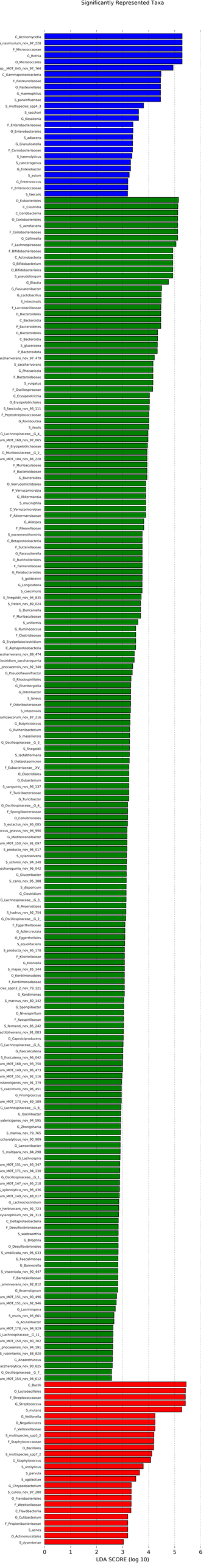
Significantly Represented Taxa
C_Actinomycetia
s_nasimurium_nov_97_228
F_Micrococcaceae
G_Rothia
O_Micrococcales
ho_MOT_045_nov_97_764
C_Gammaproteobacteria
F_Pasteurellaceae
O_Pasteurellales
O_Haemophilus
S_parainfluenzae
S_multispecies_spp4_3
S_sacchari
G_Kosakonia
F_Enterobacteriaceae
O_Enterobacterales
S_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_haemolyticus
S_cancerogenus
G_Enterobacter
S_avium
G_Enterococcus
F_Enterococcaceae
S_faecalis
O_Eubacteriales
C_Clostridia
C_Coriobacteria
O_Coriobacteriales
S_aerofaciens
F_Coriobacteriaceae
G_Collinsella
F_Lachnospiraceae
F_Bifidobacteriaceae
C_Actinobacteria
G_Bifidobacterium
O_Bifidobacteriales
S_pseudolongum
G_Blautia
G_Fusicatenibacter
G_Lactobacillus
S_intestinalis
F_Lactobacillaceae
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
S_glucerasea
P_Bacteroidota
cchariivorans_nov_97_479
S_saccharivorans
G_Phocaeicola
F_Bacteroidaceae
S_vulgatus
F_Oscillospiraceae
C_Erysipelotrichia
O_Erysipelotrichales
S_Tseacisia_nov_93_111
F_Peptostreptococcaceae
G_Romboutsia
S_ilealis
G_Lachnospiraceae__G_4_
um_MOT_169_nov_97_065
F_Erysipelotrichaceae
G_Muribaculaceae__G_2_
um_MOT_104_nov_96_228
F_Muribaculaceae
F_Bacteroidaceae
G_Bacteroides
O_Verrucomicrobiales
P_Verrucomicrobia
G_Akkermansia
S_muciniphila
C_Verrucomicrobiae
F_Akkermansiaceae
G_Alistipes
F_Rikenellaceae
S_excrementihominis
C_Betaproteobacteria
F_Sutterellaceae
G_Parasutterella
O_Burkholderiales
F_Tannerellaceae
G_Parabacteroides
S_goldsteinii
G_Longicatena
S_caecimuris
S_finegoldii_nov_94_835
S_fretri_nov_89_024
G_Duncaniella
F_Muribaculaceae
S_uniformis
G_Ruminococcus
F_Clostridiaceae
G_Erysipelatoclostridium
C_Alphaproteobacteria
cchariivorans_nov_89_474
lostridium_saccharogumia
phocaeensis_nov_92_340
G_Pseudoflavonifractor
O_Rhodospirillales
G_Eisenbergiella
G_Odoribacter
S_laneus
F_Odoribacteriaceae
S_intestinalis
ulicaecorum_nov_87_216
G_Butyricicoccus
G_Ruthenibacterium
S_massiliensis
G_Oscillospiraceae__G_3_
S_finegoldii
S_lactaltifermans
S_thetaiotaomicron
F_Eubacteriaceae__XV_
O_Clostridiales
G_Eubacterium
S_sanguinis_nov_96_137
F_Turicibacteraceae
G_Turicibacter
G_Oscillospiraceae__G_4_
F_Spongibacteraceae
O_Cellvibrionales
S_eutactus_nov_95_085
ccus_gnavus_nov_94_990
G_Mediterraneibacter
um_MOT_150_nov_91_097
S_producta_nov_96_017
S_xylanisolvens
S_schinkii_nov_94_340
cchanguyna_nov_96_042
G_Glucerbacter
S_canis_nov_95_388
S_disporicum
G_Clostridium
G_Lachnospiraceae__G_3_
G_Anaerostipes
S_hadrus_nov_92_754
G_Oscillospiraceae__G_2_
F_Eggerthellaceae
G_Adlercreutzia
O_Eggerthellales
S_equolifaciens
S_producta_nov_95_178
F_Klosiellaceae
G_Klosiella
S_majae_nov_85_144
O_Kordiimonadales
F_Kordiimonadaceae
cies_spp13_2_nov_79_121
G_Kordiimonas
S_marinus_nov_80_142
F_Azospirillaceae
S_fermenti_nov_85_242
acitoliivorans_nov_91_083
G_Caproicipro
G_Lachnospiraceae__G_6_
G_Faecalicatena
S_fissicatena_nov_96_042
um_MOT_168_nov_93_750
um_MOT_149_nov_96_473
um_MOT_151_nov_92_116
stanoliigenes_nov_91_379
S_caecimuris_nov_96_451
G_Frisingicoccus
um_MOT_173_nov_89_189
G_Lachnospiraceae__G_8_
G_Oscillibacter
valericigenes_nov_94_595
G_Zhongshania
S_marina_nov_79_765
cchariolyticus_nov_90_909
G_Lawsonibacter
S_multipara_nov_84_298
G_Lachnospira
um_MOT_151_nov_93_347
um_MOT_171_nov_94_130
G_Oscillospiraceae__G_1_
um_MOT_147_nov_95_218
_xylanolytica_nov_96_436
um_MOT_149_nov_88_017
G_Lachnoclostridium
G_hellriovaria_nov_92_723
xylanophilum_nov_91_313
C_Deltaproteobacteria
F_Desulfovibrionaceae
S_wadsworthia
G_Bilophila
O_Desulfovibrionales
S_umbilicata_nov_96_033
G_Faecalimonas
G_Barnesiella
S_viscericola_nov_90_447
F_Barnesiellaceae
_aminivorans_nov_92_812
G_Anaerotignum
um_MOT_151_nov_90_496
um_MOT_151_nov_92_946
G_Lacrimispora
S_muris_nov_95_661
G_Acutalibacter
um_MOT_178_nov_94_929
_Lachnospiraceae__G_11_
um_MOT_150_nov_90_702
phocaeensis_nov_94_191
S_rubinfantis_nov_88_820
G_Anaerotruncus
cchariolytica_nov_90_625
G_Oscillospiraceae__G_7_
um_MOT_154_nov_94_612
C_Bacilli
O_Lactobacillales
F_Streptococcaceae
G_Streptococcus
S_mutans
G_Veillonella
O_Negativicutes
F_Veillonellaceae
S_multispecies_spp5_2
F_Staphylococcaceae
O_Bacillales
S_multispecies_spp7_2
G_Staphylococcus
S_urealyticus
S_parvula
S_agalactiae
G_Chryseobacterium
S_culicis_nov_97_280
O_Flavobacteriales
F_Weeksellaceae
C_Flavobacteria
G_Cutibacterium
F_Propionibacteriaceae
S_acnes
O_Actinomycetales
S_dysenteriae
0
1
2
3
4
5
6
LDA SCORE (log 10)