

## Species

- SP20 Capnocytophaga leadbetteri
- SP200 Staphylococcus hominis
- SP201 Kingella sp.\_HMT\_012
- SP202 Capnocytophaga sp.\_HMT\_380
- SP203 Lachnoanaerobaculum umeaense
- SP204 Johnsonella ignava
- SP205 Veillonella tobetsuensis
- SP206 Moraxella catarrhalis
- SP207 Catonella morbi
- SP208 Streptococcus anginosus
- SP209 Leptotrichia trevisanii
- SP21 Weeksellaceae\_[G-1] sp.\_HMT\_900
- SP210 Actinomyces dentalis
- SP211 Treponema vincentii
- SP212 Lachnospiraceae\_[G-2] bacterium\_HMT\_088
- SP213 Absconditabacteria\_(SR1)\_[G-1] bacterium\_HMT\_345
- SP214 Capnocytophaga sp.\_HMT\_878
- SP215 Peptoniphilaceae\_[G-1] bacterium\_HMT\_113
- SP216 Actinomyces massiliensis
- SP217 Neisseria flava
- SP218 Haemophilus sputorum
- SP219 Saccharibacteria\_(TM7)\_[G-3] bacterium\_HMT\_351
- SP22 Abiotrophia defectiva
- SP220 Capnocytophaga sp.\_HMT\_901
- SP221 Capnocytophaga sp.\_HMT\_412
- SP222 Streptococcus lactarius
- SP224 Oribacterium parvum
- SP225 Neisseria flavescens
- SP226 Rothia mucilaginosa
- SP227 Streptococcus massiliensis
- SP228 Selenomonas sp.\_HMT\_919
- SP229 Pseudoramibacter alactolyticus
- SP23 Prevotella oulorum
- SP230 Schaalia lingnae
- SP231 Streptococcus pseudopneumoniae
- SP232 Treponema sp.\_HMT\_231
- SP233 Capnocytophaga ochracea
- SP234 Treponema denticola
- SP235 Campylobacter showae

- SP61 Porphyromonas catoniae
- SP62 Prevotella nanceiensis
- SP63 Streptococcus chosunense
- SP64 Lachnoanaerobaculum saburreum
- SP65 Prevotella oris
- SP66 Streptococcus salivarius
- SP67 Streptococcus sp.\_HMT\_061
- SP68 Peptostreptococcaceae\_[G-7] bacterium\_HMT\_922
- SP69 Ottowia sp.\_HMT\_894
- SP7 Prevotella jejuni
- SP70 Mitsuokella sp.\_HMT\_521
- SP71 Treponema sp.\_HMT\_237
- SP72 Fusobacterium sp.\_HMT\_248
- SP73 Prevotella sp.\_HMT\_317
- SP74 Porphyromonas sp.\_HMT\_930
- SP75 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_349
- SP76 Porphyromonas pasteri
- SP77 Lachnospiraceae\_[G-3] bacterium\_HMT\_100
- SP78 Porphyromonas sp.\_HMT\_284
- SP79 Campylobacter gracilis
- SP8 Capnocytophaga sputigena
- SP82 Fusobacterium hwasookii
- SP85 Delftia lacustris
- SP86 Streptococcus constellatus
- SP87 Capnocytophaga granulosa
- SP88 Kingella denitrificans
- SP89 Streptococcus oralis\_subsp.\_denticani\_clade\_058
- SP9 Oribacterium sp.\_HMT\_078
- SP90 Streptococcus mutans
- SP91 Peptostreptococcaceae\_[G-4] bacterium\_HMT\_369
- SP92 Capnocytophaga sp.\_HMT\_332
- SP93 Actinomyces sp.\_HMT\_169
- SP94 Alloprevotella sp.\_HMT\_473
- SP95 Cardiobacterium hominis
- SP96 Veillonella sp.\_HMT\_780
- SP97 Capnocytophaga sp.\_HMT\_864
- SP98 Veillonella dispar
- SP99 Streptococcus mitis
- SPN1 Eikenella corrodens\_nov\_94.274%

- SPN297 Vespertiliibacter pulmonis\_nov\_89.162%
- SPN298 Veillonella sp.\_HMT\_780\_nov\_97.628%
- SPN299 Nonlabens tegetincola\_nov\_84.758%
- SPN3 Neisseria elongata\_nov\_88.710%
- SPN30 Haemophilus pittmaniae\_nov\_90.233%
- SPN300 Leptotrichia shahii\_nov\_88.362%
- SPN301 Weeksellaceae\_[G-1] sp.\_HMT\_900\_nov\_97.942%
- SPN302 Xylanimonas cellulositytica\_nov\_86.864%
- SPN303 Glutamicibacter creatinolyticus\_nov\_78.571%
- SPN304 Fusobacterium hwasookii\_nov\_97.783%
- SPN305 Veillonellaceae\_[G-1] bacterium\_HMT\_145\_nov\_96.356%
- SPN306 Prevotella salivae\_nov\_96.538%
- SPN307 Actinomyces sp.\_HMT\_171\_nov\_96.813%
- SPN308 Actinomyces howellii\_nov\_87.347%
- SPN309 Selenomonas sp.\_HMT\_138\_nov\_97.024%
- SPN31 Porphyromonas catoniae\_nov\_97.951%
- SPN310 Cryobacterium melibiosiphilum\_nov\_84.725%
- SPN311 Allobranchiobius huperziae\_nov\_86.091%
- SPN312 Pectinatus haikarae\_nov\_84.211%
- SPN313 Veillonella rogosa\_nov\_89.526%
- SPN314 Hydrogenophilus thermoluteolus\_nov\_87.090%
- SPN315 Streptobacillus notomytis\_nov\_94.068%
- SPN316 Corynebacterium matruchotii\_nov\_97.959%
- SPN317 Abiotrophia defectiva\_nov\_95.382%
- SPN318 Streptococcus pyogenes\_nov\_85.312%
- SPN319 Prevotella oris\_nov\_93.256%
- SPN32 Jeotgalibacillus soli\_nov\_81.312%
- SPN320 Prevotella marshii\_nov\_97.951%
- SPN33 Mobiluncus curtisii\_nov\_96.694%
- SPN34 Lautropia mirabilis\_nov\_95.010%
- SPN35 Porphyromonas catoniae\_nov\_97.741%
- SPN36 Nocardia casuarinae\_nov\_80.785%
- SPN37 Capnocytophaga sp.\_HMT\_901\_nov\_90.466%
- SPN38 Capnocytophaga sp.\_HMT\_878\_nov\_97.877%
- SPN39 Neisseria mucosa\_nov\_95.910%
- SPN4 Selenomonas sp.\_HMT\_388\_nov\_86.391%
- SPN40 Neisseria oralis\_nov\_95.277%
- SPN41 Ralstonia pickettii\_nov\_80.040%
- SPN42 Renibacterium salmoninarum\_nov\_89.300%