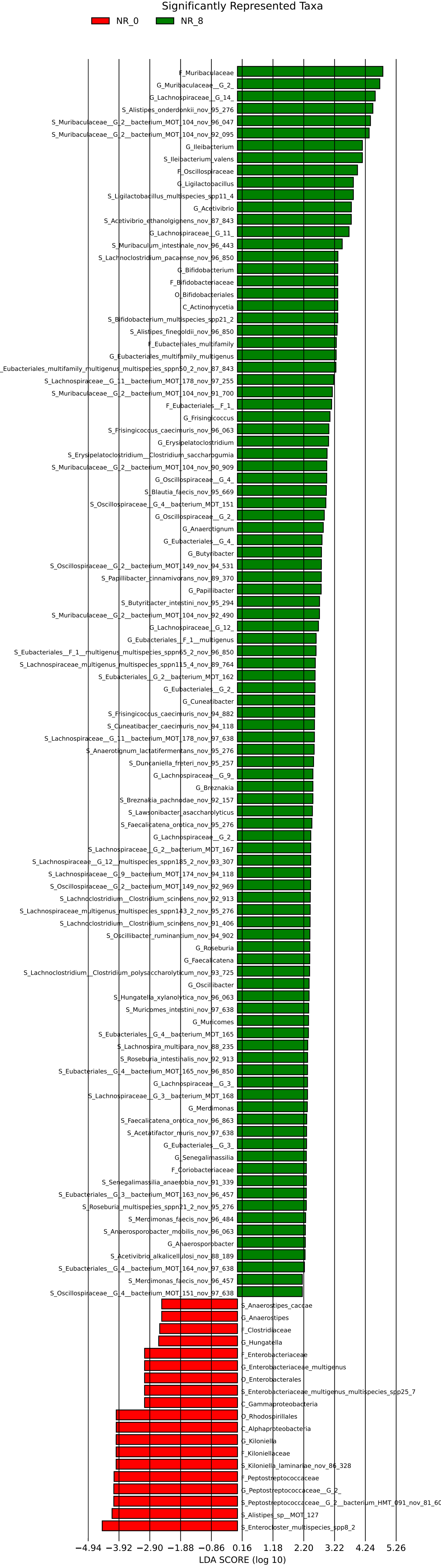
Significantly Represented Taxa
NR_0
NR_8
F_Muribaculaceae
G_Muribaculaceae__G_2_
G_Lachnospiraceae__G_14_
S_Alistipes_onderdonkii_nov_95_276
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_095
G_Ileibacterium
S_Ileibacterium_valens
F_Oscillospiraceae
G_Ligilactobacillus
S_Ligilactobacillus_multispecies_spp11_4
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
G_Lachnospiraceae__G_11_
S_Muribaculum_intestinale_nov_96_443
S_Lachnoclostridium_pacaense_nov_96_850
G_Bifidobacterium
F_Bifidobacteriaceae
O_Bifidobacteriales
C_Actinomycetia
S_Bifidobacterium_multispecies_spp21_2
S_Alistipes_finegoldii_nov_96_850
F_Eubacteriales_multifamily
G_Eubacteriales_multifamily_multigenus
_Eubacteriales_multifamily_multigenus_multispecies_sppn50_2_nov_87_843
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_255
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_91_700
F_Eubacteriales__F_1_
G_Frisingicoccus
S_Frisingicoccus_caecimuris_nov_96_063
G_Erysipelatoclostridium
S_Erysipelatoclostridium__Clostridium_saccharogumia
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_90_909
G_Oscillospiraceae__G_4_
S_Blautia_faecis_nov_95_669
S_Oscillospiraceae__G_4__bacterium_MOT_151
G_Oscillospiraceae__G_2_
G_Anaerotignum
G_Eubacteriales__G_4_
G_Butyribacter
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
S_Papillibacter_cinnamivorans_nov_89_370
G_Papillibacter
S_Butyribacter_intestini_nov_95_294
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_490
G_Lachnospiraceae__G_12_
G_Eubacteriales__F_1__multigenus
S_Eubacteriales__F_1__multigenus_multispecies_sppn65_2_nov_96_850
S_Lachnospiraceae_multigenus_multispecies_sppn115_4_nov_89_764
S_Eubacteriales__G_2__bacterium_MOT_162
G_Eubacteriales__G_2_
G_Cuneatibacter
S_Frisingicoccus_caecimuris_nov_94_882
S_Cuneatibacter_caecimuris_nov_94_118
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_638
S_Anaerotignum_lactatifermentans_nov_95_276
S_Duncaniella_freteri_nov_95_257
G_Lachnospiraceae__G_9_
G_Breznakia
S_Breznakia_pachnodae_nov_92_157
S_Lawsonibacter_asaccharolyticus
S_Faecalicatena_orotica_nov_95_276
G_Lachnospiraceae__G_2_
S_Lachnospiraceae__G_2__bacterium_MOT_167
S_Lachnospiraceae__G_12__multispecies_sppn185_2_nov_93_307
S_Lachnospiraceae__G_9__bacterium_MOT_174_nov_94_118
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_92_969
S_Lachnoclostridium__Clostridium_scindens_nov_92_913
S_Lachnospiraceae_multigenus_multispecies_sppn143_2_nov_95_276
S_Lachnoclostridium__Clostridium_scindens_nov_91_406
S_Oscillibacter_ruminantium_nov_94_902
G_Roseburia
G_Faecalicatena
S_Lachnoclostridium__Clostridium_polysaccharolyticum_nov_93_725
G_Oscillibacter
S_Hungatella_xylanolytica_nov_96_063
S_Muricomes_intestini_nov_97_638
G_Muricomes
S_Eubacteriales__G_4__bacterium_MOT_165
S_Lachnospira_multipara_nov_88_235
S_Roseburia_intestinalis_nov_92_913
S_Eubacteriales__G_4__bacterium_MOT_165_nov_96_850
G_Lachnospiraceae__G_3_
S_Lachnospiraceae__G_3__bacterium_MOT_168
G_Merdimonas
S_Faecalicatena_orotica_nov_96_863
S_Acetatifactor_muris_nov_97_638
G_Eubacteriales__G_3_
G_Senegalimassilia
F_Coriobacteriaceae
S_Senegalimassilia_anaerobia_nov_91_339
S_Eubacteriales__G_3__bacterium_MOT_163_nov_96_457
S_Roseburia_multispecies_sppn21_2_nov_95_276
S_Merdimonas_faecis_nov_96_484
S_Anaerosporobacter_mobilis_nov_96_063
G_Anaerosporobacter
S_Acetivibrio_alkalicellulosi_nov_88_189
S_Eubacteriales__G_4__bacterium_MOT_164_nov_97_638
S_Merdimonas_faecis_nov_96_457
S_Oscillospiraceae__G_4__bacterium_MOT_151_nov_97_638
S_Anaerostipes_caccae
G_Anaerostipes
F_Clostridiaceae
G_Hungatella
F_Enterobacteriaceae
G_Enterobacteriaceae_multigenus
O_Enterobacterales
S_Enterobacteriaceae_multigenus_multispecies_spp25_7
C_Gammaproteobacteria
O_Rhodospirillales
C_Alphaproteobacteria
G_Kiloniella
F_Kiloniellaceae
S_Kiloniella_laminariae_nov_86_328
F_Peptostreptococcaceae
G_Peptostreptococcaceae__G_2_
S_Peptostreptococcaceae__G_2__bacterium_HMT_091_nov_81_60
S_Alistipes_sp__MOT_127
S_Enterocloster_multispecies_spp8_2
−4.94
−3.92
−2.90
−1.88
−0.86
0.16
1.18
2.20
3.22
4.24
5.26
LDA SCORE (log 10)