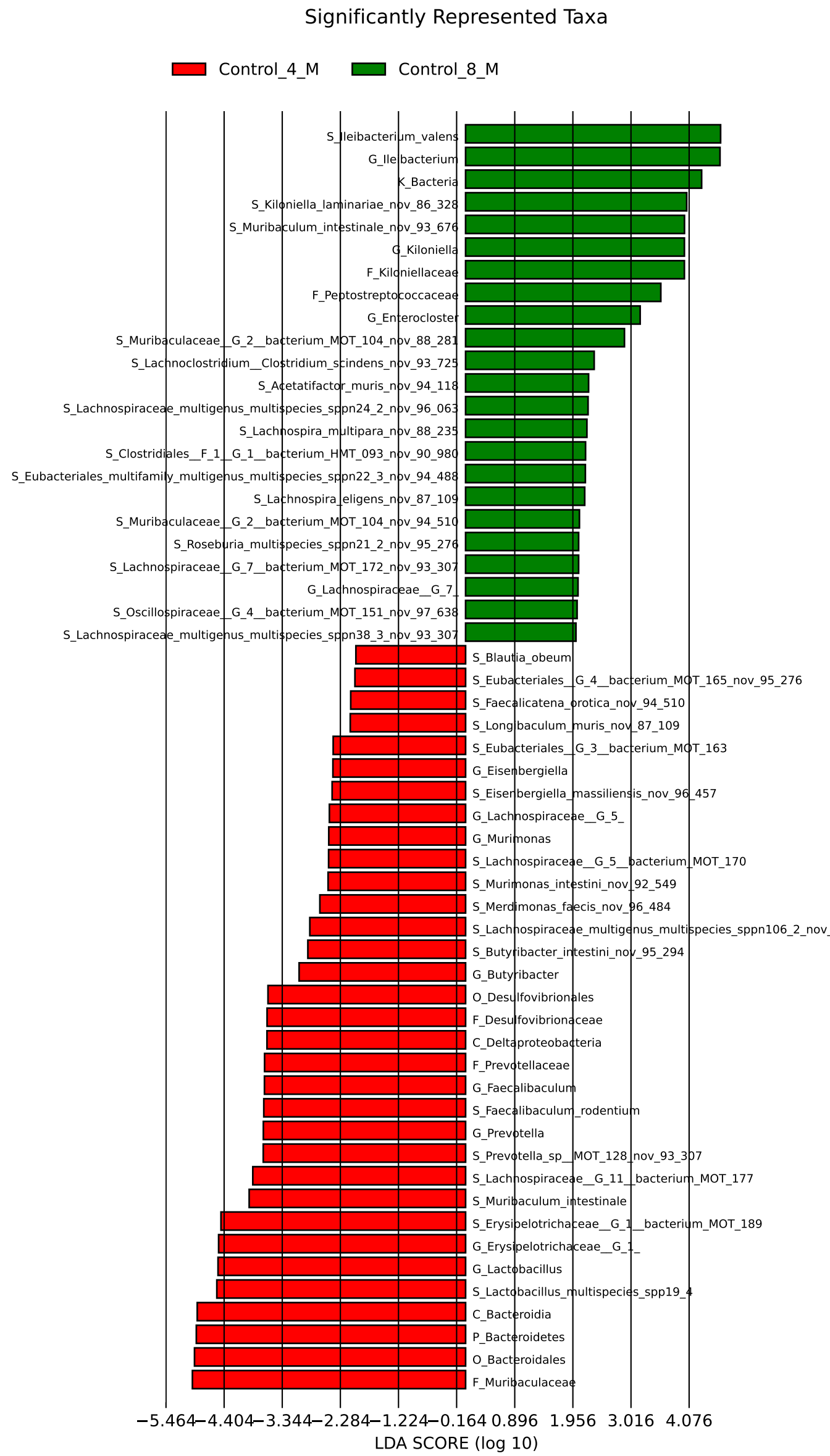
Significantly Represented Taxa
Control_4_M
Control_8_M
S_Ileibacterium_valens
G_Ileibacterium
K_Bacteria
S_Kiloniella_laminariae_nov_86_328
S_Muribaculum_intestinale_nov_93_676
G_Kiloniella
F_Kiloniellaceae
F_Peptostreptococcaceae
G_Enterocloster
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_88_281
S_Lachnoclostridium__Clostridium_scindens_nov_93_725
S_Acetatifactor_muris_nov_94_118
S_Lachnospiraceae_multigenus_multispecies_sppn24_2_nov_96_063
S_Lachnospira_multipara_nov_88_235
S_Clostridiales__F_1__G_1__bacterium_HMT_093_nov_90_980
S_Eubacteriales_multifamily_multigenus_multispecies_sppn22_3_nov_94_488
S_Lachnospira_eligens_nov_87_109
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_94_510
S_Roseburia_multispecies_sppn21_2_nov_95_276
S_Lachnospiraceae__G_7__bacterium_MOT_172_nov_93_307
G_Lachnospiraceae__G_7_
S_Oscillospiraceae__G_4__bacterium_MOT_151_nov_97_638
S_Lachnospiraceae_multigenus_multispecies_sppn38_3_nov_93_307
S_Blautia_obeum
S_Eubacteriales__G_4__bacterium_MOT_165_nov_95_276
S_Faecalicatena_orotica_nov_94_510
S_Longibaculum_muris_nov_87_109
S_Eubacteriales__G_3__bacterium_MOT_163
G_Eisenbergiella
S_Eisenbergiella_massiliensis_nov_96_457
G_Lachnospiraceae__G_5_
G_Murimonas
S_Lachnospiraceae__G_5__bacterium_MOT_170
S_Murimonas_intestini_nov_92_549
S_Merdimonas_faecis_nov_96_484
S_Lachnospiraceae_multigenus_multispecies_sppn106_2_nov_
S_Butyribacter_intestini_nov_95_294
G_Butyribacter
O_Desulfovibrionales
F_Desulfovibrionaceae
C_Deltaproteobacteria
F_Prevotellaceae
G_Faecalibaculum
S_Faecalibaculum_rodentium
G_Prevotella
S_Prevotella_sp__MOT_128_nov_93_307
S_Lachnospiraceae__G_11__bacterium_MOT_177
S_Muribaculum_intestinale
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
G_Erysipelotrichaceae__G_1_
G_Lactobacillus
S_Lactobacillus_multispecies_spp19_4
C_Bacteroidia
P_Bacteroidetes
O_Bacteroidales
F_Muribaculaceae
−5.464
−4.404
−3.344
−2.284
−1.224
−0.164
0.896
1.956
3.016
4.076
LDA SCORE (log 10)