

Species

- SP100 *Lactobacillus johnsonii*
- SP11 *Bacteroides acidifaciens*
- SP14 *Sporosarcina* sp._MOT-205
- SP15 *Enterococcus hirae*
- SP16 *Mammaliicoccus lentus*
- SP18 *Jeotgalicoccus halotolerans*
- SP19 *Sporosarcina* sp._MOT-204
- SP2 *Ligilactobacillus murinus*
- SP20 *Corynebacterium mastitidis*
- SP21 *Streptococcus acidominimus*
- SP22 *Bradyrhizobium pachyrhizi*
- SP23 *Delftia acidovorans*
- SP25 *Muribacter* sp._MOT-143
- SP26 *Lactococcus lactis*
- SP3 *Streptococcus danieliae*
- SP33 *Staphylococcus ureilyticus*
- SP34 *Rodentibacter pneumotropicus*
- SP4 *Limosilactobacillus reuteri*
- SP42 *Mammaliicoccus sciuri*
- SP44 *Acinetobacter lwoffii*
- SP47 *Atopostipes* sp._MOT-201
- SP5 *Acinetobacter johnsonii*
- SP54 *Staphylococcus saprophyticus*
- SP56 *Bifidobacterium pseudolongum*
- SP57 *Erysipelotrichaceae*_[G-1] bacterium_MOT-189
- SP59 *Pediococcus pentosaceus*
- SP64 *Corynebacterium stationis*
- SP74 *Porphyromonas* sp._MOT-131
- SP76 *Enterococcus faecalis*
- SP8 *Atopostipes* sp._MOT-202
- SP81 *Streptococcus thoraltensis*
- SP89 *Helicobacter ganmani*
- SP9 *Faecalibaculum rodentium*
- SP93 *Weeksellaceae*_[G_1] bacterium_MOT-126
- SP99 *Muribacter muris*
- SPN105 *Savagea faecisuis*_nov_92.585%
- SPN110 *Clostridium collagenovorans*_nov_80.258%
- SPN142 *Gloeobacter kilaeensis*_nov_84.668%
- SPN144 *Sporosarcina* sp._MOT-205_nov_95.565%
- SPN156 *Prevotella* sp._HMT_820_nov_89.980%
- SPN165 *Kineothrix alysoides*_nov_86.667%
- SPN176 *Muribaculaceae*_[G-1] bacterium_MOT-129_nov_89.431%
- SPN187 *Lachnospiraceae*_[G-11] bacterium_MOT-177_nov_96.066%
- SPN198 *Muribaculaceae*_[G-1] bacterium_MOT-129_nov_83.636%
- SPN212 *Lachnospiraceae*_[G-12] bacterium_MOT-180_nov_89.613%
- SPN223 *Prevotella shahii*_nov_86.667%
- SPN233 *Lachnospiraceae*_[G-7] bacterium_MOT-172_nov_93.075%
- SPN3 *Caproicibacter fermentans*_nov_89.662%
- SPN32 *Prevotella* sp._HMT_317_nov_90.041%
- SPP1 *Staphylococcus edaphicus*_saprophyticus
- SPP3 *Staphylococcus saprophyticus_xylosus*
- SPP5 *Pasteurella_Rodentibacter caecimuris_heylii*
- SPPN4 *Gloeobacter multispecies_sppn4_2_nov_83.144%*