

Species

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| SP14 | Bacillus sp._Oral_Taxon_B77 | SPN1426 | Lacnociostridium sp._str._M62/1_nov_88.565% | SPN2289 | Lacnociostridium bolteae_nov_91.458% |
| SP15 | Rothia aeria | SPN1438 | Acetivibrio ethanolgignens_nov_87.071% | SPN2301 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_84.940% |
| SP159 | Propionibacterium sp._oral_taxon_194 | SPN1449 | Lachnociostridium fissicatena_nov_94.824% | SPN2302 | Saccharopolyspora dendranthemae_nov_80.802% |
| SP16 | Prevotella sp._oral_taxon_313 | SPN1471 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.657% | SPN2312 | Eubacterium ramulus_nov_87.344% |
| SP18 | Streptococcus dentisani | SPN1477 | Lacnociostridium sp._str._M62/1_nov_90.417% | SPN2324 | Fusicatenibacter saccharivorans_nov_89.669% |
| SP19 | Veillonella atypica | SPN1483 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_86.800% | SPN2335 | TM7_[G-1] sp._oral_taxon_348_nov_95.133% |
| SP2 | Peptostreptococcus stomatis | SPN1484 | Streptococcus troglodytidis_nov_87.209% | SPN2338 | Fusicatenibacter saccharivorans_nov_89.256% |
| SP21 | Enterococcus faecalis | SPN1495 | Lacnociostridium asparagiforme_nov_93.983% | SPN2357 | Manihot esculenta_Oral_Taxon_C60_nov_97.945% |
| SP22 | Lactobacillus reuteri_genosp._2 | SPN1506 | Lachnociostridium polysaccharolyticum_nov_92.340% | SPN2370 | Eubacterium ramulus_nov_88.636% |
| SP23 | Abiotrophia defectiva | SPN1512 | Streptococcus danieliae_nov_94.926% | SPN2381 | Oscillibacter valericigenes_nov_95.031% |
| SP25 | Escherichia coli | SPN1527 | Eubacterium ventriosum_nov_93.361% | SPN2393 | Roseburia inulinivorans_nov_90.760% |
| SP27 | Streptococcus tigurinus | SPN1538 | Alphaproteobacteria_[G] sp._Oral_Taxon_A28_nov_92.634% | SPN2403 | Dorea formicigenerans_nov_94.387% |
| SP28 | Streptococcus mitis | SPN1549 | Lacnociostridium bolteae_nov_91.213% | SPN2414 | Triticum aestivum_Oral_Taxon_D31_nov_97.925% |
| SP3 | Streptococcus sp._oral_taxon_064 | SPN1560 | Anaerostipes hadrus_nov_86.242% | SPN2415 | Anaerocolumna xylanovorans_nov_86.922% |
| SP31 | Lautropia mirabilis | SPN1569 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.193% | SPN2426 | Eubacterium ventriosum_nov_91.564% |
| SP32 | Manihot esculenta_Oral_Taxon_C60 | SPN1575 | unclassified_Ruminococcaceae sp._str._D16_nov_92.577% | SPN2437 | Blautia coccoides_nov_90.683% |
| SP33 | Mucispirillum schaedleri | SPN1581 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_86.139% | SPN2449 | Lacnociostridium sp._SS3/4_nov_90.408% |
| SP35 | Triticum aestivum_Oral_Taxon_D31 | SPN1591 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_86.600% | SPN2452 | Ruminococcus champanellensis_nov_91.471% |
| SP36 | Porphyromonas sp._oral_taxon_279 | SPN1593 | Tyzzereella lactatifermentans_nov_91.992% | SPN2461 | Lachnociostridium xylanolyticum_nov_90.249% |
| SP37 | Selenomonas sp._oral_taxon_136 | SPN1612 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_84.692% | SPN2473 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.571% |
| SP4 | Streptococcus cristatus | SPN1619 | Desulfovibrio legallii_nov_92.415% | SPN2483 | Lachnociostridium herbivorans_nov_92.961% |
| SP40 | Enterococcus casseliflavus | SPN1622 | Lachnociostridium xylanolyticum_nov_93.827% | SPN2496 | Ruminococcaceae_[G-2] sp._oral_taxon_085_nov_87.789% |
| SP43 | Campylobacter concisus | SPN1634 | Roseburia inulinivorans_nov_88.753% | SPN2508 | Lacnociostridium sp._SS3/4_nov_89.938% |
| SP45 | Lactobacillus intestinalis | SPN1642 | Eubacterium coprostanoligenes_nov_91.489% | SPN2520 | Eubacterium siraeum_nov_91.239% |
| SP46 | Selenomonas sp._oral_taxon_149 | SPN1652 | Lachnociostridium polysaccharolyticum_nov_85.892% | SPN2531 | Ruminococcus gauvreauii_nov_89.419% |
| SP48 | Streptococcus parasanguinis_II | SPN1663 | Streptococcus infantis_nov_97.379% | SPN2532 | Blautia coccoides_nov_94.617% |
| SP49 | Calycanthus floridus_Oral_Taxon_D07 | SPN1693 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.714% | SPN2555 | Butyrivibrio pullicaecorum_nov_91.020% |
| SP51 | Prevotella melaninogenica | SPN1695 | Lacnociostridium hathewayi_nov_89.388% | SPN2567 | Oscillibacter valericigenes_nov_93.634% |
| SP56 | Streptococcus infantis | SPN1707 | Eubacterium xylanophilum_nov_91.129% | SPN2570 | Blautia producta_nov_95.228% |
| SP6 | Lactobacillus murinus | SPN1719 | Blautia gnavus_nov_92.739% | SPN2579 | Bacteroides thetaiotaomicron_nov_94.715% |
| SP60 | Fusobacterium periodonticum | SPN1727 | Ruminococcaceae_[G-2] sp._oral_taxon_085_nov_90.126% | SPN2591 | Lacnociostridium hathewayi_nov_89.571% |
| SP61 | Fusobacterium nucleatum_subsp._polymorphum | SPN1732 | Oribacterium parvum_nov_89.855% | SPN2604 | Enterorhabdus caecimuris_nov_89.270% |
| SP63 | Neisseria sicca | SPN1744 | Streptococcus rattii_nov_84.633% | SPN2649 | Streptococcus acidominimus_nov_94.632% |
| SP64 | Streptococcus sp._oral_taxon_423 | SPN1754 | Coprococcus catus_nov_89.809% | SPN269 | Ruminococcaceae_[G-2] sp._oral_taxon_085_nov_89.873% |
| SP67 | Capnocytophaga sputigena | SPN1771 | Abiotrophia defectiva_nov_93.281% | SPN30 | Ruminococcaceae_[G-2] sp._oral_taxon_085_nov_89.518% |
| SP69 | Enterorhabdus caecimuris | SPN1783 | Fusicatenibacter saccharivorans_nov_90.795% | SPN333 | Barnesiella viscericola_nov_85.714% |
| SP7 | Phyllobacterium myrsinacearum | SPN1793 | Anaerostipes hadrus_nov_86.392% | SPN383 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_83.773% |
| SP70 | Staphylococcus sciuri | SPN1812 | Streptococcus alactolyticus_nov_95.391% | SPN451 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.288% |
| SP71 | Catonella morbi | SPN1825 | Streptococcus danieliae_nov_92.600% | SPN498 | Lachnociostridium xylanolyticum_nov_92.100% |
| SP77 | Alkalibacterium pelagium | SPN1835 | Catonella morbi_nov_85.832% | SPN519 | Barnesiella viscericola_nov_85.317% |
| SP8 | Propionibacterium acnes | SPN1855 | Oscillibacter valericigenes_nov_91.959% | SPN563 | Oribacterium sinus_nov_86.722% |
| SP82 | Actinomyces naeslundii | SPN1867 | Lachnociostridium polysaccharolyticum_nov_89.419% | SPN610 | Actinomyces sp._oral_taxon_175_nov_97.764% |
| SP87 | Streptococcus oralis | SPN1878 | Odoribacter splanchnicus_nov_92.608% | SPN676 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_87.149% |
| SP9 | Rhizobium rhizogenes_Oral_Taxon_D34 | SPN1890 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_87.302% | SPN723 | Lacnociostridium bolteae_nov_95.010% |
| SP95 | Streptococcus parasanguinis_I | SPN1916 | Fusicatenibacter saccharivorans_nov_89.121% | SPN791 | Alistipes senegalensis_nov_93.522% |
| SP98 | Prevotella histicola | SPN1926 | Streptococcus danieliae_nov_96.632% | SPN837 | Lachnociostridium bolteae_nov_94.375% |
| SPN1020 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_82.200% | SPN1929 | Lachnociostridium xylanolyticum_nov_94.583% | SPN900 | Blautia gnavus_nov_89.855% |
| SPN1040 | Streptococcus danieliae_nov_98.943% | SPN1950 | Lacnociostridium sp._str._M62/1_nov_87.603% | SPN954 | Blautia producta_nov_95.859% |
| SPN1069 | Blautia producta_nov_88.866% | SPN1960 | Streptococcus danieliae_nov_93.235% | SPP18 | Streptococcus salivarius_vestibularis |
| SPN110 | Prevotella shahii_nov_87.903% | SPN1961 | Streptococcus danieliae_nov_93.277% | SPP36 | Bacteroides sp._str._4136_uniformis |
| SPN111 | Marvinbryantia formatexigens_nov_91.295% | SPN1994 | Roseburia inulinivorans_nov_88.889% | SPPN106 | Lachnociostridium scindens |
| SPN1130 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_86.172% | SPN1995 | Tepidibacter mesophilus_nov_82.353% | SPPN127 | Lacnociostridium bolteae |
| SPN1163 | Calycanthus floridus_Oral_Taxon_D07_nov_97.717% | SPN2007 | Lachnociostridium fissicatena_nov_95.228% | SPN132 | Clostridium glycyrrhizinolyticum |
| SPN1165 | Roseburia inulinivorans_nov_89.004% | SPN2017 | Lacnociostridium hathewayi_nov_89.184% | SPPN143 | multigenus multispecies_sppn143_3_nov_92.083% |
| SPN1176 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_81.526% | SPN2027 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_84.479% | SPPN148 | multigenus multispecies_sppn148_3_nov_95.258% |
| SPN1183 | Lacnociostridium hathewayi_nov_87.800% | SPN2030 | Algimonas porphyrae_nov_79.724% | SPPN156 | Lachnociostridium scindens |
| SPN1196 | Eubacterium hallii_nov_84.440% | SPN2040 | Anaerocolumna xylanovorans_nov_88.189% | SPPN159 | Lacnociostridium scindens |
| SPN1207 | Streptococcus danieliae_nov_92.647% | SPN2052 | Parasutterella excrementihominis_nov_91.200% | SPPN165 | multigenus multispecies_sppn165_2_nov_89.530% |
| SPN1228 | Barnesiella viscericola_nov_82.400% | SPN2062 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.714% | SPPN173 | Lacnociostridium bolteae |
| SPN1237 | Bacteroides massiliensis_nov_93.509% | SPN2073 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_84.661% | SPPN174 | multigenus multispecies_sppn174_2_nov_95.474% |
| SPN1248 | Anaerostipes sp._str._3256FAA_nov_88.636% | SPN2074 | Enterorhabdus caecimuris_nov_89.722% | SPN179 | Roseburia multispecies_sppn179_2_nov_89.549% |
| SPN1260 | Lacnociostridium sp._SS3/4_nov_84.742% | SPN2085 | Lacnociostridium sp._SS3/4_nov_89.959% | SPPN182 | Bacteroides capillosus |
| SPN1270 | Streptococcus danieliae_nov_96.829% | SPN2103 | Lachnoanaerobaculum sp._oral_taxon_083_nov_88.333% | SPPN185 | Ruminiclostridium methylpentosum |
| SPN1271 | Streptococcus tigurinus_nov_96.016% | SPN2105 | Oscillibacter valericigenes_nov_94.398% | SPPN188 | Blautia multispecies_sppn188_2_nov_95.436% |
| SPN1281 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_84.584% | SPN2115 | Tyzzereella colinum_nov_87.013% | SPPN191 | Ruminiclostridium methylpentosum |
| SPN1287 | Barnesiella viscericola_nov_85.455% | SPN2128 | Streptococcus danieliae_nov_92.842% | SPPN193 | Roseburia multispecies_sppn193_2_nov_88.844% |
| SPN1291 | Bacteroidetes_[G-7] sp._oral_taxon_911_nov_85.542% | SPN2139 | Dorea longicatena_nov_94.583% | SPPN204 | Bacteroides capillosus |