

Species

SP102 *Mediterraneibacter* [Ruminococcus] lactaris
SP103 *Bifidobacterium* scardovii
SP104 *Bifidobacterium* longum
SP105 *Prevotella* bivia
SP106 *Finexgoldia* magna
SP108 *Gemella* haemolysans
SP109 *Collinsella* aerofaciens
SP110 *Peptoniphilus* grossensis
SP111 *Anaerococcus* octavius
SP118 *Veillonella* atypica
SP119 *Solobacterium* moorei
SP120 *Serratia* nematodiphila
SP121 *Bacteroides* fragilis
SP122 *Clostridium* chromiireducens
SP123 *Phocaeicola* vulgatus
SP124 *Staphylococcus* saccharolyticus
SP125 *Clostridium* paraputrificum
SP126 *Anaerococcus* vaginalis
SP127 *Clostridium* tertium
SP130 *Gemella* sanguinis
SP131 *Schaalia* sp._HMT_172
SP134 *Rothia* aeria
SP135 *Granulicatella* adiacens
SP136 *Haemophilus* parainfluenzae
SP137 *Peptostreptococcus* anaerobius
SP138 *Peptoniphilus* lacrimalis
SP139 *Anaerococcus* lactolyticus
SP140 *Paratractidigestivibacter* faecalis
SP144 *Stutzerimonas* [Pseudomonas] urumqiensis
SP145 *Bacteroides* faecis
SP146 *Mediterraneibacter* [Ruminococcus] gnavus
SP147 *Cutibacterium* acnes
SP149 *Staphylococcus* lugdunensis
SP150 *Corynebacterium* amycolatum
SP152 *Serratia* surfactantfaciens
SP153 *Capnocytophaga* sputigena
SP154 *Prevotella* corporis
SP155 *Acinetobacter* baumannii
SP156 *Veillonella* dispar
SP157 *Gemella* morbillorum
SP159 *Porphyromonas* pasteri
SP160 *Citrobacter* koseri
SP162 *Lactobacillus* gasseri
SP173 *Faecalibacterium* prausnitzii
SP176 *Pinus* sylvestris
SP179 *Prevotella* timonensis
SP182 *Bifidobacterium* bifidum
SP183 *Rothia* dentocariosa
SP184 *Bergeyella* sp._HMT_931
SP186 *Parabacteroides* distasonis
SP187 *Pantoea* conspicua
SP189 *Actinomyces* oris
SP192 *Prevotella* melaninogenica
SP193 *Bacteroides* thetaiotaomicron
SP194 *Peptoniphilus* coxii
SP195 *Actinomyces* urogenitalis
SP196 *Schaalia* odontolyticus
SP197 *Varibaculum* anthropi
SP198 *Parvimonas* micra
SP199 *Phocaeicola* dorei
SP2 *Veillonella* parvula
SP203 *Streptococcus* sanguinis
SP204 *Stenotrophomonas* maltophilia
SP205 *Varibaculum* cambriense
SP206 *Oribacterium* sinus
SP211 *Clostridium* butyricum
SP212 *Peptostreptococcus* stomatis
SP217 *Granulicatella* elegans
SP219 *Saccharibacteria*_(TM7)_[G-3] bacterium_HMT_351
SP22 *Streptococcus* sp._HMT_074

SP229 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_352
SP234 *Staphylococcus* simulans
SP235 *Anaerococcus* obesiensis
SP238 *Neisseria* flavescens
SP239 *Peptostreptococcaceae*_[XII][G-6] [Eubacterium]_nodatum
SP240 *Limosilactobacillus* mucosae
SP243 *Serratia* marcescens
SP244 *Actinomyces* massiliensis
SP245 *Klebsiella* variicola
SP250 *Eggerthella* lenta
SP252 *Erysipelatoclostridium* [Clostridium]_innocuum
SP257 *Gardnerella* vaginalis
SP265 *Prevotella* histicola
SP266 *Anaerococcus* sp._HMT_294
SP284 *Corynebacterium* kroppenstedtii
SP287 *Hymenobacter* rigui
SP288 *Bifidobacterium* dentium
SP294 *Peptostreptococcaceae*_[XII][G-1] [Eubacterium]_sulci
SP295 *Stomatobaculum* sp._HMT_097
SP298 *Corynebacterium* pilbarens
SP305 *Saccharibacteria*_(TM7)_[G-6] bacterium_HMT_870
SP307 *Dermabacter* hominis
SP309 *Peptostreptococcaceae*_[XII][G-9] [Eubacterium]_brachy
SP312 *Capnocytophaga* leadbetteri
SP325 *Actinomyces* israelii
SP327 *Actinomyces* ihuae
SP329 *Veillonella* denticariosi
SP341 *Bacteroides* caccae
SP342 *Leptotrichia* sp._HMT_215
SP344 *Faecalimonas* umblicata
SP353 *Oribacterium* parvum
SP354 *Porphyromonas* endodontalis
SP357 *Winkia* neuii
SP359 *Actinomyces* naeslundii
SP365 *Ruminococcaceae*_[G-1] bacterium_HMT_075
SP367 *Cutibacterium* avidum
SP370 *Leptotrichia* sp._HMT_392
SP400 *Bacteroides* stercoris
SP41 *Veillonella* rogosa
SP42 *Anaerococcus* hydrogenalis
SP43 *Anaerostipes* caccae
SP430 *Corynebacterium* durum
SP44 *Schaalia* lingnae [Not_Validly_Published]
SP45 *Proteus* mirabilis
SP451 *Ruminococcaceae*_[G-2] bacterium_HMT_085
SP458 *Bifidobacterium* wadsworthia
SP49 *Klebsiella* pneumoniae
SP51 *Rothia* mucilaginosa
SP52 *Enterobacter* bugandensis
SP53 *Enterococcus* faecalis
SP54 *Fusobacterium* nucleatum
SP55 *Streptococcus* parasanguinis_clade_411
SP56 *Staphylococcus* capitis
SP58 *Clostridium* perfringens
SP63 *Staphylococcus* hominis
SP64 *Streptococcus* lactarius
SP65 *Streptococcus* peroris
SP66 *Megamonas* funiformis
SP67 *Enterobacter* mori
SP68 *Streptococcus* sp._HMT_056
SP75 *Staphylococcus* haemolyticus
SP79 *Abiotrophia* defectiva
SP80 *Veillonella* sp._HMT_780
SP81 *Blautia* wexlerae
SP82 *Stenotrophomonas* geniculata
SP87 *Lautropia* mirabilis
SP88 *Raoultella* planticola
SP89 *Lawsonella* clevelandensis
SP90 *Bifidobacterium* breve
SP95 *Actinomyces* graevenitzi

SPN133 *Peptoniphilus* sp._HMT_375_nov_87.810%
SPN178 *Gemella* haemolysans_nov_97.425%
SPN18 *Enterococcus* faecalis_nov_96.788%
SPN204 *Gemella* haemolysans_nov_94.030%
SPN233 *Peptoniphilus* coxii_nov_97.955%
SPN27 *Veillonella* atypica_nov_96.137%
SPN301 *Clostridium* saccharobutylicum_nov_96.110%
SPN329 *Clostridium* chromiireducens_nov_97.959%
SPN36 *Enterococcus* faecalis_nov_89.507%
SPN45 *Mediterraneibacter* [Ruminococcus] gnavus_nov_92.601%
SPN496 *Megamonas* funiformis_nov_97.849%
SPN525 *Clostridium* saccharobutylicum_nov_96.606%
SPN56 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_346_nov_97.964%
SPN65 *Clostridium* tarantellae_nov_97.052%
SPN86 *Clostridium* saccharobutylicum_nov_97.052%
SPN9 *Rothia* mucilaginosa_nov_97.991%
SPP1 *Schaalia* odontolyticus_sp._HMT_180
SPP10 *Streptococcus* parasanguinis_clade_411_sinensis_sp._HMT_056
SPP121 *Streptococcus* australis_rubneri_sp._HMT_066
SPP127 *Klebsiella* pneumoniae_quasipneumoniae
SPP141 *Staphylococcus* capitis_caprae
SPP142 *Staphylococcus* capitis_caprae_epidermidis
SPP148 *Enterobacter* asburiae_cancerogenus_cloacae_hormaechei
SPP149 *Streptococcus* cristatus_cristatus_clade_578
SPP155 *Streptococcus* lactarius_peroris
SPP16 *Streptococcus* cristatus_cristatus_clade_578_downii_gwangjuense_i ...(18 species)
SPP161 *Streptococcus* salivarius_vestibularis
SPP165 *Escherichia*_Shigella_coli_fergusonii_flexneri_sonnei
SPP17 *Streptococcus* cristatus_downii_gwangjuense_infantis_infantis_cla ...(17 species)
SPP172 *Enterococcus* casseliflavus_gallinarum
SPP18 *Hoeflea*_Mesorhizobium_camelthorni_halophila
SPP180 *Staphylococcus* argenteus_aureus_roterodami_simiae
SPP182 *Bacillus* inaquosorum_spizizenii_subtilis_tequilensis
SPP19 *Anaerococcus* prevotii_tetradius
SPP21 *Anaerococcus* mediterraneensis_murdochii
SPP25 *Bifidobacterium* adolescentis_faecale
SPP28 *Peptoniphilus* gorbachii_lacydonensis_sp._HMT_187
SPP33 *Streptococcus* parasanguinis_clade_411_sp._HMT_057
SPP37 *Fusobacterium* nucleatum_nucleatum_subsp_vincentii
SPP39 *Streptococcus* parasanguinis_parasanguinis_clade_721_sp._HMT_066
SPP41 *Atopobium*_Lancefieldella_parvula_parvulum
SPP42 *Colibacter*_Megaspheera_massiliensis_sp._HMT_123
SPP44 *Atopobium*_Fannyhessea_vaginae
SPP46 *Pseudomonas* oryzihabitans_psychrotolerans
SPP47 *Clostridium* disporicum_saudiense
SPP50 *Lactococcus* cremoris_lactis
SPP52 *Fusobacterium* canifelinum_nucleatum_nucleatum_subsp_polymorphum
SPP53 *Hungateella* effluvii_hathewayi
SPP54 *Bifidobacterium* catenulatum_pseudocatenulatum
SPP55 *Olsenella* sp._HMT_939_urinifantis
SPP56 *Streptococcus* australis_rubneri_sp._HMT_066_sp._HMT_074
SPP61 *Staphylococcus* argensis_pettenkoferi
SPP63 *Alipia*_Bradyrhizobium_archetypum_austrialiense_broomeae_elkanii_embraeps ...(18 species)
SPP64 *Haemophilus* seminalis_sp._HMT_908
SPP68 *Neisseria* flava_macacae_mucosa_sicca
SPP76 *Acinetobacter*_Prolinoborus_fasciculus_woffii
SPP79 *Haemophilus* haemolyticus_sp._HMT_036
SPP8 *Streptococcus* parasanguinis_clade_411_sinensis
SPP83 *Lactobacillus* crispatus_gallinarum
SPP85 *Veillonella* parvula_tobetsuensis
SPP87 *Veillonella* parvula_rogosae_tobetsuensis
SPP90 *Lachnoanaerobaculum* gingivalis_umeaense
SPP98 *Parvimonas*_Peptostreptococcus_Candidatus_massiliensis_sp._HMT_110_sp._HMT_393
SPPN104 *Enterococcus* multispecies_sppn104_2_nov_96.360%
SPPN109 *Schaalia* multispecies_sppn109_2_nov_87.419%
SPPN13 *Streptococcus* multispecies_sppn13_18_nov_97.859%
SPPN143 *Streptococcus* multispecies_sppn143_2_nov_97.849%
SPPN151 *Streptococcus* multispecies_sppn151_17_nov_92.505%
SPPN171 *Streptococcus* multispecies_sppn171_17_nov_94.658%
SPPN184 *Streptococcus* multispecies_sppn184_17_nov_97.634%