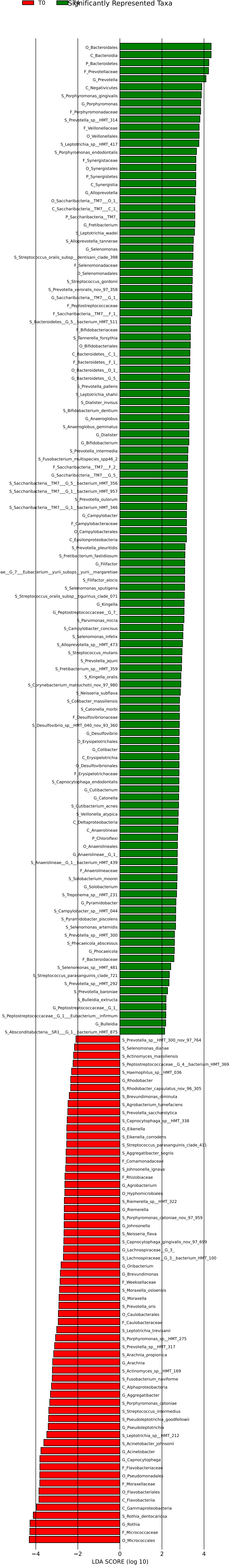
T0
T4
Significantly Represented Taxa
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
F_Prevotellaceae
G_Prevotella
C_Negativicutes
S_Porphyromonas_gingivalis
G_Porphyromonas
F_Porphyromonadaceae
S_Prevotella_sp__HMT_314
F_Veillonellaceae
O_Veillonellales
S_Leptotrichia_sp__HMT_417
S_Porphyromonas_endodontalis
F_Synergistaceae
O_Synergistales
P_Synergistetes
C_Synergistia
G_Alloprevotella
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
G_Fretibacterium
S_Leptotrichia_wadei
S_Alloprevotella_tannerae
G_Selenomonas
S_Streptococcus_oralis_subsp__dentisani_clade_398
F_Selenomonadaceae
O_Selenomonadales
S_Streptococcus_gordonii
S_Prevotella_veroralis_nov_97_358
G_Saccharibacteria__TM7___G_1_
F_Peptostreptococcaceae
F_Saccharibacteria__TM7___F_1_
S_Bacteroidetes__G_5__bacterium_HMT_511
F_Bifidobacteriaceae
S_Tannerella_forsythia
O_Bifidobacteriales
C_Bacteroidetes__C_1_
F_Bacteroidetes__F_1_
O_Bacteroidetes__O_1_
G_Bacteroidetes__G_5_
S_Prevotella_pallens
S_Leptotrichia_shahii
S_Dialister_invisus
S_Bifidobacterium_dentium
G_Anaeroglobus
S_Anaeroglobus_geminatus
G_Dialister
G_Bifidobacterium
S_Prevotella_intermedia
S_Fusobacterium_multispecies_spp46_2
F_Saccharibacteria__TM7___F_2_
G_Saccharibacteria__TM7___G_5_
S_Saccharibacteria__TM7___G_5__bacterium_HMT_356
S_Saccharibacteria__TM7___G_1__bacterium_HMT_957
S_Prevotella_oulorum
S_Saccharibacteria__TM7___G_1__bacterium_HMT_346
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Prevotella_pleuritidis
S_Fretibacterium_fastidiosum
G_Filifactor
eptostreptococcaceae__G_7___Eubacterium__yurii_subsps__yurii__margaretiae
S_Filifactor_alocis
S_Selenomonas_sputigena
S_Streptococcus_oralis_subsp__tigurinus_clade_071
G_Kingella
G_Peptostreptococcaceae__G_7_
S_Parvimonas_micra
S_Campylobacter_concisus
S_Selenomonas_infelix
S_Alloprevotella_sp__HMT_473
S_Streptococcus_mutans
S_Prevotella_jejuni
S_Fretibacterium_sp__HMT_359
S_Kingella_oralis
S_Corynebacterium_matruchotii_nov_97_980
S_Neisseria_subflava
S_Colibacter_massiliensis
S_Catonella_morbi
F_Desulfovibrionaceae
S_Desulfovibrio_sp__HMT_040_nov_93_360
G_Desulfovibrio
O_Erysipelotrichales
G_Colibacter
C_Erysipelotrichia
O_Desulfovibrionales
F_Erysipelotrichaceae
S_Capnocytophaga_endodontalis
G_Cutibacterium
G_Catonella
S_Cutibacterium_acnes
S_Veillonella_atypica
C_Deltaproteobacteria
C_Anaerolineae
P_Chloroflexi
O_Anaerolineales
G_Anaerolineae__G_1_
S_Anaerolineae__G_1__bacterium_HMT_439
F_Anaerolineaceae
S_Solobacterium_moorei
G_Solobacterium
S_Treponema_sp__HMT_231
G_Pyramidobacter
S_Campylobacter_sp__HMT_044
S_Pyramidobacter_piscolens
S_Selenomonas_artemidis
S_Prevotella_sp__HMT_300
S_Phocaeicola_abscessus
G_Phocaeicola
F_Bacteroidaceae
S_Selenomonas_sp__HMT_481
S_Streptococcus_parasanguinis_clade_721
S_Prevotella_sp__HMT_292
S_Prevotella_baroniae
S_Bulleidia_extructa
G_Peptostreptococcaceae__G_1_
S_Peptostreptococcaceae__G_1___Eubacterium__infirmum
G_Bulleidia
S_Absconditabacteria__SR1___G_1__bacterium_HMT_875
S_Prevotella_sp__HMT_300_nov_97_764
S_Selenomonas_dianae
S_Actinomyces_massiliensis
S_Peptostreptococcaceae__G_4__bacterium_HMT_369
S_Haemophilus_sp__HMT_036
G_Rhodobacter
S_Rhodobacter_capsulatus_nov_96_305
S_Brevundimonas_diminuta
S_Agrobacterium_tumefaciens
S_Prevotella_saccharolytica
S_Capnocytophaga_sp__HMT_338
G_Eikenella
S_Eikenella_corrodens
S_Streptococcus_parasanguinis_clade_411
S_Aggregatibacter_segnis
F_Comamonadaceae
S_Johnsonella_ignava
F_Rhizobiaceae
G_Agrobacterium
O_Hyphomicrobiales
S_Riemerella_sp__HMT_322
G_Riemerella
S_Porphyromonas_catoniae_nov_97_959
G_Johnsonella
S_Neisseria_flava
S_Capnocytophaga_gingivalis_nov_97_699
G_Lachnospiraceae__G_3_
S_Lachnospiraceae__G_3__bacterium_HMT_100
G_Oribacterium
G_Brevundimonas
F_Weeksellaceae
S_Moraxella_osloensis
G_Moraxella
S_Prevotella_oris
O_Caulobacterales
F_Caulobacteraceae
S_Leptotrichia_trevisanii
S_Porphyromonas_sp__HMT_275
S_Prevotella_sp__HMT_317
S_Arachnia_propionica
G_Arachnia
S_Actinomyces_sp__HMT_169
S_Fusobacterium_naviforme
C_Alphaproteobacteria
G_Aggregatibacter
S_Porphyromonas_catoniae
S_Streptococcus_intermedius
S_Pseudoleptotrichia_goodfellowii
G_Pseudoleptotrichia
S_Leptotrichia_sp__HMT_212
S_Acinetobacter_johnsonii
G_Acinetobacter
G_Capnocytophaga
F_Flavobacteriaceae
O_Pseudomonadales
F_Moraxellaceae
O_Flavobacteriales
C_Flavobacteriia
C_Gammaproteobacteria
S_Rothia_dentocariosa
G_Rothia
F_Micrococcaceae
O_Micrococcales
−4
−2
0
2
4
LDA SCORE (log 10)