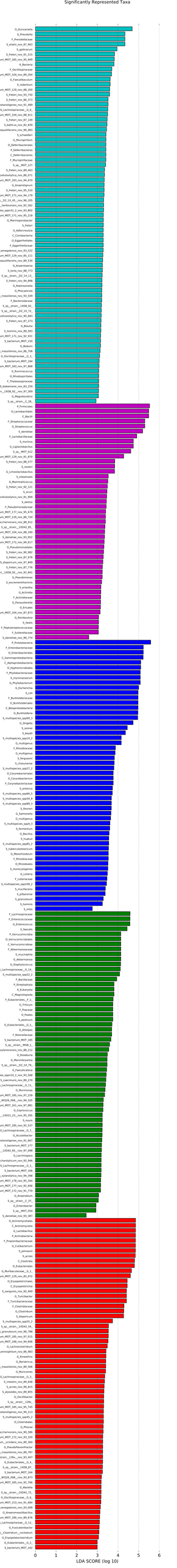
Significantly Represented Taxa
G_Duncaniella
G_Prevotella
F_Prevotellaceae
S_shahii_now_87_903
S_gallinarum
S_freteri_now_91_515
un_MOT_185_now_95_949
K_Bacteria
F_Oscillospiraceae
un_MOT_166_now_89_264
G_Faecalibaculum
S_rodentium
un_MOT_129_now_88_200
S_freteri_now_93_750
S_freteri_now_98_373
ostanolingenes_now_91_489
G_Lachnospiraceae_G_6_
un_MOT_104_now_88_911
S_freteri_now_97_149
S_faditicus_now_82_839
equifaciens_now_96_983
S_schaedleri
G_Mucispirillum
O_Defferibacterales
F_Defferibacteres
C_Defferibacteres
F_Mucispirillaceae
S_sp._MOT_127
S_freteri_now_98_483
celulosilytica_now_86_072
un_MOT_163_now_94_670
G_Anaerotignum
S_freteri_now_95_510
un_MOT_171_now_94_179
__02_14_45_now_96_205
_herbivorans_now_92_562
_sapes_sp2r3_2_now_93_802
un_MOT_171_now_95_218
G_Marinisporobacter
S_freteri
G_Alkercreutzia
C_Coriobacteria
O_Eggerthellales
F_Eggerthellaceae
senegalensis_now_93_522
un_MOT_129_now_89_311
equifaciens_now_98_530
G_Anaerotaenia
S_torta_now_88_773
S_sp._strain__02_14_13_
S_freteri_now_96_898
G_Paludimonella
G_Phocaeicola
_massiliensis_now_91_509
F_Bacteroidaceae
S_sp._strain__1408_92_
S_sp._strain__02_14_72_
celulosilytica_now_90_683
S_freteri_now_97_273
G_Blautia
S_hominis_now_89_583
un_MOT_171_now_92_931
S_bacterium_MOT_150
G_Bebum
_massiliensis_now_86_708
G_Oscillospiraceae_G_5_
S_bacterium_MOT_184
un_MOT_163_now_97_868
G_Ruminococcus
O_Rhodospirillales
F_Thalassospiraceae
S_blakemore_now_83_259
_1408_92_now_97_309
G_Magnetovibrio
S_sp._strain_C_28_
F_Fonsiculaceae
O_Lactobacillales
C_Bacilli
F_Streptococcaceae
G_Streptococcus
S_danielae
F_Lactobacillaceae
S_murinus
G_Ligilactobacillus
S_sp._MOT_812
un_MOT_129_now_91_870
S_freteri_now_88_577
S_avium
G_Limosilactobacillus
S_intestinalis
G_Mammaliicoccus
S_freteri_now_92_121
S_aciuri
celulosilytica_now_91_909
S_aestus
F_Pseudomonadaceae
un_MOT_177_now_95_679
un_MOT_129_now_98_720
ichaoriavora_now_88_912
S_sp._strain__1D42_95_
un_MOT_104_now_88_340
S_danielae_now_93_952
un_MOT_172_now_94_617
O_Pseudomonadales
S_freteri_now_90_982
S_freteri_now_97_976
S_disporicum_now_97_849
S_freteri_now_97_776
_1408_92_now_92_841
G_Pseudomonas
S_excrementihomnis
S_xylanilytica
G_Actinotia
F_Actinotaceae
G_Parautisella
O_Micrales
un_MOT_104_now_87_873
G_Rombutsia
S_lealis
F_Peptostreptococcaceae
F_Sutterellaceae
S_danielae_now_96_774
P_Proteobacteria
F_Enterobacteriaceae
O_Enterobacteriales
C_Gammaproteobacteria
C_Alphaproteobacteria
O_Hyphomicrobiales
F_Phyllobacteriaceae
S_myrsinacearum
G_Phyllobacterium
G_Escherichia
S_coli
F_Burkholderiaceae
O_Burkholderiales
C_Betaproteobacteria
G_Burkholderia
S_multispecies_sp49_3
G_Shigella
S_sonnei
S_boydii
S_multispecies_sp414_2
G_multigenus
F_Rhizobiaceae
G_multigenus
S_ferguson
S_chisuanse
S_multispecies_sp27_2
G_Corynebacterales
G_Corynebacterium
F_Corynebacteriaceae
S_enterica
S_multispecies_sp66_5
S_multispecies_sp59_8
S_multispecies_sp89_3
S_flexneri
G_Salmonella
G_multigenus
S_multispecies_sp4_3
S_fermentum
G_Bacillus
S_huakui
S_multispecies_sp85_2
S_tuberculosistarium
G_Mesorhizobium
F_Rhizobiaceae
O_Rhizobiales
S_monocytogenes
G_Listeria
F_Listeriaceae
S_multispecies_sp105_2
S_mucifaciens
S_pilosiversae
S_granulosum
S_hominis
S_mitis
F_Lachnospiraceae
F_Enterococcaceae
G_Enterococcus
S_faecalis
P_Verrucomicrobia
O_Verrucomicrobiales
C_Verrucomicrobiae
F_Akkermansiaceae
S_muciniphila
G_Akkermansia
G_Staphylococcus
S_multispecies_sp22_2
S_multispecies_sp22_2
F_Bacillaceae
P_Streptophyta
K_Eukaryota
C_Magnoliopsida
F_Eubacteriales_F_5_
G_Triticum
F_Poaceae
O_Poales
S_aestivum
G_Eubacteriales_G_1_
G_Alistipes
F_Rikenellaceae
S_bacterium_MOT_185
S_sp._strain_ _MOR_1_
oryanovorans_now_88_212
G_Roseburia
G_Marvinbryantia
S_sp._strain_D2_14_79
S_Faecalicatena
_sapes_sp2r10_2_now_93_568
S_caecimuris_now_89_270
G_Lachnospiraceae_G_11_
G_Muribaculum
un_MOT_185_now_97_228
S_WY20_X68_now_94_105
un_MOT_165_now_97_881
G_Coprococcus
S_1XD21_23_ _now_95_595
S_muris
un_MOT_185_now_92_537
G_Lachnospiraceae_G_7_
G_Acutalbacter
ostanolingenes_now_91_667
S_bacterium_MOT_177
S_1XD42_95_ _now_97_098
G_Lacrimispora
chanshyicum_now_92_946
G_Lachnospiraceae_G_1_
S_bacterium_MOT_166
celulosilytica_now_98_398
un_MOT_178_now_95_344
un_MOT_172_now_92_608
un_MOT_172_now_93_770
G_Anaerotruncus
S_sp._strain_C_37_
G_Enterobacter
S_sp._MOT_050
S_danielae_now_93_387
G_Actinomycetales
G_Actinomycetales
G_Lactobacillus
P_Actinobacteria
F_Propionibacteriaceae
G_Cutibacterium
S_johnsonii
S_acnes
C_Clostridia
O_Eubacteriales
G_Muribaculaceae_G_1_
un_MOT_129_now_89_972
O_Erysipelotrichales
C_Erysipelotrichia
S_sanguinis_now_95_949
G_Turicibacter
F_Turicibacteraceae
F_Clostridiaceae
G_Clostridium
S_disporicum
S_multispecies_sp85_2
S_sp._strain_ _1XD42_54_
S_granulosum_now_96_788
un_MOT_165_now_97_015
un_MOT_168_now_94_606
G_Lachnoclostridium
_aminophilum_now_86_983
G_Kineothrix
G_Barnesiella
_massiliensis_now_89_506
G_Muricomes
G_Lachnospiraceae_G_3_
S_intestini_now_89_648
S_acnes_now_98_815
S_alysoides_now_89_855
G_Oscillibacter
S_sp._strain_ _128x_
un_MOT_185_now_95_745
ostanolingenes_now_90_213
S_multispecies_sp45_2
O_Clostridiales
G_Phocea
ichaoriavora_now_90_586
un_MOT_172_now_93_300
_sn_scindens_now_89_300
G_Paludibacteroides
_massiliensis_now_89_787
strain_ _136x_now_93_407
G_Eubacteriales_G_4_
S_sp._strain_ _1XD8_87_
S_bacterium_MOT_164
_WY20_X68_now_93_873
un_MOT_185_now_92_766
G_Alistella
S_sp._strain_ _1XD42_72_
G_Oscillospiraceae_G_6_
un_MOT_153_now_91_684
_senegalensis_now_91_008
G_Anaeromassilibacillus
un_MOT_180_now_89_676
G_Lachnospiraceae_G_12_
G_Faecalibacter
G_Clostridium_cocleatum
G_Erysipelatoclostridium
G_Eubacteriales_G_2_
S_bacterium_MOT_182
0
1
2
3
4
5
6
LDA SCORE (log 10)