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- SP147 Neisseria subflava
- SP148 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356
- SP149 Leptotrichia wadei
- SP15 Slackia exigua
- SP150 Prevotella sp._HMT_315
- SP151 Streptococcus australis
- SP152 Leptotrichia sp._HMT_392
- SP153 Selenomonas diana
- SP154 Prevotella nigrescens
- SP155 Selenomonas noxia
- SP156 Leptotrichia sp._HMT_463
- SP157 Streptococcus mitis
- SP159 Prevotella veroralis
- SP16 Veillonellaceae_[G-1] bacterium_HMT_155
- SP160 Actinomyces sp._HMT_897
- SP161 Tannerella sp._HMT_286
- SP162 Peptococcus sp._HMT_167
- SP163 Selenomonas sp._HMT_920
- SP165 Stomatobaculum sp._HMT_097
- SP166 Prevotella salivae
- SP167 Capnocytophaga gingivalis
- SP168 Bacteroidales_[G-2] bacterium_HMT_274
- SP169 Selenomonas sp._HMT_892
- SP17 Prevotella jejuni
- SP170 Eggerthia cateniformis
- SP172 Riemerella sp._HMT_322
- SP173 Rothia dentocariosa
- SP174 Selenomonas sp._HMT_137
- SP177 Solobacterium moorei
- SP18 Schaalia meyeri
- SP182 Abiotrophia defectiva
- SP183 Fusobacterium sp._HMT_370
- SP185 Prevotella melaninogenica
- SP186 Lachnoanaerobaculum sp._HMT_496
- SP187 Lachnoanaerobaculum sp._HMT_083
- SP188 Treponema sp._HMT_237
- SP189 Peptostreptococcus stomatis
- SP19 Johnsonella sp._HMT_166
- SP190 Selenomonas sp._HMT_136
- SP191 Eubacterium sp._HMT_295

- SP311 Streptococcus oralis_subsp._agathinus_clade_071
- SP312 Streptococcus vestibularis
- SP313 Aggregatibacter sp._HMT_898
- SP315 Lancefieldella rimae
- SP316 Actinomyces dentalis
- SP318 Lachnoanaerobaculum gingivalis
- SP32 Schaalia odontolytica
- SP321 Peptoniphilaceae_[G-1] bacterium_HMT_113
- SP322 Actinomyces timonensis
- SP323 Schaalia cardiffensis
- SP324 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869
- SP325 Treponema sp._HMT_234
- SP326 Saccharibacteria_(TM7)_[G-4] bacterium_HMT_355
- SP327 Tannerella sp._HMT_916
- SP328 Ottowia sp._HMT_894
- SP329 Pseudoramibacter alactolyticus
- SP331 Peptoanaerobacter [Eubacterium] yurii
- SP334 Prevotella marshii
- SP336 Centipeda periodontii
- SP338 Stomatobaculum sp._HMT_910
- SP339 Leptotrichia sp._HMT_212
- SP34 Haemophilus haemolyticus
- SP342 Rothia aeria
- SP343 Veillonellaceae_[G-1] bacterium_HMT_135
- SP346 Tannerella serpentiformis
- SP347 Actinomyces oricola
- SP349 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347
- SP350 Neisseria flava
- SP351 Peptoanaerobacter stomatis
- SP353 Prevotella sp._HMT_475
- SP355 Treponema sp._HMT_238
- SP356 Treponema sp._HMT_253
- SP358 Treponema sp._HMT_268
- SP359 Capnocytophaga sp._HMT_323
- SP36 Veillonellaceae_[G-1] bacterium_HMT_132
- SP360 Capnocytophaga sp._HMT_412
- SP361 Streptococcus chosunense
- SP362 Peptostreptococcaceae_[G-6] [Eubacterium] minutum
- SP363 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349
- SP368 Campylobacter sp._HMT_299

- SP83 Lancefieldella parvula
- SP84 Schaalia sp._HMT_180
- SP85 Prevotella sp._HMT_300
- SP86 Eikenella corrodens
- SP88 Prevotella sp._HMT_317
- SP89 Alloprevotella sp._HMT_473
- SP9 Fusobacterium nucleatum_subsp._animalis
- SP90 Gemella sanguinis
- SP91 Actinomyces sp._HMT_448
- SP92 Selenomonas sp._HMT_919
- SP93 Lachnoanaerobaculum umeaense
- SP94 Neisseria oralis
- SP96 Treponema sp._HMT_230
- SP97 Treponema denticola
- SP98 Bacteroidetes_[G-3] bacterium_HMT_280
- SP99 Campylobacter showae
- SPN103 Actinomyces israelii_nov_96.838%
- SPN109 Cardiobacterium hominis_nov_97.510%
- SPN111 Prevotella veroralis_nov_97.959%
- SPN121 Porphyromonas catoniae_nov_97.951%
- SPN128 Actinomyces israelii_nov_94.882%
- SPN136 Corynebacterium matruchotii_nov_97.769%
- SPN145 Selenomonas sp._HMT_481_nov_97.822%
- SPN152 Hoppeia youngheungensis_nov_80.457%
- SPN161 Actinomyces sp._HMT_171_nov_96.813%
- SPN162 Prevotella sp._HMT_300_nov_97.955%
- SPN172 Actinomyces sp._HMT_897_nov_97.769%
- SPN178 Desulfovibrio legallii_nov_94.343%
- SPN184 Tannerella forsythia_nov_97.308%
- SPN192 Sneathia sanguinegens_nov_90.578%
- SPN199 Lacrimispora xylanolytica_nov_88.613%
- SPN207 Corynebacterium durum_nov_97.484%
- SPN210 Actinomyces sp._HMT_175_nov_97.751%
- SPN213 Corynebacterium matruchotii_nov_97.955%
- SPN215 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348_nov_95.089%
- SPN223 Kingella oralis_nov_97.536%
- SPN232 Tepidisphaera mucosa_nov_92.461%
- SPN233 Selenomonas infelix_nov_97.822%
- SPN242 Actinomyces naeslundii_nov_97.963%
- SPN251 Prevotella sp._HMT_297