0
20
40
60
Value
Group
Saliva Healthy Control
Feces Healthy Control
Prevotella melaninogenica
Gemella haemolysans
Abiotrophia defectiva
Bergeyella sp._HMT_322
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Streptococcus sanguinis_sp._HMT_074
Neisseria subflava
Oribacterium sinus
Veillonella rogosae_sp._str._NVG100cf
Prevotella pallens
Prevotella salivae
Fusobacterium periodonticum
Capnocytophaga leadbetteri
Absconditabacteria_(SR1)_[G−1] bacterium_HMT_345
Schaalia sp._HMT_172
Granulicatella elegans
Rothia aeria
Neisseria flavescens|subflava_perflava
Lachnospira pectinoschiza_nov_96.372%
Bacteroides caccae
Bacteroides_Phocaeicola dorei_sp._Oral_Taxon_D25_sp._str._219_
Oscillibacter ruminantium_nov_92.601%
Oscillibacter ruminantium_nov_94.144%
Eubacterium coprostanoligenes_nov_95.711%
Lacnoclostridium sp._str._L2_50
Ruminococcus callidus_nov_94.369%
Dorea longicatena
Coprococcus comes
Eubacterium ventriosum
Barnesiella intestinihominis
Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
Blautia obeum
Eubacterium coprostanoligenes_nov_92.777%
Lacrimispora xylanolytica_nov_97.285%
Catenibacterium mitsuokai
Actinomyces sp._HMT_169_sp._str._ChDCB197
Ruminococcaceae_[G−1] bacterium_HMT_075
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Solobacterium moorei
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Mobiluncus multispecies_sppn34_2_nov_97.817%
Schaalia sp._HMT_180
Neisseria flavescens_flavescens|subflava
Capnocytophaga gingivalis
Actinomyces graevenitzii
Atopobium_Lancefieldella parvula_parvulum
Veillonella dispar_parvula
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
Gemella sanguinis
Haemophilus parainfluenzae
Monoglobus pectinilyticus_nov_90.112%
Faecalibacterium prausnitzii_nov_97.964%
Ruminococcus bromii
Lacnoclostridium sp._SS3/4
Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
Roseburia intestinalis
Collinsella aerofaciens
Duncaniella freteri_nov_87.069%
Alistipes putredinis
Kineothrix alysoides_nov_96.825%
Fusicatenibacter saccharivorans_nov_97.279%
Agathobaculum butyriciproducens
Oscillibacter ruminantium_nov_95.701%
Dialister succinatiphilus_nov_96.567%
Eubacterium ramulus
Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
Parabacteroides distasonis
Blautia obeum_wexlerae
Bifidobacterium longum
Eubacterium hallii
Bacteroides koreensis_kribbi
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Lactobacillus rogosae
Blautia luti
Eubacterium rectale
Subdoligranulum variabile
Roseburia faecalis_faecis
Phocaeicola vulgatus
Bacteroides sp._str._4136_uniformis
Fusicatenibacter saccharivorans
Lachnospira eligens_nov_94.785%
Lachnoclostridium pacaense_nov_96.145%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Lachnospira eligens
Ruminococcus multispecies_sppn1_2_nov_95.023%
Blautia faecis
Roseburia multispecies_sppn106_3_nov_96.833%
Bacteroides_Phocaeicola plebeius
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Peptostreptococcus stomatis
Porphyromonas pasteri
Granulicatella adiacens_paradiacens
Rothia dentocariosa
Rothia mucilaginosa
Faecalibacterium prausnitzii
Prevotella copri
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_i
Species
F6488.S06
F6488.S07
F6488.S11
F6488.S54
F6488.S56
F6488.S52
F6488.S60
F6488.S53
F6488.S55
F6488.S58
F6488.S57
F6488.S59
F6488.S61
F6488.S16
F6488.S17
F6488.S03
F6488.S08
F6488.S05
F6488.S10
F6488.S15
F6488.S02
F6488.S04
F6488.S01
F6488.S13
F6488.S14
F6488.S12
F6488.S09
Samples