0
20
40
60
80
Value
Group
Saliva Periodontitis
Feces Periodontitis
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Haemophilus parainfluenzae
Bacteroidales_[G−2] sp._Oral_Taxon_274
Rothia dentocariosa
Abiotrophia defectiva
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Schaalia sp._HMT_172
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Peptostreptococcus stomatis
Tannerella forsythia
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Fusicatenibacter saccharivorans
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Mycoplasma faucium
Methanobrevibacter smithii
Gemella sanguinis
Porphyromonas endodontalis
Rothia mucilaginosa
Porphyromonas sp._HMT_285
Lactobacillus rogosae
Eubacterium rectale
Romboutsia timonensis
Coprococcus eutactus
Prevotella copri
Gemella haemolysans
Lachnospira eligens
Lacnoclostridium sp._SS3/4
Coprococcus comes
Ruminococcus bromii
Roseburia inulinivorans
Catenibacterium mitsuokai
Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
Coprococcus catus
Granulicatella elegans
Prevotella stercorea
Akkermansia muciniphila
Phocaeicola vulgatus
Porphyromonas pasteri
Faecalibacterium prausnitzii
Streptococcus anginosus
Ruminococcaceae_[G−1] bacterium_HMT_075
Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
Parvimonas micra
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Dorea longicatena
Subdoligranulum variabile
Collinsella aerofaciens
Solobacterium moorei
Blautia faecis
Gracilibacteria_(GN02)_[G−2] bacterium_HMT_873
Bifidobacterium longum
Oscillibacter ruminantium_nov_94.144%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Prevotella stercorea_nov_97.403%
Ruminococcus callidus_nov_94.369%
Eubacterium coprostanoligenes_nov_97.059%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Alloprevotella rava_nov_92.857%
Lachnospira eligens_nov_94.785%
Butyrivibrio sp._HMT_090_nov_92.795%
Prevotellamassilia timonensis_nov_97.397%
Eubacterium coprostanoligenes_nov_92.777%
Succinivibrio dextrinosolvens_nov_97.079%
Eubacterium coprostanoligenes_nov_95.711%
Prevotella oralis_nov_92.641%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.422%
Bergeyella sp._HMT_907_nov_97.397%
Dialister succinatiphilus_nov_96.360%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346_nov_97.968%
Lacrimispora xylanolytica_nov_97.285%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Lachnoclostridium pacaense_nov_96.145%
Sporobacter termitidis_nov_92.584%
Ruminoclostridium methylpentosum_nov_89.462%
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Agathobaculum desmolans_nov_97.738%
Oscillibacter ruminantium_nov_92.601%
Holdemanella biformis_nov_97.854%
Lactobacillus_Ligilactobacillus ruminis
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
Actinomyces_Schaalia odontolytica_odontolyticus
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari ..
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Bifidobacterium adolescentis_faecale
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Gemella morbillorum_moribillum
Blautia obeum_wexlerae
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Veillonella dispar_parvula
Veillonella parvula_tobetsuensis
Ruminococcus multispecies_sppn1_2_nov_95.023%
Christensenella multispecies_sppn2_2_nov_91.216%
Clostridium multispecies_sppn27_2_nov_97.959%
Species
F6488.S18
F6488.S19
F6488.S20
F6488.S21
F6488.S22
F6488.S23
F6488.S24
F6488.S25
F6488.S26
F6488.S27
F6488.S28
F6488.S29
F6488.S30
F6488.S31
F6488.S32
F6488.S33
F6488.S34
F6488.S35
F6488.S62
F6488.S63
F6488.S64
F6488.S65
F6488.S66
F6488.S67
F6488.S68
F6488.S69
F6488.S70
F6488.S71
Samples