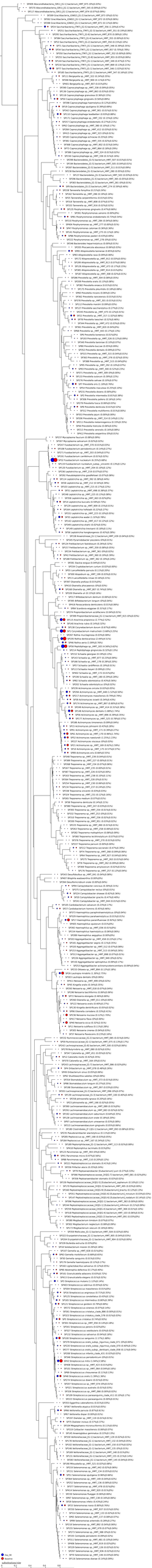

SP409 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0%|0.03%)
SP270 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0.09%|0.07%)
SP117 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0.07%|0.09%)
SP269 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873 (0%|0.01%)
SP262 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872 (0.03%|0.06%)
SP288 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_871 (0.11%|0.06%)
SP24 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (1.20%|0.47%)
SP251 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0.19%|0.06%)
SP295 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.08%|0.10%)
SP384 Saccharibacteria_(TM7)_[G-4] bacterium_HMT_355 (0.00%|0.01%)
SP53 Saccharibacteria_(TM7)_[G-2] bacterium_HMT_350 (0.15%|0.10%)
SP71 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0.39%|0.35%)
SP34 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0.70%|0.78%)
SP350 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0.00%|0.00%)
SP255 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955 (0.05%|0.06%)
SP144 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0.47%|0.57%)
SP216 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869 (0.13%|0.21%)
SP77 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (1.02%|0.91%)
SP185 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347 (0.16%|0.15%)
SP111 Bergeyella sp._HMT_322 (0.24%|0.35%)
SP266 Bergeyella sp._HMT_900 (0.11%|0.07%)
SP401 Bergeyella sp._HMT_907 (0.01%|0%)
SP290 Capnocytophaga sp._HMT_338 (0.09%|0.05%)
SP136 Capnocytophaga sp._HMT_863 (0.03%|0.05%)
SP118 Capnocytophaga granulosa (0.36%|0.15%)
SP50 Capnocytophaga gingivalis (0.56%|0.89%)
SP298 Capnocytophaga haemolytica (0.12%|0.00%)
SP19 Capnocytophaga sputigena (0.39%|0.69%)
SP342 Capnocytophaga sp._HMT_901 (0.01%|0.01%)
SP110 Capnocytophaga leadbetteri (0.93%|0.88%)
SP171 Capnocytophaga sp._HMT_332 (0.15%|0.10%)
SP257 Capnocytophaga endodontalis (0.27%|0.21%)
SP62 Capnocytophaga sp._HMT_380 (0.10%|0.17%)
SP284 Capnocytophaga sp._HMT_412 (0.03%|0.01%)
SP415 Capnocytophaga sp._HMT_323 (0%|0.01%)
SP183 Capnocytophaga ochracea (0.25%|0.19%)
SP381 Capnocytophaga sp._HMT_335 (0.02%|0.02%)
SP368 Capnocytophaga sp._HMT_902 (0.01%|0.04%)
SP73 Capnocytophaga sp._HMT_864 (0.18%|0.29%)
SP386 Capnocytophaga sp._HMT_324 (0.01%|0.02%)
SP156 Capnocytophaga sp._HMT_336 (0.31%|0.16%)
SP299 Bacteroidetes_[G-5] bacterium_HMT_507 (0.01%|0.01%)
SP194 Bacteroidetes_[G-5] bacterium_HMT_505 (0.04%|0.01%)
SP287 Bacteroidetes_[G-5] bacterium_HMT_511 (0.01%|0.09%)
SP238 Bacteroidetes_[G-3] bacterium_HMT_280 (0.05%|0.03%)
SP137 Bacteroidetes_[G-3] bacterium_HMT_503 (0.03%|0.01%)
SP296 Bacteroidetes_[G-3] bacterium_HMT_365 (0.05%|0.05%)
SP383 Bacteroidetes_[G-3] bacterium_HMT_281 (0%|0.01%)
SP9 Bacteroidales_[G-2] bacterium_HMT_274 (0.59%|0.46%)
SP228 Tannerella forsythia (0.21%|0.24%)
SP162 Tannerella sp._HMT_286 (0.19%|0.16%)
SP25 Tannerella serpentiformis (0.01%|0.03%)
SP316 Tannerella sp._HMT_808 (0.07%|0.07%)
SP333 Tannerella sp._HMT_916 (0.02%|0.02%)
SP135 Porphyromonas gingivalis (0.47%|0.66%)
SP391 Porphyromonas uenonis (0.00%|0%)
SP81 Porphyromonas endodontalis (0.73%|0.54%)
SP210 Porphyromonas sp._HMT_277 (0.00%|0.01%)
SP404 Porphyromonas sp._HMT_284 (0.04%|0.06%)
SP47 Porphyromonas catoniae (0.36%|0.36%)
SP141 Porphyromonas sp._HMT_275 (0.11%|0.18%)
SP86 Porphyromonas pasteri (0.43%|0.69%)
SP222 Porphyromonas sp._HMT_278 (0%|0.02%)
SP246 Bacteroides heparinolyticus (0.00%|0.01%)
SP205 Phocaeicola abscessus (0.06%|0.03%)
SP85 Alloprevotella tannerae (0.85%|0.62%)
SP83 Alloprevotella rava (0.09%|0.06%)
SP172 Alloprevotella sp._HMT_912 (0.03%|0.05%)
SP199 Alloprevotella sp._HMT_913 (0.07%|0.06%)
SP126 Alloprevotella sp._HMT_473 (0.17%|0.15%)
SP385 Alloprevotella sp._HMT_914 (0.01%|0%)
SP187 Alloprevotella sp._HMT_308 (0.02%|0.01%)
SP286 Prevotella sp._HMT_304 (0.08%|0.05%)
SP209 Prevotella oralis (0.13%|0.08%)
SP382 Prevotella enoeca (0.01%|0.01%)
SP175 Prevotella pleuritidis (0.18%|0.08%)
SP60 Prevotella micans (0.06%|0.10%)
SP362 Prevotella nanceiensis (0.01%|0.01%)
SP378 Prevotella sp._HMT_301 (0.01%|0.01%)
SP112 Prevotella marshii (0.09%|0.03%)
SP127 Prevotella saccharolytica (0.27%|0.21%)
SP245 Prevotella sp._HMT_475 (0.12%|0.01%)
SP22 Prevotella sp._HMT_317 (1.02%|0.98%)
SP78 Prevotella loescheii (0.52%|0.46%)
SP344 Prevotella sp._HMT_472 (0.03%|0.05%)
SP361 Prevotella sp._HMT_820 (0.04%|0%)
SP68 Prevotella sp._HMT_443 (0.17%|0.13%)
SP6 Prevotella koreensis (0.01%|0%)
SP193 Prevotella sp._HMT_526 (0.12%|0.09%)
SP340 Prevotella baroniae (0.10%|0.07%)
SP80 Prevotella buccae (0.05%|0.09%)
SP352 Prevotella dentalis (0.09%|0.07%)
SP315 Prevotella sp._HMT_315 (0.04%|0.01%)
SP363 Prevotella sp._HMT_376 (0.02%|0.05%)
SP388 Prevotella sp._HMT_515 (0.04%|0%)
SP95 Prevotella sp._HMT_292 (0.10%|0.07%)
SP93 Prevotella sp._HMT_300 (0.52%|0.23%)
SP371 Prevotella sp._HMT_309 (0%|0.00%)
SP133 Prevotella oulorum (0.39%|0.22%)
SP276 Prevotella salivae (0.10%|0.07%)
SP7 Prevotella oris (1.18%|0.76%)
SP54 Prevotella maculosa (0.25%|0.10%)
SP231 Prevotella disiens (0.00%|0%)
SP2 Prevotella intermedia (0.81%|0.38%)
SP306 Prevotella pallens (0.16%|0.14%)
SP279 Prevotella fusca (0.06%|0.03%)
SP8 Prevotella denticola (0.61%|0.52%)
SP312 Prevotella multiformis (0.01%|0.00%)
SP353 Prevotella jejuni (0.00%|0.00%)
SP206 Prevotella sp._HMT_314 (0.14%|0.11%)
SP211 Prevotella melaninogenica (0.33%|0.35%)
SP44 Prevotella histicola (0.06%|0.03%)
SP213 Prevotella veroralis (0.16%|0.08%)
SP412 Prevotella vespertina (0%|0.01%)
SP237 Mycoplasma faucium (0.08%|0.09%)
SP307 Mycoplasma salivarium (0.02%|0.02%)
SP247 Fusobacterium sp._HMT_370 (0.01%|0.03%)
SP108 Fusobacterium periodonticum (0.14%|0.27%)
SP271 Fusobacterium sp._HMT_203 (0.04%|0.07%)
SP191 Fusobacterium canifelinum (0.01%|0.02%)
SP32 Fusobacterium nucleatum (6.35%|5.68%)
SP119 Fusobacterium nucleatum_subsp._vincentii (0.13%|0.12%)
SP120 Fusobacterium sp._HMT_204 (0.10%|0.12%)
SP336 Leptotrichia sp._HMT_219 (0.07%|0.07%)
SP292 Pseudoleptotrichia goodfellowii (0.07%|0.08%)
SP114 Leptotrichia sp._HMT_392 (0.38%|0.44%)
SP28 Leptotrichia sp._HMT_212 (0.54%|0.39%)
SP103 Leptotrichia sp._HMT_215 (0.17%|0.22%)
SP31 Leptotrichia sp._HMT_498 (0.38%|0.37%)
SP248 Leptotrichia sp._HMT_225 (0.13%|0.08%)
SP358 Leptotrichia sp._HMT_463 (0.02%|0%)
SP14 Leptotrichia buccalis (0.59%|0.73%)
SP129 Leptotrichia sp._HMT_909 (0.15%|0.01%)
SP164 Leptotrichia hofstadii (0.22%|0.27%)
SP331 Leptotrichia sp._HMT_223 (0.10%|0.05%)
SP35 Leptotrichia wadei (1.32%|0.78%)
SP142 Leptotrichia sp._HMT_417 (0.12%|0.12%)
SP349 Leptotrichia shahii (0.02%|0.03%)
SP149 Leptotrichia trevisanii (0.18%|0.11%)
SP36 Leptotrichia hongkongensis (0.42%|0.28%)
SP158 Anaerolineae_[G-1] bacterium_HMT_439 (0.04%|0.03%)
SP345 Pyramidobacter piscolens (0%|0.01%)
SP124 Fretibacterium fastidiosum (0.26%|0.32%)
SP157 Fretibacterium sp._HMT_359 (0.08%|0.05%)
SP234 Fretibacterium sp._HMT_361 (0%|0.02%)
SP42 Fretibacterium sp._HMT_360 (0.19%|0.39%)
SP188 Fretibacterium sp._HMT_362 (0.19%|0.24%)
SP281 Slackia exigua (0.04%|0.01%)
SP334 Cryptobacterium curtum (0.02%|0.00%)
SP55 Lancefieldella parvula (0.11%|0.20%)
SP300 Atopobium sp._HMT_199 (0.03%|0.01%)
SP13 Lancefieldella rimae (0.16%|0.21%)
SP397 Olsenella profusa (0.01%|0%)
SP410 Olsenella phocaeensis (0%|0.03%)
SP168 Olsenella sp._HMT_807 (0.74%|0.59%)
SP328 Olsenella uli (0.15%|0.16%)
SP317 Bifidobacterium dentium (0.06%|0.01%)
SP395 Bifidobacterium longum (0%|0.04%)
SP416 Parascardovia denticolens (0.01%|0.06%)
SP84 Scardovia wiggsiae (0.31%|0.17%)
SP374 Propionibacterium acidifaciens (0.04%|0.01%)
SP399 Propionibacteriaceae_[G-1] bacterium_HMT_915 (0%|0.03%)
SP115 Arachnia propionica (2.77%|2.52%)
SP140 Arachnia rubra (0.14%|0.32%)
SP138 Corynebacterium durum (0.67%|0.49%)
SP1 Corynebacterium matruchotii (3.68%|3.25%)
SP267 Rothia mucilaginosa (0.03%|0.08%)
SP195 Rothia dentocariosa (2.44%|4.51%)
SP46 Rothia aeria (1.09%|0.76%)
SP58 Peptidiphaga sp._HMT_183 (4.56%|3.62%)
SP214 Peptidiphaga gingivicola (0.32%|0.13%)
SP152 Schaalia georgiae (0.19%|0.13%)
SP143 Schaalia sp._HMT_877 (0.23%|0.09%)
SP166 Schaalia sp._HMT_178 (0.28%|0.20%)
SP57 Schaalia cardiffensis (0.19%|0.01%)
SP113 Schaalia meyeri (0.09%|0.15%)
SP402 Schaalia sp._HMT_172 (0.01%|0%)
SP106 Schaalia sp._HMT_180 (0.29%|0.28%)
SP82 Schaalia odontolytica (0.43%|0.56%)
SP202 Schaalia odontolyticus (0%|0.02%)
SP239 Actinomyces oricola (0.02%|0.05%)
SP294 Actinomyces sp._HMT_448 (1.52%|0.50%)
SP217 Actinomyces massiliensis (0.79%|0.79%)
SP39 Actinomyces israelii (0.34%|0.43%)
SP174 Actinomyces sp._HMT_897 (0.80%|0.67%)
SP249 Actinomyces sp._HMT_414 (0.31%|0.38%)
SP148 Actinomyces dentalis (1.88%|1.74%)
SP56 Actinomyces sp._HMT_896 (0.39%|0.35%)
SP177 Actinomyces sp._HMT_525 (0.78%|0.57%)
SP186 Actinomyces timonensis (0.09%|0.04%)
SP15 Actinomyces johnsonii (0.43%|0.28%)
SP91 Actinomyces sp._HMT_171 (0.39%|0.27%)
SP61 Actinomyces sp._HMT_170 (0.96%|1.74%)
SP52 Actinomyces naeslundii (1.25%|1.23%)
SP227 Actinomyces viscosus (0%|0.03%)
SP21 Actinomyces sp._HMT_169 (0.62%|1.59%)
SP10 Actinomyces sp._HMT_175 (0.57%|0.57%)
SP89 Actinomyces oris (0.66%|0.55%)
SP348 Treponema sp._HMT_239 (0.04%|0.01%)
SP389 Treponema sp._HMT_227 (0.00%|0.01%)
SP268 Treponema sp._HMT_234 (0.07%|0.06%)
SP167 Treponema sp._HMT_235 (0.00%|0.01%)
SP387 Treponema sp._HMT_226 (0.02%|0.00%)
SP163 Treponema sp._HMT_238 (0.10%|0.11%)
SP264 Treponema sp._HMT_228 (0%|0.01%)
SP254 Treponema sp._HMT_230 (0.06%|0.02%)
SP134 Treponema sp._HMT_237 (0.24%|0.23%)
SP338 Treponema vincentii (0.03%|0.03%)
SP324 Treponema sp._HMT_231 (0.12%|0.16%)
SP265 Treponema medium (0.07%|0.01%)
SP38 Treponema denticola (0.14%|0.21%)
SP380 Treponema sp._HMT_247 (0.03%|0.01%)
SP393 Treponema sp._HMT_254 (0.01%|0.01%)
SP332 Treponema sp._HMT_253 (0%|0.01%)
SP310 Treponema sp._HMT_256 (0.02%|0.01%)
SP200 Treponema sp._HMT_951 (0.03%|0.02%)
SP225 Treponema sp._HMT_490 (0.01%|0.01%)
SP263 Treponema sp._HMT_258 (0.06%|0.01%)
SP282 Treponema maltophilum (0.08%|0.09%)
SP360 Treponema lecithinolyticum (0.07%|0.07%)
SP376 Treponema sp._HMT_270 (0.01%|0.02%)
SP337 Treponema parvum (0.00%|0.00%)
SP212 Treponema socranskii (0.47%|0.39%)
SP74 Treponema sp._HMT_268 (0.09%|0.06%)
SP64 Treponema sp._HMT_927 (0.04%|0.04%)
SP275 Treponema sp._HMT_263 (0.01%|0.04%)
SP76 Treponema sp._HMT_262 (0.09%|0.06%)
SP369 Treponema amylovorum (0.02%|0.01%)
SP176 Treponema sp._HMT_257 (0.12%|0.04%)
SP343 Desulfobulbus sp._HMT_041 (0.04%|0.08%)
SP407 Bilophila wadsworthia (0.00%|0%)
SP394 Desulfomicrobium orale (0.00%|0.04%)
SP94 Campylobacter concisus (0.30%|0.31%)
SP375 Campylobacter rectus (0%|0.01%)
SP224 Campylobacter showae (0.62%|0.36%)
SP20 Campylobacter gracilis (0.47%|0.40%)
SP241 Campylobacter sp._HMT_044 (0.01%|0.03%)
SP105 Cardiobacterium valvarum (0.23%|0.17%)
SP27 Cardiobacterium hominis (0.45%|0.46%)
SP373 Haemophilus paraphrohaemolyticus (0%|0.04%)
SP329 Haemophilus parahaemolyticus (0.01%|0.01%)
SP17 Haemophilus parainfluenzae (0.92%|2.56%)
SP405 Haemophilus sputorum (0.01%|0%)
SP392 Haemophilus sp._HMT_036 (0.01%|0%)
SP154 Haemophilus haemolyticus (0.06%|0.06%)
SP289 Haemophilus aegyptius (0.00%|0%)
SP203 Aggregatibacter sp._HMT_458 (0.22%|0.17%)
SP181 Aggregatibacter segnis (0.11%|0.35%)
SP320 Aggregatibacter sp._HMT_512 (0.07%|0.08%)
SP153 Aggregatibacter sp._HMT_513 (0.04%|0.26%)
SP313 Aggregatibacter sp._HMT_898 (0.07%|0.01%)
SP226 Aggregatibacter sp._HMT_949 (0%|0.02%)
SP150 Aggregatibacter aphrophilus (0.09%|0.17%)
SP253 Aggregatibacter actinomycetemcomitans (0.09%|0.04%)
SP233 Ottowia sp._HMT_894 (0.12%|0.18%)
SP26 Lautropia mirabilis (1.30%|2.75%)
SP303 Lautropia dentalis (0%|0.06%)
SP411 Neisseria sp._HMT_499 (0%|0.02%)
SP45 Kingella oralis (0.34%|0.35%)
SP293 Neisseria sp._HMT_018 (0.07%|0.04%)
SP196 Neisseria bacilliformis (0.06%|0.06%)
SP72 Neisseria elongata (0.46%|0.80%)
SP260 Eikenella sp._HMT_011 (0%|0.00%)
SP122 Neisseria oralis (0.40%|0.27%)
SP130 Kingella denitrificans (0.05%|0.02%)
SP96 Eikenella corrodens (0.33%|0.41%)
SP236 Neisseria mucosa (0.12%|1.70%)
SP417 Neisseria flava (0%|0.00%)
SP40 Neisseria sicca (0.32%|2.01%)
SP11 Neisseria subflava (0.11%|0.28%)
SP305 Neisseria cinerea (0.04%|0.03%)
SP37 Neisseria flavescens (0.23%|0.16%)
SP232 Ruminococcaceae_[G-2] bacterium_HMT_085 (0.01%|0.04%)
SP59 Ruminococcaceae_[G-1] bacterium_HMT_075 (0.13%|0.17%)
SP242 Lachnospiraceae_[G-8] bacterium_HMT_500 (0.02%|0.06%)
SP278 Butyrivibrio sp._HMT_080 (0.02%|0.03%)
SP347 Catonella sp._HMT_451 (0.05%|0.02%)
SP12 Catonella morbi (0.42%|0.35%)
SP370 Catonella sp._HMT_164 (0%|0.01%)
SP243 Lachnospiraceae_[G-7] bacterium_HMT_086 (0.02%|0%)
SP4 Oribacterium sp._HMT_078 (0.48%|0.26%)
SP49 Oribacterium sinus (0.05%|0.06%)
SP92 Shuttleworthia satelles (0%|0.09%)
SP354 Stomatobaculum sp._HMT_373 (0.01%|0.05%)
SP98 Stomatobaculum longum (0.27%|0.25%)
SP198 Stomatobaculum sp._HMT_097 (0.03%|0.03%)
SP201 Lachnospiraceae_[G-2] bacterium_HMT_096 (0%|0.07%)
SP139 Lachnospiraceae_[G-3] bacterium_HMT_100 (0.40%|0.44%)
SP258 Johnsonella ignava (0.20%|0.16%)
SP23 Johnsonella sp._HMT_166 (0.02%|0.05%)
SP390 Lachnoanaerobaculum sp._HMT_089 (0.01%|0%)
SP291 Lachnoanaerobaculum sp._HMT_083 (0.02%|0.02%)
SP192 Lachnoanaerobaculum saburreum (0.64%|0.35%)
SP128 Lachnoanaerobaculum orale (0.18%|0.20%)
SP97 Lachnoanaerobaculum umeaense (0.03%|0.08%)
SP215 Lachnoanaerobaculum gingivalis (0.05%|0.06%)
SP180 Clostridiales_[F-1][G-1] bacterium_HMT_093 (0.08%|0.05%)
SP235 Pseudoramibacter alactolyticus (0.11%|0.04%)
SP285 Peptococcus sp._HMT_168 (0.03%|0.05%)
SP184 Peptococcus sp._HMT_167 (0.24%|0.12%)
SP189 Peptoniphilaceae_[G-1] bacterium_HMT_113 (0.02%|0.08%)
SP418 Peptoniphilus lacrimalis (0.01%|0%)
SP123 Parvimonas sp._HMT_393 (0%|0.00%)
SP41 Parvimonas micra (0.97%|0.56%)
SP88 Parvimonas sp._HMT_110 (0.20%|0.12%)
SP221 Peptostreptococcus stomatis (0.41%|0.24%)
SP256 Filifactor alocis (0.25%|0.24%)
SP79 Peptoanaerobacter [Eubacterium] yurii (0.27%|0.33%)
SP366 Peptostreptococcaceae_[XI][G-7] bacterium_HMT_081 (0.02%|0%)
SP356 Peptoanaerobacter stomatis (0.02%|0.02%)
SP159 Peptostreptococcaceae_[XI][G-5] [Eubacterium]_saphenum (0.10%|0.11%)
SP170 Peptostreptococcaceae_[XI][G-3] bacterium_HMT_495 (0.03%|0.00%)
SP160 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0.12%|0.13%)
SP351 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_minutum (0.03%|0.03%)
SP147 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatum (0.14%|0.12%)
SP396 Peptostreptococcaceae_[XI][G-1] bacterium_HMT_383 (0.00%|0.00%)
SP244 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_infirmum (0.21%|0.09%)
SP259 Peptostreptococcaceae_[XI][G-4] bacterium_HMT_369 (0.01%|0.10%)
SP318 Peptostreptococcaceae_[XI][G-4] bacterium_HMT_103 (0.06%|0.01%)
SP365 Peptostreptococcaceae_[XI][G-2] bacterium_HMT_091 (0.03%|0%)
SP280 Mogibacterium timidum (0.07%|0.07%)
SP302 Mogibacterium neglectum (0.08%|0.09%)
SP173 Mogibacterium vescum (0.16%|0.06%)
SP359 Mollicutes_[G-1] bacterium_HMT_504 (0.02%|0.00%)
SP322 Erysipelotrichaceae_[G-1] bacterium_HMT_905 (0.04%|0.03%)
SP204 Erysipelotrichaceae_[G-1] bacterium_HMT_904 (0.02%|0.00%)
SP339 Bulleidia extructa (0.03%|0%)
SP18 Solobacterium moorei (0.26%|0.18%)
SP107 Gemella sp._HMT_928 (0.01%|0%)
SP63 Gemella morbillorum (0.68%|0.65%)
SP30 Gemella sanguinis (0.01%|0.03%)
SP179 Gemella haemolysans (0.12%|0.17%)
SP182 Ligilactobacillus salivarius (0.12%|0.05%)
SP90 Abiotrophia defectiva (0.17%|0.40%)
SP161 Granulicatella adiacens (0.63%|0.72%)
SP413 Granulicatella elegans (0.01%|0.02%)
SP3 Streptococcus mutans (1.12%|0.14%)
SP403 Streptococcus sobrinus (0.05%|0.02%)
SP304 Streptococcus massiliensis (0.01%|0%)
SP16 Streptococcus anginosus (0.71%|0.35%)
SP125 Streptococcus constellatus (0.43%|0.12%)
SP165 Streptococcus intermedius (0.80%|0.39%)
SP121 Streptococcus gordonii (0.79%|0.69%)
SP272 Streptococcus sinensis (0.07%|0.14%)
SP261 Streptococcus cristatus_clade_886 (0.00%|0.01%)
SP323 Streptococcus cristatus_clade_578 (0.01%|0.03%)
SP5 Streptococcus cristatus (0.74%|0.93%)
SP301 Streptococcus sp._HMT_056 (0.14%|0.18%)
SP341 Streptococcus peroris (0.02%|0%)
SP327 Streptococcus vestibularis (0.03%|0.02%)
SP132 Streptococcus salivarius (0.04%|0.18%)
SP100 Streptococcus sanguinis (1.17%|1.68%)
SP379 Streptococcus oralis_subsp._tigurinus_clade_071 (0%|0.00%)
SP220 Streptococcus oralis_subsp._tigurinus_clade_070 (0.40%|0.23%)
SP277 Streptococcus oralis_subsp._dentisani_clade_058 (0.11%|0.04%)
SP208 Streptococcus infantis_clade_431 (0.02%|0.02%)
SP346 Streptococcus periodonticum (0%|0.01%)
SP43 Streptococcus mitis (1.94%|3.58%)
SP408 Streptococcus sp._HMT_423 (0.01%|0%)
SP101 Streptococcus sp._HMT_064 (0.04%|0.16%)
SP69 Streptococcus chosunense (0.11%|0.13%)
SP48 Streptococcus oralis (1.39%|1.36%)
SP274 Streptococcus downii (0.01%|0.05%)
SP207 Streptococcus sp._HMT_074 (0%|0.01%)
SP321 Streptococcus australis (0.01%|0.03%)
SP326 Streptococcus sp._HMT_066 (0.00%|0.02%)
SP109 Streptococcus parasanguinis_clade_411 (0.10%|0.17%)
SP223 Streptococcus parasanguinis (0.00%|0.01%)
SP355 Eggerthia catenaformis (0.01%|0.03%)
SP367 Veillonella atypica (0.01%|0.05%)
SP66 Veillonella parvula (0.87%|0.61%)
SP67 Veillonella dispar (0.00%|0.01%)
SP197 Dialister sp._HMT_119 (0.01%|0.01%)
SP75 Dialister invisus (0.57%|0.37%)
SP104 Megasphaera micronuciformis (0.11%|0.05%)
SP319 Colibacter massiliensis (0.06%|0.07%)
SP145 Anaeroglobus geminatus (0.22%|0.13%)
SP330 Veillonellaceae_[G-1] bacterium_HMT_129 (0.02%|0.01%)
SP178 Veillonellaceae_[G-1] bacterium_HMT_132 (0.04%|0.05%)
SP102 Veillonellaceae_[G-1] bacterium_HMT_150 (0.07%|0.02%)
SP146 Veillonellaceae_[G-1] bacterium_HMT_135 (0%|0.01%)
SP169 Veillonellaceae_[G-1] bacterium_HMT_148 (0.01%|0%)
SP131 Veillonellaceae_[G-1] bacterium_HMT_155 (0.21%|0.14%)
SP283 Veillonellaceae_[G-1] bacterium_HMT_145 (0.04%|0.05%)
SP190 Mitsuokella sp._HMT_521 (0.02%|0.08%)
SP219 Selenomonas sp._HMT_442 (0.02%|0.06%)
SP151 Selenomonas sp._HMT_134 (0.05%|0.01%)
SP51 Selenomonas sputigena (0.96%|0.66%)
SP364 Selenomonas sp._HMT_149 (0.04%|0.02%)
SP252 Selenomonas sp._HMT_136 (0.08%|0.02%)
SP377 Selenomonas sp._HMT_478 (0.01%|0%)
SP414 Selenomonas sp._HMT_920 (0.01%|0%)
SP230 Selenomonas flueggei (0.06%|0.06%)
SP87 Selenomonas sp._HMT_138 (0.16%|0.06%)
SP29 Selenomonas infelix (0.43%|0.24%)
SP33 Selenomonas noxia (0.80%|0.76%)
SP250 Selenomonas sp._HMT_937 (0.01%|0.03%)
SP314 Selenomonas sp._HMT_133 (0.02%|0.01%)
SP65 Selenomonas sp._HMT_137 (0.08%|0.07%)
SP99 Selenomonas artemidis (0.28%|0.27%)
SP218 Selenomonas sp._HMT_892 (0.04%|0.06%)
SP155 Selenomonas sp._HMT_479 (0.07%|0.04%)
SP273 Selenomonas sp._HMT_388 (0%|0.01%)
SP335 Centipeda periodontii (0.09%|0.02%)
SP311 Selenomonas sp._HMT_481 (0.10%|0.04%)
SP406 Selenomonas sp._HMT_919 (0.02%|0%)
SP229 Selenomonas sp._HMT_126 (0.02%|0.01%)
SP297 Selenomonas sp._HMT_936 (0.02%|0.01%)
SP70 Selenomonas sp._HMT_146 (0.03%|0.04%)
SP116 Selenomonas dianae (0.14%|0.09%)
Day_90
Baseline
substitutions/site
0
0.2
0.4
0.6
0.8