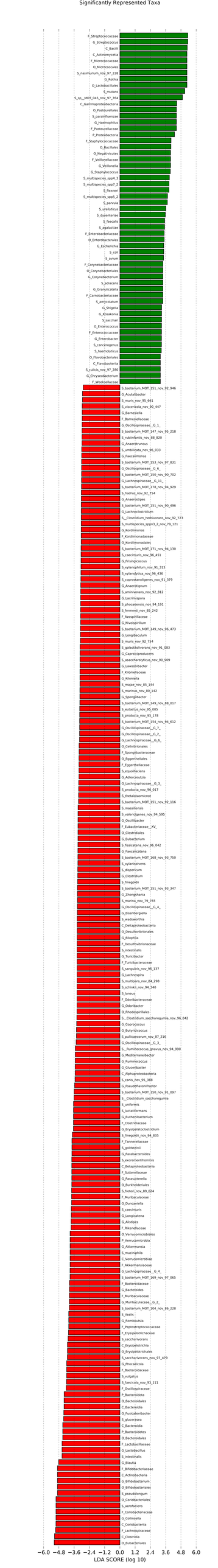
Significantly Represented Taxa
F_Streptococcaceae
G_Streptococcus
C_Bacilli
C_Actinomycetia
F_Micrococcaceae
O_Micrococcales
S_nasimurium_nov_97_228
G_Rothia
O_Lactobacillales
S_mutans
S_sp._M07_045_nov_97_764
C_Gammaproteobacteria
O_Pasteurellales
S_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
P_Proteobacteria
F_Staphylococcaceae
O_Bacillales
O_Negativicutes
F_Veillonellaceae
G_Veillonella
G_Staphylococcus
S_multispecies_spp4_3
S_multispecies_spp7_2
S_flexneri
S_multispecies_spp5_2
S_parvula
S_ureilyticus
S_dysenteriae
S_faecalis
S_agalactiae
F_Enterobacteriaceae
O_Enterobacterales
G_Escherichia
S_coli
S_avium
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
S_adiacens
G_Granulicatella
F_Carnobacteriaceae
S_amycolatum
G_Shigella
G_Kosakonia
S_sacchari
G_Enterococcus
F_Enterococcaceae
G_Enterobacter
S_cancerogenus
S_haemolyticus
O_Flavobacteriales
C_Flavobacteria
S_culicis_nov_97_280
G_Chryseobacterium
F_Weeksellaceae
S_bacterium_MOT_151_nov_92_946
G_Acutalibacter
S_muris_nov_95_661
S_viscericola_nov_90_447
G_Barnesiella
F_Barnesiellaceae
G_Oscillospiraceae_G_1_
S_bacterium_MOT_147_nov_95_218
S_rubinfantis_nov_88_820
G_Anaerotruncus
S_umbilicata_nov_86_033
G_Faecalimonas
S_bacterium_MOT_153_nov_97_831
G_Oscillospiraceae_G_6_
S_bacterium_MOT_150_nov_90_702
G_Lachnospiraceae_G_11_
S_bacterium_MOT_178_nov_94_929
S_hadrus_nov_92_754
G_Anaerostipes
S_bacterium_MOT_151_nov_90_496
G_Lachnoclostridium
S_Clostridium_nebriovarans_nov_92_723
S_multispecies_spp3_2_nov_79_121
G_Kordiimonas
F_Kordiimonadaceae
O_Kordiimonadales
S_bacterium_MOT_171_nov_94_130
S_caecimuris_nov_96_451
G_Frisingicoccus
S_xylanophilum_nov_91_313
S_xylanolytica_nov_96_436
S_coprostanoligenes_nov_91_379
G_Anaerotignum
S_aminivorans_nov_92_812
G_Lacrimispora
S_phocaeensis_nov_94_191
S_fermentii_nov_85_242
F_Acrospirillaceae
G_Niveispirillum
S_bacterium_MOT_149_nov_96_473
G_Longibaculum
S_muris_nov_92_754
S_galactitolivorans_nov_91_083
G_Caproiciproducens
S_saacharolyticus_nov_90_909
G_Lawsonibacter
F_Kiloniellaceae
G_Kiloniella
S_majae_nov_85_144
S_marinis_nov_80_142
G_Spongibacter
S_bacterium_MOT_149_nov_88_017
S_eutactus_nov_95_085
S_pirdusta_nov_95_178
S_bacterium_MOT_154_nov_94_612
G_Oscillospiraceae_G_7_
G_Oscillospiraceae_G_2_
G_Lachnospiraceae_G_8_
O_Cellvibrionales
F_Spongibacteraceae
O_Eggerthellales
F_Eggerthellaceae
S_equolifaciens
G_Adlercreutzia
G_Lachnospiraceae_G_3_
S_producta_nov_96_017
S_thetaiotaomicron
S_bacterium_MOT_151_nov_92_116
S_massiliensis
S_valericigenes_nov_94_595
G_Oscillibacter
F_Eubacteriaceae_XV_
O_Clostridiales
G_Eubacterium
S_fissicatena_nov_96_042
G_Faecalicatena
S_bacterium_MOT_168_nov_93_750
S_xylanisolvens
S_disporicum
G_Clostridium
S_finegoldii
S_bacterium_MOT_151_nov_93_347
G_Zhongshania
S_marina_nov_79_765
G_Oscillospiraceae_G_4_
G_Eisenbergiella
S_wadsworthia
C_Deltaproteobacteria
O_Desulfovibrionales
G_Bilophila
F_Desulfovibrionaceae
S_intestinalis
G_Turicibacter
F_Turicibacteraceae
S_sanguinis_nov_96_137
G_Lachnospira
S_multipara_nov_84_298
S_schinki_nov_94_340
S_taneus
F_Odoribacteraceae
G_Odoribacter
O_Rhodospirillales
S_Clostridium_saccharogumia_nov_96_042
G_Coprococcus
G_Butyricicoccus
S_pullicaecorum_nov_87_216
G_Oscillospiraceae_G_3_
S_Ruminococcus_gravus_nov_94_990
G_Mediterraneibacter
G_Ruminococcus
G_Gluceilibacter
C_Alphaproteobacteria
S_canis_nov_95_388
G_Pseudoflavonifractor
S_bacterium_MOT_150_nov_91_097
S_Clostridium_saccharogumia
S_uniformis
S_lactaltiformans
G_Ruthenibacterium
F_Clostridiaceae
G_Erysipelatoclostridium
S_finegoldii_nov_94_835
F_Tannerellaceae
S_siddhiensis
G_Parabacteroides
S_excrementihominis
C_Betaproteobacteria
F_Sutterellaceae
G_Parasutterella
O_Burkholderiales
S_freteri_nov_89_924
F_Muribaculaceae
G_Duncaniella
S_caecimuris
G_Longicatena
G_Alistipes
F_Rikenellaceae
O_Verrucomicrobiales
P_Verrucomicrobia
G_Akkermansia
S_muciniphila
C_Verrucomicrobiae
F_Akkermansiaceae
G_Lachnospiraceae_G_4_
S_bacterium_MOT_169_nov_97_065
F_Bacteroidaceae
G_Bacteroides
F_Muribaculaceae
G_Muribaculaceae_G_2_
S_bacterium_MOT_104_nov_86_228
S_ilealis
G_Romboutsia
F_Peptostreptococcaceae
F_Erysipelotrichaceae
S_saccharivorans
C_Erysipelotrichia
O_Erysipelotrichales
S_saccharivorans_nov_97_479
G_Phocaeicola
F_Bacteroidaceae
S_vulgatus
S_faecicola_nov_93_111
F_Oscillospiraceae
F_Bacteroidota
O_Bacteroidales
C_Bacteroidia
G_Fusicatenibacter
S_glucerasea
C_Bacteroidia
P_Bacteroidetes
O_Bacteroidales
F_Lactobacillaceae
G_Lactobacillus
S_intestinalis
G_Blautia
F_Bifidobacteriaceae
C_Actinobacteria
G_Bifidobacterium
O_Bifidobacteriales
S_pseudolongum
O_Coriobacteriales
S_perofaciens
F_Coriobacteriaceae
G_Collinsella
C_Coriobacteria
F_Lachnospiraceae
C_Clostridia
O_Eubacteriales
−6.0
−4.8
−3.6
−2.4
−1.2
0.0
1.2
2.4
3.6
4.8
6.0
LDA SCORE (log 10)