

es

- SP222 Lactobacillus plantarum
- SP223 Pseudomonas cedrina
- SP224 Scardovia wiggsiae
- SP225 Corynebacterium propinquum
- SP226 Veillonella sp.\_HMT\_780
- SP229 Mitsukella sp.\_HMT\_131
- SP23 Streptococcus sp.\_HMT\_061
- SP230 Fusobacterium canifilinum
- SP231 Actinomyces sp.\_HMT\_448
- SP232 Prevotella sp.\_HMT\_526
- SP233 Cardiobacterium valvarum
- SP234 Lactococcus lactis
- SP235 Fusobacterium sp.\_HMT\_203
- SP236 Haemophilus sp.\_HMT\_908
- SP237 Staphylococcus aureus
- SP238 Prevotella oralis
- SP239 Prevotella sp.\_HMT\_515
- SP24 Streptococcus mitis
- SP240 Capnocytophaga sp.\_HMT\_335
- SP241 Lactobacillus ultunensis
- SP242 Schaalia sp.\_HMT\_172
- SP243 Kingella denitrificans
- SP244 Peptostreptococcaceae\_[XII][G-9] [Eubacterium]\_brachy
- SP245 Streptococcus sobrinus
- SP246 Bacteroidetes\_[G-5] bacterium\_HMT\_511
- SP247 Prevotella sp.\_HMT\_301
- SP248 Prevotella pleuritidis
- SP249 Actinomyces sp.\_HMT\_170
- SP25 Streptococcus sp.\_HMT\_074
- SP250 Prevotella sp.\_HMT\_300
- SP251 Pseudomonas pseudoalcaligenes
- SP252 Prevotella sp.\_HMT\_443
- SP253 Selenomonas flueggei
- SP254 Leptotrichia sp.\_AF189244.1
- SP255 Haemophilus influenzae
- SP256 Selenomonas sp.\_HMT\_134
- SP257 Prevotella sp.\_HMT\_376
- SP258 Saccharibacteria\_(TM7)\_[G-6] bacterium\_HMT\_870
- SP259 Prevotella buccae
- SP26 Leptotrichia buccalis
- SP260 Porphyromonas catoniae
- SP261 Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_369
- SP262 Prevotella marshii
- SP263 Mogibacterium timidum
- SP264 Bulleidia extracta
- SP265 Parascardovia denticolens
- SP266 Haemophilus sp.\_Oral\_Taxon\_G21
- SP267 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_348
- SP268 Schaalia odontolyticus
- SP269 Actinomyces johnsonii
- SP27 Lachnoanaerobaculum orale
- SP270 Corynebacterium kroppenstedtii
- SP271 Slackia exigua
- SP272 Treponema sp.\_HMT\_268
- SP273 Prevotella sp.\_HMT\_292
- SP274 Treponema lecithinolyticum
- SP275 Peptostreptococcaceae\_[XII][G-5] [Eubacterium]\_saphenum
- SP276 Mogibacterium neglectum
- SP277 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_352
- SP28 Streptococcus cristatus
- SP280 Alloscardovia urinalis
- SP281 Haemophilus aegyptius
- SP282 Veillonellaceae\_[G-1] bacterium\_HMT\_132
- SP283 Veillonella sp.\_Oral\_Taxon\_C09
- SP284 Streptococcus oralis\_subsp.\_dentisani\_clade\_058
- SP285 Leptotrichia trevisanii
- SP286 Selenomonas sp.\_Oral\_Taxon\_E97
- SPN106 Prevotella sp.\_HMT\_443\_nov\_95.678%
- SPN107 Streptococcus parasanguinis\_I\_nov\_92.710%
- SPN108 Haemophilus haemolyticus\_nov\_87.647%
- SPN109 Actinomyces sp.\_HMT\_175\_nov\_97.633%
- SPN11 Actinomyces sp.\_HMT\_171\_nov\_96.737%
- SPN110 Mycoplasma orale\_nov\_96.538%
- SPN111 Lachnospiraceae\_[G-3] bacterium\_HMT\_100\_nov\_95.992%
- SPN112 Actinomyces sp.\_HMT\_175\_nov\_97.053%
- SPN113 Streptococcus parasanguinis\_II\_nov\_97.373%
- SPN114 Lactobacillus gasseri\_nov\_94.803%
- SPN115 Anoxybacillus flavithermus\_nov\_97.373%
- SPN116 Bacteroides coprocola\_nov\_85.102%
- SPN117 Leptotrichia trevisanii\_nov\_94.779%
- SPN118 Porphyromonas pasteri\_nov\_90.566%
- SPN119 Propionibacterium acidifaciens\_nov\_91.816%
- SPN12 Selenomonas sp.\_HMT\_136\_nov\_97.917%
- SPN120 Prevotella melaninogenica\_nov\_97.159%
- SPN121 Megasphaera elsdenii\_nov\_82.149%
- SPN122 Selenomonas sp.\_HMT\_478\_nov\_89.163%
- SPN123 Saccharophagus degradans\_nov\_77.863%
- SPN124 Haemophilus haemolyticus\_nov\_88.583%
- SPN125 Anaeroglobus geminatus\_nov\_97.164%
- SPN126 Veillonella parvula\_nov\_95.037%
- SPN127 Negativicoccus massiliensis\_nov\_76.234%
- SPN128 Megasphaera micronuciformis\_nov\_96.000%
- SPN129 Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_103\_nov\_83.366%
- SPN13 Neptuniibacter caesariensis\_nov\_79.057%
- SPN130 Paenibacillus apiarius\_nov\_80.224%
- SPN131 Kingella kingae\_nov\_96.875%
- SPN132 Calycanthus floridus\_Oral\_Taxon\_D07\_nov\_94.658%
- SPN133 Selenomonas sp.\_HMT\_478\_nov\_97.897%
- SPN134 Peptostreptococcaceae\_[XII][G-2] bacterium\_HMT\_091\_nov\_97.189%
- SPN135 Capnocytophaga sputigena\_nov\_95.136%
- SPN136 Leptotrichia buccalis\_nov\_95.618%
- SPN137 Streptococcus mitis\_nov\_95.301%
- SPN138 Actinomyces graevenitzi\_nov\_96.825%
- SPN139 Prevotella sp.\_HMT\_305\_nov\_97.426%
- SPN14 Acinetobacter septicus\_nov\_97.137%
- SPN140 Alloprevotella sp.\_HMT\_473\_nov\_89.152%
- SPN141 Streptococcus dentasini\_nov\_84.888%
- SPN142 Ureibacillus composti\_nov\_85.714%
- SPN143 Kingella oralis\_nov\_94.737%
- SPN144 Streptococcus parasanguinis\_I\_nov\_96.987%
- SPN145 Tatumella punctata\_nov\_81.579%
- SPN146 Streptococcus sp.\_str\_C300\_nov\_91.729%
- SPN147 Alloscardovia urinalis\_nov\_95.565%
- SPN148 Catonella morbi\_nov\_95.372%
- SPN149 Porphyromonas sp.\_HMT\_930\_nov\_96.024%
- SPN15 Prevotella baroniae\_nov\_94.925%
- SPN150 Gemella haemolysans\_nov\_96.030%
- SPN151 Actinomyces odontolyticus\_nov\_83.804%
- SPN152 Shuttleworthia satelles\_nov\_95.753%
- SPN153 Rothia aeria\_nov\_96.920%
- SPN154 Haemophilus parainfluenzae\_nov\_97.529%
- SPN155 Streptococcus mitis\_nov\_93.928%
- SPN156 Actinomyces graevenitzi\_nov\_96.457%
- SPN157 Dialister pneumosintes\_nov\_91.712%
- SPN158 Capnocytophaga gingivalis\_nov\_92.843%
- SPN159 Actinomyces israelii\_nov\_97.015%
- SPN16 Lactobacillus plantarum\_nov\_96.558%
- SPN160 Brachymonas chironomi\_nov\_91.762%
- SPN161 Campylobacter showae\_nov\_93.613%
- SPN162 Streptococcus peroris\_nov\_87.664%
- SPN163 Gemella haemolysans\_nov\_93.220%
- SPN164 Veillonellaceae\_[G-1] bacterium\_HMT\_918\_nov\_96.296%
- SPN165 Streptococcus vestibularis\_nov\_97.561%
- SPN166 Cohnella plantaginis\_nov\_78.165%
- SPN402 Corynebacterium matruchotii\_nov\_95.455%
- SPN403 Capnocytophaga sp.\_HMT\_864\_nov\_97.782%
- SPN404 Bifidobacterium dentium\_nov\_97.614%
- SPN405 Abiotrophia defectiva\_nov\_96.468%
- SPN406 Actinomyces gerencseriae\_nov\_96.816%
- SPN407 Streptococcus sanguinis\_nov\_97.749%
- SPN408 Actinomyces viscosus\_nov\_97.393%
- SPN409 Chryseobacterium hispalense\_nov\_97.282%
- SPN41 Lactobacillus ultunensis\_nov\_94.891%
- SPN410 Aggregatibacter segnis\_nov\_97.723%
- SPN411 Triticum aestivum\_Oral\_Taxon\_D31\_nov\_91.026%
- SPN412 Haemophilus sp.\_HMT\_036\_nov\_97.826%
- SPN413 Porphyromonas endodontalis\_nov\_96.395%
- SPN414 Bifidobacterium longum\_nov\_96.647%
- SPN415 Lactobacillus salivarius\_nov\_95.818%
- SPN416 Selenomonas sp.\_HMT\_126\_nov\_97.723%
- SPN417 Kingella oralis\_nov\_97.338%
- SPN418 Treponema amylovorum\_nov\_97.839%
- SPN419 Haemophilus parainfluenzae\_nov\_97.519%
- SPN42 Mycoplasma equigenitalis\_nov\_96.907%
- SPN420 Leptotrichia sp.\_HMT\_215\_nov\_95.218%
- SPN421 Alloscardovia urinalis\_nov\_96.781%
- SPN422 Flavobacterium phragmitis\_nov\_79.732%
- SPN423 Lactobacillus iners\_nov\_84.568%
- SPN424 Leptotrichia wadei\_nov\_97.194%
- SPN425 Bergeyella sp.\_HMT\_931\_nov\_97.183%
- SPN426 Prevotella intermedia\_nov\_96.422%
- SPN427 Actinomyces sp.\_Oral\_Taxon\_848\_nov\_96.000%
- SPN428 NA felis\_nov\_95.446%
- SPN429 Tannerella sp.\_HMT\_286\_nov\_96.008%
- SPN43 Prevotella oulorum\_nov\_96.780%
- SPN430 Leptotrichia sp.\_AF189244.1\_nov\_88.623%
- SPN431 Bacteroidetes\_[G-5] bacterium\_HMT\_511\_nov\_96.982%
- SPN432 Granulicatella paradiacens\_nov\_96.629%
- SPN433 Prevotella enoea\_nov\_96.693%
- SPN434 Selenomonas sp.\_Oral\_Taxon\_137\_nov\_95.037%
- SPN435 Prevotella oris\_nov\_96.415%
- SPN436 Prevotella pallens\_nov\_97.348%
- SPN437 Leptotrichia sp.\_HMT\_215\_nov\_97.280%
- SPN438 Capnocytophaga sp.\_HMT\_326\_nov\_97.955%
- SPN439 Actinomyces viscosus\_nov\_97.765%
- SPN44 Prevotella oulorum\_nov\_89.792%
- SPN440 Neisseriaceae\_[G-1] bacterium\_HMT\_174\_nov\_97.830%
- SPN441 Actinomyces urogenitalis\_nov\_79.623%
- SPN442 Arthrobacter pityocampae\_nov\_88.512%
- SPN443 Neisseria flavescens|subflava\_nov\_94.896%
- SPN444 Streptococcus parasanguinis\_I\_nov\_96.992%
- SPN445 Porphyromonas pasteri\_nov\_96.205%
- SPN446 Prevotella sp.\_HMT\_306\_nov\_97.047%
- SPN447 Streptococcus halotolerans\_nov\_88.503%
- SPN448 Alloprevotella sp.\_HMT\_308\_nov\_97.250%
- SPN449 Lactobacillus salivarius\_nov\_93.333%
- SPN45 Prevotella dentalis\_nov\_97.529%
- SPN450 Streptomyces polyrhachis\_nov\_81.698%
- SPN451 Alkaliflexus imshenetskii\_nov\_80.337%
- SPN452 Selenomonas noxia\_nov\_96.875%
- SPN453 Cardiobacterium valvarum\_nov\_96.813%
- SPN454 Colibacter massiliensis\_nov\_96.545%
- SPN455 Arachnia rubra\_nov\_97.718%
- SPN456 Streptococcus parasanguinis\_I\_nov\_92.495%
- SPN457 Bacteroidetes\_[G-3] bacterium\_HMT\_365\_nov\_97.628%
- SPN458 Solobacterium moorei\_nov\_95.627%
- SPN459 Actinomyces sp.\_Oral\_Taxon\_178\_nov\_97.148%
- SPN46 Actinomyces sp.\_Oral\_Taxon\_848\_nov\_97.714%
- SPN460 Anaerocolumna xylanovorans\_nov\_82.746%
- SPN461 Granulicatella elegans\_nov\_97.566%
- SPN462 Selenomonas sputigena\_nov\_95.353%