

Species

SP269 Oscillospira muciniphila
SP329 Akkermansia muciniphila
SP33 Nitrospira moscoviensis
SP34 Lactobacillus johnsonii
SP348 Devosia insulae
SP349 Lactobacillus intestinalis
SP35 Kineothrix alysoides
SP36 Thiobacillus thioparus
SP37 Eubacteriales_[G-4] bacterium_MOT-164
SP374 Ramlibacter monticola
SP375 Lawsonibacter asaccharolyticus
SP38 Bifidobacterium pseudolongum
SP391 Phocaeicola sartorii
SP393 Lachnospiraceae_[G-12] bacterium_MOT-179
SP421 Muribaculaceae_[G-2] bacterium_MOT-104
SP427 Alistipes putredinis
SP429 Lachnospiraceae_[G-3] bacterium_MOT-168
SP44 Lachnoclostridium [Clostridium] scindens
SP484 Acidovorax wautersii
SP488 Erysipelatoclostridium [Clostridium] innocuum
SP490 Ruthenibacterium lactatiformans
SP550 Erysipelatoclostridium [Clostridium] cocleatum
SP553 Carnobacteriaceae_[G-1] bacterium_MOT-198
SP558 Ligilactobacillus murinus
SP56 Adlercreutzia mucosicola
SP561 Bacteroides acidifaciens
SP569 Alistipes sp._MOT-127
SP581 Faecalicatena orotica
SP6 Paenibacillus acris
SP622 Anaerostipes caccae
SP659 Sphingomonas sediminicola
SP667 Cryobacterium psychrotolerans
SP67 Erysipelotrichaceae_[G-1] bacterium_MOT-189
SP68 Rhodoferrax saidenbachensis
SP691 Lachnospiraceae_[G-11] bacterium_MOT-177
SP693 Paenibacillus endophyticus
SP7 Nocardioides kribbensis
SP71 Pedobacter duraqueae
SP72 Oscillospiraecae_[G-7] bacterium_MOT-154
SP730 Eubacteriales_[G-2] bacterium_MOT-162
SP746 Lachnospiraceae_[G-11] bacterium_MOT-178
SP772 Oscillospiraecae_[G-3] bacterium_MOT-150
SP792 Ruminococcus bromii
SP812 Muribaculaceae_[G-1] bacterium_MOT-129
SP824 Lachnospiraceae_[G-14] bacterium_MOT-184
SP827 Corynebacterium mastitidis
SP85 Lachnospiraceae_[G-2] bacterium_MOT-167
SP86 Rodentibacter pneumotropicus
SP88 Lachnospiraceae_[G-14] bacterium_MOT-185
SP896 Desulfovibrio fairfieldensis
SP911 Oscillospiraecae_[G-6] bacterium_MOT-153
SP951 Herminiimonas contaminans
SP981 Lachnospiraceae_[G-11] bacterium_MOT-176
SPN1 Chondromyces lanuginosus_nov_93.636%
SPN1007 Alistipes senegalensis_nov_93.686%
SPN1008 Adlercreutzia equolifaciens_nov_92.029%
SPN1018 Adlercreutzia equolifaciens_nov_94.498%
SPN1030 Aminipla butyrlica_nov_88.544%
SPN1039 Adlercreutzia caecimuris_nov_94.156%
SPN1051 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.800%
SPN1061 Caproicibacter fermentans_nov_89.895%
SPN1072 Turicibacter sanguinis_nov_95.923%
SPN1073 Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.540%
SPN1085 Sulfuriflexus mobilis_nov_90.271%
SPN1097 Azorarcus olearius_nov_91.591%
SPN110 Paraeggerthella hongkongensis_nov_93.510%
SPN1118 Bacteroidetes_[G-3] bacterium_HMT_436_nov_86.585%
SPN1120 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.761%
SPN1127 Oscillospiraecae_[G-1] bacterium_MOT-147_nov_96.466%
SPN1139 Aggregatilinea lenta_nov_86.811%
SPN1453 Pseudomonas aeruginosa_DSM_21829_bacterium_HMT_001_nov_82.358%
SPN1453 Pseudomonas aeruginosa_DSM_21829_bacterium_HMT_001_nov_82.358%

SPN140 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_95.453%
SPN2151 Parabacteroides distansoni_nov_97.701%
SPN2163 Mucispirillum schaedleri_nov_96.833%
SPN2174 Aminipla butyrlica_nov_91.627%
SPN218 Odoribacter splanchnicus_nov_92.562%
SPN2187 Eubacteriales_[G-1] bacterium_MOT-158_nov_95.204%
SPN2207 Eubacterium ventriosum_nov_97.108%
SPN2212 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_87.381%
SPN2218 Holdemania massiliensis_nov_92.483%
SPN2230 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.414%
SPN2231 Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.724%
SPN2252 Desulfovibrio fairfieldensis_nov_89.379%
SPN2262 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.449%
SPN2275 Nitrospira multiformis_nov_93.864%
SPN228 Caproiciproducens galactitolivorans_nov_89.737%
SPN2287 Culturomica massiliensis_nov_91.055%
SPN2297 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.800%
SPN2308 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
SPN2326 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.874%
SPN2331 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.922%
SPN2337 Eubacteriales_[G-4] bacterium_MOT-165_nov_97.356%
SPN2343 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
SPN2354 Eubacteriales_[G-3] bacterium_MOT-163_nov_93.765%
SPN2365 Povalibacter uvarum_nov_90.909%
SPN2376 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.523%
SPN238 Eubacteriales_[G-3] bacterium_MOT-163_nov_81.065%
SPN2388 Kineothrix alysoides_nov_96.154%
SPN2400 Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
SPN2410 Ruminococcus albus_nov_92.530%
SPN2422 Duncaniella freteri_nov_87.398%
SPN2433 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_94.231%
SPN244 Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.578%
SPN2445 Duncaniella freteri_nov_90.612%
SPN2448 Amedibacillus dolichus_nov_92.290%
SPN2465 Ruminoclostridium cellulolyticum_nov_89.499%
SPN2474 Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.264%
SPN2478 Odoribacter splanchnicus_nov_93.807%
SPN248 Oscillospiraecae_[G-4] bacterium_MOT-151_nov_96.651%
SPN2486 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.548%
SPN2498 Bacteroides uniformis_nov_95.893%
SPN2509 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
SPN2511 Glucerbacter canis_nov_93.501%
SPN2521 Longibaculum muris_nov_92.308%
SPN2532 Enterocloster asparagiformis_nov_96.875%
SPN2543 Butyrivibrio pullicaeorum_nov_94.737%
SPN2555 Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.034%
SPN2556 Paraburkholderia hospita_nov_91.667%
SPN2566 Eubacteriales_[G-4] bacterium_MOT-165_nov_94.724%
SPN2577 Parasutterella excrementihominis_nov_94.578%
SPN2588 Kilonella majae_nov_88.836%
SPN259 Eubacteriales_[G-1] bacterium_MOT-158_nov_96.635%
SPN2594 Parabacteroides merdae_nov_93.265%
SPN2600 Holophaga foetida_nov_85.102%
SPN2610 Erysipelatoclostridium [Clostridium] saccharogumia_nov_93.213%
SPN2622 Anaerotruncus colihominis_nov_95.904%
SPN2633 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.414%
SPN2643 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_93.525%
SPN2655 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.925%
SPN2665 Alistipes putredinis_nov_97.471%
SPN2666 Odoribacter splanchnicus_nov_92.184%
SPN2676 Duncaniella freteri_nov_93.293%
SPN2688 Paludibaculum fermentans_nov_92.048%
SPN2694 Duncaniella freteri_nov_87.375%
SPN2698 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.449%
SPN2708 Kineothrix alysoides_nov_97.356%
SPN271 Dokdonella fugitiva_nov_97.955%
SPN2719 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_96.394%
SPN2731 Muricomes intestini_nov_97.831%
SPN2742 Lachnoclostridium [Clostridium] herbivorans_nov_92.917%
SPN2752 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.449%
SPN2772 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.339%
SPN2772 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.339%

SPN372 Pseudomonas aeruginosa_DSM_21829_bacterium_MOT-175_nov_89.474%
SPN573 Schlegella aquatica_nov_90.971%
SPN584 Papillibacter cinnamivorans_nov_93.795%
SPN596 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_92.089%
SPN60 Actinomarinicola tropica_nov_91.607%
SPN606 Harryflintia acetispora_nov_93.079%
SPN617 Anaerocolumna cellulolytica_nov_90.000%
SPN628 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.501%
SPN640 Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.333%
SPN649 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.208%
SPN660 Marvinbryantia formatexigens_nov_91.435%
SPN672 Povalibacter uvarum_nov_94.104%
SPN681 Alistipes putredinis_nov_96.782%
SPN683 Roseburia faecis_nov_94.231%
SPN696 Eisenbergiella massiliensis_nov_94.286%
SPN705 Muribaculum intestinale_nov_94.533%
SPN715 Piscinibacter defluvi_nov_97.500%
SPN72 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.188%
SPN726 Aggregatilinea lenta_nov_87.081%
SPN735 Adlercreutzia mucosicola_nov_97.826%
SPN746 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.608%
SPN747 Luteitalea pratensis_nov_92.135%
SPN757 Lacrimispora xylanolytica_nov_93.789%
SPN767 Hydrogenoanaerobacterium saccharovorans_nov_93.735%
SPN779 Paludibaculum fermentans_nov_91.388%
SPN789 Coprococcus catus_nov_95.192%
SPN791 Lacrimispora indolis_nov_90.625%
SPN800 Lachnoclostridium [Clostridium] populeti_nov_96.386%
SPN809 Adlercreutzia muris_nov_88.745%
SPN819 Pelotomaculum terephthalicum_nov_86.667%
SPN830 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_94.606%
SPN84 Butyrivibrio crossotus_nov_97.108%
SPN841 Vitreimonas flagellata_nov_95.465%
SPN852 Saccharibacteria (TM7)_[G-3] bacterium_HMT_351_nov_97.122%
SPN853 Prevotella sp._MOT-128_nov_91.954%
SPN864 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_91.213%
SPN876 Duncaniella freteri_nov_89.474%
SPN887 Adlercreutzia caecimuris_nov_92.291%
SPN899 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_94.820%
SPN900 Anaerotruncus lactatifermentans_nov_97.847%
SPN923 Culturomica massiliensis_nov_89.613%
SPN932 Dongia mobilis_nov_96.145%
SPN943 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_90.612%
SPN95 Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.284%
SPN953 Chryseolinea soli_nov_95.392%
SPN962 Muribaculum intestinale_nov_93.822%
SPN963 Arsenicibacter rosenii_nov_97.936%
SPN974 Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.223%
SPN98 Kineothrix alysoides_nov_97.837%
SPN985 Duncaniella freteri_nov_90.283%
SPN989 Muribaculaceae_[G-1] bacterium_MOT-129_nov_91.705%
SPN99 Anaeroplasmabactoclasticum_nov_92.874%
SPN997 Eubacteriales_[G-3] bacterium_MOT-163_nov_94.005%
SPP102 Staphylococcus capitis_caprae_epidermidis
SPP110 Afipia_Bradyrhizobium archetypum_austrialiense_broomeae_elkanii_embraopens...(18 species)
SPP115 Lachnospiraceae_[G-12] bacterium_MOT-179_bacterium_MOT-180
SPP12 Arthrobacter_Pseudarthrobacter humicola_oryzae_oxydans_pascens_penanthenrivorans...(7 species)
SPP122 Pasteurella_Rodentibacter caecimuris_hyllii
SPP147 Ligilactobacillus animalis_apodemi_murinus
SPP148 Bacteroides acidifaciens_acidofaciens
SPP159 Sphingomonas aquatilis_melonis
SPP164 Chryseobacterium_Epilithomonas ginsenosidimitans_zeae
SPP166 Blautia hansenii_hominis_marasmi
SPP178 Escherichia_Pseudoescherichia_Shigella coli_fergusonii_flexnerii_sonnei_vulneris
SPP188 Methylobacterium aminovorans_extorquens_podarium_populi_pseudosasa...(9 species)
SPP189 Pseudomonas arsenicocydans_mandeli_proseki_silesiensis
SPP195 Enterocloster boiteae_clostridioformis
SPP2 Hungateella effluvi_hathewayi
SPP209 Clostridium bowmanii_esthereticum
SPP23 Arthrobacter_Paeniglutamicibacter_Pseudarthrobacter cryotolerans_globiformis_psychrotolerans_scleromae...(5 species)
SPP23 Arthrobacter_Paeniglutamicibacter_Pseudarthrobacter cryotolerans_globiformis_psychrotolerans_scleromae...(5 species)
SPP23 Arthrobacter_Paeniglutamicibacter_Pseudarthrobacter cryotolerans_globiformis_psychrotolerans_scleromae...(5 species)