

Species

SP135 Leucobacter chromiresistens
SP14 Acinetobacter johnsonii
SP140 Enterobacter cloacae
SP142 Mammaliacoccus lentus
SP143 Chryseobacterium gambrini
SP148 Ralstonia sp._HMT_406
SP149 Atlantibacter hermannii
SP15 Proteus mirabilis
SP152 Actinomyces sp._HMT_170
SP155 Schaalila sp._HMT_180
SP158 Mammaliacoccus sciuri
SP16 Streptococcus oralis_subsp._tigurinus_clade_071
SP162 Granulicatella adiacens
SP163 Schaalila odontolytica
SP164 Capnocytophaga granulosa
SP166 Mycoplasma faucium
SP167 Lactobacillus johnsonii
SP169 Acinetobacter radioresistens
SP17 Acidovorax temperans
SP173 Fretibacterium fastidiosum
SP174 Streptococcus gordonii
SP175 Anaeroglobus geminatus
SP176 Treponema vincentii
SP177 Treponema sp._HMT_231
SP178 Enterobacter cancerogenus
SP179 Eubacteriales_[G-3] bacterium_MOT-163
SP18 Streptococcus mitis
SP182 Scardovia wiggisiae
SP185 Veillonella rogosa
SP186 Peptoniphilus sp._HMT_187
SP19 Cutibacterium acnes
SP190 Lachnospiraceae_[G-3] bacterium_HMT_100
SP193 Cardiobacterium valvarum
SP198 Oscillospiraceae_[G-7] bacterium_MOT-154
SP199 Aggregatibacter sp._HMT_898
SP2 Klebsiella pneumoniae
SP20 Rhodococcus qingshengii
SP200 Lactobacillus jensenii
SP201 Acinetobacter septicus
SP202 Megasphaera sp._HMT_123
SP203 Porphyromonas pasteri
SP208 Staphylococcus warneri
SP21 Bacillus haynesii
SP218 Treponema denticola
SP219 Haemophilus parainfluenzae
SP220 Oscillospiraceae_[G-4] bacterium_MOT-151
SP221 Leptotrichia buccalis
SP225 Prevotella sp._HMT_376
SP226 Roseomonas mucosa
SP23 Streptococcus sp._HMT_423
SP230 Porphyromonas catoniae
SP231 Acinetobacter modestus
SP232 Sphingomonas leidyi
SP236 Bacillus subtilis
SP237 Johnsonella sp._HMT_166
SP238 Tannerella serpentiniformis
SP239 Acutalibacter muris
SP24 Fiiifactor alocis
SP240 Arthrobacter psychrolactophilus
SP242 Treponema sp._HMT_268
SP248 Alloprevotella tannerae
SP250 Peptoniphilaceae_[G-1] bacterium_HMT_113
SP251 Anaerolineae_[G-1] bacterium_HMT_439
SP254 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
SP256 Brevundimonas variabilis
SP258 Corynebacterium macginleyi
SP259 Porphyromonas endodontalis
SP261 Bacteroides intestinalis
SP262 Parabacteroides distasonis
SP263 Methylobacterium goeingense

SP68 Pelomonas saccharophila
SP69 Epilithimonas hominis
SP7 Staphylococcus caprae
SP70 Parvimonas micra
SP71 Shigella sonnei
SP72 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346
SP73 Faecalibaculum rodentium
SP74 Sphingomonas yabuuchiiae
SP75 Bulleidia exstructa
SP8 Enterococcus faecalis
SP81 Tannerella forsythia
SP82 Neisseriaceae_[G-1] bacterium_HMT_174
SP83 Streptococcus oralis
SP85 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
SP86 Microbacterium maritipicum
SP88 Acinetobacter lwoffii
SP9 Eubacteriales_[G-4] bacterium_MOT-164
SP92 Enterobacter hormaechei
SP93 Adlercreutzia caecimuris
SP95 Streptococcus constellatus
SP96 Prevotella baroniae
SP97 Rothia mucilaginosa
SP99 Enterococcus gallinarum
SPN1 Mollicutes_[G-2] bacterium_MOT-187_nov_90.607%
SPN104 Phoea massiliensis_nov_91.471%
SPN106 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.842%
SPN107 Duncaniella freteri_nov_93.712%
SPN108 Adlercreutzia equofaciens_nov_94.577%
SPN112 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.781%
SPN114 Paludicola psychrotolerans_nov_87.759%
SPN117 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.097%
SPN118 Duncaniella freteri_nov_89.919%
SPN12 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_91.736%
SPN120 Mollicutes_[G-2] bacterium_MOT-187_nov_95.285%
SPN123 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_89.918%
SPN125 Kineothrix alysoides_nov_89.562%
SPN126 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_91.803%
SPN127 Selenomonas sp._HMT_137_nov_97.624%
SPN128 Miniimonas arenae_nov_95.736%
SPN129 Pseudoflavonifractor phocaensis_nov_95.859%
SPN13 Lawsonibacter asaccharolyticus_nov_91.116%
SPN130 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_94.929%
SPN131 Acutalibacter muris_nov_94.227%
SPN132 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.933%
SPN133 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.781%
SPN134 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_87.097%
SPN135 Kineothrix alysoides_nov_89.792%
SPN136 Roseburia intestinalis_nov_90.229%
SPN137 Sphaerobacter thermophilus_nov_82.516%
SPN138 Mycobacterium grossiae_nov_97.826%
SPN139 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_95.519%
SPN14 Streptococcus azizii_nov_95.171%
SPN140 Eubacteriales_[G-4] bacterium_MOT-165_nov_92.781%
SPN141 Thermoanaerobaculum aquaticum_nov_78.020%
SPN142 Oribacterium parvum_nov_89.770%
SPN143 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.254%
SPN144 Beduini massiliensis_nov_86.598%
SPN145 Agathobaculum desmolans_nov_91.649%
SPN146 Acetivibrio cellulolyticus_nov_83.190%
SPN147 Anaerotruncus rubinfantis_nov_93.347%
SPN148 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_87.097%
SPN149 Lawsonibacter asaccharolyticus_nov_93.996%
SPN150 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.759%
SPN159 Lawsonibacter asaccharolyticus_nov_91.116%
SPN16 Ithubacter massiliensis_nov_94.572%
SPN173 Lachnospiraceae_[G-2] bacterium_HMT_096_nov_91.632%
SPN178 Streptococcus xylanophilum_nov_91.075%
SPN180 Gelidibacter algens_nov_96.042%
SPN184 Mollicutes_[G-2] bacterium_MOT-187_nov_94.892%
SPN19 Lacrimispora xylanolytica_nov_94.363%

SPN375 Acutalibacter muris_nov_96.289%
SPN378 Lachnospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
SPN379 Culturomica massiliensis_nov_89.817%
SPN382 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.572%
SPN384 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
SPN385 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
SPN390 Eubacteriales_[G-1] bacterium_MOT-159_nov_94.268%
SPN393 Duncaniella freteri_nov_89.697%
SPN398 Turicibacter sanguinis_nov_95.923%
SPN4 Hydrogenoanaerobacterium saccharovorans_nov_90.041%
SPN40 Acetivibrio cellulolyticus_nov_83.801%
SPN402 Duncaniella freteri_nov_89.775%
SPN403 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
SPN407 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_90.369%
SPN41 Lachnospiraceae_[G-2] bacterium_MOT-167_nov_88.773%
SPN410 Breznakia pachnodae_nov_82.824%
SPN411 Hathewayia proteolytica_nov_83.514%
SPN414 Acetivibrio cellulolyticus_nov_83.761%
SPN416 Acetivibrio cellulolyticus_nov_83.153%
SPN42 Phoea massiliensis_nov_89.914%
SPN420 Anaerotruncus rubinfantis_nov_92.708%
SPN425 Glucerbacter canis_nov_93.305%
SPN426 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
SPN429 Erysipelatoclostridium_[Clostridium] innocuum_nov_88.270%
SPN43 Adlercreutzia caecimuris_nov_92.291%
SPN431 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.979%
SPN432 Marinisporobacter balticus_nov_82.692%
SPN433 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.179%
SPN437 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.228%
SPN438 Pseudoflavonifractor capillosus_nov_89.897%
SPN440 Flavonifractor plautii_nov_92.308%
SPN443 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_91.213%
SPN444 Sporobacter terminalis_nov_87.580%
SPN447 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.100%
SPN45 Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.854%
SPN450 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.228%
SPN452 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.157%
SPN453 Longibaculum muris_nov_93.361%
SPN455 Faecalibaculum acidiformans_nov_89.600%
SPN459 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.305%
SPN462 Christensenella massiliensis_nov_88.041%
SPN463 Adlercreutzia caecimuris_nov_89.009%
SPN466 Cryobacterium psychrophilum_nov_77.169%
SPN467 Christensenella hongkongensis_nov_86.308%
SPN47 Butyrivibrio pullicaecorum_nov_85.093%
SPN471 Phoea massiliensis_nov_90.426%
SPN473 Roseburia hominis_nov_91.476%
SPN475 Chryseobacterium yeoncheonense_nov_97.484%
SPN476 Eubacteriales_[G-1] bacterium_MOT-159_nov_92.161%
SPN477 Hydrogenoanaerobacterium saccharovorans_nov_88.589%
SPN479 Agrococcus versicolor_nov_83.227%
SPN49 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.561%
SPN50 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_84.086%
SPN53 Eubacteriales_[G-3] bacterium_MOT-163_nov_87.952%
SPN56 Flavonifractor plautii_nov_93.555%
SPN59 Ithubacter massiliensis_nov_90.644%
SPN6 Adlercreutzia caecimuris_nov_88.961%
SPN60 Neglectibacter timonensis_nov_94.490%
SPN66 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_87.423%
SPN68 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_96.881%
SPN7 Eisenbergiella massiliensis_nov_87.578%
SPN70 Tannerella forsythia_nov_97.516%
SPN71 Hydrogenoanaerobacterium saccharovorans_nov_87.759%
SPN73 Eubacteriales_[G-3] bacterium_MOT-163_nov_85.825%
SPN74 Anaeromassilibacillus senegalensis_nov_92.489%
SPN76 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.996%
SPN79 Duncaniella freteri_nov_90.612%
SPN81 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.466%
SPN86 Butyrivibrio pullicaecorum_nov_85.644%
SPN88 Prevotella veroralis_nov_97.751%