

Species

SP121 Sphingomonas zeae
SP123 Cetobacterium somerae
SP124 Treponema denticola
SP126 Lactobacillus johnsonii
SP127 Enterobacter hormaechei
SP129 Streptococcus gordonii
SP13 Streptococcus oralis
SP130 Shigella flexneri
SP132 Fusobacterium periodonticum
SP133 Eubacteriales_[G-1] bacterium_MOT-158
SP134 Bifidobacterium longum
SP135 Pseudoramibacter alactolyticus
SP136 Agromyces mediolanus
SP137 Alloprevotella tannerae
SP138 Peptostreptococcaceae_[XII][G-4] bacterium_HMT_369
SP139 Lachnoclostridium [Clostridium] symbiosum
SP14 Streptococcus sp._HMT_064
SP140 Bradyrhizobium pachyrhizi
SP141 Peptococcaceae_[G-1] bacterium_MOT-146
SP142 Peptidiphaga sp._HMT_183
SP143 Leucobacter chromiirensis
SP144 Acinetobacter lwofii
SP145 Deinococcus geothermalis
SP146 Selenomonas flueggei
SP147 Catonella sp._HMT_164
SP148 Prevotella nigrescens
SP149 Adlercreutzia caecimuris
SP15 Acinetobacter johnsonii
SP151 Filifactor aloisii
SP152 Mucispirillum schaedleri
SP156 Methylobacterium brachiatum
SP157 Atlantibacter hermannii
SP158 Streptococcus acidominimus
SP159 Streptococcus oralis_subsp._denticani_clade_058
SP16 Haemophilus haemolyticus
SP161 Aggregatibacter aphrophilus
SP164 Corynebacterium aurumucosum
SP165 Streptococcus oralis_subsp._tigurinus_clade_070
SP166 Streptococcus mutans
SP168 Butyrivibrio sp._HMT_080
SP170 Aggregatibacter sp._HMT_512
SP171 Johnsonella sp._HMT_166
SP172 Oscillospiraceae_[G-6] bacterium_MOT-153
SP176 Prevotella sp._HMT_304
SP177 Lachnospiraceae_[G-3] bacterium_MOT-168
SP179 Desulfobulbus sp._HMT_041
SP18 Janibacter melonis
SP180 Prevotella melaninogenica
SP181 Actinomyces sp._HMT_896
SP185 Dialister pneumosintes
SP186 Corynebacterium matruchotii
SP187 Ruthenibacterium lactatiformans
SP189 Clostridium disporicum
SP19 Limosilactobacillus reuteri
SP191 Schaalia sp._HMT_178
SP192 Thermicanus aegyptius
SP194 Duncaniella freteri
SP195 Staphylococcus warneri
SP196 Rothia dentocariosa
SP197 Actinomyces sp._HMT_448
SP198 Leptotrichia buccalis
SP20 Akkermansia muciniphila
SP201 Actinomyces sp._HMT_414
SP21 Alishewanella agri
SP22 Streptococcus sinensis
SP23 Kosakonia sacchari
SP24 Erysipelotrichaceae_[G-1] bacterium_MOT-189
SP25 Porphyromonas endodontalis
SP26 Cutibacterium acnes
SP27 Streptococcus danieliae

SP67 Porphyromonas gingivalis
SP7 Neisseriaceae_[G-1] bacterium_MOT-031
SP70 Muribaculaceae_[G-1] bacterium_MOT-129
SP71 Stenotrophomonas [Pseudomonas] hibiscicola
SP73 Parabacteroides goldsteinii
SP74 Acidovorax temperans
SP75 Alloprevotella sp._HMT_473
SP76 Rhodococcus qingshengii
SP77 Faecalibaculum rodentium
SP78 Bifidobacterium pseudolongum
SP79 Enterobacter cancerogenus
SP8 Phocaeicola sartorii
SP80 Lawsonella clevelandensis
SP81 Campylobacter sp._HMT_044
SP82 Acidovorax ebureus
SP83 Alistipes sp._MOT-127
SP84 Escherichia fergusonii
SP91 Staphylococcus saccharolyticus
SP92 Leptotrichia sp._HMT_219
SP93 Streptococcus cristatus_clade_578
SP94 Muribaculaceae_[G-2] bacterium_MOT-104
SP95 Streptococcus chosunense
SP96 Blautia hominis
SP97 Enterococcus casseliflavus
SP98 Streptococcus sanguinis
SPN100 Duncaniella freteri_nov_91.853%
SPN107 Muribaculaceae_[G-2] bacterium_MOT-104_nov_85.686%
SPN108 Anaeromassilibacillus senegalensis_nov_93.390%
SPN109 Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.431%
SPN11 Lacrimispora xylanolytica_nov_93.789%
SPN118 Duncaniella freteri_nov_90.184%
SPN120 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.279%
SPN126 Alloprevotella sp._HMT_473_nov_89.366%
SPN13 Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.576%
SPN131 Duncaniella freteri_nov_92.915%
SPN132 Leptotrichia hofstadii_nov_96.970%
SPN137 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.640%
SPN142 Marinisporobacter balticus_nov_82.692%
SPN144 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.621%
SPN149 Alistipes senegalensis_nov_93.648%
SPN153 Anaerostipes caccae_nov_96.328%
SPN156 Alistipes putredinis_nov_94.444%
SPN161 Lachnospiraceae_[G-3] bacterium_MOT-168_nov_92.902%
SPN164 Duncaniella freteri_nov_90.612%
SPN170 Erysipelatoclostridium [Clostridium] innocuum_nov_88.270%
SPN174 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.427%
SPN175 Alistipes finegoldii_nov_93.608%
SPN185 Hathewayia proteolytica_nov_83.514%
SPN191 Duncaniella freteri_nov_93.699%
SPN194 Muribaculaceae_[G-1] bacterium_MOT-129_nov_90.816%
SPN196 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
SPN199 Beduini massiliensis_nov_87.705%
SPN206 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.600%
SPN207 Phocaea massiliensis_nov_90.129%
SPN210 Duncaniella freteri_nov_87.424%
SPN217 Acutalibacter muris_nov_96.289%
SPN218 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.582%
SPN222 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.400%
SPN220 Longibaculum muris_nov_86.957%
SPN228 Parabacteroides distasonis_nov_97.938%
SPN229 Flavonifractor plautii_nov_92.308%
SPN23 Duncaniella freteri_nov_86.842%
SPN238 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.290%
SPN240 Blautia luti_nov_94.561%
SPN242 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.822%
SPN246 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.655%
SPN249 Duncaniella freteri_nov_88.330%
SPN254 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_91.718%
SPN256 Duncaniella freteri_nov_87.071%
SPN259 Duncaniella freteri_nov_93.712%

SPN331 Lawsonibacter asaccharolyticus_nov_90.722%
SPN332 Duncaniella freteri_nov_89.069%
SPN333 Alistipes senegalensis_nov_93.089%
SPN334 Faecalicatena orotica_nov_92.484%
SPN335 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.931%
SPN336 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_93.515%
SPN337 Actinomyces sp._HMT_169_nov_97.992%
SPN338 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_94.411%
SPN339 Eubacterium coprostanoligenes_nov_91.631%
SPN340 Natroniella acetigena_nov_80.992%
SPN341 Muribaculaceae_[G-1] bacterium_MOT-129_nov_84.929%
SPN342 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.623%
SPN343 Hydrogenoanaerobacterium saccharovorans_nov_90.249%
SPN344 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.572%
SPN345 Phocaea massiliensis_nov_90.426%
SPN346 Neglectibacter timonensis_nov_95.325%
SPN347 Butyrivibrio proteoclasticus_nov_83.594%
SPN348 Eubacteriales_[G-3] bacterium_MOT-163_nov_84.913%
SPN349 Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.089%
SPN350 Bacteroidetes_[G-3] bacterium_HMT_436_nov_86.585%
SPN351 Duncaniella freteri_nov_88.978%
SPN352 Lachnospiraceae_[G-5] bacterium_MOT-170_nov_97.904%
SPN353 Phocaea massiliensis_nov_87.179%
SPN354 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_95.417%
SPN355 Lachnospiraceae_[G-13] bacterium_MOT-181_nov_91.189%
SPN356 Eisenbergiella massiliensis_nov_84.867%
SPN357 Sporobacter termitidis_nov_87.580%
SPN358 Lachnospiraceae_[G-11] bacterium_MOT-177_nov_88.296%
SPN359 Christensenella hongkongensis_nov_86.308%
SPN360 Lachnospiraceae_[G-14] bacterium_MOT-182_nov_90.254%
SPN361 Lachnoclostridium phytofermentans_nov_90.417%
SPN362 Blautia marasmi_nov_93.711%
SPN363 Muribaculaceae_[G-1] bacterium_MOT-129_nov_83.636%
SPN364 Butyrivibrio pullicaecorum_nov_85.093%
SPN365 Selenomonas sp._HMT_137_nov_97.228%
SPN366 Mollicutes_[G-2] bacterium_MOT-187_nov_95.481%
SPN367 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_83.582%
SPN368 Eubacterium xylanophilum_nov_91.075%
SPN369 Ihubacter massiliensis_nov_90.644%
SPN370 Prevotella shahii_nov_87.602%
SPN371 Anaerotruncus rubiinfantis_nov_92.500%
SPN372 Eubacteriales_[G-3] bacterium_MOT-163_nov_85.944%
SPN373 Duncaniella freteri_nov_89.379%
SPN374 Anaerotruncus faecicola_nov_85.396%
SPN375 Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.643%
SPN376 Duncaniella freteri_nov_85.685%
SPN377 Glucibacter canis_nov_93.305%
SPN378 Eubacteriales_[G-4] bacterium_MOT-164_nov_92.373%
SPN379 Parafannyhessea umbonata_nov_92.161%
SPN380 Veillonellaceae_[G-1] bacterium_HMT_129_nov_96.356%
SPN381 Adlercreutzia muris_nov_97.186%
SPN382 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.802%
SPN383 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_91.423%
SPN384 Hydrogenoanaerobacterium saccharovorans_nov_88.797%
SPN385 Kineothrix alysioides_nov_86.667%
SPN390 Lawsonibacter asaccharolyticus_nov_90.329%
SPN42 Lachnoclostridium [Clostridium] aminophilum_nov_87.318%
SPN45 Duncaniella freteri_nov_92.653%
SPN5 Duncaniella freteri_nov_86.290%
SPN52 Chryseobacterium yeoncheonense_nov_97.484%
SPN53 Pyrinomonas methylaliphatogenes_nov_96.809%
SPN58 Alloprevotella sp._HMT_473_nov_89.634%
SPN64 Muricomes intestini_nov_89.583%
SPN67 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.373%
SPN71 Muribaculaceae_[G-2] bacterium_MOT-104_nov_86.000%
SPN75 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_89.293%
SPN86 Lachnoclostridium [Clostridium] polysaccharolyticum_nov_92.083%
SPN88 Duncaniella freteri_nov_91.039%
SPN92 Duncaniella freteri_nov_91.429%
SPN93 Duncaniella freteri_nov_87.169%