

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%