

Species

SP146 Staphylococcus similaris
SP151 Pseudonocardia profundimaris
SP155 Mammaliococcus sciuri
SP16 Blautia luti
SP160 Corynebacterium falsenii
SP161 Kocuria palustris
SP162 Phocaeicola vulgaris
SP165 Metasolibacillus fluoroglycofenilyticus
SP166 Psychrobacter faecalis
SP167 Bacteroides uniformis
SP17 Parasutterella excrementihominis
SP170 Mycolicibacterium vanbaalenii
SP172 Staphylococcus pasteurii
SP176 Longicatena caecimuris
SP177 Apotostipes sp._MOT-200
SP178 Sphingomonas fonticola
SP179 Methylobacterium platani
SP180 Ralstonia sp._HMT_406
SP189 Collinsella aerofaciens
SP19 Sphingomonas silvisoli
SP190 Sporosarcina siberiensis
SP191 Shigella flexneri
SP196 Roseburia hominis
SP197 Marmoricola pocheonensis
SP199 Akkermansia muciniphila
SP20 Bergeyella sp._HMT_422
SP200 Methylobacterium variabile
SP201 Micrococcus aloeverae
SP202 Bacteroides intestinalis
SP205 Corynebacterium pilbarensis
SP206 Rothia aerolata
SP207 Enterococcus faecalis
SP208 Odoribacter laneus
SP210 Microbacterium aurum
SP211 Enterobacter cancerogenus
SP217 Monoglobus pectinilyticus
SP22 Corynebacterium tuberculostearicum
SP226 Corynebacterium amycolatum
SP227 Parabacteroides distenii
SP231 Gordonia sediminis
SP235 Peribacillus frigoritolerans
SP236 Sphingomonas zeicaulis
SP238 Leptothrix sp._HMT_266
SP240 Erysipelatoclostridium [Clostridium] saccharogumia
SP241 Gordonia soli
SP247 Bacillus halotolerans
SP249 Cutibacterium acnes
SP258 Sphingomonas lutea
SP26 Bacteroides xylanisolvens
SP260 Clostridium tunisiense
SP261 Acinetobacter johnsonii
SP262 Kocuria sediminis
SP27 Enterococcus gallinarum
SP275 Acidovorax ebreus
SP28 Streptococcus mutans
SP281 Streptococcus sanguinis
SP291 Comamonas aquatilis
SP3 Lachnospiraceae_[G-3] bacterium_MOT-168
SP30 Haemophilus parainfluenzae
SP35 Escherichia coli
SP37 Romboutsia ilealis
SP38 Staphylococcus warneri
SP39 Eubacteriales_[G-1] bacterium_MOT-158
SP40 Romboutsia timonensis
SP41 Staphylococcus hominis
SP43 Citrobacter koseri
SP44 Bacteroides thetaiotaomicron
SP45 Mycolicibacterium italicum
SP51 Veillonella parvula
SP52 Blautia glucerasea
SP54 Streptococcus faecalis

SPN1102 Dysosmiodacter welchii_nov_00.902%
SPN1161 Sporobacter termitidis_nov_88.223%
SPN1167 Arthrospira platensis_nov_90.138%
SPN1171 Schlegella brevitalea_nov_97.510%
SPN1195 Arthrospira platensis_nov_87.963%
SPN13 Duncaniella freteri_nov_89.024%
SPN14 Glucibacter canis_nov_94.561%
SPN153 Blautia producta_nov_95.178%
SPN163 Spongiibacter marinus_nov_80.142%
SPN169 Craurococcus roseus_nov_95.602%
SPN178 Lactobacillus rogosae_nov_94.990%
SPN187 Kineothrix alysioides_nov_91.441%
SPN192 Coprococcus eutactus_nov_95.085%
SPN206 Microlunatus aurantiacus_nov_92.903%
SPN216 Lachnospirillum [Clostridium] herbivorans_nov_92.723%
SPN220 Lachnospirillum [Clostridium] herbivorans_nov_92.917%
SPN230 Arthrospira platensis_nov_88.046%
SPN235 Roseomonas aquatica_nov_97.912%
SPN24 Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%
SPN243 Agathobaculum desmolans_nov_86.345%
SPN251 Ruminococcus flavefaciens_nov_94.760%
SPN256 Anaerocolumna cellulolytica_nov_88.655%
SPN260 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_90.496%
SPN269 Turicibacter sanguinis_nov_96.137%
SPN277 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.347%
SPN284 Eubacterium coprostanoligenes_nov_91.379%
SPN292 Eubacterium ventriosum_nov_93.320%
SPN304 Longibaculum muris_nov_89.506%
SPN314 Eubacterium coprostanoligenes_nov_92.424%
SPN320 Corynebacterium mycetoides_nov_97.629%
SPN327 Pseudoflavonifractor phocaeensis_nov_91.511%
SPN334 Cuneatibacter caecimuris_nov_92.308%
SPN340 Eubacterium coprostanoligenes_nov_90.129%
SPN341 Rothia nasimurium_nov_97.228%
SPN345 Paludicola psychrotolerans_nov_88.115%
SPN367 Glucibacter canis_nov_95.388%
SPN384 Lachnospirillum [Clostridium] polysaccharolyticum_nov_90.437%
SPN40 Clostridium swelfunianum_nov_94.714%
SPN401 Eubacterium xylanophilum_nov_90.505%
SPN407 Microvirga subterranea_nov_95.150%
SPN408 Lawsonibacter asaccharolyticus_nov_90.909%
SPN415 Blautia faecicola_nov_93.111%
SPN416 Sporobacter termitidis_nov_87.124%
SPN418 Sporichthya polymorpha_nov_93.305%
SPN425 Raoultibacter timonensis_nov_90.476%
SPN440 Butyrivibrio pullicaecorum_nov_87.216%
SPN447 Glucibacter canis_nov_93.946%
SPN448 Dyadobacter sediminis_nov_97.240%
SPN449 Eubacteriales_[G-1] bacterium_MOT-158_nov_91.507%
SPN458 Arthrospira platensis_nov_88.249%
SPN464 Lawsonibacter asaccharolyticus_nov_90.083%
SPN474 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_92.975%
SPN483 Amaricoccus kaplicensis_nov_95.833%
SPN484 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.971%
SPN493 Arthrospira platensis_nov_89.425%
SPN499 Rubellimicrobium mesophilum_nov_95.592%
SPN51 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.555%
SPN513 Frisingicoccus caecimuris_nov_96.451%
SPN518 Lachnospira multipara_nov_84.298%
SPN520 Blautia producta_nov_89.834%
SPN537 Longibaculum muris_nov_92.754%
SPN54 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_87.885%
SPN543 Anaerostrophobacter polymeriproducens_nov_95.218%
SPN549 Coprococcus eutactus_nov_92.704%
SPN558 Oscillochloris trichoides_nov_85.202%
SPN563 Actinidia eriantha_nov_97.297%
SPN571 Oscillibacter valericigenes_nov_96.250%
SPN577 Eubacterium xylanophilum_nov_89.738%
SPN582 Eubacterium coprostanoligenes_nov_92.672%
SPN594 Nocardioides terrigena_nov_95.217%
SPN600 Blautia luti_nov_95.188%

SPN796 Eubacterium
SPN797 Anaerotruncus faecicola_nov_85.644%
SPN798 Butyrivibrio intestinalis_nov_93.111%
SPN799 Arthrospira platensis_nov_91.244%
SPN800 Ihubacter massiliensis_nov_94.792%
SPN801 Mediterraneibacter [Ruminococcus] torques_nov_92.693%
SPN802 Treuepera radiocitrix_nov_90.260%
SPN804 Lachnospiraceae_[G-8] bacterium_MOT-173_nov_89.167%
SPN805 Deinococcus papagoneensis_nov_89.083%
SPN806 Slackia piriformis_nov_85.129%
SPN807 Blautia luti_nov_96.429%
SPN808 Actinomycetozoa succinea_nov_97.881%
SPN809 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.058%
SPN810 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.946%
SPN811 Kallotenue papyrolyticum_nov_86.830%
SPN812 Blautia glucerasea_nov_94.142%
SPN813 Craurococcus roseus_nov_94.064%
SPN815 Propionibacteriaceae_[G-2] bacterium_HMT_192_nov_95.238%
SPN816 Pseudoflavonifractor phocaeensis_nov_91.097%
SPN817 Arthrospira platensis_nov_90.805%
SPN818 Blautia luti_nov_97.059%
SPN819 Kallotenue papyrolyticum_nov_85.177%
SPN820 Sphingomonas carri_nov_96.767%
SPN821 Mediterraneibacter [Ruminococcus] gnavus_nov_90.377%
SPN822 Zhongshania marina_nov_79.765%
SPN823 Acutalibacter muris_nov_95.670%
SPN824 Eubacteriales_[G-1] bacterium_MOT-159_nov_92.797%
SPN825 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_93.737%
SPN826 Spirosoma lacussanchae_nov_91.614%
SPN827 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.228%
SPN828 Lachnospiraceae_[G-8] bacterium_MOT-173_nov_89.189%
SPN829 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.258%
SPN830 Lihuaxuella thermophila_nov_97.419%
SPN831 Anaerotruncus aminivorans_nov_92.812%
SPN832 Niveispirillum fermenti_nov_85.242%
SPN833 Treuepera radiocitrix_nov_89.325%
SPN834 Phocaea massiliensis_nov_88.602%
SPN836 Neglectibacter timonensis_nov_92.449%
SPN837 Serratia rubidaea_nov_82.306%
SPN838 Arthrospira platensis_nov_88.991%
SPN839 Fusicatenibacter saccharivorans_nov_97.689%
SPN84 Sphingomonas suadeae_nov_96.970%
SPN840 Blautia luti_nov_93.697%
SPN841 Oribacterium parvum_nov_89.770%
SPN842 Paludicola psychrotolerans_nov_87.037%
SPN843 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.417%
SPN844 Caldovatus sediminis_nov_92.414%
SPN845 Alistipes finegoldii_nov_94.835%
SPN847 Arthrospira platensis_nov_90.345%
SPN848 Drancourtella massiliensis_nov_92.276%
SPN849 Faecalimonas umbilicata_nov_96.242%
SPN850 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.549%
SPN851 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_96.042%
SPN852 Kallotenue papyrolyticum_nov_84.956%
SPN861 Lachnospiraceae_[G-4] bacterium_MOT-169_nov_97.484%
SPN867 Blautia luti_nov_97.899%
SPN869 Arthrospira platensis_nov_88.736%
SPN879 Chryseobacterium culicis_nov_97.280%
SPN880 Blautia producta_nov_96.226%
SPN886 Lacrimispora saccharolytica_nov_90.625%
SPN895 Arboricoccus pini_nov_88.276%
SPN905 Gordonia soli_nov_97.198%
SPN926 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_90.760%
SPN938 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_93.595%
SPN948 Eubacterium xylanophilum_nov_91.313%
SPN954 Eisenbergiella massiliensis_nov_86.486%
SPN96 Oscillospiraceae_[G-7] bacterium_MOT-154_nov_92.918%
SPN960 Arthrospira platensis_nov_86.605%
SPN985 Arthrospira platensis_nov_88.073%
SPN997 Roseiflexus castenholzii_nov_84.876%
SPN10 Streptococcus faecalis