



S076
S077
S078
S079
S080
S081
S082
S083
S084
S085
S086
S087
S088
S089
S090
S046
S047
S048
S049
S050
S051
S052
S053
S054
S055
S056
S057
S058
S059
S060

Description2
■ PT_VP_PP
■ PT_NP_PP

- Veillonellaceae_[G-1] bacterium_HMT_145
- Rothia mucilaginosa
- Micrococcus cohnii
- Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum
- Campylobacter gracilis
- Selenomonas noxia
- Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy
- Streptococcus sanguinis
- Treponema socranskii
- Corynebacterium durum
- Stomatobaculum sp._HMT_373
- Oribacterium sp._HMT_078
- Lachnospiraceae_[G-8] bacterium_HMT_500
- Enterococcus faecalis
- Olsenella profusa
- Olsenella sp._HMT_807
- Stenotrophobacter terrae
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
- Peptostreptococcaceae_[XII][G-6] [Eubacterium]_nodatum
- Parvimonas sp._HMT_110
- Peptostreptococcaceae_[XII][G-6] [Eubacterium]_minutum
- Porphyromonas gingivalis
- Peptoniphilaceae_[G-1] bacterium_HMT_113
- Haemophilus parainfluenzae
- Johnsonella ignava
- Fretibacterium sp._HMT_360
- Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- Veillonella dispar
- Mogibacterium timidum
- Lautropia mirabilis
- Schaalia odontolytica
- Gemella morbillorum
- Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
- Streptococcus sinensis
- Streptococcus anginosus
- Actinomyces sp._HMT_525
- Lachnoanaerobaculum sp._HMT_089
- Fretibacterium sp._HMT_359
- Treponema denticola
- Bacteroidales_[G-2] bacterium_HMT_274
- Desulfobulbus sp._HMT_041
- Streptococcus intermedius
- Fusobacterium sp._HMT_203
- Granulicatella adiacens
- Actinomyces sp._HMT_448
- Fretibacterium fastidiosum
- Pseudoramibacter alactolyticus
- Fusobacterium nucleatum_subsp._vincentii
- Anaerolineae_[G-1] bacterium_HMT_439
- Tannerella forsythia
- Porphyromonas endodontalis
- Peptidiphaga sp._HMT_183
- Limosilactobacillus mucosae
- Rothia dentocariosa
- Schaalia meyeri
- Peptostreptococcus stomatis
- Streptococcus lactarius
- Schaalia cardiffensis
- Fusobacterium nucleatum
- Filifactor alocis
- Aggregatibacter paraphrophilus
- Streptococcus parasanguinis_clade_411
- Johnsonella sp._HMT_166
- Paracoccus speluncae
- Parvimonas micra
- Selenomonas sputigena
- Peptostreptococcaceae_[XII][G-4] bacterium_HMT_103
- Streptococcus gordonii
- Oribacterium sp._HMT_078_nov_97.537%
- Desulfogranum japonicum_nov_86.848%
- Pseudacidobacterium ailaai_nov_91.605%
- Abyssohalobacter fermentans_nov_87.042%
- Desulfobulbus sp._HMT_041_nov_97.907%
- Pseudolabrys taiwanensis_nov_96.782%
- Parvimonas micra_nov_95.426%
- Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy_nov_96.05
- Peptostreptococcaceae_[XII][G-6] [Eubacterium]_nodatum_nov_95.5
- Parvimonas micra_nov_93.971%
- Schnuerera ultunensis_nov_92.840%
- Pseudacidobacterium ailaai_nov_94.321%
- Streptococcus cristatus_cristatus_clade_578_downii_gwangjuense
- Fusobacterium canifelinum_nucleatum_nucleatum_subsp._polymor
- Streptococcus cristatus_downii_gwangjuense_infantis_infantis_cla
- Acinetobacter_Prolinoborus fasciculus_lwoffii
- Streptococcus salivarius_vestibularis
- Paracoccus aestuarii_beibuensis_hibisci_marinus_pueri
- Brevundimonas bullata_halotolerans
- Aerococcus urinaeequi_viridans
- Neisseria flava_macacae_mucosa_sicca
- Parvimonas_Peptostreptococcus Candidatus massiliensis_sp._HMT
- Staphylococcus capitis_caprae_epidermidis
- Peptoanaerobacter_Peptostreptococcaceae_[XII][G-7] [Eubacterium]
- Peptoanaerobacter_Peptostreptococcaceae_[XII][G-7] [Eubacterium]
- Pseudomonas alcaliphila_chengduensis_oleovorans_toyotomiensis
- Streptococcus cristatus_cristatus_clade_578
- Fusobacterium naviforme_nucleatum_sp._HMT_204
- Streptococcus anginosus_constellatus
- Streptococcus cristatus_cristatus_clade_578_downii_gwangjuense
- Brevundimonas albigilva_nasdae_vesicularis
- Fusobacterium nucleatum_nucleatum_subsp._vincentii

Species

Samples