

Species

SP155 Mammaliacoccus sciuri	SPN1167 Arthrospira platensis_nov_90.138%	SPN798 Butyrivacter intestini_nov_93.111%
SP16 Blautia luti	SPN1171 Schlegelella brevitalea_nov_97.510%	SPN799 Arthrospira platensis_nov_91.244%
SP160 Corynebacterium falsenii	SPN1195 Arthrospira platensis_nov_87.963%	SPN800 Ihubacter massiliensis_nov_94.792%
SP161 Kocuria palustris	SPN13 Duncaniella freteri_nov_89.024%	SPN801 Mediterraneibacter [Ruminococcus] torques_nov_92.693%
SP162 Phocaeicola vulgatus	SPN14 Glucerbacter canis_nov_94.561%	SPN802 Truepera radiovictrix_nov_90.260%
SP165 Metasolibacillus fluoroglycofenilyticus	SPN153 Blautia producta_nov_95.178%	SPN804 Lachnospiraceae [G-8] bacterium_MOT-173_nov_89.167%
SP166 Psychrobacter faecalis	SPN163 Spongibacter marinus_nov_80.142%	SPN805 Deinococcus papagonensis_nov_89.083%
SP167 Bacteroides uniformis	SPN169 Craurococcus roseus_nov_95.602%	SPN806 Slackia piriformis_nov_85.129%
SP17 Parasutterella excrementihominis	SPN178 Lactobacillus rogosae_nov_94.990%	SPN807 Blautia luti_nov_96.429%
SP170 Mycolicibacterium vanbaalenii	SPN187 Kineothrix alysoides_nov_91.441%	SPN808 Actinomycetospora succinea_nov_97.881%
SP172 Staphylococcus pasteurii	SPN192 Coprococcus eutactus_nov_95.085%	SPN809 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.058%
SP176 Longicatena caecimuris	SPN206 Microcylindrus aurantiacus_nov_92.903%	SPN810 Oscillospiraceae [G-4] bacterium_MOT-151_nov_92.946%
SP177 Atopostipes sp._MOT-200	SPN216 Lachnoclostridium [Clostridium] herbivorans_nov_92.723%	SPN811 Kallotenue papyrolyticum_nov_86.830%
SP178 Sphingomonas fonticola	SPN220 Lachnoclostridium [Clostridium] herbivorans_nov_92.917%	SPN812 Blautia glucerata_nov_94.142%
SP179 Methylobacterium platani	SPN230 Arthrospira platensis_nov_88.046%	SPN813 Craurococcus roseus_nov_94.064%
SP180 Ralstonia sp._HMT_406	SPN235 Roseomonas aquatica_nov_97.912%	SPN815 Propionibacteriaceae [G-2] bacterium_HMT_192_nov_95.238%
SP189 Collinsella aerofaciens	SPN24 Erysipelatoclostridium [Clostridium] saccharogumia_nov_96.042%	SPN816 Pseudoflavonifractor phocaensis_nov_91.097%
SP19 Sphingomonas silvisoli	SPN243 Agathobaculum desmolans_nov_86.345%	SPN817 Arthrospira platensis_nov_90.805%
SP190 Sporosarcina sibiriensis	SPN251 Ruminococcus flavefaciens_nov_94.760%	SPN818 Blautia luti_nov_97.059%
SP191 Shigella flexneri	SPN256 Anaerocolumna cellulolytica_nov_88.655%	SPN819 Kallotenue papyrolyticum_nov_85.177%
SP196 Roseburia hominis	SPN260 Oscillospiraceae [G-4] bacterium_MOT-151_nov_90.496%	SPN820 Sphingomonas carri_nov_96.767%
SP197 Marmoricola pocheonensis	SPN269 Turicibacter sanguinis_nov_96.137%	SPN821 Mediterraneibacter [Ruminococcus] gnavus_nov_90.377%
SP199 Akkermansia muciniphila	SPN277 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.347%	SPN822 Zhongshania marina_nov_79.765%
SP20 Bergeyella sp._HMT_422	SPN284 Eubacterium coprostanoligenes_nov_91.379%	SPN823 Acetivibrio muris_nov_95.670%
SP200 Methylobacterium variabile	SPN292 Eubacterium ventriosum_nov_93.320%	SPN824 Eubacteriales [G-1] bacterium_MOT-159_nov_92.797%
SP201 Micrococcus aloweverae	SPN304 Longibaculum muris_nov_89.506%	SPN825 Lachnospiraceae [G-3] bacterium_MOT-168_nov_93.737%
SP202 Bacteroides intestinalis	SPN314 Eubacterium coprostanoligenes_nov_92.424%	SPN826 Spirosoma lacussanchae_nov_91.614%
SP205 Corynebacterium pilbarensis	SPN320 Corynebacterium mycetoides_nov_97.629%	SPN827 Muribaculaceae [G-2] bacterium_MOT-104_nov_86.228%
SP206 Rothia aerolata	SPN327 Pseudoflavonifractor phocaensis_nov_91.511%	SPN828 Lachnospiraceae [G-8] bacterium_MOT-173_nov_89.189%
SP207 Enterococcus faecalis	SPN334 Cuneatibacter caecimuris_nov_92.308%	SPN829 Oscillospiraceae [G-2] bacterium_MOT-149_nov_96.258%
SP208 Odoribacter laneus	SPN340 Eubacterium coprostanoligenes_nov_90.129%	SPN830 Lihuxuella thermophila_nov_97.419%
SP210 Microbacterium aurum	SPN341 Rothia nasimurium_nov_97.228%	SPN831 Anaerostipes aminivorans_nov_92.812%
SP211 Enterobacter cancerogenus	SPN345 Paludicola psychrotolerans_nov_88.115%	SPN832 Niveispirillum fermenti_nov_85.242%
SP217 Monoglobus pectinilyticus	SPN367 Glucerbacter canis_nov_95.388%	SPN833 Truepera radiovictrix_nov_89.325%
SP22 Corynebacterium tuberculoostearicum	SPN384 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.437%	SPN834 Phocaea massiliensis_nov_88.602%
SP226 Corynebacterium amycolatum	SPN40 Clostridium swelfunianum_nov_94.714%	SPN836 Neglectibacter timonensis_nov_92.449%
SP227 Parabacteroides goldsteinii	SPN401 Eubacterium xylanophilum_nov_90.505%	SPN837 Serratia rubidaea_nov_82.306%
SP231 Gordonia sediminis	SPN407 Microvirga subterranea_nov_95.150%	SPN838 Arthrospira platensis_nov_88.991%
SP235 Peribacillus frigoritolerans	SPN408 Lawsonibacter asaccharolyticus_nov_90.909%	SPN839 Fusicatenibacter saccharivorans_nov_97.689%
SP236 Sphingomonas zeicola	SPN415 Blautia faecicola_nov_93.111%	SPN84 Sphingomonas suadeae_nov_96.970%
SP238 Leptothrix sp._HMT_266	SPN416 Sporobacter termitidis_nov_87.124%	SPN840 Blautia luti_nov_93.697%
SP240 Erysipelatoclostridium [Clostridium] saccharogumia	SPN418 Sporichthya polymorpha_nov_93.305%	SPN841 Oribacterium parvum_nov_89.770%
SP241 Gordonia soli	SPN425 Raoultibacter timonensis_nov_90.476%	SPN842 Paludicola psychrotolerans_nov_87.037%
SP247 Bacillus halotolerans	SPN440 Butyrivibrio pullicaecorum_nov_87.216%	SPN843 Oscillospiraceae [G-4] bacterium_MOT-151_nov_95.417%
SP249 Cutibacterium acnes	SPN447 Glucerbacter canis_nov_93.946%	SPN844 Caldivatus sediminis_nov_92.414%
SP258 Sphingomonas lutea	SPN448 Dyadobacter sediminis_nov_97.240%	SPN845 Alistipes finegoldii_nov_94.835%
SP26 Bacteroides xylanisolvens	SPN449 Eubacteriales [G-1] bacterium_MOT-158_nov_91.507%	SPN847 Arthrospira platensis_nov_90.345%
SP260 Clostridium tunisiense	SPN458 Arthrospira platensis_nov_88.249%	SPN848 Drancourtella massiliensis_nov_92.276%
SP261 Acinetobacter johnsonii	SPN464 Lawsonibacter asaccharolyticus_nov_90.083%	SPN849 Faecalimonas umbilicata_nov_96.242%
SP262 Kocuria sediminis	SPN474 Oscillospiraceae [G-1] bacterium_MOT-147_nov_92.975%	SPN850 Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.549%
SP27 Enterococcus gallinarum	SPN483 Amaricoccus kaplicensis_nov_95.833%	SPN851 Oscillospiraceae [G-4] bacterium_MOT-151_nov_96.042%
SP275 Acidovorax ebreus	SPN484 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.971%	SPN852 Kallotenue papyrolyticum_nov_84.956%
SP28 Streptococcus mutans	SPN493 Arthrospira platensis_nov_89.425%	SPN861 Lachnospiraceae [G-4] bacterium_MOT-169_nov_97.484%
SP281 Streptococcus sanguinis	SPN499 Rubellimicrobium mesophilum_nov_95.592%	SPN867 Blautia luti_nov_97.899%
SP291 Comamonas aquatilis	SPN51 Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.555%	SPN869 Arthrospira platensis_nov_88.736%
SP3 Lachnospiraceae [G-3] bacterium_MOT-168	SPN513 Frisingicoccus caecimuris_nov_96.451%	SPN879 Chryseobacterium culicis_nov_97.280%
SP30 Haemophilus parainfluenzae	SPN518 Lachnospira multipara_nov_84.298%	SPN880 Blautia producta_nov_96.226%
SP35 Escherichia coli	SPN520 Blautia producta_nov_89.834%	SPN886 Lacrimispora saccharolytica_nov_90.625%
SP37 Romboutsia ilealis	SPN537 Longibaculum muris_nov_92.754%	SPN895 Arboricoccus pini_nov_88.276%
SP38 Staphylococcus warneri	SPN54 Oscillospiraceae [G-2] bacterium_MOT-149_nov_87.885%	SPN905 Gordonia soli_nov_97.198%
SP39 Eubacteriales [G-1] bacterium_MOT-158	SPN543 Anaerobacillus xylanophilus polymeriproducens_nov_95.218%	SPN926 Oscillospiraceae [G-1] bacterium_MOT-147_nov_90.760%
SP40 Romboutsia timonensis	SPN549 Coprococcus eutactus_nov_92.704%	SPN938 Oscillospiraceae [G-1] bacterium_MOT-147_nov_93.595%
SP41 Staphylococcus hominis	SPN558 Oscillochloris trichoides_nov_85.202%	SPN948 Eubacterium xylanophilum_nov_91.313%
SP43 Citrobacter koseri	SPN563 Actinidia eriantha_nov_97.297%	SPN954 Eisenbergiella massiliensis_nov_86.486%
SP44 Bacteroides thetaiotaomicron	SPN571 Oscillibacter valerigenes_nov_96.250%	SPN96 Oscillospiraceae [G-7] bacterium_MOT-154_nov_92.918%
SP45 Mycolicibacterium iranicum	SPN577 Eubacterium xylanophilum_nov_89.738%	SPN960 Arthrospira platensis_nov_86.605%
SP51 Veillonella parvula	SPN582 Eubacterium coprostanoligenes_nov_92.672%	SPN985 Arthrospira platensis_nov_88.073%