0
20
40
60
80
100
Value
Group
Saliva Healthy Control
Feces Healthy Control
Roseburia intestinalis
Schaalia sp._HMT_180
Haemophilus parainfluenzae
Rothia aeria
Rothia dentocariosa
Abiotrophia defectiva
Fusobacterium periodonticum
Schaalia sp._HMT_172
Alistipes onderdonkii
Peptostreptococcus stomatis
Prevotella salivae
Bacteroides caccae
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Blautia obeum
Bergeyella sp._HMT_322
Fusicatenibacter saccharivorans
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Agathobaculum butyriciproducens
Oribacterium sinus
Barnesiella intestinihominis
Prevotella melaninogenica
Gemella sanguinis
Absconditabacteria_(SR1)_[G−1] bacterium_HMT_345
Parabacteroides distasonis
Actinomyces graevenitzii
Alistipes putredinis
Rothia mucilaginosa
Lactobacillus rogosae
Eubacterium rectale
Capnocytophaga gingivalis
Prevotella copri
Gemella haemolysans
Blautia luti
Stomatobaculum sp._HMT_097
Lachnospira eligens
Lacnoclostridium sp._SS3/4
Coprococcus comes
Prevotella pallens
Ruminococcus bromii
Catenibacterium mitsuokai
Capnocytophaga leadbetteri
Granulicatella elegans
Eubacterium ventriosum
Lacnoclostridium sp._str._L2_50
Capnocytophaga sputigena
Phocaeicola vulgatus
Porphyromonas pasteri
Eubacterium hallii
Faecalibacterium prausnitzii
Neisseria subflava
Ruminococcaceae_[G−1] bacterium_HMT_075
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Dorea longicatena
Subdoligranulum variabile
Collinsella aerofaciens
Solobacterium moorei
Blautia faecis
Bifidobacterium longum
Oscillibacter ruminantium_nov_94.144%
Ruminococcus callidus_nov_94.369%
Kineothrix alysoides_nov_96.825%
Lachnospira pectinoschiza_nov_96.372%
Lachnospira eligens_nov_94.785%
Eubacterium coprostanoligenes_nov_92.777%
Eubacterium coprostanoligenes_nov_95.711%
Dialister succinatiphilus_nov_96.567%
Duncaniella freteri_nov_87.069%
Oscillibacter ruminantium_nov_95.701%
Fusicatenibacter saccharivorans_nov_97.279%
Lacrimispora xylanolytica_nov_97.285%
Monoglobus pectinilyticus_nov_90.112%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Lachnoclostridium pacaense_nov_96.145%
Oscillibacter ruminantium_nov_92.601%
Bacteroides sp._str._4136_uniformis
Anaerostipes_Lacnoclostridium_unclassified_Lachnospiraceae hadr
Neisseria flavescens_flavescens|subflava
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_
Neisseria flavescens|subflava_perflava
Streptococcus sanguinis_sp._HMT_074
Atopobium_Lancefieldella parvula_parvulum
Roseburia faecalis_faecis
Granulicatella adiacens_paradiacens
Blautia_unclassified_Lachnospiraceae sp._str._cont1.79_torques
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari ..
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Blautia obeum_wexlerae
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Parasutterella_Sutterella excrementihominis_sp._str._cont1.66
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Veillonella rogosae_sp._str._NVG100cf
Bacteroides koreensis_kribbi
Veillonella dispar_parvula
Bacteroides_Phocaeicola plebeius
Actinomyces sp._HMT_169_sp._str._ChDCB197
Ruminococcus multispecies_sppn1_2_nov_95.023%
Roseburia multispecies_sppn106_3_nov_96.833%
Mobiluncus multispecies_sppn34_2_nov_97.817%
Species
F6488.S01
F6488.S02
F6488.S03
F6488.S04
F6488.S05
F6488.S06
F6488.S07
F6488.S08
F6488.S09
F6488.S10
F6488.S11
F6488.S12
F6488.S13
F6488.S14
F6488.S15
F6488.S16
F6488.S17
F6488.S52
F6488.S53
F6488.S54
F6488.S55
F6488.S56
F6488.S57
F6488.S58
F6488.S59
F6488.S60
F6488.S61
Samples