Significantly Represented Taxa
BL
ligSPM
C_Actinobacteria
P_Actinobacteria
S_acnes
O_Actinomycetales
F_Propionibacteriaceae
G_Propionibacterium
O_Rhizobiales
C_Alphaproteobacteria
F_Moraxellaceae
G_Acinetobacter
O_Pseudomonadales
O_Propionibacteriales
S_acnes
G_Cutibacterium
F_Propionibacteriaceae
S_johnsonii
G_Bradyrhizobium
F_Bradyrhizobiaceae
S_lupini
S_finegoldii_nov_92_600
S_sp__Oral_Taxon_A28_nov_92_634
O_Alphaproteobacteria__O_
G_Alphaproteobacteria__G_
F_Clostridiaceae
S_sp__str__D16_nov_94_605
F_Alphaproteobacteria__F_
S_sp__str__D16_nov_90_249
G_Barnesiella
F_Barnesiellaceae
O_Anaeroplasmatales
S_bacterium_HMT_911_nov_84_854
G_Anaerotruncus
G_Acetivibrio
S_bacterium_HMT_911_nov_78_835
C_Mollicutes
F_Anaeroplasmataceae
G_Anaeroplasma
S_bacterium_HMT_911_nov_85_127
P_Tenericutes
G_Lachnoclostridium
S_bacterium_HMT_911_nov_85_853
F_NA
G_Faecalicatena
S_sanguinis_nov_95_833
G_Turicibacter
S_orotica_nov_92_218
S_bacterium_HMT_351_nov_94_004
S_bacterium_HMT_351_nov_92_706
G_Saccharibacteria__TM7___G_3_
S_cellulolyticus_nov_84_058
S_senegalensis_nov_94_073
S_bacterium_HMT_911_nov_84_423
S_sp__str__cont1_66_nov_97_744
S_bacterium_HMT_911_nov_84_660
S_sp__str__cont1_66
O_Saccharibacteria__TM7___O_1_
F_Alcaligenaceae
G_Sutterella
S_finegoldii_nov_93_690
S_bacterium_HMT_911_nov_85_742
G_unclassified_Ruminococcaceae
C_Saccharibacteria__TM7___C_1_
F_multifamily
S_bacterium_HMT_911_nov_86_127
G_Gabonia
S_bacterium_HMT_911_nov_85_910
F_Lachnospiraceae
P_Verrucomicrobia
S_massiliensis_nov_86_704
G_Akkermansia
O_Verrucomicrobiales
S_muciniphila
F_Akkermansiaceae
S_senegalensis_nov_93_846
C_Verrucomicrobiae
P_Saccharibacteria__TM7_
F_Saccharibacteria__TM7___F_1_
G_Alistipes
F_Rikenellaceae
S_bacterium_HMT_911_nov_85_352
S_bacterium_HMT_911_nov_83_268
F_Bacteroidetes__F_1_
G_Bacteroidetes__G_7_
O_Bacteroidetes__O_1_
C_Bacteroidetes__C_1_
P_Bacteroidetes
−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)