

## Species

- SP103 Porphyromonas sp.\_MOT-131
- SP104 Pediococcus pentosaceus
- SP106 Streptococcus acidominimus
- SP11 Streptococcus danieliae
- SP110 Eubacteriales\_[G-4] bacterium\_MOT-164
- SP114 Enterococcus faecalis
- SP12 Corynebacterium stationis
- SP121 Mammaliicoccus lentus
- SP129 Bifidobacterium pseudolongum
- SP133 Mollicutes\_[G-1] bacterium\_MOT-186
- SP135 Lactococcus lactis
- SP138 Lactobacillus johnsonii
- SP14 Streptococcus thoraltensis
- SP149 Stenotrophomonas [Pseudomonas] hibiscicola
- SP15 Staphylococcus ureilyticus
- SP168 Weissella cibaria
- SP17 Staphylococcus saprophyticus
- SP172 Anaerococcus sp.\_HMT\_290
- SP2 Triticum aestivum
- SP22 Muribacter sp.\_MOT-143
- SP23 Rodentibacter pneumotropicus
- SP25 Streptococcus thermophilus
- SP29 Moraxella osloensis
- SP3 Enterococcus gallinarum
- SP33 Limosilactobacillus reuteri
- SP34 Staphylococcus equorum
- SP36 Massilia aurea
- SP37 Psychrobacter alimentarius
- SP38 Dubosiella newyorkensis
- SP41 Atopostipes sp.\_MOT-201
- SP42 Ligilactobacillus animalis
- SP43 Lachnospiraceae\_[G-14] bacterium\_MOT-185
- SP44 Ligilactobacillus murinus
- SP46 Eubacteriales\_[G-2] bacterium\_MOT-162
- SP47 Cutibacterium granulosum
- SP48 Corynebacterium mastitidis
- SP50 Cutibacterium acnes
- SP52 Clostridium disporicum
- SP54 Stenotrophomonas maltophilia
- SP56 Streptomyces aculeolatus
- SP57 Lachnospiraceae\_[G-11] bacterium\_MOT-177
- SP64 Pseudomonas helleri
- SP65 Lactocaseibacillus rhamnosus
- SP69 Jeotgalicoccus halotolerans
- SP71 Levilactobacillus brevis
- SP72 Gemella sp.\_MOT-033
- SP73 Corynebacterium ammoniagenes
- SP75 Secundilactobacillus paracollinoides
- SP77 Leptothrix sp.\_HMT\_025
- SP8 Weissella paramesenteroides
- SP84 Delftia acidovorans
- SP87 Carnobacteriaceae\_[G-1] bacterium\_MOT-197
- SP90 Rothia nasimurium
- SP92 Pelomonas saccharophila
- SP96 Ralstonia sp.\_HMT\_406
- SP97 Actinidia eriantha
- SP99 Akkermansia muciniphila
- SPN109 Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_88.105%
- SPN110 Duncaniella freteri\_nov\_93.293%
- SPN122 Turicibacter sanguinis\_nov\_95.923%
- SPN151 Enterococcus faecalis\_nov\_95.825%
- SPN159 Fusobacterium varium\_nov\_96.696%
- SPN188 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_88.867%
- SPN21 Oribacterium parvum\_nov\_89.770%
- SPN226 Duncaniella freteri\_nov\_89.775%
- SPN272 Alistipes senegalensis\_nov\_93.443%
- SPN276 Erysipelotrichaceae\_[G-1] bacterium\_MOT-189\_nov\_88.798%
- SPN283 Peptococcus sp.\_HMT\_168\_nov\_84.866%
- SPN286 Duncaniella freteri\_nov\_89.718%
- SPN29 Fusicatenibacter saccharivorans\_nov\_90.526%
- SPN293 Oscillospiraceae\_[G-6] bacterium\_MOT-153\_nov\_91.631%
- SPN3 Actinidia eriantha\_nov\_97.011%
- SPN302 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.000%
- SPN311 Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_92.719%
- SPN319 Fusobacterium perfoetens\_nov\_91.126%
- SPN47 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_88.423%
- SPN54 Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_85.887%
- SPN66 Duncaniella freteri\_nov\_88.934%
- SPN72 Duncaniella freteri\_nov\_93.699%
- SPP10 Latilactobacillus curvatus\_graminis
- SPP11 Sphingomonas aquatilis\_melonis
- SPP9 Staphylococcus saprophyticus\_xylosus