



Group

Saliva Healthy Control

Saliva Parkinson + Periodont

- Roseburia intestinalis
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Schaalia sp._HMT_180
- Haemophilus parainfluenzae
- Bacteroidales_[G-2] sp._Oral_Taxon_274
- Rothia aeria
- Rothia dentocariosa
- Abiotrophia defectiva
- Bacteroides fragilis
- Fusobacterium periodonticum
- Butyrivibrio crossotus
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Schaalia sp._HMT_172
- Neisseria elongata
- Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- Peptostreptococcus stomatis
- Tannerella forsythia
- Holdemanella biformis
- Prevotella salivae
- Bacteroides caccae
- Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351
- Blautia obeum
- Bergeyella sp._HMT_322
- Veillonellaceae_[G-1] bacterium_HMT_155
- Prevotella histicola
- Fusicatenibacter saccharivorans
- Peptostreptococcaceae_[XII][G-1] [Eubacterium]_sulci
- Mycoplasma faucium
- Agathobaculum butyriciproducens
- Oribacterium sinus
- Barnesiella intestinihominis
- Megasphaera elsdenii
- Prevotella melaninogenica
- Ruminococcaceae_[G-2] bacterium_HMT_085
- Odoribacter splanchnicus
- Methanobrevibacter smithii
- Erysipelotrichaceae_[G-1] bacterium_HMT_905
- Gemella sanguinis
- Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
- Campylobacter concisus
- Parabacteroides distasonis
- Ruminococcus callidus
- Parabacteroides merdae
- Actinomyces graevenitzi
- Alistipes putredinis
- Porphyromonas endodontalis
- Rothia mucilaginosa
- Porphyromonas sp._HMT_285
- Mediterraneibacter faecis
- Lactobacillus rogosae
- Eubacterium rectale
- Romboutsia timonensis
- Capnocytophaga gingivalis
- Gemella haemolysans
- Corynebacterium durum
- Stomatobaculum sp._HMT_097
- Prevotella pallens
- Capnocytophaga leadbetteri
- Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- Granulicatella elegans
- TM7_[G] sp._Oral_Taxon_A56
- Corynebacterium argentoratense
- Veillonella atypica
- Porphyromonas pasteri
- Selenomonas noxia
- 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
- Solobacterium moorei
- Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873
- Bifidobacterium longum
- Bergeyella sp._HMT_907_nov_97.397%
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346_nov_97.968%
- Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.072%
- 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
- 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
- Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
- Gemella morbillorum_morbillum
- Neisseria flava_macacae_mucosa_sicca_sp._TM101
- Actinomyces_Schaalia odontolyticus_sp._HMT_180
- Selenomonas sp._HMT_136_sp._Oral_Taxon_149
- Veillonella rogosae_sp._str._NVG100cf
- Veillonella dispar_parvula
- Veillonella parvula_tobetsuensis
- Oribacterium parvum_sp._str._ACB1
- Mogibacterium diversum_neglectum_vescum
- Actinomyces sp._HMT_169_sp._str._ChDCB197
- Streptococcus parasanguinis_parasanguinis_l_parasanguinis_clade
- Mobiluncus multispecies_sppn34_2_nov_97.817%

Species

Samples