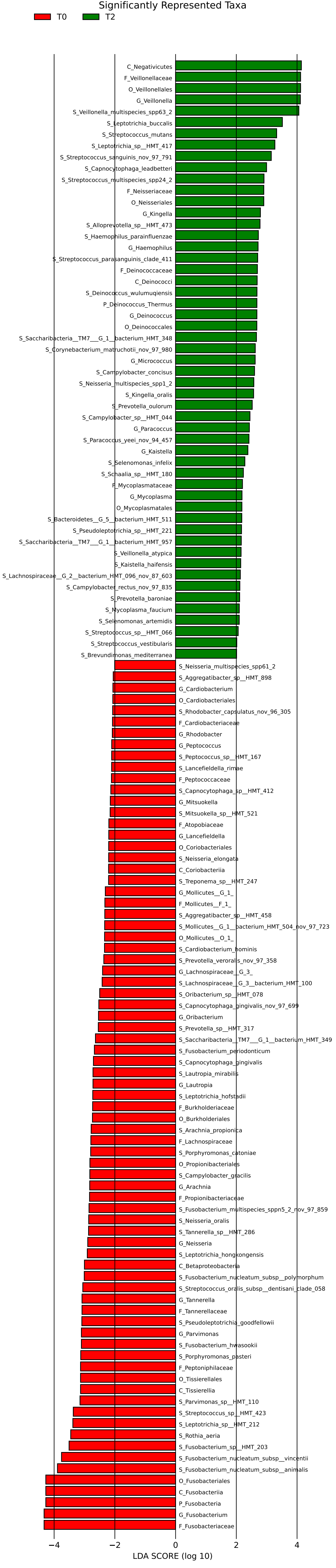
Significantly Represented Taxa
T0
T2
C_Negativicutes
F_Veillonellaceae
O_Veillonellales
G_Veillonella
S_Veillonella_multispecies_spp63_2
S_Leptotrichia_buccalis
S_Streptococcus_mutans
S_Leptotrichia_sp__HMT_417
S_Streptococcus_sanguinis_nov_97_791
S_Capnocytophaga_leadbetteri
S_Streptococcus_multispecies_spp24_2
F_Neisseriaceae
O_Neisseriales
G_Kingella
S_Alloprevotella_sp__HMT_473
S_Haemophilus_parainfluenzae
G_Haemophilus
S_Streptococcus_parasanguinis_clade_411
F_Deinococcaceae
C_Deinococci
S_Deinococcus_wulumuqiensis
P_Deinococcus_Thermus
G_Deinococcus
O_Deinococcales
S_Saccharibacteria__TM7___G_1__bacterium_HMT_348
S_Corynebacterium_matruchotii_nov_97_980
G_Micrococcus
S_Campylobacter_concisus
S_Neisseria_multispecies_spp1_2
S_Kingella_oralis
S_Prevotella_oulorum
S_Campylobacter_sp__HMT_044
G_Paracoccus
S_Paracoccus_yeei_nov_94_457
G_Kaistella
S_Selenomonas_infelix
S_Schaalia_sp__HMT_180
F_Mycoplasmataceae
G_Mycoplasma
O_Mycoplasmatales
S_Bacteroidetes__G_5__bacterium_HMT_511
S_Pseudoleptotrichia_sp__HMT_221
S_Saccharibacteria__TM7___G_1__bacterium_HMT_957
S_Veillonella_atypica
S_Kaistella_haifensis
S_Lachnospiraceae__G_2__bacterium_HMT_096_nov_87_603
S_Campylobacter_rectus_nov_97_835
S_Prevotella_baroniae
S_Mycoplasma_faucium
S_Selenomonas_artemidis
S_Streptococcus_sp__HMT_066
S_Streptococcus_vestibularis
S_Brevundimonas_mediterranea
S_Neisseria_multispecies_spp61_2
S_Aggregatibacter_sp__HMT_898
G_Cardiobacterium
O_Cardiobacteriales
S_Rhodobacter_capsulatus_nov_96_305
F_Cardiobacteriaceae
G_Rhodobacter
G_Peptococcus
S_Peptococcus_sp__HMT_167
S_Lancefieldella_rimae
F_Peptococcaceae
S_Capnocytophaga_sp__HMT_412
G_Mitsuokella
S_Mitsuokella_sp__HMT_521
F_Atopobiaceae
G_Lancefieldella
O_Coriobacteriales
S_Neisseria_elongata
C_Coriobacteriia
S_Treponema_sp__HMT_247
G_Mollicutes__G_1_
F_Mollicutes__F_1_
S_Aggregatibacter_sp__HMT_458
S_Mollicutes__G_1__bacterium_HMT_504_nov_97_723
O_Mollicutes__O_1_
S_Cardiobacterium_hominis
S_Prevotella_veroralis_nov_97_358
G_Lachnospiraceae__G_3_
S_Lachnospiraceae__G_3__bacterium_HMT_100
S_Oribacterium_sp__HMT_078
S_Capnocytophaga_gingivalis_nov_97_699
G_Oribacterium
S_Prevotella_sp__HMT_317
S_Saccharibacteria__TM7___G_1__bacterium_HMT_349
S_Fusobacterium_periodonticum
S_Capnocytophaga_gingivalis
S_Lautropia_mirabilis
G_Lautropia
S_Leptotrichia_hofstadii
F_Burkholderiaceae
O_Burkholderiales
S_Arachnia_propionica
F_Lachnospiraceae
S_Porphyromonas_catoniae
O_Propionibacteriales
S_Campylobacter_gracilis
G_Arachnia
F_Propionibacteriaceae
S_Fusobacterium_multispecies_sppn5_2_nov_97_859
S_Neisseria_oralis
S_Tannerella_sp__HMT_286
G_Neisseria
S_Leptotrichia_hongkongensis
C_Betaproteobacteria
S_Fusobacterium_nucleatum_subsp__polymorphum
S_Streptococcus_oralis_subsp__dentisani_clade_058
G_Tannerella
F_Tannerellaceae
S_Pseudoleptotrichia_goodfellowii
G_Parvimonas
S_Fusobacterium_hwasookii
S_Porphyromonas_pasteri
F_Peptoniphilaceae
O_Tissierellales
C_Tissierellia
S_Parvimonas_sp__HMT_110
S_Streptococcus_sp__HMT_423
S_Leptotrichia_sp__HMT_212
S_Rothia_aeria
S_Fusobacterium_sp__HMT_203
S_Fusobacterium_nucleatum_subsp__vincentii
S_Fusobacterium_nucleatum_subsp__animalis
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
−4
−2
0
2
4
LDA SCORE (log 10)