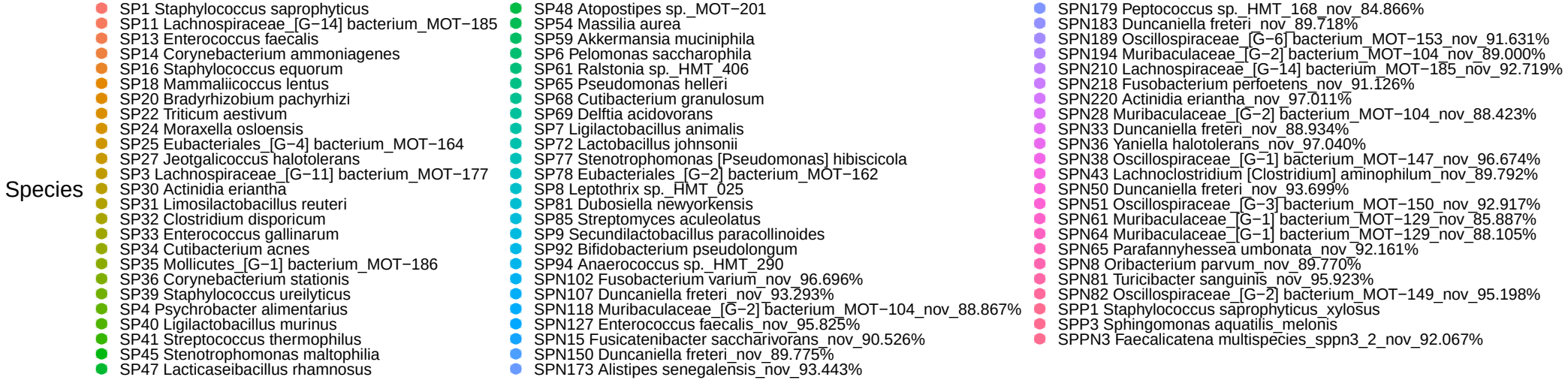
Species
SP1 Staphylococcus saprophyticus
SP11 Lachnospiraceae_[G−14] bacterium_MOT−185
SP13 Enterococcus faecalis
SP14 Corynebacterium ammoniagenes
SP16 Staphylococcus equorum
SP18 Mammaliicoccus lentus
SP20 Bradyrhizobium pachyrhizi
SP22 Triticum aestivum
SP24 Moraxella osloensis
SP25 Eubacteriales_[G−4] bacterium_MOT−164
SP27 Jeotgalicoccus halotolerans
SP3 Lachnospiraceae_[G−11] bacterium_MOT−177
SP30 Actinidia eriantha
SP31 Limosilactobacillus reuteri
SP32 Clostridium disporicum
SP33 Enterococcus gallinarum
SP34 Cutibacterium acnes
SP35 Mollicutes_[G−1] bacterium_MOT−186
SP36 Corynebacterium stationis
SP39 Staphylococcus ureilyticus
SP4 Psychrobacter alimentarius
SP40 Ligilactobacillus murinus
SP41 Streptococcus thermophilus
SP45 Stenotrophomonas maltophilia
SP47 Lacticaseibacillus rhamnosus
SP48 Atopostipes sp._MOT−201
SP54 Massilia aurea
SP59 Akkermansia muciniphila
SP6 Pelomonas saccharophila
SP61 Ralstonia sp._HMT_406
SP65 Pseudomonas helleri
SP68 Cutibacterium granulosum
SP69 Delftia acidovorans
SP7 Ligilactobacillus animalis
SP72 Lactobacillus johnsonii
SP77 Stenotrophomonas [Pseudomonas] hibiscicola
SP78 Eubacteriales_[G−2] bacterium_MOT−162
SP8 Leptothrix sp._HMT_025
SP81 Dubosiella newyorkensis
SP85 Streptomyces aculeolatus
SP9 Secundilactobacillus paracollinoides
SP92 Bifidobacterium pseudolongum
SP94 Anaerococcus sp._HMT_290
SPN102 Fusobacterium varium_nov_96.696%
SPN107 Duncaniella freteri_nov_93.293%
SPN118 Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.867%
SPN127 Enterococcus faecalis_nov_95.825%
SPN15 Fusicatenibacter saccharivorans_nov_90.526%
SPN150 Duncaniella freteri_nov_89.775%
SPN173 Alistipes senegalensis_nov_93.443%
SPN179 Peptococcus sp._HMT_168_nov_84.866%
SPN183 Duncaniella freteri_nov_89.718%
SPN189 Oscillospiraceae_[G−6] bacterium_MOT−153_nov_91.631%
SPN194 Muribaculaceae_[G−2] bacterium_MOT−104_nov_89.000%
SPN210 Lachnospiraceae_[G−14] bacterium_MOT−185_nov_92.719%
SPN218 Fusobacterium perfoetens_nov_91.126%
SPN220 Actinidia eriantha_nov_97.011%
SPN28 Muribaculaceae_[G−2] bacterium_MOT−104_nov_88.423%
SPN33 Duncaniella freteri_nov_88.934%
SPN36 Yaniella halotolerans_nov_97.040%
SPN38 Oscillospiraceae_[G−1] bacterium_MOT−147_nov_96.674%
SPN43 Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
SPN50 Duncaniella freteri_nov_93.699%
SPN51 Oscillospiraceae_[G−3] bacterium_MOT−150_nov_92.917%
SPN61 Muribaculaceae_[G−1] bacterium_MOT−129_nov_85.887%
SPN64 Muribaculaceae_[G−1] bacterium_MOT−129_nov_88.105%
SPN65 Parafannyhessea umbonata_nov_92.161%
SPN8 Oribacterium parvum_nov_89.770%
SPN81 Turicibacter sanguinis_nov_95.923%
SPN82 Oscillospiraceae_[G−2] bacterium_MOT−149_nov_95.198%
SPP1 Staphylococcus saprophyticus_xylosus
SPP3 Sphingomonas aquatilis_melonis
SPPN3 Faecalicatena multispecies_sppn3_2_nov_92.067%