



- 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
Lacnospiridium 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_
Neisseria flavescens[subflava_perflava
Streptococcus sanguinis_sp._HMT_074
Atopobium_Lancefieldella parvula_parvulum
Roseburia faecalis_faecis
Granulicatella adiacens_padiaciens
Lacnospiridium 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
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
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%

Species

Samples