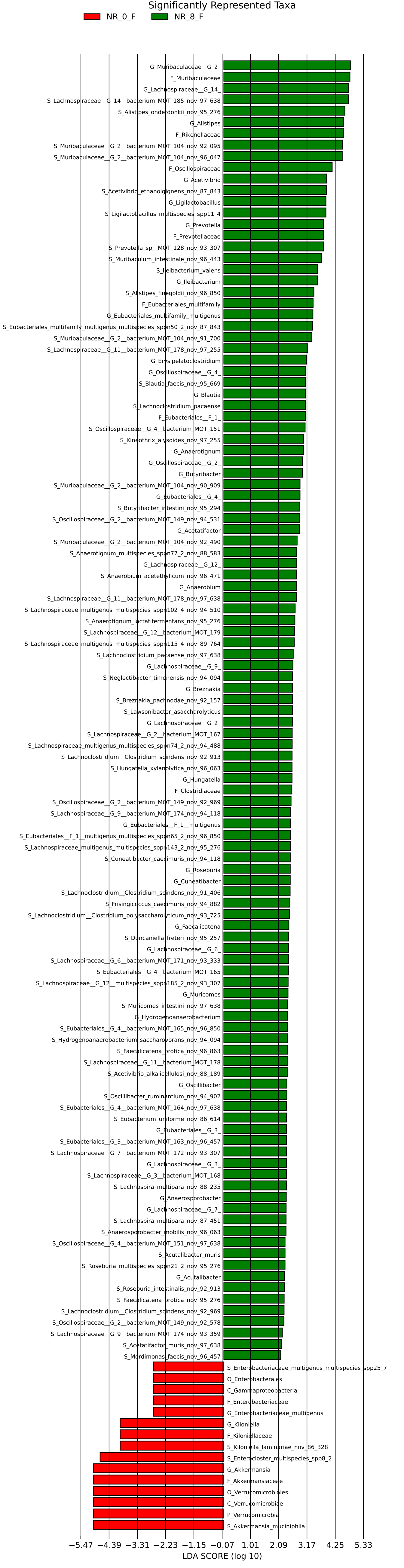
Significantly Represented Taxa
NR_0_F
NR_8_F
G_Muribaculaceae__G_2_
F_Muribaculaceae
G_Lachnospiraceae__G_14_
S_Lachnospiraceae__G_14__bacterium_MOT_185_nov_97_638
S_Alistipes_onderdonkii_nov_95_276
G_Alistipes
F_Rikenellaceae
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_095
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
F_Oscillospiraceae
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
G_Ligilactobacillus
S_Ligilactobacillus_multispecies_spp11_4
G_Prevotella
F_Prevotellaceae
S_Prevotella_sp__MOT_128_nov_93_307
S_Muribaculum_intestinale_nov_96_443
S_Ileibacterium_valens
G_Ileibacterium
S_Alistipes_finegoldii_nov_96_850
F_Eubacteriales_multifamily
G_Eubacteriales_multifamily_multigenus
S_Eubacteriales_multifamily_multigenus_multispecies_sppn50_2_nov_87_843
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_91_700
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_255
G_Erysipelatoclostridium
G_Oscillospiraceae__G_4_
S_Blautia_faecis_nov_95_669
G_Blautia
S_Lachnoclostridium_pacaense
F_Eubacteriales__F_1_
S_Oscillospiraceae__G_4__bacterium_MOT_151
S_Kineothrix_alysoides_nov_97_255
G_Anaerotignum
G_Oscillospiraceae__G_2_
G_Butyribacter
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_90_909
G_Eubacteriales__G_4_
S_Butyribacter_intestini_nov_95_294
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
G_Acetatifactor
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_490
S_Anaerotignum_multispecies_sppn77_2_nov_88_583
G_Lachnospiraceae__G_12_
S_Anaerobium_acetethylicum_nov_96_471
G_Anaerobium
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_638
S_Lachnospiraceae_multigenus_multispecies_sppn102_4_nov_94_510
S_Anaerotignum_lactatifermentans_nov_95_276
S_Lachnospiraceae__G_12__bacterium_MOT_179
S_Lachnospiraceae_multigenus_multispecies_sppn115_4_nov_89_764
S_Lachnoclostridium_pacaense_nov_97_638
G_Lachnospiraceae__G_9_
S_Neglectibacter_timonensis_nov_94_094
G_Breznakia
S_Breznakia_pachnodae_nov_92_157
S_Lawsonibacter_asaccharolyticus
G_Lachnospiraceae__G_2_
S_Lachnospiraceae__G_2__bacterium_MOT_167
S_Lachnospiraceae_multigenus_multispecies_sppn74_2_nov_94_488
S_Lachnoclostridium__Clostridium_scindens_nov_92_913
S_Hungatella_xylanolytica_nov_96_063
G_Hungatella
F_Clostridiaceae
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_92_969
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_94_118
G_Eubacteriales__F_1__multigenus
S_Eubacteriales__F_1__multigenus_multispecies_sppn65_2_nov_96_850
S_Lachnospiraceae_multigenus_multispecies_sppn143_2_nov_95_276
S_Cuneatibacter_caecimuris_nov_94_118
G_Roseburia
G_Cuneatibacter
S_Lachnoclostridium__Clostridium_scindens_nov_91_406
S_Frisingicoccus_caecimuris_nov_94_882
S_Lachnoclostridium__Clostridium_polysaccharolyticum_nov_93_725
G_Faecalicatena
S_Duncaniella_freteri_nov_95_257
G_Lachnospiraceae__G_6_
S_Lachnospiraceae__G_6__bacterium_MOT_171_nov_93_333
S_Eubacteriales__G_4__bacterium_MOT_165
S_Lachnospiraceae__G_12__multispecies_sppn185_2_nov_93_307
G_Muricomes
S_Muricomes_intestini_nov_97_638
G_Hydrogenoanaerobacterium
S_Eubacteriales__G_4__bacterium_MOT_165_nov_96_850
S_Hydrogenoanaerobacterium_saccharovorans_nov_94_094
S_Faecalicatena_protica_nov_96_863
S_Lachnospiraceae__G_11__bacterium_MOT_178
S_Acetivibrio_alkalicellulosi_nov_88_189
G_Oscillibacter
S_Oscillibacter_ruminantium_nov_94_902
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_638
S_Eubacterium_uniforme_nov_86_614
G_Eubacteriales__G_3_
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Lachnospiraceae__G_7__bacterium_MOT_172_nov_93_307
G_Lachnospiraceae__G_3_
S_Lachnospiraceae__G_3__bacterium_MOT_168
S_Lachnospira_multipara_nov_88_235
G_Anaerosporobacter
G_Lachnospiraceae__G_7_
S_Lachnospira_multipara_nov_87_451
S_Anaerosporobacter_mobilis_nov_96_063
S_Oscillospiraceae__G_4__bacterium_MOT_151_nov_97_638
S_Acutalibacter_muris
S_Roseburia_multispecies_sppn21_2_nov_95_276
G_Acutalibacter
S_Roseburia_intestinalis_nov_92_913
S_Faecalicatena_protica_nov_95_276
S_Lachnoclostridium__Clostridium_scindens_nov_92_969
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_92_578
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_93_359
S_Acetatifactor_muris_nov_97_638
S_Merdimonas_faecis_nov_96_457
S_Enterobacteriaceae_multigenus_multispecies_spp25_7
O_Enterobacterales
C_Gammaproteobacteria
F_Enterobacteriaceae
G_Enterobacteriaceae_multigenus
G_Kiloniella
F_Kiloniellaceae
S_Kiloniella_laminariae_nov_86_328
S_Enterocloster_multispecies_spp8_2
G_Akkermansia
F_Akkermansiaceae
O_Verrucomicrobiales
C_Verrucomicrobiae
P_Verrucomicrobia
S_Akkermansia_muciniphila
−5.47
−4.39
−3.31
−2.23
−1.15
−0.07
1.01
2.09
3.17
4.25
5.33
LDA SCORE (log 10)