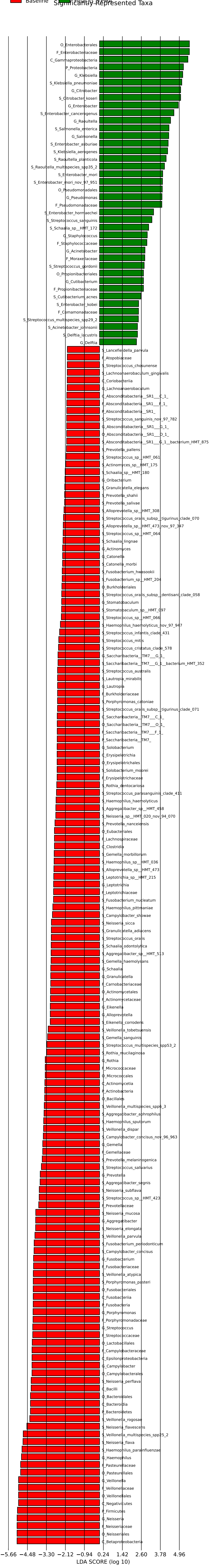
Baseline
Significantly Represented Taxa
O_Enterobacterales
F_Enterobacteriaceae
C_Gammaproteobacteria
P_Proteobacteria
G_Klebsiella
S_Klebsiella_pneumoniae
G_Citrobacter
S_Citrobacter_koseri
G_Enterobacter
S_Enterobacter_cancerogenus
G_Raoultella
S_Salmonella_enterica
G_Salmonella
S_Enterobacter_asburiae
S_Klebsiella_aerogenes
S_Raoultella_planticola
S_Raoultella_multispecies_spp35_2
S_Enterobacter_mori
S_Enterobacter_mori_nov_97_951
O_Pseudomonadales
G_Pseudomonas
F_Pseudomonadaceae
S_Enterobacter_hormaechei
S_Streptococcus_sanguinis
S_Schaalia_sp__HMT_172
G_Staphylococcus
F_Staphylococcaceae
G_Acinetobacter
F_Moraxellaceae
S_Streptococcus_gordonii
O_Propionibacteriales
G_Cutibacterium
F_Propionibacteriaceae
S_Cutibacterium_acnes
S_Enterobacter_kobei
F_Comamonadaceae
S_Streptococcus_multispecies_spp29_2
S_Acinetobacter_johnsonii
S_Delftia_lacustris
G_Delftia
S_Lancefieldella_parvula
F_Atopobiaceae
S_Streptococcus_chosunense
S_Lachnoanaerobaculum_gingivalis
C_Coriobacteriia
G_Lachnoanaerobaculum
C_Absconditabacteria__SR1___C_1_
F_Absconditabacteria__SR1___F_1_
P_Absconditabacteria__SR1_
S_Streptococcus_sanguinis_nov_97_782
G_Absconditabacteria__SR1___G_1_
O_Absconditabacteria__SR1___O_1_
S_Absconditabacteria__SR1___G_1__bacterium_HMT_875
S_Prevotella_pallens
S_Streptococcus_sp__HMT_061
S_Actinomyces_sp__HMT_175
S_Schaalia_sp__HMT_180
G_Oribacterium
S_Granulicatella_elegans
S_Prevotella_shahii
S_Prevotella_salivae
S_Alloprevotella_sp__HMT_308
S_Streptococcus_oralis_subsp__tigurinus_clade_070
S_Alloprevotella_sp__HMT_473_nov_97_347
S_Streptococcus_sp__HMT_064
S_Schaalia_lingnae
G_Actinomyces
G_Catonella
S_Catonella_morbi
S_Fusobacterium_hwasookii
S_Fusobacterium_sp__HMT_204
O_Burkholderiales
S_Streptococcus_oralis_subsp__dentisani_clade_058
G_Stomatobaculum
S_Stomatobaculum_sp__HMT_097
S_Streptococcus_sp__HMT_066
S_Haemophilus_haemolyticus_nov_97_947
S_Streptococcus_infantis_clade_431
S_Streptococcus_mitis
S_Streptococcus_cristatus_clade_578
G_Saccharibacteria__TM7___G_1_
S_Saccharibacteria__TM7___G_1__bacterium_HMT_352
S_Streptococcus_australis
S_Lautropia_mirabilis
G_Lautropia
F_Burkholderiaceae
S_Porphyromonas_catoniae
S_Streptococcus_oralis_subsp__tigurinus_clade_071
C_Saccharibacteria__TM7___C_1_
O_Saccharibacteria__TM7___O_1_
F_Saccharibacteria__TM7___F_1_
P_Saccharibacteria__TM7_
G_Solobacterium
C_Erysipelotrichia
O_Erysipelotrichales
S_Solobacterium_moorei
F_Erysipelotrichaceae
S_Rothia_dentocariosa
S_Streptococcus_parasanguinis_clade_411
S_Haemophilus_haemolyticus
S_Aggregatibacter_sp__HMT_458
S_Neisseria_sp__HMT_020_nov_94_070
S_Prevotella_nanceiensis
O_Eubacteriales
F_Lachnospiraceae
C_Clostridia
S_Gemella_morbillorum
S_Haemophilus_sp__HMT_036
S_Alloprevotella_sp__HMT_473
S_Leptotrichia_sp__HMT_215
G_Leptotrichia
F_Leptotrichiaceae
S_Fusobacterium_nucleatum
S_Haemophilus_pittmaniae
S_Campylobacter_showae
S_Neisseria_sicca
S_Granulicatella_adiacens
S_Streptococcus_oralis
S_Schaalia_odontolytica
S_Aggregatibacter_sp__HMT_513
S_Gemella_haemolysans
G_Schaalia
G_Granulicatella
F_Carnobacteriaceae
O_Actinomycetales
F_Actinomycetaceae
G_Eikenella
G_Alloprevotella
S_Eikenella_corrodens
S_Veillonella_tobetsuensis
S_Gemella_sanguinis
S_Streptococcus_multispecies_spp53_2
S_Rothia_mucilaginosa
G_Rothia
F_Micrococcaceae
O_Micrococcales
C_Actinomycetia
P_Actinobacteria
O_Bacillales
S_Veillonella_multispecies_spp6_3
S_Aggregatibacter_aphrophilus
S_Haemophilus_sputorum
S_Veillonella_dispar
S_Campylobacter_concisus_nov_96_963
G_Gemella
F_Gemellaceae
S_Prevotella_melaninogenica
S_Streptococcus_salivarius
G_Prevotella
S_Aggregatibacter_segnis
S_Neisseria_subflava
S_Streptococcus_sp__HMT_423
F_Prevotellaceae
S_Neisseria_mucosa
G_Aggregatibacter
S_Neisseria_elongata
S_Veillonella_parvula
S_Fusobacterium_periodonticum
S_Campylobacter_concisus
G_Fusobacterium
F_Fusobacteriaceae
S_Veillonella_atypica
S_Porphyromonas_pasteri
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Porphyromonas
F_Porphyromonadaceae
G_Streptococcus
F_Streptococcaceae
O_Lactobacillales
F_Campylobacteraceae
C_Epsilonproteobacteria
G_Campylobacter
O_Campylobacterales
S_Neisseria_perflava
C_Bacilli
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes
S_Veillonella_rogosae
S_Neisseria_flavescens
S_Veillonella_multispecies_spp25_2
S_Neisseria_flava
S_Haemophilus_parainfluenzae
G_Haemophilus
F_Pasteurellaceae
O_Pasteurellales
G_Veillonella
F_Veillonellaceae
O_Veillonellales
C_Negativicutes
P_Firmicutes
G_Neisseria
F_Neisseriaceae
O_Neisseriales
C_Betaproteobacteria
−5.66
−4.48
−3.30
−2.12
−0.94
0.24
1.42
2.60
3.78
4.96
LDA SCORE (log 10)