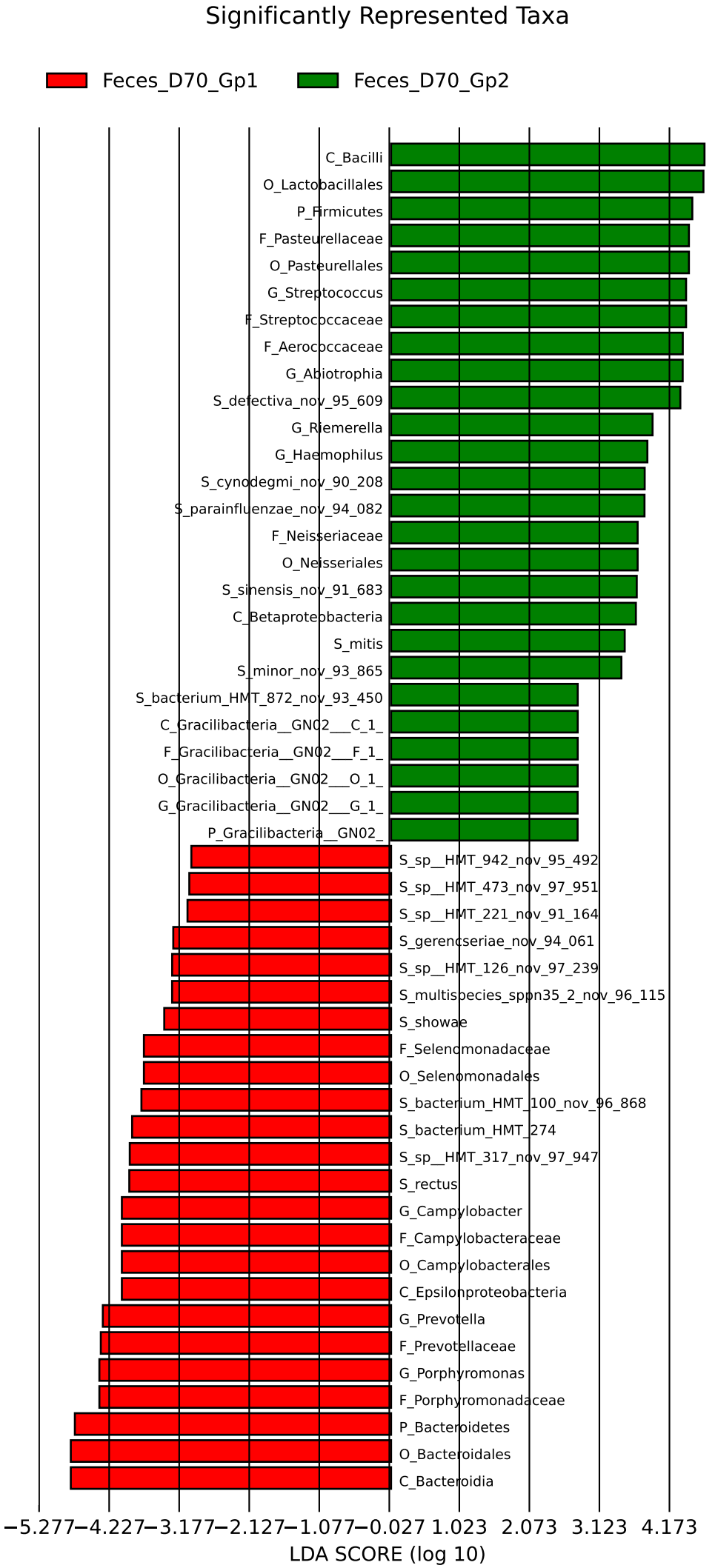
Significantly Represented Taxa
Feces_D70_Gp1
Feces_D70_Gp2
C_Bacilli
O_Lactobacillales
P_Firmicutes
F_Pasteurellaceae
O_Pasteurellales
G_Streptococcus
F_Streptococcaceae
F_Aerococcaceae
G_Abiotrophia
S_defectiva_nov_95_609
G_Riemerella
G_Haemophilus
S_cynodegmi_nov_90_208
S_parainfluenzae_nov_94_082
F_Neisseriaceae
O_Neisseriales
S_sinensis_nov_91_683
C_Betaproteobacteria
S_mitis
S_minor_nov_93_865
S_bacterium_HMT_872_nov_93_450
C_Gracilibacteria__GN02___C_1_
F_Gracilibacteria__GN02___F_1_
O_Gracilibacteria__GN02___O_1_
G_Gracilibacteria__GN02___G_1_
P_Gracilibacteria__GN02_
S_sp__HMT_942_nov_95_492
S_sp__HMT_473_nov_97_951
S_sp__HMT_221_nov_91_164
S_gerencseriae_nov_94_061
S_sp__HMT_126_nov_97_239
S_multispecies_sppn35_2_nov_96_115
S_showae
F_Selenomonadaceae
O_Selenomonadales
S_bacterium_HMT_100_nov_96_868
S_bacterium_HMT_274
S_sp__HMT_317_nov_97_947
S_rectus
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
G_Prevotella
F_Prevotellaceae
G_Porphyromonas
F_Porphyromonadaceae
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
−5.277
−4.227
−3.177
−2.127
−1.077
−0.027
1.023
2.073
3.123
4.173
LDA SCORE (log 10)