

SP100 Clostridium chromiireducens	SP83 Anaerotignum sp. strain [1XD42-85]	SPN538 Muricomes intestini nov. 89.648%
SP109 Anaerobium sp. strain [C-37]	SP84 Achromobacter pulmonis	SPN539 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 92.946%
SP11 Shigella sonnei	SP86 Acidipropionibacterium thoenii	SPN550 Streptococcus danieliae nov. 96.787%
SP111 Coprococcus sp. strain [1XD21-23]	SP89 Alistipes sp. MOT-127	SPN551 Duncaniella freteri nov. 87.976%
SP112 Pseudomonas aestus	SP9 Ligilactobacillus murinus	SPN562 Duncaniella freteri nov. 87.976%
SP116 Finegoldia magna	SP95 Erysipelatoclostridium [Clostridium] cocleatum	SPN574 Duncaniella freteri nov. 94.898%
SP118 Robinsoniella sp. strain [D2-1X-13]	SP96 Actinidia eriantha	SPN586 Lachnoclostridium [Clostridium] herbivorans nov. 92.562%
SP12 Enterococcus gallinarum	SP99 Enterobacter sp. MOT-050	SPN598 Lachnoclostridium [Clostridium] scindens nov. 89.300%
SP120 Ruminococcus sp. strain [C-28]	SPN1010 Duncaniella freteri nov. 89.463%	SPN610 Eubacterium coprostanoligenes nov. 91.489%
SP121 Duncaniella freteri	SPN1028 Streptococcus danieliae nov. 97.379%	SPN621 Lachnospiraceae [G-3] bacterium MOT-168 nov. 94.606%
SP127 Stenotrophomonas maltophilia	SPN1053 Actinomyces polynesiensis nov. 97.183%	SPN624 Lachnospiraceae [G-3] bacterium MOT-168 nov. 93.776%
SP129 Eubacteriales [G-4] bacterium MOT-164	SPN109 Alistipes senegalensis nov. 93.483%	SPN633 Phocaea massiliensis nov. 89.787%
SP131 Eubacteriales [G-2] bacterium MOT-162	SPN1093 Actinomyces polynesiensis nov. 94.153%	SPN643 Anaeromassilibacillus senegalensis nov. 93.008%
SP139 Paracoccus mangrovi	SPN1137 Streptococcus danieliae nov. 93.952%	SPN655 Anaerotignum sp. strain [D2-1X-45] nov. 96.205%
SP149 Adlercreutzia muris	SPN1167 Duncaniella freteri nov. 88.577%	SPN667 Duncaniella freteri nov. 91.952%
SP15 Escherichia coli	SPN121 Muribaculaceae [G-2] bacterium MOT-104 nov. 88.911%	SPN668 Eubacteriales [G-1] bacterium MOT-161 nov. 97.881%
SP16 Phyllobacterium myrsinacearum	SPN1211 Cutibacterium granulosum nov. 96.788%	SPN679 Lachnoclostridium [Clostridium] polysaccharolyticum nov. 90.496%
SP163 Oscillospiraceae [G-3] bacterium MOT-150	SPN1242 Lachnospiraceae [G-7] bacterium MOT-172 nov. 91.770%	SPN691 Adlercreutzia equolifaciens nov. 89.530%
SP169 Trichococcus pasteurii	SPN1283 Duncaniella freteri nov. 89.780%	SPN70 Preotella shahii nov. 87.903%
SP17 Enterococcus camelliae	SPN1323 Wolinella succinogenes nov. 80.044%	SPN703 Clostridium disporicum nov. 97.849%
SP18 Streptococcus sp. MOT-012	SPN133 Lachnospiraceae [G-6] bacterium MOT-171 nov. 94.179%	SPN715 Faecalicatena fissicatena nov. 94.824%
SP182 Streptococcus chosunense	SPN1396 Saliinibacillus xinjiangensis nov. 86.160%	SPN72 Streptococcus danieliae nov. 93.387%
SP185 Cutibacterium granulosum	SPN1437 Lachnospiraceae [G-12] bacterium MOT-180 nov. 89.676%	SPN727 Lachnospiraceae [G-14] bacterium MOT-185 nov. 92.537%
SP19 Escherichia fergusonii	SPN145 Duncaniella freteri nov. 95.510%	SPN738 Lachnospiraceae [G-6] bacterium MOT-171 nov. 95.218%
SP193 Lachnoclostridium sp. strain [1XD42-54]	SPN149 Streptococcus danieliae nov. 95.582%	SPN741 Muribaculaceae [G-2] bacterium MOT-104 nov. 89.264%
SP199 Mesorhizobium huakuii	SPN1509 Duncaniella freteri nov. 86.373%	SPN750 Cutibacterium acnes nov. 96.815%
SP2 Streptococcus danieliae	SPN1555 Alkalibacterium olivapoviliticus nov. 97.595%	SPN763 Lachnospiraceae [G-11] bacterium MOT-178 nov. 95.344%
SP202 Oscillibacter sp. strain [128x]	SPN157 Glucerberacter canis nov. 93.542%	SPN775 Fusicatenibacter saccharivorans nov. 88.912%
SP21 Clostridium disporicum	SPN160 Eisenbergiella massiliensis nov. 85.744%	SPN787 Muribaculaceae [G-2] bacterium MOT-104 nov. 89.703%
SP22 Lactobacillus johnsonii	SPN1628 Muribaculaceae [G-1] bacterium MOT-129 nov. 88.200%	SPN788 Adlercreutzia equolifaciens nov. 96.983%
SP222 Lachnospiraceae [G-1] bacterium MOT-166	SPN1674 Lacrimispora xylanolytica nov. 94.398%	SPN799 Lachnospiraceae [G-14] bacterium MOT-185 nov. 92.766%
SP223 Alkalibacterium pelagium	SPN169 Duncaniella freteri nov. 92.121%	SPN811 Lachnospiraceae [G-14] bacterium MOT-183 nov. 97.868%
SP24 Shigella boydii	SPN1744 Duncaniella freteri nov. 91.515%	SPN822 Adlercreutzia caecimuris nov. 89.270%
SP25 Lachnospiraceae [G-14] bacterium MOT-185	SPN1791 Lachnoclostridium [Clostridium] aminophilum nov. 86.983%	SPN834 Streptococcus danieliae nov. 96.774%
SP26 Lachnospiraceae [G-11] bacterium MOT-177	SPN181 Acetivibrio sp. strain [D16-59] nov. 90.812%	SPN84 Lachnospiraceae [G-14] bacterium MOT-185 nov. 95.949%
SP27 Anaerotignum sp. strain [D2-1X-72]	SPN1858 Duncaniella freteri nov. 86.922%	SPN845 Lachnospiraceae [G-14] bacterium MOT-183 nov. 94.670%
SP3 Faecalibaculum rodentium	SPN194 Muribaculaceae [G-1] bacterium MOT-129 nov. 85.972%	SPN855 Eubacterium coprostanoligenes nov. 90.213%
SP30 Limosilactobacillus reuteri	SPN195 Oscillospiraceae [G-6] bacterium MOT-153 nov. 91.684%	SPN857 Lachnospiraceae [G-11] bacterium MOT-177 nov. 95.679%
SP31 Enterococcus faecalis	SPN206 Duncaniella freteri nov. 90.982%	SPN868 Anaerocolumna cellulolytica nov. 91.909%
SP312 Tessaracoccus aquimaris	SPN218 Phocaeicola massiliensis nov. 93.509%	SPN891 Marinisporobacter balticus nov. 82.839%
SP4 Cutibacterium acnes	SPN229 Lachnospiraceae [G-14] bacterium MOT-185 nov. 95.745%	SPN902 Duncaniella freteri nov. 88.668%
SP40 Mammaliicoccus lentus	SPN241 Muribaculaceae [G-1] bacterium MOT-129 nov. 86.720%	SPN903 Murimonas sp. strain [WYJ26-X68] nov. 94.105%
SP41 Acutalibacter muris	SPN253 Bariatricus massiliensis nov. 89.506%	SPN913 Eisenbergiella sp. strain [1XD8-92] nov. 92.841%
SP42 Staphylococcus ureilyticus	SPN274 Lachnospiraceae [G-14] bacterium MOT-185 nov. 97.228%	SPN925 Anaerocolumna cellulolytica nov. 90.683%
SP43 Corynebacterium tuberculostearicum	SPN277 Eubacteriales [G-1] bacterium MOT-159 nov. 81.420%	SPN937 Beduini massiliensis nov. 86.708%
SP44 Listeria monocytogenes	SPN284 Pseudoflavonifractor sp. strain [136x] nov. 93.407%	SPN949 Phocaea massiliensis nov. 89.787%
SP46 Kineothrix sp. strain [1XD8-87]	SPN295 Muribaculaceae [G-2] bacterium MOT-104 nov. 87.873%	SPN96 Lachnospiraceae [G-14] bacterium MOT-185 nov. 97.015%
SP47 Enterococcus olivae	SPN308 Muribaculaceae [G-1] bacterium MOT-129 nov. 91.870%	SPN961 Pilibacter termitis nov. 90.581%
SP5 Mammaliicoccus sciuri	SPN309 Blautia hominis nov. 89.583%	SPN974 Duncaniella freteri nov. 93.750%
SP50 Actinomyces polynesiensis	SPN31 Anaerotignum sp. strain [1XD42-85] nov. 97.098%	SPN975 Lachnospiraceae [G-11] bacterium MOT-177 nov. 92.608%
SP53 Mucispirillum schaedleri	SPN321 Kineothrix alysooides nov. 89.855%	SPN986 Anaerocolumna xylanovorans nov. 88.212%
SP59 Eisenbergiella sp. strain [1XD8-92]	SPN332 Coprococcus sp. strain [1XD21-23] nov. 95.595%	SPN998 Lachnospiraceae [G-7] bacterium MOT-172 nov. 94.617%
SP6 Marvinbryantia sp. strain [D2-1X-79]	SPN344 Murimonas sp. strain [WYJ26-X68] nov. 93.873%	SPP109 Staphylococcus capitis epidermidis
SP60 Mordavella sp. strain [D2-1X-71]	SPN356 Duncaniella freteri nov. 87.298%	SPP14 Escherichia Shigella albertii sonnei
SP61 Streptococcus mitis	SPN368 Lachnospiraceae [G-6] bacterium MOT-171 nov. 92.931%	SPP18 Enterococcus casseliflavus gallinarum
SP62 Shigella flexneri	SPN380 Lachnospiraceae [G-7] bacterium MOT-172 nov. 93.320%	SPP22 Staphylococcus saprophyticus xylosus
SP63 Triticum aestivum	SPN383 Cellulomonas bogoriensis nov. 97.872%	SPP27 Agrobacterium Rhizobium lusitanum rhizogenes
SP64 Corynebacterium pilbarensis	SPN392 Anaerocolumna cellulolytica nov. 86.071%	SPP4 Streptococcus mitis pneumoniae pseudopneumoniae
SP67 Lactobacillus intestinalis	SPN404 Eisenbergiella sp. strain [1XD8-92] nov. 97.309%	SPP45 Lactobacillus johnsonii taiwanensis
SP69 Oscillospiraceae [G-7] bacterium MOT-154	SPN417 Eubacterium coprostanoligenes nov. 91.667%	SPP55 Achromobacter aegriificans insuavis
SP7 Salmonella enterica	SPN430 Turicibacter sanguinis nov. 95.949%	SPP59 Burkholderia Caenibaculum aenigmatica baiandianus cepacia contaminans diffu ...(8 sp
SP70 Romboutsia ilealis	SPN431 Alistipes senegalensis nov. 93.522%	SPP66 Burkholderia aenigmatica cepacia contaminans lata multivorans
SP71 Paracoccus laevigulosivorans	SPN442 Magnetovibrio blakemorei nov. 83.259%	SPP69 Burkholderia contaminans lata multivorans
SP72 Staphylococcus hominis	SPN454 Muribaculaceae [G-1] bacterium MOT-129 nov. 85.111%	SPP85 Bacillus halotolerans spizizenii
SP73 Corynebacterium mucifaciens	SPN466 Duncaniella freteri nov. 87.273%	SPP89 Staphylococcus argenteus aureus roterodami
SP74 Absiella sp. strain [1XD42-72]	SPN478 Fusicatenibacter saccharivorans nov. 90.586%	SPPN10 Faecalicatena multispecies sppn10_2 nov. 93.568%
SP75 Roseburia sp. strain [MGB-1]	SPN490 Streptococcus danieliae nov. 96.976%	SPPN31 Faecalicatena multispecies sppn31_2 nov. 93.802%