

cies

- SP132 *Fusobacterium nucleatum*_subsp._animalis
- SP133 *Megasphaera micronuciformis*
- SP134 *Lachnoanaerobaculum umeaense*
- SP136 *Tannerella* sp._HMT_286
- SP137 *Streptococcus australis*
- SP138 *Actinomyces naeslundii*
- SP139 *Selenomonas noxia*
- SP14 *Aggregatibacter* sp._HMT_458
- SP140 *Fusobacterium hwasookii*
- SP142 *Aggregatibacter* sp._HMT_513
- SP143 *Actinomyces* sp._HMT_170
- SP144 *Stomatobaculum longum*
- SP145 *Prevotella pallens*
- SP146 *Haemophilus* sp._HMT_036
- SP147 *Prevotella nigrescens*
- SP148 *Leptotrichia hongkongensis*
- SP15 *Leptotrichia wadei*
- SP150 *Streptococcus infantis*_clade_431
- SP152 *Prevotella oris*
- SP153 *Selenomonas* sp._HMT_134
- SP155 *Absconditabacteria*_(SR1)_[G-1] bacterium_HMT_875
- SP156 *Oribacterium* sp._HMT_078
- SP157 *Gemella morbillorum*
- SP158 *Capnocytophaga* sp._HMT_332
- SP159 *Streptococcus cristatus*_clade_578
- SP16 *Veillonella dispar*
- SP160 *Streptococcus sinensis*
- SP161 *Streptococcus oralis*_subsp._dentisani_clade_058
- SP163 *Porphyromonas* sp._HMT_284
- SP164 *Lancefieldella parvula*
- SP165 *Capnocytophaga* sp._HMT_901
- SP167 *Prevotella denticola*
- SP168 *Haemophilus parahaemolyticus*
- SP169 *Streptococcus mutans*
- SP17 *Veillonella parvula*
- SP170 *Stomatobaculum* sp._HMT_097
- SP172 *Actinomyces johnsonii*
- SP173 *Peptoanaerobacter* [Eubacterium] *yurii*
- SP174 *Prevotella intermedia*

- SP26 *Pseudoleptotrichia* sp._HMT_221
- SP261 *Saccharibacteria*_(TM7)_[G-3] bacterium_HMT_351
- SP264 *Streptococcus* sp._HMT_074
- SP268 *Gracilibacteria*_(GN02)_[G-1] bacterium_HMT_872
- SP272 *Capnocytophaga* sp._HMT_863
- SP275 *Selenomonas* sp._HMT_920
- SP277 *Ruminococcaceae*_[G-2] bacterium_HMT_085
- SP28 *Pseudoleptotrichia* sp._HMT_219
- SP281 *Lautropia mirabilis*
- SP282 *Prevotella pleuritidis*
- SP283 *Streptococcus sobrinus*
- SP284 *Capnocytophaga* sp._HMT_336
- SP285 *Streptococcus oralis*_subsp._tigurinus_clade_071
- SP287 *Selenomonas* sp._HMT_937
- SP289 *Porphyromonas* sp._HMT_275
- SP29 *Haemophilus haemolyticus*
- SP290 *Prevotella micans*
- SP291 *Actinomyces* sp._HMT_525
- SP292 *Neisseria perflava*
- SP294 *Solobacterium moorei*
- SP297 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_957
- SP298 *Schaalia* sp._HMT_172
- SP3 *Porphyromonas* sp._HMT_930
- SP30 *Prevotella salivae*
- SP301 *Streptococcus oralis*_subsp._dentisani_clade_398
- SP304 *Mogibacterium diversum*
- SP305 *Neisseria mucosa*
- SP31 *Porphyromonas catoniae*
- SP313 *Streptococcus pseudopneumoniae*
- SP32 *Actinomyces oris*
- SP324 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_348
- SP329 *Leptotrichia* sp._HMT_218
- SP33 *Lachnoanaerobaculum gingivalis*
- SP34 *Fusobacterium nucleatum*
- SP350 *Streptococcus constellatus*
- SP356 *Streptococcus peroris*
- SP359 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_869
- SP36 *Catonella morbi*
- SP361 *Rothia dentocariosa*

- SP85 *Aggregatibacter* sp._HMT_512
- SP86 *Campylobacter gracilis*
- SP87 *Lachnoanaerobaculum saburreum*
- SP88 *Lachnoanaerobaculum orale*
- SP89 *Leptotrichia* sp._HMT_212
- SP9 *Leptotrichia* sp._HMT_392
- SP90 *Leptotrichia* sp._HMT_417
- SP91 *Streptococcus periodonticum*
- SP92 *Prevotella melaninogenica*
- SP93 *Schaalia* sp._HMT_180
- SP95 *Haemophilus parainfluenzae*
- SP96 *Bacteroidales*_[G-2] bacterium_HMT_274
- SP97 *Leptotrichia hofstadii*
- SP99 *Actinomyces* sp._HMT_897
- SPN105 *Pseudoleptotrichia goodfellowii*_nov_91.416%
- SPN109 *Actinomyces* sp._HMT_175_nov_97.751%
- SPN120 *Aggregatibacter* sp._HMT_458_nov_97.947%
- SPN132 *Prevotella veroralis*_nov_97.342%
- SPN140 *Actinomyces naeslundii*_nov_97.963%
- SPN149 *Streptococcus cristatus*_nov_97.576%
- SPN152 *Selenomonas noxia*_nov_97.441%
- SPN153 *Parvimonas micra*_nov_97.053%
- SPN154 *Granulicatella elegans*_nov_97.988%
- SPN155 *Fusobacterium hwasookii*_nov_97.783%
- SPN156 *Prevotella* sp._HMT_305_nov_93.878%
- SPN157 *Alloprevotella* sp._HMT_913_nov_97.955%
- SPN158 *Aggregatibacter* sp._HMT_949_nov_96.524%
- SPN159 *Selenomonas* sp._HMT_138_nov_97.233%
- SPN165 *Corynebacterium durum*_nov_97.689%
- SPN176 *Streptococcus gordonii*_nov_97.018%
- SPN18 *Corynebacterium matruchotii*_nov_97.755%
- SPN184 *Kingella oralis*_nov_97.741%
- SPN192 *Lachnospiraceae*_[G-2] bacterium_HMT_088_nov_93.096%
- SPN201 *Cardiobacterium hominis*_nov_97.510%
- SPN211 *Moraxella oblonga*_nov_93.776%
- SPN213 *Porphyromonas catoniae*_nov_97.746%
- SPN220 *Porphyromonas catoniae*_nov_97.951%
- SPN231 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_348_nov_95.08
- SPN249 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_347_nov_95.75