

Species

SP100 <i>Lactobacillus johnsonii</i>	SP42 <i>Mammaliococcus sciuri</i>	SPN110 <i>Clostridium collagenovorans</i> _nov_80.258%
SP11 <i>Bacteroides acidifaciens</i>	SP44 <i>Acinetobacter lwoffii</i>	SPN142 <i>Gloeobacter kilaueensis</i> _nov_84.668%
SP14 <i>Sporosarcina</i> sp._MOT-205	SP47 <i>Atopostipes</i> sp._MOT-201	SPN144 <i>Sporosarcina</i> sp._MOT-205_nov_95.565%
SP15 <i>Enterococcus hirae</i>	SP5 <i>Acinetobacter johnsonii</i>	SPN156 <i>Prevotella</i> sp._HMT_820_nov_89.980%
SP16 <i>Mammaliococcus lentus</i>	SP54 <i>Staphylococcus saprophyticus</i>	SPN165 <i>Kineothrix alysoidea</i> _nov_86.667%
SP18 <i>Jeotgaliococcus halotolerans</i>	SP56 <i>Bifidobacterium pseudolongum</i>	SPN176 <i>Muribaculaceae</i> _ [G-1] bacterium_MOT-129_nov_89.431%
SP19 <i>Sporosarcina</i> sp._MOT-204	SP57 <i>Erysipelotrichaceae</i> _ [G-1] bacterium_MOT-189	SPN187 <i>Lachnospiraceae</i> _ [G-11] bacterium_MOT-177_nov_96.066%
SP2 <i>Ligilactobacillus murinus</i>	SP59 <i>Pediococcus pentosaceus</i>	SPN198 <i>Muribaculaceae</i> _ [G-1] bacterium_MOT-129_nov_83.636%
SP20 <i>Corynebacterium mastitidis</i>	SP64 <i>Corynebacterium stationis</i>	SPN212 <i>Lachnospiraceae</i> _ [G-12] bacterium_MOT-180_nov_89.613%
SP21 <i>Streptococcus acidominimus</i>	SP74 <i>Porphyromonas</i> sp._MOT-131	SPN223 <i>Prevotella shahii</i> _nov_86.667%
SP22 <i>Bradyrhizobium pachyrhizi</i>	SP76 <i>Enterococcus faecalis</i>	SPN233 <i>Lachnospiraceae</i> _ [G-7] bacterium_MOT-172_nov_93.075%
SP23 <i>Delftia acidovorans</i>	SP8 <i>Atopostipes</i> sp._MOT-202	SPN3 <i>Caproicibacter fermentans</i> _nov_89.662%
SP25 <i>Muribacter</i> sp._MOT-143	SP81 <i>Streptococcus thoraltensis</i>	SPN32 <i>Prevotella</i> sp._HMT_317_nov_90.041%
SP26 <i>Lactococcus lactis</i>	SP89 <i>Helicobacter ganmani</i>	SPP1 <i>Staphylococcus edaphicus_saprophyticus</i>
SP3 <i>Streptococcus danieliae</i>	SP9 <i>Faecalibaculum rodentium</i>	SPP3 <i>Staphylococcus saprophyticus_xylosum</i>
SP33 <i>Staphylococcus ureilyticus</i>	SP93 <i>Weeksellaceae</i> _ [G_1] bacterium_MOT-126	SPP5 <i>Pasteurella_Rodentibacter caecimuris_heylii</i>
SP34 <i>Rodentibacter pneumotropicus</i>	SP99 <i>Muribacter muris</i>	SPN4 <i>Gloeobacter multispecies_sppn4_2_nov_83.144%</i>
SP4 <i>Limosilactobacillus reuteri</i>	SPN105 <i>Savagea faecisuis</i> _nov_92.585%	