Significantly Represented Taxa

negative positive

P_Firmicutes
C_Clostridia
O_Clostridiales
G_Rothia
F_Micrococcaceae
O_Micrococcales
S_mucilaginosa
F_Peptostreptococcaceae__XI_
F_Lachnospiraceae__XIV_
S_multispecies_spp34_7
S_adiacens
S_multispecies_spp57_3
S_multispecies_spp5_4
G_Oribacterium
F_Actinomycetaceae
O_Actinomycetales
G_Schaalia
S_sinus
S_stomatis
G_Peptostreptococcus
G_Stomatobaculum
C_Actinobacteria
F_Peptoniphilaceae
S_micra
G_Parvimonas
S_sp__HMT_172
S_sp__HMT_097
G_Mogibacterium
G_Clostridioides
S_difficile
S_graevenitzii
S_multispecies_spp101_2
G_multigenus
F_multifamily
O_Coriobacteriales
C_Coriobacteriia
G_Romboutsia
S_timonensis
S_multispecies_spp100_4
O_Erysipelotrichales
C_Erysipelotrichia
F_Erysipelotrichaceae
S_moorei
G_Solobacterium
F_Peptostreptococcaceae
S__Eubacterium__brachy
G_Peptostreptococcaceae__XI__G_9_
S_multispecies_spp116_3
G_Peptostreptococcaceae__XI__G_1_
S_asaccharolyticum
S__Eubacterium__sulci
F_Synergistaceae
O_Synergistales
G_Fretibacterium
C_Synergistia
P_Synergistetes
S_lingnae__Not_Validly_Published_
S_sp__HMT_360
G_Peptostreptococcaceae__XI__G_2_
S_bacterium_HMT_091
S_morbi
S_sp__HMT_894
F_Comamonadaceae
G_Ottowia
G_Pseudoramibacter
F_Eubacteriaceae
S_alactolyticus
S_satelles
G_Shuttleworthia
S_sp__HMT_455
G_Peptidiphaga
S_hongkongensis
G_Pseudostreptobacillus
S_bacterium_HMT_505
S_bacterium_HMT_507
S_sp__HMT_912
S_fusca
S_sp__HMT_907
S_multispecies_spp1_2
S_sp__HMT_246
G_Bacteroidetes__G_5_
S_sp__HMT_012
S_mutans
S_amylovorum
S_multispecies_spp129_2
S_anginosus
G_Kingella
S_actinomycetemcomitans
S_multispecies_spp122_2
G_Kocuria
O_Enterobacterales
S_multispecies_spp123_7
G_multigenus
F_Enterobacteriaceae
S_multispecies_spp61_2
S_oralis
S_multispecies_spp115_2
S_sp__HMT_930

−4 −3 −2 −1 0 1 2 3 4 5

LDA SCORE (log 10)