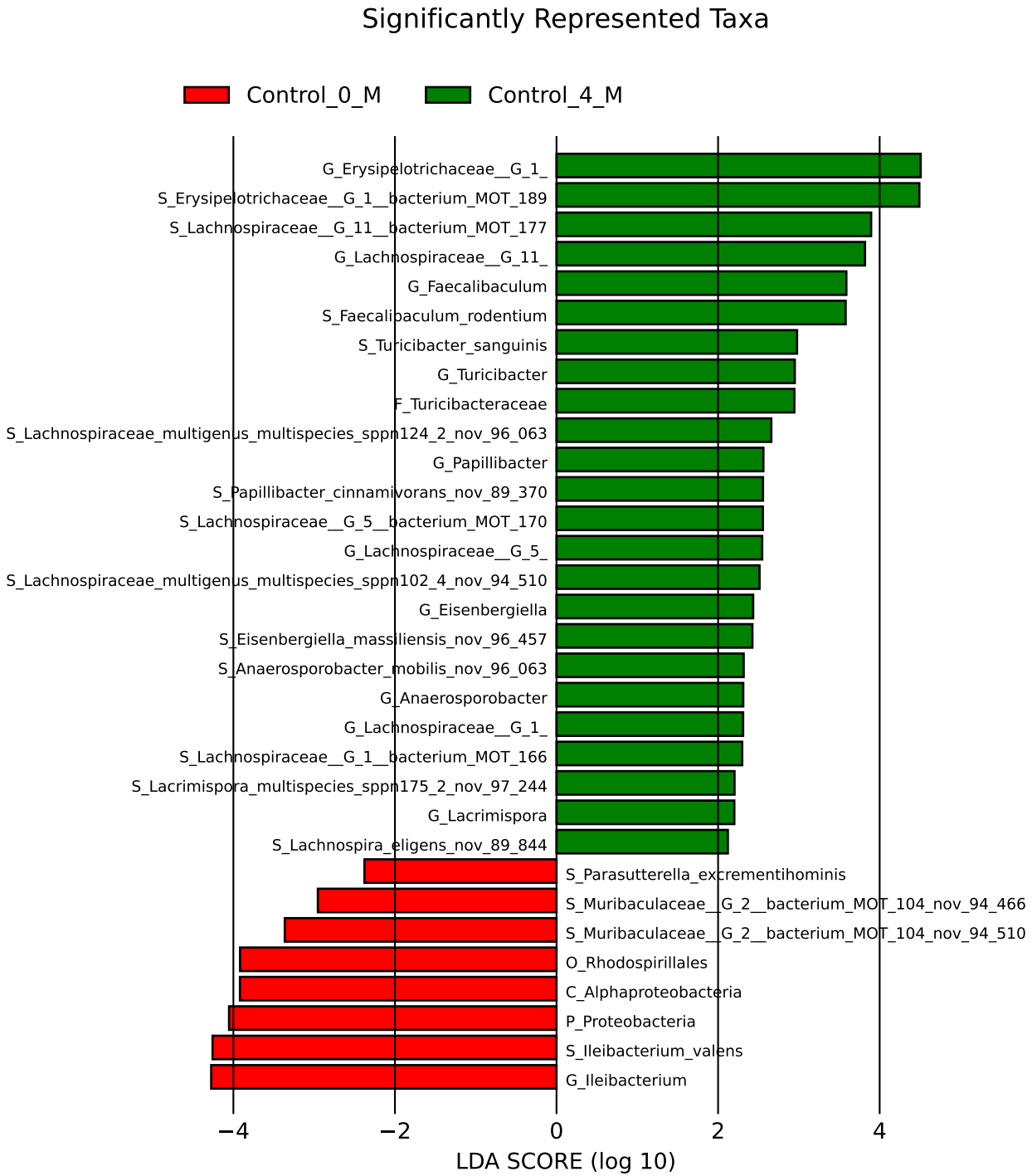
Significantly Represented Taxa
Control_0_M
Control_4_M
G_Erysipelotrichaceae__G_1_
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
S_Lachnospiraceae__G_11__bacterium_MOT_177
G_Lachnospiraceae__G_11_
G_Faecalibaculum
S_Faecalibaculum_rodentium
S_Turicibacter_sanguinis
G_Turicibacter
F_Turicibacteraceae
S_Lachnospiraceae_multigenus_multispecies_sppn124_2_nov_96_063
G_Papillibacter
S_Papillibacter_cinnamivorans_nov_89_370
S_Lachnospiraceae__G_5__bacterium_MOT_170
G_Lachnospiraceae__G_5_
S_Lachnospiraceae_multigenus_multispecies_sppn102_4_nov_94_510
G_Eisenbergiella
S_Eisenbergiella_massiliensis_nov_96_457
S_Anaerosporobacter_mobilis_nov_96_063
G_Anaerosporobacter
G_Lachnospiraceae__G_1_
S_Lachnospiraceae__G_1__bacterium_MOT_166
S_Lacrimispora_multispecies_sppn175_2_nov_97_244
G_Lacrimispora
S_Lachnospira_eligens_nov_89_844
S_Parasutterella_excrementihominis
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_466
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_510
O_Rhodospirillales
C_Alphaproteobacteria
P_Proteobacteria
S_Ileibacterium_valens
G_Ileibacterium
−4
−2
0
2
4
LDA SCORE (log 10)