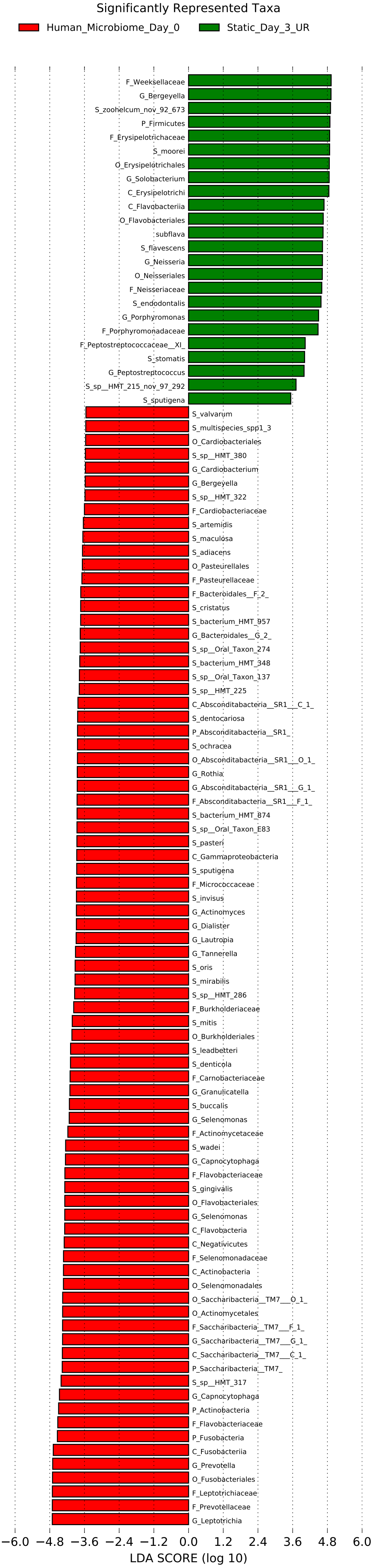
Significantly Represented Taxa
Human_Microbiome_Day_0
Static_Day_3_UR
F_Weeksellaceae
G_Bergeyella
S_zoohelcum_nov_92_673
P_Firmicutes
F_Erysipelotrichaceae
S_moorei
O_Erysipelotrichales
G_Solobacterium
C_Erysipelotrichi
C_Flavobacteriia
O_Flavobacteriales
subflava
S_flavescens
G_Neisseria
O_Neisseriales
F_Neisseriaceae
S_endodontalis
G_Porphyromonas
F_Porphyromonadaceae
F_Peptostreptococcaceae__XI_
S_stomatis
G_Peptostreptococcus
S_sp__HMT_215_nov_97_292
S_sputigena
S_valvarum
S_multispecies_spp1_3
O_Cardiobacteriales
S_sp__HMT_380
G_Cardiobacterium
G_Bergeyella
S_sp__HMT_322
F_Cardiobacteriaceae
S_artemidis
S_maculosa
S_adiacens
O_Pasteurellales
F_Pasteurellaceae
F_Bacteroidales__F_2_
S_cristatus
S_bacterium_HMT_957
G_Bacteroidales__G_2_
S_sp__Oral_Taxon_274
S_bacterium_HMT_348
S_sp__Oral_Taxon_137
S_sp__HMT_225
C_Absconditabacteria__SR1__C_1_
S_dentocariosa
P_Absconditabacteria__SR1_
S_ochracea
O_Absconditabacteria__SR1__O_1_
G_Rothia
G_Absconditabacteria__SR1__G_1_
F_Absconditabacteria__SR1__F_1_
S_bacterium_HMT_874
S_sp__Oral_Taxon_E83
S_pasteri
C_Gammaproteobacteria
S_sputigena
F_Micrococcaceae
S_invisus
G_Actinomyces
G_Dialister
G_Lautropia
G_Tannerella
S_oris
S_mirabilis
S_sp__HMT_286
F_Burkholderiaceae
S_mitis
O_Burkholderiales
S_leadbetteri
S_denticola
F_Carnobacteriaceae
G_Granulicatella
S_buccalis
G_Selenomonas
F_Actinomycetaceae
S_wadei
G_Capnocytophaga
F_Flavobacteriaceae
S_gingivalis
O_Flavobacteriales
G_Selenomonas
C_Flavobacteria
C_Negativicutes
F_Selenomonadaceae
C_Actinobacteria
O_Selenomonadales
O_Saccharibacteria__TM7___O_1_
O_Actinomycetales
F_Saccharibacteria__TM7___F_1_
G_Saccharibacteria__TM7___G_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
S_sp__HMT_317
G_Capnocytophaga
P_Actinobacteria
F_Flavobacteriaceae
P_Fusobacteria
C_Fusobacteriia
G_Prevotella
O_Fusobacteriales
F_Leptotrichiaceae
F_Prevotellaceae
G_Leptotrichia
−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)