

Species

SP21 Tepidococcaceae [G-1] bacterium_MOT-146

SP22 Prevotella sp._MOT-128

SP23 Lactobacillus taiwanensis

SP24 Acutalibacter muris

SP27 Roseburia faecis

SP28 Flavonifractor plautii

SP29 Oscillospiraceae [G-2] bacterium_MOT-149

SP3 Lachnospiraceae [G-14] bacterium_MOT-183

SP30 Robinsoniella perniensis

SP31 Lachnospiraceae [G-9] bacterium_MOT-174

SP32 Eubacteriales [G-4] bacterium_MOT-164

SP33 Erysipelatoclostridium [Clostridium] coeleatum

SP34 Alistipes timonensis

SP35 Eubacteriales [G-2] bacterium_MOT-162

SP36 Lawsonibacter asaccharolyticus

SP37 Eubacteriales [G-1] bacterium_MOT-159

SP38 Eubacteriales [G-1] bacterium_MOT-158

SP39 Neglectibacter timonensis

SP4 Bacteroides stercorisoris

SP40 Eubacteriales [G-3] bacterium_MOT-163

SP42 Lachnospiraceae [G-2] bacterium_MOT-167

SP43 Oscillospiraceae [G-7] bacterium_MOT-154

SP44 Clostridium tertium

SP5 Muribaculum intestinale

SP6 Lachnospiraceae [G-14] bacterium_MOT-184

SP7 Oscillospiraceae [G-6] bacterium_MOT-153

SP8 Parabacteroides goldsteinii

SP9 Akkermansia muciniphila

SPN1 Prevotellamassilia timonensis_nov_92.641%

SPN10 Parabacteroides goldsteinii_nov_97.614%

SPN100 Anaerotaenia torta_nov_97.273%

SPN101 Oscillospiraceae [G-3] bacterium_MOT-150_nov_90.745%

SPN102 Kineothrix alysoides_nov_95.227%

SPN103 Anaerotruncus rubifantiss_nov_92.760%

SPN104 Muribaculaceae [G-2] bacterium_MOT-104_nov_91.991%

SPN105 Meditteraneibacter [Ruminococcus] gnavus_nov_93.424%

SPN106 Alistipes putredinis_nov_95.887%

SPN107 Oscillibacter valericigenis_nov_95.260%

SPN108 Kineothrix alysoides_nov_97.279%

SPN109 Pseudobutyryvibrio ruminis_nov_91.176%

SPN11 Lachnospiraceae [G-9] bacterium_MOT-174_nov_96.364%

SPN110 Muribaculaceae [G-1] bacterium_MOT-129_nov_91.522%

SPN111 Saccharofermentans acetigenes_nov_88.764%

SPN112 Lacinimicrobia indolis_nov_90.724%

SPN113 Kineothrix alysoides_nov_93.651%

SPN114 Lawsonibacter asaccharolyticus_nov_97.973%

SPN115 Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.35%

SPN116 Muribaculaceae [G-2] bacterium_MOT-104_nov_90.870%

SPN117 Muribaculaceae [G-2] bacterium_MOT-104_nov_93.043%

SPN118 Muribaculaceae [G-2] bacterium_MOT-104_nov_91.106%

SPN119 Alistipes putredinis_nov_95.879%

SPN20 Oscillospiraceae [G-4] bacterium_MOT-151_nov_93.708%

SPN20 Odoribacter splanchnicus_nov_93.939%

SPN121 Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.506%

SPN122 Ruthenibacterium lactatiformans_nov_97.045%

SPN123 Lachnospiraceae [G-10] bacterium_MOT-175_nov_95.47%

SPN124 Lachnospiraceae [G-11] bacterium_MOT-178_nov_97.97%

SPN125 Lachnospiraceae [G-9] bacterium_MOT-174_nov_96.388%

SPN126 Lachnospiraceae [G-14] bacterium_MOT-185_nov_96.59%

SPN127 Muribaculaceae [G-2] bacterium_MOT-104_nov_90.022%

SPN128 Faecalicatena orotica_nov_95.238%

SPN129 Enterocloster asparagiformis_nov_94.344%

SPN13 Prevotellamassilia timonensis_nov_94.168%

SPN130 Eisenbergiella massiliensis_nov_96.599%

SPN131 Neglectibacter timonensis_nov_97.500%

SPN132 Maitiella massiliensis_nov_92.094%

SPN133 Anaerotruncus rubifantiss_nov_92.517%

SPN134 Neglectibacter timonensis_nov_97.727%

SPN135 Butyricicoccus pullicaecorum_nov_94.820%

SPN136 Lachnospiraceae [G-11] bacterium_MOT-176_nov_95.94%

SPN137 Eubacterium aerogenes-like_nov_95.701%

SPN149 Ruminococcus albus_nov_92.900%

SPN15 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_92.568%

SPN150 Alistipes putredinis_nov_96.529%

SPN151 Lawsonibacter asaccharolyticus_nov_97.297%

SPN152 Clostridium oryzae_nov_88.889%

SPN153 Eisenbergiella massiliensis_nov_95.260%

SPN154 Rhodospirillum rubrum_nov_88.036%

SPN155 Lawsonibacter asaccharolyticus_nov_95.730%

SPN156 Eubacteriales_[G-3] bacterium_MOT-163_nov_95.023%

SPN157 Anaerotrignum lactatifermentans_nov_97.523%

SPN158 Phoea massiliensis_nov_95.682%

SPN159 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.171%

SPN16 Harryflintia acetispora_nov_96.388%

SPN160 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.485%

SPN161 Lachnospiraceae_[G-6] bacterium_MOT-171_nov_96.606%

SPN162 Muribaculum intestinale_nov_93.737%

SPN163 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_90.337%

SPN164 Falcatimonas natans_nov_92.955%

SPN165 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.692%

SPN166 Butyrivibrio pullicaeorum_nov_94.382%

SPN167 Kineothrix alysoides_nov_95.465%

SPN168 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.396%

SPN169 Eubacteriales_[G-3] bacterium_MOT-163_nov_93.679%

SPN17 Marvinbryantia formatexiensis_nov_91.403%

SPN170 Lachnospiraceae_[G-12] bacterium_MOT-180_nov_93.665%

SPN171 Duncaniella freteri_nov_88.462%

SPN172 Lachnoclostridium_[Clostridium] populeti_nov_94.331%

SPN173 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.227%

SPN174 Lachnoclostridium_[Clostridium] populeti_nov_96.145%

SPN175 Anaerotruncus rubinfantis_nov_93.182%

SPN176 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_97.065%

SPN177 Eubacteriales_[G-4] bacterium_MOT-164_nov_96.606%

SPN178 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.270%

SPN179 Eubacteriales_[G-4] bacterium_MOT-165_nov_97.059%

SPN18 Lachnospiraceae_[G-10] bacterium_MOT-175_nov_90.693%

SPN180 Anaerotruncus colihominis_nov_94.091%

SPN181 Mageelbaccilus indolicus_nov_87.678%

SPN182 Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.796%

SPN183 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.873%

SPN184 Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.238%

SPN185 Lachnoclostridium_[Clostridium] populeti_nov_92.955%

SPN186 Roseburia faecis_nov_95.475%

SPN187 Muricomes intestini_nov_94.331%

SPN188 Neglectibacter timonensis_nov_97.959%

SPN189 Eubacteriales_[G-2] bacterium_MOT-162_nov_95.692%

SPN19 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_94.344%

SPN190 Harryflintia acetispora_nov_93.468%

SPN191 Kineothrix alysoides_nov_96.372%

SPN192 Gracilibacter thermotolerans_nov_88.315%

SPN193 Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%

SPN194 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_96.629%

SPN195 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.919%

SPN196 Paludicola psychrotolerans_nov_94.533%

SPN197 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.291%

SPN198 Oscillibacter ruminantium_nov_93.919%

SPN199 Anaerostipes butyraticus_nov_97.285%

SPN2 Acetatifactor muris_nov_92.551%

SPN20 Roseburia intestinalis_nov_95.475%

SPN200 Alistipes sp._MOT-127_nov_91.775%

SPN201 Murimonas intestini_nov_96.591%

SPN202 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_96.323%

SPN203 Eubacteriales_[G-4] bacterium_MOT-165_nov_97.065%

SPN204 Kineothrix alysoides_nov_97.059%

SPN205 Lachnoclostridium_[Clostridium] populeti_nov_92.986%

SPN206 Kineothrix alysoides_nov_96.136%

SPN207 Anaerocolumna jejuensis_nov_92.358%

SPN208 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_96.847%

SPN209 Eubacteriales_[G-1] bacterium_MOT-160_nov_95.270%

SPN21 Christensenella hongkongensis_nov_86.353%

SPN210 Mediobacterium_[Ruminococcus] grnavus_nov_97.045%

SPN211 Blautia obeum_nov_96.825%

SPN223 *Elaiobacterium massiliensis*_nov_94.362%

SPN224 *Mediterraneibacterium* [Ruminococcus] *gnavus*_nov_97.285%

SPN225 *Eubacteriales*_[G-4] *bacterium*_MOT-165_nov_95.918%

SPN226 *Lachnospiraceae*_[G-9] *bacterium*_MOT-174_nov_96.136%

SPN23 *Peptostreptococcaceae*_[XII][G-2] *bacterium*_HMT_091_nov_93.002%

SPN24 *Harryflintia acetispora*_nov_92.517%

SPN25 *Peptococcus* sp._HMT_168_nov_89.979%

SPN26 *Kineothrix alisoides*_nov_93.682%

SPN27 *Parabacteroides goldsteinii*_nov_93.074%

SPN3 *Saccharofermentans acetigenes*_nov_88.739%

SPN32 *Coprococcus catus*_nov_94.570%

SPN38 *Roseburia faecis*_nov_97.964%

SPN4 *Bilophila wadsworthia*_nov_91.684%

SPN49 *Duncanella freteri*_nov_93.103%

SPN5 *Oscillibacter valericigenes*_nov_95.475%

SPN53 *Eubacterium ventriosum*_nov_96.825%

SPN6 *Oscillospiraceae*_[G-1] *bacterium*_MOT-147_nov_95.937%

SPN60 *Muribaculaceae*_[G-2] *bacterium*_MOT-104_nov_93.074%

SPN63 *Muribaculaceae*_[G-2] *bacterium*_MOT-104_nov_92.441%

SPN7 *Lachnospiraceae*_[G-2] *bacterium*_MOT-167_nov_97.968%

SPN72 *Lacrimispora xylanolytica*_nov_97.285%

SPN74 *Hydrogenoanaerobacterium saccharovorans*_nov_93.636%

SPN78 *Culturomedia massiliensis*_nov_93.709%

SPN79 *Kineothrix alisoides*_nov_95.928%

SPN8 *Gracilibacter thermolators*_nov_87.668%

SPN80 *Lachnospiraceae*_[G-10] *bacterium*_MOT-175_nov_96.372%

SPN81 *Oscillospiraceae*_[G-4] *bacterium*_MOT-151_nov_96.847%

SPN82 *Muribaculaceae*_[G-2] *bacterium*_MOT-104_nov_92.191%

SPN83 *Lachnospiraceae*_[G-10] *bacterium*_MOT-175_nov_92.174%

SPN84 *Lachnospiraceae*_pacense_nov_96.825%

SPN85 *Muribaculaceae*_[G-2] *bacterium*_MOT-104_nov_89.462%

SPN86 *Eubacterium coprostanigenes*_nov_95.485%

SPN87 *Muribaculaceae*_[G-2] *bacterium*_MOT-104_nov_91.974%

SPN88 *Lachnospiraceae*_[G-12] *bacterium*_MOT-179_nov_92.534%

SPN89 *Lachnospiraceae*_[G-14] *bacterium*_MOT-184_nov_94.989%

SPN9 *Eisenbergiella tayi*_nov_94.318%

SPN90 *Pseudoflavivibractor capillosus*_nov_95.721%

SPN91 *Anaerotrignum lactatifermentans*_nov_95.270%

SPN92 *Lachnospiraceae*_[G-9] *bacterium*_MOT-174_nov_95.238%

SPN93 *Caecibacterium sporiformans*_nov_95.045%

SPN94 *Alistipes timonensis*_nov_97.831%

SPN95 *Oscillospiraceae*_[G-2] *bacterium*_MOT-149_nov_95.946%

SPN96 *Oscillospiraceae*_[G-2] *bacterium*_MOT-149_nov_94.157%

SPN97 *Alistipes senegalensis*_nov_95.228%

SPN98 *Oscillospiraceae*_[G-4] *bacterium*_MOT-151_nov_91.723%

SPN99 *Lachnospiraceae*_[G-11] *bacterium*_MOT-177_nov_97.523%

SPP1 *Clostridium disporicum*_saudiense

SPP2 *Enterobacter asburiae*_cancerogenus_cloacae_hormaechei

SPP3 *Bacteroides acidifaciens acidofaciens*

SPP4 *Lachnospiraceae*_[G-12] *bacterium*_MOT-179_bacterium_MOT-180

SPP5 *Blautia hansenii_hominis_marasmi*

SPPN1 *multigenus multispecies_sppn1_2_nov_95.918%*

SPPN10 *multigenus multispecies_sppn10_2_nov_95.918%*

SPPN11 *multigenus multispecies_sppn11_2_nov_95.465%*

SPPN12 *Alistipes multispecies_sppn12_2_nov_96.304%*

SPPN13 *multigenus multispecies_sppn13_5_nov_94.570%*

SPPN14 *multigenus multispecies_sppn14_2_nov_82.889%*

SPPN15 *Eubacteriales*_[G-1] *multispecies_sppn15_2_nov_97.511%*

SPPN16 *multigenus multispecies_sppn16_2_nov_96.833%*

SPPN17 *multigenus multispecies_sppn17_2_nov_95.928%*

SPPN18 *multigenus multispecies_sppn18_2_nov_92.063%*

SPPN19 *multigenus multispecies_sppn19_3_nov_96.818%*

SPPN2 *Bacteroidetes*_[G-3] *multispecies_sppn2_2_nov_87.554%*

SPPN20 *multigenus multispecies_sppn20_3_nov_95.455%*

SPPN21 *Roseburia multispecies_sppn21_3_nov_95.711%*

SPPN22 *multigenus multispecies_sppn22_2_nov_95.465%*

SPPN23 *multigenus multispecies_sppn23_2_nov_96.818%*

SPPN24 *multigenus multispecies_sppn24_2_nov_93.878%*

SPPN25 *multigenus multispecies_sppn25_3_nov_96.145%*

SPPN26 *Anaerotrignum multispecies_sppn26_2_nov_94.808%*

SPPN27 *multigenus multispecies_sppn27_3_nov_92.665%*