NR_0_F
NR_4_F
NR_8_F
Significantly Represented Taxa
O_Eubacteriales
C_Clostridia
F_Lachnospiraceae
G_Lachnospiraceae__G_14_
um_MOT_185_nov_97_638
G_Muribaculaceae__G_2_
F_Muribaculaceae
G_Alistipes
F_Rikenellaceae
s_onderdonkii_nov_95_276
um_MOT_104_nov_92_095
um_MOT_104_nov_96_047
F_Oscillospiraceae
G_Lachnospiraceae__G_11_
G_Acetivibrio
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
S_Turicibacter_sanguinis
F_Turicibacteraceae
G_Turicibacter
pes_finegoldii_nov_96_850
es_multifamily_multigenus
_Eubacteriales_multifamily
ies_sppn50_2_nov_87_843
G_Anaerotignum
chnoclostridium_pacaense
Blautia_faecis_nov_95_669
G_Blautia
um_MOT_178_nov_97_255
G_Frisingicoccus
G_Oscillospiraceae__G_4_
us_caecimuris_nov_96_063
um_MOT_185_nov_94_882
ies_sppn60_2_nov_95_276
G_Oscillospiraceae__G_2_
G_4__bacterium_MOT_151
G_Erysipelatoclostridium
G_Butyribacter
ies_sppn77_2_nov_88_583
um_MOT_149_nov_94_531
_14__bacterium_MOT_183
cter_intestini_nov_95_294
es_sppn102_4_nov_94_510
_Lachnospiraceae__G_12_
G_Acetatifactor
G_Anaerotruncus
_12__bacterium_MOT_179
ium_scindens_nov_93_701
_11__bacterium_MOT_176
G_Eubacteriales__G_4_
cetethylicum_nov_96_471
G_Anaerobium
atifermentans_nov_95_276
ies_sppn99_3_nov_94_094
um_MOT_176_nov_95_669
es_sppn115_4_nov_89_764
G_Lachnospira
G_Papillibacter
innamivorans_nov_89_370
um_MOT_178_nov_97_638
es_sppn106_2_nov_94_882
terella_excrementihominis
s_rubiinfantis_nov_93_307
G_Lachnospiraceae__G_9_
es_sppn133_2_nov_96_063
s_colihominis_nov_94_466
es_sppn124_2_nov_96_063
G_Romboutsia
S_Romboutsia_timonensis
um_pacaense_nov_97_638
es_sppn143_2_nov_95_276
_xylanolytica_nov_96_063
G_2__bacterium_MOT_167
G_Lachnospiraceae__G_2_
er_timonensis_nov_94_094
onibacter_asaccharolyticus
mes_intestini_nov_97_638
G_Muricomes
G_Hungatella
F_Clostridiaceae
um_MOT_164_nov_97_244
G_Sporobacter
ter_termitidis_nov_94_094
um_MOT_174_nov_94_118
ies_sppn74_2_nov_94_488
monas_faecis_nov_96_484
G_Roseburia
G_Pseudobutyrivibrio
G_Oscillospiraceae__G_3_
vibrio_ruminis_nov_91_732
tifactor_muris_nov_92_549
ccharolyticum_nov_93_725
rio_crossotus_nov_95_294
G_Butyrivibrio
G_Faecalicatena
atena_orotica_nov_96_863
ium_scindens_nov_92_913
um_MOT_164_nov_97_638
_ruminantium_nov_94_902
G_Oscillibacter
G_Peptococcaceae__G_1_
F_Peptococcaceae
er_caecimuris_nov_94_118
G_Cuneatibacter
cies_sppn1_2_nov_93_676
scillospiraceae_multigenus
ia_intestinalis_nov_95_276
spira_eligens_nov_89_844
um_uniforme_nov_86_614
um_MOT_165_nov_96_850
um_MOT_150_nov_93_701
otignum_lactatifermentans
ydrogenoanaerobacterium
S_Acetatifactor_muris
ccharovorans_nov_94_094
ira_multipara_nov_87_451
G_Eubacteriales__G_3_
alkalicellulosi_nov_88_189
um_MOT_163_nov_96_457
G_Lachnospiraceae__G_6_
um_MOT_171_nov_93_333
us_caecimuris_nov_94_882
um_MOT_174_nov_93_359
um_MOT_149_nov_92_969
G_Anaerocolumna
mna_jejuensis_nov_94_882
ium_scindens_nov_91_406
um_MOT_185_nov_94_488
tifactor_muris_nov_97_638
es_sppn185_2_nov_93_307
ira_multipara_nov_88_235
G_Lachnospiraceae__G_5_
G_5__bacterium_MOT_170
G_Lachnospiraceae__G_7_
G_Lacrimispora
um_MOT_172_nov_93_307
es_sppn175_2_nov_97_244
G_Anaerosporobacter
G_Acetitomaculum
acter_mobilis_nov_96_063
um_MOT_146_nov_96_850
dium_populeti_nov_93_359
ulum_ruminis_nov_90_551
G_Marvinbryantia
um_MOT_149_nov_92_578
ormatexigens_nov_93_307
ies_sppn13_2_nov_87_008
ies_sppn21_2_nov_95_276
um_MOT_151_nov_97_638
um_MOT_150_nov_96_457
ium_scindens_nov_92_969
G_Lachnospiraceae__G_3_
G_3__bacterium_MOT_168
atena_orotica_nov_95_276
G_Acutalibacter
ia_intestinalis_nov_92_913
S_Acutalibacter_muris
G_Akkermansia
F_Akkermansiaceae
O_Verrucomicrobiales
C_Verrucomicrobiae
P_Verrucomicrobia
_Akkermansia_muciniphila
G_Ligilactobacillus
llus_multispecies_spp11_4
C_Gammaproteobacteria
F_Enterobacteriaceae
nus_multispecies_spp25_7
O_Enterobacterales
obacteriaceae_multigenus
um_MOT_104_nov_91_700
S_Anaerostipes_caccae
G_Anaerostipes
F_Peptostreptococcaceae
ium_scindens_nov_95_276
G_Clostridiales__F_1__G_1_
um_HMT_093_nov_89_804
F_Clostridiales__F_1_
0
1
2
3
4
5
LDA SCORE (log 10)