

Species

SP110 Turicimonas muris	SPN1144 Duncaniella freteri nov. 87.976%	SPN1905 Lachnospiraceae [G-11] 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 cellulositytica nov. 87.164%
SP119 Eubacteriales [G-1] bacterium MOT-159	SPN1180 Thalassobacillus devorans nov. 86.372%	SPN1948 Eubacterium coprostanoligenes nov. 91.026%
SP12 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-11] bacterium MOT-178 nov. 92.944%	SPN2020 Saccharopolyspora dendranthemae 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	SPN1282 Duncaniella freteri nov. 88.668%	SPN2044 Salimoesphilobacter 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-11] 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 tyroglycidis nov. 87.209%	SPN2113 Anaerocolumna cellulositytica 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 Centipeda periodontis nov. 95.850%
SP24 Oscillibacter sp. strain [128x]	SPN1393 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 xylanolytica 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 Alicococcus persicus nov. 79.245%
SP43 Ruminococcus sp. strain [C-28]	SPN1516 Lachnospiraceae [Clostridium] polysaccharolyticum nov. 85.892%	SPN2286 Butyricicoccus 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 Actinomyces 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 populi	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 rattus 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 Alginomas 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-11] 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 Fusicatenaibacter saccharivorans nov. 90.795%	SPN324 Alistipes senegalensis nov. 93.522%
SP69 Adlercreutzia caecimuris	SPN1642 Fusicatenaibacter 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 histiocola	SPN1656 Acetivibrio sp. strain [D16-59] nov. 90.792%	SPN438 Mediterraneibacter [Ruminococcus] gnavus nov. 89.855%
SP75 Peptostreptococcus stomatis	SPN1668 Pseudoflavonifactor sp. strain [136x] nov. 93.187%	SPN484 Anaerocolumna xylanovorans nov. 88.189%
SP76 Absiella 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 Phoecea 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 Mammaliococcus sciuri	SPN1752 Streptococcus danieliae nov. 93.968%	SPN88 Anaerocolumna cellulositytica nov. 86.071%
SP91 Actinomyces naeslundii	SPN1760 Eubacterium ramulus nov. 89.256%	SPN915 Muribaculaceae [G-1] bacterium MOT-129 nov. 85.111%