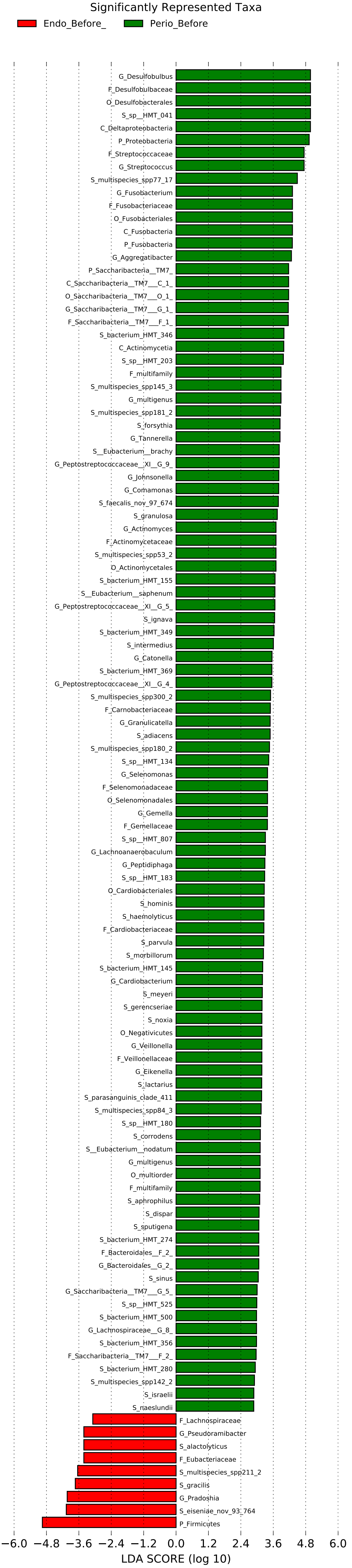
Significantly Represented Taxa
Endo_Before_
Perio_Before
G_Desulfobulbus
F_Desulfobulbaceae
O_Desulfobacterales
S_sp__HMT_041
C_Deltaproteobacteria
P_Proteobacteria
F_Streptococcaceae
G_Streptococcus
S_multispecies_spp77_17
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
P_Fusobacteria
G_Aggregatibacter
P_Saccharibacteria__TM7_
C_Saccharibacteria__TM7___C_1_
O_Saccharibacteria__TM7___O_1_
G_Saccharibacteria__TM7___G_1_
F_Saccharibacteria__TM7___F_1_
S_bacterium_HMT_346
C_Actinomycetia
S_sp__HMT_203
F_multifamily
S_multispecies_spp145_3
G_multigenus
S_multispecies_spp181_2
S_forsythia
G_Tannerella
S__Eubacterium__brachy
G_Peptostreptococcaceae__XI__G_9_
G_Johnsonella
G_Comamonas
S_faecalis_nov_97_674
S_granulosa
G_Actinomyces
F_Actinomycetaceae
S_multispecies_spp53_2
O_Actinomycetales
S_bacterium_HMT_155
S__Eubacterium__saphenum
G_Peptostreptococcaceae__XI__G_5_
S_ignava
S_bacterium_HMT_349
S_intermedius
G_Catonella
S_bacterium_HMT_369
G_Peptostreptococcaceae__XI__G_4_
S_multispecies_spp300_2
F_Carnobacteriaceae
G_Granulicatella
S_adiacens
S_multispecies_spp180_2
S_sp__HMT_134
G_Selenomonas
F_Selenomonadaceae
O_Selenomonadales
G_Gemella
F_Gemellaceae
S_sp__HMT_807
G_Lachnoanaerobaculum
G_Peptidiphaga
S_sp__HMT_183
O_Cardiobacteriales
S_hominis
S_haemolyticus
F_Cardiobacteriaceae
S_parvula
S_morbillorum
S_bacterium_HMT_145
G_Cardiobacterium
S_meyeri
S_gerencseriae
S_noxia
O_Negativicutes
G_Veillonella
F_Veillonellaceae
G_Eikenella
S_lactarius
S_parasanguinis_clade_411
S_multispecies_spp84_3
S_sp__HMT_180
S_corrodens
S__Eubacterium__nodatum
G_multigenus
O_multiorder
F_multifamily
S_aphrophilus
S_dispar
S_sputigena
S_bacterium_HMT_274
F_Bacteroidales__F_2_
G_Bacteroidales__G_2_
S_sinus
G_Saccharibacteria__TM7___G_5_
S_sp__HMT_525
S_bacterium_HMT_500
G_Lachnospiraceae__G_8_
S_bacterium_HMT_356
F_Saccharibacteria__TM7___F_2_
S_bacterium_HMT_280
S_multispecies_spp142_2
S_israelii
S_naeslundii
F_Lachnospiraceae
G_Pseudoramibacter
S_alactolyticus
F_Eubacteriaceae
S_multispecies_spp211_2
S_gracilis
G_Pradoshia
S_eiseniae_nov_93_764
P_Firmicutes
−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)