Significantly Represented Taxa

ADULT-NON-SHEDDER ADULT-SHEDDER CHILD-NON-SH

K_Bacteria
F_Veillonellaceae
O_Veillonellales
S_oralis
S_sp__HMT_930
S_invisus
S_hongkongensis
G_Kocuria
S_multispecies_spp42_2
F_Coriobacteriaceae
G_Atopobium
K_Eukaryota
P_Streptophyta
C_Pinopsida
O_Pinales
S_sylvestris
G_Pinus
F_Pinaceae
G_Lautropia
F_Burkholderiaceae
O_Burkholderiales
S_mirabilis
S_multispecies_spp59_2
S_multispecies_spp113_6
S_shahii
S_actinomycetemcomitans
G_multigenus
S_multispecies_spp159_7
O_Enterobacterales
S_sp__HMT_012
S_sp__HMT_847
F_Enterobacteriaceae
F_Poaceae
S_multispecies_spp112_4
O_Poales
G_Triticum
C_Magnoliopsida
S_aestivum
G_Kingella
S_multispecies_spp62_2
S_aurantiaca
S_veroralis
S_oralis
S_trevisanii
O_Clostridiales
F_Lachnospiraceae__XIV_
G_Oribacterium
G_Schaalia
S_sp__HMT_172
G_Mogibacterium
S_multispecies_spp82_4
S_bacterium_HMT_091
S_multispecies_spp145_3
reptococcaceae__XI__G_2_
S_morbi
S_bacterium_HMT_155
G_Veillonellaceae__G_1_
G_Butyrivibrio
S_parvum
S_asaccharolyticum
S_anginosus
S_bacterium_HMT_274
G_Bacteroidales__G_2_
F_Bacteroidales__F_2_
S_gracilis
F_Bacteroidaceae
S_multispecies_spp165_2
G_multigenus
S_sp__HMT_199
G_Catonella
S_sp__HMT_455
F_Synergistaceae
P_Synergistetes
G_Fretibacterium
O_Synergistales
C_Synergistia
G_Lancefieldella
S_rimae
C_Clostridia
S_multispecies_spp162_4
G_Neisseria
S_multispecies_spp161_3
S_melaninogenica
eptostreptococcaceae__XI_
S_multispecies_spp91_3
S_multispecies_spp133_2
G_Romboutsia
S_timonensis
O_Coriobacteriales
C_Coriobacteriia
F_multifamily
S_multispecies_spp21_2
G_multigenus
S_stomatis
G_Peptostreptococcus
nditabacteria__SR1___F_1_
nditabacteria__SR1___C_1_
_Absconditabacteria__SR1_
nditabacteria__SR1___O_1_
nditabacteria__SR1___G_1_
F_Peptostreptococcaceae
S_gingivalis
S_sp__HMT_206
S_sp__HMT_215
S_moorei
G_Solobacterium
O_Erysipelotrichales
C_Erysipelotrichia
F_Erysipelotrichaceae
S_sp__HMT_097
S__Eubacterium__sulci
G_Megasphaera
S_micronuciformis
reptococcaceae__XI__G_1_
S_multispecies_spp32_4
F_multifamily
G_multigenus
S_multispecies_spp13_7
S_bacterium_HMT_875
S_multispecies_spp16_2
S_baroniae
e__Not_Validly_Published_
S_sp__HMT_914
S_shahii

0 1 2 3 4 5

LDA SCORE (log 10)