

Species

| | | |
|--|---|--|
| SP225 <i>Coproccoccus copri</i> _HMT_122 | SPN1596 <i>Prevotella stercorea</i> _nov_95.885% | SPN339 <i>Methanosphaera cuniculi</i> _nov_95.652% |
| SP285 <i>Treponema denticola</i> | SPN1603 <i>Prevotella copri</i> _nov_95.732% | SPN340 <i>Haemophilus</i> sp._HMT_036_nov_94.683% |
| SP287 <i>Absconditabacteria</i> _(SR1)_[G-1]_bacterium_HMT_875 | SPN1605 <i>Streptococcus australis</i> _nov_95.152% | SPN341 <i>Granulicatella elegans</i> _nov_96.774% |
| SP288 <i>Prevotella falsenii</i> | SPN1613 <i>Bacteroidetes</i> _[G-7]_bacterium_HMT_911_nov_89.528% | SPN352 <i>Streptococcus gordonii</i> _nov_93.638% |
| SP291 <i>Selenomonas</i> sp._HMT_892 | SPN1632 <i>Blautia luti</i> _nov_97.059% | SPN364 <i>Streptococcus gordonii</i> _nov_94.036% |
| SP293 <i>Leptotrichia trevisanii</i> | SPN1642 <i>Coproccoccus catus</i> _nov_97.098% | SPN376 <i>Capnocytophaga</i> sp._HMT_878_nov_97.028% |
| SP294 <i>Johnsonella ignava</i> | SPN1653 <i>Treponema</i> sp._HMT_247_nov_94.939% | SPN387 <i>Eikenella</i> sp._HMT_011_nov_96.728% |
| SP30 <i>Abiotrophia defectiva</i> | SPN1665 <i>Bacteroidetes</i> _[G-3]_bacterium_HMT_899_nov_88.912% | SPN40 <i>Dysosmobacter welbionis</i> _nov_86.307% |
| SP306 <i>Actinomyces</i> sp._HMT_896 | SPN167 <i>Actinobacillus lignieresii</i> _nov_97.342% | SPN400 <i>Aggregatibacter</i> sp._HMT_898_nov_94.490% |
| SP308 <i>Aerococcus viridans</i> | SPN1675 <i>Aggregatibacter aphrophilus</i> _nov_95.697% | SPN412 <i>Lachnospiraceae</i> _[G-3]_bacterium_HMT_100_nov_95.407% |
| SP310 <i>Filifactor alocis</i> | SPN1676 <i>Anaerobutyricum hallii</i> _nov_97.053% | SPN424 <i>Capnocytophaga</i> sp._HMT_332_nov_95.763% |
| SP32 <i>Sphingomonas paucimobilis</i> | SPN1687 <i>Blautia obeum</i> _nov_96.753% | SPN432 <i>Mogibacterium neglectum</i> _nov_89.855% |
| SP320 <i>Fibrobacter intestinalis</i> | SPN1694 <i>Leptotrichia</i> sp._HMT_225_nov_96.957% | SPN433 <i>Capnocytophaga cynodegmi</i> _nov_90.208% |
| SP33 <i>Limosilactobacillus reuteri</i> | SPN17 <i>Lachnospiraceae</i> _[G-14]_bacterium_MOT-185_nov_91.845% | SPN437 <i>Capnocytophaga</i> sp._HMT_412_nov_95.597% |
| SP335 <i>Bifidobacterium bifidum</i> | SPN1705 <i>Solobacterium moorei</i> _nov_89.441% | SPN438 <i>Porphyromonas</i> sp._HMT_930_nov_95.473% |
| SP337 <i>Streptococcus downii</i> | SPN1713 <i>Oscillibacter valericigenes</i> _nov_87.137% | SPN450 <i>Neisseria flava</i> _nov_97.947% |
| SP344 <i>Kocuria carniphila</i> | SPN1721 <i>Peptostreptococcaceae</i> _[G-7]_bacterium_HMT_081_nov_96.414% | SPN461 <i>Campylobacter rectus</i> _nov_96.957% |
| SP349 <i>Bacteroidales</i> _[G-2]_bacterium_HMT_274 | SPN1733 <i>Actinomyces dentalis</i> _nov_96.192% | SPN472 <i>Lachnospiraceae</i> _[G-3]_bacterium_HMT_100_nov_96.868% |
| SP351 <i>Leptotrichia</i> sp._HMT_225 | SPN1740 <i>Bergeyella zoohelcum</i> _nov_92.798% | SPN484 <i>Aggregatibacter</i> sp._HMT_898_nov_97.331% |
| SP352 <i>Riemerella</i> sp._HMT_322 | SPN1749 <i>Blautia luti</i> _nov_97.059% | SPN496 <i>Selenomonas</i> sp._HMT_892_nov_96.832% |
| SP356 <i>Micrococcus aloeverae</i> | SPN1759 <i>Butyrivibrio fibrisolvens</i> _nov_91.546% | SPN50 <i>Alloprevotella</i> sp._HMT_473_nov_97.951% |
| SP359 <i>Porphyromonas gingivalis</i> | SPN176 <i>Anaerobutyricum hallii</i> _nov_97.684% | SPN508 <i>Propionivibrio dicarboxylicus</i> _nov_93.802% |
| SP366 <i>Campylobacter showae</i> | SPN1769 <i>Streptococcus oralis</i> _subsp._tigurinus_clade_071_nov_96.162% | SPN520 <i>Peptostreptococcaceae</i> _[G-7]_bacterium_HMT_081_nov_94.105% |
| SP367 <i>Kocuria rhizophila</i> | SPN1770 <i>Lachnospiraceae</i> _[G-8]_bacterium_HMT_500_nov_97.263% | SPN533 <i>Lachnospiraceae</i> _[G-3]_bacterium_HMT_100_nov_95.407% |
| SP372 <i>Treponema maltophilum</i> | SPN1780 <i>Dielma fastidiosa</i> _nov_81.162% | SPN543 <i>Bacteroidetes</i> _[G-3]_bacterium_HMT_899_nov_87.730% |
| SP38 <i>Fretibacterium fastidiosum</i> | SPN1788 <i>Neisseria cinerea</i> _nov_97.541% | SPN546 <i>Alloprevotella</i> sp._HMT_308_nov_95.483% |
| SP383 <i>Peptostreptococcus stomatis</i> | SPN1800 <i>Corynebacterium durum</i> _nov_93.865% | SPN557 <i>Neisseria cinerea</i> _nov_97.951% |
| SP389 <i>Streptococcus sinensis</i> | SPN1809 <i>Clostridium sardiniense</i> _nov_97.619% | SPN558 <i>Lachnospiraceae</i> _[G-2]_bacterium_HMT_088_nov_94.958% |
| SP399 <i>Streptococcus periodonticum</i> | SPN1820 <i>Selenomonas</i> sp._HMT_892_nov_97.228% | SPN57 <i>Sporobacter termitidis</i> _nov_85.837% |
| SP4 <i>Prevotella copri</i> | SPN1832 <i>Dialister succinatiphilus</i> _nov_93.738% | SPN570 <i>Capnocytophaga sputigena</i> _nov_95.754% |
| SP40 <i>Fusobacterium nucleatum</i> | SPN1840 <i>Capnocytophaga gingivalis</i> _nov_95.992% | SPN582 <i>Streptococcus</i> sp._HMT_066_nov_95.142% |
| SP409 <i>Tannerella forsythia</i> | SPN1844 <i>Prevotella oulorum</i> _nov_89.837% | SPN594 <i>Prevotella</i> sp._HMT_317_nov_97.746% |
| SP41 <i>Porphyromonas pasteri</i> | SPN1850 <i>Bacteroidetes</i> _[G-3]_bacterium_HMT_280_nov_89.876% | SPN606 <i>Streptococcus troglodytidis</i> _nov_92.079% |
| SP412 <i>Actinomyces</i> sp._HMT_170 | SPN186 <i>Prevotella oulorum</i> _nov_89.837% | SPN615 <i>Pseudostreptobacillus hongkongensis</i> _nov_92.797% |
| SP42 <i>Coproccoccus eutactus</i> | SPN1860 <i>Lachnospiraceae</i> _[G-5]_bacterium_MOT-170_nov_92.453% | SPN627 <i>Peptococcus</i> sp._HMT_167_nov_97.250% |
| SP423 <i>Capnocytophaga</i> sp._HMT_878 | SPN1872 <i>Aggregatibacter aphrophilus</i> _nov_96.524% | SPN639 <i>Prevotella saccharolytica</i> _nov_93.661% |
| SP432 <i>Streptococcus cristatus</i> _clade_578 | SPN1874 <i>Porphyromonas pasteri</i> _nov_97.347% | SPN651 <i>Riemerella</i> sp._HMT_322_nov_97.942% |
| SP433 <i>Streptococcus oralis</i> | SPN1881 <i>Blautia luti</i> _nov_96.218% | SPN66 <i>Flavonifractor plautii</i> _nov_93.996% |
| SP446 <i>Cardiobacterium valvarum</i> | SPN1891 <i>Oscillibacter valericigenes</i> _nov_88.501% | SPN663 <i>Porphyromonas endodontalis</i> _nov_96.327% |
| SP453 <i>Chlorobacterium</i> sp._HMT_100 | SPN1900 <i>Porphyromonas</i> sp._HMT_100_nov_95.407% | SPN664 <i>Porphyromonas</i> sp._HMT_100_nov_95.407% |