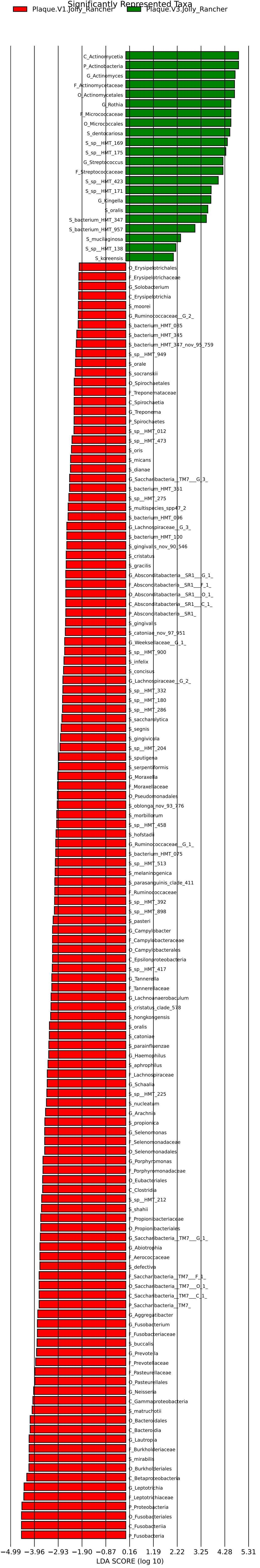
Significantly Represented Taxa
Plaque.V1.Jolly_Rancher
Plaque.V3.Jolly_Rancher
C_Actinomycetia
P_Actinobacteria
G_Actinomyces
F_Actinomycetaceae
O_Actinomycetales
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_dentocariosa
S_sp__HMT_169
S_sp__HMT_175
G_Streptococcus
F_Streptococcaceae
S_sp__HMT_423
S_sp__HMT_171
G_Kingella
S_oralis
S_bacterium_HMT_347
S_bacterium_HMT_957
S_mucilaginosa
S_sp__HMT_138
S_koreensis
O_Erysipelotrichales
F_Erysipelotrichaceae
G_Solobacterium
C_Erysipelotrichia
S_moorei
G_Ruminococcaceae__G_2_
S_bacterium_HMT_085
S_bacterium_HMT_345
S_bacterium_HMT_347_nov_95_759
S_sp__HMT_949
S_orale
S_socranskii
O_Spirochaetales
F_Treponemataceae
C_Spirochaetia
G_Treponema
P_Spirochaetes
S_sp__HMT_012
S_sp__HMT_473
S_oris
S_micans
S_dianae
G_Saccharibacteria__TM7___G_3_
S_bacterium_HMT_351
S_sp__HMT_275
S_multispecies_spp47_2
S_bacterium_HMT_096
G_Lachnospiraceae__G_3_
S_bacterium_HMT_100
S_gingivalis_nov_90_546
S_cristatus
S_gracilis
G_Absconditabacteria__SR1___G_1_
F_Absconditabacteria__SR1___F_1_
O_Absconditabacteria__SR1___O_1_
C_Absconditabacteria__SR1___C_1_
P_Absconditabacteria__SR1_
S_gingivalis
S_catoniae_nov_97_951
G_Weeksellaceae__G_1_
S_sp__HMT_900
S_infelix
S_concisus
G_Lachnospiraceae__G_2_
S_sp__HMT_332
S_sp__HMT_180
S_sp__HMT_286
S_saccharolytica
S_segnis
S_gingivicola
S_sp__HMT_204
S_sputigena
S_serpentiformis
G_Moraxella
F_Moraxellaceae
O_Pseudomonadales
S_oblonga_nov_93_776
S_morbillorum
S_sp__HMT_458
S_hofstadii
G_Ruminococcaceae__G_1_
S_bacterium_HMT_075
S_sp__HMT_513
S_melaninogenica
S_parasanguinis_clade_411
F_Ruminococcaceae
S_sp__HMT_392
S_sp__HMT_898
S_pasteri
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_sp__HMT_417
G_Tannerella
F_Tannerellaceae
G_Lachnoanaerobaculum
S_cristatus_clade_578
S_hongkongensis
S_oralis
S_catoniae
S_parainfluenzae
G_Haemophilus
S_aphrophilus
F_Lachnospiraceae
G_Schaalia
S_sp__HMT_225
S_nucleatum
G_Arachnia
S_propionica
G_Selenomonas
F_Selenomonadaceae
O_Selenomonadales
G_Porphyromonas
F_Porphyromonadaceae
O_Eubacteriales
C_Clostridia
S_sp__HMT_212
S_shahii
F_Propionibacteriaceae
O_Propionibacteriales
G_Saccharibacteria__TM7___G_1_
G_Abiotrophia
F_Aerococcaceae
S_defectiva
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
G_Aggregatibacter
G_Fusobacterium
F_Fusobacteriaceae
S_buccalis
G_Prevotella
F_Prevotellaceae
F_Pasteurellaceae
O_Pasteurellales
G_Neisseria
C_Gammaproteobacteria
S_matruchotii
O_Bacteroidales
C_Bacteroidia
G_Lautropia
F_Burkholderiaceae
S_mirabilis
O_Burkholderiales
C_Betaproteobacteria
G_Leptotrichia
F_Leptotrichiaceae
P_Proteobacteria
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
−4.99
−3.96
−2.93
−1.90
−0.87
0.16
1.19
2.22
3.25
4.28
5.31
LDA SCORE (log 10)