

Species

SP105 Pseudomonas putida	SP42 Phocaeicola sarotaii	SPN467 Leptothrix holstadii nov_96.970%
SP111 Streptococcus acidominimus	SP5 Faecalibaculum rodentium	SPN47 Muricomes intestini nov_89.583%
SP112 Comamonas testosteroni	SP50 Streptococcus gordonii	SPN476 Duncaniella freteri nov_92.653%
SP116 Acidovorax monticola	SP51 Streptococcus thoraltensis	SPN479 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.000%
SP117 Streptococcus mutans	SP53 Parabacteroides goldsteini	SPN48 Duncaniella freteri nov_87.221%
SP119 Citrobacter koseri	SP56 Peptostreptococcaceae_[X1][G-4] bacterium_HMT_369	SPN483 Alloprevotella sp._HMT_473_nov_89.634%
SP120 Enterobacter hormaechei	SP58 Eubacteriales_[G-4] bacterium_MOT-164	SPN489 Lawsonibacter asaccharolyticus nov_91.116%
SP121 Mammaliococcus sciuri	SP6 Ligilactobacillus murinus	SPN49 Lawsonibacter asaccharolyticus nov_90.329%
SP125 Mammaliococcus lentus	SP61 Cutibacterium acnes	SPN492 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.373%
SP129 Bacteroides acidifaciens	SP64 Methylobacterium brachiatum	SPN495 Hathewayia proteolytica nov_83.297%
SP13 Lachnospiraceae_[G-14] bacterium_MOT-185	SP69 Sediminibacterium aquarii	SPN5 Beduini massiliensis nov_87.705%
SP136 Escherichia fergusonii	SP70 Cetobacterium somerae	SPN50 Duncaniella freteri nov_87.400%
SP138 Actinomyces sp._HMT_448	SP71 Klebsiella aerogenes	SPN507 Duncaniella freteri nov_91.853%
SP139 Corynebacterium matruchotii	SP73 Robinsoniella peoriensis	SPN51 Magnetovibrio blakemorei nov_83.371%
SP14 Neisseriaceae_[G-1] bacterium_MOT-031	SP76 Muribaculum intestinale	SPN511 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
SP141 Fusobacterium nucleatum_subsp._vincentii	SP77 Lachnospiraceae_[G-13] bacterium_MOT-181	SPN512 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.621%
SP145 Streptococcus sp._MOT-012	SP8 Priestia aryabhatai	SPN514 Parasutterella excrementihominis nov_97.972%
SP146 Fusobacterium nucleatum	SP84 Sphingomonas yabuuchiae	SPN52 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.792%
SP149 Porphyromonas endodontalis	SP86 Acinetobacter radioresistens	SPN520 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
SP15 Erysipelotrichaceae_[G-1] bacterium_MOT-189	SP89 Acidovorax temperans	SPN521 Acetivibrio cellulolyticus nov_83.405%
SP155 Methylobacterium goeingense	SP9 Rhodococcus qingshengii	SPN53 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.347%
SP157 Magnetospirillum magnetotacticum	SP93 Streptococcus oralis	SPN531 Alistipes putredinis nov_94.444%
SP158 Parabacteroides distansoni	SP96 Enterococcus gallinarum	SPN533 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_92.902%
SP16 Klebsiella pneumoniae	SP97 Corynebacterium mastitidis	SPN536 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_93.096%
SP161 Proteus mirabilis	SP98 Parasutterella excrementihominis	SPN539 Thermodesulfobium acidiphilum nov_80.255%
SP164 Lysinibacillus sphaericus	SP99 Priestia megaterium	SPN542 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.549%
SP168 Enterobacter mori	SPN11 Erysipelatoclostridium_[Clostridium] innocuum nov_88.270%	SPN548 Duncaniella freteri nov_88.330%
SP17 Bacillus subtilis	SPN13 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.600%	SPN55 Muribaculaceae_[G-2] bacterium_MOT-104_nov_87.174%
SP170 Lachnospiraceae_[G-11] bacterium_MOT-177	SPN153 Duncaniella freteri nov_87.169%	SPN56 Prevotella sp._HMT_317_nov_90.244%
SP173 Hungatella hathewayi	SPN155 Streptococcus azizii nov_95.171%	SPN57 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.800%
SP18 Muribaculaceae_[G-2] bacterium_MOT-104	SPN16 Anaeroplasmata abactoclasticum nov_86.538%	SPN58 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.279%
SP19 Microbacterium maritropicum	SPN18 Ralstonia solanacearum nov_96.296%	SPN59 Duncaniella freteri nov_89.697%
SP197 Enterobacter asburiae	SPN183 Chryseobacterium yooncheonense nov_97.484%	SPN60 Neglectibacter timonensis nov_95.325%
SP199 Sphingobacterium multivorum	SPN2 Duncaniella freteri nov_87.424%	SPN61 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.319%
SP20 Kosakonia sacchari	SPN20 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.718%	SPN62 Acinetobacter johnsonii nov_97.737%
SP203 Citrobacter amalonaticus	SPN205 Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.686%	SPN63 Parafannyhessea umbonata nov_92.161%
SP204 Actinidia eriantha	SPN215 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%	SPN64 Lacrimispora xylanolytica nov_94.363%
SP206 Lachnospiraceae_[G-1] bacterium_MOT-166	SPN222 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.577%	SPN66 Blautia luti nov_94.561%
SP208 Agrobacterium vitis	SPN237 Duncaniella freteri nov_90.612%	SPN67 Acetivibrio cellulolyticus nov_82.289%
SP209 Corynebacterium macginleyi	SPN25 Longibaculum muris nov_86.957%	SPN68 Phocaea massiliensis nov_86.966%
SP22 Leucobacter chromiirensis	SPN256 Duncaniella freteri nov_90.184%	SPN69 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.340%
SP221 Massilia pinisoli	SPN272 Duncaniella freteri nov_93.712%	SPN70 Sediminibacterium aquarii nov_92.355%
SP232 Acidovorax ebreus	SPN279 Alistipes senegalensis nov_93.648%	SPN71 Duncaniella freteri nov_89.919%
SP233 Klebsiella oxytoca	SPN282 Duncaniella freteri nov_87.475%	SPN72 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.561%
SP235 Clostridium disporicum	SPN29 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.822%	SPN73 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_97.495%
SP237 Shigella flexneri	SPN30 Parabacteroides distansoni nov_97.938%	SPN74 Flavonifractor plautii nov_94.410%
SP239 Streptococcus oralis_subsp._tigurinus_clade_071	SPN306 Duncaniella freteri nov_93.699%	SPN76 Longibaculum muris nov_90.289%
SP244 Ileibacterium valens	SPN31 Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%	SPN77 Sporobacter termitidis nov_87.580%
SP249 Lactobacillus johnsonii	SPN323 Muribaculaceae_[G-1] bacterium_MOT-129_nov_84.929%	SPN78 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.561%
SP251 Epilithonimonas hominis	SPN33 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.290%	SPN79 Adlercreutzia muris nov_88.961%
SP253 Muribaculaceae_[G-1] bacterium_MOT-129	SPN34 Blautia schinkii nov_93.711%	SPN80 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.931%
SP255 Acutalibacter muris	SPN35 Actinidia eriantha nov_97.011%	SPN81 Pyrinomonas methylaliphatogenes nov_96.809%
SP256 Aggregatibacter sp._HMT_512	SPN353 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%	SPN82 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.305%
SP257 Stenotrophomonas maltophilia	SPN36 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%	SPN83 Duncaniella freteri nov_89.135%
SP266 Alistipes sp._MOT-127	SPN361 Lysinibacillus sphaericus nov_97.988%	SPN84 Eubacterium xylanophilum nov_91.075%
SP269 Staphylococcus caprae	SPN37 Eubacterium xylanophilum nov_89.940%	SPN85 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_93.800%
SP271 Lactobacillus intestinalis	SPN370 Duncaniella freteri nov_86.290%	SPN87 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.634%
SP273 Dialister invisus	SPN379 Prevotella shahii nov_87.602%	SPN88 Lawsonibacter asaccharolyticus nov_90.722%
SP275 Chryseobacterium gambrini	SPN38 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.511%	SPN89 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_97.694%
SP276 Bifidobacterium pseudolongum	SPN39 Faecalicatena orotica nov_92.484%	SPN90 Hydrogenoanaerobacterium saccharovorans nov_88.773%
SP285 Bacteroides thetaiotaomicron	SPN398 Muribaculaceae_[G-1] bacterium_MOT-129_nov_83.636%	SPP1 Staphylococcus saprophyticus_xylusos
SP286 Mucispirillum schaedleri	SPN40 Desulfovibrio fairfieldensis nov_96.349%	SPP10 Pseudomonas cedrina_bactis
SP289 Oscillospiraceae_[G-3] bacterium_MOT-150	SPN401 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.576%	SPP11 Pasteurella_Rodentibacter caecimuris_heylii
SP3 Staphylococcus hominis	SPN408 Glucibacter canis nov_93.305%	SPP13 Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro...(6 species)
SP303 Staphylococcus warneri	SPN41 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.854%	SPP20 Bradyrhizobium archetypum_austrialiense_cajani_japonicum_liaoningae...(8 species)
SP307 Shigella sonnei	SPN414 Alistipes senegalensis nov_93.673%	SPP3 Staphylococcus capitis_epidermidis
SP317 Atlantibacter hermanningii	SPN415 Enterobacter mori nov_97.951%	SPP34 Staphylococcus argenteus_aureus_roterodami
SP321 Escherichia coli	SPN42 Bacteroides uniformis nov_95.893%	SPN35 Acinetobacter calcoaceticus_nitii