RC_NP_HPT RC_NP_PD

Significantly Represented Taxa

P_Firmicutes
C_Bacilli
O_Lactobacillales
S_faecalis
F_Enterococcaceae
G_Enterococcus
F_Peptoniphilaceae
G_Parvimonas
S_micra
F_Lachnospiraceae__XIV_
F_Aerococcaceae
G_Aerococcus
S_multispecies_spp162_2
C_Coriobacteriia
O_Coriobacteriales
G_Stomatobaculum
S_sp__HMT_373
F_Coriobacteriaceae
G_Atopobium
G_Oribacterium
S_sp__HMT_078_nov_97_537
F_Bacillaceae
S_eiseniae_nov_93_764
G_Pradoshia
S_timidum
G_Paracoccus
F_Rhodobacteraceae
G_Mogibacterium
S_sp__HMT_199
O_Rhodobacterales
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_odontolytica
C_Actinobacteria
S_stomatis
G_Peptostreptococcus
S_alocis
G_Filifactor
F_Desulfobulbaceae
O_Desulfobacterales
G_Desulfobulbus
S_multispecies_spp78_2
S_sp__HMT_041
F_Atopobiaceae
G_Olsenella
S_multispecies_spp36_3
S_multispecies_spp145_5
S_multispecies_spp57_4
G_Peptostreptococcaceae__XI__G_9_
S__Eubacterium__brachy
S_multispecies_spp66_2
S_uli
S_multispecies_spp63_2
S_sediminicola
G_Prevotella
G_Peptostreptococcaceae__XI__G_5_
S__Eubacterium__saphenum
F_Prevotellaceae
S_concisus
O_Caulobacterales
F_Caulobacteraceae
G_Brevundimonas
S_gordonii
S_invisus
F_Weeksellaceae
S_multispecies_spp129_2
S_multispecies_spp132_2
C_Flavobacteriia
O_Flavobacteriales
G_Chryseobacterium
S_multispecies_spp93_3
S_fermentans_nov_87_042
F_Clostridiaceae
S_multispecies_spp37_5
G_Abyssisolibacter
S_sp__HMT_212
G_Neobacillus
G_Janthinobacterium
S_multispecies_spp53_3
S_sp__HMT_041_nov_97_907
F_Enterobacteriaceae
G_Enterobacter
S_multispecies_spp103_5
G_Peptostreptococcaceae__XI__G_2_
S_desulfuricans
S_mori
S_multispecies_spp138_2
S_bacterium_HMT_091
S_mucilaginosa
G_Schnuerera
S_ultunensis_nov_92_840
F_Tissierellaceae
S_multispecies_spp23_4
S_multispecies_spp18_2
F_Actinomycetaceae
G_Actinomyces
S_multispecies_spp94_2
S_multispecies_spp149_8
G_Neisseria
S_sp__HMT_373_nov_94_581
S_sputigena
S_saprophyticus
F_Oxalobacteraceae
S_multispecies_spp13_3
S_sp__HMT_169
S_atriviolacea
G_Massilia
S_aeruginosa
S_anginosus
F_Rhizobiaceae
S_rosettiformans
G_Rhizobium
S_longum
S_histicola
S_parvula
S_melaninogenica
S_nucleatum_subsp__animalis
S_atypica
G_Lachnoanaerobaculum
S_multispecies_spp35_3
S_dispar
S_gordonii_nov_95_814
S_denticariosi
S_denticola
S_adiacens
F_Carnobacteriaceae
G_Granulicatella
S_sp__HMT_417
S_multispecies_spp52_2
O_Negativicutes
F_Veillonellaceae
G_Veillonella
F_Bacteroidales__F_2_
S_bacterium_HMT_274
G_Bacteroidales__G_2_
S_multispecies_spp72_3
S_multispecies_spp108_2
S_warneri
C_Tissierellia
S_atypica_nov_90_000
O_Tissierellales
S_orale
S_multispecies_spp70_2
O_Hyphomicrobiales
G_Dialister
S_bacterium_HMT_369
G_Peptostreptococcaceae__XI__G_4_
S_multispecies_spp1_3
G_Desulfovibrio
C_Deltaproteobacteria
S_gracilis
C_Epsilonproteobacteria
O_Campylobacterales
G_Campylobacter
F_Campylobacteraceae
F_Desulfovibrionaceae
S_multispecies_spp55_2
O_Desulfovibrionales
F_Corynebacteriaceae
O_Corynebacteriales
G_Corynebacterium
S_multispecies_spp2_2
S_helmanticensis
S_sinensis
G_Anaeroglobus
S_geminatus
G_Stenotrophomonas
S_pavanii_nov_97_902
F_Xanthomonadaceae
O_Xanthomonadales
G_Fusobacterium
F_Fusobacteriaceae
O_Fusobacteriales
C_Fusobacteria
P_Fusobacteria
S_multispecies_spp115_2
F_Moraxellaceae
G_Moraxella
O_Pseudomonadales
S_multispecies_spp81_6
C_Gammaproteobacteria
P_Proteobacteria

−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)