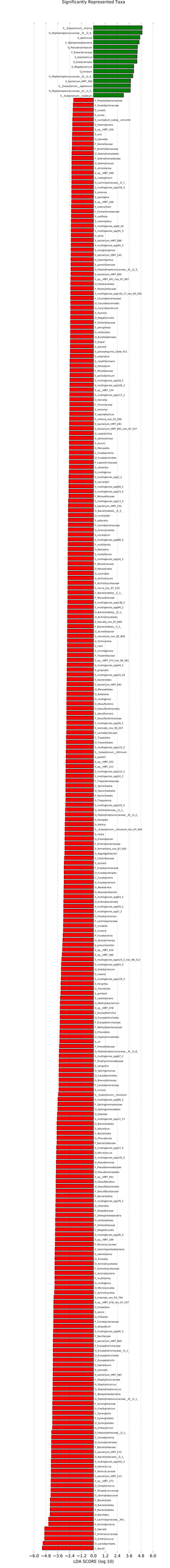
Significantly Represented Taxa
s_Eubacterium_brachy
G_Peptostreptococcaceae_XI_G_5
s_Gelria_sp
C_Alphaproteobacteria
G_Pseudoramibacter
F_Eubacteriaceae
s_shelfordii
O_Eubacteriales
C_Negativicutes
s_timidum
G_Peptostreptococcaceae_XI_G_6
s_Bacteroides_sartorii_DA5
s_Eubacterium_saphenum
G_Peptostreptococcaceae_XI_G_5
s_Eubacterium_minutum
F_Propionibacteriaceae
F_Cholelacibacteraceae
s_sinaii
s_acnes
s_rectalium_subsp._vincentii
s_kanoi/yoanis
s_sp._HMT_359
s_uria
G_Gemella
F_Gemellaceae
F_Burkholderiaceae
O_Selenomonadales
F_Selenomonadaceae
G_Selenomonas
s_artrosiacea
s_sp._HMT_448
s_hallopilum
G_Lachnospiraceae_G_7
s_multiparus_ss/104_3
s_oralis
s_sp._HMT_359
s_uria
s_sp._HMT_386
s_multiparus_ss/65_5
s_hongkongensis
s_bacterium_HMT_145
G_Haemophilus
s_parainfluenzae
G_Peptostreptococcaceae_XI_G_3
s_bacterium_HMT_855
s_sp._HMT_841/nov_97_907
O_Pedosporales
F_Pedosporaceae
s_multiparus_ss/pb14_17/nov_95_592
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
s_hominis
O_Negativicutes
F_Veillonellaceae
s_aeruginosa
G_Helicobacter
O_Burkholderiales
s_dispar
s_weinerii
s_parasanguinis_clade_411
s_virginiae
s_nisellii/mans
G_Moraxella
F_Moraxellaceae
G_Porphyromonas
s_multiparus_ss/18_2
s_multiparus_ss/100_3
s_sp._HMT_134
s_multiparus_ss/117_2
G_Serratia
F_Yersiniaceae
s_lactarius
s_sapinigfrifrus
s_oliouo_nov_93_300
s_bacterium_HMT_281
s_bacterium_HMT_091/nov_97_537
G_Lepptotrichia
s_denticanosa
s_durum
G_Morasella
C_Fusobacteria
O_Fusobacteriales
F_Leptotrichiaceae
s_volserisis
G_multigenus
s_multiparus_ss/2_2
s_succinatii
s_multiparus_ss/8_2
s_multiparus_ss/23_4
F_Moraxellaceae
s_multiparus_ss/13_3
s_bacterium_HMT_155
G_Bacteroides_G_3
G_multiorder
s_adiacens
F_Carnobacteriaceae
G_Granulicatella
s_rectalium
s_multiparus_ss/88_4
F_multigenig
G_Neisseria
s_mordiform
s_multiparus_ss/16_2
F_Neisseriaceae
O_Neisseriales
G_Lactopia
G_Actinomyces
F_Actinomycetaceae
s_nicra_nov_97_516
G_Bacteroides_G_1
F_Moraxellaceae
s_multiparus_ss/138_2
s_multiparus_ss/84_2
G_Bacteroides_G_1
G_Actinomycetales
s_faecalis_nov_97_669
F_Bacteroides_G_1
G_Acinetobacter
s_ulrunensis_nov_92_840
G_Schnuerera
s_noni
s_muciaginosa
F_Flavonifractaceae
s_sp._HMT_373_nov_94_581
s_multiparus_ss/94_2
s_gingivalis
s_multiparus_ss/10_20
s_bacteroides
s_bacterium_HMT_091
G_Morasella
G_Ralstonia
G_multigenus
G_Desulfovibrio
O_Desulfovibrionales
s_bisulfuricans
F_Desulfovibrionaceae
s_multiparus_ss/19_2
s_stomatis_nov_95_037
F_Lachnospiraceae
G_Treponella
O_Treponeales
s_multiparus_ss/121_2
s_Eubacterium_infirmum
s_podoci
s_sp._HMT_203
s_sp._HMT_212
s_multiparus_ss/112_2
s_multiparus_ss/12_2
F_Treponemataceae
G_Spirochaeta
O_Spirochaetales
F_Spirochaetaceae
G_Treponema
s_multiparus_ss/103_5
O_Veillonellaceae_G_1
G_Peptostreptococcaceae_XI_G_2
s_stingala
G_Rothia
s_Eubacterium_rimulum_nov_97_044
s_nodatum
G_Enterobacter
F_Enterobacteriaceae
s_terrestris_nov_97_042
G_Aggregatibacter
F_Clostridiaceae
s_stuteri
F_Fusobacteriaceae
O_Fusobacteriales
F_Fusobacteria
G_Fusobacterium
G_Neobacillus
G_Atopobiobacter
s_multiparus_ss/63_2
O_Enterobacteriales
s_multiparus_ss/59_2
s_multiparus_ss/7_2
G_Porphyromonas
F_Lachnospiraceae
s_notabilis
s_mutans
F_Fusobacteria
G_Stutzerimonas
s_pneumoniae
s_sp._HMT_416
s_sp._HMT_360
s_multiparus_ss/ss/23_2_nov_96_512
s_multiparus_ss/89_3
G_Solobacterium
s_moorei
s_multiparus_ss/129_2
G_Scrofilia
G_Tannerella
s_gordonii
s_odontolyticus
G_Methylobacterium
s_sp._HMT_078
G_Erysipelotrichia
G_Erysipelotrichales
F_Erysipelotrichaceae
F_Methylobacteriaceae
G_Prevotella
G_Oryhanomicrobiales
s_ali
F_Prevotellaceae
G_Peptostreptococcaceae_XI_G_6
s_multiparus_ss/67_2
F_Porphyromonadaceae
s_sanguinis
G_Sphingomonas
O_Caulobacterales
G_Brevundimonas
F_Caulobacteraceae
s_minuta
S_Eubacterium_rimulum
s_multiparus_ss/68_2
F_Sphingomonadaceae
O_Sphingomonadales
G_Dialister
s_multiparus_ss/12_17
G_Bacteroidales
s_siloceous
G_Bacteroides
G_Phocaeicola
F_Bacteroidaceae
s_multiparus_ss/57_4
G_Micrococcus
s_multiparus_ss/45_5
G_Pseudomonas
F_Pseudomonadaceae
G_Pseudomonadales
s_sp._HMT_041
G_Desulfovibrio
O_Desulfobacterales
F_Desulfobulbaceae
F_Bacteroidota
s_multiparus_ss/78_2
G_Oranella
F_Atopobiaceae
G_Desporobacteria
O_Veillonellales
F_Veillonellaceae
G_Negativicutes
s_multiparus_ss/99_3
s_sp._HMT_199
F_Micrococcaceae
F_Carnimonadaceae
s_odontolyica
G_Schaalia
O_Actinomycetales
F_Actinomycetaceae
C_Actinobacteria
F_multigenig
G_multigenus
O_Micrococcales
C_Actinomycetia
s_sinensiae_nov_93_764
s_sp._HMT_278_nov_97_537
G_Haloshia
s_micro
G_Filifactor
F_Ornithobacteraceae
G_Atopobium
s_multiparus_ss/40_3
F_Bacteraceae
s_bacterium_HMT_904
F_Erysipelotrichaceae
G_Erysipelotrichaceae_G_1
G_Erysipelothrichales
C_Erysipelotrichi
S_Solobacium
s_stomatis
s_bacterium_HMT_383
F_Staphylococcaceae
G_Staphylococcus
G_Peptostreptococcus
F_Betapolimobacteria
G_Peptostreptococcaceae_XI_G_1
F_Synergistaceae
G_Fretibacterium
G_Synergistia
F_Synergistaceae
O_Synergistales
G_Cristachorum
G_Porphyrihaloaceae_G_1
C_Cytobacteria
O_Cytobacteriales
F_Bacteroidaceae
s_bacterium_HMT_272
G_Bacteroidaceae_G_1
s_multiparus_ss/162_2
G_Aerococcus
F_Aerococcaceae
s_bacterium_HMT_113
s_sp._HMT_371
G_Streptococcus
F_Streptococcaceae
G_Stomatobaculum
F_Bacteroidia
O_Bacteroidales
F_Bacteroidales
O_Bacillales
F_Lachnospiraceae_XIV_
G_Actinobacteria
G_Saccalis
F_Enterococcaceae
G_Enterococcus
O_Lactobacillales
C_Bacilli
−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)