

Species

SP100 Acidovorax temperans
SP101 Methyloversatilis universalis
SP103 Granulicatella adiacens
SP104 Streptococcus agalactiae
SP107 Mammaliococcus sciuri
SP109 Faecalibaculum rodentium
SP110 Blautia luti
SP113 Parasutereilla excrementihominis
SP114 Streptococcus salivarius
SP117 Actinidia eriantha
SP118 Enterococcus avium
SP119 Staphylococcus rostri
SP12 Eubacteriales_[G-4] bacterium_MOT-164
SP128 Bacteroides intestinalis
SP130 Shigella flexneri
SP138 Pseudoxanthomonas mexicana
SP139 Fusicatenibacter saccharivorans
SP14 Akkermansia muciniphila
SP145 Eubacteriales_[G-1] bacterium_MOT-159
SP146 Bacteroides caccae
SP148 Parabacteroides goldsteinii
SP15 Herbaspirillum huttiense
SP151 Alistipes finegoldii
SP154 Blautia producta
SP16 Streptococcus chosunense
SP160 Anaerostipes caccae
SP161 Phascolarctobacterium faecium
SP166 Streptococcus sp._HMT_064
SP169 Microlunatus antarcticus
SP176 Acinetobacter bereziniae
SP179 Microlunatus spumicola
SP18 Ralstonia pickettii
SP183 Blautia glucerasea
SP189 Pinus sylvestris
SP195 Dermacoccus nishinomiyaensis
SP196 Bacteroides thetaiotaomicron
SP205 Sphingobium limneticum
SP207 Enterococcus faecalis
SP21 Phyllobacterium myrsinacearum
SP210 Muribaculum intestinale
SP219 Staphylococcus haemolyticus
SP22 Escherichia coli
SP222 Agrobacterium tumefaciens
SP224 Collinsella aerofaciens
SP23 Bifidobacterium pseudolongum
SP230 Aerococcus urinaequi
SP238 Caulobacter hibisci
SP24 Veillonella dispar
SP248 Corynebacterium amycolatum
SP25 Lactobacillus intestinalis
SP259 Prevotella sp._MOT-128
SP26 Shigella sonnei
SP266 Bacteroides cellulosilyticus
SP269 Bacteroides acidifaciens
SP285 Proteus mirabilis
SP299 Longicatena caecimuris
SP3 Corynebacterium mastitidis
SP31 Haemophilus parainfluenzae
SP333 Citrobacter koseri
SP345 Brachybacterium conglomeratum
SP36 Lactobacillus vaginalis
SP364 Alcaligenes faecalis
SP38 Adlercreutzia equolifaciens
SP396 Salmonella enterica
SP4 Stutzerimonas stutzeri
SP40 Veillonella parvula
SP41 Phocaecola vulgatus
SP43 Clostridium disporicum
SP439 Micrococcus antarcticus
SP44 Kocuria rhizophila
SP49 Enterococcus gallinarum
SP57 Brevundimonas nasdae
SP6 Frisingicoccus caecimuris
SP61 Bifidobacterium animalis
SP63 Ligilactobacillus murinus
SP64 Sphingomonas kyeonggiensis
SP65 Limosilactobacillus reuteri
SP69 Streptococcus mitis
SP7 Lachnospiraceae_[G-14] bacterium_MOT-185
SP76 Stenotrophomonas maltophilia
SP79 Eubacteriales_[G-2] bacterium_MOT-162
SP8 Acinetobacter johnsonii
SP80 Streptococcus lutetiensis
SP83 Staphylococcus urelyticus
SP86 Shigella dysenteriae
SP88 Enterococcus hirae
SP89 Streptococcus azizii
SP9 Clostridium saudiense
SP92 Streptococcus thoraltensis
SP95 Romboutsia ilealis
SP98 Bacteroides xyloisolvans
SPN1007 Coprococcus eutactus_nov_92.949%
SPN1019 Frisingicoccus caecimuris_nov_96.868%
SPN1030 Anaerobutyricum hallii_nov_93.895%
SPN1043 Roseburia faecis_nov_97.917%
SPN1054 Faecalicatena fissicatena_nov_96.042%
SPN1065 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.033%
SPN1077 Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.484%
SPN1079 Rothia nasimurium_nov_97.228%
SPN1121 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_95.717%
SPN1177 Blautia glucerasea_nov_96.429%
SPN1186 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.602%
SPN1219 Ruminococcus champanellensis_nov_91.810%
SPN1276 Duncaniella freteri_nov_91.700%
SPN1296 Eubacteriales_[G-4] bacterium_MOT-164_nov_94.055%
SPN1327 Cuneatibacter caecimuris_nov_93.528%
SPN1359 Duncaniella freteri_nov_89.024%
SPN1445 Roseburia inulinivorans_nov_90.249%
SPN1452 Anaerostipes hadrus_nov_92.961%
SPN1542 Butyricicoccus faecihominis_nov_90.909%
SPN1545 Turicibacter sanguinis_nov_95.923%
SPN158 Frisingicoccus caecimuris_nov_97.912%
SPN162 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_87.885%
SPN1627 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
SPN1661 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_92.704%
SPN1706 Prevotella sp._HMT_317_nov_90.244%
SPN1775 Paludicola psychrotolerans_nov_88.115%
SPN1776 Allobaculum stercoricanis_nov_92.739%
SPN1777 Coprococcus eutactus_nov_95.717%
SPN1839 Streptococcus plurinimalium_nov_94.280%
SPN1884 Sphingibacter marinus_nov_79.905%
SPN1900 Blautia faecicola_nov_93.111%
SPN214 Monoglobus pectinilyticus_nov_92.887%
SPN228 Duncaniella freteri_nov_90.631%
SPN24 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.662%
SPN240 Butyricicoccus pullicaecorum_nov_87.010%
SPN252 Blautia schinkii_nov_94.130%
SPN264 Streptococcus sp._MOT-045_nov_97.764%
SPN265 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.382%
SPN276 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
SPN282 Staphylococcus auricularis_nov_79.960%
SPN288 Eubacterium coprostanoligenes_nov_93.966%
SPN299 Faecalibaculum rodentium_nov_96.715%
SPN311 Duncaniella freteri_nov_86.061%
SPN323 Oscillibacter valericiogenes_nov_95.010%
SPN326 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.154%
SPN336 Ruthenibacterium lactatiformans_nov_89.669%
SPN349 Roseburia intestinalis_nov_95.218%
SPN361 Glucerbacter canis_nov_95.388%
SPN372 Mediterraneibacter [Ruminococcus] gnavus_nov_95.198%
SPN384 Eubacterium coprostanoligenes_nov_94.130%
SPN394 Eubacterium coprostanoligenes_nov_94.130%

SPN397 Zhongshania marina_nov_80.423%
SPN408 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%
SPN419 Eubacterium ramulus_nov_93.971%
SPN431 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_97.831%
SPN444 Eubacteriales_[G-1] bacterium_MOT-158_nov_91.507%
SPN454 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.363%
SPN465 Blautia glucerasea_nov_97.899%
SPN476 Microlunatus spumicola_nov_97.408%
SPN488 Ruminococcus albus_nov_87.474%
SPN499 Anaerocolumna cellulosilytica_nov_87.657%
SPN500 Dysosmobacter welbionis_nov_86.902%
SPN513 Saccharibacteria_(TM7)_[G-7] bacterium_HMT_954_nov_81.099%
SPN524 Eubacteriales_[G-4] bacterium_MOT-164_nov_94.268%
SPN535 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.275%
SPN54 Faecalibaculum rodentium_nov_85.455%
SPN548 Alistipes finegoldii_nov_94.835%
SPN559 Eubacteriales_[G-1] bacterium_MOT-144_nov_88.199%
SPN571 Agathobaculum desmolans_nov_86.345%
SPN593 Cuneatibacter caecimuris_nov_92.308%
SPN604 Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.230%
SPN616 Blautia luti_nov_95.798%
SPN617 Blautia hominis_nov_91.004%
SPN628 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_96.017%
SPN64 Ruminococcus albus_nov_91.045%
SPN640 Duncaniella freteri_nov_86.345%
SPN649 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_92.259%
SPN661 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_95.851%
SPN671 Breznakia pachnodae_nov_83.112%
SPN683 Caproiciproducens galactitolivorans_nov_91.083%
SPN695 Eubacteriales_[G-1] bacterium_MOT-158_nov_90.870%
SPN708 Lacrimispora saccharolytica_nov_90.625%
SPN720 Neglectibacter timonensis_nov_93.252%
SPN731 Fusicatenibacter saccharivorans_nov_97.689%
SPN732 Anaeromassilibacillus senegalensis_nov_93.843%
SPN744 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.347%
SPN756 Blautia producta_nov_97.077%
SPN768 Blautia faecicola_nov_93.319%
SPN779 Glucerbacter canis_nov_94.561%
SPN791 Blautia schinkii_nov_95.597%
SPN801 Oscillospiraceae_[G-7] bacterium_MOT-154_nov_94.828%
SPN812 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_97.275%
SPN824 Eubacteriales_[G-3] bacterium_MOT-163_nov_85.106%
SPN835 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_93.878%
SPN846 Parafannyhessea umbonata_nov_91.915%
SPN848 Eubacteriales_[G-1] bacterium_MOT-158_nov_97.660%
SPN859 Blautia schinkii_nov_95.789%
SPN871 Lacrimispora xylanolytica_nov_96.646%
SPN883 Dechloromonas agitata_nov_97.286%
SPN894 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_96.646%
SPN906 Ruminococcus albus_nov_92.225%
SPN91 Blautia luti_nov_97.899%
SPN918 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.398%
SPN929 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.531%
SPN939 Amaricoccus kaplicensis_nov_96.737%
SPN950 Eisenbergiella massiliensis_nov_86.486%
SPN960 Olsenella phocaeensis_nov_91.966%
SPN961 Duncaniella freteri_nov_85.371%
SPN972 Ruthenibacterium lactatiformans_nov_90.515%
SPN995 Bacteroides uniformis_nov_96.099%
SPP1 Escherichia_Shigella_fergusonii_flexneri
SPP11 Sphingomonas aquatilis_melonis
SPP17 Veillonella dispar_parvula
SPP2 Staphylococcus saprophyticus_xylosus
SPP20 Staphylococcus argenteus_aureus_roterodami
SPP26 Aeromonas caviae_enteropelogenes_hydrophila
SPP36 Streptococcus parasanguinis_parasanguinis_clade_721
SPP6 Blautia coccoides_producta
SPP7 Caulobacter segnis_vibrioides
SPPN12 Roseburia multispecies_sppn12_2_nov_92.133%
SPPN22 Blautia multispecies_sppn22_2_nov_96.226%