

Species

- SP103 *Staphylococcus caprae*
- SP104 *Epilithonimonas hominis*
- SP106 *Atlantibacter hermannii*
- SP11 *Rhodococcus qingshengii*
- SP111 *Priestia megaterium*
- SP114 *Staphylococcus warneri*
- SP117 *Nocardia coeliaca*
- SP12 *Sphingomonas aquatica*
- SP126 *Raoultella planticola*
- SP13 *Klebsiella pneumoniae*
- SP131 *Comamonas aquatilis*
- SP134 *Acinetobacter lwoffii*
- SP14 *Ligilactobacillus murinus*
- SP141 *Klebsiella quasipneumoniae*
- SP143 *Bacteroides thetaiotaomicron*
- SP144 *Parasutterella excrementihominis*
- SP149 *Enterobacter kobei*
- SP15 *Comamonas testosteroni*
- SP155 *Massilia pinisoli*
- SP16 *Faecalibaculum rodentium*
- SP17 *Microbacterium maritopicum*
- SP18 *Stenotrophomonas maltophilia*
- SP19 *Cutibacterium acnes*
- SP2 *Streptococcus danieliae*
- SP21 *Bacillus subtilis*
- SP23 *Enterococcus gallinarum*
- SP26 *Enterobacter hormaechei*
- SP27 *Agrobacterium vitis*
- SP28 *Sediminibacterium aquarii*
- SP3 *Pseudomonas putida*
- SP33 *Mammaliicoccus sciuri*
- SP35 *Corynebacterium mucifaciens*
- SP36 *Bifidobacterium pseudolongum*
- SP39 *Acinetobacter radioresistens*
- SP4 *Acinetobacter johnsonii*
- SP41 *Enterobacter mori*
- SP44 *Bradyrhizobium lupini*
- SP5 *Bacillus halotolerans*
- SP50 *Acidovorax temperans*
- SP51 *Sphingomonas yabuuchiae*
- SP58 *Lysinibacillus sphaericus*
- SP59 *Enterococcus casseliflavus*
- SP6 *Leucobacter chromiirestiens*
- SP60 *Acinetobacter haemolyticus*
- SP62 *Sphingobacterium multivorum*
- SP63 *Klebsiella oxytoca*
- SP67 *Brochothrix thermosphacta*
- SP68 *Robinsoniella peoriensis*
- SP69 *Magnetospirillum magnetotacticum*
- SP74 *Streptococcus thoraltensis*
- SP77 *Mammaliicoccus lentus*
- SP78 *Brachybacterium conglomeratum*
- SP79 *Chryseobacterium gambrini*
- SP82 *Enterococcus faecalis*
- SP83 *Citrobacter amalonaticus*
- SP84 *Acidovorax monticola*
- SP85 *Staphylococcus ureilyticus*
- SP86 *Actinidia eriantha*
- SP89 *Corynebacterium macginleyi*
- SP9 *Priestia aryabhatai*
- SP95 *Citrobacter koseri*
- SP97 *Klebsiella aerogenes*
- SP99 *Staphylococcus hominis*
- SPN10 *Sediminibacterium aquarii_nov_92.355%*
- SPN13 *Chryseobacterium yeoncheonense_nov_97.484%*
- SPN20 *Anaerostipes caccae_nov_96.328%*
- SPN24 *Lysinibacillus sphaericus_nov_97.988%*
- SPN31 *Siccibacter turicensis_nov_97.955%*
- SPN42 *Massilia agri_nov_97.713%*
- SPN51 *Enterobacter mori_nov_97.951%*
- SPN52 *Acinetobacter johnsonii_nov_97.732%*
- SPN68 *Thermodesulfobium acidophilum_nov_80.255%*
- SPN69 *Ralstonia solanacearum_nov_96.296%*
- SPN70 *Acinetobacter johnsonii_nov_97.737%*
- SPP1 *Staphylococcus argenteus_aureus_roterodami*
- SPP12 *Pseudomonas cedrina_lactis*
- SPP13 *Staphylococcus capitis_epidermidis*
- SPP14 *Sphingomonas aquatilis_melonis*
- SPP15 *Bradyrhizobium archetypum_australiense_cajani_japonicum_liaoningense...(8 species)*
- SPP17 *Bradyrhizobium cajani_japonicum_liaoningense_lupini*
- SPP19 *Staphylococcus saprophyticus_xylosum*
- SPP5 *Acinetobacter calcoaceticus_pittii*
- SPPN3 *Gloeobacter multispecies_sppn3_2_nov_83.592%*
- SPPN4 *Terrimonas multispecies_sppn4_2_nov_90.928%*