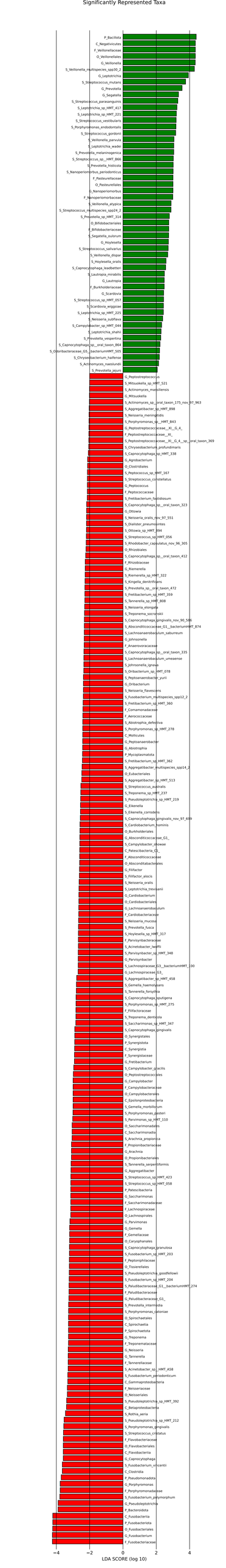
Significantly Represented Taxa
P_Bacillota
C_Negativicutes
F_Veillonellaceae
O_Veillonellales
G_Veillonella
S_Veillonella_multispecies_spp30_2
G_Leptotrichia
S_Streptococcus_mutans
G_Prevotella
G_Segatella
S_Streptococcus_parasanguinis
S_Leptotrichia_sp_HMT_417
S_Leptotrichia_sp_HMT_221
S_Streptococcus_vestibularis
S_Porphyromonas_endodontalis
S_Streptococcus_gordonii
S_Veillonella_parvula
S_Leptotrichia_wadei
S_Prevotella_melaninogenica
S_Streptococcus_sp._HMT_866
S_Prevotella_histicola
S_Nanoperiomorbus_periodonticus
F_Pasteurellaceae
O_Pasteurellales
G_Nanoperiomorbus
F_Nanoperiomorbaceae
S_Veillonella_atypica
S_Streptococcus_multispecies_spp24_2
S_Prevotella_sp_HMT_314
O_Bifidobacteriales
F_Bifidobacteriaceae
S_Segatella_oulorum
G_Hoylesella
S_Streptococcus_salivarius
S_Veillonella_dispar
S_Hoylesella_oralis
S_Capnocytophaga_leadbetteri
S_Lautropia_mirabilis
G_Lautropia
F_Burkholderiaceae
G_Scardovia
S_Streptococcus_sp_HMT_057
S_Scardovia_wiggsiae
S_Leptotrichia_sp_HMT_225
S_Neisseria_subflava
S_Campylobacter_sp_HMT_044
S_Leptotrichia_shahii
S_Prevotella_vespertina
S_Capnocytophaga_sp._oral_taxon_864
S_Oribacteraceae_G5_ bacteriumHMT_505
S_Chryseobacterium_hallenses
S_Actinomyces_naeslundii
S_Prevotella_jejuni
G_Peptostreptococcus
S_Mitsuokella_sp_HMT_521
S_Actinomyces_massiliensis
G_Mitsuokella
S_Actinomyces_sp._oral_taxon_175_nov_97_963
S_Aggregatibacter_sp_HMT_898
S_Neisseria_meningitidis
S_Porphyromonas_sp._HMT_843
G_Peptostreptococcaceae__XI_G_4_
F_Peptostreptococcaceae__XI_
S_Peptostreptococcaceae__XI_G_4_ sp._oral_taxon_369
S_Chryseobacterium_profundimaris
S_Capnocytophaga_sp_HMT_338
G_Agrobacterium
G_Clostridiales
S_Peptococcus_sp_HMT_167
S_Streptococcus_constellatus
G_Peptococcus
F_Peptococcaceae
S_Fretibacterium_fastidiosum
S_Capnocytophaga_sp._oral_taxon_323
G_Ottowia
S_Neisseria_oralis_nov_97_551
S_Dialister_pneumosintes
S_Ottowia_sp_HMT_394
S_Streptococcus_sp_HMT_056
S_Rhodobacter_capsulatus_nov_96_305
O_Rhizobiales
S_Capnocytophaga_sp._oral_taxon_412
F_Rhizobiaceae
G_Riemerella
S_Riemerella_sp_HMT_322
S_Kingella_denitrificans
S_Prevotella_sp._oral_taxon_472
S_Fretibacterium_sp_HMT_359
S_Tannerella_sp_HMT_808
S_Neisseria_elongata
S_Treponema_socranskii
S_Capnocytophaga_gingivalis_nov_90_586
S_Absconditicoccaceae_G1_ bacteriumHMT_874
S_Lachnoanaerobaculum_saburreum
G_Johnsonella
F_Anaerovoracaceae
S_Capnocytophaga_sp._oral_taxon_335
S_Lachnoanaerobaculum_umeaense
S_Johnsonella_ignava
S_Oribacterium_sp_HMT_078
S_Peptoanaerobacter_yurii
G_Oribacterium
S_Neisseria_flavescens
S_Fusobacterium_multispecies_spp12_2
S_Fretibacterium_sp_HMT_360
F_Comamonadaceae
F_Aerococcaceae
S_Alloistophia_defectiva
S_Porphyromonas_sp_HMT_278
C_Mollicutes
G_Peptoanaerobacter
G_Alloistophia
P_Mycoplasmatota
S_Fretibacterium_sp_HMT_362
S_Aggregatibacter_multispecies_spp14_2
O_Eubacteriales
S_Aggregatibacter_sp_HMT_513
S_Streptococcus_australis
S_Treponema_sp_HMT_237
S_Pseudobjotrichia_sp_HMT_219
G_Eikenella
S_Eikenella_corrodens
S_Capnocytophaga_gingivalis_nov_97_699
S_Cardiobacterium_hominis
O_Burkholderiales
G_Absconditicoccaceae_G1_
S_Campylobacter_showae
G_Patescibacteria_G1_
F_Absconditicoccaceae
O_Absconditabacteriales
G_Filifactor
S_Filifactor_alocis
S_Neisseria_oralis
S_Leptotrichia_trevisanii
G_Cardiobacterium
O_Cardiobacteriales
G_Lachnoanaerobaculum
F_Cardiobacteriaceae
S_Neisseria_mucosa
S_Prevotella_fusca
S_Hoylesella_sp_HMT_317
F_Pervisynbacteraceae
S_Acinetobacter_lwoffii
S_Pervisynbacter_sp_HMT_348
G_Pervisynbacter
S_Lachnospiraceae_G3_ bacteriumHMT_100
G_Lachnospiraceae_G3_
S_Aggregatibacter_sp_HMT_458
S_Gemella_haemolysans
S_Tannerella_forsythia
S_Capnocytophaga_sputigena
S_Porphyromonas_sp_HMT_275
F_Filifactoraceae
S_Treponema_denticola
S_Saccharimonas_sp_HMT_347
S_Capnocytophaga_gingivalis
O_Synergistales
P_Synergistota
C_Synergistia
F_Synergistaceae
G_Fretibacterium
S_Campylobacter_gracilis
O_Peptostreptococcales
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Gemella_morbillorum
S_Porphyromonas_gasteri
S_Parvimonas_sp_HMT_110
O_Saccharimonadales
G_Saccharimonadia
S_Arachnia_propionica
F_Propionibacteriaceae
G_Arachnia
O_Propionibacteriales
S_Tannerella_serpentiformis
G_Aggregatibacter
S_Streptococcus_sp_HMT_423
S_Streptococcus_sp_HMT_058
P_Patescibacteria
G_Saccharimonas
F_Saccharimonadaceae
F_Lachnospiraceae
O_Lachnospirales
G_Parvimonas
G_Gemella
F_Gemellaceae
O_Caryophanales
S_Capnocytophaga_granulosa
S_Fusobacterium_sp_HMT_203
F_Peptoniphilaceae
O_Tissierellales
S_Pseudobjotrichia_goodfellowii
S_Fusobacterium_sp_HMT_204
S_Paludibacteraceae_G1_ bacteriumHMT_274
F_Paludibacteraceae
G_Paludibacteraceae_G1_
S_Prevotella_intermedia
S_Porphyromonas_catoniae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetota
G_Treponema
F_Treponemataceae
G_Neisseria
G_Tannerella
F_Tannerellaceae
S_Acinetobacter_sp._HMT_858
S_Fusobacterium_periodonticum
C_Gammaproteobacteria
F_Neisseriaceae
O_Neisseriales
S_Pseudobjotrichia_sp_HMT_392
C_Betaproteobacteria
S_Rothia_aeria
S_Pseudobjotrichia_sp_HMT_212
S_Porphyromonas_gingivalis
S_Streptococcus_cristatus
F_Flavobacteriaceae
O_Flavobacteriales
C_Flavobacteriia
G_Capnocytophaga
S_Fusobacterium_vincentii
C_Clostridia
P_Pseudomonadota
G_Porphyromonas
F_Porphyromonadaceae
S_Fusobacterium_polymorphum
G_Pseudobjotrichia
P_Bacteroidota
C_Fusobacteriia
P_Fusobacteriota
O_Fusobacteriales
G_Fusobacterium
F_Fusobacteriaceae
−4
−2
0
2
4
LDA SCORE (log 10)