

Species

SP14 Acinetobacter johnsonii	SP69 Epilimnionomonas hominis	SPN378 Lachnociostrium [Clostridium] schindens nov_89.234%
SP140 Enterobacter cloacae	SP7 Staphylococcus caprae	SPN379 Culturomedia massiliensis nov_89.817%
SP142 Mammaliococcus lentus	SP70 Parvimonas micra	SPN382 Lachnospiraceae [G-6] bacterium MOT-171 nov_94.572%
SP143 Chryseobacterium gambrini	SP71 Shigella sonnei	SPN384 Oscillospiraceae [G-1] bacterium MOT-147 nov_96.674%
SP148 Ralstonia sp._HMT_406	SP72 Saccharibacteria (TM7) [G-1] bacterium_HMT_346	SPN385 Muribaculaceae [G-2] bacterium MOT-104 nov_89.000%
SP149 Atlantibacter hermanningii	SP73 Faecalibaculum rodentium	SPN390 Eubacteriales [G-1] bacterium MOT-159 nov_94.268%
SP15 Proteus mirabilis	SP74 Sphingomonas yabuuchiae	SPN393 Duncaniella freteri nov_89.697%
SP152 Actinomyces sp._HMT_170	SP75 Bulleidia extructa	SPN398 Turicibacter sanguinis nov_95.923%
SP155 Schaalia sp._HMT_180	SP8 Enterococcus faecalis	SPN4 Hydrogenoanaerobacterium saccharovorans nov_90.041%
SP158 Mammaliococcus sciuri	SP81 Tannerella forsythia	SPN40 Acetivibrio cellulolyticus nov_83.801%
SP16 Streptococcus oralis subsp._tigurinus_clade_071	SP82 Neisseriaceae [G-1] bacterium_HMT_174	SPN402 Duncaniella freteri nov_89.775%
SP162 Granulicatella adiacens	SP83 Streptococcus oralis	SPN403 Oscillospiraceae [G-2] bacterium MOT-149 nov_95.198%
SP163 Schaalia odontolytica	SP85 Saccharibacteria (TM7) [G-5] bacterium_HMT_356	SPN407 Lachnospiraceae [G-10] bacterium MOT-175 nov_90.369%
SP164 Capnocytophaga granulosa	SP86 Microbacterium maritipicum	SPN41 Lachnospiraceae [G-2] bacterium MOT-167 nov_88.773%
SP166 Mycoplasma faucium	SP88 Acinetobacter lwoffii	SPN410 Breznakia pachnoda nov_82.824%
SP167 Lactobacillus johnsonii	SP9 Eubacteriales [G-4] bacterium MOT-164	SPN411 Hathewayia proteolytica nov_83.514%
SP169 Acinetobacter radioresistens	SP92 Enterobacter hormaechei	SPN414 Acetivibrio cellulolyticus nov_83.761%
SP17 Acidovorax temperans	SP93 Adlercreutzia caecimuris	SPN416 Acetivibrio cellulolyticus nov_83.153%
SP173 Fretibacterium fastidiosum	SP95 Streptococcus constellatus	SPN42 Phoea massiliensis nov_89.914%
SP174 Streptococcus gordonii	SP96 Prevotella baroniae	SPN420 Anaerotruncus rubiinfantis nov_92.708%
SP175 Anaeroglobus gluminatus	SP97 Rothia mucilaginosa	SPN425 Glucerbacter canis nov_93.305%
SP176 Treponema vincentii	SP99 Enterococcus gallinarum	SPN426 Oscillospiraceae [G-6] bacterium MOT-153 nov_91.631%
SP177 Treponema sp._HMT_231	SPN1 Mollicutes [G-2] bacterium MOT-187 nov_90.607%	SPN429 Erysipelatoclostridium [Clostridium] innocuum nov_88.270%
SP178 Enterobacter cancerogenus	SPN104 Phoea massiliensis nov_91.471%	SPN43 Adlercreutzia caecimuris nov_92.291%
SP179 Eubacteriales [G-3] bacterium MOT-163	SPN106 Oscillospiraceae [G-4] bacterium MOT-151 nov_95.842%	SPN431 Lachnospiraceae [G-6] bacterium MOT-171 nov_94.979%
SP18 Streptococcus mitis	SPN107 Duncaniella freteri nov_93.712%	SPN432 Marinisporobacter balticus nov_82.692%
SP182 Scardovia wiggsiae	SPN108 Adlercreutzia equolifaciens nov_94.577%	SPN433 Oscillospiraceae [G-4] bacterium MOT-151 nov_94.179%
SP185 Veillonella rogosae	SPN112 Oscillospiraceae [G-2] bacterium MOT-149 nov_94.781%	SPN437 Eubacteriales [G-4] bacterium MOT-164 nov_97.228%
SP186 Peptoniphilus sp._HMT_187	SPN114 Paludicola psychrotolerans nov_87.759%	SPN438 Pseudoflavonifractor capillosus nov_89.897%
SP19 Cutibacterium acnes	SPN117 Lachnospiraceae [G-7] bacterium MOT-172 nov_91.097%	SPN440 Flavonifractor plautii nov_92.308%
SP190 Lachnospiraceae [G-3] bacterium_HMT_100	SPN118 Duncaniella freteri nov_89.919%	SPN443 Lachnospiraceae [G-6] bacterium MOT-171 nov_91.213%
SP193 Cardiobacterium valvarum	SPN12 Oscillospiraceae [G-3] bacterium MOT-150 nov_91.736%	SPN444 Sporobacter termitidis nov_87.580%
SP198 Oscillospiraceae [G-7] bacterium MOT-154	SPN120 Mollicutes [G-2] bacterium MOT-187 nov_95.285%	SPN447 Oscillospiraceae [G-4] bacterium MOT-151 nov_92.100%
SP199 Aggregatibacter sp._HMT_898	SPN123 Lachnospiraceae [G-9] bacterium MOT-174 nov_89.918%	SPN45 Lachnospiraceae [G-14] bacterium MOT-183 nov_97.854%
SP2 Klebsiella pneumoniae	SPN125 Kineothrix alysoides nov_89.562%	SPN450 Eubacteriales [G-4] bacterium MOT-164 nov_97.228%
SP20 Rhodococcus qingshengii	SPN126 Lachnospiraceae [G-10] bacterium MOT-175 nov_91.803%	SPN452 Muribaculaceae [G-2] bacterium MOT-104 nov_89.157%
SP200 Lactobacillus jensenii	SPN127 Selenomonas sp._HMT_137 nov_97.624%	SPN453 Longibaculum muris nov_93.361%
SP201 Acinetobacter septicus	SPN128 Miniimonas arenae nov_95.736%	SPN455 Faecalicoccus acidiformans nov_89.600%
SP202 Megaspheera sp._HMT_123	SPN129 Pseudoflavonifractor phocaeensis nov_95.859%	SPN459 Lachnospiraceae [G-6] bacterium MOT-171 nov_93.305%
SP203 Porphyromonas pasteri	SPN13 Lawsonibacter asaccharolyticus nov_91.116%	SPN462 Christensenella massiliensis nov_88.041%
SP208 Staphylococcus warneri	SPN130 Lachnospiraceae [G-11] bacterium MOT-178 nov_94.929%	SPN463 Adlercreutzia caecimuris nov_89.009%
SP21 Bacillus haynesii	SPN131 Acutalibacter muris nov_94.227%	SPN466 Cryobacterium psychrophilum nov_77.169%
SP218 Treponema denticola	SPN132 Lachnospiraceae [G-6] bacterium MOT-171 nov_93.933%	SPN467 Christensenella hongkongensis nov_86.308%
SP219 Haemophilus parainfluenzae	SPN133 Lachnospiraceae [G-14] bacterium MOT-185 nov_96.781%	SPN47 Butyricicoccus pullicaecorum nov_85.093%
SP220 Oscillospiraceae [G-4] bacterium MOT-151	SPN134 Lachnospiraceae [G-12] bacterium MOT-179 nov_97.097%	SPN471 Phoea massiliensis nov_90.426%
SP221 Leptotrichia buccalis	SPN135 Kineothrix alysoides nov_89.792%	SPN473 Roseburia hominis nov_91.476%
SP225 Prevotella sp._HMT_376	SPN136 Roseburia intestinalis nov_90.229%	SPN475 Chryseobacterium yeoncheonense nov_97.484%
SP226 Roseomonas mucosa	SPN137 Sphaerobacter thermophilus nov_82.516%	SPN476 Eubacteriales [G-1] bacterium MOT-159 nov_92.161%
SP23 Streptococcus sp._HMT_423	SPN138 Mycobacterium grossiae nov_97.826%	SPN477 Hydrogenoanaerobacterium saccharovorans nov_88.589%
SP230 Porphyromonas catoniae	SPN139 Lachnospiraceae [G-11] bacterium MOT-178 nov_95.519%	SPN479 Agrococcus versicolor nov_83.227%
SP231 Acinetobacter modestus	SPN14 Streptococcus azizii nov_95.171%	SPN49 Lachnospiraceae [G-6] bacterium MOT-171 nov_94.561%
SP232 Sphingomonas leidy	SPN140 Eubacteriales [G-4] bacterium MOT-165 nov_92.781%	SPN50 Clostridiales [F-1][G-1] bacterium_HMT_093 nov_84.086%
SP236 Bacillus subtilis	SPN141 Thermoanaerobaculum aquaticum nov_78.020%	SPN53 Eubacteriales [G-3] bacterium MOT-163 nov_87.952%
SP237 Johnsonella sp._HMT_166	SPN142 Oribacterium parvum nov_89.770%	SPN56 Flavonifractor plautii nov_93.555%
SP238 Tannerella serpentiniformis	SPN143 Lachnospiraceae [G-14] bacterium MOT-182 nov_90.254%	SPN59 Ihubacter massiliensis nov_90.644%
SP239 Acutalibacter muris	SPN144 Beduini massiliensis nov_86.598%	SPN6 Adlercreutzia muris nov_88.961%
SP24 Filifactor alocis	SPN145 Agathobaculum desmolans nov_91.649%	SPN60 Neglectibacter timonensis nov_94.490%
SP240 Arthrobacter psychrolactophilus	SPN146 Acetivibrio cellulolyticus nov_83.190%	SPN66 Oscillospiraceae [G-3] bacterium MOT-150 nov_87.423%
SP242 Treponema sp._HMT_268	SPN147 Anaerotruncus rubiinfantis nov_93.347%	SPN68 Oscillospiraceae [G-4] bacterium MOT-151 nov_96.881%
SP248 Alloprevotella tannerae	SPN148 Lachnoclostridium [Clostridium] polysaccharolyticum nov_90.437%	SPN7 Eisenbergiella massiliensis nov_87.578%
SP250 Peptoniphilaceae [G-1] bacterium_HMT_113	SPN149 Lawsonibacter asaccharolyticus nov_93.996%	SPN70 Tannerella forsythia nov_97.516%
SP251 Anaerolineae [G-1] bacterium_HMT_439	SPN150 Lachnospiraceae [G-6] bacterium MOT-171 nov_94.759%	SPN71 Hydrogenoanaerobacterium saccharovorans nov_87.759%
SP254 Saccharibacteria (TM7) [G-1] bacterium_HMT_349	SPN159 Lawsonibacter asaccharolyticus nov_91.116%	SPN73 Eubacteriales [G-3] bacterium MOT-163 nov_85.825%
SP256 Brevundimonas variabilis	SPN16 Ihubacter massiliensis nov_94.572%	SPN74 Anaeromassilibacillus senegalensis nov_92.489%
SP258 Corynebacterium magcinleyi	SPN173 Lachnospiraceae [G-2] bacterium_HMT_096 nov_91.632%	SPN76 Lachnospiraceae [G-14] bacterium MOT-185 nov_96.996%
SP259 Porphyromonas endodontalis	SPN18 Eubacterium xylanophilum nov_91.075%	SPN79 Duncaniella freteri nov_90.612%
SP261 Bacteroides intestinalis	SPN180 Gelidibacter algens nov_96.042%	SPN81 Oscillospiraceae [G-2] bacterium MOT-149 nov_96.466%
SP262 Parabacteroides distans	SPN184 Mollicutes [G-2] bacterium MOT-187 nov_94.892%	SPN86 Butyricicoccus pullicaecorum nov_85.644%