



Source	
Saliva	
Feces	
Species	
Roseburia intestinalis	
Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346	
Schaalia sp._HMT_180	
Haemophilus parainfluenzae	
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	
Bergeyella sp._HMT_322	
Fusicatenibacter saccharivorans	
Peptostreptococcaceae_[XII][G-1] [Eubacterium]_sulci	
Mycoplasma faucium	
Agathobaculum butyriciproducens	
Oribacterium sinus	
Prevotella melaninogenica	
Ruminococcaceae_[G-2] bacterium_HMT_085	
Methanobrevibacter smithii	
Gemella sanguinis	
Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345	
Ruminococcus callidus	
Actinomyces graevenitzi	
Porphyromonas endodontalis	
Rothia mucilaginosa	
Lactobacillus rogosae	
Eubacterium rectale	
Coprococcus eutactus	
Capnocytophaga gingivalis	
Prevotella copri	
Gemella haemolysans	
Blautia luti	
Lachnospira eligens	
Lacnocolostridium sp._SS3/4	
Coprococcus comes	
Prevotella pallens	
Ruminococcus bromii	
Catenibacterium mitsuokai	
Granulicatella elegans	
Prevotella stercora	
Akkermansia muciniphila	
Phocaeicola vulgatus	
Porphyromonas pasteri	
Faecalibacterium prausnitzii	
Neisseria subflava	
Streptococcus anginosus	
Ruminococcaceae_[G-1] bacterium_HMT_075	
Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy	
Parvimonas micra	
Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352	
Actinomyces odontolyticus	
Dorea longicatena	
Subdoligranulum variabile	
Collinsella aerofaciens	
Solobacterium moorei	
Blautia faecis	
Bifidobacterium longum	
Oscillibacter ruminantium_nov_94.144%	
Ruminococcus callidus_nov_94.369%	
Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_89.038%	
Lachnospira eligens_nov_94.785%	
Sporobacter termitidis_nov_90.562%	
Eubacterium coprostanoligenes_nov_92.777%	
Succinivibrio dextrinosolvens_nov_97.079%	
Eubacterium coprostanoligenes_nov_95.711%	
Duncaniella freteri_nov_87.069%	
Dialister succinatiphilus_nov_96.360%	
Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%	
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%	
Sporobacter termitidis_nov_92.584%	
Oscillibacter ruminantium_nov_92.601%	
Bacteroides sp._str_4136_uniformis	
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_padiaciens	
Lacnocolostridium sp._SM4/1_sp_str_M62/1	
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_moribillum	
Blautia obeum_wexlerae	
Actinomyces_Schaalia odontolyticus_sp_HMT_180	
Selenomonas sp._HMT_136_sp_Oral_Taxon_149	
Veillonella dispar_parvula	
Veillonella parvula_tobetsuensis	
Actinomyces sp._HMT_169_sp_str_ChDCB197	
Streptococcus parasanguinis_parasanguinis_l_parasanguinis_clade	
Ruminococcus multispecies_sppn1_2_nov_95.023%	
Roseburia multispecies_sppn106_3_nov_96.833%	

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Samples