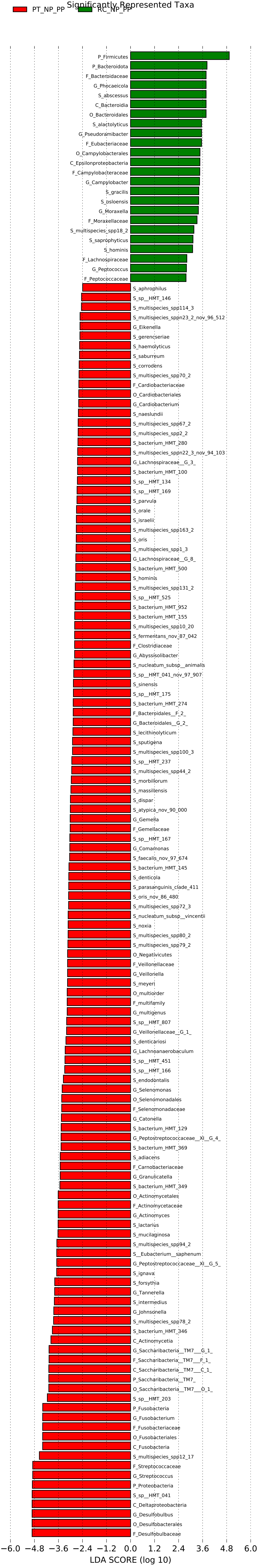
Significantly Represented Taxa
PT_NP_PP
RC_NP_PP
P_Firmicutes
P_Bacteroidota
F_Bacteroidaceae
G_Phocaeicola
S_abscessus
C_Bacteroidia
O_Bacteroidales
S_alactolyticus
G_Pseudoramibacter
F_Eubacteriaceae
O_Campylobacterales
C_Epsilonproteobacteria
F_Campylobacteraceae
G_Campylobacter
S_gracilis
S_osloensis
G_Moraxella
F_Moraxellaceae
S_multispecies_spp18_2
S_saprophyticus
S_hominis
F_Lachnospiraceae
G_Peptococcus
F_Peptococcaceae
S_aphrophilus
S_sp__HMT_146
S_multispecies_spp114_3
S_multispecies_sppn23_2_nov_96_512
G_Eikenella
S_gerencseriae
S_haemolyticus
S_saburreum
S_corrodens
S_multispecies_spp70_2
F_Cardiobacteriaceae
O_Cardiobacteriales
G_Cardiobacterium
S_naeslundii
S_multispecies_spp67_2
S_multispecies_spp2_2
S_bacterium_HMT_280
S_multispecies_sppn22_3_nov_94_103
G_Lachnospiraceae__G_3_
S_bacterium_HMT_100
S_sp__HMT_134
S_sp__HMT_169
S_parvula
S_orale
S_israelii
S_multispecies_spp163_2
S_oris
S_multispecies_spp1_3
G_Lachnospiraceae__G_8_
S_bacterium_HMT_500
S_hominis
S_multispecies_spp131_2
S_sp__HMT_525
S_bacterium_HMT_952
S_bacterium_HMT_155
S_multispecies_spp10_20
S_fermentans_nov_87_042
F_Clostridiaceae
G_Abyssisolibacter
S_nucleatum_subsp__animalis
S_sp__HMT_041_nov_97_907
S_sinensis
S_sp__HMT_175
S_bacterium_HMT_274
F_Bacteroidales__F_2_
G_Bacteroidales__G_2_
S_lecithinolyticum
S_sputigena
S_multispecies_spp100_3
S_sp__HMT_237
S_multispecies_spp44_2
S_morbillorum
S_massiliensis
S_dispar
S_atypica_nov_90_000
G_Gemella
F_Gemellaceae
S_sp__HMT_167
G_Comamonas
S_faecalis_nov_97_674
S_bacterium_HMT_145
S_denticola
S_parasanguinis_clade_411
S_oris_nov_86_480
S_multispecies_spp72_3
S_nucleatum_subsp__vincentii
S_noxia
S_multispecies_spp80_2
S_multispecies_spp79_2
O_Negativicutes
F_Veillonellaceae
G_Veillonella
S_meyeri
O_multiorder
F_multifamily
G_multigenus
S_sp__HMT_807
G_Veillonellaceae__G_1_
S_denticariosi
G_Lachnoanaerobaculum
S_sp__HMT_451
S_sp__HMT_166
S_endodontalis
G_Selenomonas
O_Selenomonadales
F_Selenomonadaceae
G_Catonella
S_bacterium_HMT_129
G_Peptostreptococcaceae__XI__G_4_
S_bacterium_HMT_369
S_adiacens
F_Carnobacteriaceae
G_Granulicatella
S_bacterium_HMT_349
O_Actinomycetales
F_Actinomycetaceae
G_Actinomyces
S_lactarius
S_mucilaginosa
S_multispecies_spp94_2
S__Eubacterium__saphenum
G_Peptostreptococcaceae__XI__G_5_
S_ignava
S_forsythia
G_Tannerella
S_intermedius
G_Johnsonella
S_multispecies_spp78_2
S_bacterium_HMT_346
C_Actinomycetia
G_Saccharibacteria__TM7___G_1_
F_Saccharibacteria__TM7___F_1_
C_Saccharibacteria__TM7___C_1_
P_Saccharibacteria__TM7_
O_Saccharibacteria__TM7___O_1_
S_sp__HMT_203
P_Fusobacteria
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
S_multispecies_spp12_17
F_Streptococcaceae
G_Streptococcus
P_Proteobacteria
S_sp__HMT_041
C_Deltaproteobacteria
G_Desulfobulbus
O_Desulfobacterales
F_Desulfobulbaceae
−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)