

- SP149 Alloprevotella sp._HMT_913
- SP15 Capnocytophaga sp._HMT_338
- SP150 Leptotrichia sp._HMT_223
- SP151 Actinomyces israelii
- SP152 Capnocytophaga sputigena
- SP153 Prevotella oralis
- SP154 Fretibacterium sp._HMT_362
- SP155 Streptococcus periodonticum
- SP156 Neisseria flavescens
- SP157 Actinomyces dentalis
- SP158 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
- SP159 Selenomonas sp._HMT_442
- SP16 Catonella morbi
- SP160 Fusobacterium sp._HMT_370
- SP161 Schaalia odontolytica
- SP163 Treponema sp._HMT_237
- SP164 Bacteroidetes_[G-3] bacterium_HMT_365
- SP165 Actinomyces sp._HMT_170
- SP166 Parvimonas sp._HMT_110
- SP167 Capnocytophaga sp._HMT_332
- SP168 Actinomyces sp._HMT_897
- SP169 Selenomonas sp._HMT_137
- SP170 Streptococcus oralis
- SP171 Capnocytophaga sp._HMT_863
- SP173 Capnocytophaga sp._HMT_335
- SP174 Streptococcus oralis_subsp._tigurinus_clade_071
- SP175 Arachnia propionica
- SP176 Tannerella sp._HMT_286
- SP177 Granulicatella elegans
- SP178 Haemophilus sp._HMT_036
- SP179 Kingella oralis
- SP18 Alloprevotella sp._HMT_914
- SP180 Alloprevotella sp._HMT_912
- SP181 Prevotella melaninogenica
- SP182 Veillonellaceae_[G-1] bacterium_HMT_132
- SP183 Colibacter massiliensis
- SP186 Oribacterium sinus
- SP187 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP188 Capnocytophaga sp._HMT_336

- SP301 Schaalia georgiae
- SP303 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875
- SP305 Prevotella multiformis
- SP306 Fretibacterium sp._HMT_360
- SP308 Aggregatibacter sp._HMT_898
- SP309 Pseudoleptotrichia goodfellowii
- SP31 Solobacterium moorei
- SP310 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP312 Selenomonas sp._HMT_937
- SP313 Porphyromonas endodontalis
- SP314 Neisseria sp._HMT_020
- SP316 Capnocytophaga sp._HMT_380
- SP319 Porphyromonas sp._HMT_284
- SP320 Bifidobacterium dentium
- SP321 Treponema lecithinolyticum
- SP323 Ottowia sp._HMT_894
- SP324 Capnocytophaga sp._HMT_864
- SP325 Prevotella sp._HMT_314
- SP326 Streptococcus thermophilus
- SP328 Prevotella sp._HMT_315
- SP33 Treponema sp._HMT_258
- SP331 Mogibacterium timidum
- SP332 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP333 Schaalia sp._HMT_877
- SP334 Lactobacillus gasseri
- SP335 Neisseria subflava
- SP337 Neisseria cinerea
- SP338 Rothia dentocariosa
- SP340 Capnocytophaga sp._HMT_412
- SP341 Mycoplasma salivarium
- SP342 Neisseria oralis
- SP343 Kingella sp._HMT_012
- SP346 Leptotrichia sp._HMT_212
- SP347 Streptococcus cristatus
- SP348 Lancefieldella parvula
- SP35 Leptotrichia sp._HMT_498
- SP350 Neisseria flava
- SP352 Eggerthia cateniformis
- SP354 Actinomyces johnsonii

- SP80 Porphyromonas pasteri
- SP81 Haemophilus parainfluenzae
- SP82 Haemophilus haemolyticus
- SP83 Aggregatibacter sp._HMT_458
- SP84 Veillonella sp._HMT_780
- SP85 Prevotella nigrescens
- SP86 Alloprevotella tannerae
- SP87 Streptococcus cristatus_clade_578
- SP88 Leptotrichia sp._HMT_225
- SP9 Corynebacterium durum
- SP90 Lachnoanaerobaculum sp._HMT_083
- SP91 Granulicatella adiacens
- SP92 Selenomonas infelix
- SP93 Leptotrichia sp._HMT_417
- SP94 Prevotella sp._HMT_472
- SP95 Veillonella dispar
- SP96 Prevotella oulorum
- SP97 Leptotrichia shahii
- SP98 Tannerella sp._HMT_808
- SP99 Prevotella fusca
- SPN100 Prevotella veroralis_nov_97.751%
- SPN108 Actinomyces sp._HMT_175_nov_97.746%
- SPN113 Schaalia meyeri_nov_97.925%
- SPN125 Aggregatibacter aphrophilus_nov_97.347%
- SPN126 Kingella oralis_nov_97.536%
- SPN138 Peptidiphaga gingivicola_nov_96.920%
- SPN149 Selenomonas noxia_nov_97.835%
- SPN152 Neisseria sp._HMT_018_nov_97.942%
- SPN159 Actinomyces sp._HMT_169_nov_97.546%
- SPN168 Prevotella oris_nov_96.926%
- SPN178 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347_nov_95.746%
- SPN188 Capnocytophaga granulosa_nov_97.890%
- SPN197 Selenomonas sp._HMT_481_nov_97.638%
- SPN208 Propionivibrio dicarboxylicus_nov_93.802%
- SPN21 Porphyromonas catoniae_nov_97.951%
- SPN217 Leptotrichia hofstadii_nov_96.970%
- SPN219 Bergeyella zoohelcum_nov_92.593%
- SPN229 Capnocytophaga gingivalis_nov_90.546%
- SPN230 Treponema sp._HMT_927_nov_91.411%