



Group	
Saliva Periodontitis	
Feces Periodontitis	
Species	
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	
Lacnocostridium sp._SS3/4	
Coprococcus comes	
Ruminococcus bromii	
Roseburia inulinivorans	
Catenibacterium mitsuokai	
Peptostreptococcaceae_[XI][G-4] bacterium_HMT_369	
Coprococcus catus	
Granulicatella elegans	
Prevotella stercora	
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 stercora_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_i	
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_vestibularis	
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110	
Bifidobacterium adolescentis_faecale	
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei	
Gemella morbillorum_morbillum	
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%	

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Samples