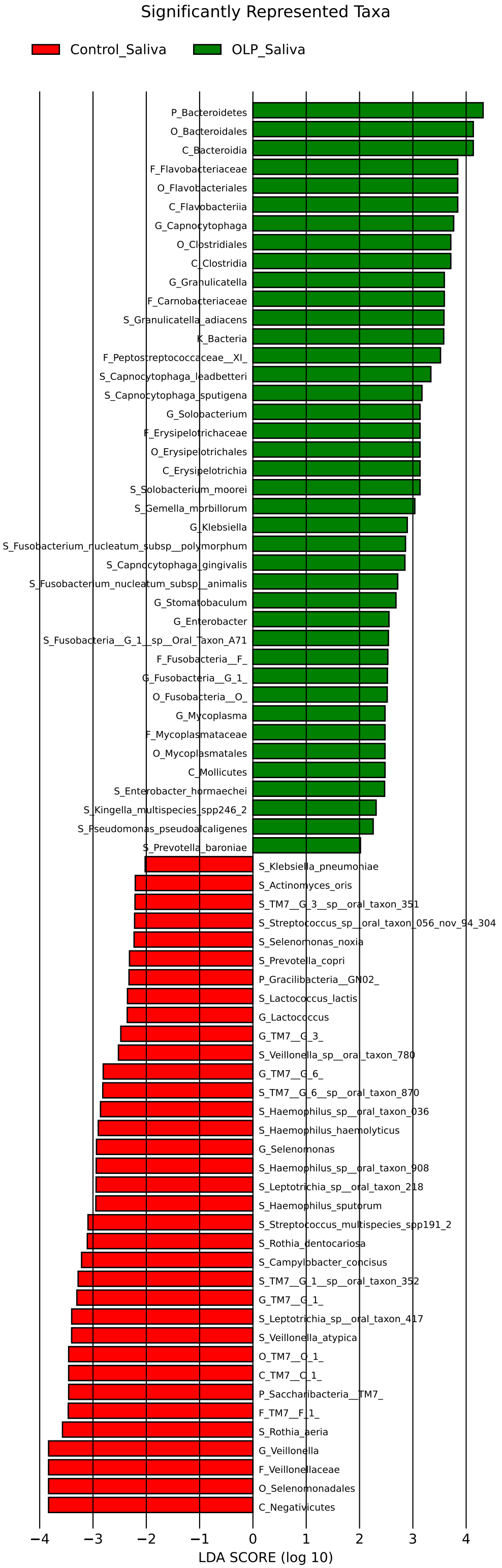
Significantly Represented Taxa
Control_Saliva
OLP_Saliva
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
F_Flavobacteriaceae
O_Flavobacteriales
C_Flavobacteriia
G_Capnocytophaga
O_Clostridiales
C_Clostridia
G_Granulicatella
F_Carnobacteriaceae
S_Granulicatella_adiacens
K_Bacteria
F_Peptostreptococcaceae__XI_
S_Capnocytophaga_leadbetteri
S_Capnocytophaga_sputigena
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_Solobacterium_moorei
S_Gemella_morbillorum
G_Klebsiella
S_Fusobacterium_nucleatum_subsp__polymorphum
S_Capnocytophaga_gingivalis
S_Fusobacterium_nucleatum_subsp__animalis
G_Stomatobaculum
G_Enterobacter
S_Fusobacteria__G_1__sp__Oral_Taxon_A71
F_Fusobacteria__F_
G_Fusobacteria__G_1_
O_Fusobacteria__O_
G_Mycoplasma
F_Mycoplasmataceae
O_Mycoplasmatales
C_Mollicutes
S_Enterobacter_hormaechei
S_Kingella_multispecies_spp246_2
S_Pseudomonas_pseudoalcaligenes
S_Prevotella_baroniae
S_Klebsiella_pneumoniae
S_Actinomyces_oris
S_TM7__G_3__sp__oral_taxon_351
S_Streptococcus_sp__oral_taxon_056_nov_94_304
S_Selenomonas_noxia
S_Prevotella_copri
P_Gracilibacteria__GN02_
S_Lactococcus_lactis
G_Lactococcus
G_TM7__G_3_
S_Veillonella_sp__oral_taxon_780
G_TM7__G_6_
S_TM7__G_6__sp__oral_taxon_870
S_Haemophilus_sp__oral_taxon_036
S_Haemophilus_haemolyticus
G_Selenomonas
S_Haemophilus_sp__oral_taxon_908
S_Leptotrichia_sp__oral_taxon_218
S_Haemophilus_sputorum
S_Streptococcus_multispecies_spp191_2
S_Rothia_dentocariosa
S_Campylobacter_concisus
S_TM7__G_1__sp__oral_taxon_352
G_TM7__G_1_
S_Leptotrichia_sp__oral_taxon_417
S_Veillonella_atypica
O_TM7__O_1_
C_TM7__C_1_
P_Saccharibacteria__TM7_
F_TM7__F_1_
S_Rothia_aeria
G_Veillonella
F_Veillonellaceae
O_Selenomonadales
C_Negativicutes
−4
−3
−2
−1
0
1
2
3
4
LDA SCORE (log 10)