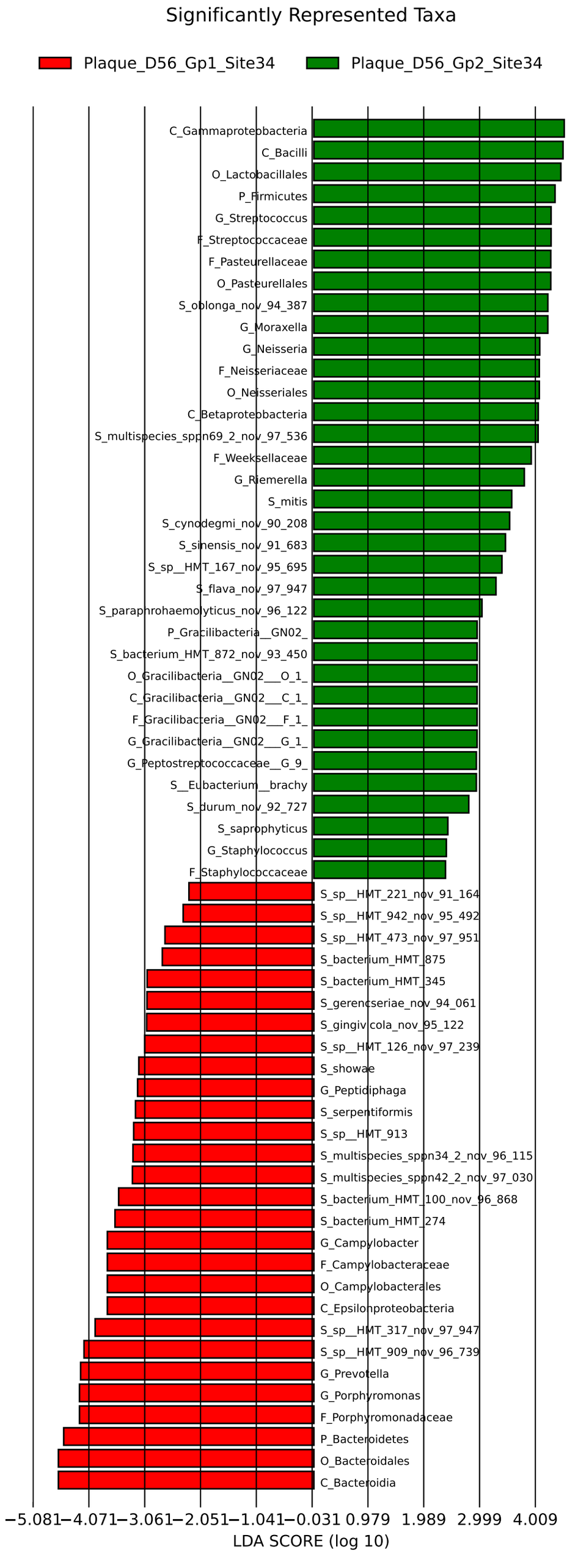
Significantly Represented Taxa
Plaque_D56_Gp1_Site34
Plaque_D56_Gp2_Site34
C_Gammaproteobacteria
C_Bacilli
O_Lactobacillales
P_Firmicutes
G_Streptococcus
F_Streptococcaceae
F_Pasteurellaceae
O_Pasteurellales
S_oblonga_nov_94_387
G_Moraxella
G_Neisseria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
S_multispecies_sppn69_2_nov_97_536
F_Weeksellaceae
G_Riemerella
S_mitis
S_cynodegmi_nov_90_208
S_sinensis_nov_91_683
S_sp__HMT_167_nov_95_695
S_flava_nov_97_947
S_paraphrohaemolyticus_nov_96_122
P_Gracilibacteria__GN02_
S_bacterium_HMT_872_nov_93_450
O_Gracilibacteria__GN02___O_1_
C_Gracilibacteria__GN02___C_1_
F_Gracilibacteria__GN02___F_1_
G_Gracilibacteria__GN02___G_1_
G_Peptostreptococcaceae__G_9_
S__Eubacterium__brachy
S_durum_nov_92_727
S_saprophyticus
G_Staphylococcus
F_Staphylococcaceae
S_sp__HMT_221_nov_91_164
S_sp__HMT_942_nov_95_492
S_sp__HMT_473_nov_97_951
S_bacterium_HMT_875
S_bacterium_HMT_345
S_gerencseriae_nov_94_061
S_gingivicola_nov_95_122
S_sp__HMT_126_nov_97_239
S_showae
G_Peptidiphaga
S_serpentiformis
S_sp__HMT_913
S_multispecies_sppn34_2_nov_96_115
S_multispecies_sppn42_2_nov_97_030
S_bacterium_HMT_100_nov_96_868
S_bacterium_HMT_274
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_sp__HMT_317_nov_97_947
S_sp__HMT_909_nov_96_739
G_Prevotella
G_Porphyromonas
F_Porphyromonadaceae
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
−5.081 −4.071 −3.061 −2.051 −1.041 −0.031 0.979 1.989 2.999 4.009
LDA SCORE (log 10)