

Species

- SP12 *Phocaeicola sartorii*
- SP16 *Clostridioides difficile*
- SP20 *Porphyromonas* sp._MOT-131
- SP24 *Adlercreutzia equolifaciens*
- SP25 *Akkermansia muciniphila*
- SP26 *Bifidobacterium pseudolongum*
- SP27 *Ileibacterium valens*
- SP28 *Muribacter* sp._MOT-143
- SP29 *Carnobacteriaceae*_[G-1] bacterium_MOT-197
- SP3 *Ligilactobacillus murinus*
- SP30 *Erysipelotrichaceae*_[G-1] bacterium_MOT-189
- SP34 *Parabacteroides distasonis*
- SP35 *Staphylococcus ureilyticus*
- SP36 *Bacteroides acidifaciens*
- SP39 *Bacteroides uniformis*
- SP40 *Bacteroides caccae*
- SP41 *Enterococcus faecalis*
- SP42 *Lachnospiraceae*_[G-3] bacterium_MOT-168
- SP46 *Weeksellaceae*_[G_1] bacterium_MOT-126
- SP47 *Cutibacterium acnes*
- SP48 *Dubosiella newyorkensis*
- SP54 *Streptococcus acidominimus*
- SP56 *Rodentibacter pneumotropicus*
- SP57 *Eubacteriales*_[G-1] bacterium_MOT-159
- SP6 *Corynebacterium mastitidis*
- SP68 *Parabacteroides goldsteinii*
- SP7 *Streptococcus danieliae*
- SP72 *Streptococcus thoraltensis*
- SP8 *Mammaliococcus lentus*
- SP82 *Lactobacillus johnsonii*
- SP86 *Erysipelatoclostridium* [*Clostridium*] *cocleatum*
- SP9 *Faecalibaculum rodentium*
- SPN106 *Lachnospiraceae*_[G-14] bacterium_MOT-185_nov_96.781%
- SPN116 *Lachnospiraceae*_[G-14] bacterium_MOT-185_nov_95.717%
- SPN117 *Muribaculaceae*_[G-1] bacterium_MOT-129_nov_89.431%
- SPN120 *Anaerostipes caccae*_nov_96.328%
- SPN123 *Lachnospiraceae*_[G-2] bacterium_HMT_096_nov_91.632%
- SPN124 *Duncaniella freteri*_nov_86.263%
- SPN129 *Alistipes putredinis*_nov_93.224%
- SPN14 *Parasutterella excrementihominis*_nov_94.578%
- SPN144 *Glucerbacter canis*_nov_93.305%
- SPN159 *Turicibacter sanguinis*_nov_95.923%
- SPN164 *Muribaculaceae*_[G-2] bacterium_MOT-104_nov_88.755%
- SPN166 *Muribaculaceae*_[G-1] bacterium_MOT-129_nov_85.887%
- SPN172 *Bacteroides uniformis*_nov_95.893%
- SPN185 *Mailhella massiliensis*_nov_90.244%
- SPN201 *Lachnospiraceae*_[G-3] bacterium_MOT-168_nov_94.792%
- SPN218 *Muribaculaceae*_[G-2] bacterium_MOT-104_nov_89.200%
- SPN226 *Ihubacter massiliensis*_nov_94.572%
- SPN235 *Eggerthella timonensis*_nov_90.929%
- SPN237 *Parafannyhessea umbonata*_nov_92.161%
- SPN247 *Lachnospiraceae*_[G-3] bacterium_MOT-168_nov_96.050%
- SPN256 *Duncaniella freteri*_nov_88.531%
- SPN28 *Oscillospiraceae*_[G-6] bacterium_MOT-153_nov_86.475%
- SPN38 *Alistipes putredinis*_nov_92.418%
- SPN48 *Duncaniella freteri*_nov_89.775%
- SPN58 *Eubacterium ventriosum*_nov_93.320%
- SPN6 *Duncaniella freteri*_nov_86.061%
- SPN64 *Alistipes senegalensis*_nov_93.443%
- SPN65 *Anaeroplasma abactoclasticum*_nov_86.538%
- SPN75 *Mucispirillum schaedleri*_nov_93.265%
- SPN85 *Duncaniella freteri*_nov_87.298%
- SPP1 *Staphylococcus saprophyticus_xylosus*
- SPPN1 *Olsenella multispecies_sppn1_2_nov_91.966%*