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| 20 Capnocytophaga leadbetteri | SP61 Porphyromonas catoniae | SPN297 Vespertiliibacter pulmonis_nov_89.162% |
| 200 Staphylococcus hominis | SP62 Prevotella nanceiensis | SPN298 Veillonella sp._HMT_780_nov_97.628% |
| 201 Kingella sp._HMT_012 | SP63 Streptococcus chosunense | SPN299 Nonlabens tegetincola_nov_84.758% |
| 202 Capnocytophaga sp._HMT_380 | SP64 Lachnoanaerobaculum saburreum | SPN3 Neisseria elongata_nov_88.710% |
| 203 Lachnoanaerobaculum umeaense | SP65 Prevotella oris | SPN30 Haemophilus pittmaniae_nov_90.233% |
| 204 Johnsonella ignava | SP66 Streptococcus salivarius | SPN300 Leptotrichia shahii_nov_88.362% |
| 205 Veillonella tobetsuensis | SP67 Streptococcus sp._HMT_061 | SPN301 Weeksellaceae_[G-1] sp._HMT_900_nov_97.942% |
| 206 Moraxella catarrhalis | SP68 Peptostreptococcaceae_[G-7] bacterium_HMT_922 | SPN302 Xylanimonas cellulositytica_nov_86.864% |
| 207 Catonella morbi | SP69 Ottowia sp._HMT_894 | SPN303 Glutamicibacter creatinolyticus_nov_78.571% |
| 208 Streptococcus anginosus | SP7 Prevotella jejuni | SPN304 Fusobacterium hwasookii_nov_97.783% |
| 209 Leptotrichia trevisanii | SP70 Mitsukella sp._HMT_521 | SPN305 Veillonellaceae_[G-1] bacterium_HMT_145_nov_96.356% |
| 21 Weeksellaceae_[G-1] sp._HMT_900 | SP71 Treponema sp._HMT_237 | SPN306 Prevotella salivae_nov_96.538% |
| 210 Actinomyces dentalis | SP72 Fusobacterium sp._HMT_248 | SPN307 Actinomyces sp._HMT_171_nov_96.813% |
| 211 Treponema vincentii | SP73 Prevotella sp._HMT_317 | SPN308 Actinomyces howellii_nov_87.347% |
| 212 Lachnospiraceae_[G-2] bacterium_HMT_088 | SP74 Porphyromonas sp._HMT_930 | SPN309 Selenomonas sp._HMT_138_nov_97.024% |
| 213 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 | SP75 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 | SPN31 Porphyromonas catoniae_nov_97.951% |
| 214 Capnocytophaga sp._HMT_878 | SP76 Porphyromonas pasteri | SPN310 Cryobacterium melibiosiphilum_nov_84.725% |
| 215 Peptoniphilaceae_[G-1] bacterium_HMT_113 | SP77 Lachnospiraceae_[G-3] bacterium_HMT_100 | SPN311 Allobranchiibius huperziae_nov_86.091% |
| 216 Actinomyces massiliensis | SP78 Porphyromonas sp._HMT_284 | SPN312 Pectinatus haikarae_nov_84.211% |
| 217 Neisseria flava | SP79 Campylobacter gracilis | SPN313 Veillonella rogosae_nov_89.526% |
| 218 Haemophilus sputorum | SP8 Capnocytophaga sputigena | SPN314 Hydrogenophilus thermoluteolus_nov_87.090% |
| 219 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 | SP82 Fusobacterium hwasookii | SPN315 Streptobacillus notomytis_nov_94.068% |
| 22 Abiotrophia defectiva | SP85 Delftia lacustris | SPN316 Corynebacterium matruchotii_nov_97.959% |
| 220 Capnocytophaga sp._HMT_901 | SP86 Streptococcus constellatus | SPN317 Abiotrophia defectiva_nov_95.382% |
| 221 Capnocytophaga sp._HMT_412 | SP87 Capnocytophaga granulosa | SPN318 Streptococcus pyogenes_nov_85.312% |
| 222 Streptococcus lactarius | SP88 Kingella denitrificans | SPN319 Prevotella oris_nov_93.256% |
| 224 Oribacterium parvum | SP89 Streptococcus oralis_subsp._dentisani_clade_058 | SPN32 Jeotgalibacillus soli_nov_81.312% |
| 225 Neisseria flavescens | SP9 Oribacterium sp._HMT_078 | SPN320 Prevotella marshii_nov_97.951% |
| 226 Rothia mucilaginosa | SP90 Streptococcus mutans | SPN33 Mobiluncus curtisii_nov_96.694% |
| 227 Streptococcus massiliensis | SP91 Peptostreptococcaceae_[G-4] bacterium_HMT_369 | SPN34 Lautropia mirabilis_nov_95.010% |
| 228 Selenomonas sp._HMT_919 | SP92 Capnocytophaga sp._HMT_332 | SPN35 Porphyromonas catoniae_nov_97.741% |
| 229 Pseudoramibacter alactolyticus | SP93 Actinomyces sp._HMT_169 | SPN36 Nocardia casuarinae_nov_80.785% |
| 23 Prevotella oulorum | SP94 Alloprevotella sp._HMT_473 | SPN37 Capnocytophaga sp._HMT_901_nov_90.466% |
| 230 Schaalia lingnae | SP95 Cardiobacterium hominis | SPN38 Capnocytophaga sp._HMT_878_nov_97.877% |
| 231 Streptococcus pseudopneumoniae | SP96 Veillonella sp._HMT_780 | SPN39 Neisseria mucosa_nov_95.910% |
| 232 Treponema sp._HMT_231 | SP97 Capnocytophaga sp._HMT_864 | SPN4 Selenomonas sp._HMT_388_nov_86.391% |
| 233 Capnocytophaga ochracea | SP98 Veillonella dispar | SPN40 Neisseria oralis_nov_95.277% |
| 234 Treponema denticola | SP99 Streptococcus mitis | SPN41 Ralstonia pickettii_nov_80.040% |
| 235 Campylobacter showae | SPN1 Eikenella corrodens_nov_94.274% | SPN42 Renibacterium salmoninarum_nov_89.300% |