Significantly Represented Taxa

ADULT_HIV-Neg_NON-SHEDDER

CHILD_HIV-Neg_NON-SHEDDER

G_Streptococcus
F_Streptococcaceae
S_sanguinis
O_Burkholderiales
G_Lautropia
F_Burkholderiaceae
S_mirabilis
S_oralis
S_sp__HMT_498
G_Actinomyces
S_multispecies_spp124_6
G_Corynebacterium
F_Corynebacteriaceae
O_Corynebacteriales
S_matruchotii
S_shahii
S_multispecies_spp82_2
G_Peptidiphaga
G_Kingella
S_aeria
S_hongkongensis
S_gingivicola
S_oralis
S_multispecies_spp125_4
S_mutans
S_veroralis
S_sp__HMT_847
S_oulorum
S_sp__HMT_012
S_gracilis
S_sp__HMT_183
S_sputigena
G_Cardiobacterium
F_Cardiobacteriaceae
O_Cardiobacteriales
S_intermedius
S_trevisanii
S_hominis
S_valvarum
S_multiformis
S_sp__HMT_078
S_granulosa
S_multispecies_spp66_2
G_Kocuria
S_bacterium_HMT_113
G_Peptoniphilaceae__G_1_
S_abscessus
F_Bacteroidaceae
G_Phocaeicola
S_bacterium_HMT_091
G_Peptostreptococcaceae__G_2_
S_sp__HMT_041
C_Deltaproteobacteria
F_Desulfobulbaceae
O_Desulfobacterales
G_Desulfobulbus
S_multispecies_spp22_3
S_multispecies_spp23_2
S_sp__HMT_455
S_sp__HMT_102
S_sp__HMT_217
G_Butyrivibrio
S_forsythia
S_sp__HMT_097
S_baroniae
S_multispecies_spp35_6
S_multispecies_spp163_2
S_sp__HMT_443
S_bacterium_HMT_375
S__Eubacterium__sulci
G_Filifactor
S_alocis
S_multispecies_spp132_2
F_Erysipelotrichaceae
O_Erysipelotrichales
C_Erysipelotrichia
G_Solobacterium
S_moorei
G_Peptostreptococcus
S_stomatis
S_gingivalis
S_bacterium_HMT_345
S_multispecies_spp24_3
S_multispecies_spp13_2
G_Absconditabacteria__SR1___G_1_
F_Absconditabacteria__SR1___F_1_
O_Absconditabacteria__SR1___O_1_
C_Absconditabacteria__SR1___C_1_
P_Absconditabacteria__SR1_
F_Peptostreptococcaceae
S_multispecies_spp130_2
S_intermedia
S_melaninogenica
S_periodonticum
S_multispecies_spp133_3
S_multispecies_spp87_4
O_Bacteroidales
C_Bacteroidia
P_Bacteroidetes

−5.27 −4.21 −3.15 −2.09 −1.03 0.03 1.09 2.15 3.21 4.27 5.33

LDA SCORE (log 10)