

Species

SP109 Streptococcaceae_sanguinis	SPN118 Clostridiaceae_spp._strain_[1X21-1X-01]_nov_99.693%	SPN1801 Eubacterium_coprostanoligenes_nov_99.9216%
SP111 Oscillospiraceae_[G-7] bacterium_MOT-154	SPN1131 Eisenbergiella sp._strain_[1XD8-92]_nov_97.309%	SPN1893 Blautia_producta_nov_95.859%
SP110 Turicimonas muris	SPN1144 Duncaniella freteri_nov_87.976%	SPN1905 Lachnospiraceae_[G-1] bacterium_MOT-176_nov_94.758%
SP111 Escherichia coli	SPN1156 Streptococcus danieliae_nov_95.582%	SPN1917 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_93.905%
SP113 Rothia aeria	SPN1168 Alistipes senegalensis_nov_93.279%	SPN1936 Anaerocolumna cellulolytica_nov_87.164%
SP119 Eubacteriales_[G-1] bacterium_MOT-159	SPN1180 Thalassobacillus devorans_nov_86.372%	SPN1948 Eubacterium coprostanoligenes_nov_91.026%
SP112 Kineothrix sp._strain_[1XD8-87]	SPN1187 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_91.898%	SPN1954 Streptococcus danieliae_nov_93.952%
SP120 Porphyromonas pasteri	SPN1193 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.179%	SPN1960 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_91.667%
SP122 Veillonella atypica	SPN1205 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.320%	SPN1972 Eubacterium ramulus_nov_89.897%
SP125 Eubacteriales_[G-2] bacterium_MOT-162	SPN1217 Duncaniella freteri_nov_86.922%	SPN1984 Streptococcus danieliae_nov_96.994%
SP126 Robinsoniella sp._strain_[D2-1X-13]	SPN1218 Lachnospiraceae_[Clostridium]_herbivorans_nov_92.577%	SPN1985 Murimonas sp._strain_[WYJ26-X68]_nov_89.520%
SP13 Anaerotignum sp._strain_[1XD42-85]	SPN1227 Alistipes senegalensis_nov_92.041%	SPN1996 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.152%
SP130 Adlercreutzia muris	SPN1238 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_94.670%	SPN2007 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.802%
SP131 Arachnia rubra	SPN1250 Streptococcus danieliae_nov_97.384%	SPN2019 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.606%
SP132 Alkalibacterium pelagium	SPN1263 Lachnospiraceae_[G-1] bacterium_MOT-178_nov_92.944%	SPN2020 Saccharopolyspora dendranthema_nov_80.802%
SP14 Marvinbryantia sp._strain_[D2-1X-79]	SPN1275 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_95.949%	SPN2032 Roseburia sp._strain_[MGB-1]_nov_94.915%
SP144 Campylobacter concisus	SPN128 Duncaniella freteri_nov_88.668%	SPN2044 Salimessophilobacter vulgaris_nov_80.085%
SP15 Streptococcus salivarius	SPN1284 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.802%	SPN205 Duncaniella freteri_nov_87.298%
SP154 Lachnospiraceae_[G-1] bacterium_MOT-178	SPN1287 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_89.357%	SPN2055 Adlercreutzia equolifaciens_nov_96.983%
SP16 Streptococcus sp._MOT-012	SPN1295 Tyzzerella [Clostridium]_colinum_nov_87.013%	SPN2064 Desulfovibrio legalii_nov_92.415%
SP165 Acetatifactor sp._strain_[1XD8-48]	SPN1300 Faecalicatena fissicatena_nov_94.824%	SPN2067 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_92.551%
SP17 Lachnospiraceae_[G-14] bacterium_MOT-185	SPN1312 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.389%	SPN2077 Streptococcus infantis_nov_97.791%
SP18 Triticum aestivum	SPN1324 Murimonas sp._strain_[WYJ26-X68]_nov_94.541%	SPN2089 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.082%
SP19 Streptococcus cristatus	SPN1334 Duncaniella freteri_nov_91.952%	SPN2101 Roseburia inulinivorans_nov_90.741%
SP20 Anaerobium sp._strain_[C-37]	SPN1335 Streptococcus troglodytidis_nov_87.209%	SPN2113 Anaerocolumna cellulolytica_nov_91.909%
SP204 Eubacteriales_[G-4] bacterium_MOT-164	SPN1346 Cuneatibacter caecimuris_nov_92.355%	SPN2124 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_94.468%
SP21 Veillonella parvula	SPN1358 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_93.139%	SPN2135 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_91.147%
SP22 Coprococcus sp._strain_[1XD21-23]	SPN1370 Glucerbacter canis_nov_93.542%	SPN2137 Eisenbergiella massiliensis_nov_85.744%
SP23 Prevotella jejuni	SPN1383 Magnetovibrio blakemorei_nov_83.259%	SPN2142 Centipedia periodontii_nov_95.850%
SP24 Oscillibacter sp._strain_[128x]	SPN139 Faecalicatena fissicatena_nov_95.228%	SPN2154 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_91.060%
SP26 Oscillospiraceae_[G-3] bacterium_MOT-150	SPN1395 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_92.340%	SPN2165 Lachnospiraceae_[G-1] bacterium_MOT-166_nov_97.930%
SP27 Thermus thermophilus	SPN1404 Streptococcus danieliae_nov_93.574%	SPN2178 Duncaniella freteri_nov_90.524%
SP29 Limosilactobacillus reuteri	SPN1407 Acetivibrio sp._strain_[D16-59]_nov_90.150%	SPN2190 Lachnospiraceae_[Clostridium]_herbivorans_nov_92.961%
SP30 Phyllobacterium myrsinacearum	SPN1418 Duncaniella freteri_nov_95.714%	SPN2201 Eubacteriales_[G-1] bacterium_MOT-158_nov_90.546%
SP31 Enterococcus faecalis	SPN1430 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.340%	SPN2214 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.881%
SP33 Cutibacterium acnes	SPN1442 Anaerotignum sp._strain_[D2-1X-45]_nov_96.205%	SPN2215 Porphyromonas pasteri_nov_97.967%
SP34 Streptococcus infantis	SPN1454 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.200%	SPN2226 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_92.608%
SP36 Eubacteriales_[G-1] bacterium_MOT-161	SPN1455 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.139%	SPN2237 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.684%
SP37 Streptococcus oralis	SPN1467 Duncaniella freteri_nov_87.149%	SPN2245 Acutalibacter muris_nov_96.099%
SP38 Lachnospiraceae_[G-11] bacterium_MOT-177	SPN1479 Lacrimispora xylandolytica_nov_93.827%	SPN2249 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_97.510%
SP39 Lachnospiraceae_[G-14] bacterium_MOT-184	SPN1492 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_95.885%	SPN2261 Coprococcus sp._strain_[1XD21-23]_nov_95.604%
SP4 Anaerotignum sp._strain_[D2-1X-72]	SPN1504 Eubacterium coprostanoligenes_nov_91.489%	SPN2273 Aliiococcus persicus_nov_79.245%
SP43 Ruminococcus sp._strain_[C-28]	SPN1516 Lachnospiraceae_[Clostridium]_polysaccharolyticum_nov_85.892%	SPN2286 Butyrivibrio sp._strain_[524-25]_nov_92.998%
SP44 Enterococcus gallinarum	SPN1524 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.625%	SPN2298 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.687%
SP49 Rothia dentocariosa	SPN1527 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.358%	SPN2309 Bacteroides thetaiotaomicron_nov_94.715%
SP5 Lautropia mirabilis	SPN1539 Eisenbergiella sp._strain_[1XD42-11]_nov_97.660%	SPN2321 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_94.672%
SP50 Mucispirillum schaedleri	SPN1550 Actinomycetes viscosus_nov_95.935%	SPN2334 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_95.949%
SP51 Eisenbergiella sp._strain_[1XD8-92]	SPN1561 Coprococcus sp._strain_[1XD21-23]_nov_94.493%	SPN2335 Adlercreutzia caecimuris_nov_89.270%
SP52 Methylobacterium populii	SPN1573 Streptococcus danieliae_nov_96.787%	SPN2366 Streptococcus danieliae_nov_93.988%
SP54 Lachnospiraceae_[G-13] bacterium_MOT-181	SPN1574 Blautia hominis_nov_89.583%	SPN2452 Duncaniella freteri_nov_90.982%
SP56 Veillonella dispar	SPN1585 Streptococcus rattii_nov_84.633%	SPN2485 Streptococcus parasanguinis_nov_97.791%
SP57 Anaerotignum sp._strain_[D16-54]	SPN1597 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.868%	SPN256 Alginomonas porphyrae_nov_79.724%
SP59 Streptococcus mitis	SPN1609 Adlercreutzia equolifaciens_nov_89.530%	SPN2569 Duncaniella freteri_nov_92.339%
SP6 Ligilactobacillus murinus	SPN1619 Abiotrophia defectiva_nov_92.885%	SPN2604 Lachnospiraceae_[G-1] bacterium_MOT-178_nov_95.547%
SP61 Streptococcus chosunense	SPN1621 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.812%	SPN2685 Duncaniella freteri_nov_89.780%
SP67 Streptococcus parasanguinis	SPN1632 Fusicatenibacter saccharivorans_nov_90.795%	SPN324 Alistipes senegalensis_nov_93.522%
SP69 Adlercreutzia caecimuris	SPN1642 Fusicatenibacter saccharivorans_nov_90.103%	SPN364 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_97.942%
SP7 Mordavella sp._strain_[D2-1X-71]	SPN1644 Streptococcus danieliae_nov_93.347%	SPN429 Streptococcus danieliae_nov_95.565%
SP73 Prevotella histicola	SPN1656 Acetivibrio sp._strain_[D16-59]_nov_90.792%	SPN438 Mediterraneibacter [Ruminococcus]_gnavus_nov_89.855%
SP75 Peptostreptococcus stomatis	SPN1668 Pseudoflavonifractor sp._strain_[136x]_nov_93.187%	SPN484 Anaerocolumna xylanovorans_nov_88.189%
SP76 Abiella sp._strain_[1XD42-72]	SPN1680 Streptococcus danieliae_nov_96.976%	SPN557 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.720%
SP78 Abiotrophia defectiva	SPN1691 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.911%	SPN601 Kineothrix sp._strain_[1XD8-87]_nov_97.859%
SP79 Faecalibaculum rodentium	SPN1692 Muricomes intestini_nov_89.648%	SPN676 Duncaniella freteri_nov_88.577%
SP8 Prevotella melaninogenica	SPN1703 Odoribacter sp._strain_[WYJ6-Z80]_nov_93.895%	SPN720 Adlercreutzia caecimuris_nov_89.722%
SP81 Lactobacillus intestinalis	SPN1714 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.264%	SPN794 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_93.008%
SP82 Fusobacterium nucleatum	SPN1726 Phocaea massiliensis_nov_88.675%	SPN840 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_94.456%
SP84 Neisseria sicca	SPN1740 Muribaculaceae_[G-2] bacterium_MOT-104_nov_87.873%	SPN87 Streptococcus danieliae_nov_97.379%
SP86 Mammallicoccus sciuri	SPN1752 Streptococcus danieliae_nov_93.968%	SPN88 Anaerocolumna cellulolytica_nov_86.071%
SP91 Actinomycetes naeslundii	SPN1760 Eubacterium ramulus_nov_89.256%	SPN915 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.111%
SP93 Alistipes sp._MOT-127	SPN1763 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.956%	SPN959 Pseudoflavonifractor sp._strain_[136x]_nov_94.066%
SP96 Mariprofundus fumaris	SPN1785 Mariprofundus fumaris_nov_82.820%	SPN964 Streptococcus mitis_pneumoniae_pneumoniae_pneumoniae