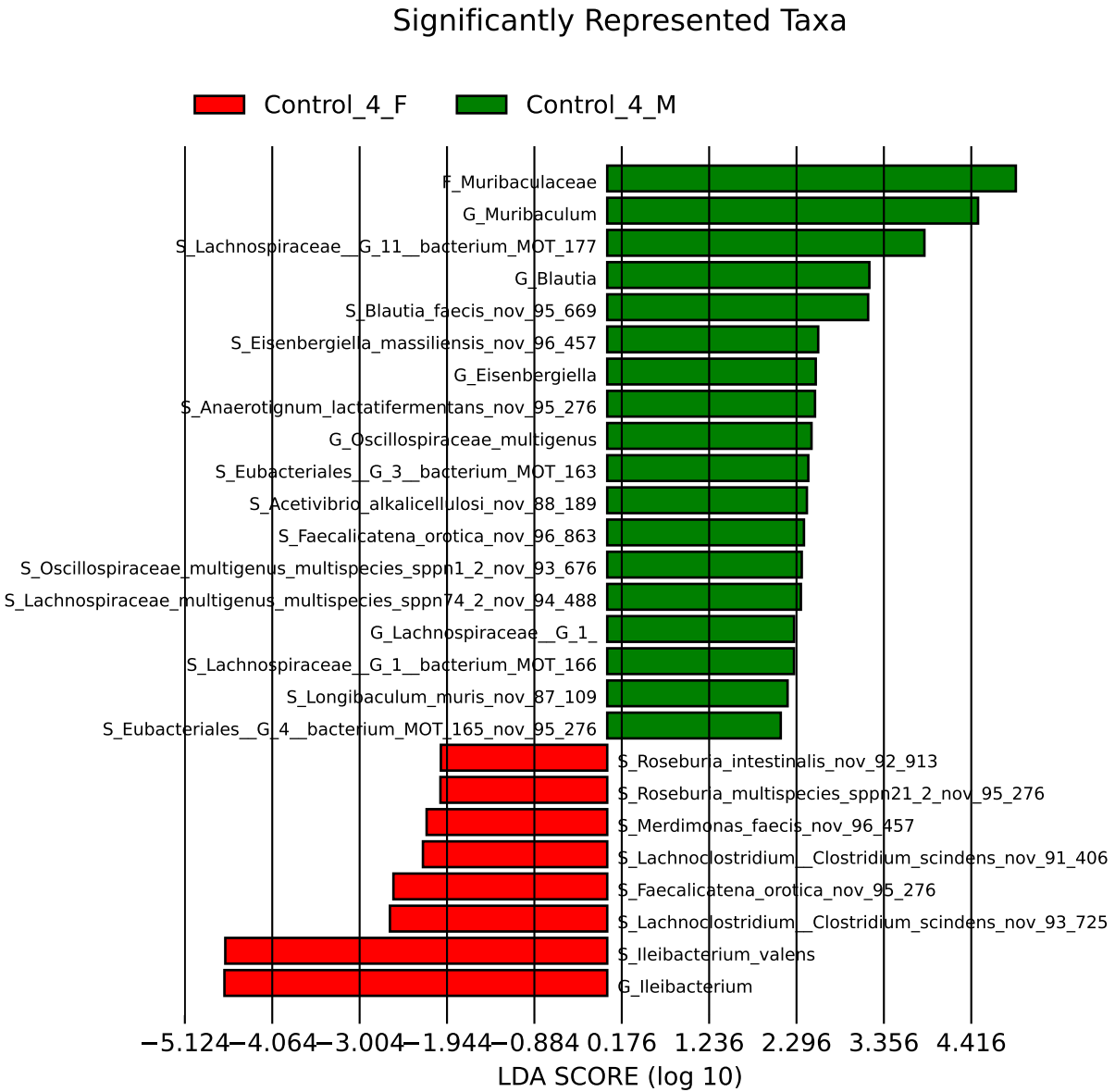
Significantly Represented Taxa
Control_4_F
Control_4_M
F_Muribaculaceae
G_Muribaculum
S_Lachnospiraceae__G_11__bacterium_MOT_177
G_Blautia
S_Blautia_faecis_nov_95_669
S_Eisenbergiella_massiliensis_nov_96_457
G_Eisenbergiella
S_Anaerotignum_lactatifermentans_nov_95_276
G_Oscillospiraceae_multigenus
S_Eubacteriales__G_3__bacterium_MOT_163
S_Acetivibrio_alkalicellulosi_nov_88_189
S_Faecalicatena_orotica_nov_96_863
S_Oscillospiraceae_multigenus_multispecies_sppn1_2_nov_93_676
S_Lachnospiraceae_multigenus_multispecies_sppn74_2_nov_94_488
G_Lachnospiraceae__G_1_
S_Lachnospiraceae__G_1__bacterium_MOT_166
S_Longibaculum_muris_nov_87_109
S_Eubacteriales__G_4__bacterium_MOT_165_nov_95_276
S_Roseburia_intestinalis_nov_92_913
S_Roseburia_multispecies_sppn21_2_nov_95_276
S_Merdimonas_faecis_nov_96_457
S_Lachnoclostridium__Clostridium_scindens_nov_91_406
S_Faecalicatena_orotica_nov_95_276
S_Lachnoclostridium__Clostridium_scindens_nov_93_725
S_Ileibacterium_valens
G_Ileibacterium
−5.124
−4.064
−3.004
−1.944
−0.884
0.176
1.236
2.296
3.356
4.416
LDA SCORE (log 10)