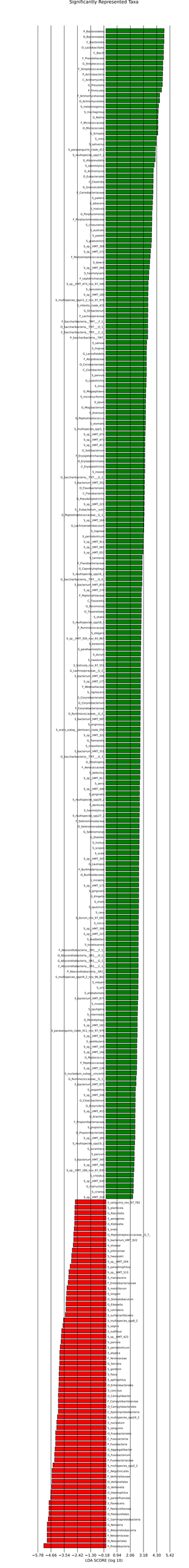
Significantly Represented Taxa
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
O_Lactobacillales
C_Bacilli
F_Prevotellaceae
G_Streptococcus
F_Streptococcaceae
F_Actinobacteria
C_Actinomycetia
G_Prevotella
F_Firmicutes
F_Actinomycetaceae
O_Actinomycetales
S_melaninogenica
S_micigliosa
G_Rothia
F_Micrococcaceae
O_Micrococcales
G_Schaalia
S_mitis
S_salivarius
S_parasanguinis_clade_411
S_multispecies_sp017_r
G_Alloprevotella
S_dentalystici
G_Actinomyces
O_Eubacteriales
C_Clostridia
G_Granulicatella
F_Carnobacteriaceae
S_pallens
S_adiacens
S_histicola
G_Porphyromonas
F_Porphyromonadaceae
S_croceum
S_nodosum
S_australis
S_pasteri
S_graevenitzii
S_sp._HMT_309
S_sp._HMT_173
F_Peptostreptococcaceae
S_dower
S_sp._HMT_066
S_haemolysans
F_Leptotrichiaceae
S_sp._HMT_473_rov_97_546
S_sanguinis
S_sp._HMT_186
S_multispecies_sp01_2_rov_97_976
S_infantis_clade_412
G_Orribacterium
F_Lachnospiraceae
F_Saccharibacteria_ _HMT_ _F_1
G_Saccharibacteria_ _THT_ _ O_1
C_Saccharibacteria_ _THT_ _ C_1
F_Saccharibacteria_ _THT_ _ F_1
S_salivae
S_ingluvei
G_Lancefieldella
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteria
S_parvula
G_Leptotrichia
S_sinus
G_Mogibacterium
S_micronuciformis
S_jalysi
G_Mogibacterium
S_diversum
G_Peptostreptococcus
S_stomatis
S_multispecies_sp03_1
S_sp._HMT_472
S_sp._HMT_473
S_sp._HMT_411
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichi
S_moorei
G_Saccharibacteria_ _THT_ _ G_1
S_bacterium_HMT_952
O_Flavobacteriales
C_Flavobacteriia
S_Pseudopriotrichia
S_sp._HMT_224
S_Eubacterium_ _saburreum
G_Peptostreptococcaceae_ _G_1
S_sp._HMT_165
G_Lachnoanaerobaculum
S_nigrescens
S_periodonticum
S_sp._HMT_931
S_sp._HMT_061
S_sp._HMT_057
S_tanneria
F_Flavobacteriaceae
C_Capnocytophagia
S_multispecies_sp056_2
G_Saccharibacteria_ _THT_ _ G_3
S_bacterium_HMT_345
S_sp._HMT_376
F_Peptostreptococcaceae
C_Tissierellia
G_Parvimonas
O_Tissierellales
S_shahii
S_multispecies_sp018_2
F_Ruminococcaceae
S_elegans
S_sp._HMT_305_rov_93_805
S_koreensis
S_parahaemolytica
S_durus
S_haesulus
S_histicola_rov_97_555
G_Lachnospiraceae_ _G_1
S_bacterium_HMT_096
S_sp._HMT_134
F_Weeksellaceae
S_ingniscens
G_Corynebacterium_1
G_Corynebacterium
F_Corynebacteriaceae
G_Ruminococcaceae_ _G_2
S_bacterium_HMT_088
S_anginosus
S_oralis_subsp._dentisani_clade_414
S_sp._HMT_324
G_Neisseria
S_massiliensis
S_bacterium_HMT_351
G_Saccharibacteria_ _THT_ _ G_2
G_Alistipes
F_Aerococcaceae
S_defectivi
S_sp._HMT_911
S_oris
S_sp._HMT_448
S_gingivae
S_multispecies_sp020
S_dentirozi
S_haemolyticus
S_multispecies_sp027
F_Selenomonadaceae
O_Selenomonadales
G_Selenomonas_6
G_Dialister
S_invisus
S_scopos
S_oralis
S_sp._HMT_307
G_Lautropia
F_Burkholderiaceae
O_Burkholderiales
S_mirabilis
S_sp._HMT_180
S_gingivalis
G_Kingella
S_oralis
S_sputorum
S_rava
S_durum_rov_97_695
S_mitis
S_sp._HMT_167
S_sp._HMT_215
S_loadbetter
S_cribrorostralis
F_Abiocondidactaceria_ _SR1_ _F_1
O_Abiocondidactaria_ _SR1_ _O_1
G_Abiocondidactaria_ _SR1_ _G_1
C_Abiocondidactaria_ _SR1_ _C_1
F_Abiocondidactaria_ _SR1_
S_multispecies_sp045_2_rov_96_841
S_meyeri
S_mira
S_periodontalis
S_bacterium_HMT_875
S_mutans
S_sputigen
S_intermedia
G_Moryella
S_sp._HMT_181
S_parasanguinis_clade_411_rov_97_917
S_sp._HMT_035
S_vestibularis
S_sp._HMT_180
S_sp._HMT_166
G_Protococcus
F_Peptococcaceae
S_sp._HMT_139
S_nucleatum_subsp._vincentii
G_Ruminococcaceae_ _G_1
S_bacterium_HMT_074
S_respertilio
S_sp._HMT_206
G_Cloacibacterium
G_Butyricicoccus
S_sp._HMT_455
G_Arachnia
F_Propionibacteriaceae
S_propionicus
G_Propionibacterium
S_sp._HMT_308
S_multispecies_sp019_2
S_turicibacter
S_parvula
S_bacterium_HMT_345
S_sp._HMT_705
S_sp._HMT_180_rov_97_930
S_cristata
S_sp._HMT_936
S_flavescens
S_catarrhalis
S_crassa
S_sp._HMT_212
S_sanguinis_rov_97_782
S_pseudopora
G_Rasculella
S_aeruginea
S_koreyensis
S_oralis
G_Peptostreptococcaceae_ _G_7_
S_bacterium_HMT_922
S_showae
S_pittmaniae
S_haesooki
S_sp._HMT_204
S_parasanguinis_clade_411
S_sp._HMT_513
S_marinoscens
F_Enterobacteriaceae
S_morbillorum
S_longum
G_Stomatobaculum
G_Ekaterella
S_corrodens
S_surfactantifaciens
S_multispecies_sp08_2
S_virgini
S_subflava
S_sp._HMT_423
S_parvula
G_periodontium
S_elysicola
F_Neisseriaceae
G_Serratia
S_gordonii
S_flavus
S_aphrophilus
O_Enterobacteriales
S_concisus
G_Campylobacter
F_Campylobacteriaceae
O_Campylobacterales
C_Epsilonproteobacteria
S_multispecies_sp024_2
S_nucleatum
S_sanguinis
O_Fusobacteriales
C_Fusobacteria
F_Fusobacteria
O_Aggregatibacter
O_Fusobacterium
F_Fusobacteriaceae
S_multispecies_sp02_2
G_Negativicutes
F_Veillonellaceae
O_Veillonellales
G_Veillonella
O_Haemophilus
S_parainfluenzae
S_flavescens
F_Pasteurellaceae
O_Pasteurellales
C_Gammaproteobacteria
G_Neisseria
C_Betaproteobacteria
O_Neisseriales
P_Proteobacteria
−5.78
−4.66
−3.54
−2.42
−1.30
−0.18
0.94
2.06
3.18
4.30
5.42
LDA SCORE (log 10)