
Species
SP1 Streptococcus danieliae
SP10 Mesorhizobium huakuii
SP100 Stenotrophomonas maltophilia
SP101 Parasutterella excrementihominis
SP104 Mucispirillum schaedleri
SP107 Robinsoniella sp._strain_[D2−1X−13]
SP11 Enterococcus gallinarum;__
SP113 Lachnospiraceae_[G−14] bacterium_MOT−185
SP114 Staphylococcus hominis;__
SP117 Paracoccus laeviglucosivorans
SP119 Enterobacter sp._MOT−050
SP12 Faecalibaculum rodentium
SP121 Adlercreutzia muris
SP122 Corynebacterium pilbarense
SP123 Erysipelatoclostridium [Clostridium]_cocleatum
SP127 Acidipropionibacterium thoenii
SP13 Cutibacterium acnes
SP130 Coprococcus sp._strain_[1XD21−23]
SP139 Eubacteriales_[G−1] bacterium_MOT−159
SP140 Absiella sp._strain_[1XD42−72]
SP141 Anaerotignum sp._strain_[1XD42−85]
SP142 Clostridium chromiireducens
SP146 Roseburia sp._strain_[MGB−1]
SP15 Lactobacillus johnsonii;__
SP156 Oscillospiraceae_[G−3] bacterium_MOT−150
SP157 Shigella flexneri;__
SP16 Limosilactobacillus fermentum
SP167 Eubacteriales_[G−4] bacterium_MOT−164
SP17 Enterococcus faecalis
SP174 Salmonella enterica
SP178 Eubacteriales_[G−2] bacterium_MOT−162
SP18 Mordavella sp._strain_[D2−1X−71]
SP185 Enterococcus olivae
SP188 Lachnoclostridium sp._strain_[1XD42−54]
SP2 Streptococcus sp._MOT−012
SP22 Oscillospiraceae_[G−7] bacterium_MOT−154
SP23 Corynebacterium tuberculostearicum
SP235 Trichococcus pasteurii;__
SP26 Staphylococcus ureilyticus;__
SP3 Ligilactobacillus murinus
SP30 Peptoniphilus lacydonensis
SP31 Shigella boydii;__
SP32 Cutibacterium granulosum
SP33 Phyllobacterium myrsinacearum
SP36 Finegoldia magna
SP37 Escherichia coli;__
SP371 Tessaracoccus aquimaris
SP4 Enterococcus camelliae
SP40 Actinomyces polynesiensis
SP41 Escherichia fergusonii;__
SP46 Kineothrix sp._strain_[1XD8−87]
SP48 Oscillibacter sp._strain_[128x]
SP49 Alkalibacterium pelagium
SP5 Shigella sonnei;__
SP50 Akkermansia muciniphila
SP51 Duncaniella freteri
SP52 Clostridium disporicum;__
SP58 Limosilactobacillus reuteri
SP59 Lactobacillus intestinalis
SP60 Romboutsia ilealis;__
SP63 Anaerobium sp._strain_[C−37]
SP64 Alistipes sp._MOT−127
SP66 Streptococcus mitis;__
SP7 Mammaliicoccus sciuri
SP70 Anaerotignum sp._strain_[D2−1X−72]
SP72 Streptococcus chosunense
SP73 Corynebacterium mucifaciens
SP75 Paracoccus mangrovi
SP76 Marvinbryantia sp._strain_[D2−1X−79]
SP77 Lachnospiraceae_[G−11] bacterium_MOT−177
SP78 Tritium costivum
SP81 Lachnospiraceae_[G−14] bacterium_MOT−184
SP83 Actinidia eriantha
SP84 Achromobacter pulmonis;__
SP85 Listeria monocytogenes
SP89 Acutalibacter muris
SP90 Lachnospiraceae_[G−1] bacterium_MOT−166
SP91 Eisenbergiella sp._strain_[1XD8−92]
SP92 Ruminococcus sp._strain_[C−28]
SP94 Pseudomonas aestus
SP98 Mammaliicoccus lentus
SPN1009 Lachnospiraceae_[G−7] bacterium_MOT−172_nov_94.617%
SPN1020 Duncaniella freteri_nov_89.463%
SPN1037 Streptococcus danieliae_nov_97.379%
SPN1063 Actinomyces polynesiensis_nov_97.183%
SPN108 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_97.015%
SPN1101 Actinomyces polynesiensis_nov_94.153%
SPN1146 Streptococcus danieliae_nov_93.952%
SPN1181 Duncaniella freteri_nov_88.577%
SPN120 Alistipes senegalensis_nov_93.483%
SPN1220 Cutibacterium granulosum_nov_96.788%
SPN1253 Lachnospiraceae_[G−7] bacterium_MOT−172_nov_91.770%
SPN1292 Duncaniella freteri_nov_89.780%
SPN132 Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.911%
SPN1332 Wolinella succinogenes_nov_80.044%
SPN1406 Salinibacillus xinjiangensis_nov_86.160%
SPN144 Lachnospiraceae_[G−6] bacterium_MOT−171_nov_94.179%
SPN1450 Lachnospiraceae_[G−12] bacterium_MOT−180_nov_89.676%
SPN150 Streptococcus danieliae_nov_95.582%
SPN1521 Duncaniella freteri_nov_86.373%
SPN156 Duncaniella freteri_nov_95.510%
SPN1565 Alkalibacterium olivapovliticus_nov_97.595%
SPN161 Eisenbergiella massiliensis_nov_85.744%
SPN1638 Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.200%
SPN168 Gluceribacter canis_nov_93.542%
SPN1682 Lacrimispora xylanolytica_nov_94.398%
SPN1754 Duncaniella freteri_nov_91.515%
SPN1799 Lachnoclostridium [Clostridium]_aminophilum_nov_86.983%
SPN180 Duncaniella freteri_nov_92.121%
SPN1869 Duncaniella freteri_nov_86.922%
SPN192 Acetivibrio sp._strain_[D16−59]_nov_90.812%
SPN204 Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.972%
SPN206 Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.684%
SPN218 Duncaniella freteri_nov_90.982%
SPN229 Phocaeicola massiliensis_nov_93.509%
SPN240 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_95.745%
SPN252 Muribaculaceae_[G−1] bacterium_MOT−129_nov_86.720%
SPN264 Bariatricus massiliensis_nov_89.506%
SPN278 Eubacteriales_[G−1] bacterium_MOT−159_nov_81.420%
SPN285 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_97.228%
SPN29 Anaerotignum sp._strain_[1XD42−85]_nov_97.098%
SPN295 Pseudoflavonifractor sp._strain_[136x]_nov_93.407%
SPN306 Muribaculaceae_[G−2] bacterium_MOT−104_nov_87.873%
SPN318 Muribaculaceae_[G−1] bacterium_MOT−129_nov_91.870%
SPN321 Blautia hominis_nov_89.583%
SPN332 Kineothrix alysoides_nov_89.855%
SPN344 Coprococcus sp._strain_[1XD21−23]_nov_95.595%
SPN356 Murimonas sp._strain_[WYJ26−X68]_nov_93.873%
SPN368 Duncaniella freteri_nov_87.298%
SPN380 Lachnospiraceae_[G−6] bacterium_MOT−171_nov_92.931%
SPN392 Lachnospiraceae_[G−7] bacterium_MOT−172_nov_93.320%
SPN396 Cellulomonas bogoriensis_nov_97.872%
SPN404 Anaerocolumna cellulosilytica_nov_86.071%
SPN416 Eisenbergiella sp._strain_[1XD8−92]_nov_97.309%
SPN428 Eubacterium coprostanoligenes_nov_91.667%
SPN442 Turicibacter sanguinis_nov_95.949%
SPN443 Alistipes senegalensis_nov_93.522%
SPN454 Magnetovibrio blakemorei_nov_83.259%
SPN466 Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.111%
SPN478 Duncaniella freteri_nov_87.273%
SPN490 Fusicatenibacter saccharivorans_nov_90.586%
SPN503 Streptococcus danieliae_nov_96.976%
SPN507 Duncaniella freteri_nov_87.776%
SPN514 Duncaniella freteri_nov_87.149%
SPN526 Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.340%
SPN538 Anaerotaenia torta_nov_88.773%
SPN540 Muricomes intestini_nov_89.648%
SPN549 Lachnoclostridium [Clostridium]_polysaccharolyticum_nov_92.946%
SPN561 Streptococcus danieliae_nov_96.787%
SPN562 Duncaniella freteri_nov_87.976%
SPN573 Duncaniella freteri_nov_87.976%
SPN585 Duncaniella freteri_nov_94.898%
SPN597 Lachnoclostridium [Clostridium]_herbivorans_nov_92.562%
SPN609 Lachnoclostridium [Clostridium]_scindens_nov_89.300%
SPN621 Eubacterium coprostanoligenes_nov_91.489%
SPN625 Lachnospiraceae_[G−3] bacterium_MOT−168_nov_93.776%
SPN633 Lachnospiraceae_[G−3] bacterium_MOT−168_nov_94.606%
SPN644 Phocea massiliensis_nov_89.787%
SPN655 Anaeromassilibacillus senegalensis_nov_93.008%
SPN666 Anaerotignum sp._strain_[D2−1X−45]_nov_96.205%
SPN677 Duncaniella freteri_nov_91.952%
SPN679 Eubacteriales_[G−1] bacterium_MOT−161_nov_97.881%
SPN690 Lachnoclostridium [Clostridium]_polysaccharolyticum_nov_90.496%
SPN702 Adlercreutzia equolifaciens_nov_89.530%
SPN714 Faecalicatena fissicatena_nov_94.824%
SPN726 Clostridium disporicum_nov_97.849%
SPN738 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.537%
SPN743 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.264%
SPN749 Lachnospiraceae_[G−6] bacterium_MOT−171_nov_95.218%
SPN762 Cutibacterium acnes_nov_96.815%
SPN773 Lachnospiraceae_[G−11] bacterium_MOT−178_nov_95.344%
SPN785 Fusicatenibacter saccharivorans_nov_88.912%
SPN798 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.703%
SPN799 Adlercreutzia equolifaciens_nov_96.983%
SPN81 Prevotella shahii_nov_87.903%
SPN810 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.766%
SPN82 Streptococcus danieliae_nov_93.387%
SPN822 Lachnospiraceae_[G−14] bacterium_MOT−183_nov_97.868%
SPN832 Adlercreutzia caecimuris_nov_89.270%
SPN844 Streptococcus danieliae_nov_96.774%
SPN855 Lachnospiraceae_[G−14] bacterium_MOT−183_nov_94.670%
SPN859 Lachnospiraceae_[G−11] bacterium_MOT−177_nov_95.679%
SPN866 Eubacterium coprostanoligenes_nov_90.213%
SPN878 Anaerocolumna cellulosilytica_nov_91.909%
SPN902 Marinisporobacter balticus_nov_82.839%
SPN913 Duncaniella freteri_nov_88.668%
SPN914 Murimonas sp._strain_[WYJ26−X68]_nov_94.105%
SPN926 Eisenbergiella sp._strain_[1XD8−92]_nov_92.841%
SPN935 Anaerocolumna cellulosilytica_nov_90.683%
SPN947 Beduini massiliensis_nov_86.708%
SPN95 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_95.949%
SPN959 Phocea massiliensis_nov_89.787%
SPN972 Pilibacter termitis_nov_90.581%
SPN975 Lachnospiraceae_[G−11] bacterium_MOT−177_nov_92.608%
SPN984 Duncaniella freteri_nov_93.750%
SPN996 Anaerocolumna xylanovorans_nov_88.212%
SPP100 Streptococcus mitis_pneumoniae_pseudopneumoniae
SPP110 Bacillus halotolerans_spizizenii
SPP18 Staphylococcus argenteus_aureus_roterodami
SPP19 Staphylococcus capitis_epidermidis
SPP3 Enterococcus casseliflavus_gallinarum
SPP33 Lactobacillus johnsonii_taiwanensis
SPP42 Agrobacterium_Rhizobium lusitanum_rhizogenes
SPP54 Staphylococcus saprophyticus_xylosus
SPP65 Achromobacter aegrifaciens_insuavis
SPP74 Burkholderia contaminans_lata_multivorans
SPP78 Burkholderia_Caenibaculum aenigmatica_baiyandianus_cepacia_contaminans_diffu ...(8 species)
SPP79 Burkholderia aenigmatica_cepacia_contaminans_lata_multivorans
SPP97 Escherichia_Shigella albertii_sonnei
SPPN31 Faecalicatena multispecies_sppn31_2_nov_93.802%
SPPN9 Faecalicatena multispecies_sppn9_2_nov_93.568%