

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 peoriensis
SP31 Lachnospiraceae [G-9] bacterium MOT-174
SP32 Eubacteriales [G-4] bacterium MOT-164
SP33 Erysipelatoclostridium [Clostridium] cocleatum
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 alysoidea nov_95.227%
SPN103 Anaerotruncus rubiinfantis nov_92.760%
SPN104 Muribaculaceae [G-2] bacterium MOT-104 nov_91.991%
SPN105 Mediterraneibacter [Ruminococcus] gnavus nov_93.424%
SPN106 Alistipes putredinis nov_95.887%
SPN107 Oscillibacter valerigenes nov_95.260%
SPN108 Kineothrix alysoidea nov_97.279%
SPN109 Pseudobutyrvibrio 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 Lacrimispora indolis nov_90.724%
SPN113 Kineothrix alysoidea nov_93.651%
SPN114 Lawsonibacter asaccharolyticus nov_97.973%
SPN115 Lachnospiraceae [G-14] bacterium MOT-185 nov_92.358%
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%
SPN12 Oscillospiraceae [G-4] bacterium MOT-151 nov_93.708%
SPN120 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_95.475%
SPN124 Lachnospiraceae [G-11] bacterium MOT-178 nov_97.978%
SPN125 Lachnospiraceae [G-9] bacterium MOT-174 nov_96.388%
SPN126 Lachnospiraceae [G-14] bacterium MOT-185 nov_96.599%
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 Mailhella massiliensis nov_92.094%
SPN133 Anaerotruncus rubiinfantis nov_92.517%
SPN134 Neglectibacter timonensis nov_97.727%
SPN135 Butyricicoccus pullicaecorum nov_94.820%

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 Anaerotignum lactatifermentans nov_97.523%
SPN158 Phocaea massiliensis nov_95.682%
SPN159 Oscillospiraceae [G-2] bacterium MOT-149 nov_96.171%
SPN16 Harryllintia 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 Butyricicoccus pullicaecorum nov_94.382%
SPN167 Kineothrix alysoidea 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 formatexigens 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 rubiinfantis 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 Mageeibacillus indolicus nov_87.668%
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 Harryllintia acetispora nov_93.468%
SPN191 Kineothrix alysoidea 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_86.323%
SPN203 Eubacteriales [G-4] bacterium MOT-165 nov_97.065%
SPN204 Kineothrix alysoidea nov_97.059%
SPN205 Lachnoclostridium [Clostridium] populeti nov_92.986%
SPN206 Kineothrix alysoidea nov_96.136%
SPN207 Anaerocolumna jejuniensis 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%

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_93.0
SPN24 Harryllintia acetispora nov_92.517%
SPN25 Peptococcus sp. HMT_168 nov_89.979%
SPN26 Kineothrix alysoidea 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 Duncaniella freteri nov_93.103%
SPN5 Oscillibacter valerigenes 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 Culturomica massiliensis nov_93.709%
SPN79 Kineothrix alysoidea nov_95.928%
SPN8 Gracilibacter thermotolerans 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 Lachnoclostridium pacaense nov_96.825%
SPN85 Muribaculaceae [G-2] bacterium MOT-104 nov_89.462%
SPN86 Eubacterium coprostanoligenes 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 Pseudoflavonifractor capillosus nov_95.721%
SPN91 Anaerotignum 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 sporicum saudiense
SPP2 Enterobacter asburiae cancerogenus cloacae hormaechei
SPP3 Bacteroides acidifaciens acidifaciens
SPP4 Lachnospiraceae [G-12] bacterium MOT-179 bacterium MOT-18
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%