

## Species

SP225	SPN1596	SPN339
SP272 Coprococcus comes	SPN1596 Prevotella stercorea_nov_95.885%	SPN339 Methanospaera cuniculi_nov_95.652%
SP285 Treponema denticola	SPN1603 Prevotella copri_nov_95.732%	SPN340 Haemophilus sp._HMT_036_nov_94.683%
SP287 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875	SPN1605 Streptococcus australis_nov_95.152%	SPN341 Granulicatella elegans_nov_96.774%
SP288 Prevotella falsenii	SPN1613 Bacteroidetes_[G-7] bacterium_HMT_911_nov_89.528%	SPN352 Streptococcus gordonii_nov_93.638%
SP291 Selenomonas sp._HMT_892	SPN1632 Blautia luti_nov_97.059%	SPN364 Streptococcus gordonii_nov_94.036%
SP293 Leptotrichia trevisanii	SPN1642 Coprococcus catus_nov_97.098%	SPN376 Capnocytophaga sp._HMT_878_nov_97.028%
SP294 Johnsonella ignava	SPN1653 Treponema sp._HMT_247_nov_94.939%	SPN387 Eikenella sp._HMT_011_nov_96.728%
SP30 Abiotrophia defectiva	SPN1665 Bacteroidetes_[G-3] bacterium_HMT_899_nov_88.912%	SPN40 Dysosmobacter welbionis_nov_86.307%
SP306 Actinomyces sp._HMT_896	SPN167 Actinobacillus lignieresii_nov_97.342%	SPN400 Aggregatibacter sp._HMT_898_nov_94.490%
SP308 Aerococcus viridans	SPN1675 Aggregatibacter aphrophilus_nov_95.697%	SPN412 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_95.407%
SP310 Filifactor alocis	SPN1676 Anaerobutyricum hallii_nov_97.053%	SPN424 Capnocytophaga sp._HMT_332_nov_95.763%
SP32 Sphingomonas paucimobilis	SPN1687 Blautia obeum_nov_96.753%	SPN432 Mogibacterium neglectum_nov_89.855%
SP320 Fibrobacter intestinalis	SPN1694 Leptotrichia sp._HMT_225_nov_96.957%	SPN433 Capnocytophaga cynodegmi_nov_90.208%
SP33 Limosilactobacillus reuteri	SPN17 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_91.845%	SPN437 Capnocytophaga sp._HMT_412_nov_95.597%
SP335 Bifidobacterium bifidum	SPN1705 Solobacterium moorei_nov_89.441%	SPN438 Porphyromonas sp._HMT_930_nov_95.473%
SP337 Streptococcus downii	SPN1713 Oscillibacter valericigenes_nov_87.137%	SPN450 Neisseria flava_nov_97.947%
SP344 Kocuria carniphila	SPN1721 Peptostreptococcaceae_[G-7] bacterium_HMT_081_nov_96.414%	SPN461 Campylobacter rectus_nov_96.957%
SP349 Bacteroidales_[G-2] bacterium_HMT_274	SPN1733 Actinomyces dentalis_nov_96.192%	SPN472 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_96.868%
SP351 Leptotrichia sp._HMT_225	SPN1740 Bergeyella zoohelcum_nov_92.798%	SPN484 Aggregatibacter sp._HMT_898_nov_97.331%
SP352 Riemerella sp._HMT_322	SPN1749 Blautia luti_nov_97.059%	SPN496 Selenomonas sp._HMT_892_nov_96.832%
SP356 Micrococcus aloeverae	SPN1759 Butyrivibrio fibrisolvens_nov_91.546%	SPN50 Alloprevotella sp._HMT_473_nov_97.951%
SP359 Porphyromonas gingivalis	SPN176 Anaerobutyricum hallii_nov_97.684%	SPN508 Propionivibrio dicarboxylicus_nov_93.802%
SP366 Campylobacter showae	SPN1769 Streptococcus oralis_subsp._tigurinus_clade_071_nov_96.162%	SPN520 Peptostreptococcaceae_[G-7] bacterium_HMT_081_nov_94.105%
SP367 Kocuria rhizophila	SPN1770 Lachnospiraceae_[G-8] bacterium_HMT_500_nov_97.263%	SPN533 Lachnospiraceae_[G-3] bacterium_HMT_100_nov_95.407%
SP372 Treponema maltophilum	SPN1780 Dielma fastidiosa_nov_81.162%	SPN543 Bacteroidetes_[G-3] bacterium_HMT_899_nov_87.730%
SP38 Fretibacterium fastidiosum	SPN1788 Neisseria cinerea_nov_97.541%	SPN546 Alloprevotella sp._HMT_308_nov_95.483%
SP383 Peptostreptococcus stomatis	SPN1800 Corynebacterium durum_nov_93.865%	SPN557 Neisseria cinerea_nov_97.951%
SP389 Streptococcus sinensis	SPN1809 Clostridium sardiniense_nov_97.619%	SPN558 Lachnospiraceae_[G-2] bacterium_HMT_088_nov_94.958%
SP399 Streptococcus periodonticum	SPN1820 Selenomonas sp._HMT_892_nov_97.228%	SPN57 Sporobacter termitidis_nov_85.837%
SP4 Prevotella copri	SPN1832 Dialister succinatiphilus_nov_93.738%	SPN570 Capnocytophaga sputigena_nov_95.754%
SP40 Fusobacterium nucleatum	SPN1840 Capnocytophaga gingivalis_nov_95.992%	SPN582 Streptococcus sp._HMT_066_nov_95.142%
SP409 Tannerella forsythia	SPN1844 Prevotella oulorum_nov_89.837%	SPN594 Prevotella sp._HMT_317_nov_97.746%
SP41 Porphyromonas pasteri	SPN1850 Bacteroidetes_[G-3] bacterium_HMT_280_nov_89.876%	SPN606 Streptococcus troglodytidis_nov_92.079%
SP412 Actinomyces sp._HMT_170	SPN186 Prevotella oulorum_nov_89.837%	SPN615 Pseudostreptobacillus hongkongensis_nov_92.797%
SP42 Coprococcus eutactus	SPN1860 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_92.453%	SPN627 Peptococcus sp._HMT_167_nov_97.250%
SP423 Capnocytophaga sp._HMT_878	SPN1872 Aggregatibacter aphrophilus_nov_96.524%	SPN639 Prevotella saccharolytica_nov_93.661%
SP432 Streptococcus cristatus_clade_578	SPN1874 Porphyromonas pasteri_nov_97.347%	SPN651 Riemerella sp._HMT_322_nov_97.942%
SP433 Streptococcus oralis	SPN1881 Blautia luti_nov_96.218%	SPN66 Flavonifractor plautii_nov_93.996%
SP446 Cardiobacterium valvarum	SPN1891 Oscillibacter valericigenes_nov_88.501%	SPN663 Porphyromonas endodontalis_nov_96.327%
SP450	SPN1899	SPN664