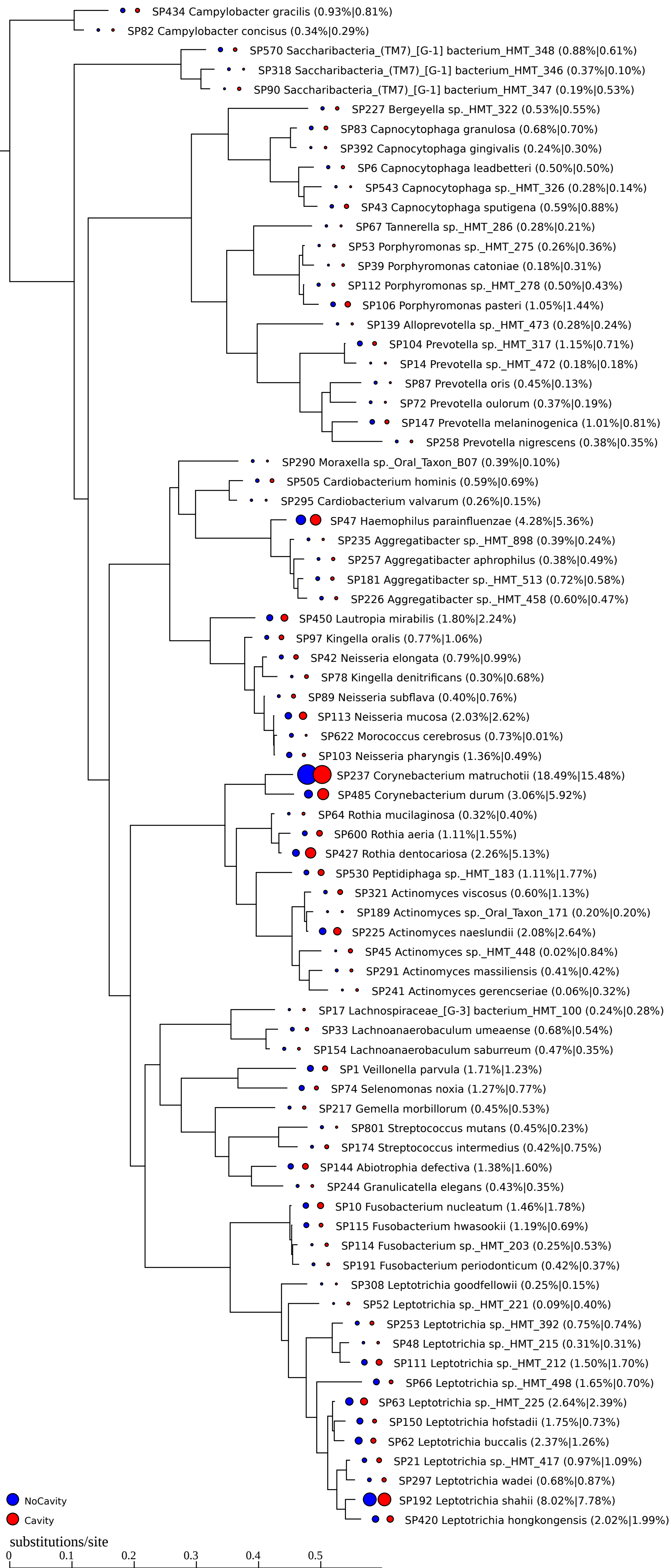
SP434 Campylobacter gracilis (0.93%|0.81%)
SP82 Campylobacter concisus (0.34%|0.29%)
SP570 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0.88%|0.61%)
SP318 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0.37%|0.10%)
SP90 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347 (0.19%|0.53%)
SP227 Bergeyella sp._HMT_322 (0.53%|0.55%)
SP83 Capnocytophaga granulosa (0.68%|0.70%)
SP392 Capnocytophaga gingivalis (0.24%|0.30%)
SP6 Capnocytophaga leadbetteri (0.50%|0.50%)
SP543 Capnocytophaga sp._HMT_326 (0.28%|0.14%)
SP43 Capnocytophaga sputigena (0.59%|0.88%)
SP67 Tannerella sp._HMT_286 (0.28%|0.21%)
SP53 Porphyromonas sp._HMT_275 (0.26%|0.36%)
SP39 Porphyromonas catoniae (0.18%|0.31%)
SP112 Porphyromonas sp._HMT_278 (0.50%|0.43%)
SP106 Porphyromonas pasteri (1.05%|1.44%)
SP139 Alloprevotella sp._HMT_473 (0.28%|0.24%)
SP104 Prevotella sp._HMT_317 (1.15%|0.71%)
SP14 Prevotella sp._HMT_472 (0.18%|0.18%)
SP87 Prevotella oris (0.45%|0.13%)
SP72 Prevotella oulorum (0.37%|0.19%)
SP147 Prevotella melaninogenica (1.01%|0.81%)
SP258 Prevotella nigrescens (0.38%|0.35%)
SP290 Moraxella sp._Oral_Taxon_B07 (0.39%|0.10%)
SP505 Cardiobacterium hominis (0.59%|0.69%)
SP295 Cardiobacterium valvarum (0.26%|0.15%)
SP47 Haemophilus parainfluenzae (4.28%|5.36%)
SP235 Aggregatibacter sp._HMT_898 (0.39%|0.24%)
SP257 Aggregatibacter aphrophilus (0.38%|0.49%)
SP181 Aggregatibacter sp._HMT_513 (0.72%|0.58%)
SP226 Aggregatibacter sp._HMT_458 (0.60%|0.47%)
SP450 Lautropia mirabilis (1.80%|2.24%)
SP97 Kingella oralis (0.77%|1.06%)
SP42 Neisseria elongata (0.79%|0.99%)
SP78 Kingella denitrificans (0.30%|0.68%)
SP89 Neisseria subflava (0.40%|0.76%)
SP113 Neisseria mucosa (2.03%|2.62%)
SP622 Morococcus cerebrosus (0.73%|0.01%)
SP103 Neisseria pharyngis (1.36%|0.49%)
SP237 Corynebacterium matruchotii (18.49%|15.48%)
SP485 Corynebacterium durum (3.06%|5.92%)
SP64 Rothia mucilaginosa (0.32%|0.40%)
SP600 Rothia aeria (1.11%|1.55%)
SP427 Rothia dentocariosa (2.26%|5.13%)
SP530 Peptidiphaga sp._HMT_183 (1.11%|1.77%)
SP321 Actinomyces viscosus (0.60%|1.13%)
SP189 Actinomyces sp._Oral_Taxon_171 (0.20%|0.20%)
SP225 Actinomyces naeslundii (2.08%|2.64%)
SP45 Actinomyces sp._HMT_448 (0.02%|0.84%)
SP291 Actinomyces massiliensis (0.41%|0.42%)
SP241 Actinomyces gerencseriae (0.06%|0.32%)
SP17 Lachnospiraceae_[G-3] bacterium_HMT_100 (0.24%|0.28%)
SP33 Lachnoanaerobaculum umeaense (0.68%|0.54%)
SP154 Lachnoanaerobaculum saburreum (0.47%|0.35%)
SP1 Veillonella parvula (1.71%|1.23%)
SP74 Selenomonas noxia (1.27%|0.77%)
SP217 Gemella morbillorum (0.45%|0.53%)
SP801 Streptococcus mutans (0.45%|0.23%)
SP174 Streptococcus intermedius (0.42%|0.75%)
SP144 Abiotrophia defectiva (1.38%|1.60%)
SP244 Granulicatella elegans (0.43%|0.35%)
SP10 Fusobacterium nucleatum (1.46%|1.78%)
SP115 Fusobacterium hwasookii (1.19%|0.69%)
SP114 Fusobacterium sp._HMT_203 (0.25%|0.53%)
SP191 Fusobacterium periodonticum (0.42%|0.37%)
SP308 Leptotrichia goodfellowii (0.25%|0.15%)
SP52 Leptotrichia sp._HMT_221 (0.09%|0.40%)
SP253 Leptotrichia sp._HMT_392 (0.75%|0.74%)
SP48 Leptotrichia sp._HMT_215 (0.31%|0.31%)
SP111 Leptotrichia sp._HMT_212 (1.50%|1.70%)
SP66 Leptotrichia sp._HMT_498 (1.65%|0.70%)
SP63 Leptotrichia sp._HMT_225 (2.64%|2.39%)
SP150 Leptotrichia hofstadii (1.75%|0.73%)
SP62 Leptotrichia buccalis (2.37%|1.26%)
SP21 Leptotrichia sp._HMT_417 (0.97%|1.09%)
SP297 Leptotrichia wadei (0.68%|0.87%)
SP192 Leptotrichia shahii (8.02%|7.78%)
SP420 Leptotrichia hongkongensis (2.02%|1.99%)
NoCavity
Cavity
substitutions/site
0
0.1
0.2
0.3
0.4
0.5