

- SP136 *Catonella morbi*
- SP137 *Aggregatibacter* sp._HMT_949
- SP138 *Fusobacterium* sp._HMT_203
- SP139 *Selenomonas* sp._HMT_138
- SP14 *Alloprevotella* sp._HMT_912
- SP140 *Arachnia propionica*
- SP141 *Actinomyces dentalis*
- SP142 *Streptococcus australis*
- SP145 *Prevotella shahii*
- SP146 *Eikenella corrodens*
- SP147 *Selenomonas* sp._HMT_892
- SP148 *Oribacterium asaccharolyticum*
- SP149 *Selenomonas noxia*
- SP15 *Pseudoleptotrichia* sp._HMT_219
- SP150 *Actinomyces* sp._HMT_897
- SP151 *Actinomyces oris*
- SP152 *Selenomonas diana*
- SP153 *Alloprevotella* sp._HMT_308
- SP154 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_352
- SP155 *Capnocytophaga endodontalis*
- SP156 *Abiotrophia defectiva*
- SP157 *Porphyromonas* sp._HMT_275
- SP158 *Actinomyces* sp._HMT_175
- SP159 *Streptococcus oralis*
- SP16 *Prevotella* sp._HMT_305
- SP160 *Alloprevotella tannerae*
- SP162 *Neisseria flava*
- SP163 *Prevotella* sp._HMT_317
- SP164 *Capnocytophaga* sp._HMT_878
- SP165 *Capnocytophaga* sp._HMT_336
- SP166 *Fusobacterium* sp._HMT_204
- SP167 *Peptostreptococcaceae*_[G-1] [Eubacterium]_sulci
- SP168 *Selenomonas* sp._HMT_137
- SP169 *Streptococcus* sp._HMT_423
- SP17 *Riemerella* sp._HMT_322
- SP171 *Aggregatibacter* sp._HMT_898
- SP173 *Campylobacter concisus*
- SP175 *Gemella sanguinis*
- SP176 *Fusobacterium nucleatum* subsp._vincentii

- SP268 *Neisseria subflava*
- SP269 *Gracilibacteria*_(GN02)_[G-1] bacterium_HMT_872
- SP27 *Schaalia* sp._HMT_178
- SP273 *Leptotrichia* sp._HMT_218
- SP274 *Haemophilus sputorum*
- SP276 *Cloacibacterium* sp._HMT_206
- SP277 *Neisseria mucosa*
- SP279 *Capnocytophaga* sp._HMT_335
- SP28 *Haemophilus parainfluenzae*
- SP285 *Schaalia meyeri*
- SP286 *Leptotrichia wadei*
- SP288 *Streptococcus* sp._HMT_056
- SP289 *Gracilibacteria*_(GN02)_[G-2] bacterium_HMT_873
- SP29 *Actinomyces massiliensis*
- SP290 *Peptostreptococcaceae*_[G-9] [Eubacterium]_brachy
- SP291 *Ruminococcaceae*_[G-2] bacterium_HMT_085
- SP295 *Lachnospiraceae*_[G-2] bacterium_HMT_088
- SP3 *Lachnoanaerobaculum orale*
- SP301 *Actinomyces johnsonii*
- SP302 *Neisseria sicca*
- SP305 *Streptococcus oralis* subsp._tigurinus_clade_071
- SP31 *Leptotrichia hongkongensis*
- SP310 *Prevotella scopos*
- SP313 *Neisseria cinerea*
- SP32 *Peptidiphaga* sp._HMT_183
- SP320 *Streptococcus* sp._HMT_074
- SP328 *Propionibacteriaceae*_[G-1] bacterium_HMT_915
- SP329 *Gracilibacteria*_(GN02)_[G-1] bacterium_HMT_871
- SP33 *Gemella haemolysans*
- SP339 *Streptococcus intermedius*
- SP34 *Capnocytophaga granulosa*
- SP343 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_346
- SP344 *Actinomyces* sp._HMT_414
- SP348 *Streptococcus mutans*
- SP352 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_348
- SP356 *Fusobacterium* sp._HMT_248
- SP358 *Arachnia rubra*
- SP359 *Dialister invisus*
- SP36 *Leptotrichia trevisanii*

- SP9 *Prevotella nigrescens*
- SP90 *Capnocytophaga sputigena*
- SP91 *Neisseria* sp._HMT_499
- SP92 *Prevotella nanceiensis*
- SP93 *Streptococcus koreensis*
- SP94 *Haemophilus haemolyticus*
- SP95 *Prevotella* sp._HMT_306
- SP96 *Alloprevotella* sp._HMT_913
- SP97 *Tannerella* sp._HMT_286
- SP98 *Peptostreptococcus stomatis*
- SPN1 *Actinomyces israelii*_nov_96.245%
- SPN100 *Actinomyces naeslundii*_nov_97.813%
- SPN102 *Corynebacterium durum*_nov_97.904%
- SPN11 *Capnocytophaga gingivalis*_nov_97.689%
- SPN112 *Porphyromonas catoniae*_nov_97.342%
- SPN120 *Prevotella jejuni*_nov_97.755%
- SPN124 *Streptococcus parasanguinis*_clade_411_nov_97.976%
- SPN129 *Pseudoleptotrichia goodfellowii*_nov_90.578%
- SPN139 *Actinomyces* sp._HMT_175_nov_97.951%
- SPN140 *Kingella oralis*_nov_97.741%
- SPN150 *Lachnospiraceae*_[G-2] bacterium_HMT_088_nov_92.887%
- SPN157 *Campylobacter concisus*_nov_97.403%
- SPN163 *Leptotrichia* sp._HMT_215_nov_97.397%
- SPN172 *Prevotella veroralis*_nov_97.342%
- SPN179 *Leptotrichia* sp._HMT_215_nov_96.963%
- SPN18 *Porphyromonas catoniae*_nov_97.951%
- SPN186 *Selenomonas* sp._HMT_137_nov_95.842%
- SPN197 *Selenomonas flueggei*_nov_97.822%
- SPN211 *Actinomyces* sp._HMT_169_nov_97.992%
- SPN222 *Corynebacterium matruchotii*_nov_97.331%
- SPN27 *Actinomyces* sp._HMT_175_nov_97.755%
- SPN304 *Streptobacillus notomitis*_nov_93.856%
- SPN31 *Capnocytophaga gingivalis*_nov_90.546%
- SPN317 *Alloprevotella* sp._HMT_914_nov_97.347%
- SPN37 *Porphyromonas catoniae*_nov_97.536%
- SPN404 *Actinomyces* sp._HMT_175_nov_97.746%
- SPN413 *Cardiobacterium hominis*_nov_97.510%
- SPN44 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_347_nov_95.759%
- SPN506 *Capnocytophaga gingivalis*_nov_96.218%