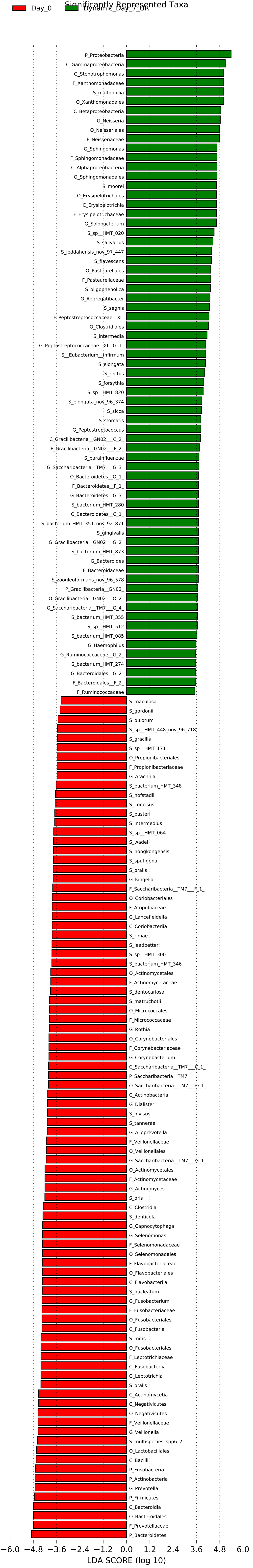
Significantly Represented Taxa
Day_0
Dynamic_Day_7_UR
P_Proteobacteria
C_Gammaproteobacteria
G_Stenotrophomonas
F_Xanthomonadaceae
S_maltophilia
O_Xanthomonadales
C_Betaproteobacteria
G_Neisseria
O_Neisseriales
F_Neisseriaceae
G_Sphingomonas
F_Sphingomonadaceae
C_Alphaproteobacteria
O_Sphingomonadales
S_moorei
O_Erysipelotrichales
C_Erysipelotrichia
F_Erysipelotrichaceae
G_Solobacterium
S_sp__HMT_020
S_salivarius
S_jeddahensis_nov_97_447
S_flavescens
O_Pasteurellales
F_Pasteurellaceae
S_oligophenolica
G_Aggregatibacter
S_segnis
F_Peptostreptococcaceae__XI_
O_Clostridiales
S_intermedia
G_Peptostreptococcaceae__XI__G_1_
S__Eubacterium__infirmum
S_elongata
S_rectus
S_forsythia
S_sp__HMT_820
S_elongata_nov_96_374
S_sicca
S_stomatis
G_Peptostreptococcus
C_Gracilibacteria__GN02___C_2_
F_Gracilibacteria__GN02___F_2_
S_parainfluenzae
G_Saccharibacteria__TM7___G_3_
O_Bacteroidetes__O_1_
F_Bacteroidetes__F_1_
G_Bacteroidetes__G_3_
S_bacterium_HMT_280
C_Bacteroidetes__C_1_
S_bacterium_HMT_351_nov_92_871
S_gingivalis
G_Gracilibacteria__GN02___G_2_
S_bacterium_HMT_873
G_Bacteroides
F_Bacteroidaceae
S_zoogleoformans_nov_96_578
P_Gracilibacteria__GN02_
O_Gracilibacteria__GN02___O_2_
G_Saccharibacteria__TM7___G_4_
S_bacterium_HMT_355
S_sp__HMT_512
S_bacterium_HMT_085
G_Haemophilus
G_Ruminococcaceae__G_2_
S_bacterium_HMT_274
G_Bacteroidales__G_2_
F_Bacteroidales__F_2_
F_Ruminococcaceae
S_maculosa
S_gordonii
S_oulorum
S_sp__HMT_448_nov_96_718
S_gracilis
S_sp__HMT_171
O_Propionibacteriales
F_Propionibacteriaceae
G_Arachnia
S_bacterium_HMT_348
S_hofstadii
S_concisus
S_pasteri
S_intermedius
S_sp__HMT_064
S_wadei
S_hongkongensis
S_sputigena
S_oralis
G_Kingella
F_Saccharibacteria__TM7___F_1_
O_Coriobacteriales
F_Atopobiaceae
G_Lancefieldella
C_Coriobacteriia
S_rimae
S_leadbetteri
S_sp__HMT_300
S_bacterium_HMT_346
O_Actinomycetales
F_Actinomycetaceae
S_dentocariosa
S_matruchotii
O_Micrococcales
F_Micrococcaceae
G_Rothia
O_Corynebacteriales
F_Corynebacteriaceae
G_Corynebacterium
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
O_Saccharibacteria__TM7___O_1_
C_Actinobacteria
G_Dialister
S_invisus
S_tannerae
G_Alloprevotella
F_Veillonellaceae
O_Veillonellales
G_Saccharibacteria__TM7___G_1_
O_Actinomycetales
F_Actinomycetaceae
G_Actinomyces
S_oris
C_Clostridia
S_denticola
G_Capnocytophaga
G_Selenomonas
F_Selenomonadaceae
O_Selenomonadales
F_Flavobacteriaceae
O_Flavobacteriales
C_Flavobacteriia
S_nucleatum
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
S_mitis
O_Fusobacteriales
F_Leptotrichiaceae
C_Fusobacteriia
G_Leptotrichia
S_oralis
C_Actinomycetia
C_Negativicutes
O_Negativicutes
F_Veillonellaceae
G_Veillonella
S_multispecies_spp6_2
O_Lactobacillales
C_Bacilli
P_Fusobacteria
P_Actinobacteria
G_Prevotella
P_Firmicutes
C_Bacteroidia
O_Bacteroidales
F_Prevotellaceae
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)