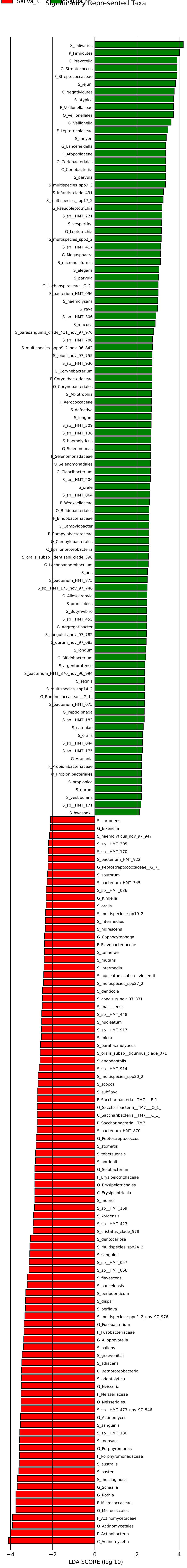
Saliva_K
Saliva_M
Significantly Represented Taxa
S_salivarius
P_Firmicutes
G_Prevotella
G_Streptococcus
F_Streptococcaceae
S_jejuni
C_Negativicutes
S_atypica
F_Veillonellaceae
O_Veillonellales
G_Veillonella
F_Leptotrichiaceae
S_meyeri
G_Lancefieldella
F_Atopobiaceae
O_Coriobacteriales
C_Coriobacteriia
S_parvula
S_multispecies_spp3_3
S_infantis_clade_431
S_multispecies_spp17_2
G_Pseudoleptotrichia
S_sp__HMT_221
S_vespertina
G_Leptotrichia
S_multispecies_spp2_2
S_sp__HMT_417
G_Megasphaera
S_micronuciformis
S_elegans
S_parvula
G_Lachnospiraceae__G_2_
S_bacterium_HMT_096
S_haemolysans
S_rava
S_sp__HMT_306
S_mucosa
S_parasanguinis_clade_411_nov_97_976
S_sp__HMT_780
S_multispecies_sppn9_2_nov_96_842
S_jejuni_nov_97_755
S_sp__HMT_930
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
G_Abiotrophia
F_Aerococcaceae
S_defectiva
S_longum
S_sp__HMT_309
S_sp__HMT_136
S_haemolyticus
G_Selenomonas
F_Selenomonadaceae
O_Selenomonadales
G_Cloacibacterium
S_sp__HMT_206
S_orale
S_sp__HMT_064
F_Weeksellaceae
O_Bifidobacteriales
F_Bifidobacteriaceae
G_Campylobacter
F_Campylobacteraceae
O_Campylobacterales
C_Epsilonproteobacteria
S_oralis_subsp__dentisani_clade_398
G_Lachnoanaerobaculum
S_oris
S_bacterium_HMT_875
S_sp__HMT_175_nov_97_746
G_Alloscardovia
S_omnicolens
G_Butyrivibrio
S_sp__HMT_455
G_Aggregatibacter
S_sanguinis_nov_97_782
S_durum_nov_97_083
S_longum
G_Bifidobacterium
S_argentoratense
S_bacterium_HMT_870_nov_96_994
S_segnis
S_multispecies_spp14_2
G_Ruminococcaceae__G_1_
S_bacterium_HMT_075
G_Peptidiphaga
S_sp__HMT_183
S_catoniae
S_oralis
S_sp__HMT_044
S_sp__HMT_175
G_Arachnia
F_Propionibacteriaceae
O_Propionibacteriales
S_propionica
S_durum
S_vestibularis
S_sp__HMT_171
S_hwasookii
S_corrodens
G_Eikenella
S_haemolyticus_nov_97_947
S_sp__HMT_305
S_sp__HMT_170
S_bacterium_HMT_922
G_Peptostreptococcaceae__G_7_
S_sputorum
S_bacterium_HMT_345
S_sp__HMT_036
G_Kingella
S_oralis
S_multispecies_spp19_2
S_intermedius
S_nigrescens
G_Capnocytophaga
F_Flavobacteriaceae
S_tannerae
S_mutans
S_intermedia
S_nucleatum_subsp__vincentii
S_multispecies_spp27_2
S_denticola
S_concisus_nov_97_831
S_massiliensis
S_sp__HMT_448
S_nucleatum
S_sp__HMT_917
S_micra
S_parahaemolyticus
S_oralis_subsp__tigurinus_clade_071
S_endodontalis
S_sp__HMT_914
S_multispecies_spp20_2
S_scopos
S_subflava
F_Saccharibacteria__TM7___F_1_
O_Saccharibacteria__TM7___O_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
S_bacterium_HMT_870
G_Peptostreptococcus
S_stomatis
S_tobetsuensis
S_gordonii
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_moorei
S_sp__HMT_169
S_koreensis
S_sp__HMT_423
S_cristatus_clade_578
S_dentocariosa
S_multispecies_spp24_2
S_sanguinis
S_sp__HMT_057
S_sp__HMT_066
S_flavescens
S_nanceiensis
S_periodonticum
S_dispar
S_perflava
S_multispecies_sppn1_2_nov_97_976
G_Fusobacterium
F_Fusobacteriaceae
G_Alloprevotella
S_pallens
S_graevenitzii
S_adiacens
C_Betaproteobacteria
S_odontolytica
G_Neisseria
F_Neisseriaceae
O_Neisseriales
S_sp__HMT_473_nov_97_546
G_Actinomyces
S_sanguinis
S_sp__HMT_180
S_rogosae
G_Porphyromonas
F_Porphyromonadaceae
S_australis
S_pasteri
S_mucilaginosa
G_Schaalia
G_Rothia
F_Micrococcaceae
O_Micrococcales
F_Actinomycetaceae
O_Actinomycetales
P_Actinobacteria
C_Actinomycetia
−4
−2
0
2
4
LDA SCORE (log 10)