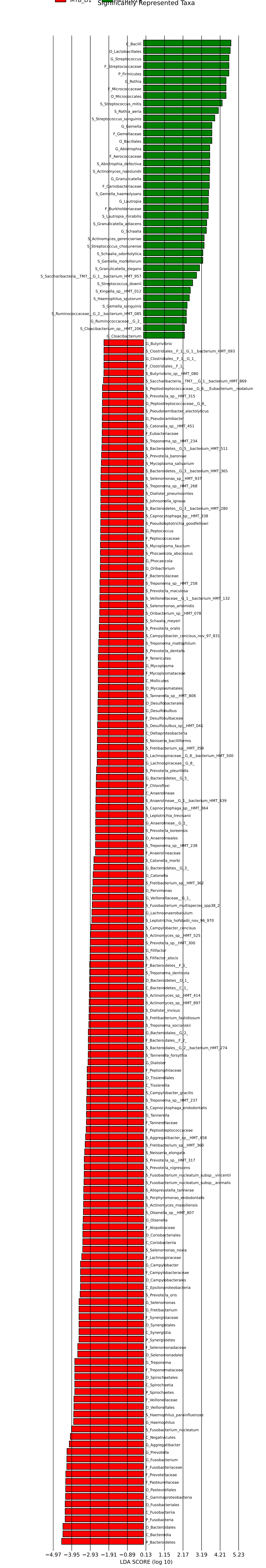
MTB_D1
MTB_D28
Significantly Represented Taxa
C_Bacilli
O_Lactobacillales
G_Streptococcus
F_Streptococcaceae
P_Firmicutes
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_Streptococcus_mitis
S_Rothia_aeria
S_Streptococcus_sanguinis
G_Gemella
F_Gemellaceae
O_Bacillales
G_Abiotrophia
F_Aerococcaceae
S_Abiotrophia_defectiva
S_Actinomyces_naeslundii
G_Granulicatella
F_Carnobacteriaceae
S_Gemella_haemolysans
G_Lautropia
F_Burkholderiaceae
S_Lautropia_mirabilis
S_Granulicatella_adiacens
G_Schaalia
S_Actinomyces_gerencseriae
S_Streptococcus_chosunense
S_Schaalia_odontolytica
S_Gemella_morbillorum
S_Granulicatella_elegans
S_Saccharibacteria__TM7___G_1__bacterium_HMT_957
S_Streptococcus_downii
S_Kingella_sp__HMT_012
S_Haemophilus_sputorum
S_Gemella_sanguinis
S_Ruminococcaceae__G_2__bacterium_HMT_085
G_Ruminococcaceae__G_2_
S_Cloacibacterium_sp__HMT_206
G_Cloacibacterium
G_Butyrivibrio
S_Clostridiales__F_1__G_1__bacterium_HMT_093
G_Clostridiales__F_1__G_1_
F_Clostridiales__F_1_
S_Butyrivibrio_sp__HMT_080
S_Saccharibacteria__TM7___G_1__bacterium_HMT_869
S_Peptostreptococcaceae__G_6___Eubacterium__nodatum
S_Prevotella_sp__HMT_315
G_Peptostreptococcaceae__G_6_
S_Pseudoramibacter_alactolyticus
G_Pseudoramibacter
S_Catonella_sp__HMT_451
F_Eubacteriaceae
S_Treponema_sp__HMT_234
S_Bacteroidetes__G_5__bacterium_HMT_511
S_Prevotella_baroniae
S_Mycoplasma_salivarium
S_Bacteroidetes__G_3__bacterium_HMT_365
S_Selenomonas_sp__HMT_937
S_Treponema_sp__HMT_268
S_Dialister_pneumosintes
S_Johnsonella_ignava
S_Bacteroidetes__G_3__bacterium_HMT_280
S_Capnocytophaga_sp__HMT_338
S_Pseudoleptotrichia_goodfellowii
G_Peptococcus
F_Peptococcaceae
S_Mycoplasma_faucium
S_Phocaeicola_abscessus
G_Phocaeicola
G_Oribacterium
F_Bacteroidaceae
S_Treponema_sp__HMT_258
S_Prevotella_maculosa
S_Veillonellaceae__G_1__bacterium_HMT_132
S_Selenomonas_artemidis
S_Oribacterium_sp__HMT_078
S_Schaalia_meyeri
S_Prevotella_oralis
S_Campylobacter_concisus_nov_97_831
S_Treponema_maltophilum
S_Prevotella_dentalis
P_Tenericutes
G_Mycoplasma
F_Mycoplasmataceae
C_Mollicutes
O_Mycoplasmatales
S_Tannerella_sp__HMT_808
O_Desulfobacterales
G_Desulfobulbus
F_Desulfobulbaceae
S_Desulfobulbus_sp__HMT_041
C_Deltaproteobacteria
S_Neisseria_bacilliformis
S_Fretibacterium_sp__HMT_359
S_Lachnospiraceae__G_8__bacterium_HMT_500
G_Lachnospiraceae__G_8_
S_Prevotella_pleuritidis
G_Bacteroidetes__G_5_
P_Chloroflexi
C_Anaerolineae
S_Anaerolineae__G_1__bacterium_HMT_439
S_Capnocytophaga_sp__HMT_864
S_Leptotrichia_trevisanii
G_Anaerolineae__G_1_
S_Prevotella_koreensis
O_Anaerolineales
S_Treponema_sp__HMT_238
F_Anaerolineaceae
S_Catonella_morbi
G_Bacteroidetes__G_3_
G_Catonella
S_Fretibacterium_sp__HMT_362
G_Parvimonas
G_Veillonellaceae__G_1_
S_Fusobacterium_multispecies_spp38_2
G_Lachnoanaerobaculum
S_Leptotrichia_hofstadii_nov_96_970
S_Campylobacter_concisus
S_Actinomyces_sp__HMT_525
S_Prevotella_sp__HMT_300
G_Filifactor
S_Filifactor_alocis
F_Bacteroidetes__F_1_
S_Treponema_denticola
O_Bacteroidetes__O_1_
C_Bacteroidetes__C_1_
S_Actinomyces_sp__HMT_414
S_Actinomyces_sp__HMT_897
S_Dialister_invisus
S_Fretibacterium_fastidiosum
S_Treponema_socranskii
G_Bacteroidales__G_2_
F_Bacteroidales__F_2_
S_Bacteroidales__G_2__bacterium_HMT_274
S_Tannerella_forsythia
G_Dialister
F_Peptoniphilaceae
O_Tissierellales
C_Tissierellia
S_Campylobacter_gracilis
S_Treponema_sp__HMT_237
S_Capnocytophaga_endodontalis
G_Tannerella
F_Tannerellaceae
F_Peptostreptococcaceae
S_Aggregatibacter_sp__HMT_458
S_Fretibacterium_sp__HMT_360
S_Neisseria_elongata
S_Prevotella_sp__HMT_317
S_Prevotella_nigrescens
S_Fusobacterium_nucleatum_subsp__vincentii
S_Fusobacterium_nucleatum_subsp__animalis
S_Alloprevotella_tannerae
S_Porphyromonas_endodontalis
S_Actinomyces_massiliensis
S_Olsenella_sp__HMT_807
G_Olsenella
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteriia
S_Selenomonas_noxia
F_Lachnospiraceae
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_Prevotella_oris
G_Selenomonas
G_Fretibacterium
F_Synergistaceae
O_Synergistales
C_Synergistia
P_Synergistetes
F_Selenomonadaceae
O_Selenomonadales
G_Treponema
F_Treponemataceae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetes
F_Veillonellaceae
O_Veillonellales
S_Haemophilus_parainfluenzae
G_Haemophilus
S_Fusobacterium_nucleatum
C_Negativicutes
G_Aggregatibacter
G_Prevotella
G_Fusobacterium
F_Fusobacteriaceae
F_Prevotellaceae
F_Pasteurellaceae
O_Pasteurellales
C_Gammaproteobacteria
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
−4.97
−3.95
−2.93
−1.91
−0.89
0.13
1.15
2.17
3.19
4.21
5.23
LDA SCORE (log 10)