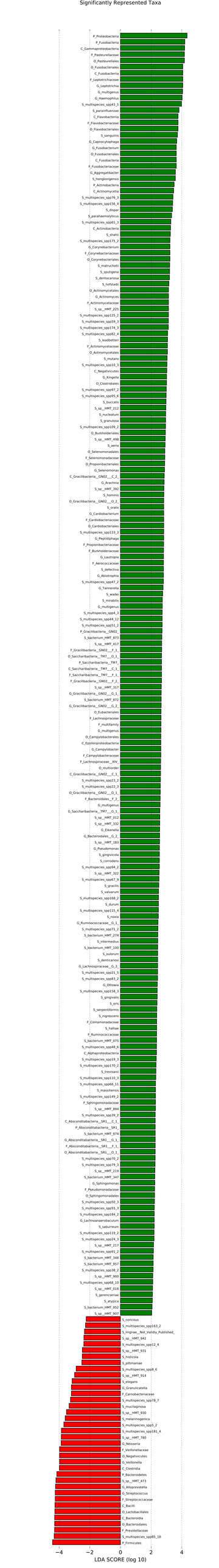
Significantly Represented Taxa
P_Proteobacteria
P_Fusobacteria
C_Gammaproteobacteria
F_Pasteurellaceae
O_Pasteurellales
O_Fusobacteriales
C_Fusobacteria
F_Leptotrichiaceae
G_Leptotrichia
G_multigenus
G_Haemophilus
S_multispecies_spp43_5
S_parainfluenzae
C_Flavobacteria
F_Flavobacteriaceae
O_Flavobacteriales
S_sanguinis
G_Capnocytophaga
G_Fusobacterium
O_Fusobacteriales
C_Fusobacteria
F_Fusobacteriaceae
G_Aggregatibacter
S_hongkongensis
P_Actinobacteria
C_Actinomycetia
S_multispecies_spp76_3
S_multispecies_spp156_4
S_dispar
S_parahaemolyticus
S_multispecies_spp61_3
C_Actinobacteria
S_shahii
S_multispecies_spp175_2
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_matruchotii
S_spurigena
S_denticariosa
S_hofstadii
O_Actinomycetales
G_Actinomyces
F_Actinomycetaceae
S_sp._HMT_225
S_multispecies_spp125_2
S_multispecies_spp59_3
S_multispecies_spp174_3
S_multispecies_spp82_4
S_leadbetteri
F_Actinomycetaceae
O_Actinomycetales
S_mutans
S_multispecies_spp10_3
C_Negativicutes
G_Kingella
O_Clostridiales
S_multispecies_spp97_2
S_multispecies_spp95_6
S_buccalis
S_sp._HMT_212
S_nuclatum
S_granulosa
S_multispecies_spp109_2
O_Burkholderiales
S_sp._HMT_498
S_aeria
G_Selenomonadales
F_Selenomonadaceae
O_Propionibacteriales
G_Selenomonas
C_Gracilibacteria__GN02__C_2_
G_Arachnia
S_sp._HMT_392
S_hominis
O_Gracilibacteria__GN02__O_2_
S_oralis
G_Cardiobacterium
F_Cardiobacteriaceae
O_Cardiobacteriales
S_multispecies_spp133_3
G_Peptidiphaga
F_Propionibacteriaceae
F_Burkholderiaceae
G_Lautropia
F_Aerococcaceae
S_defectiva
G_Abiotrophia
S_multispecies_spp47_2
G_Tannerella
S_wadei
S_mirabilis
G_multigenus
S_multispecies_spp4_3
S_multispecies_spp84_12
S_multispecies_spp51_2
P_Gracilibacteria__GN02__
S_bacterium_HMT_873
S_sp._HMT_417
F_Gracilibacteria__GN02__F_1_
O_Saccharibacteria__TM7__O_1_
F_Saccharibacteria__TM7__F_1_
C_Saccharibacteria__TM7__C_1_
F_Saccharibacteria__TM7__F_1_
F_Gracilibacteria__GN02__F_2_
S_sp._HMT_317
G_Gracilibacteria__GN02__G_1_
S_bacterium_HMT_872
G_Gracilibacteria__GN02__G_2_
O_Eubacteriales
F_Lachnospiraceae
F_multifamily
G_multigenus
O_Campylobacterales
C_Epsilonproteobacteria
G_Campylobacter
F_Campylobacteraceae
F_Lachnospiraceae__XIV_
O_multiorder
C_Gracilibacteria__GN02__C_1_
S_multispecies_spp23_2
S_multispecies_spp22_3
O_Gracilibacteria__GN02__O_1_
F_Bacteroidales__F_2_
G_multigenus
G_Saccharibacteria__TM7__G_1_
S_sp._HMT_012
S_sp._HMT_332
G_Eikenella
G_Bacteroidales__G_2_
S_sp._HMT_183
G_Pseudomonas
S_gingivicola
S_corrodens
S_multispecies_spp94_2
S_sp._HMT_322
S_multispecies_spp67_9
S_gracilis
S_valvarum
S_multispecies_spp168_2
S_durum
S_multispecies_spp115_4
S_noxia
G_Ruminococcaceae__G_1_
S_multispecies_spp71_2
S_bacterium_HMT_274
S_intermedius
S_bacterium_HMT_100
S_nutorum
S_denticariosa
G_Lachnospiraceae__G_3_
S_multispecies_spp21_5
S_multispecies_spp83_2
G_Ottowia
S_multispecies_spp158_3
S_gingivalis
S_oris
S_serpentiformis
S_nigrescens
F_Comamonadaceae
S_hallae
F_Ruminococcaceae
S_bacterium_HMT_075
S_multispecies_spp46_6
C_Alphaproteobacteria
S_multispecies_spp19_3
S_multispecies_spp170_2
S_trevisanii
S_multispecies_spp110_3
S_multispecies_spp66_11
S_massiliensis
S_multispecies_spp149_2
F_Sphingomonadaceae
S_sp._HMT_894
S_multispecies_spp39_2
C_Absconditabacteria__SR1__C_1_
P_Absconditabacteria__SR1__
S_bacterium_HMT_874
G_Absconditabacteria__SR1__G_1_
F_Absconditabacteria__SR1__F_1_
O_Absconditabacteria__SR1__O_1_
S_multispecies_spp70_2
S_multispecies_spp79_3
S_sp._HMT_219
S_bacterium_HMT_347
G_Sphingomonas
F_Pseudomonadaceae
O_Sphingomonadales
S_multispecies_spp50_3
S_multispecies_spp55_3
S_multispecies_spp184_2
G_Lachnoanaerobaculum
S_saburreum
S_multispecies_spp119_2
S_multispecies_spp24_3
S_sp._HMT_217
S_multispecies_spp91_2
S_bacterium_HMT_348
S_bacterium_HMT_957
S_multispecies_spp38_2
S_sp._HMT_900
S_multispecies_spp68_10
S_sp._HMT_018
S_gerencseriae
S_atypica
S_bacterium_HMT_952
S_sp._HMT_907
S_conclusus
S_multispecies_spp163_2
S_lingnae_Not_Validly_Published_
S_sp._HMT_942
S_multispecies_spp32_4
S_sp._HMT_931
S_histicola
S_pittmaniae
S_multispecies_spp8_6
S_sp._HMT_914
S_elegans
G_Granulicatella
F_Carnobacteriaceae
S_multispecies_spp78_7
S_mucilaginosa
S_sp._HMT_930
S_melaninogenica
S_multispecies_spp5_2
S_multispecies_spp181_4
S_sp._HMT_780
G_Neisseria
F_Veillonellaceae
O_Negativicutes
G_Veillonella
C_Clostridia
P_Bacteroidetes
S_sp._HMT_473
G_Alloprevotella
G_Streptococcus
F_Streptococcaceae
C_Bacilli
O_Lactobacillales
C_Bacteroidia
O_Bacteroidales
F_Prevotellaceae
S_multispecies_spp85_18
P_Firmicutes
−4
−2
0
2
4
LDA SCORE (log 10)