

Species

- SP2379 Kocuria palustris
- SP282 Fusobacterium sp._HMT_204
- SP285 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345
- SP292 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP294 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874
- SP295 Catonella sp._HMT_164
- SP296 Corynebacterium matruchotii
- SP30 Leptotrichia sp._HMT_392
- SP308 Fusobacterium periodonticum
- SP310 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875
- SP322 Corynebacterium suicordis
- SP323 Johnsonella ignava
- SP346 Desulfohalobium sp._HMT_041
- SP358 Abiotrophia defectiva
- SP36 Ligilactobacillus animalis
- SP360 Psychrobacter sanguinis
- SP361 Treponema denticola
- SP375 Fusicatenibacter saccharivorans
- SP392 Neisseria oralis
- SP398 Rothia endophytica
- SP40 Arachnia propionica
- SP415 Ruminococcus champanellensis
- SP419 Porphyromonas sp._HMT_277
- SP42 Methanobrevibacter smithii
- SP425 Lactobacillus johnsonii
- SP431 Pseudoleptotrichia goodfellowii
- SP434 Prevotella sp._HMT_317
- SP438 Streptococcus thoraltensis
- SP441 Streptococcus sp._HMT_423
- SP443 Streptococcus sinensis
- SP445 Actinomyces sp._HMT_170
- SP45 Lachnoanaerobaculum sp._HMT_083
- SP459 Terrisporobacter mayombeii
- SP460 Cardiobacterium valvarum
- SP47 Fusobacterium nucleatum
- SP475 Actinomyces viscosus
- SP490 Porphyromonas sp._HMT_930
- SP492 Bacteroidetes_[G-5] bacterium_HMT_511
- SP496 Streptococcus oralis
- SP499 Ruminococcaceae_[G-1] bacterium_HMT_075

- SPN1763 Selenomonas sp._HMT_478_nov_96.640%
- SPN177 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_92.453%
- SPN1775 Treponema sp._HMT_264_nov_83.534%
- SPN1778 Haemophilus sp._HMT_036_nov_95.501%
- SPN1797 Peptostreptococcaceae_[G-5] bacterium_HMT_493_nov_95.417%
- SPN1808 Eubacteriales_[G-3] bacterium_MOT-163_nov_86.680%
- SPN1819 Selenomonas sp._HMT_892_nov_97.446%
- SPN1821 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.292%
- SPN1833 Gemella haemolysans_nov_97.984%
- SPN1844 Ruminococcus champanellensis_nov_90.086%
- SPN1853 Porphyromonas pasteri_nov_96.939%
- SPN1854 Prevotella sp._HMT_304_nov_95.492%
- SPN1865 Catenibacterium mitsuokai_nov_97.934%
- SPN1876 Anaerocolumna cellulositytica_nov_87.657%
- SPN1885 Peptidiphaga gingivicola_nov_95.297%
- SPN189 Actinobacillus lignieresii_nov_97.342%
- SPN1895 Prevotella brevis_nov_89.184%
- SPN1897 Comamonas faecalis_nov_91.803%
- SPN1908 Haemophilus haemolyticus_nov_94.908%
- SPN1920 Bacteroidetes_[G-7] bacterium_HMT_911_nov_89.528%
- SPN1932 Acinetobacter tandoii_nov_97.308%
- SPN1943 Actinomyces gerencseriae_nov_94.061%
- SPN1953 Solobacterium moorei_nov_89.441%
- SPN196 Dialister succinatiphilus_nov_93.738%
- SPN1965 Catonella morbi_nov_97.699%
- SPN1966 Butyrivibrio fibrilvolens_nov_91.546%
- SPN1977 Prevotella saccharolytica_nov_93.661%
- SPN1988 Streptococcus anginosus_nov_94.433%
- SPN1999 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_91.494%
- SPN2006 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.126%
- SPN2015 Haemophilus haemolyticus_nov_95.910%
- SPN2018 Desulfovibrio legallii_nov_94.343%
- SPN2028 Syntrophotalea carbinolica_nov_83.497%
- SPN204 Kineothrix alysoidea_nov_91.770%
- SPN2041 Acinetobacter indicus_nov_97.938%
- SPN2051 Alloprevotella sp._HMT_913_nov_97.751%
- SPN2062 Prevotella sp._MOT-128_nov_86.992%
- SPN2073 Aggregatibacter actinomycetemcomitans_nov_97.541%
- SPN2074 Blautia luti_nov_97.059%
- SPN2085 Treponema sp._HMT_236_nov_97.363%

- SPN817 Corynebacterium durum_nov_91.968%
- SPN818 Prevotella sp._HMT_293_nov_89.024%
- SPN819 Bacteroidetes_[G-3] bacterium_HMT_365_nov_87.373%
- SPN820 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_91.232%
- SPN821 Coprococcus comes_nov_97.275%
- SPN822 Enterocloster bolteae_nov_91.649%
- SPN823 Prevotella copri_nov_97.755%
- SPN824 Clostridium cellulovorans_nov_82.505%
- SPN825 Alloprevotella sp._HMT_473_nov_95.918%
- SPN826 Streptococcus suis_nov_93.387%
- SPN827 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_88.797%
- SPN828 Oscillibacter valericigenes_nov_96.050%
- SPN829 Capnocytophaga ochracea_nov_94.792%
- SPN830 Peptostreptococcaceae_[G-3] bacterium_HMT_950_nov_88.727%
- SPN831 Prevotella oulorum_nov_89.837%
- SPN832 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_88.706%
- SPN833 Prevotella copri_nov_96.741%
- SPN835 Prevotella copri_nov_93.469%
- SPN836 Prevotella saccharolytica_nov_95.092%
- SPN837 Coprococcus eutactus_nov_95.085%
- SPN838 Fusobacterium sp._HMT_370_nov_97.629%
- SPN839 Haemophilus haemolyticus_nov_94.251%
- SPN840 Tannerella forsythia_nov_88.613%
- SPN841 Oscillibacter valericigenes_nov_88.017%
- SPN842 Streptococcus peroris_nov_94.949%
- SPN843 Fournierella massiliensis_nov_91.340%
- SPN844 Bacteroidetes_[G-3] bacterium_HMT_899_nov_86.939%
- SPN846 Sporobacter termitidis_nov_86.325%
- SPN847 Corynebacterium massiliense_nov_95.319%
- SPN848 Clostridiales_[F-1][G-2] bacterium_HMT_402_nov_84.289%
- SPN849 Eubacteriales_[G-1] bacterium_MOT-144_nov_87.397%
- SPN850 Pseudoleptotrichia goodfellowii_nov_92.241%
- SPN851 Aggregatibacter aphrophilus_nov_96.115%
- SPN852 Mollicutes_[G-2] bacterium_HMT_906_nov_82.922%
- SPN853 Slackia equolifaciens_nov_88.362%
- SPN854 Eubacteriales_[G-1] bacterium_MOT-144_nov_89.278%
- SPN857 Fretibacterium sp._HMT_362_nov_97.817%
- SPN858 Prevotella oulorum_nov_89.837%
- SPN859 Schaalia sp._HMT_178_nov_94.908%
- SPN860 Oscillibacter valericigenes_nov_86.392%