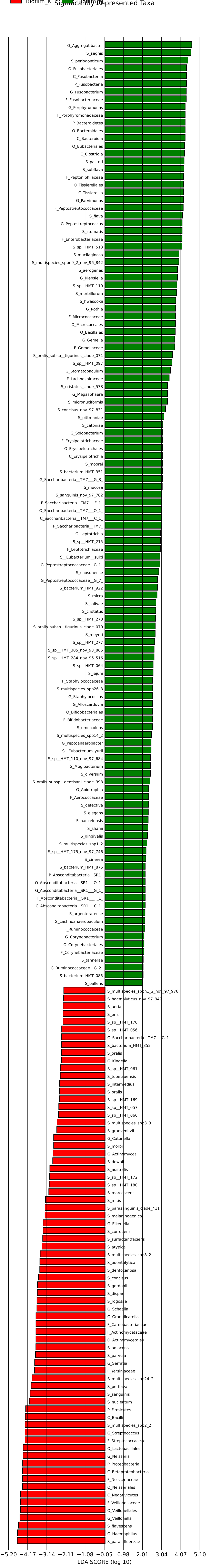
Biofilm_K
Biofilm_M
Significantly Represented Taxa
G_Aggregatibacter
S_segnis
S_periodonticum
O_Fusobacteriales
C_Fusobacteriia
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
G_Porphyromonas
F_Porphyromonadaceae
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
O_Eubacteriales
C_Clostridia
S_pasteri
S_subflava
F_Peptoniphilaceae
O_Tissierellales
C_Tissierellia
G_Parvimonas
F_Peptostreptococcaceae
S_flava
G_Peptostreptococcus
S_stomatis
F_Enterobacteriaceae
S_sp__HMT_513
S_mucilaginosa
S_multispecies_sppn9_2_nov_96_842
S_aerogenes
G_Klebsiella
S_sp__HMT_110
S_morbillorum
S_hwasookii
G_Rothia
F_Micrococcaceae
O_Micrococcales
O_Bacillales
G_Gemella
F_Gemellaceae
S_oralis_subsp__tigurinus_clade_071
S_sp__HMT_097
G_Stomatobaculum
F_Lachnospiraceae
S_cristatus_clade_578
G_Megasphaera
S_micronuciformis
S_concisus_nov_97_831
S_pittmaniae
S_catoniae
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_moorei
S_bacterium_HMT_351
G_Saccharibacteria__TM7___G_3_
S_mucosa
S_sanguinis_nov_97_782
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
G_Leptotrichia
S_sp__HMT_215
F_Leptotrichiaceae
S__Eubacterium__sulci
G_Peptostreptococcaceae__G_1_
S_chosunense
G_Peptostreptococcaceae__G_7_
S_bacterium_HMT_922
S_micra
S_salivae
S_cristatus
S_sp__HMT_278
S_oralis_subsp__tigurinus_clade_070
S_meyeri
S_sp__HMT_277
S_sp__HMT_305_nov_93_865
S_sp__HMT_284_nov_96_516
S_sp__HMT_064
S_jejuni
F_Staphylococcaceae
S_multispecies_spp26_3
G_Staphylococcus
G_Alloscardovia
O_Bifidobacteriales
F_Bifidobacteriaceae
S_omnicolens
S_multispecies_spp14_2
G_Peptoanaerobacter
S__Eubacterium_yurii
S_sp__HMT_110_nov_97_684
G_Mogibacterium
S_diversum
S_oralis_subsp__dentisani_clade_398
G_Abiotrophia
F_Aerococcaceae
S_defectiva
S_elegans
S_nanceiensis
S_shahii
S_gingivalis
S_multispecies_spp1_2
S_sp__HMT_175_nov_97_746
S_cinerea
S_bacterium_HMT_875
P_Absconditabacteria__SR1_
O_Absconditabacteria__SR1___O_1_
G_Absconditabacteria__SR1___G_1_
F_Absconditabacteria__SR1___F_1_
C_Absconditabacteria__SR1___C_1_
S_argentoratense
G_Lachnoanaerobaculum
F_Ruminococcaceae
G_Corynebacterium
O_Corynebacteriales
F_Corynebacteriaceae
S_tannerae
G_Ruminococcaceae__G_2_
S_bacterium_HMT_085
S_pallens
S_multispecies_sppn1_2_nov_97_976
S_haemolyticus_nov_97_947
S_aeria
S_oris
S_sp__HMT_170
S_sp__HMT_056
G_Saccharibacteria__TM7___G_1_
S_bacterium_HMT_352
S_oralis
G_Kingella
S_sp__HMT_061
S_tobetsuensis
S_intermedius
S_oralis
S_sp__HMT_169
S_sp__HMT_057
S_sp__HMT_066
S_multispecies_spp3_3
S_graevenitzii
G_Catonella
S_morbi
G_Actinomyces
S_downii
S_australis
S_sp__HMT_172
S_sp__HMT_180
S_marcescens
S_mitis
S_parasanguinis_clade_411
S_melaninogenica
G_Eikenella
S_corrodens
S_surfactantfaciens
S_atypica
S_multispecies_spp8_2
S_odontolytica
S_dentocariosa
S_concisus
S_gordonii
S_dispar
S_rogosae
G_Schaalia
G_Granulicatella
F_Carnobacteriaceae
F_Actinomycetaceae
O_Actinomycetales
S_adiacens
S_parvula
G_Serratia
F_Yersiniaceae
S_multispecies_spp24_2
S_perflava
S_sanguinis
S_nucleatum
P_Firmicutes
C_Bacilli
S_multispecies_spp2_2
G_Streptococcus
F_Streptococcaceae
O_Lactobacillales
G_Neisseria
P_Proteobacteria
C_Betaproteobacteria
F_Neisseriaceae
O_Neisseriales
C_Negativicutes
F_Veillonellaceae
O_Veillonellales
G_Veillonella
S_flavescens
G_Haemophilus
S_parainfluenzae
−5.20
−4.17
−3.14
−2.11
−1.08
−0.05
0.98
2.01
3.04
4.07
5.10
LDA SCORE (log 10)