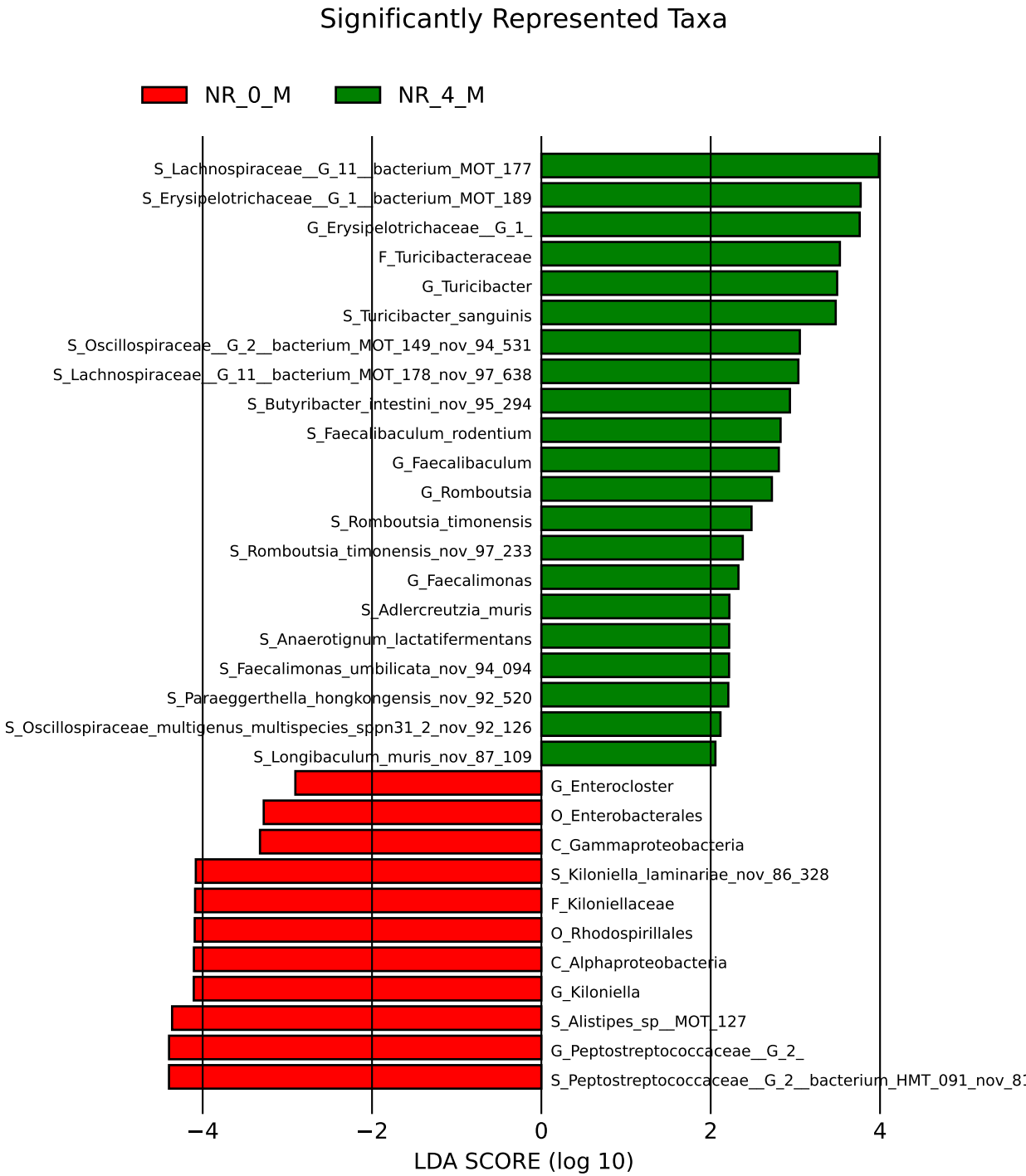
Significantly Represented Taxa
NR_0_M
NR_4_M
S_Lachnospiraceae__G_11__bacterium_MOT_177
S_Erysipelotrichaceae__G_1__bacterium_MOT_189
G_Erysipelotrichaceae__G_1_
F_Turicibacteraceae
G_Turicibacter
S_Turicibacter_sanguinis
S_Oscillospiraceae__G_2__bacterium_MOT_149_nov_94_531
S_Lachnospiraceae__G_11__bacterium_MOT_178_nov_97_638
S_Butyribacter_intestini_nov_95_294
S_Faecalibaculum_rodentium
G_Faecalibaculum
G_Romboutsia
S_Romboutsia_timonensis
S_Romboutsia_timonensis_nov_97_233
G_Faecalimonas
S_Adlercreutzia_muris
S_Anaerotignum_lactatifermentans
S_Faecalimonas_umbilicata_nov_94_094
S_Paraeggerthella_hongkongensis_nov_92_520
S_Oscillospiraceae_multigenus_multispecies_sppn31_2_nov_92_126
S_Longibaculum_muris_nov_87_109
G_Enterocloster
O_Enterobacterales
C_Gammaproteobacteria
S_Kiloniella_laminariae_nov_86_328
F_Kiloniellaceae
O_Rhodospirillales
C_Alphaproteobacteria
G_Kiloniella
S_Alistipes_sp__MOT_127
G_Peptostreptococcaceae__G_2_
S_Peptostreptococcaceae__G_2__bacterium_HMT_091_nov_8
−4
−2
0
2
4
LDA SCORE (log 10)