SP459 Methanobrevibacter oralis (0.02%|0%|0%|0%|0%|0.06%|0%|0.02%|0%
SP282 Treponema sp._HMT_234 (0.05%|0.00%|0.03%|0%|0.03%|0
SP160 Treponema sp._HMT_230 (0.04%|0.01%|0.04%|0%|0.00%
SP250 Treponema sp._HMT_238 (0.05%|0.07%|0.04%|0%|0.04%
SP18 Treponema sp._HMT_237 (0.11%|0.15%|0.24%|0%|0.28%
SP139 Treponema medium (0.02%|0.06%|0%|0.03%|0.06%
SP114 Treponema sp._HMT_231 (0.06%|0.13%|0.09%|0%|0.09%
SP234 Treponema vincentii (0.03%|0.01%|0.01%|0%|0.01%|0.01%
SP28 Treponema sp._HMT_951 (0.01%|0.02%|0.02%|0%|0.02%
SP307 Treponema sp._HMT_247 (0.03%|0.10%|0.02%|0%|0.02%
SP84 Treponema denticola (0.09%|0.09%|0.23%|0%|0.27%|0.20
SP246 Treponema sp._HMT_258 (0.01%|0.02%|0.03%|0%|0.0
SP264 Treponema lecithinolyticum (0.11%|0.07%|0.16%|0%
SP203 Treponema maltophilum (0.07%|0.03%|0.02%|0%|0.0
SP343 Treponema sp._HMT_270 (0.04%|0%|0.01%|0%|0.05
SP317 Treponema sp._HMT_268 (0.06%|0.01%|0.02%|
SP72 Treponema socranskii (0.28%|0.56%|0.50%|0%
SP421 Treponema sp._HMT_263 (0.01%|0.01%|0.02%
SP91 Treponema sp._HMT_262 (0.03%|0.07%|0.13%|0%
SP272 Treponema amylovorum (0.01%|0.01%|0.02%|0%
SP120 Treponema sp._HMT_257 (0.06%|0.08%|0.08%|0
SP297 Desulfobulbus sp._HMT_041 (0.08%|0.01%|0.07%|0
SP44 Campylobacter concisus (0.22%|0.05%|0.17%|
SP46 Campylobacter showae (0.18%|0.16%|0.25%
SP409 Campylobacter rectus (0%|0.01%|0.01%|0%
SP228 Campylobacter sp._HMT_044 (0.01%|0.07%
SP151 Campylobacter gracilis (0.74%|0.77%|0.71
SP131 Lautropia mirabilis (0.75%|0.69%|0.90%|0.18%|0
SP392 Ottowia sp._HMT_894 (0.04%|0.05%|0%|0.0
SP232 Kingella oralis (0.50%|0.47%|0.82%|1.01%|0.49%
SP231 Neisseria bacilliformis (0.07%|0.04%|0.02%|1.82%
SP295 Neisseria elongata (0.27%|0.07%|0.30%|0.1
SP263 Neisseria oralis (0%|0.06%|0%|0%|0.05%|0
SP48 Eikenella corrodens (0.14%|0.14%|0.24%|0.0
SP333 Kingella denitrificans (0.01%|0%|0.01%|0.0
SP205 Neisseria mucosa (0.15%|0.03%|0.21%|0.15
SP318 Neisseria sicca (0.42%|0.09%|0.33%|0%|0.12%|
SP372 Neisseria subflava (0.00%|0.01%|0.05%|0.00
SP301 Neisseria perflava (0.01%|0.07%|0%|0%|0%|
SP202 Neisseria flavescens (0.02%|0.01%|0.16%|0%
SP356 Cardiobacterium valvarum (0.08%|0.03%|0.04%|0%
SP257 Cardiobacterium hominis (0.15%|0.12%|0.26%|0%|
SP63 Haemophilus parainfluenzae (2.83%|1.22%|2.26%
SP395 Haemophilus parahaemolyticus (0%|0%|0.02%
SP357 Haemophilus haemolyticus (0.04%|0.13%|0.0
SP401 Haemophilus sp._HMT_036 (0.00%|0.11%|0.0
SP68 Aggregatibacter sp._HMT_458 (0.12%|0.26%|0.
SP313 Aggregatibacter sp._HMT_512 (0.06%|0.09%|
SP149 Aggregatibacter segnis (0.11%|0.01%|0.06%
SP54 Aggregatibacter sp._HMT_513 (0.03%|0.03%|0
SP184 Aggregatibacter sp._HMT_898 (0%|0%|0.05%
SP152 Aggregatibacter aphrophilus (0.07%|0%|0.3
SP278 Aggregatibacter sp._HMT_949 (0.01%|0.02%
SP26 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_
SP245 Saccharibacteria_(TM7)_[G-6] bacterium_HMT
SP83 Saccharibacteria_(TM7)_[G-3] bacterium_H
SP340 Saccharibacteria_(TM7)_[G-2] bacterium_HM
SP386 Saccharibacteria_(TM7)_[G-1] bacterium_HM
SP166 Saccharibacteria_(TM7)_[G-1] bacterium_HMT
SP125 Saccharibacteria_(TM7)_[G-1] bacterium_HMT
SP135 Saccharibacteria_(TM7)_[G-1] bacterium_HM
SP183 Saccharibacteria_(TM7)_[G-1] bacterium_HM
SP344 Gracilibacteria_(GN02)_[G-1] bacterium_HM
SP201 Absconditabacteria_(SR1)_[G-1
SP410 Absconditabacteria_(SR1)_[G-1] b
SP196 Bergeyella sp._HMT_900 (0.06%|0.04%|
SP70 Bergeyella sp._HMT_322 (0.18%|0.16%|0
SP291 Capnocytophaga sp._HMT_338 (0.06%|0.
SP82 Capnocytophaga gingivalis (0.40%|0.19%
SP8 Capnocytophaga granulosa (0.26%|0.33%|0
SP50 Capnocytophaga sp._HMT_332 (0.08%|0.0
SP92 Capnocytophaga leadbetteri (0.41%|0.23%
SP216 Capnocytophaga sputigena (0.23%|0.19
SP101 Capnocytophaga endodontalis (0.23%|0
SP225 Capnocytophaga sp._HMT_336 (0.08%|0
SP178 Capnocytophaga sp._HMT_324 (0.01%|0
SP45 Capnocytophaga sp._HMT_380 (0.09%|0.
SP140 Capnocytophaga sp._HMT_864 (0.03%|
SP353 Capnocytophaga sp._HMT_902 (0.04%|
SP299 Capnocytophaga sp._HMT_335 (0.01%|
SP172 Capnocytophaga ochracea (0.07%|0.1
SP391 Capnocytophaga sp._HMT_412 (0.01%
SP81 Bacteroidetes_[G-5] bacterium_HMT_511
SP309 Bacteroidetes_[G-3] bacterium_HMT_3
SP240 Bacteroidetes_[G-3] bacterium_H
SP53 Bacteroidetes_[G-3] bacterium_HMT_2
SP14 Bacteroidales_[G-2] bacterium_HMT_27
SP193 Tannerella forsythia (0.37%|0.16%|0.28%
SP283 Tannerella serpentiformis (0.01%|0.03
SP212 Tannerella sp._HMT_286 (0.12%|0.08%
SP89 Tannerella sp._HMT_916 (0%|0.02%|0.0
SP30 Tannerella sp._HMT_808 (0.10%|0.17%|
SP86 Porphyromonas gingivalis (0.10%|0.20
SP132 Porphyromonas endodontalis (1.09
SP273 Porphyromonas sp._HMT_284 (0%|0%
SP113 Porphyromonas sp._HMT_277 (0.01%
SP24 Porphyromonas catoniae (0.07%|0.06
SP40 Porphyromonas sp._HMT_275 (0.09%|0
SP266 Porphyromonas sp._HMT_278 (0.00%
SP31 Porphyromonas pasteri (0.06%|0.30%
SP293 Phocaeicola abscessus (0.07%|0.0
SP36 Alloprevotella tannerae (0.35%|0
SP199 Alloprevotella rava (0.14%|0.14%
SP269 Alloprevotella sp._HMT_912 (0.0
SP105 Alloprevotella sp._HMT_913 (0.0
SP238 Alloprevotella sp._HMT_473 (0.
SP336 Alloprevotella sp._HMT_308 (0.
SP394 Alloprevotella sp._HMT_914 (0%
SP29 Prevotella sp._HMT_304 (0.07%|0.04
SP171 Prevotella micans (0.01%|0.03%
SP277 Prevotella nanceiensis (0.02%|0
SP11 Prevotella pleuritidis (0.24%|0.2
SP331 Prevotella enoeca (0.00%|0.02%
SP405 Prevotella oralis (0.14%|0.15%|
SP328 Prevotella sp._HMT_301 (0.02%|0
SP209 Prevotella marshii (0.03%|0.09
SP268 Prevotella saccharolytica (0.06
SP265 Prevotella sp._HMT_475 (0.01
SP118 Prevotella sp._HMT_317 (0.2
SP384 Prevotella sp._HMT_472 (0.0
SP119 Prevotella loescheii (0.09%|
SP335 Prevotella sp._HMT_820 (0%|0.0
SP5 Prevotella sp._HMT_443 (0.02%|0.0
SP20 Prevotella sp._HMT_526 (0.02
SP329 Prevotella buccae (0.03%|0.1
SP284 Prevotella baroniae (0.07%|0.
SP143 Prevotella dentalis (0.02%|0.0
SP210 Prevotella sp._HMT_315 (0.0
SP168 Prevotella sp._HMT_376 (0.0
SP23 Prevotella sp._HMT_292 (0.03
SP69 Prevotella sp._HMT_300 (0.14%
SP111 Prevotella oulorum (0.21%|0.20
SP124 Prevotella salivae (0.11%|0.16
SP129 Prevotella maculosa (0.14%|
SP32 Prevotella oris (0.57%|1.06%|0
SP73 Prevotella intermedia (0.29%|
SP148 Prevotella pallens (0.06%|0.0
SP259 Prevotella fusca (0.11%|0.02%|
SP59 Prevotella denticola (0.51%|1
SP64 Prevotella multiformis (0.23%
SP108 Prevotella jejuni (0.08%|0.01%|
SP17 Prevotella sp._HMT_314 (0.08%|
SP223 Prevotella melaninogenica (0.3
SP249 Prevotella histicola (0.07%|0.02
SP102 Prevotella veroralis (0.24%|0.2
SP247 Prevotella vespertina (0%|0.03
SP271 Mycoplasma faucium (0.05%|0.08%|0.14%|0%|
SP381 Mycoplasma salivarium (0.03%|0.07%|0.01%|
SP312 Eggerthia catenaformis (0.11%|0.01%|0.01%|0%|
SP258 Mollicutes_[G-1] bacterium_HMT_504 (0%|0
SP128 Erysipelotrichaceae_[G-1] bacterium_HMT_905 (0.
SP390 Erysipelotrichaceae_[G-1] bacterium_HMT_904 (0
SP316 Bulleidia extructa (0.08%|0.02%|0.07%|0%|0.11%|
SP371 Solobacterium moorei (0.13%|0.20%|0.06%|0%|0.2
SP303 Gemella bergeri (0.02%|0.00%|0.03%|0%|0.03%|0
SP77 Gemella morbillorum (0.28%|0.36%|0.47%|0%|0.61%
SP237 Gemella sanguinis (0.04%|0.06%|0.00%|0.70%|0.0
SP354 Gemella haemolysans (0.08%|0.32%|0.11%|0%|0.2
SP27 Ligilactobacillus salivarius (0%|0.37%|0%|0%|0.01%|
SP294 Lactobacillus ultunensis (0%|0.01%|0%|0%|0.01%
SP56 Abiotrophia defectiva (0.06%|0.07%|0.14%|0%|0.14%
SP34 Granulicatella elegans (0%|0.05%|0.03%|0%|0.09%|
SP2 Granulicatella adiacens (0.90%|0.86%|0.81%|4.23%|1.
SP4 Streptococcus mutans (2.51%|5.04%|2.10%|0.44%|2
SP21 Streptococcus sobrinus (0.12%|0.02%|0.01%|0%|0.
SP147 Streptococcus intermedius (1.29%|1.00%|1.27%|0
SP222 Streptococcus constellatus (0.21%|0.12%|0.17%|0
SP180 Streptococcus anginosus (0.58%|0.44%|0.81%|0.
SP363 Streptococcus massiliensis (0.02%|0.02%|0%|0%
SP155 Streptococcus gordonii (2.01%|2.40%|1.14%|0.55%
SP332 Streptococcus sp._HMT_056 (0.02%|0%|0.07%|0%
SP87 Streptococcus salivarius (0.29%|0.40%|0.48%|0%
SP39 Streptococcus vestibularis (0.32%|0.04%|0.48%|0
SP123 Streptococcus cristatus (0.80%|1.66%|1.00%|1.33
SP7 Streptococcus sanguinis (2.98%|2.90%|3.57%|1.94%
SP117 Streptococcus oralis_subsp._tigurinus_clade_070
SP200 Streptococcus oralis_subsp._dentisani_clade_058
SP156 Streptococcus chosunense (0.06%|0.04%|0.01%|
SP57 Streptococcus mitis (4.37%|7.88%|3.89%|1.73%|5
SP9 Streptococcus sp._HMT_064 (1.47%|1.00%|0.24%|6
SP12 Streptococcus oralis (2.21%|2.05%|2.27%|17.36%
SP226 Streptococcus australis (0.05%|0.01%|0%|0.0
SP133 Streptococcus sp._HMT_066 (0.02%|0.04%|0.02
SP37 Streptococcus parasanguinis_clade_411 (0.70%|
SP185 Streptococcus parasanguinis (0.01%|0.02%|0.0
SP76 Lachnospiraceae_[G-8] bacterium_HMT_500 (0.02%|0
SP454 Butyrivibrio sp._HMT_080 (0%|0%|0.02%|0%|0.03%
SP19 Catonella morbi (0.22%|0.16%|0.27%|0%|0.27%|0.2
SP319 Catonella sp._HMT_451 (0.01%|0.00%|0.02%|0%
SP221 Oribacterium sinus (0.09%|0.03%|0.03%|0%|0.08%
SP186 Oribacterium sp._HMT_078 (0.24%|0.43%|0.40%|0
SP326 Shuttleworthia satelles (0.06%|0.18%|0.11%|0%|0.
SP256 Stomatobaculum sp._HMT_910 (0.01%|0.01%|0.01%
SP241 Stomatobaculum sp._HMT_097 (0.02%|0.02%|0.02
SP3 Stomatobaculum longum (0.11%|0.27%|0.35%|0%|0.
SP373 Lachnospiraceae_[G-2] bacterium_HMT_096 (0.09%|0
SP99 Lachnoanaerobaculum sp._HMT_496 (0.04%|0.01%|
SP60 Lachnospiraceae_[G-3] bacterium_HMT_100 (0.12
SP380 Johnsonella sp._HMT_166 (0.06%|0.01%|0%|0%
SP467 Johnsonella ignava (0.03%|0%|0.00%|0%|0%
SP305 Lachnoanaerobaculum sp._HMT_083 (0.0%|0.02%
SP67 Lachnoanaerobaculum saburreum (0.32%|0.42%|0.
SP198 Lachnoanaerobaculum umeaense (0.04%|0.01%|
SP224 Lachnoanaerobaculum orale (0.14%|0.12%|0.09%
SP218 Ruminococcaceae_[G-1] bacterium_HMT_075 (0
SP275 Ruminococcaceae_[G-2] bacterium_HMT_085 (0.
SP423 Pseudoramibacter alactolyticus (0.04%|0%|0.02%|
SP217 Peptococcus sp._HMT_168 (0%|0.03%|0.00%|0%|0
SP47 Peptococcus sp._HMT_167 (0.05%|0.08%|0.30%|0%
SP162 Clostridiales_[F-1][G-1] bacterium_HMT_093 (0.
SP213 Peptoniphilus lacrimalis (0.01%|0.02%|0%|0.0
SP112 Parvimonas sp._HMT_110 (0.06%|0.29%|0.43%|0%
SP93 Parvimonas micra (1.37%|1.19%|1.32%|0%|1.77%|1
SP107 Filifactor alocis (0.19%|0.13%|0.22%|0%|0.32%|0
SP157 Peptoanaerobacter [Eubacterium] yurii (0.0
SP267 Peptoanaerobacter stomatis (0.02%|0.01%|
SP365 Peptostreptococcaceae_[XI][G-7] bacterium_
SP208 Peptostreptococcus stomatis (0.23%|0.21%|0.21
SP292 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_b
SP116 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_
SP323 Peptostreptococcaceae_[XI][G-5] [Eubacterium
SP351 Peptostreptococcaceae_[XI][G-6] [Eubacte
SP287 Peptostreptococcaceae_[XI][G-4] bacterium_H
SP440 Peptostreptococcaceae_[XI][G-4] bacterium_H
SP321 Peptostreptococcaceae_[XI][G-2] bacterium_H
SP165 Mogibacterium timidum (0.22%|0.17%|0.25%|
SP298 Mogibacterium neglectum (0.05%|0.09%|0.12
SP174 Mogibacterium vescum (0.03%|0.05%|0.04%|
SP43 Fusobacterium nucleatum (4.98%|6.25%|7.02%|0%
SP233 Fusobacterium sp._HMT_370 (0.03%|0%|0.04%|0
SP296 Fusobacterium periodonticum (0.05%|0.02%|0.10
SP375 Fusobacterium sp._HMT_204 (0.03%|0.05%|0.12%
SP58 Fusobacterium nucleatum_subsp._vincentii (0.65%
SP182 Fusobacterium canifelinum (0.02%|0.01%|0.07%|
SP304 Fusobacterium sp._HMT_203 (0.03%|0.09%|0.10%
SP254 Leptotrichia sp._HMT_221 (0.02%|0.03%|0.08%|0
SP376 Leptotrichia sp._HMT_219 (0%|0.01%|0.05%|0
SP310 Leptotrichia sp._HMT_392 (0.05%|0.03%|0.10
SP192 Leptotrichia sp._HMT_212 (0.08%|0.11%|0.06
SP62 Leptotrichia sp._HMT_215 (0.13%|0.12%|0.16%
SP35 Leptotrichia sp._HMT_498 (0.14%|0.18%|0.06
SP110 Leptotrichia sp._HMT_225 (0.00%|0%|0.04%|0
SP10 Leptotrichia buccalis (0.35%|0.22%|0.07%|0
SP227 Leptotrichia sp._HMT_223 (0.03%|0.04%|0.07
SP74 Leptotrichia hofstadii (0.03%|0.08%|0.06%|0
SP288 Leptotrichia sp._HMT_417 (0.27%|0.09%|0.13
SP85 Leptotrichia wadei (1.11%|0.63%|0.99%|0%|1.
SP163 Leptotrichia shahii (0.08%|0.06%|0.04%|0
SP100 Leptotrichia trevisanii (0%|0.00%|0.03%|0%
SP167 Leptotrichia hongkongensis (0.31%|0.55%|0.
SP447 Actinidia eriantha (0%|0.01%|0%|0%|0%|0.1
SP38 Veillonella sp._HMT_780 (0.01%|0.05%|0.01%|0
SP52 Veillonella parvula (0.86%|1.72%|0.91%|1.95%|1.4
SP80 Veillonella atypica (0.05%|0.07%|0.18%|0.23%|
SP462 Dialister sp._HMT_119 (0%|0.05%|0.02%|0%|0.0
SP25 Dialister invisus (0.67%|1.12%|0.54%|0%|0.95%|1
SP320 Megasphaera micronuciformis (0.09%|0.20%|0.19%
SP79 Anaeroglobus geminatus (0.17%|0.27%|0.28%|0%
SP378 Megasphaera sp._HMT_123 (0%|0.01%|0.01%|0
SP94 Colibacter massiliensis (0.04%|0.20%|0.03%|0%
SP229 Veillonellaceae_[G-1] bacterium_HMT_155 (0.08%
SP314 Veillonellaceae_[G-1] bacterium_HMT_145 (0.06
SP175 Veillonellaceae_[G-1] bacterium_HMT_129 (0.03%|
SP382 Veillonellaceae_[G-1] bacterium_HMT_150 (0.13%
SP352 Veillonellaceae_[G-1] bacterium_HMT_132 (0.02%
SP300 Mitsuokella sp._HMT_521 (0.08%|0.06%|0.09%|0
SP230 Selenomonas sp._HMT_442 (0.01%|0%|0%|0%|0.
SP127 Selenomonas sp._HMT_134 (0.04%|0.12%|0.06%|
SP13 Selenomonas sputigena (0.31%|0.42%|0.53%|0%
SP370 Selenomonas sp._HMT_937 (0.00%|0.00%|0.01%|
SP161 Selenomonas noxia (0.21%|0.35%|0.29%|0%|0.39
SP214 Selenomonas sp._HMT_133 (0.01%|0.01%|0.01%
SP103 Selenomonas artemidis (0.06%|0.07%|0.09%|0
SP177 Selenomonas sp._HMT_137 (0.02%|0.06%|0.03%
SP75 Selenomonas sp._HMT_136 (0.02%|0.01%|0.02%
SP126 Selenomonas sp._HMT_149 (0%|0.04%|0.01%|0%
SP399 Selenomonas sp._HMT_920 (0%|0.05%|0.00%|0
SP188 Selenomonas flueggei (0.07%|0.01%|0.04%|0
SP359 Selenomonas sp._HMT_892 (0.02%|0.02%|0.07%
SP159 Selenomonas sp._HMT_138 (0.02%|0.06%|0.05%
SP255 Selenomonas dianae (0.05%|0.02%|0.01%|0%|0.
SP338 Selenomonas sp._HMT_479 (0.02%|0.02%|0.01%|
SP164 Selenomonas infelix (0.08%|0.03%|0.05%|0%|0.1
SP396 Selenomonas sp._HMT_481 (0.04%|0.01%|0%|0%
SP377 Centipeda periodontii (0.02%|0.01%|0%|0%|0.0
SP279 Selenomonas sp._HMT_919 (0.01%|0.02%|0.03%
SP416 Selenomonas sp._HMT_146 (0.02%|0.02%|0.06%
SP158 Selenomonas sp._HMT_126 (0%|0.01%|0%|0%|0
SP360 Anaerolineae_[G-1] bacterium_HMT_439 (0.0
SP65 Fretibacterium fastidiosum (0.08%|0.06%|0.25%|0%
SP239 Fretibacterium sp._HMT_359 (0.09%|0.03%|0.09%
SP315 Fretibacterium sp._HMT_362 (0.06%|0.02%|0.09%
SP260 Fretibacterium sp._HMT_360 (0.17%|0.04%|0.15%
SP349 Cryptobacterium curtum (0.02%|0.03%|0.11%
SP195 Slackia exigua (0.11%|0.02%|0.01%|0%|0.09
SP207 Olsenella profusa (0%|0%|0%|0.16%|0%|0%
SP136 Olsenella sp._HMT_807 (1.04%|1.07%|0.73%|0%
SP281 Olsenella uli (0.11%|0.06%|0.10%|0.11%|0.1
SP347 Atopobium sp._HMT_416 (0%|0%|0%|0%|0%|0%
SP97 Lancefieldella parvula (0.24%|0.71%|0.33%|0%
SP461 Atopobium sp._HMT_199 (0.06%|0%|0.04%|0
SP42 Lancefieldella rimae (0.41%|0.94%|0.60%|0%|
SP95 Bifidobacterium dentium (0.05%|0.21%|0.38%
SP411 Bifidobacterium longum (0.03%|0%|0%|0%|0.
SP437 Parascardovia denticolens (0.01%|0%|0.01%
SP236 Scardovia wiggsiae (0.23%|0.74%|1.09%|0%
SP219 Propionibacterium acidifaciens (0.05%|0.01%|
SP235 Arachnia propionica (0.17%|0.25%|0.15%|0%
SP429 Arachnia rubra (0%|0.02%|0%|0%|0.01%|0.04
SP96 Corynebacterium durum (0.24%|0.24%|0.24%
SP6 Corynebacterium matruchotii (1.94%|1.32%|1.5
SP41 Rothia mucilaginosa (0.48%|0.10%|0.25%|0%|0.
SP61 Rothia aeria (0.52%|0.25%|0.44%|0%|0.11%|0
SP66 Rothia dentocariosa (18.77%|12.36%|12.92%|1
SP251 Peptidiphaga sp._HMT_183 (0.47%|2.28%|0.4
SP286 Peptidiphaga gingivicola (0.64%|0.07%|0.42
SP122 Schaalia sp._HMT_178 (0.22%|0.16%|0.35%
SP191 Schaalia georgiae (0.01%|0.03%|0.06%|0%
SP51 Schaalia meyeri (0.14%|0.21%|0.53%|0%|0.
SP130 Schaalia cardiffensis (0.04%|0.00%|0.01%
SP244 Schaalia lingnae_[Not_Validly_Published] (
SP456 Schaalia odontolytica (0.02%|0%|0%|0%
SP141 Schaalia odontolytica (0.25%|0.10%|0.09%
SP121 Schaalia sp._HMT_180 (0.55%|0.60%|0.48
SP153 Actinomyces sp._HMT_448 (3.29%|2.01%|2
SP253 Actinomyces massiliensis (0.19%|0.22%|0
SP109 Actinomyces israelii (0.42%|0.35%|0.75%
SP1 Actinomyces sp._HMT_897 (0.33%|0.59%|0
SP206 Actinomyces sp._HMT_414 (0.60%|0.07
SP346 Actinomyces dentalis (0.61%|0.59%|0.7
SP145 Actinomyces sp._HMT_896 (0.07%|0.08
SP137 Actinomyces sp._HMT_525 (0.24%|0.28
SP88 Actinomyces timonensis (0.06%|0.03%|0.10
SP242 Actinomyces johnsonii (0.49%|0.11%|0.25
SP189 Actinomyces sp._HMT_171 (0.26%|0.22%|0
SP49 Actinomyces sp._HMT_170 (0.49%|0.32%|0.3
SP179 Actinomyces naeslundii (0.32%|0.91%|0.34
SP146 Actinomyces viscosus (0.13%|0.07%|0.13%
SP22 Actinomyces sp._HMT_169 (2.54%|1.56%|2.
SP173 Actinomyces oris (0.99%|0.71%|0.81%|2.1
SP190 Actinomyces sp._HMT_175 (1.05%|0.53%|
24_Weeks.B.Plaque_2
12_Weeks.A.Plaque_2
12_Weeks.B.Plaque_1
Baseline.NA.Plaque_1
Baseline.B.Plaque_2
Baseline.A.Plaque_2
12_Weeks.B.Plaque_2
Baseline.B.Plaque_1
24_Weeks.A.Plaque_1
Baseline.NA.Plaque_2
Baseline.A.Plaque_1
12_Weeks.A.Plaque_1
24_Weeks.B.Plaque_1
24_Weeks.A.Plaque_2
substitutions/site
0
0.25
0.5
0.75
1
1.25