Feces_Group_8
Feces_Group_9
Significantly Represented Taxa
P_Firmicutes
C_Clostridia
O_Eubacteriales
C_Bacteroidia
O_Bacteroidales
P_Bacteroidota
S_murinus
G_Ligilactobacillus
F_Oscillospiraceae
G_Phocaeicola
S_vulgatus
F_Bacteroidaceae
F_Muribaculaceae
S_freteri_nov_89_024
G_Duncaniella
S_faecicola_nov_93_111
S_champanellensis_nov_91_810
O_Erysipelotrichales
C_Erysipelotrichia
G_Oscillospiraceae__G_6_
G_Ruminococcus
S_bacterium_MOT_153_nov_97_831
S_timonensis_nov_90_476
G_Raoultibacter
G_Oscillospiraceae__G_1_
O_Clostridiales
G_Eubacterium
F_Eubacteriaceae__XV_
S_psychrotolerans_nov_88_115
G_Lacrimispora
S_saccharolytica_nov_90_625
S_bacterium_MOT_177_nov_93_582
G_Lachnoclostridium
S_bacterium_MOT_147_nov_90_760
S_bacterium_MOT_151_nov_92_946
S_aminivorans_nov_92_812
G_Oscillospiraceae__G_2_
G_Oscillospiraceae__G_4_
F_Eubacteriales__F_1_
G_Sporobacter
S__Ruminococcus_torques_nov_92_693
G_Eubacteriales__G_1_
S_xylanophilum_nov_91_313
S_bacterium_MOT_151_nov_95_417
S_bacterium_MOT_168
G_Kineothrix
G_Lachnospiraceae__G_11_
G_Phocea
S_massiliensis_nov_88_602
G_Ihubacter
S_massiliensis_nov_94_792
F_EubacterialesFamilyXIII_IncertaeSedis
G_Lachnospiraceae__G_14_
S_saccharivorans_nov_89_263
G_Lawsonibacter
G_Lachnospiraceae__G_6_
S_bacterium_MOT_171_nov_94_549
S__Clostridium_herbivorans_nov_92_917
S_bacterium_MOT_149_nov_87_885
S_termitidis_nov_88_223
G_Pseudoflavonifractor
G_Anaerotignum
S_bacterium_MOT_151_nov_96_042
S_multispecies_sppn24_2_nov_95_228
S_multispecies_sppn28_2_nov_95_198
S_bacterium_MOT_153
S_valericigenes_nov_96_250
G_Oscillibacter
S__Ruminococcus_torques_nov_93_515
S_canis_nov_94_561
S_welbionis_nov_86_902
G_Dysosmobacter
O_Cellvibrionales
F_Spongiibacteraceae
S_bacterium_MOT_177_nov_95_851
S_champanellensis_nov_90_713
S_bacterium_MOT_149_nov_93_555
S_bacterium_MOT_159_nov_92_797
S_intestinalis
G_Lachnospiraceae__G_3_
S_bacterium_MOT_151_nov_93_347
S_bacterium_MOT_151_nov_93_971
S_alysoides_nov_91_441
G_Oscillospiraceae__G_3_
G_Paludicola
S_bacterium_MOT_150_nov_96_458
G_Faecalicatena
S_termitidis_nov_87_124
G_Spongiibacter
S_marinus_nov_80_142
G_Anaerocolumna
S_massiliensis_nov_92_276
G_Drancourtella
S_cellulosilytica_nov_88_655
S_bacterium_MOT_170_nov_92_259
G_Lachnospiraceae__G_5_
S_fissicatena_nov_93_946
S_asaccharolyticus_nov_90_083
C_Gammaproteobacteria
S_capillosus_nov_91_060
S_phocaeensis_nov_91_511
S_bacterium_MOT_168_nov_93_946
S_xylanophilum_nov_90_505
S_asaccharolyticus_nov_90_123
S_ventriosum_nov_93_320
G_Lachnospiraceae__G_12_
S_bacterium_MOT_180_nov_90_184
G_Cuneatibacter
S_caecimuris_nov_92_308
S_asaccharolyticus_nov_90_909
G_Acutalibacter
S_muris_nov_95_670
S_intestini_nov_93_111
G_Butyribacter
S_coprostanoligenes_nov_92_424
S_faecicola_nov_85_644
S__Clostridium_polysaccharolyticum_nov_90_437
S_producta_nov_89_834
S_bacterium_MOT_182_nov_92_489
S__Clostridium_saccharogumia_nov_96_042
S__Ruminococcus_gnavus_nov_90_377
S_bacterium_MOT_161_nov_92_308
S_bacterium_MOT_149_nov_96_258
S_bacterium_MOT_150_nov_95_188
S__Clostridium_innocuum_nov_87_078
S_bacterium_MOT_149_nov_96_058
S_viscericola_nov_90_447
F_Barnesiellaceae
G_Barnesiella
S_psychrotolerans_nov_87_037
S_bacterium_MOT_168_nov_93_737
S__Clostridium_polysaccharolyticum_nov_87_500
S_alysoides_nov_87_083
G_Butyricicoccus
S_pullicaecorum_nov_87_216
S_eutactus_nov_92_704
G_Eisenbergiella
S_umbilicata_nov_96_242
G_Faecalimonas
S_schinkii_nov_94_130
S_timonensis
S_massiliensis
S__Ruminococcus_gnavus_nov_95_188
S_albus_nov_91_045
F_Streptococcaceae
G_Streptococcus
S_eutactus_nov_95_085
S_piriformis_nov_85_129
G_Slackia
S_producta_nov_95_178
S_muris_nov_92_754
S_canis_nov_95_388
G_Coprococcus
S_mutans
S_saccharivorans
G_Monoglobus
S_pectinilyticus
S_multispecies_spp10_2
S_flavefaciens_nov_94_760
F_Muribaculaceae
S_bacterium_MOT_104_nov_86_228
G_Muribaculaceae__G_2_
S_saccharivorans_nov_97_689
S_producta_nov_96_226
G_Lachnospiraceae__G_4_
S_bacterium_MOT_169_nov_97_484
S_glucerasea
G_Fusicatenibacter
O_Verrucomicrobiales
C_Verrucomicrobiae
F_Akkermansiaceae
S_muciniphila
G_Akkermansia
P_Verrucomicrobia
S_multispecies_spp19_2
S_luti_nov_96_429
C_Coriobacteriia
S_aerofaciens
G_Collinsella
F_Coriobacteriaceae
O_Coriobacteriales
F_Bifidobacteriaceae
G_Bifidobacterium
C_Actinobacteria
O_Bifidobacteriales
S_pseudolongum
P_Actinobacteria
−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)