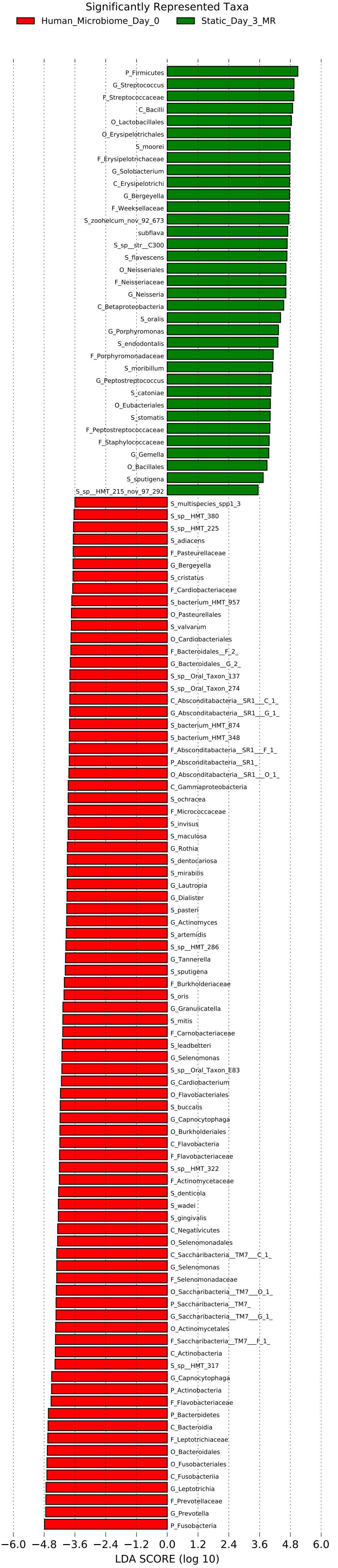
Significantly Represented Taxa
Human_Microbiome_Day_0
Static_Day_3_MR
P_Firmicutes
G_Streptococcus
F_Streptococcaceae
C_Bacilli
O_Lactobacillales
O_Erysipelotrichales
S_moorei
F_Erysipelotrichaceae
G_Solobacterium
C_Erysipelotrichi
G_Bergeyella
F_Weeksellaceae
S_zoohelcum_nov_92_673
subflava
S_sp__str__C300
S_flavescens
O_Neisseriales
F_Neisseriaceae
G_Neisseria
C_Betaproteobacteria
S_oralis
G_Porphyromonas
S_endodontalis
F_Porphyromonadaceae
S_moribillum
G_Peptostreptococcus
S_catoniae
O_Eubacteriales
S_stomatis
F_Peptostreptococcaceae
F_Staphylococcaceae
G_Gemella
O_Bacillales
S_sputigena
S_sp__HMT_215_nov_97_292
S_multispecies_spp1_3
S_sp__HMT_380
S_sp__HMT_225
S_adiacens
F_Pasteurellaceae
G_Bergeyella
S_cristatus
F_Cardiobacteriaceae
S_bacterium_HMT_957
O_Pasteurellales
S_valvarum
O_Cardiobacteriales
F_Bacteroidales__F_2_
G_Bacteroidales__G_2_
S_sp__Oral_Taxon_137
S_sp__Oral_Taxon_274
C_Absconditabacteria__SR1___C_1_
G_Absconditabacteria__SR1___G_1_
S_bacterium_HMT_874
S_bacterium_HMT_348
F_Absconditabacteria__SR1___F_1_
P_Absconditabacteria__SR1_
O_Absconditabacteria__SR1___O_1_
C_Gammaproteobacteria
S_ochracea
F_Micrococcaceae
S_invisus
S_maculosa
G_Rothia
S_dentocariosa
S_mirabilis
G_Lautropia
G_Dialister
S_pasteri
G_Actinomyces
S_artemidis
S_sp__HMT_286
G_Tannerella
S_sputigena
F_Burkholderiaceae
S_oris
G_Granulicatella
S_mitis
F_Carnobacteriaceae
S_leadbetteri
G_Selenomonas
S_sp__Oral_Taxon_E83
G_Cardiobacterium
O_Flavobacteriales
S_buccalis
G_Capnocytophaga
O_Burkholderiales
C_Flavobacteria
F_Flavobacteriaceae
S_sp__HMT_322
F_Actinomycetaceae
S_denticola
S_wadei
S_gingivalis
C_Negativicutes
O_Selenomonadales
C_Saccharibacteria__TM7___C_1_
G_Selenomonas
F_Selenomonadaceae
O_Saccharibacteria__TM7___O_1_
P_Saccharibacteria__TM7_
G_Saccharibacteria__TM7___G_1_
O_Actinomycetales
F_Saccharibacteria__TM7___F_1_
C_Actinobacteria
S_sp__HMT_317
G_Capnocytophaga
P_Actinobacteria
F_Flavobacteriaceae
P_Bacteroidetes
C_Bacteroidia
F_Leptotrichiaceae
O_Bacteroidales
O_Fusobacteriales
C_Fusobacteriia
G_Leptotrichia
F_Prevotellaceae
G_Prevotella
P_Fusobacteria
−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)