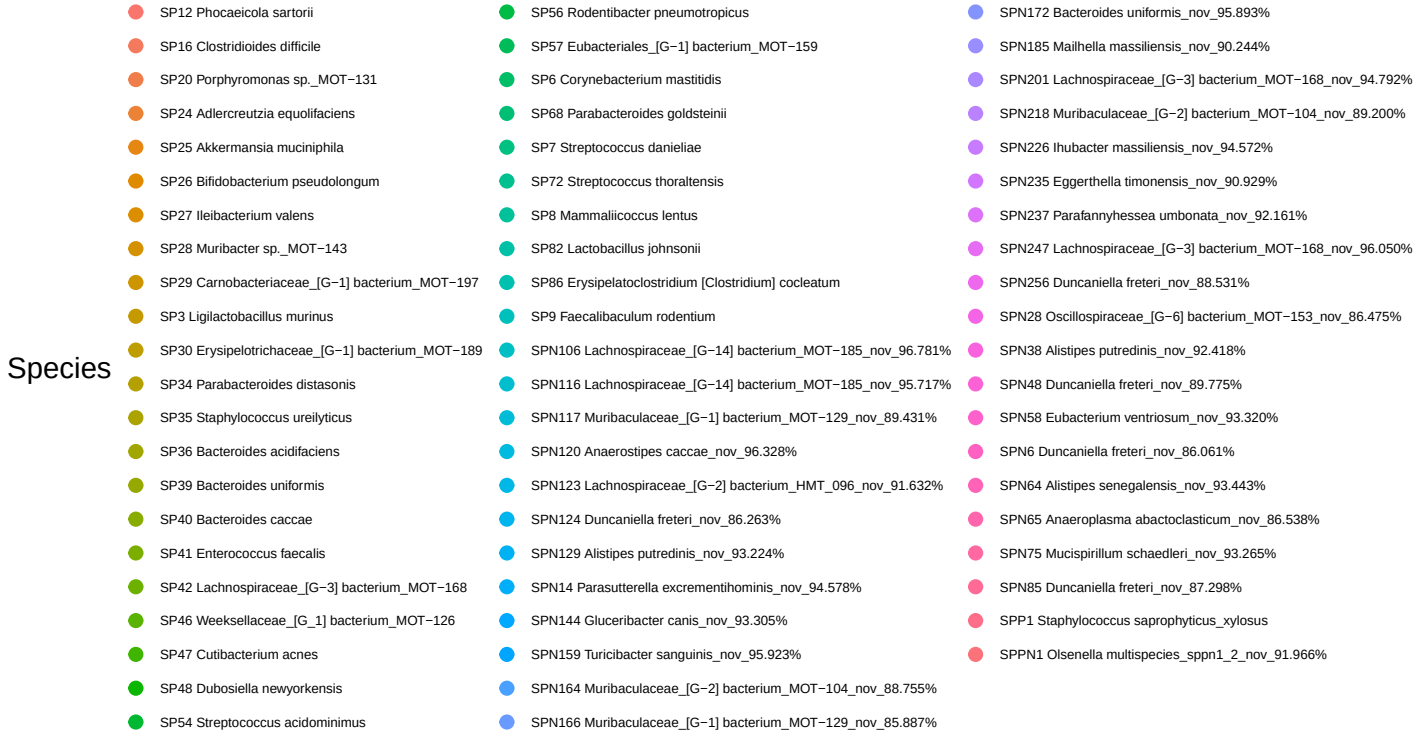

Species
SP12 Phocaeicola sartorii
SP16 Clostridioides difficile
SP20 Porphyromonas sp._MOT−131
SP24 Adlercreutzia equolifaciens
SP25 Akkermansia muciniphila
SP26 Bifidobacterium pseudolongum
SP27 Ileibacterium valens
SP28 Muribacter sp._MOT−143
SP29 Carnobacteriaceae_[G−1] bacterium_MOT−197
SP3 Ligilactobacillus 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 Mammaliicoccus 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 Gluceribacter 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%