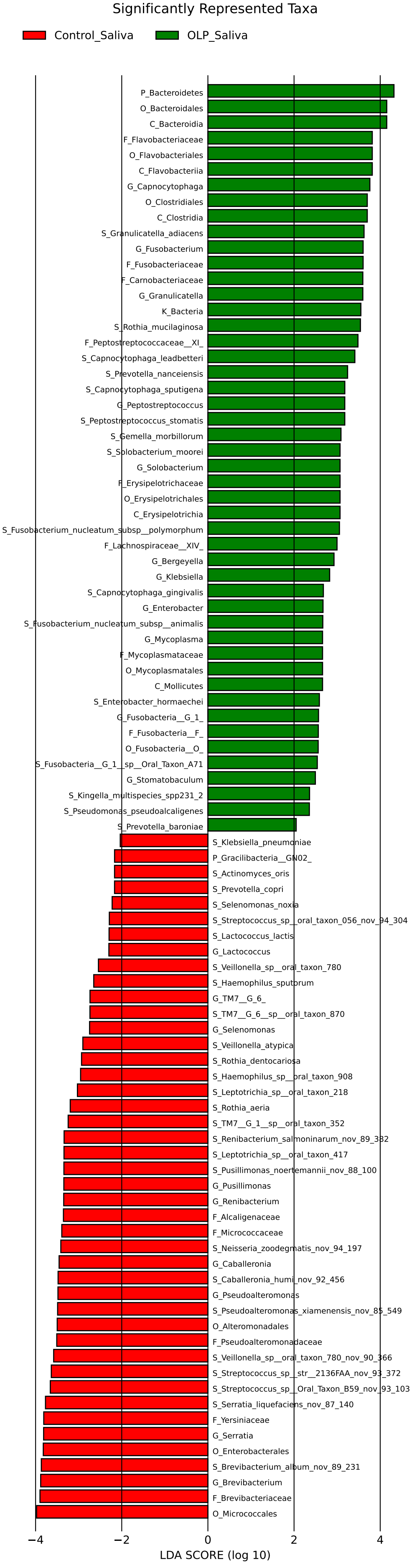
Significantly Represented Taxa
Control_Saliva
OLP_Saliva
P_Bacteroidetes
O_Bacteroidales
C_Bacteroidia
F_Flavobacteriaceae
O_Flavobacteriales
C_Flavobacteriia
G_Capnocytophaga
O_Clostridiales
C_Clostridia
S_Granulicatella_adiacens
G_Fusobacterium
F_Fusobacteriaceae
F_Carnobacteriaceae
G_Granulicatella
K_Bacteria
S_Rothia_mucilaginosa
F_Peptostreptococcaceae__XI_
S_Capnocytophaga_leadbetteri
S_Prevotella_nanceiensis
S_Capnocytophaga_sputigena
G_Peptostreptococcus
S_Peptostreptococcus_stomatis
S_Gemella_morbillorum
S_Solobacterium_moorei
G_Solobacterium
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
S_Fusobacterium_nucleatum_subsp__polymorphum
F_Lachnospiraceae__XIV_
G_Bergeyella
G_Klebsiella
S_Capnocytophaga_gingivalis
G_Enterobacter
S_Fusobacterium_nucleatum_subsp__animalis
G_Mycoplasma
F_Mycoplasmataceae
O_Mycoplasmatales
C_Mollicutes
S_Enterobacter_hormaechei
G_Fusobacteria__G_1_
F_Fusobacteria__F_
O_Fusobacteria__O_
S_Fusobacteria__G_1__sp__Oral_Taxon_A71
G_Stomatobaculum
S_Kingella_multispecies_spp231_2
S_Pseudomonas_pseudoalcaligenes
S_Prevotella_baroniae
S_Klebsiella_pneumoniae
P_Gracilibacteria__GN02_
S_Actinomyces_oris
S_Prevotella_copri
S_Selenomonas_noxia
S_Streptococcus_sp__oral_taxon_056_nov_94_304
S_Lactococcus_lactis
G_Lactococcus
S_Veillonella_sp__oral_taxon_780
S_Haemophilus_sputorum
G_TM7__G_6_
S_TM7__G_6__sp__oral_taxon_870
G_Selenomonas
S_Veillonella_atypica
S_Rothia_dentocariosa
S_Haemophilus_sp__oral_taxon_908
S_Leptotrichia_sp__oral_taxon_218
S_Rothia_aeria
S_TM7__G_1__sp__oral_taxon_352
S_Renibacterium_salmoninarum_nov_89_382
S_Leptotrichia_sp__oral_taxon_417
S_Pusillimonas_noertemannii_nov_88_100
G_Pusillimonas
G_Renibacterium
F_Alcaligenaceae
F_Micrococcaceae
S_Neisseria_zoodegmatis_nov_94_197
G_Caballeronia
S_Caballeronia_humi_nov_92_456
G_Pseudoalteromonas
S_Pseudoalteromonas_xiamenensis_nov_85_549
O_Alteromonadales
F_Pseudoalteromonadaceae
S_Veillonella_sp__oral_taxon_780_nov_90_366
S_Streptococcus_sp__str__2136FAA_nov_93_372
S_Streptococcus_sp__Oral_Taxon_B59_nov_93_103
S_Serratia_liquefaciens_nov_87_140
F_Yersiniaceae
G_Serratia
O_Enterobacterales
S_Brevibacterium_album_nov_89_231
G_Brevibacterium
F_Brevibacteriaceae
O_Micrococcales
−4
−2
0
2
4
LDA SCORE (log 10)