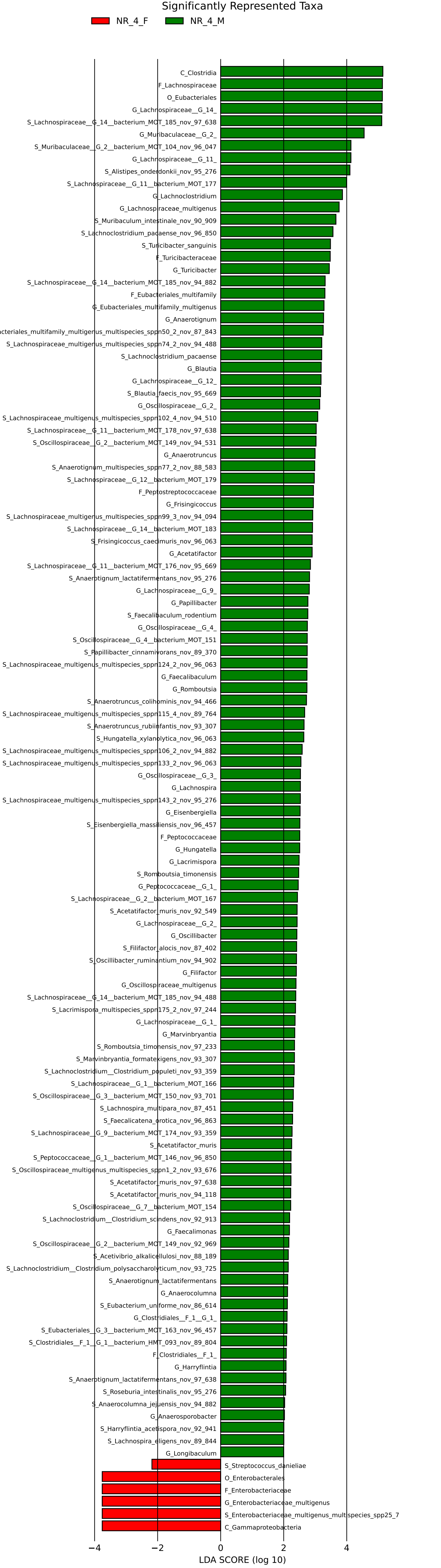
Significantly Represented Taxa
NR_4_F
NR_4_M
C_Clostridia
F_Lachnospiraceae
O_Eubacteriales
G_Lachnospiraceae__G_14_
S_Lachnospiraceae__G_14__bacterium_MOT_185_nov_97_638
G_Muribaculaceae__G_2_
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
G_Lachnospiraceae__G_11_
S_Alistipes_onderdonkii_nov_95_276
S_Lachnospiraceae__G_11__bacterium_MOT_177
G_Lachnoclostridium
G_Lachnospiraceae_multigenus
S_Muribaculum_intestinale_nov_90_909
S_Lachnoclostridium_pacaense_nov_96_850
S_Turicibacter_sanguinis
F_Turicibacteraceae
G_Turicibacter
S_Lachnospiraceae__G_14__bacterium_MOT_185_nov_94_882
F_Eubacteriales_multifamily
G_Eubacteriales_multifamily_multigenus
G_Anaerotignum
acteriales_multifamily_multigenus_multispecies_sppn50_2_nov_87_843
S_Lachnospiraceae_multigenus_multispecies_sppn74_2_nov_94_488
S_Lachnoclostridium_pacaense
G_Blautia
G_Lachnospiraceae__G_12_
S_Blautia_faecis_nov_95_669
G_Oscillospiraceae__G_2_
S_Lachnospiraceae_multigenus_multispecies_sppn102_4_nov_94_510
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_638
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
G_Anaerotruncus
S_Anaerotignum_multispecies_sppn77_2_nov_88_583
S_Lachnospiraceae__G_12__bacterium_MOT_179
F_Peptostreptococcaceae
G_Frisingicoccus
S_Lachnospiraceae_multigenus_multispecies_sppn99_3_nov_94_094
S_Lachnospiraceae__G_14__bacterium_MOT_183
S_Frisingicoccus_caecimuris_nov_96_063
G_Acetatifactor
S_Lachnospiraceae__G_11__bacterium_MOT_176_nov_95_669
S_Anaerotignum_lactatifermentans_nov_95_276
G_Lachnospiraceae__G_9_
G_Papillibacter
S_Faecalibaculum_rodentium
G_Oscillospiraceae__G_4_
S_Oscillospiraceae__G_4__bacterium_MOT_151
S_Papillibacter_cinnamivorans_nov_89_370
S_Lachnospiraceae_multigenus_multispecies_sppn124_2_nov_96_063
G_Faecalibaculum
G_Romboutsia
S_Anaerotruncus_colihominis_nov_94_466
S_Lachnospiraceae_multigenus_multispecies_sppn115_4_nov_89_764
S_Anaerotruncus_rubiinfantis_nov_93_307
S_Hungatella_xylanolytica_nov_96_063
S_Lachnospiraceae_multigenus_multispecies_sppn106_2_nov_94_882
S_Lachnospiraceae_multigenus_multispecies_sppn133_2_nov_96_063
G_Oscillospiraceae__G_3_
G_Lachnospira
S_Lachnospiraceae_multigenus_multispecies_sppn143_2_nov_95_276
G_Eisenbergiella
S_Eisenbergiella_massiliensis_nov_96_457
F_Peptococcaceae
G_Hungatella
G_Lacrimispora
S_Romboutsia_timonensis
G_Peptococcaceae__G_1_
S_Lachnospiraceae__G_2__bacterium_MOT_167
S_Acetatifactor_muris_nov_92_549
G_Lachnospiraceae__G_2_
G_Oscillibacter
S_Filifactor_alocis_nov_87_402
S_Oscillibacter_ruminantium_nov_94_902
G_Filifactor
G_Oscillospiraceae_multigenus
S_Lachnospiraceae__G_14__bacterium_MOT_185_nov_94_488
S_Lacrimispora_multispecies_sppn175_2_nov_97_244
G_Lachnospiraceae__G_1_
G_Marvinbryantia
S_Romboutsia_timonensis_nov_97_233
S_Marvinbryantia_formatexigens_nov_93_307
S_Lachnoclostridium__Clostridium_populeti_nov_93_359
S_Lachnospiraceae__G_1__bacterium_MOT_166
S_Oscillospiraceae__G_3__bacterium_MOT_150_nov_93_701
S_Lachnospira_multipara_nov_87_451
S_Faecalicatena_orotica_nov_96_863
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_93_359
S_Acetatifactor_muris
S_Peptococcaceae__G_1__bacterium_MOT_146_nov_96_850
S_Oscillospiraceae_multigenus_multispecies_sppn1_2_nov_93_676
S_Acetatifactor_muris_nov_97_638
S_Acetatifactor_muris_nov_94_118
S_Oscillospiraceae__G_7__bacterium_MOT_154
S_Lachnoclostridium__Clostridium_scindens_nov_92_913
G_Faecalimonas
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_92_969
S_Acetivibrio_alkalicellulosi_nov_88_189
S_Lachnoclostridium__Clostridium_polysaccharolyticum_nov_93_725
S_Anaerotignum_lactatifermentans
G_Anaerocolumna
S_Eubacterium_uniforme_nov_86_614
G_Clostridiales__F_1__G_1_
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Clostridiales__F_1__G_1__bacterium_HMT_093_nov_89_804
F_Clostridiales__F_1_
G_Harryflintia
S_Anaerotignum_lactatifermentans_nov_97_638
S_Roseburia_intestinalis_nov_95_276
S_Anaerocolumna_jejuensis_nov_94_882
G_Anaerosporobacter
S_Harryflintia_acetispora_nov_92_941
S_Lachnospira_eligens_nov_89_844
G_Longibaculum
S_Streptococcus_danieliae
O_Enterobacterales
F_Enterobacteriaceae
G_Enterobacteriaceae_multigenus
S_Enterobacteriaceae_multigenus_multispecies_spp25_7
C_Gammaproteobacteria
−4
−2
0
2
4
LDA SCORE (log 10)