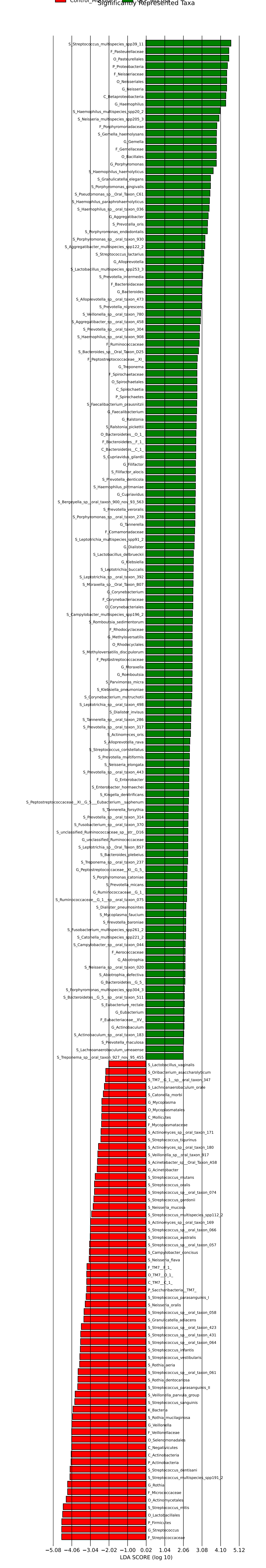
Control_Mucosa
CD_Mucosa
Significantly Represented Taxa
S_Streptococcus_multispecies_spp39_11
F_Pasteurellaceae
O_Pasteurellales
P_Proteobacteria
F_Neisseriaceae
O_Neisseriales
G_Neisseria
C_Betaproteobacteria
G_Haemophilus
S_Haemophilus_multispecies_spp20_2
S_Neisseria_multispecies_spp205_3
F_Porphyromonadaceae
S_Gemella_haemolysans
G_Gemella
F_Gemellaceae
O_Bacillales
G_Porphyromonas
S_Haemophilus_haemolyticus
S_Granulicatella_elegans
S_Porphyromonas_gingivalis
S_Pseudomonas_sp__Oral_Taxon_C61
S_Haemophilus_paraphrohaemolyticus
S_Haemophilus_sp__oral_taxon_036
G_Aggregatibacter
S_Prevotella_oris
S_Porphyromonas_endodontalis
S_Porphyromonas_sp__oral_taxon_930
S_Aggregatibacter_multispecies_spp122_2
S_Streptococcus_lactarius
G_Alloprevotella
S_Lactobacillus_multispecies_spp253_3
S_Prevotella_intermedia
F_Bacteroidaceae
G_Bacteroides
S_Alloprevotella_sp__oral_taxon_473
S_Prevotella_nigrescens
S_Veillonella_sp__oral_taxon_780
S_Aggregatibacter_sp__oral_taxon_458
S_Prevotella_sp__oral_taxon_304
S_Haemophilus_sp__oral_taxon_908
F_Ruminococcaceae
S_Bacteroides_sp__Oral_Taxon_D25
F_Peptostreptococcaceae__XI_
G_Treponema
F_Spirochaetaceae
O_Spirochaetales
C_Spirochaetia
P_Spirochaetes
S_Faecalibacterium_prausnitzii
G_Faecalibacterium
G_Ralstonia
S_Ralstonia_pickettii
O_Bacteroidetes__O_1_
F_Bacteroidetes__F_1_
C_Bacteroidetes__C_1_
S_Cupriavidus_gilardii
G_Filifactor
S_Filifactor_alocis
S_Prevotella_denticola
S_Haemophilus_pittmaniae
G_Cupriavidus
S_Bergeyella_sp__oral_taxon_900_nov_93_563
S_Prevotella_veroralis
S_Porphyromonas_sp__oral_taxon_278
G_Tannerella
F_Comamonadaceae
S_Leptotrichia_multispecies_spp91_2
G_Dialister
S_Lactobacillus_delbrueckii
G_Klebsiella
S_Leptotrichia_buccalis
S_Leptotrichia_sp__oral_taxon_392
S_Moraxella_sp__Oral_Taxon_B07
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_Campylobacter_multispecies_spp196_2
S_Romboutsia_sedimentorum
F_Rhodocyclaceae
G_Methyloversatilis
O_Rhodocyclales
S_Methyloversatilis_discipulorum
F_Peptostreptococcaceae
G_Moraxella
G_Romboutsia
S_Parvimonas_micra
S_Klebsiella_pneumoniae
S_Corynebacterium_matruchotii
S_Leptotrichia_sp__oral_taxon_498
S_Dialister_invisus
S_Tannerella_sp__oral_taxon_286
S_Prevotella_sp__oral_taxon_317
S_Actinomyces_oris
S_Alloprevotella_rava
S_Streptococcus_constellatus
S_Prevotella_multiformis
S_Neisseria_elongata
S_Prevotella_sp__oral_taxon_443
G_Enterobacter
S_Enterobacter_hormaechei
S_Kingella_denitrificans
S_Peptostreptococcaceae__XI__G_5___Eubacterium__saphenum
S_Tannerella_forsythia
S_Prevotella_sp__oral_taxon_314
S_Fusobacterium_sp__oral_taxon_370
S_unclassified_Ruminococcaceae_sp__str__D16
G_unclassified_Ruminococcaceae
S_Leptotrichia_sp__Oral_Taxon_B57
S_Bacteroides_plebeius
S_Treponema_sp__oral_taxon_237
G_Peptostreptococcaceae__XI__G_5_
S_Porphyromonas_catoniae
S_Prevotella_micans
G_Ruminococcaceae__G_1_
S_Ruminococcaceae__G_1__sp__oral_taxon_075
S_Dialister_pneumosintes
S_Mycoplasma_faucium
S_Prevotella_baroniae
S_Fusobacterium_multispecies_spp261_2
S_Catonella_multispecies_spp221_2
S_Campylobacter_sp__oral_taxon_044
F_Aerococcaceae
G_Abiotrophia
S_Neisseria_sp__oral_taxon_020
S_Abiotrophia_defectiva
G_Bacteroidetes__G_5_
S_Porphyromonas_multispecies_spp304_3
S_Bacteroidetes__G_5__sp__oral_taxon_511
S_Eubacterium_rectale
G_Eubacterium
F_Eubacteriaceae__XV_
G_Actinobaculum
S_Actinobaculum_sp__oral_taxon_183
S_Prevotella_maculosa
S_Lachnoanaerobaculum_umeaense
S_Treponema_sp__oral_taxon_927_nov_95_455
S_Lactobacillus_vaginalis
S_Oribacterium_asaccharolyticum
S_TM7__G_1__sp__oral_taxon_347
S_Lachnoanaerobaculum_orale
S_Catonella_morbi
G_Mycoplasma
O_Mycoplasmatales
C_Mollicutes
F_Mycoplasmataceae
S_Actinomyces_sp__oral_taxon_171
S_Streptococcus_tigurinus
S_Actinomyces_sp__oral_taxon_180
S_Veillonella_sp__oral_taxon_917
S_Acinetobacter_sp__Oral_Taxon_A58
G_Acinetobacter
S_Streptococcus_mutans
S_Streptococcus_oralis
S_Streptococcus_sp__oral_taxon_074
S_Streptococcus_gordonii
S_Neisseria_mucosa
S_Streptococcus_multispecies_spp112_2
S_Actinomyces_sp__oral_taxon_169
S_Streptococcus_sp__oral_taxon_066
S_Streptococcus_australis
S_Streptococcus_sp__oral_taxon_057
S_Campylobacter_concisus
S_Neisseria_flava
F_TM7__F_1_
O_TM7__O_1_
C_TM7__C_1_
P_Saccharibacteria__TM7_
S_Streptococcus_parasanguinis_I
S_Neisseria_oralis
S_Streptococcus_sp__oral_taxon_058
S_Granulicatella_adiacens
S_Streptococcus_sp__oral_taxon_423
S_Streptococcus_sp__oral_taxon_431
S_Streptococcus_sp__oral_taxon_064
S_Streptococcus_infantis
S_Streptococcus_vestibularis
S_Rothia_aeria
S_Streptococcus_sp__oral_taxon_061
S_Rothia_dentocariosa
S_Streptococcus_parasanguinis_II
S_Veillonella_parvula_group
S_Streptococcus_sanguinis
K_Bacteria
S_Rothia_mucilaginosa
G_Veillonella
F_Veillonellaceae
O_Selenomonadales
C_Negativicutes
C_Actinobacteria
P_Actinobacteria
S_Streptococcus_dentisani
S_Streptococcus_multispecies_spp191_2
G_Rothia
F_Micrococcaceae
O_Actinomycetales
S_Streptococcus_mitis
O_Lactobacillales
P_Firmicutes
G_Streptococcus
F_Streptococcaceae
−5.08
−4.06
−3.04
−2.02
−1.00
0.02
1.04
2.06
3.08
4.10
5.12
LDA SCORE (log 10)