

## Species

SP101 <i>Enterobacter mori</i>	SP44 <i>Eubacteriales</i> _[G-3] <i>bacterium</i> _MOT-163	SPN223 <i>Blautia producta</i> _nov_91.152%
SP103 <i>Granulicatella adiacens</i>	SP47 <i>Enterococcus haemoperoxidus</i>	SPN235 <i>Blautia faecicola</i> _nov_93.887%
SP107 <i>Leifsonia aquatica</i>	SP48 <i>Bifidobacterium breve</i>	SPN247 <i>Saccharibacteria</i> _(TM7)_[G-3] <i>bacterium</i> _HMT_351_nov_96.074%
SP11 <i>Mammaliicoccus sciuri</i>	SP5 <i>Acinetobacter bereziniae</i>	SPN259 <i>Ruminococcus albus</i> _nov_94.425%
SP115 <i>Atopostipes</i> sp._MOT-201	SP58 <i>Phocaeicola vulgatus</i>	SPN26 <i>Blautia schinkii</i> _nov_95.124%
SP116 <i>Klebsiella grimontii</i>	SP6 <i>Romboutsia ilealis</i>	SPN260 <i>Adlercreutzia equolifaciens</i> _nov_92.569%
SP122 <i>Escherichia marmotae</i>	SP60 <i>Gordonia polyisoprenivorans</i>	SPN272 <i>Duncaniella freteri</i> _nov_92.077%
SP123 <i>Cutibacterium acnes</i>	SP63 <i>Streptococcus parasanguinis</i>	SPN277 <i>Lachnoclostridium</i> [ <i>Clostridium</i> ] <i>scindens</i> _nov_94.719%
SP127 <i>Enterobacter ludwigii</i>	SP64 <i>Streptococcus mitis</i>	SPN284 <i>Lachnospiraceae</i> _[G-3] <i>bacterium</i> _MOT-168_nov_97.598%
SP131 <i>Erysipelatoclostridium</i> [ <i>Clostridium</i> ] <i>innocuum</i>	SP7 <i>Rothia nasimurium</i>	SPN296 <i>Arthrospira platensis</i> _nov_90.832%
SP136 <i>Erysipelatoclostridium</i> [ <i>Clostridium</i> ] <i>saccharogumia</i>	SP70 <i>Streptococcus</i> sp._MOT-045	SPN308 <i>Sporobacter termitidis</i> _nov_94.306%
SP146 <i>Neisseria elongata</i>	SP72 <i>Streptococcus mutans</i>	SPN320 <i>Muribaculaceae</i> _[G-2] <i>bacterium</i> _MOT-104_nov_89.925%
SP15 <i>Staphylococcus ureilyticus</i>	SP75 <i>Bifidobacterium pseudolongum</i>	SPN33 <i>Muribaculaceae</i> _[G-1] <i>bacterium</i> _MOT-129_nov_89.815%
SP152 <i>Mammaliicoccus lentus</i>	SP76 <i>Blautia glucerasea</i>	SPN334 <i>Qipengyuania sediminis</i> _nov_97.809%
SP155 <i>Streptococcus agalactiae</i>	SP80 <i>Bacteroides uniformis</i>	SPN346 <i>Faecalimonas umbilicata</i> _nov_93.562%
SP16 <i>Lactobacillus intestinalis</i>	SP81 <i>Escherichia coli</i>	SPN358 <i>Muribaculaceae</i> _[G-1] <i>bacterium</i> _MOT-129_nov_91.040%
SP165 <i>Enterobacter hormaechei</i>	SP82 <i>Limosilactobacillus reuteri</i>	SPN369 <i>Anaerocolumna aminovalerica</i> _nov_93.182%
SP168 <i>Acinetobacter courvalinii</i>	SP83 <i>Faecalibaculum rodentium</i>	SPN38 <i>Sporosarcina</i> sp._MOT-205_nov_97.019%
SP170 <i>Extibacter muris</i>	SP84 <i>Coprobaillus cateniformis</i>	SPN381 <i>Blautia schinkii</i> _nov_95.261%
SP174 <i>Hungatella hathewayi</i>	SP86 <i>Enterococcus avium</i>	SPN382 <i>Eubacteriales</i> _[G-4] <i>bacterium</i> _MOT-165_nov_93.797%
SP178 <i>Brachybacterium conglomeratum</i>	SP90 <i>Proteus mirabilis</i>	SPN398 <i>Thermacetogenium phaeum</i> _nov_88.464%
SP18 <i>Corynebacterium stationis</i>	SP92 <i>Citrobacter koseri</i>	SPN482 <i>Muribaculaceae</i> _[G-1] <i>bacterium</i> _MOT-129_nov_91.102%
SP182 <i>Monoglobus pectinilyticus</i>	SP93 <i>Providencia rettgeri</i>	SPN494 <i>Muribaculaceae</i> _[G-2] <i>bacterium</i> _MOT-104_nov_91.473%
SP186 <i>Akkermansia muciniphila</i>	SP95 <i>Yaniella halotolerans</i>	SPN50 <i>Mobilitalea sibirica</i> _nov_95.271%
SP19 <i>Clostridium disporicum</i>	SP97 <i>Rodentibacter rattii</i>	SPN506 <i>Sporosarcina</i> sp._MOT-204_nov_95.599%
SP194 <i>Enterococcus faecalis</i>	SP98 <i>Staphylococcus equorum</i>	SPN518 <i>Acetivibrio cellulolyticus</i> _nov_92.505%
SP195 <i>Streptococcus acidominimus</i>	SP99 <i>Veillonella parvula</i>	SPN519 <i>Lederbergia lenta</i> _nov_95.190%
SP198 <i>Lactobacillus johnsonii</i>	SPN1 <i>Vallitalea pronyensis</i> _nov_88.227%	SPN541 <i>Eubacteriales</i> _[G-4] <i>bacterium</i> _MOT-165_nov_96.886%
SP2 <i>Enterococcus gallinarum</i>	SPN111 <i>Streptococcus</i> sp._MOT-045_nov_97.609%	SPN553 <i>Ileibacterium valens</i> _nov_94.552%
SP20 <i>Staphylococcus epidermidis</i>	SPN112 <i>Rubrivivax gelatinosus</i> _nov_96.350%	SPN565 <i>Anaerostipes caccae</i> _nov_97.708%
SP200 <i>Corynebacterium lowii</i>	SPN124 <i>Frisingicoccus caecimuris</i> _nov_97.456%	SPN62 <i>Eubacterium coprostanoligenes</i> _nov_94.236%
SP209 <i>Streptococcus cristatus</i>	SPN13 <i>Globicatella sanguinis</i> _nov_93.157%	SPN74 <i>Neglectibacter timonensis</i> _nov_96.238%
SP21 <i>Haemophilus parainfluenzae</i>	SPN136 <i>Turicibacter sanguinis</i> _nov_97.503%	SPN86 <i>Eubacteriales</i> _[G-4] <i>bacterium</i> _MOT-165_nov_92.483%
SP22 <i>Aerococcus urinaeequi</i>	SPN137 <i>Lysinibacter cavernae</i> _nov_97.439%	SPN98 <i>Eubacterium coprostanoligenes</i> _nov_94.452%
SP23 <i>Jeotgalicoccus halotolerans</i>	SPN14 <i>Ruminococcus champanellensis</i> _nov_93.519%	SPP1 <i>Staphylococcus capitis</i> _epidermidis
SP24 <i>Staphylococcus haemolyticus</i>	SPN149 <i>Blautia faecicola</i> _nov_94.993%	SPP11 <i>Escherichia</i> _Shigella <i>fergusonii</i> _flexneri
SP28 <i>Ligilactobacillus murinus</i>	SPN155 <i>Murimonas intestini</i> _nov_94.767%	SPP2 <i>Staphylococcus argenteus</i> _aureus_roterodami
SP3 <i>Enterococcus faecium</i>	SPN161 <i>Prevotella loescheii</i> _nov_92.529%	SPP6 <i>Shigella flexneri</i> _sonnei
SP33 <i>Corynebacterium amycolatum</i>	SPN173 <i>Eubacteriales</i> _[G-4] <i>bacterium</i> _MOT-165_nov_92.892%	SPP7 <i>Staphylococcus saprophyticus</i> _xylosus