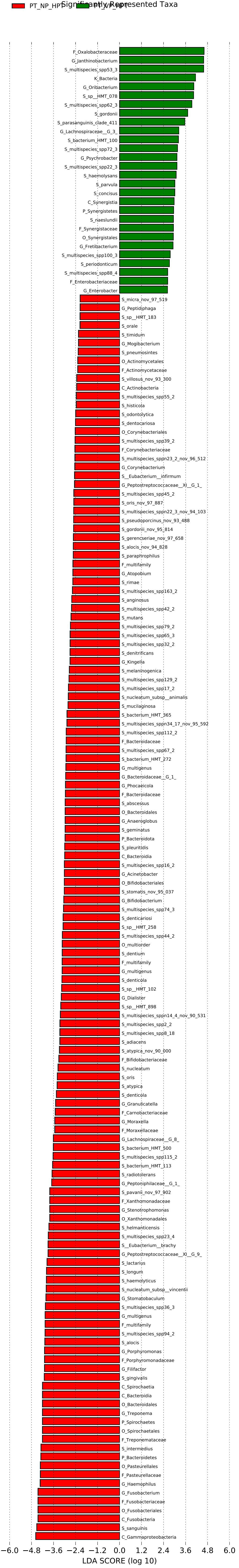
PT_NP_HPT
PT_VP_HPT
Significantly Represented Taxa
F_Oxalobacteraceae
G_Janthinobacterium
S_multispecies_spp53_3
K_Bacteria
G_Oribacterium
S_sp__HMT_078
S_multispecies_spp62_3
S_gordonii
S_parasanguinis_clade_411
G_Lachnospiraceae__G_3_
S_bacterium_HMT_100
S_multispecies_spp72_3
G_Psychrobacter
S_multispecies_spp22_3
S_haemolysans
S_parvula
S_concisus
C_Synergistia
P_Synergistetes
S_naeslundii
F_Synergistaceae
O_Synergistales
G_Fretibacterium
S_multispecies_spp100_3
S_periodonticum
S_multispecies_spp88_4
F_Enterobacteriaceae
G_Enterobacter
S_micra_nov_97_519
G_Peptidiphaga
S_sp__HMT_183
S_orale
S_timidum
G_Mogibacterium
S_pneumosintes
O_Actinomycetales
F_Actinomycetaceae
S_villosus_nov_93_300
C_Actinobacteria
S_multispecies_spp55_2
S_histicola
S_odontolytica
S_dentocariosa
O_Corynebacteriales
S_multispecies_spp39_2
F_Corynebacteriaceae
S_multispecies_sppn23_2_nov_96_512
G_Corynebacterium
S__Eubacterium__infirmum
G_Peptostreptococcaceae__XI__G_1_
S_multispecies_spp45_2
S_oris_nov_97_887
S_multispecies_sppn22_3_nov_94_103
S_pseudoporcinus_nov_93_488
S_gordonii_nov_95_814
S_gerencseriae_nov_97_658
S_alocis_nov_94_828
S_paraphrophilus
F_multifamily
G_Atopobium
S_rimae
S_multispecies_spp163_2
S_anginosus
S_multispecies_spp42_2
S_mutans
S_multispecies_spp79_2
S_multispecies_spp65_3
S_multispecies_spp32_2
S_denitrificans
G_Kingella
S_melaninogenica
S_multispecies_spp129_2
S_multispecies_spp17_2
S_nucleatum_subsp__animalis
S_mucilaginosa
S_bacterium_HMT_365
S_multispecies_sppn34_17_nov_95_592
S_multispecies_spp112_2
F_Bacteroidaceae
S_multispecies_spp67_2
S_bacterium_HMT_272
G_multigenus
G_Bacteroidaceae__G_1_
G_Phocaeicola
F_Bacteroidaceae
S_abscessus
O_Bacteroidales
G_Anaeroglobus
S_geminatus
P_Bacteroidota
S_pleuritidis
C_Bacteroidia
S_multispecies_spp16_2
G_Acinetobacter
O_Bifidobacteriales
S_stomatis_nov_95_037
G_Bifidobacterium
S_multispecies_spp74_3
S_denticariosi
S_sp__HMT_258
S_multispecies_spp44_2
O_multiorder
S_dentium
F_multifamily
G_multigenus
S_denticola
S_sp__HMT_102
G_Dialister
S_sp__HMT_898
S_multispecies_sppn14_4_nov_90_531
S_multispecies_spp2_2
S_multispecies_spp8_18
S_adiacens
S_atypica_nov_90_000
F_Bifidobacteriaceae
S_nucleatum
S_oris
S_atypica
S_denticola
G_Granulicatella
F_Carnobacteriaceae
G_Moraxella
F_Moraxellaceae
G_Lachnospiraceae__G_8_
S_bacterium_HMT_500
S_multispecies_spp115_2
S_bacterium_HMT_113
S_radiotolerans
G_Peptoniphilaceae__G_1_
S_pavanii_nov_97_902
F_Xanthomonadaceae
G_Stenotrophomonas
O_Xanthomonadales
S_helmanticensis
S_multispecies_spp23_4
S__Eubacterium__brachy
G_Peptostreptococcaceae__XI__G_9_
S_lactarius
S_longum
S_haemolyticus
S_nucleatum_subsp__vincentii
G_Stomatobaculum
S_multispecies_spp36_3
G_multigenus
F_multifamily
S_multispecies_spp94_2
S_alocis
G_Porphyromonas
F_Porphyromonadaceae
G_Filifactor
S_gingivalis
C_Spirochaetia
C_Bacteroidia
O_Bacteroidales
G_Treponema
P_Spirochaetes
O_Spirochaetales
F_Treponemataceae
S_intermedius
P_Bacteroidetes
O_Pasteurellales
F_Pasteurellaceae
G_Haemophilus
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
S_sanguinis
C_Gammaproteobacteria
−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)