

Species

- SP295 Leptotrichia sp._HMT_212
- SP300 Streptococcus oralis
- SP313 Sphingomonas hankookensis
- SP316 Bacteroidales_[G-2] bacterium_HMT_274
- SP318 Kocuria palustris
- SP33 Blautia luti
- SP335 Actinomyces viscosus
- SP336 Veillonellaceae_[G-1] bacterium_HMT_135
- SP34 Fusobacterium periodonticum
- SP341 Streptococcus periodonticum
- SP350 Streptococcus downii
- SP352 Terrisporobacter mayombei
- SP36 Micrococcus flavus
- SP362 Fretribacterium fastidiosum
- SP364 Porphyromonas sp._HMT_277
- SP368 Streptococcus sinensis
- SP371 Lachnospiraceae_[G-8] bacterium_HMT_500
- SP372 Fibrobacter intestinalis
- SP38 Fusobacterium hwasookii
- SP380 Fretribacterium sp._HMT_362
- SP388 Selenomonas sp._HMT_442
- SP4 Alloprevotella sp._HMT_913
- SP40 Staphylococcus saprophyticus
- SP400 Tannerella sp._HMT_808
- SP406 Weeksellaceae_[G-1] sp._HMT_900
- SP416 Streptococcus oralis_subsp_dentisani_clade_058
- SP418 Desulfomicrobium orale
- SP423 Haemophilus parahaemolyticus
- SP427 Alloprevotella sp._HMT_914
- SP44 Cardiobacterium hominis
- SP441 Bacteroides heparinolyticus
- SP450 Capnocytophaga sp._HMT_878
- SP452 Alloprevotella sp._HMT_473
- SP453 Mogibacterium timidum
- SP456 Actinomyces johnsonii
- SP459 Peptostreptococcus stomatis
- SP464 Actinomyces sp._HMT_896
- SP482 Tannerella sp._HMT_916
- SP483 Porphyromonas gingivalis
- SP487 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955

- SPN176 Capnocytophaga sp._HMT_332_nov_94.480%
- SPN1763 Selenomonas sp._HMT_478_nov_96.640%
- SPN1765 Mogibacterium neglectum_nov_89.855%
- SPN1774 Treponema sp._HMT_264_nov_83.534%
- SPN1795 Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_95.417%
- SPN1808 Aggregatibacter actinomycetemcomitans_nov_95.910%
- SPN1809 Eubacteriales_[G-3] bacterium_MOT-163_nov_86.680%
- SPN1821 Selenomonas sp._HMT_892_nov_97.446%
- SPN1831 Gemella haemolysans_nov_97.984%
- SPN1841 Catonella morbi_nov_97.699%
- SPN1842 Ruminococcus champanellensis_nov_90.086%
- SPN1851 Prevotella sp._HMT_304_nov_95.492%
- SPN1862 Catenibacterium mitsuokai_nov_97.934%
- SPN187 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_92.453%
- SPN1873 Anaerocolumna cellulositytica_nov_87.657%
- SPN1882 Prevotella brevis_nov_89.184%
- SPN1884 Peptidiphaga gingivicola_nov_95.297%
- SPN1896 Comamonas faecalis_nov_91.803%
- SPN1908 Haemophilus haemolyticus_nov_94.908%
- SPN1919 Bacteroidetes_[G-7] bacterium_HMT_911_nov_89.528%
- SPN1931 Acinetobacter tandooi_nov_97.308%
- SPN1941 Actinomyces gerenceriae_nov_94.061%
- SPN195 Actinobacillus lignieresii_nov_97.342%
- SPN1953 Porphyromonas pasteri_nov_96.939%
- SPN1954 Solobacterium moorei_nov_89.441%
- SPN1965 Butyrivibrio fibrisolvens_nov_91.546%
- SPN1976 Prevotella saccharolytica_nov_93.661%
- SPN1988 Streptococcus anginosus_nov_94.433%
- SPN1996 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_91.494%
- SPN20 Leptotrichia sp._HMT_212_nov_93.043%
- SPN2002 Haemophilus sp._HMT_036_nov_95.501%
- SPN2006 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.126%
- SPN2016 Desulfovibrio legallii_nov_94.343%
- SPN2028 Syntrophotalea carbinolica_nov_83.497%
- SPN203 Dialister succinatiphilus_nov_93.738%
- SPN2039 Acinetobacter indicus_nov_97.938%
- SPN2050 Alloprevotella sp._HMT_913_nov_97.751%
- SPN2061 Aggregatibacter actinomycetemcomitans_nov_97.541%
- SPN2062 Prevotella sp._MOT-128_nov_86.992%
- SPN2071 Blautia luti_nov_97.059%

- SPN805 Prevotella sp._HMT_293_nov_89.024%
- SPN806 Bacteroidetes_[G-3] bacterium_HMT_365_nov_87.373%
- SPN807 Corynebacterium durum_nov_91.968%
- SPN808 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_91.232%
- SPN809 Coprococcus comes_nov_97.275%
- SPN81 Oscillibacter valericigenes_nov_88.501%
- SPN810 Alloprevotella sp._HMT_473_nov_95.918%
- SPN812 Streptococcus sanguinis_nov_96.371%
- SPN813 Enterocloster boltae_nov_91.649%
- SPN814 Clostridium cellulovorans_nov_82.505%
- SPN815 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_88.797%
- SPN816 Oscillibacter valericigenes_nov_96.050%
- SPN817 Capnocytophaga ochracea_nov_94.792%
- SPN818 Peptostreptococcaceae_[G-3] bacterium_HMT_950_nov_88.727%
- SPN819 Prevotella oulorum_nov_89.837%
- SPN820 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_88.706%
- SPN821 Prevotella copri_nov_96.741%
- SPN822 Prevotella copri_nov_97.755%
- SPN823 Prevotella copri_nov_93.469%
- SPN824 Pseudoleptotrichia goodfellowii_nov_92.241%
- SPN825 Coprococcus eutactus_nov_95.085%
- SPN826 Prevotella saccharolytica_nov_95.092%
- SPN827 Tannerella forsythia_nov_88.613%
- SPN828 Oscillibacter valericigenes_nov_88.017%
- SPN829 Haemophilus haemolyticus_nov_94.251%
- SPN830 Streptococcus peroris_nov_94.949%
- SPN831 Fournierella massiliensis_nov_91.340%
- SPN832 Bacteroidetes_[G-3] bacterium_HMT_899_nov_86.939%
- SPN834 Sporobacter termitidis_nov_86.325%
- SPN835 Corynebacterium massiliense_nov_95.319%
- SPN836 Streptococcus gordonii_nov_94.036%
- SPN837 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_84.289%
- SPN838 Eubacteriales_[G-1] bacterium_MOT-144_nov_87.397%
- SPN839 Mollicutes_[G-2] bacterium_HMT_906_nov_82.922%
- SPN840 Slackia equolifaciens_nov_88.362%
- SPN841 Eubacteriales_[G-1] bacterium_MOT-144_nov_89.278%
- SPN842 Fretribacterium sp._HMT_362_nov_97.817%
- SPN844 Prevotella oulorum_nov_89.837%
- SPN845 Schaalia sp._HMT_178_nov_94.908%
- SPN846 Oscillibacter valericigenes_nov_86.392%