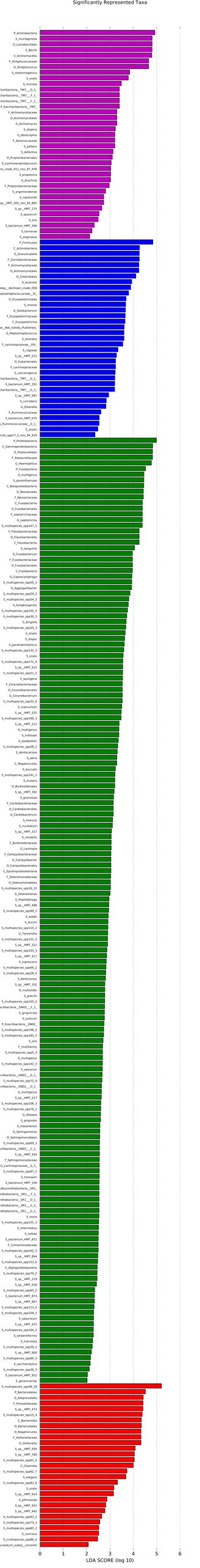
Significantly Represented Taxa
P_Actinobacteria_
S_muciniphila
O_Lactobacillales
C_Bacilli
C_Actinomycetia
F_Streptococcaceae
G_Streptococcus
S_melanogenica
S_orale
S_mucosa
harbacteria_TM7__G_1_
harbacteria_TM7__F_1_
harbacteria_TM7__C_1_
P_Saccharibacteria_TM7_
F_Actinomycetaceae
O_Actinomycetales
G_Actinomyces
S_atypica
O_Bifidobacteria
F_Aerococcaceae
S_pallens
S_defectiva
O_Propionibacteriales
G_Lachnoanaerobaculum
ns_clade_811_ns_37_976
S_propionica
G_Arachnia
F_Propionibacteriaceae
S_argentoratense
S_naeslundii
sp__HMT_305_ noc_93_865
S_sp__HMT_175
S_sputorum
S_oris
S_bacterium_HMT_346
S_tannerae
S_anginosus
P_Firmicutes
C_Actinobacteria
G_Granulicatella
F_Carnobacteriaceae
F_Actinomycetaceae
O_Actinomycetales
O_Clostridiales
S_avitarla
obsp__dentiani_clade_098
eptostreptococcaceae_XI_
O_Erysipelotrichales
S_moore
G_Solobacterium
F_Erysipelotrichaceae
C_Erysipelotrichia
ae_Not_Validly_Published_
G_Peptostreptococcus
S_stomatis
F_Lachnospiraceae_XIV_
S_ringosae
S_sp__HMT_215
O_Eubacteriales
F_Lachnospiraceae
S_cancerogenus
harbacteria_TM7__G_1_
S_bacterium_HMT_352
harbacteria_TM7__G_3_
S_sp__HMT_097
S_corrodens
G_Eikenella
F_Ruminococcaceae
S_bacterium_HMT_075
F_Ruminococcaceae_G_1_
S_shahii
cies_sp97_2_noc_94_919
P_Proteobacteria
C_Gammaproteobacteria
O_Pasteurellales
F_Pasteurellaceae
G_Haemophilus
P_Fusobacteria
G_multigenus
S_parainfluenzae
C_Betaproteobacteria
O_Neisseriales
F_Neisseriaceae
C_Fusobacteria
O_Fusobacteriales
F_Leptotrichiaceae
G_Leptotrichia
S_multispecies_spp147_5
F_Flavobacteriaceae
O_Flavobacteriales
C_Flavobacteria
S_sanguinis
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
G_Capnocytophaga
S_multispecies_spp50_3
G_Aggregatibacter
S_multispecies_spp28_2
S_multispecies_spp54_3
S_hongkongensis
S_multispecies_spp102_4
S_multispecies_spp30_3
G_Kingella
S_multispecies_spp35_3
S_shahii
S_dispar
S_parahaemolyticus
S_multispecies_spp132_3
S_oralis
S_multispecies_spp171_4
S_sp__HMT_012
S_multispecies_spp51_2
S_sputigena
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
S_multispecies_spp32_6
S_matruchotii
S_sp__HMT_225
S_multispecies_spp186_3
S_sp__HMT_212
G_multigenus
S_hofstadii
S_bisdbattiae
S_multispecies_spp38_2
S_denticanina
S_aerris
C_Negativicutes
S_buccalis
S_multispecies_spp143_2
S_mutans
O_Burkholderiales
S_sp__HMT_392
S_granulosa
F_Cardiobacteriaceae
O_Cardiobacteriales
G_Cardiobacterium
S_hominis
S_nucleatum
S_sp__HMT_317
S_mirabilis
F_Burkholderiaceae
G_Lautropia
F_Campylobacteraceae
G_Campylobacter
O_Campylobacterales
C_Epsilonproteobacteria
F_Selenomonadaceae
O_Selenomonadales
S_multispecies_spp16_12
G_Selenomonas
G_Peptidphaga
S_sp__HMT_498
S_multispecies_spp99_2
S_wadei
S_durum
S_multispecies_spp114_2
G_Tannerella
S_multispecies_spp131_2
S_sp__HMT_327
S_multispecies_spp103_3
S_sp__HMT_417
S_ruminicola
S_multispecies_spp66_2
S_multispecies_spp28_4
S_denticanum
S_sp__HMT_332
O_multorder
S_gracilis
S_multispecies_spp105_2
cilibacteria_GN02__F_1_
S_gingivalis
S_zuelzerae
P_Gracilibacteria_GN02_
S_multispecies_spp198_2
S_multispecies_spp182_2
S_oris
F_multifamily
S_multispecies_spp5_3
G_multigenus
S_multispecies_spp142_3
S_valveruni
cilibacteria_GN02__G_1_
S_multispecies_spp72_9
cilibacteria_GN02__O_1_
G_multigenus
S_sp__HMT_237
S_multispecies_spp106_3
S_multispecies_spp76_2
G_Ottowia
S_gingivalis
S_massiliensis
G_Sphingomonas
O_Sphingomonadales
S_multispecies_spp69_3
cilibacteria_GN02__C_1_
S_sp__HMT_183
F_Sphingomonadaceae
G_Lachnospiraceae_G_3_
S_multispecies_spp87_2
S_trevisanii
S_bacterium_HMT_100
Abscondibacteria_SR1_
ndibacteria_SR1__F_1_
ndibacteria_SR1__O_1_
ndibacteria_SR1__C_1_
ndibacteria_SR1__G_1_
S_noeni
S_multispecies_spp101_3
S_intermedius
S_hallae
S_bacterium_HMT_872
F_Comamonadaceae
S_multispecies_spp162_3
S_sp__HMT_894
S_multispecies_spp153_6
C_Alphaproteobacteria
S_multispecies_spp79_2
S_sp__HMT_219
S_sp__HMT_018
S_multispecies_spp95_2
S_bacterium_HMT_874
S_sp__HMT_907
S_multispecies_spp113_2
S_multispecies_spp108_2
S_saburreum
S_sp__HMT_472
S_multispecies_spp164_2
S_serpentiformis
S_maculosa
S_multispecies_spp39_2
S_sp__HMT_900
S_multispecies_spp90_3
S_saccharolytica
S_multispecies_spp36_3
S_bacterium_HMT_952
S_greenscarlae
S_multispecies_spp48_14
P_Bacteroidetes
G_Alloprevotella
F_Prevotellaceae
S_sp__HMT_473
S_multispecies_spp15_4
C_Bacteroidia
O_Bacteroidales
O_Negativicutes
F_Veillonellaceae
G_Veillonella
S_sp__HMT_930
S_sp__HMT_780
S_multispecies_spp61_2
C_Clostridia
S_multispecies_spp62_7
S_elegans
S_multispecies_spp83_8
S_oralis
S_sp__HMT_914
S_pittmaniae
S_sp__HMT_931
S_sp__HMT_942
S_multispecies_spp93_2
S_multispecies_spp74_3
S_multispecies_spp85_2
S_concisus
S_multispecies_spp86_4
acteriaceae_subsp__vincentii
0
1
2
3
4
5
6
LDA SCORE (log 10)