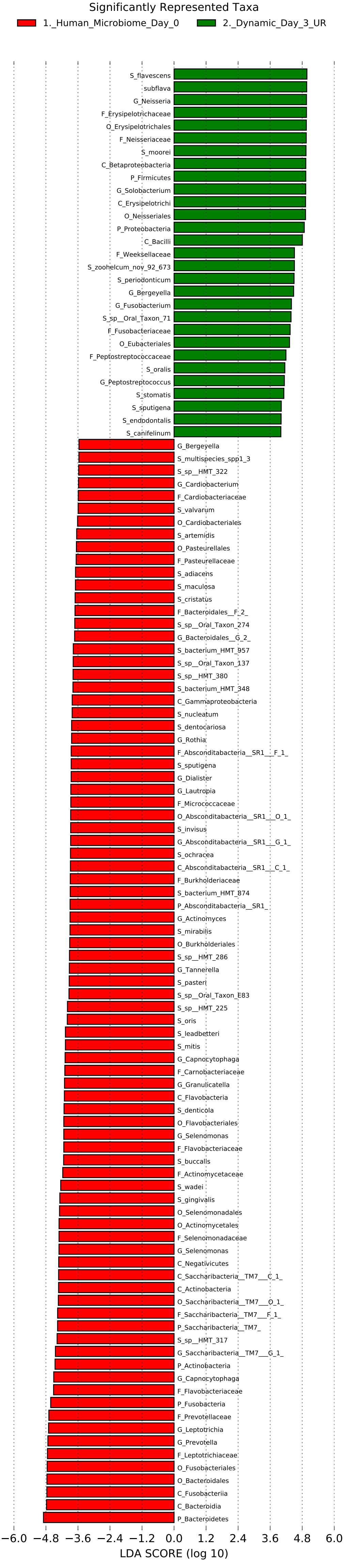
Significantly Represented Taxa
1._Human_Microbiome_Day_0
2._Dynamic_Day_3_UR
S_flavescens
subflava
G_Neisseria
F_Erysipelotrichaceae
O_Erysipelotrichales
F_Neisseriaceae
S_moorei
C_Betaproteobacteria
P_Firmicutes
G_Solobacterium
C_Erysipelotrichi
O_Neisseriales
P_Proteobacteria
C_Bacilli
F_Weeksellaceae
S_zoohelcum_nov_92_673
S_periodonticum
G_Bergeyella
G_Fusobacterium
S_sp__Oral_Taxon_71
F_Fusobacteriaceae
O_Eubacteriales
F_Peptostreptococcaceae
S_oralis
G_Peptostreptococcus
S_stomatis
S_sputigena
S_endodontalis
S_canifelinum
G_Bergeyella
S_multispecies_spp1_3
S_sp__HMT_322
G_Cardiobacterium
F_Cardiobacteriaceae
S_valvarum
O_Cardiobacteriales
S_artemidis
O_Pasteurellales
F_Pasteurellaceae
S_adiacens
S_maculosa
S_cristatus
F_Bacteroidales__F_2_
S_sp__Oral_Taxon_274
G_Bacteroidales__G_2_
S_bacterium_HMT_957
S_sp__Oral_Taxon_137
S_sp__HMT_380
S_bacterium_HMT_348
C_Gammaproteobacteria
S_nucleatum
S_dentocariosa
G_Rothia
F_Absconditabacteria__SR1__F_1_
S_sputigena
G_Dialister
G_Lautropia
F_Micrococcaceae
O_Absconditabacteria__SR1__O_1_
S_invisus
G_Absconditabacteria__SR1__G_1_
S_ochracea
C_Absconditabacteria__SR1__C_1_
F_Burkholderiaceae
S_bacterium_HMT_874
P_Absconditabacteria__SR1_
G_Actinomyces
S_mirabilis
O_Burkholderiales
S_sp__HMT_286
G_Tannerella
S_pasteri
S_sp__Oral_Taxon_E83
S_sp__HMT_225
S_oris
S_leadbetteri
S_mitis
G_Capnocytophaga
F_Carnobacteriaceae
G_Granulicatella
C_Flavobacteria
S_denticola
O_Flavobacteriales
G_Selenomonas
F_Flavobacteriaceae
S_buccalis
F_Actinomycetaceae
S_wadei
S_gingivalis
O_Selenomonadales
O_Actinomycetales
F_Selenomonadaceae
G_Selenomonas
C_Negativicutes
C_Saccharibacteria__TM7___C_1_
C_Actinobacteria
O_Saccharibacteria__TM7___O_1_
F_Saccharibacteria__TM7___F_1_
P_Saccharibacteria__TM7_
S_sp__HMT_317
G_Saccharibacteria__TM7___G_1_
P_Actinobacteria
G_Capnocytophaga
F_Flavobacteriaceae
P_Fusobacteria
F_Prevotellaceae
G_Leptotrichia
G_Prevotella
F_Leptotrichiaceae
O_Fusobacteriales
O_Bacteroidales
C_Fusobacteriia
C_Bacteroidia
P_Bacteroidetes
−6.0 −4.8 −3.6 −2.4 −1.2 0.0 1.2 2.4 3.6 4.8 6.0
LDA SCORE (log 10)