0
20
40
60
80
Value
Group
Saliva Periodontitis
Feces Periodontitis
Coprococcus eutactus
Ruminococcus multispecies_sppn1_2_nov_95.023%
Lacrimispora xylanolytica_nov_97.285%
Succinivibrio dextrinosolvens_nov_97.079%
Alloprevotella rava_nov_92.857%
Eubacterium coprostanoligenes_nov_95.711%
Ruminoclostridium methylpentosum_nov_89.462%
Roseburia faecalis_faecis
Ruminococcus callidus_nov_94.369%
Butyrivibrio sp._HMT_090_nov_92.795%
Lactobacillus rogosae
Agathobaculum desmolans_nov_97.738%
Blautia obeum_wexlerae
Lachnospira eligens_nov_94.785%
Blautia faecis
Roseburia inulinivorans
Eubacterium coprostanoligenes_nov_92.777%
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Fusicatenibacter saccharivorans
Lachnoclostridium pacaense_nov_96.145%
Bifidobacterium longum
Methanobrevibacter smithii
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_91.422%
Oscillibacter ruminantium_nov_94.144%
Oscillibacter ruminantium_nov_92.601%
Christensenella multispecies_sppn2_2_nov_91.216%
Sporobacter termitidis_nov_92.584%
Clostridium multispecies_sppn27_2_nov_97.959%
Prevotella stercorea_nov_97.403%
Romboutsia timonensis
Prevotella oralis_nov_92.641%
Mycoplasma faucium
Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
Parvimonas micra
Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
Tannerella forsythia
Gracilibacteria_(GN02)_[G−2] bacterium_HMT_873
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346_nov_97.968%
Bacteroidales_[G−2] sp._Oral_Taxon_274
Actinomyces_Schaalia odontolytica_odontolyticus
Abiotrophia defectiva
Haemophilus parainfluenzae
Veillonella dispar_parvula
Gemella sanguinis
Rothia dentocariosa
Streptococcus sanguinis_sp._HMT_074
Streptococcus anginosus
Phocaeicola vulgatus
Eubacterium rectale
Lactobacillus_Ligilactobacillus ruminis
Holdemanella biformis_nov_97.854%
Dialister succinatiphilus_nov_96.360%
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Subdoligranulum variabile
Coprococcus comes
Dorea longicatena
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_93.946%
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Ruminococcus bromii
Ruminococcaceae_[G−2] bacterium_HMT_085_nov_92.809%
Coprococcus catus
Eubacterium coprostanoligenes_nov_97.059%
Gemella morbillorum_moribillum
Granulicatella elegans
Bergeyella sp._HMT_907_nov_97.397%
Prevotella stercorea
Prevotellamassilia timonensis_nov_97.397%
Lachnospira eligens
Catenibacterium mitsuokai
Lacnoclostridium sp._SS3/4
Akkermansia muciniphila
Ruminococcaceae_[G−1] bacterium_HMT_075
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Porphyromonas endodontalis
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Porphyromonas pasteri
Peptostreptococcus stomatis
Neisseria flavescens|subflava_perflava
Porphyromonas sp._HMT_285
Veillonella parvula_tobetsuensis
Neisseria flavescens_flavescens|subflava
Solobacterium moorei
Atopobium_Lancefieldella parvula_parvulum
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibulari ..
Schaalia sp._HMT_172
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Gemella haemolysans
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Rothia mucilaginosa
Granulicatella adiacens_paradiacens
Prevotella copri
Faecalibacterium prausnitzii
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_
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