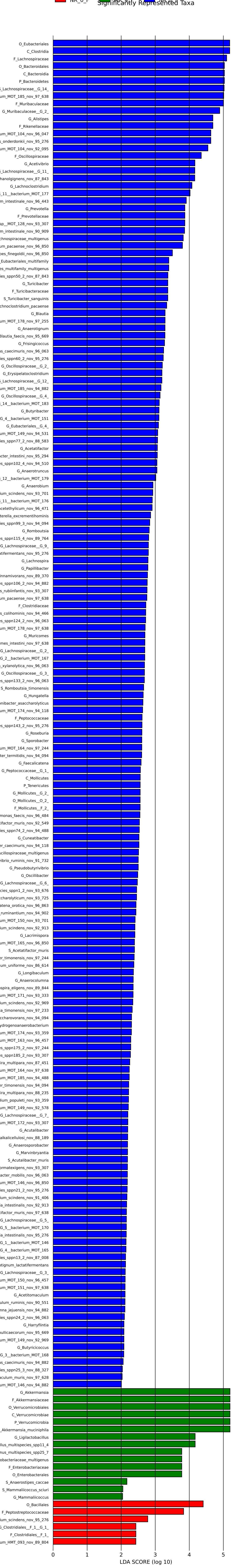
Significantly Represented Taxa
O_Eubacteriales
C_Clostridia
F_Lachnospiraceae
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
G_Lachnospiraceae__G_14_
um_MOT_185_nov_97_638
F_Muribaculaceae
G_Muribaculaceae__G_2_
G_Alistipes
F_Rikenellaceae
um_MOT_104_nov_96_047
s_onderdonkii_nov_95_276
um_MOT_104_nov_92_095
F_Oscillospiraceae
G_Acetivibrio
G_Lachnospiraceae__G_11_
hanolgignens_nov_87_843
G_Lachnoclostridium
G_11__bacterium_MOT_177
m_intestinale_nov_96_443
G_Prevotella
F_Prevotellaceae
sp__MOT_128_nov_93_307
m_intestinale_nov_90_909
chnospiraceae_multigenus
um_pacaense_nov_96_850
pes_finegoldii_nov_96_850
_Eubacteriales_multifamily
es_multifamily_multigenus
ies_sppn50_2_nov_87_843
G_Turicibacter
F_Turicibacteraceae
S_Turicibacter_sanguinis
chnoclostridium_pacaense
G_Blautia
um_MOT_178_nov_97_255
G_Anaerotignum
Blautia_faecis_nov_95_669
G_Frisingicoccus
us_caecimuris_nov_96_063
ies_sppn60_2_nov_95_276
G_Oscillospiraceae__G_2_
G_Erysipelatoclostridium
G_Lachnospiraceae__G_12_
um_MOT_185_nov_94_882
G_Oscillospiraceae__G_4_
G_14__bacterium_MOT_183
G_Butyribacter
G_4__bacterium_MOT_151
G_Eubacteriales__G_4_
um_MOT_149_nov_94_531
ies_sppn77_2_nov_88_583
G_Acetatifactor
acter_intestini_nov_95_294
es_sppn102_4_nov_94_510
G_Anaerotruncus
G_12__bacterium_MOT_179
G_Anaerobium
ium_scindens_nov_93_701
G_11__bacterium_MOT_176
acetethylicum_nov_96_471
terella_excrementihominis
ies_sppn99_3_nov_94_094
G_Romboutsia
es_sppn115_4_nov_89_764
G_Lachnospiraceae__G_9_
atifermentans_nov_95_276
G_Lachnospira
G_Papillibacter
innamivorans_nov_89_370
es_sppn106_2_nov_94_882
s_rubiinfantis_nov_93_307
um_pacaense_nov_97_638
F_Clostridiaceae
s_colihominis_nov_94_466
es_sppn124_2_nov_96_063
um_MOT_178_nov_97_638
G_Muricomes
mes_intestini_nov_97_638
G_Lachnospiraceae__G_2_
G_2__bacterium_MOT_167
_xylanolytica_nov_96_063
G_Oscillospiraceae__G_3_
es_sppn133_2_nov_96_063
S_Romboutsia_timonensis
G_Hungatella
onibacter_asaccharolyticus
um_MOT_174_nov_94_118
F_Peptococcaceae
es_sppn143_2_nov_95_276
G_Roseburia
G_Sporobacter
um_MOT_164_nov_97_244
ter_termitidis_nov_94_094
G_Faecalicatena
G_Peptococcaceae__G_1_
C_Mollicutes
P_Tenericutes
G_Mollicutes__G_2_
O_Mollicutes__O_2_
F_Mollicutes__F_2_
monas_faecis_nov_96_484
tifactor_muris_nov_92_549
ies_sppn74_2_nov_94_488
G_Cuneatibacter
er_caecimuris_nov_94_118
scillospiraceae_multigenus
vibrio_ruminis_nov_91_732
G_Pseudobutyrivibrio
G_Oscillibacter
G_Lachnospiraceae__G_6_
cies_sppn1_2_nov_93_676
ccharolyticum_nov_93_725
atena_orotica_nov_96_863
_ruminantium_nov_94_902
um_MOT_150_nov_93_701
ium_scindens_nov_92_913
G_Lacrimispora
um_MOT_165_nov_96_850
S_Acetatifactor_muris
er_timonensis_nov_97_244
um_uniforme_nov_86_614
G_Longibaculum
G_Anaerocolumna
spira_eligens_nov_89_844
um_MOT_171_nov_93_333
ium_scindens_nov_92_969
ia_timonensis_nov_97_233
ccharovorans_nov_94_094
ydrogenoanaerobacterium
um_MOT_174_nov_93_359
um_MOT_163_nov_96_457
es_sppn175_2_nov_97_244
es_sppn185_2_nov_93_307
ira_multipara_nov_87_451
um_MOT_164_nov_97_638
um_MOT_185_nov_94_488
er_timonensis_nov_94_094
ira_multipara_nov_88_235
dium_populeti_nov_93_359
um_MOT_149_nov_92_578
G_Lachnospiraceae__G_7_
um_MOT_172_nov_93_307
G_Acutalibacter
alkalicellulosi_nov_88_189
G_Anaerosporobacter
G_Marvinbryantia
S_Acutalibacter_muris
ormatexigens_nov_93_307
acter_mobilis_nov_96_063
um_MOT_146_nov_96_850
ies_sppn21_2_nov_95_276
ium_scindens_nov_91_406
ia_intestinalis_nov_92_913
tifactor_muris_nov_97_638
G_Lachnospiraceae__G_5_
G_5__bacterium_MOT_170
ia_intestinalis_nov_95_276
G_1__bacterium_MOT_146
G_4__bacterium_MOT_165
ies_sppn13_2_nov_87_008
otignum_lactatifermentans
G_Lachnospiraceae__G_3_
um_MOT_150_nov_96_457
um_MOT_151_nov_97_638
G_Acetitomaculum
ulum_ruminis_nov_90_551
nna_jejuensis_nov_94_882
ies_sppn24_2_nov_96_063
G_Harryflintia
ullicaecorum_nov_95_669
um_MOT_149_nov_92_969
G_Butyricicoccus
G_3__bacterium_MOT_168
us_caecimuris_nov_94_882
ies_sppn25_3_nov_88_327
aculum_muris_nov_97_628
um_MOT_146_nov_94_882
G_Akkermansia
F_Akkermansiaceae
O_Verrucomicrobiales
C_Verrucomicrobiae
P_Verrucomicrobia
_Akkermansia_muciniphila
G_Ligilactobacillus
llus_multispecies_spp11_4
nus_multispecies_spp25_7
obacteriaceae_multigenus
F_Enterobacteriaceae
O_Enterobacterales
S_Anaerostipes_caccae
S_Mammaliicoccus_sciuri
G_Mammaliicoccus
O_Bacillales
F_Peptostreptococcaceae
ium_scindens_nov_95_276
G_Clostridiales__F_1__G_1_
F_Clostridiales__F_1_
um_HMT_093_nov_89_804
0
1
2
3
4
5
LDA SCORE (log 10)