

Species

- SP272 Coprococcus comes
- SP285 Treponema denticola
- SP287 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875
- SP288 Prevotella falsenii
- SP291 Selenomonas sp._HMT_892
- SP293 Leptotrichia trevisanii
- SP294 Johnsonella ignava
- SP30 Abiotrophia defectiva
- SP306 Actinomyces sp._HMT_896
- SP308 Aerococcus viridans
- SP310 Filifactor alocis
- SP32 Sphingomonas paucimobilis
- SP320 Fibrobacter intestinalis
- SP33 Limosilactobacillus reuteri
- SP335 Bifidobacterium bifidum
- SP337 Streptococcus downii
- SP344 Kocuria carniphila
- SP349 Bacteroidales_[G-2] bacterium_HMT_274
- SP351 Leptotrichia sp._HMT_225
- SP352 Riemerella sp._HMT_322
- SP356 Micrococcus aloeverae
- SP359 Porphyromonas gingivalis
- SP366 Campylobacter showae
- SP367 Kocuria rhizophila
- SP372 Treponema maltophilum
- SP38 Fretibacterium fastidiosum
- SP383 Peptostreptococcus stomatis
- SP389 Streptococcus sinensis
- SP399 Streptococcus periodonticum
- SP4 Prevotella copri
- SP40 Fusobacterium nucleatum
- SP409 Tannerella forsythia
- SP41 Porphyromonas pasteri
- SP412 Actinomyces sp._HMT_170
- SP42 Coprococcus eutactus
- SP423 Capnocytophaga sp._HMT_878
- SP432 Streptococcus cristatus_clade_578
- SP433 Streptococcus oralis
- SP446 Cardiobacterium valvarum

- SPN1596 Prevotella stercorea_nov_95.885%
- SPN1603 Prevotella copri_nov_95.732%
- SPN1605 Streptococcus australis_nov_95.152%
- SPN1613 Bacteroidetes_[G-7] bacterium_HMT_911_nov_89.528%
- SPN1632 Blautia luti_nov_97.059%
- SPN1642 Coprococcus catus_nov_97.098%
- SPN1653 Treponema sp._HMT_247_nov_94.939%
- SPN1665 Bacteroidetes_[G-3] bacterium_HMT_899_nov_88.912%
- SPN167 Actinobacillus lignieresii_nov_97.342%
- SPN1675 Aggregatibacter aphrophilus_nov_95.697%
- SPN1676 Anaerobutyricum hallii_nov_97.053%
- SPN1687 Blautia obeum_nov_96.753%
- SPN1694 Leptotrichia sp._HMT_225_nov_96.957%
- SPN17 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_91.845%
- SPN1705 Solobacterium moorei_nov_89.441%
- SPN1713 Oscillibacter valericigenes_nov_87.137%
- SPN1721 Peptostreptococcaceae_[G-7] bacterium_HMT_081_nov_96.414%
- SPN1733 Actinomyces dentalis_nov_96.192%
- SPN1740 Bergeyella zoohelcum_nov_92.798%
- SPN1749 Blautia luti_nov_97.059%
- SPN1759 Butyrivibrio fibrisolvens_nov_91.546%
- SPN176 Anaerobutyricum hallii_nov_97.684%
- SPN1769 Streptococcus oralis_subsp._tigurinus_clade_071_nov_96.162%
- SPN1770 Lachnospiraceae_[G-8] bacterium_HMT_500_nov_97.263%
- SPN1780 Dielma fastidiosa_nov_81.162%
- SPN1788 Neisseria cinerea_nov_97.541%
- SPN1800 Corynebacterium durum_nov_93.865%
- SPN1809 Clostridium sardiniense_nov_97.619%
- SPN1820 Selenomonas sp._HMT_892_nov_97.228%
- SPN1832 Dialister succinatiphilus_nov_93.738%
- SPN1840 Capnocytophaga gingivalis_nov_95.992%
- SPN1844 Prevotella oulorum_nov_89.837%
- SPN1850 Bacteroidetes_[G-3] bacterium_HMT_280_nov_89.876%
- SPN186 Prevotella oulorum_nov_89.837%
- SPN1860 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_92.453%
- SPN1872 Aggregatibacter aphrophilus_nov_96.524%
- SPN1874 Porphyromonas pasteri_nov_97.347%
- SPN1881 Blautia luti_nov_96.218%
- SPN1891 Oscillibacter valericigenes_nov_88.501%

- SPN339 Methanospira cuniculi_nov_95.652%
- SPN340 Haemophilus sp._HMT_036_nov_94.683%
- SPN341 Granulicatella elegans_nov_96.774%
- SPN352 Streptococcus gordonii_nov_93.638%
- SPN364 Streptococcus gordonii_nov_94.036%
- SPN376 Capnocytophaga sp._HMT_878_nov_97.028%
- SPN387 Eikenella sp._HMT_011_nov_96.728%
- SPN40 Dysosmobacter welbionis_nov_86.307%
- SPN400 Aggregatibacter sp._HMT_898_nov_94.490%
- SPN412 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_95.407%
- SPN424 Capnocytophaga sp._HMT_332_nov_95.763%
- SPN432 Mogibacterium neglectum_nov_89.855%
- SPN433 Capnocytophaga cynodegmi_nov_90.208%
- SPN437 Capnocytophaga sp._HMT_412_nov_95.597%
- SPN438 Porphyromonas sp._HMT_930_nov_95.473%
- SPN450 Neisseria flava_nov_97.947%
- SPN461 Campylobacter rectus_nov_96.957%
- SPN472 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_96.868%
- SPN484 Aggregatibacter sp._HMT_898_nov_97.331%
- SPN496 Selenomonas sp._HMT_892_nov_96.832%
- SPN50 Alloprevotella sp._HMT_473_nov_97.951%
- SPN508 Propionivibrio dicarboxylicus_nov_93.802%
- SPN520 Peptostreptococcaceae_[G-7] bacterium_HMT_081_nov_94.105%
- SPN533 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_95.407%
- SPN543 Bacteroidetes_[G-3] bacterium_HMT_899_nov_87.730%
- SPN546 Alloprevotella sp._HMT_308_nov_95.483%
- SPN557 Neisseria cinerea_nov_97.951%
- SPN558 Lachnospiraceae_[G-2] bacterium_HMT_088_nov_94.958%
- SPN57 Sporobacter termitidis_nov_85.837%
- SPN570 Capnocytophaga sputigena_nov_95.754%
- SPN582 Streptococcus sp._HMT_066_nov_95.142%
- SPN594 Prevotella sp._HMT_317_nov_97.746%
- SPN606 Streptococcus troglodytidis_nov_92.079%
- SPN615 Pseudostreptobacillus hongkongensis_nov_92.797%
- SPN627 Peptococcus sp._HMT_167_nov_97.250%
- SPN639 Prevotella saccharolytica_nov_93.661%
- SPN651 Riemerella sp._HMT_322_nov_97.942%
- SPN66 Flavonifractor plautii_nov_93.996%
- SPN663 Porphyromonas endodontalis_nov_96.327%