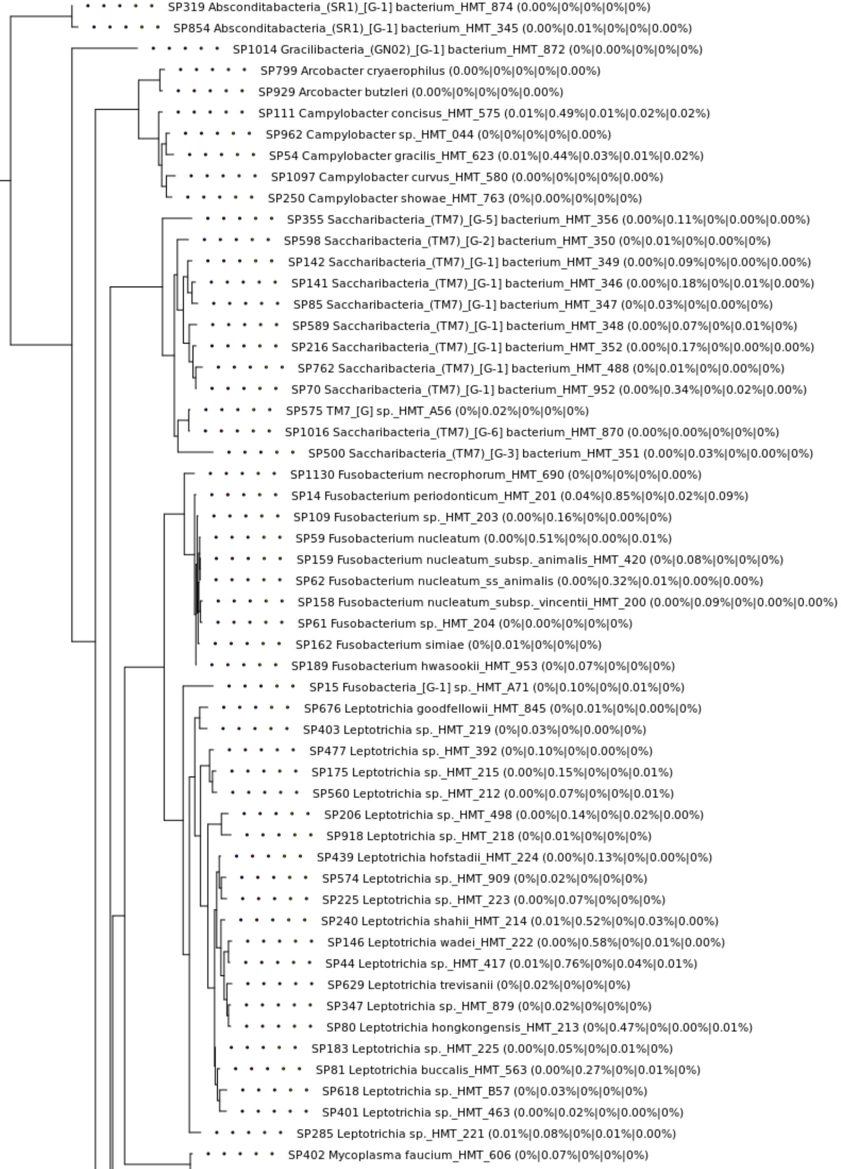
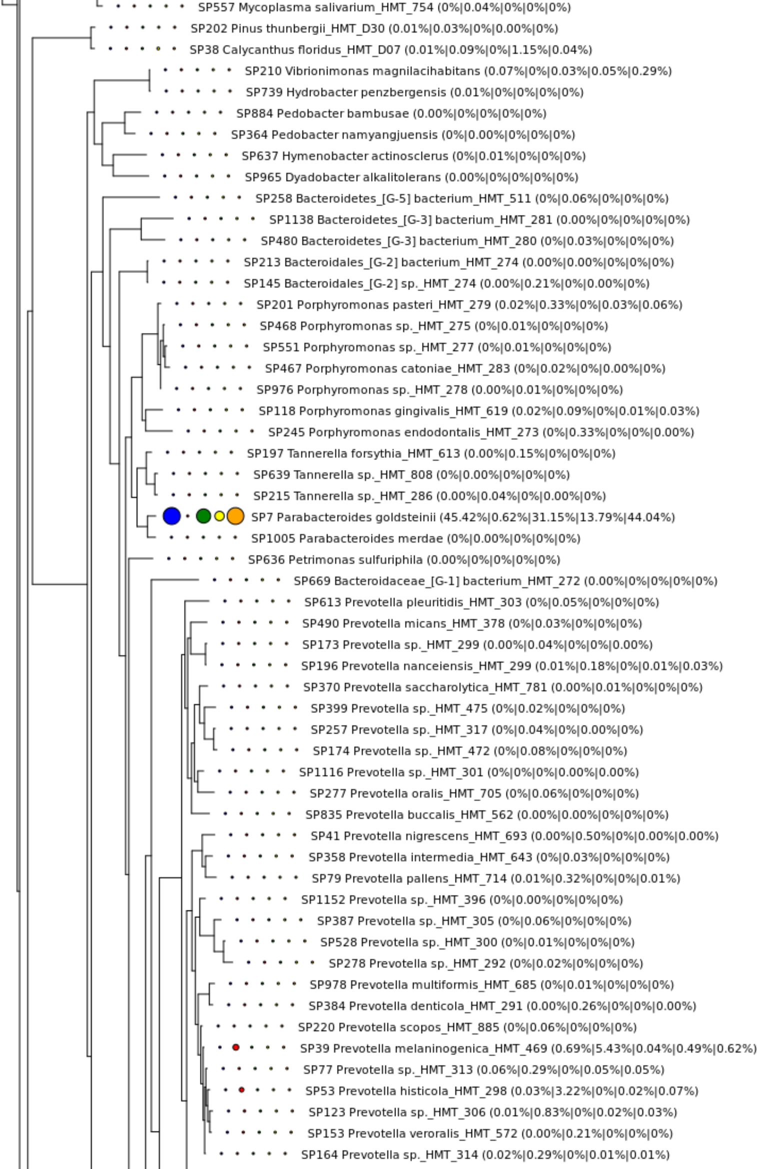
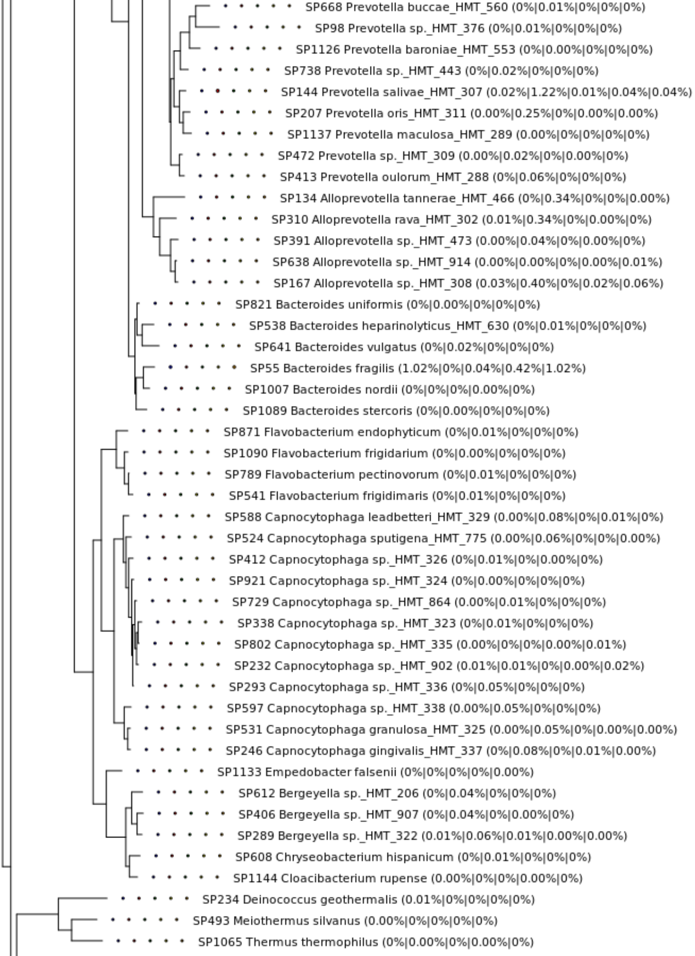
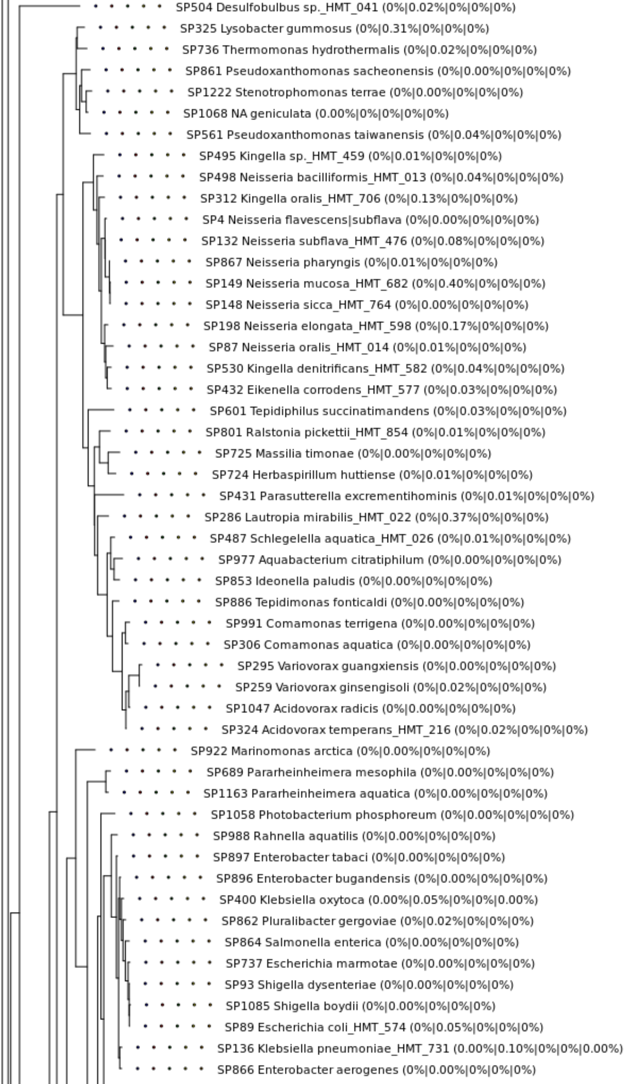
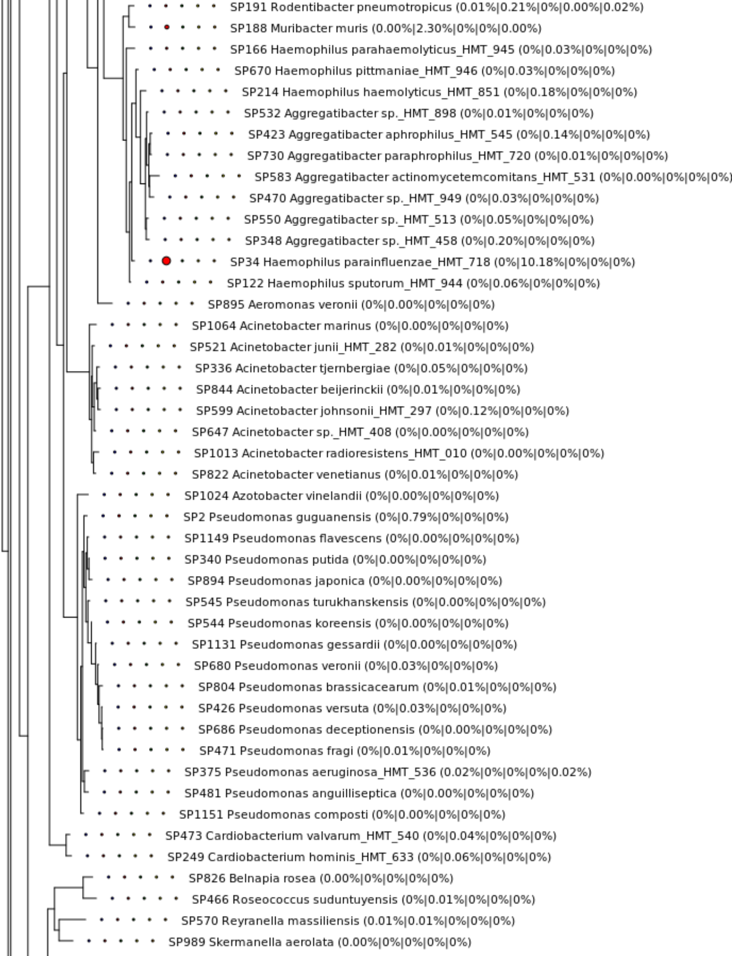
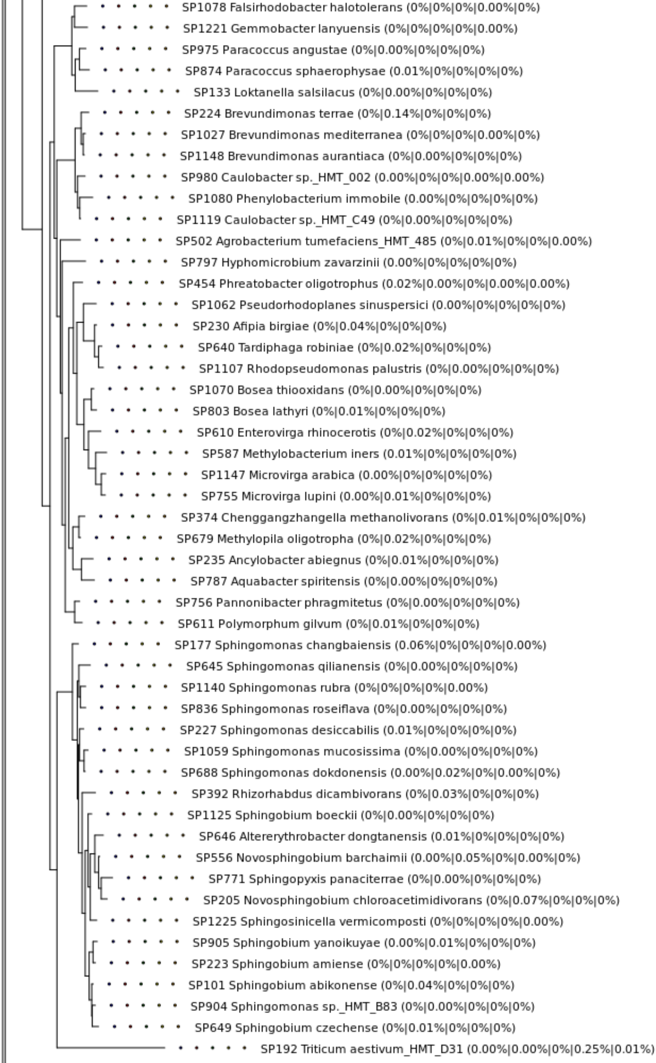

SP319 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0.00%|0%|0%|0%|0%)
SP854 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0.00%|0.01%|0%|0%|0.00%)
SP1014 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872 (0%|0.00%|0%|0%|0%)
SP799 Arcobacter cryaerophilus (0.00%|0%|0%|0%|0.00%)
SP929 Arcobacter butzleri (0.00%|0%|0%|0%|0.00%)
SP111 Campylobacter concisus_HMT_575 (0.01%|0.49%|0.01%|0.02%|0.02%)
SP962 Campylobacter sp._HMT_044 (0%|0%|0%|0%|0.00%)
SP54 Campylobacter gracilis_HMT_623 (0.01%|0.44%|0.03%|0.01%|0.02%)
SP1097 Campylobacter curvus_HMT_580 (0.00%|0%|0%|0%|0.00%)
SP250 Campylobacter showae_HMT_763 (0%|0.00%|0%|0%|0%)
SP355 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0.00%|0.11%|0%|0.00%|0.00%)
SP598 Saccharibacteria_(TM7)_[G-2] bacterium_HMT_350 (0%|0.01%|0%|0.00%|0%)
SP142 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0.00%|0.09%|0%|0.00%|0.00%)
SP141 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0.00%|0.18%|0%|0.01%|0.00%)
SP85 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347 (0%|0.03%|0%|0.00%|0%)
SP589 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0.00%|0.07%|0%|0.01%|0%)
SP216 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0.00%|0.17%|0%|0.00%|0.00%)
SP762 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_488 (0%|0.01%|0%|0.00%|0%)
SP70 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_952 (0.00%|0.34%|0%|0.02%|0.00%)
SP575 TM7_[G] sp._HMT_A56 (0%|0.02%|0%|0%|0%)
SP1016 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.00%|0.00%|0%|0%|0%)
SP500 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0.00%|0.03%|0%|0.00%|0%)
SP1130 Fusobacterium necrophorum_HMT_690 (0%|0%|0%|0%|0.00%)
SP14 Fusobacterium periodonticum_HMT_201 (0.04%|0.85%|0%|0.02%|0.09%)
SP109 Fusobacterium sp._HMT_203 (0.00%|0.16%|0%|0.00%|0%)
SP59 Fusobacterium nucleatum (0.00%|0.51%|0%|0.00%|0.01%)
SP159 Fusobacterium nucleatum_subsp._animalis_HMT_420 (0%|0.08%|0%|0%|0%)
SP62 Fusobacterium nucleatum_ss_animalis (0.00%|0.32%|0.01%|0.00%|0.00%)
SP158 Fusobacterium nucleatum_subsp._vincentii_HMT_200 (0.00%|0.09%|0%|0.00%|0.00%)
SP61 Fusobacterium sp._HMT_204 (0%|0.00%|0%|0%|0%)
SP162 Fusobacterium simiae (0%|0.01%|0%|0%|0%)
SP189 Fusobacterium hwasookii_HMT_953 (0%|0.07%|0%|0%|0%)
SP15 Fusobacteria_[G-1] sp._HMT_A71 (0%|0.10%|0%|0.01%|0%)
SP676 Leptotrichia goodfellowii_HMT_845 (0%|0.01%|0%|0.00%|0%)
SP403 Leptotrichia sp._HMT_219 (0%|0.03%|0%|0.00%|0%)
SP477 Leptotrichia sp._HMT_392 (0%|0.10%|0%|0.00%|0%)
SP175 Leptotrichia sp._HMT_215 (0.00%|0.15%|0%|0%|0.01%)
SP560 Leptotrichia sp._HMT_212 (0.00%|0.07%|0%|0%|0.01%)
SP206 Leptotrichia sp._HMT_498 (0.00%|0.14%|0%|0.02%|0.00%)
SP918 Leptotrichia sp._HMT_218 (0%|0.01%|0%|0%|0%)
SP439 Leptotrichia hofstadii_HMT_224 (0.00%|0.13%|0%|0.00%|0%)
SP574 Leptotrichia sp._HMT_909 (0%|0.02%|0%|0%|0%)
SP225 Leptotrichia sp._HMT_223 (0.00%|0.07%|0%|0%|0%)
SP240 Leptotrichia shahii_HMT_214 (0.01%|0.52%|0%|0.03%|0.00%)
SP146 Leptotrichia wadei_HMT_222 (0.00%|0.58%|0%|0.01%|0.00%)
SP44 Leptotrichia sp._HMT_417 (0.01%|0.76%|0%|0.04%|0.01%)
SP629 Leptotrichia trevisanii (0%|0.02%|0%|0%|0%)
SP347 Leptotrichia sp._HMT_879 (0%|0.02%|0%|0%|0%)
SP80 Leptotrichia hongkongensis_HMT_213 (0%|0.47%|0%|0.00%|0.01%)
SP183 Leptotrichia sp._HMT_225 (0.00%|0.05%|0%|0.01%|0%)
SP81 Leptotrichia buccalis_HMT_563 (0.00%|0.27%|0%|0.01%|0%)
SP618 Leptotrichia sp._HMT_B57 (0%|0.03%|0%|0%|0%)
SP401 Leptotrichia sp._HMT_463 (0.00%|0.02%|0%|0.00%|0%)
SP285 Leptotrichia sp._HMT_221 (0.01%|0.08%|0%|0.01%|0.00%)
SP402 Mycoplasma faucium_HMT_606 (0%|0.07%|0%|0%|0%)

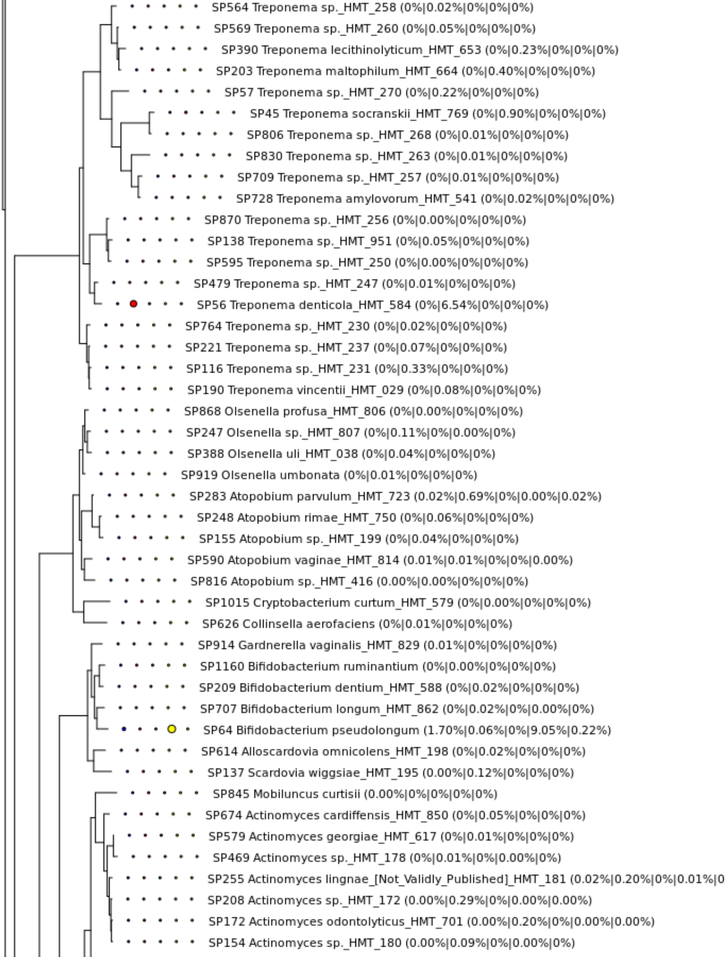

SP557 Mycoplasma salivarium_HMT_754 (0%|0.04%|0%|0%|0%)
SP202 Pinus thunbergii_HMT_D30 (0.01%|0.03%|0%|0.00%|0%)
SP38 Calycanthus floridus_HMT_D07 (0.01%|0.09%|0%|1.15%|0.04%)
SP210 Vibrionimonas magnilacihabitans (0.07%|0%|0.03%|0.05%|0.29%)
SP739 Hydrobacter penzbergensis (0.01%|0%|0%|0%|0%)
SP884 Pedobacter bambusae (0.00%|0%|0%|0%|0%)
SP364 Pedobacter namyangjuensis (0%|0.00%|0%|0%|0%)
SP637 Hymenobacter actinosclerus (0%|0.01%|0%|0%|0%)
SP965 Dyadobacter alkalitolerans (0.00%|0%|0%|0%|0%)
SP258 Bacteroidetes_[G-5] bacterium_HMT_511 (0%|0.06%|0%|0%|0%)
SP1138 Bacteroidetes_[G-3] bacterium_HMT_281 (0.00%|0%|0%|0%|0%)
SP480 Bacteroidetes_[G-3] bacterium_HMT_280 (0%|0.03%|0%|0%|0%)
SP213 Bacteroidales_[G-2] bacterium_HMT_274 (0.00%|0.00%|0%|0%|0%)
SP145 Bacteroidales_[G-2] sp._HMT_274 (0.00%|0.21%|0%|0.00%|0%)
SP201 Porphyromonas pasteri_HMT_279 (0.02%|0.33%|0%|0.03%|0.06%)
SP468 Porphyromonas sp._HMT_275 (0%|0.01%|0%|0%|0%)
SP551 Porphyromonas sp._HMT_277 (0%|0.01%|0%|0%|0%)
SP467 Porphyromonas catoniae_HMT_283 (0%|0.02%|0%|0.00%|0%)
SP976 Porphyromonas sp._HMT_278 (0.00%|0.01%|0%|0%|0%)
SP118 Porphyromonas gingivalis_HMT_619 (0.02%|0.09%|0%|0.01%|0.03%)
SP245 Porphyromonas endodontalis_HMT_273 (0%|0.33%|0%|0%|0.00%)
SP197 Tannerella forsythia_HMT_613 (0.00%|0.15%|0%|0%|0%)
SP639 Tannerella sp._HMT_808 (0%|0.00%|0%|0%|0%)
SP215 Tannerella sp._HMT_286 (0.00%|0.04%|0%|0.00%|0%)
SP7 Parabacteroides goldsteinii (45.42%|0.62%|31.15%|13.79%|44.04%)
SP1005 Parabacteroides merdae (0%|0.00%|0%|0%|0%)
SP636 Petrimonas sulfuriphila (0.00%|0%|0%|0%|0%)
SP669 Bacteroidaceae_[G-1] bacterium_HMT_272 (0.00%|0%|0%|0%|0%)
SP613 Prevotella pleuritidis_HMT_303 (0%|0.05%|0%|0%|0%)
SP490 Prevotella micans_HMT_378 (0%|0.03%|0%|0%|0%)
SP173 Prevotella sp._HMT_299 (0.00%|0.04%|0%|0%|0.00%)
SP196 Prevotella nanceiensis_HMT_299 (0.01%|0.18%|0%|0.01%|0.03%)
SP370 Prevotella saccharolytica_HMT_781 (0.00%|0.01%|0%|0%|0%)
SP399 Prevotella sp._HMT_475 (0%|0.02%|0%|0%|0%)
SP257 Prevotella sp._HMT_317 (0%|0.04%|0%|0.00%|0%)
SP174 Prevotella sp._HMT_472 (0%|0.08%|0%|0%|0%)
SP1116 Prevotella sp._HMT_301 (0%|0%|0%|0.00%|0.00%)
SP277 Prevotella oralis_HMT_705 (0%|0.06%|0%|0%|0%)
SP835 Prevotella buccalis_HMT_562 (0.00%|0.00%|0%|0%|0%)
SP41 Prevotella nigrescens_HMT_693 (0.00%|0.50%|0%|0.00%|0.00%)
SP358 Prevotella intermedia_HMT_643 (0%|0.03%|0%|0%|0%)
SP79 Prevotella pallens_HMT_714 (0.01%|0.32%|0%|0%|0.01%)
SP1152 Prevotella sp._HMT_396 (0%|0.00%|0%|0%|0%)
SP387 Prevotella sp._HMT_305 (0%|0.06%|0%|0%|0%)
SP528 Prevotella sp._HMT_300 (0%|0.01%|0%|0%|0%)
SP278 Prevotella sp._HMT_292 (0%|0.02%|0%|0%|0%)
SP978 Prevotella multiformis_HMT_685 (0%|0.01%|0%|0%|0%)
SP384 Prevotella denticola_HMT_291 (0.00%|0.26%|0%|0%|0.00%)
SP220 Prevotella scopos_HMT_885 (0%|0.06%|0%|0%|0%)
SP39 Prevotella melaninogenica_HMT_469 (0.69%|5.43%|0.04%|0.49%|0.62%)
SP77 Prevotella sp._HMT_313 (0.06%|0.29%|0%|0.05%|0.05%)
SP53 Prevotella histicola_HMT_298 (0.03%|3.22%|0%|0.02%|0.07%)
SP123 Prevotella sp._HMT_306 (0.01%|0.83%|0%|0.02%|0.03%)
SP153 Prevotella veroralis_HMT_572 (0.00%|0.21%|0%|0%|0%)
SP164 Prevotella sp._HMT_314 (0.02%|0.29%|0%|0.01%|0.01%)

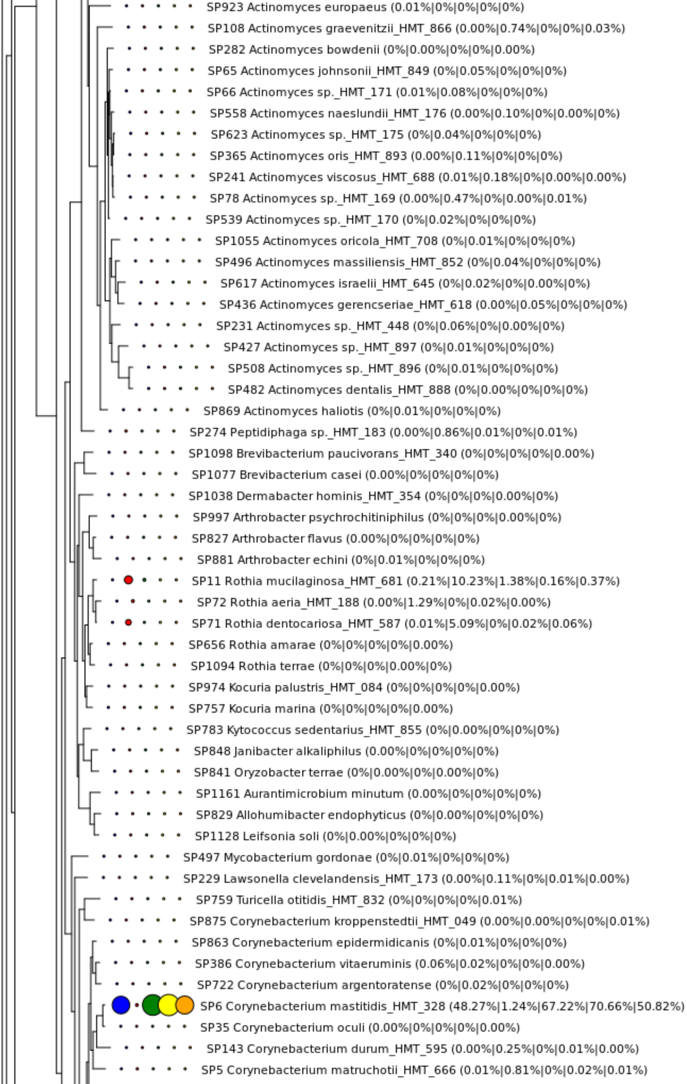

SP668 Prevotella buccae_HMT_560 (0%|0.01%|0%|0%|0%)
SP98 Prevotella sp._HMT_376 (0%|0.01%|0%|0%|0%)
SP1126 Prevotella baroniae_HMT_553 (0%|0.00%|0%|0%|0%)
SP738 Prevotella sp._HMT_443 (0%|0.02%|0%|0%|0%)
SP144 Prevotella salivae_HMT_307 (0.02%|1.22%|0.01%|0.04%|0.04%)
SP207 Prevotella oris_HMT_311 (0.00%|0.25%|0%|0.00%|0.00%)
SP1137 Prevotella maculosa_HMT_289 (0.00%|0%|0%|0%|0%)
SP472 Prevotella sp._HMT_309 (0.00%|0.02%|0%|0.00%|0%)
SP413 Prevotella oulorum_HMT_288 (0%|0.06%|0%|0%|0%)
SP134 Alloprevotella tannerae_HMT_466 (0%|0.34%|0%|0%|0.00%)
SP310 Alloprevotella rava_HMT_302 (0.01%|0.34%|0%|0.00%|0%)
SP391 Alloprevotella sp._HMT_473 (0.00%|0.04%|0%|0.00%|0%)
SP638 Alloprevotella sp._HMT_914 (0.00%|0.00%|0%|0.00%|0.01%)
SP167 Alloprevotella sp._HMT_308 (0.03%|0.40%|0%|0.02%|0.06%)
SP821 Bacteroides uniformis (0%|0.00%|0%|0%|0%)
SP538 Bacteroides heparinolyticus_HMT_630 (0%|0.01%|0%|0%|0%)
SP641 Bacteroides vulgatus (0%|0.02%|0%|0%|0%)
SP55 Bacteroides fragilis (1.02%|0%|0.04%|0.42%|1.02%)
SP1007 Bacteroides nordii (0%|0%|0%|0.00%|0%)
SP1089 Bacteroides stercoris (0%|0.00%|0%|0%|0%)
SP871 Flavobacterium endophyticum (0%|0.01%|0%|0%|0%)
SP1090 Flavobacterium frigidarium (0%|0.00%|0%|0%|0%)
SP789 Flavobacterium pectinovorum (0%|0.01%|0%|0%|0%)
SP541 Flavobacterium frigidimaris (0%|0.01%|0%|0%|0%)
SP588 Capnocytophaga leadbetteri_HMT_329 (0.00%|0.08%|0%|0.01%|0%)
SP524 Capnocytophaga sputigena_HMT_775 (0.00%|0.06%|0%|0%|0.00%)
SP412 Capnocytophaga sp._HMT_326 (0%|0.01%|0%|0.00%|0%)
SP921 Capnocytophaga sp._HMT_324 (0%|0.00%|0%|0%|0%)
SP729 Capnocytophaga sp._HMT_864 (0.00%|0.01%|0%|0%|0%)
SP338 Capnocytophaga sp._HMT_323 (0%|0.01%|0%|0%|0%)
SP802 Capnocytophaga sp._HMT_335 (0.00%|0%|0%|0.00%|0.01%)
SP232 Capnocytophaga sp._HMT_902 (0.01%|0.01%|0%|0.00%|0.02%)
SP293 Capnocytophaga sp._HMT_336 (0%|0.05%|0%|0%|0%)
SP597 Capnocytophaga sp._HMT_338 (0.00%|0.05%|0%|0%|0%)
SP531 Capnocytophaga granulosa_HMT_325 (0.00%|0.05%|0%|0.00%|0.00%)
SP246 Capnocytophaga gingivalis_HMT_337 (0%|0.08%|0%|0.01%|0.00%)
SP1133 Empedobacter falsenii (0%|0%|0%|0%|0.00%)
SP612 Bergeyella sp._HMT_206 (0%|0.04%|0%|0%|0%)
SP406 Bergeyella sp._HMT_907 (0%|0.04%|0%|0.00%|0%)
SP289 Bergeyella sp._HMT_322 (0.01%|0.06%|0.01%|0.00%|0.00%)
SP608 Chryseobacterium hispanicum (0%|0.01%|0%|0%|0%)
SP1144 Cloacibacterium rupense (0.00%|0%|0%|0.00%|0%)
SP234 Deinococcus geothermalis (0.01%|0%|0%|0%|0%)
SP493 Meiothermus silvanus (0.00%|0%|0%|0%|0%)
SP1065 Thermus thermophilus (0%|0.00%|0%|0.00%|0%)

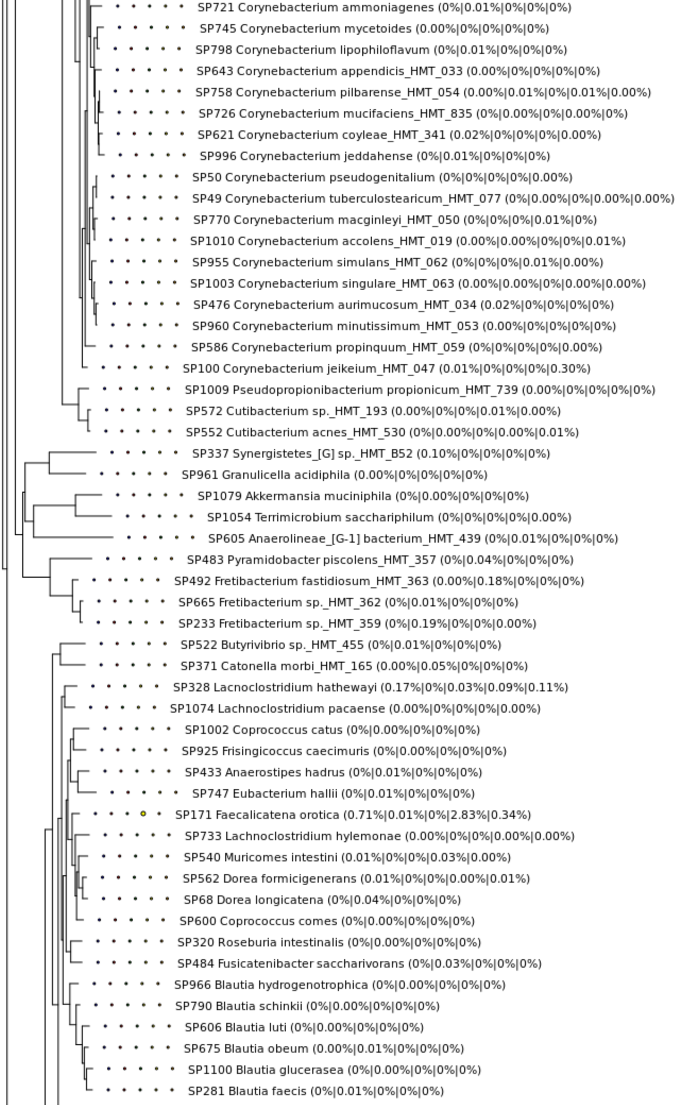

SP504 Desulfobulbus sp._HMT_041 (0%|0.02%|0%|0%|0%)
SP325 Lysobacter gummosus (0%|0.31%|0%|0%|0%)
SP736 Thermomonas hydrothermalis (0%|0.02%|0%|0%|0%)
SP861 Pseudoxanthomonas sacheonensis (0%|0.00%|0%|0%|0%)
SP1222 Stenotrophomonas terrae (0%|0.00%|0%|0%|0%)
SP1068 NA geniculata (0.00%|0%|0%|0%|0%)
SP561 Pseudoxanthomonas taiwanensis (0%|0.04%|0%|0%|0%)
SP495 Kingella sp._HMT_459 (0%|0.01%|0%|0%|0%)
SP498 Neisseria bacilliformis_HMT_013 (0%|0.04%|0%|0%|0%)
SP312 Kingella oralis_HMT_706 (0%|0.13%|0%|0%|0%)
SP4 Neisseria flavescens|subflava (0%|0.00%|0%|0%|0%)
SP132 Neisseria subflava_HMT_476 (0%|0.08%|0%|0%|0%)
SP867 Neisseria pharyngis (0%|0.01%|0%|0%|0%)
SP149 Neisseria mucosa_HMT_682 (0%|0.40%|0%|0%|0%)
SP148 Neisseria sicca_HMT_764 (0%|0.00%|0%|0%|0%)
SP198 Neisseria elongata_HMT_598 (0%|0.17%|0%|0%|0%)
SP87 Neisseria oralis_HMT_014 (0%|0.01%|0%|0%|0%)
SP530 Kingella denitrificans_HMT_582 (0%|0.04%|0%|0%|0%)
SP432 Eikenella corrodens_HMT_577 (0%|0.03%|0%|0%|0%)
SP601 Tepidiphilus succinatimandens (0%|0.03%|0%|0%|0%)
SP801 Ralstonia pickettii_HMT_854 (0%|0.01%|0%|0%|0%)
SP725 Massilia timonae (0%|0.00%|0%|0%|0%)
SP724 Herbaspirillum huttiense (0%|0.01%|0%|0%|0%)
SP431 Parasutterella excrementihominis (0%|0.01%|0%|0%|0%)
SP286 Lautropia mirabilis_HMT_022 (0%|0.37%|0%|0%|0%)
SP487 Schlegelella aquatica_HMT_026 (0%|0.01%|0%|0%|0%)
SP977 Aquabacterium citratiphilum (0%|0.00%|0%|0%|0%)
SP853 Ideonella paludis (0%|0.00%|0%|0%|0%)
SP886 Tepidimonas fonticaldi (0%|0.00%|0%|0%|0%)
SP991 Comamonas terrigena (0%|0.00%|0%|0%|0%)
SP306 Comamonas aquatica (0%|0.00%|0%|0%|0%)
SP295 Variovorax guangxiensis (0%|0.00%|0%|0%|0%)
SP259 Variovorax ginsengisoli (0%|0.02%|0%|0%|0%)
SP1047 Acidovorax radicis (0%|0.00%|0%|0%|0%)
SP324 Acidovorax temperans_HMT_216 (0%|0.02%|0%|0%|0%)
SP922 Marinomonas arctica (0%|0.00%|0%|0%|0%)
SP689 Pararheinheimera mesophila (0%|0.00%|0%|0%|0%)
SP1163 Pararheinheimera aquatica (0%|0.00%|0%|0%|0%)
SP1058 Photobacterium phosphoreum (0%|0.00%|0%|0%|0%)
SP988 Rahnella aquatilis (0%|0.00%|0%|0%|0%)
SP897 Enterobacter tabaci (0%|0.00%|0%|0%|0%)
SP896 Enterobacter bugandensis (0%|0.00%|0%|0%|0%)
SP400 Klebsiella oxytoca (0.00%|0.05%|0%|0%|0.00%)
SP862 Pluralibacter gergoviae (0%|0.02%|0%|0%|0%)
SP864 Salmonella enterica (0%|0.00%|0%|0%|0%)
SP737 Escherichia marmotae (0%|0.00%|0%|0%|0%)
SP93 Shigella dysenteriae (0%|0.00%|0%|0%|0%)
SP1085 Shigella boydii (0%|0.00%|0%|0%|0%)
SP89 Escherichia coli_HMT_574 (0%|0.05%|0%|0%|0%)
SP136 Klebsiella pneumoniae_HMT_731 (0.00%|0.10%|0%|0%|0.00%)
SP866 Enterobacter aerogenes (0%|0.00%|0%|0%|0%)

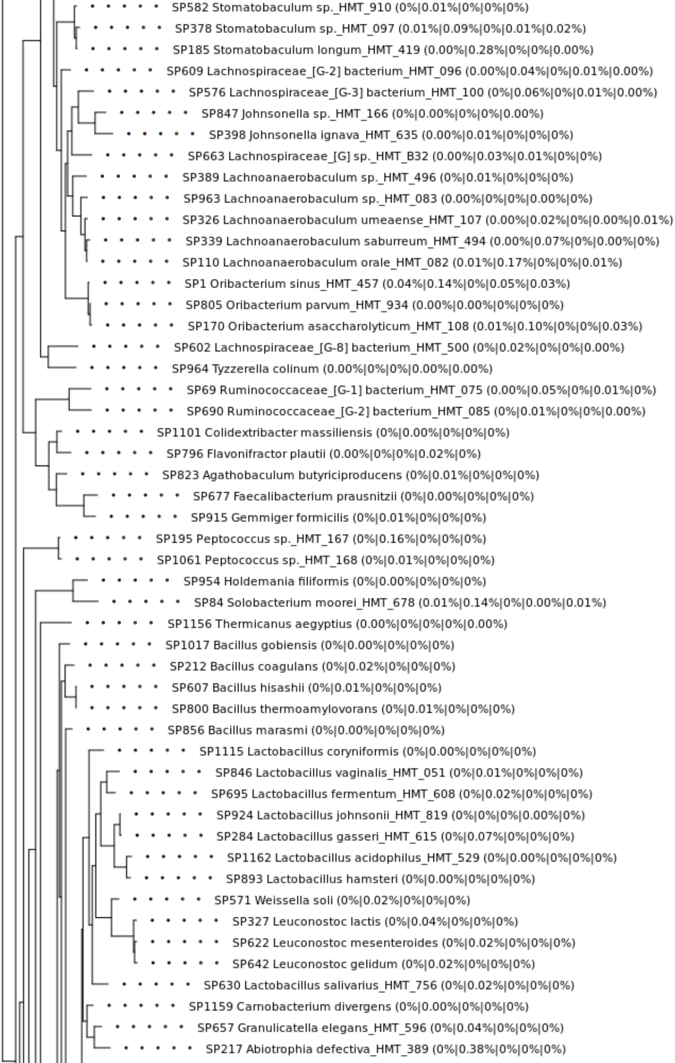

SP191 Rodentibacter pneumotropicus (0.01%|0.21%|0%|0.00%|0.02%)
SP188 Muribacter muris (0.00%|2.30%|0%|0%|0.00%)
SP166 Haemophilus parahaemolyticus_HMT_945 (0%|0.03%|0%|0%|0%)
SP670 Haemophilus pittmaniae_HMT_946 (0%|0.03%|0%|0%|0%)
SP214 Haemophilus haemolyticus_HMT_851 (0%|0.18%|0%|0%|0%)
SP532 Aggregatibacter sp._HMT_898 (0%|0.01%|0%|0%|0%)
SP423 Aggregatibacter aphrophilus_HMT_545 (0%|0.14%|0%|0%|0%)
SP730 Aggregatibacter paraphrophilus_HMT_720 (0%|0.01%|0%|0%|0%)
SP583 Aggregatibacter actinomycetemcomitans_HMT_531 (0%|0.00%|0%|0%|0%)
SP470 Aggregatibacter sp._HMT_949 (0%|0.03%|0%|0%|0%)
SP550 Aggregatibacter sp._HMT_513 (0%|0.05%|0%|0%|0%)
SP348 Aggregatibacter sp._HMT_458 (0%|0.20%|0%|0%|0%)
SP34 Haemophilus parainfluenzae_HMT_718 (0%|10.18%|0%|0%|0%)
SP122 Haemophilus sputorum_HMT_944 (0%|0.06%|0%|0%|0%)
SP895 Aeromonas veronii (0%|0.00%|0%|0%|0%)
SP1064 Acinetobacter marinus (0%|0.00%|0%|0%|0%)
SP521 Acinetobacter junii_HMT_282 (0%|0.01%|0%|0%|0%)
SP336 Acinetobacter tjernbergiae (0%|0.05%|0%|0%|0%)
SP844 Acinetobacter beijerinckii (0%|0.01%|0%|0%|0%)
SP599 Acinetobacter johnsonii_HMT_297 (0%|0.12%|0%|0%|0%)
SP647 Acinetobacter sp._HMT_408 (0%|0.00%|0%|0%|0%)
SP1013 Acinetobacter radioresistens_HMT_010 (0%|0.00%|0%|0%|0%)
SP822 Acinetobacter venetianus (0%|0.01%|0%|0%|0%)
SP1024 Azotobacter vinelandii (0%|0.00%|0%|0%|0%)
SP2 Pseudomonas guguanensis (0%|0.79%|0%|0%|0%)
SP1149 Pseudomonas flavescens (0%|0.00%|0%|0%|0%)
SP340 Pseudomonas putida (0%|0.00%|0%|0%|0%)
SP894 Pseudomonas japonica (0%|0.00%|0%|0%|0%)
SP545 Pseudomonas turukhanskensis (0%|0.00%|0%|0%|0%)
SP544 Pseudomonas koreensis (0%|0.00%|0%|0%|0%)
SP1131 Pseudomonas gessardii (0%|0.00%|0%|0%|0%)
SP680 Pseudomonas veronii (0%|0.03%|0%|0%|0%)
SP804 Pseudomonas brassicacearum (0%|0.01%|0%|0%|0%)
SP426 Pseudomonas versuta (0%|0.03%|0%|0%|0%)
SP686 Pseudomonas deceptionensis (0%|0.00%|0%|0%|0%)
SP471 Pseudomonas fragi (0%|0.01%|0%|0%|0%)
SP375 Pseudomonas aeruginosa_HMT_536 (0.02%|0%|0%|0%|0.02%)
SP481 Pseudomonas anguilliseptica (0%|0.00%|0%|0%|0%)
SP1151 Pseudomonas composti (0%|0.00%|0%|0%|0%)
SP473 Cardiobacterium valvarum_HMT_540 (0%|0.04%|0%|0%|0%)
SP249 Cardiobacterium hominis_HMT_633 (0%|0.06%|0%|0%|0%)
SP826 Belnapia rosea (0.00%|0%|0%|0%|0%)
SP466 Roseococcus suduntuyensis (0%|0.01%|0%|0%|0%)
SP570 Reyranella massiliensis (0.01%|0.01%|0%|0%|0%)
SP989 Skermanella aerolata (0.00%|0%|0%|0%|0%)

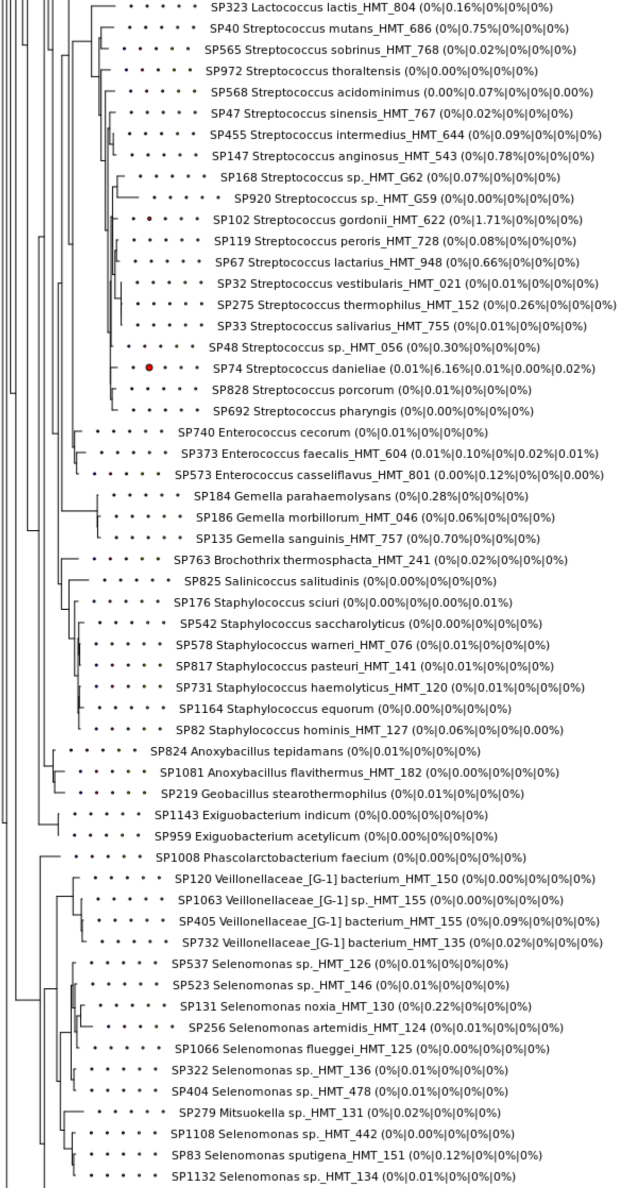

SP1078 Falsirhodobacter halotolerans (0%|0%|0%|0.00%|0%)
SP1221 Gemmobacter lanyuensis (0%|0%|0%|0%|0.00%)
SP975 Paracoccus angustae (0%|0.00%|0%|0%|0%)
SP874 Paracoccus sphaerophysae (0.01%|0%|0%|0%|0%)
SP133 Loktanella salsilacus (0%|0.00%|0%|0%|0%)
SP224 Brevundimonas terrae (0%|0.14%|0%|0%|0%)
SP1027 Brevundimonas mediterranea (0%|0%|0%|0.00%|0%)
SP1148 Brevundimonas aurantiaca (0%|0.00%|0%|0%|0%)
SP980 Caulobacter sp._HMT_002 (0.00%|0%|0%|0.00%|0.00%)
SP1080 Phenylobacterium immobile (0.00%|0%|0%|0%|0%)
SP1119 Caulobacter sp._HMT_C49 (0%|0.00%|0%|0%|0%)
SP502 Agrobacterium tumefaciens_HMT_485 (0%|0.01%|0%|0%|0.00%)
SP797 Hyphomicrobium zavarzinii (0.00%|0%|0%|0%|0%)
SP454 Phreatobacter oligotrophus (0.02%|0.00%|0%|0.00%|0.00%)
SP1062 Pseudorhodoplanes sinuspersici (0.00%|0%|0%|0%|0%)
SP230 Afipia birgiae (0%|0.04%|0%|0%|0%)
SP640 Tardiphaga robiniae (0%|0.02%|0%|0%|0%)
SP1107 Rhodopseudomonas palustris (0%|0.00%|0%|0%|0%)
SP1070 Bosea thiooxidans (0%|0.00%|0%|0%|0%)
SP803 Bosea lathyri (0%|0.01%|0%|0%|0%)
SP610 Enterovirga rhinocerotis (0%|0.02%|0%|0%|0%)
SP587 Methylobacterium iners (0.01%|0%|0%|0%|0%)
SP1147 Microvirga arabica (0.00%|0%|0%|0%|0%)
SP755 Microvirga lupini (0.00%|0.01%|0%|0%|0%)
SP374 Chenggangzhangella methanolivorans (0%|0.01%|0%|0%|0%)
SP679 Methylopila oligotropha (0%|0.02%|0%|0%|0%)
SP235 Ancylobacter abiegnus (0%|0.01%|0%|0%|0%)
SP787 Aquabacter spiritensis (0%|0.00%|0%|0%|0%)
SP756 Pannonibacter phragmitetus (0%|0.00%|0%|0%|0%)
SP611 Polymorphum gilvum (0%|0.01%|0%|0%|0%)
SP177 Sphingomonas changbaiensis (0.06%|0%|0%|0%|0.00%)
SP645 Sphingomonas qilianensis (0%|0.00%|0%|0%|0%)
SP1140 Sphingomonas rubra (0%|0%|0%|0%|0.00%)
SP836 Sphingomonas roseiflava (0%|0.00%|0%|0%|0%)
SP227 Sphingomonas desiccabilis (0.01%|0%|0%|0%|0%)
SP1059 Sphingomonas mucosissima (0%|0.00%|0%|0%|0%)
SP688 Sphingomonas dokdonensis (0.00%|0.02%|0%|0.00%|0%)
SP392 Rhizorhabdus dicambivorans (0%|0.03%|0%|0%|0%)
SP1125 Sphingobium boeckii (0%|0.00%|0%|0%|0%)
SP646 Altererythrobacter dongtanensis (0.01%|0%|0%|0%|0%)
SP556 Novosphingobium barchaimii (0.00%|0.05%|0%|0.00%|0%)
SP771 Sphingopyxis panaciterrae (0%|0.00%|0%|0%|0%)
SP205 Novosphingobium chloroacetimidivorans (0%|0.07%|0%|0%|0%)
SP1225 Sphingosinicella vermicomposti (0%|0%|0%|0%|0.00%)
SP905 Sphingobium yanoikuyae (0.00%|0.01%|0%|0%|0%)
SP223 Sphingobium amiense (0%|0%|0%|0%|0.00%)
SP101 Sphingobium abikonense (0%|0.04%|0%|0%|0%)
SP904 Sphingomonas sp._HMT_B83 (0%|0.00%|0%|0%|0%)
SP649 Sphingobium czechense (0%|0.01%|0%|0%|0%)
SP192 Triticum aestivum_HMT_D31 (0.00%|0.00%|0%|0.25%|0.01%)

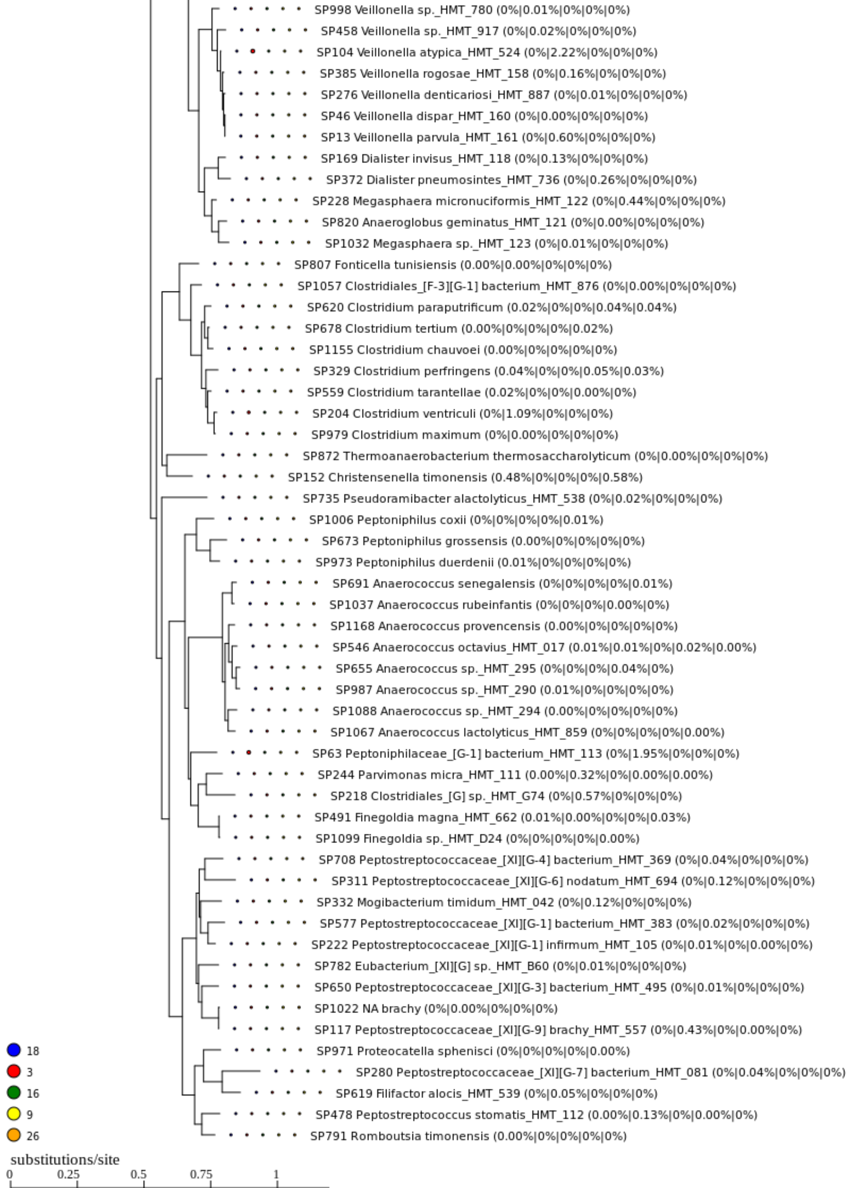

SP564 Treponema sp._HMT_258 (0%|0.02%|0%|0%|0%)
SP569 Treponema sp._HMT_260 (0%|0.05%|0%|0%|0%)
SP390 Treponema lecithinolyticum_HMT_653 (0%|0.23%|0%|0%|0%)
SP203 Treponema maltophilum_HMT_664 (0%|0.40%|0%|0%|0%)
SP57 Treponema sp._HMT_270 (0%|0.22%|0%|0%|0%)
SP45 Treponema socranskii_HMT_769 (0%|0.90%|0%|0%|0%)
SP806 Treponema sp._HMT_268 (0%|0.01%|0%|0%|0%)
SP830 Treponema sp._HMT_263 (0%|0.01%|0%|0%|0%)
SP709 Treponema sp._HMT_257 (0%|0.01%|0%|0%|0%)
SP728 Treponema amylovorum_HMT_541 (0%|0.02%|0%|0%|0%)
SP870 Treponema sp._HMT_256 (0%|0.00%|0%|0%|0%)
SP138 Treponema sp._HMT_951 (0%|0.05%|0%|0%|0%)
SP595 Treponema sp._HMT_250 (0%|0.00%|0%|0%|0%)
SP479 Treponema sp._HMT_247 (0%|0.01%|0%|0%|0%)
SP56 Treponema denticola_HMT_584 (0%|6.54%|0%|0%|0%)
SP764 Treponema sp._HMT_230 (0%|0.02%|0%|0%|0%)
SP221 Treponema sp._HMT_237 (0%|0.07%|0%|0%|0%)
SP116 Treponema sp._HMT_231 (0%|0.33%|0%|0%|0%)
SP190 Treponema vincentii_HMT_029 (0%|0.08%|0%|0%|0%)
SP868 Olsenella profusa_HMT_806 (0%|0.00%|0%|0%|0%)
SP247 Olsenella sp._HMT_807 (0%|0.11%|0%|0.00%|0%)
SP388 Olsenella uli_HMT_038 (0%|0.04%|0%|0%|0%)
SP919 Olsenella umbonata (0%|0.01%|0%|0%|0%)
SP283 Atopobium parvulum_HMT_723 (0.02%|0.69%|0%|0.00%|0.02%)
SP248 Atopobium rimae_HMT_750 (0%|0.06%|0%|0%|0%)
SP155 Atopobium sp._HMT_199 (0%|0.04%|0%|0%|0%)
SP590 Atopobium vaginae_HMT_814 (0.01%|0.01%|0%|0%|0.00%)
SP816 Atopobium sp._HMT_416 (0.00%|0.00%|0%|0%|0%)
SP1015 Cryptobacterium curtum_HMT_579 (0%|0.00%|0%|0%|0%)
SP626 Collinsella aerofaciens (0%|0.01%|0%|0%|0%)
SP914 Gardnerella vaginalis_HMT_829 (0.01%|0%|0%|0%|0%)
SP1160 Bifidobacterium ruminantium (0%|0.00%|0%|0%|0%)
SP209 Bifidobacterium dentium_HMT_588 (0%|0.02%|0%|0%|0%)
SP707 Bifidobacterium longum_HMT_862 (0%|0.02%|0%|0.00%|0%)
SP64 Bifidobacterium pseudolongum (1.70%|0.06%|0%|9.05%|0.22%)
SP614 Alloscardovia omnicolens_HMT_198 (0%|0.02%|0%|0%|0%)
SP137 Scardovia wiggsiae_HMT_195 (0.00%|0.12%|0%|0%|0%)
SP845 Mobiluncus curtisii (0.00%|0%|0%|0%|0%)
SP674 Actinomyces cardiffensis_HMT_850 (0%|0.05%|0%|0%|0%)
SP579 Actinomyces georgiae_HMT_617 (0%|0.01%|0%|0%|0%)
SP469 Actinomyces sp._HMT_178 (0%|0.01%|0%|0.00%|0%)
SP255 Actinomyces lingnae_[Not_Validly_Published]_HMT_181 (0.02%|0.20%|0%|0.01%|0
SP208 Actinomyces sp._HMT_172 (0.00%|0.29%|0%|0.00%|0.00%)
SP172 Actinomyces odontolyticus_HMT_701 (0.00%|0.20%|0%|0.00%|0.00%)
SP154 Actinomyces sp._HMT_180 (0.00%|0.09%|0%|0.00%|0%)


SP923 Actinomyces europaeus (0.01%|0%|0%|0%|0%)
SP108 Actinomyces graevenitzii_HMT_866 (0.00%|0.74%|0%|0%|0.03%)
SP282 Actinomyces bowdenii (0%|0.00%|0%|0%|0.00%)
SP65 Actinomyces johnsonii_HMT_849 (0%|0.05%|0%|0%|0%)
SP66 Actinomyces sp._HMT_171 (0.01%|0.08%|0%|0%|0%)
SP558 Actinomyces naeslundii_HMT_176 (0.00%|0.10%|0%|0.00%|0%)
SP623 Actinomyces sp._HMT_175 (0%|0.04%|0%|0%|0%)
SP365 Actinomyces oris_HMT_893 (0.00%|0.11%|0%|0%|0%)
SP241 Actinomyces viscosus_HMT_688 (0.01%|0.18%|0%|0.00%|0.00%)
SP78 Actinomyces sp._HMT_169 (0.00%|0.47%|0%|0.00%|0.01%)
SP539 Actinomyces sp._HMT_170 (0%|0.02%|0%|0%|0%)
SP1055 Actinomyces oricola_HMT_708 (0%|0.01%|0%|0%|0%)
SP496 Actinomyces massiliensis_HMT_852 (0%|0.04%|0%|0%|0%)
SP617 Actinomyces israelii_HMT_645 (0%|0.02%|0%|0.00%|0%)
SP436 Actinomyces gerencseriae_HMT_618 (0.00%|0.05%|0%|0%|0%)
SP231 Actinomyces sp._HMT_448 (0%|0.06%|0%|0.00%|0%)
SP427 Actinomyces sp._HMT_897 (0%|0.01%|0%|0%|0%)
SP508 Actinomyces sp._HMT_896 (0%|0.01%|0%|0%|0%)
SP482 Actinomyces dentalis_HMT_888 (0%|0.00%|0%|0%|0%)
SP869 Actinomyces haliotis (0%|0.01%|0%|0%|0%)
SP274 Peptidiphaga sp._HMT_183 (0.00%|0.86%|0.01%|0%|0.01%)
SP1098 Brevibacterium paucivorans_HMT_340 (0%|0%|0%|0%|0.00%)
SP1077 Brevibacterium casei (0.00%|0%|0%|0%|0%)
SP1038 Dermabacter hominis_HMT_354 (0%|0%|0%|0.00%|0%)
SP997 Arthrobacter psychrochitiniphilus (0%|0%|0%|0.00%|0%)
SP827 Arthrobacter flavus (0.00%|0%|0%|0%|0%)
SP881 Arthrobacter echini (0%|0.01%|0%|0%|0%)
SP11 Rothia mucilaginosa_HMT_681 (0.21%|10.23%|1.38%|0.16%|0.37%)
SP72 Rothia aeria_HMT_188 (0.00%|1.29%|0%|0.02%|0.00%)
SP71 Rothia dentocariosa_HMT_587 (0.01%|5.09%|0%|0.02%|0.06%)
SP656 Rothia amarae (0%|0%|0%|0%|0.00%)
SP1094 Rothia terrae (0%|0%|0%|0.00%|0%)
SP974 Kocuria palustris_HMT_084 (0%|0%|0%|0%|0.00%)
SP757 Kocuria marina (0%|0%|0%|0%|0.00%)
SP783 Kytococcus sedentarius_HMT_855 (0%|0.00%|0%|0%|0%)
SP848 Janibacter alkaliphilus (0.00%|0%|0%|0%|0%)
SP841 Oryzobacter terrae (0%|0.00%|0%|0.00%|0%)
SP1161 Aurantimicrobium minutum (0.00%|0%|0%|0%|0%)
SP829 Allohumibacter endophyticus (0%|0.00%|0%|0%|0%)
SP1128 Leifsonia soli (0%|0.00%|0%|0%|0%)
SP497 Mycobacterium gordonae (0%|0.01%|0%|0%|0%)
SP229 Lawsonella clevelandensis_HMT_173 (0.00%|0.11%|0%|0.01%|0.00%)
SP759 Turicella otitidis_HMT_832 (0%|0%|0%|0%|0.01%)
SP875 Corynebacterium kroppenstedtii_HMT_049 (0.00%|0.00%|0%|0%|0.01%)
SP863 Corynebacterium epidermidicanis (0%|0.01%|0%|0%|0%)
SP386 Corynebacterium vitaeruminis (0.06%|0.02%|0%|0%|0.00%)
SP722 Corynebacterium argentoratense (0%|0.02%|0%|0%|0%)
SP6 Corynebacterium mastitidis_HMT_328 (48.27%|1.24%|67.22%|70.66%|50.82%)
SP35 Corynebacterium oculi (0.00%|0%|0%|0%|0.00%)
SP143 Corynebacterium durum_HMT_595 (0.00%|0.25%|0%|0.01%|0.00%)
SP5 Corynebacterium matruchotii_HMT_666 (0.01%|0.81%|0%|0.02%|0.01%)


SP721 Corynebacterium ammoniagenes (0%|0.01%|0%|0%|0%)
SP745 Corynebacterium mycetoides (0.00%|0%|0%|0%|0%)
SP798 Corynebacterium lipophiloflavum (0%|0.01%|0%|0%|0%)
SP643 Corynebacterium appendicis_HMT_033 (0.00%|0%|0%|0%|0%)
SP758 Corynebacterium pilbarense_HMT_054 (0.00%|0.01%|0%|0.01%|0.00%)
SP726 Corynebacterium mucifaciens_HMT_835 (0%|0.00%|0%|0.00%|0%)
SP621 Corynebacterium coyleae_HMT_341 (0.02%|0%|0%|0%|0.00%)
SP996 Corynebacterium jeddahense (0%|0.01%|0%|0%|0%)
SP50 Corynebacterium pseudogenitalium (0%|0%|0%|0%|0.00%)
SP49 Corynebacterium tuberculostearicum_HMT_077 (0%|0.00%|0%|0.00%|0.00%)
SP770 Corynebacterium macginleyi_HMT_050 (0%|0%|0%|0.01%|0%)
SP1010 Corynebacterium accolens_HMT_019 (0.00%|0.00%|0%|0%|0.01%)
SP955 Corynebacterium simulans_HMT_062 (0%|0%|0%|0.01%|0.00%)
SP1003 Corynebacterium singulare_HMT_063 (0.00%|0.00%|0%|0.00%|0.00%)
SP476 Corynebacterium aurimucosum_HMT_034 (0.02%|0%|0%|0%|0%)
SP960 Corynebacterium minutissimum_HMT_053 (0.00%|0%|0%|0%|0%)
SP586 Corynebacterium propinquum_HMT_059 (0%|0%|0%|0%|0.00%)
SP100 Corynebacterium jeikeium_HMT_047 (0.01%|0%|0%|0%|0.30%)
SP1009 Pseudopropionibacterium propionicum_HMT_739 (0.00%|0%|0%|0%|0%)
SP572 Cutibacterium sp._HMT_193 (0.00%|0%|0%|0.01%|0.00%)
SP552 Cutibacterium acnes_HMT_530 (0%|0.00%|0%|0.00%|0.01%)
SP337 Synergistetes_[G] sp._HMT_B52 (0.10%|0%|0%|0%|0%)
SP961 Granulicella acidiphila (0.00%|0%|0%|0%|0%)
SP1079 Akkermansia muciniphila (0%|0.00%|0%|0%|0%)
SP1054 Terrimicrobium sacchariphilum (0%|0%|0%|0%|0.00%)
SP605 Anaerolineae_[G-1] bacterium_HMT_439 (0%|0.01%|0%|0%|0%)
SP483 Pyramidobacter piscolens_HMT_357 (0%|0.04%|0%|0%|0%)
SP492 Fretibacterium fastidiosum_HMT_363 (0.00%|0.18%|0%|0%|0%)
SP665 Fretibacterium sp._HMT_362 (0%|0.01%|0%|0%|0%)
SP233 Fretibacterium sp._HMT_359 (0%|0.19%|0%|0%|0.00%)
SP522 Butyrivibrio sp._HMT_455 (0%|0.01%|0%|0%|0%)
SP371 Catonella morbi_HMT_165 (0.00%|0.05%|0%|0%|0%)
SP328 Lacnoclostridium hathewayi (0.17%|0%|0.03%|0.09%|0.11%)
SP1074 Lachnoclostridium pacaense (0.00%|0%|0%|0%|0.00%)
SP1002 Coprococcus catus (0%|0.00%|0%|0%|0%)
SP925 Frisingicoccus caecimuris (0%|0.00%|0%|0%|0%)
SP433 Anaerostipes hadrus (0%|0.01%|0%|0%|0%)
SP747 Eubacterium hallii (0%|0.01%|0%|0%|0%)
SP171 Faecalicatena orotica (0.71%|0.01%|0%|2.83%|0.34%)
SP733 Lachnoclostridium hylemonae (0.00%|0%|0%|0.00%|0.00%)
SP540 Muricomes intestini (0.01%|0%|0%|0.03%|0.00%)
SP562 Dorea formicigenerans (0.01%|0%|0%|0.00%|0.01%)
SP68 Dorea longicatena (0%|0.04%|0%|0%|0%)
SP600 Coprococcus comes (0%|0.00%|0%|0%|0%)
SP320 Roseburia intestinalis (0%|0.00%|0%|0%|0%)
SP484 Fusicatenibacter saccharivorans (0%|0.03%|0%|0%|0%)
SP966 Blautia hydrogenotrophica (0%|0.00%|0%|0%|0%)
SP790 Blautia schinkii (0%|0.00%|0%|0%|0%)
SP606 Blautia luti (0%|0.00%|0%|0%|0%)
SP675 Blautia obeum (0.00%|0.01%|0%|0%|0%)
SP1100 Blautia glucerasea (0%|0.00%|0%|0%|0%)
SP281 Blautia faecis (0%|0.01%|0%|0%|0%)


SP582 Stomatobaculum sp._HMT_910 (0%|0.01%|0%|0%|0%)
SP378 Stomatobaculum sp._HMT_097 (0.01%|0.09%|0%|0.01%|0.02%)
SP185 Stomatobaculum longum_HMT_419 (0.00%|0.28%|0%|0%|0.00%)
SP609 Lachnospiraceae_[G-2] bacterium_HMT_096 (0.00%|0.04%|0%|0.01%|0.00%)
SP576 Lachnospiraceae_[G-3] bacterium_HMT_100 (0%|0.06%|0%|0.01%|0.00%)
SP847 Johnsonella sp._HMT_166 (0%|0.00%|0%|0%|0.00%)
SP398 Johnsonella ignava_HMT_635 (0.00%|0.01%|0%|0%|0%)
SP663 Lachnospiraceae_[G] sp._HMT_B32 (0.00%|0.03%|0.01%|0%|0%)
SP389 Lachnoanaerobaculum sp._HMT_496 (0%|0.01%|0%|0%|0%)
SP963 Lachnoanaerobaculum sp._HMT_083 (0.00%|0%|0%|0.00%|0%)
SP326 Lachnoanaerobaculum umeaense_HMT_107 (0.00%|0.02%|0%|0.00%|0.01%)
SP339 Lachnoanaerobaculum saburreum_HMT_494 (0.00%|0.07%|0%|0.00%|0%)
SP110 Lachnoanaerobaculum orale_HMT_082 (0.01%|0.17%|0%|0%|0.01%)
SP1 Oribacterium sinus_HMT_457 (0.04%|0.14%|0%|0.05%|0.03%)
SP805 Oribacterium parvum_HMT_934 (0.00%|0.00%|0%|0%|0%)
SP170 Oribacterium asaccharolyticum_HMT_108 (0.01%|0.10%|0%|0%|0.03%)
SP602 Lachnospiraceae_[G-8] bacterium_HMT_500 (0%|0.02%|0%|0%|0.00%)
SP964 Tyzzerella colinum (0.00%|0%|0%|0.00%|0.00%)
SP69 Ruminococcaceae_[G-1] bacterium_HMT_075 (0.00%|0.05%|0%|0.01%|0%)
SP690 Ruminococcaceae_[G-2] bacterium_HMT_085 (0%|0.01%|0%|0%|0.00%)
SP1101 Colidextribacter massiliensis (0%|0.00%|0%|0%|0%)
SP796 Flavonifractor plautii (0.00%|0%|0%|0.02%|0%)
SP823 Agathobaculum butyriciproducens (0%|0.01%|0%|0%|0%)
SP677 Faecalibacterium prausnitzii (0%|0.00%|0%|0%|0%)
SP915 Gemmiger formicilis (0%|0.01%|0%|0%|0%)
SP195 Peptococcus sp._HMT_167 (0%|0.16%|0%|0%|0%)
SP1061 Peptococcus sp._HMT_168 (0%|0.01%|0%|0%|0%)
SP954 Holdemania filiformis (0%|0.00%|0%|0%|0%)
SP84 Solobacterium moorei_HMT_678 (0.01%|0.14%|0%|0.00%|0.01%)
SP1156 Thermicanus aegyptius (0.00%|0%|0%|0%|0.00%)
SP1017 Bacillus gobiensis (0%|0.00%|0%|0%|0%)
SP212 Bacillus coagulans (0%|0.02%|0%|0%|0%)
SP607 Bacillus hisashii (0%|0.01%|0%|0%|0%)
SP800 Bacillus thermoamylovorans (0%|0.01%|0%|0%|0%)
SP856 Bacillus marasmi (0%|0.00%|0%|0%|0%)
SP1115 Lactobacillus coryniformis (0%|0.00%|0%|0%|0%)
SP846 Lactobacillus vaginalis_HMT_051 (0%|0.01%|0%|0%|0%)
SP695 Lactobacillus fermentum_HMT_608 (0%|0.02%|0%|0%|0%)
SP924 Lactobacillus johnsonii_HMT_819 (0%|0%|0%|0.00%|0%)
SP284 Lactobacillus gasseri_HMT_615 (0%|0.07%|0%|0%|0%)
SP1162 Lactobacillus acidophilus_HMT_529 (0%|0.00%|0%|0%|0%)
SP893 Lactobacillus hamsteri (0%|0.00%|0%|0%|0%)
SP571 Weissella soli (0%|0.02%|0%|0%|0%)
SP327 Leuconostoc lactis (0%|0.04%|0%|0%|0%)
SP622 Leuconostoc mesenteroides (0%|0.02%|0%|0%|0%)
SP642 Leuconostoc gelidum (0%|0.02%|0%|0%|0%)
SP630 Lactobacillus salivarius_HMT_756 (0%|0.02%|0%|0%|0%)
SP1159 Carnobacterium divergens (0%|0.00%|0%|0%|0%)
SP657 Granulicatella elegans_HMT_596 (0%|0.04%|0%|0%|0%)
SP217 Abiotrophia defectiva_HMT_389 (0%|0.38%|0%|0%|0%)


SP323 Lactococcus lactis_HMT_804 (0%|0.16%|0%|0%|0%)
SP40 Streptococcus mutans_HMT_686 (0%|0.75%|0%|0%|0%)
SP565 Streptococcus sobrinus_HMT_768 (0%|0.02%|0%|0%|0%)
SP972 Streptococcus thoraltensis (0%|0.00%|0%|0%|0%)
SP568 Streptococcus acidominimus (0.00%|0.07%|0%|0%|0.00%)
SP47 Streptococcus sinensis_HMT_767 (0%|0.02%|0%|0%|0%)
SP455 Streptococcus intermedius_HMT_644 (0%|0.09%|0%|0%|0%)
SP147 Streptococcus anginosus_HMT_543 (0%|0.78%|0%|0%|0%)
SP168 Streptococcus sp._HMT_G62 (0%|0.07%|0%|0%|0%)
SP920 Streptococcus sp._HMT_G59 (0%|0.00%|0%|0%|0%)
SP102 Streptococcus gordonii_HMT_622 (0%|1.71%|0%|0%|0%)
SP119 Streptococcus peroris_HMT_728 (0%|0.08%|0%|0%|0%)
SP67 Streptococcus lactarius_HMT_948 (0%|0.66%|0%|0%|0%)
SP32 Streptococcus vestibularis_HMT_021 (0%|0.01%|0%|0%|0%)
SP275 Streptococcus thermophilus_HMT_152 (0%|0.26%|0%|0%|0%)
SP33 Streptococcus salivarius_HMT_755 (0%|0.01%|0%|0%|0%)
SP48 Streptococcus sp._HMT_056 (0%|0.30%|0%|0%|0%)
SP74 Streptococcus danieliae (0.01%|6.16%|0.01%|0.00%|0.02%)
SP828 Streptococcus porcorum (0%|0.01%|0%|0%|0%)
SP692 Streptococcus pharyngis (0%|0.00%|0%|0%|0%)
SP740 Enterococcus cecorum (0%|0.01%|0%|0%|0%)
SP373 Enterococcus faecalis_HMT_604 (0.01%|0.10%|0%|0.02%|0.01%)
SP573 Enterococcus casseliflavus_HMT_801 (0.00%|0.12%|0%|0%|0.00%)
SP184 Gemella parahaemolysans (0%|0.28%|0%|0%|0%)
SP186 Gemella morbillorum_HMT_046 (0%|0.06%|0%|0%|0%)
SP135 Gemella sanguinis_HMT_757 (0%|0.70%|0%|0%|0%)
SP763 Brochothrix thermosphacta_HMT_241 (0%|0.02%|0%|0%|0%)
SP825 Salinicoccus salitudinis (0%|0.00%|0%|0%|0%)
SP176 Staphylococcus sciuri (0%|0.00%|0%|0.00%|0.01%)
SP542 Staphylococcus saccharolyticus (0%|0.00%|0%|0%|0%)
SP578 Staphylococcus warneri_HMT_076 (0%|0.01%|0%|0%|0%)
SP817 Staphylococcus pasteuri_HMT_141 (0%|0.01%|0%|0%|0%)
SP731 Staphylococcus haemolyticus_HMT_120 (0%|0.01%|0%|0%|0%)
SP1164 Staphylococcus equorum (0%|0.00%|0%|0%|0%)
SP82 Staphylococcus hominis_HMT_127 (0%|0.06%|0%|0%|0.00%)
SP824 Anoxybacillus tepidamans (0%|0.01%|0%|0%|0%)
SP1081 Anoxybacillus flavithermus_HMT_182 (0%|0.00%|0%|0%|0%)
SP219 Geobacillus stearothermophilus (0%|0.01%|0%|0%|0%)
SP1143 Exiguobacterium indicum (0%|0.00%|0%|0%|0%)
SP959 Exiguobacterium acetylicum (0%|0.00%|0%|0%|0%)
SP1008 Phascolarctobacterium faecium (0%|0.00%|0%|0%|0%)
SP120 Veillonellaceae_[G-1] bacterium_HMT_150 (0%|0.00%|0%|0%|0%)
SP1063 Veillonellaceae_[G-1] sp._HMT_155 (0%|0.00%|0%|0%|0%)
SP405 Veillonellaceae_[G-1] bacterium_HMT_155 (0%|0.09%|0%|0%|0%)
SP732 Veillonellaceae_[G-1] bacterium_HMT_135 (0%|0.02%|0%|0%|0%)
SP537 Selenomonas sp._HMT_126 (0%|0.01%|0%|0%|0%)
SP523 Selenomonas sp._HMT_146 (0%|0.01%|0%|0%|0%)
SP131 Selenomonas noxia_HMT_130 (0%|0.22%|0%|0%|0%)
SP256 Selenomonas artemidis_HMT_124 (0%|0.01%|0%|0%|0%)
SP1066 Selenomonas flueggei_HMT_125 (0%|0.00%|0%|0%|0%)
SP322 Selenomonas sp._HMT_136 (0%|0.01%|0%|0%|0%)
SP404 Selenomonas sp._HMT_478 (0%|0.01%|0%|0%|0%)
SP279 Mitsuokella sp._HMT_131 (0%|0.02%|0%|0%|0%)
SP1108 Selenomonas sp._HMT_442 (0%|0.00%|0%|0%|0%)
SP83 Selenomonas sputigena_HMT_151 (0%|0.12%|0%|0%|0%)
SP1132 Selenomonas sp._HMT_134 (0%|0.01%|0%|0%|0%)


SP998 Veillonella sp._HMT_780 (0%|0.01%|0%|0%|0%)
SP458 Veillonella sp._HMT_917 (0%|0.02%|0%|0%|0%)
SP104 Veillonella atypica_HMT_524 (0%|2.22%|0%|0%|0%)
SP385 Veillonella rogosae_HMT_158 (0%|0.16%|0%|0%|0%)
SP276 Veillonella denticariosi_HMT_887 (0%|0.01%|0%|0%|0%)
SP46 Veillonella dispar_HMT_160 (0%|0.00%|0%|0%|0%)
SP13 Veillonella parvula_HMT_161 (0%|0.60%|0%|0%|0%)
SP169 Dialister invisus_HMT_118 (0%|0.13%|0%|0%|0%)
SP372 Dialister pneumosintes_HMT_736 (0%|0.26%|0%|0%|0%)
SP228 Megasphaera micronuciformis_HMT_122 (0%|0.44%|0%|0%|0%)
SP820 Anaeroglobus geminatus_HMT_121 (0%|0.00%|0%|0%|0%)
SP1032 Megasphaera sp._HMT_123 (0%|0.01%|0%|0%|0%)
SP807 Fonticella tunisiensis (0.00%|0.00%|0%|0%|0%)
SP1057 Clostridiales_[F-3][G-1] bacterium_HMT_876 (0%|0.00%|0%|0%|0%)
SP620 Clostridium paraputrificum (0.02%|0%|0%|0.04%|0.04%)
SP678 Clostridium tertium (0.00%|0%|0%|0%|0.02%)
SP1155 Clostridium chauvoei (0.00%|0%|0%|0%|0%)
SP329 Clostridium perfringens (0.04%|0%|0%|0.05%|0.03%)
SP559 Clostridium tarantellae (0.02%|0%|0%|0.00%|0%)
SP204 Clostridium ventriculi (0%|1.09%|0%|0%|0%)
SP979 Clostridium maximum (0%|0.00%|0%|0%|0%)
SP872 Thermoanaerobacterium thermosaccharolyticum (0%|0.00%|0%|0%|0%)
SP152 Christensenella timonensis (0.48%|0%|0%|0%|0.58%)
SP735 Pseudoramibacter alactolyticus_HMT_538 (0%|0.02%|0%|0%|0%)
SP1006 Peptoniphilus coxii (0%|0%|0%|0%|0.01%)
SP673 Peptoniphilus grossensis (0.00%|0%|0%|0%|0%)
SP973 Peptoniphilus duerdenii (0.01%|0%|0%|0%|0%)
SP691 Anaerococcus senegalensis (0%|0%|0%|0%|0.01%)
SP1037 Anaerococcus rubeinfantis (0%|0%|0%|0.00%|0%)
SP1168 Anaerococcus provencensis (0.00%|0%|0%|0%|0%)
SP546 Anaerococcus octavius_HMT_017 (0.01%|0.01%|0%|0.02%|0.00%)
SP655 Anaerococcus sp._HMT_295 (0%|0%|0%|0.04%|0%)
SP987 Anaerococcus sp._HMT_290 (0.01%|0%|0%|0%|0%)
SP1088 Anaerococcus sp._HMT_294 (0.00%|0%|0%|0%|0%)
SP1067 Anaerococcus lactolyticus_HMT_859 (0%|0%|0%|0%|0.00%)
SP63 Peptoniphilaceae_[G-1] bacterium_HMT_113 (0%|1.95%|0%|0%|0%)
SP244 Parvimonas micra_HMT_111 (0.00%|0.32%|0%|0.00%|0.00%)
SP218 Clostridiales_[G] sp._HMT_G74 (0%|0.57%|0%|0%|0%)
SP491 Finegoldia magna_HMT_662 (0.01%|0.00%|0%|0%|0.03%)
SP1099 Finegoldia sp._HMT_D24 (0%|0%|0%|0%|0.00%)
SP708 Peptostreptococcaceae_[XI][G-4] bacterium_HMT_369 (0%|0.04%|0%|0%|0%)
SP311 Peptostreptococcaceae_[XI][G-6] nodatum_HMT_694 (0%|0.12%|0%|0%|0%)
SP332 Mogibacterium timidum_HMT_042 (0%|0.12%|0%|0%|0%)
SP577 Peptostreptococcaceae_[XI][G-1] bacterium_HMT_383 (0%|0.02%|0%|0%|0%)
SP222 Peptostreptococcaceae_[XI][G-1] infirmum_HMT_105 (0%|0.01%|0%|0.00%|0%)
SP782 Eubacterium_[XI][G] sp._HMT_B60 (0%|0.01%|0%|0%|0%)
SP650 Peptostreptococcaceae_[XI][G-3] bacterium_HMT_495 (0%|0.01%|0%|0%|0%)
SP1022 NA brachy (0%|0.00%|0%|0%|0%)
SP117 Peptostreptococcaceae_[XI][G-9] brachy_HMT_557 (0%|0.43%|0%|0.00%|0%)
SP971 Proteocatella sphenisci (0%|0%|0%|0%|0.00%)
SP280 Peptostreptococcaceae_[XI][G-7] bacterium_HMT_081 (0%|0.04%|0%|0%|0%)
SP619 Filifactor alocis_HMT_539 (0%|0.05%|0%|0%|0%)
SP478 Peptostreptococcus stomatis_HMT_112 (0.00%|0.13%|0%|0.00%|0%)
SP791 Romboutsia timonensis (0.00%|0%|0%|0%|0%)
18
3
16
9
26
substitutions/site
0
0.25
0.5
0.75
1