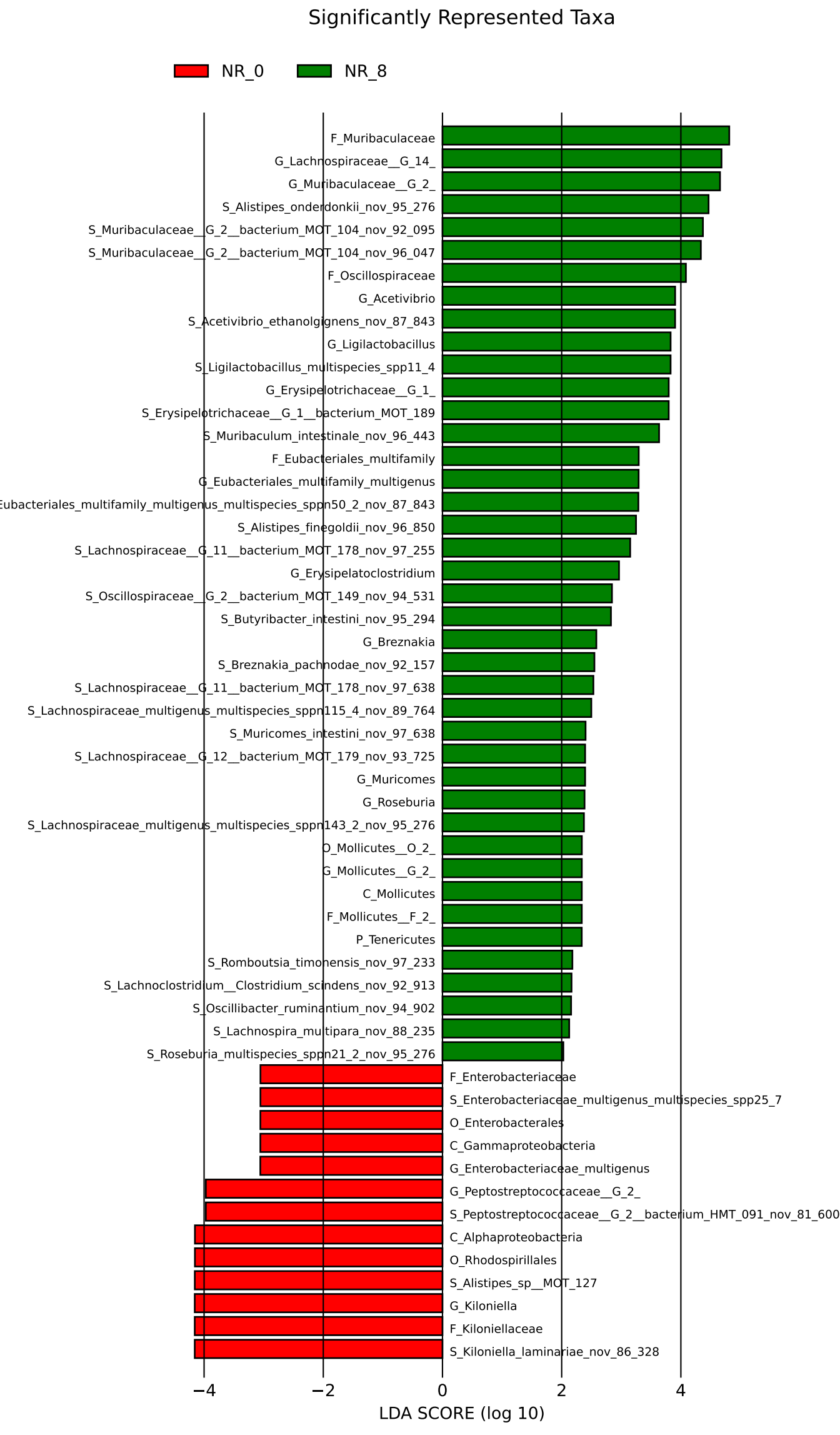
Significantly Represented Taxa
NR_0
NR_8
F_Muribaculaceae
G_Lachnospiraceae__G_14_
G_Muribaculaceae__G_2_
S_Alistipes_onderdonkii_nov_95_276
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_92_095
S_Muribaculaceae__G_2__bacterium_MOT_104_nov_96_047
F_Oscillospiraceae
G_Acetivibrio
S_Acetivibrio_ethanolgignens_nov_87_843
G_Ligilactobacillus
S_Ligilactobacillus_multispecies_spp11_4
G_Erysipelotrichaceae__G_1_
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
S_Muribaculum_intestinale_nov_96_443
F_Eubacteriales_multifamily
G_Eubacteriales_multifamily_multigenus
Eubacteriales_multifamily_multigenus_multispecies_sppn50_2_nov_87_843
S_Alistipes_finegoldii_nov_96_850
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_255
G_Erysipelatoclostridium
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
S_Butyribacter_intestini_nov_95_294
G_Breznakia
S_Breznakia_pachnodae_nov_92_157
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_638
S_Lachnospiraceae_multigenus_multispecies_sppn115_4_nov_89_764
S_Muricomes_intestini_nov_97_638
S_Lachnospiraceae__G_12__bacterium_MOT_179_nov_93_725
G_Muricomes
G_Roseburia
S_Lachnospiraceae_multigenus_multispecies_sppn143_2_nov_95_276
O_Mollicutes__O_2_
G_Mollicutes__G_2_
C_Mollicutes
F_Mollicutes__F_2_
P_Tenericutes
S_Romboutsia_timonensis_nov_97_233
S_Lachnoclostridium__Clostridium_scindens_nov_92_913
S_Oscillibacter_ruminantium_nov_94_902
S_Lachnospira_multipara_nov_88_235
S_Roseburia_multispecies_sppn21_2_nov_95_276
F_Enterobacteriaceae
S_Enterobacteriaceae_multigenus_multispecies_spp25_7
O_Enterobacterales
C_Gammaproteobacteria
G_Enterobacteriaceae_multigenus
G_Peptostreptococcaceae__G_2_
S_Peptostreptococcaceae__G_2__bacterium_HMT_091_nov_81_600
C_Alphaproteobacteria
O_Rhodospirillales
S_Alistipes_sp__MOT_127
G_Kiloniella
F_Kiloniellaceae
S_Kiloniella_laminariae_nov_86_328
−4
−2
0
2
4
LDA SCORE (log 10)