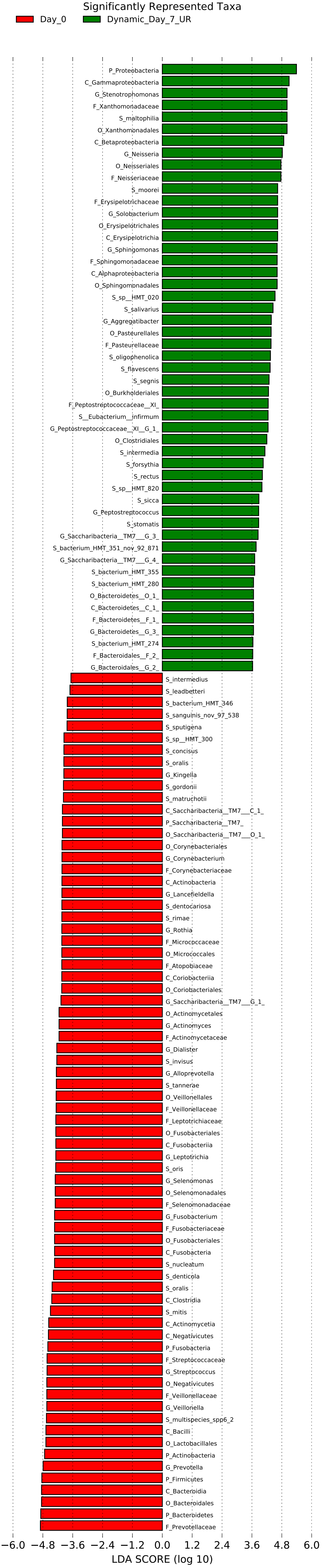
Significantly Represented Taxa
Day_0
Dynamic_Day_7_UR
P_Proteobacteria
C_Gammaproteobacteria
G_Stenotrophomonas
F_Xanthomonadaceae
S_maltophilia
O_Xanthomonadales
C_Betaproteobacteria
G_Neisseria
O_Neisseriales
F_Neisseriaceae
S_moorei
F_Erysipelotrichaceae
G_Solobacterium
O_Erysipelotrichales
C_Erysipelotrichia
G_Sphingomonas
F_Sphingomonadaceae
C_Alphaproteobacteria
O_Sphingomonadales
S_sp__HMT_020
S_salivarius
G_Aggregatibacter
O_Pasteurellales
F_Pasteurellaceae
S_oligophenolica
S_flavescens
S_segnis
O_Burkholderiales
F_Peptostreptococcaceae__XI_
S__Eubacterium__infirmum
G_Peptostreptococcaceae__XI__G_1_
O_Clostridiales
S_intermedia
S_forsythia
S_rectus
S_sp__HMT_820
S_sicca
G_Peptostreptococcus
S_stomatis
G_Saccharibacteria__TM7___G_3_
S_bacterium_HMT_351_nov_92_871
G_Saccharibacteria__TM7___G_4_
S_bacterium_HMT_355
S_bacterium_HMT_280
O_Bacteroidetes__O_1_
C_Bacteroidetes__C_1_
F_Bacteroidetes__F_1_
G_Bacteroidetes__G_3_
S_bacterium_HMT_274
F_Bacteroidales__F_2_
G_Bacteroidales__G_2_
S_intermedius
S_leadbetteri
S_bacterium_HMT_346
S_sanguinis_nov_97_538
S_sputigena
S_sp__HMT_300
S_concisus
S_oralis
G_Kingella
S_gordonii
S_matruchotii
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
O_Saccharibacteria__TM7___O_1_
O_Corynebacteriales
G_Corynebacterium
F_Corynebacteriaceae
C_Actinobacteria
G_Lancefieldella
S_dentocariosa
S_rimae
G_Rothia
F_Micrococcaceae
O_Micrococcales
F_Atopobiaceae
C_Coriobacteriia
O_Coriobacteriales
G_Saccharibacteria__TM7___G_1_
O_Actinomycetales
G_Actinomyces
F_Actinomycetaceae
G_Dialister
S_invisus
G_Alloprevotella
S_tannerae
O_Veillonellales
F_Veillonellaceae
F_Leptotrichiaceae
O_Fusobacteriales
C_Fusobacteriia
G_Leptotrichia
S_oris
G_Selenomonas
O_Selenomonadales
F_Selenomonadaceae
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
S_nucleatum
S_denticola
S_oralis
C_Clostridia
S_mitis
C_Actinomycetia
C_Negativicutes
P_Fusobacteria
F_Streptococcaceae
G_Streptococcus
O_Negativicutes
F_Veillonellaceae
G_Veillonella
S_multispecies_spp6_2
C_Bacilli
O_Lactobacillales
P_Actinobacteria
G_Prevotella
P_Firmicutes
C_Bacteroidia
O_Bacteroidales
P_Bacteroidetes
F_Prevotellaceae
−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)