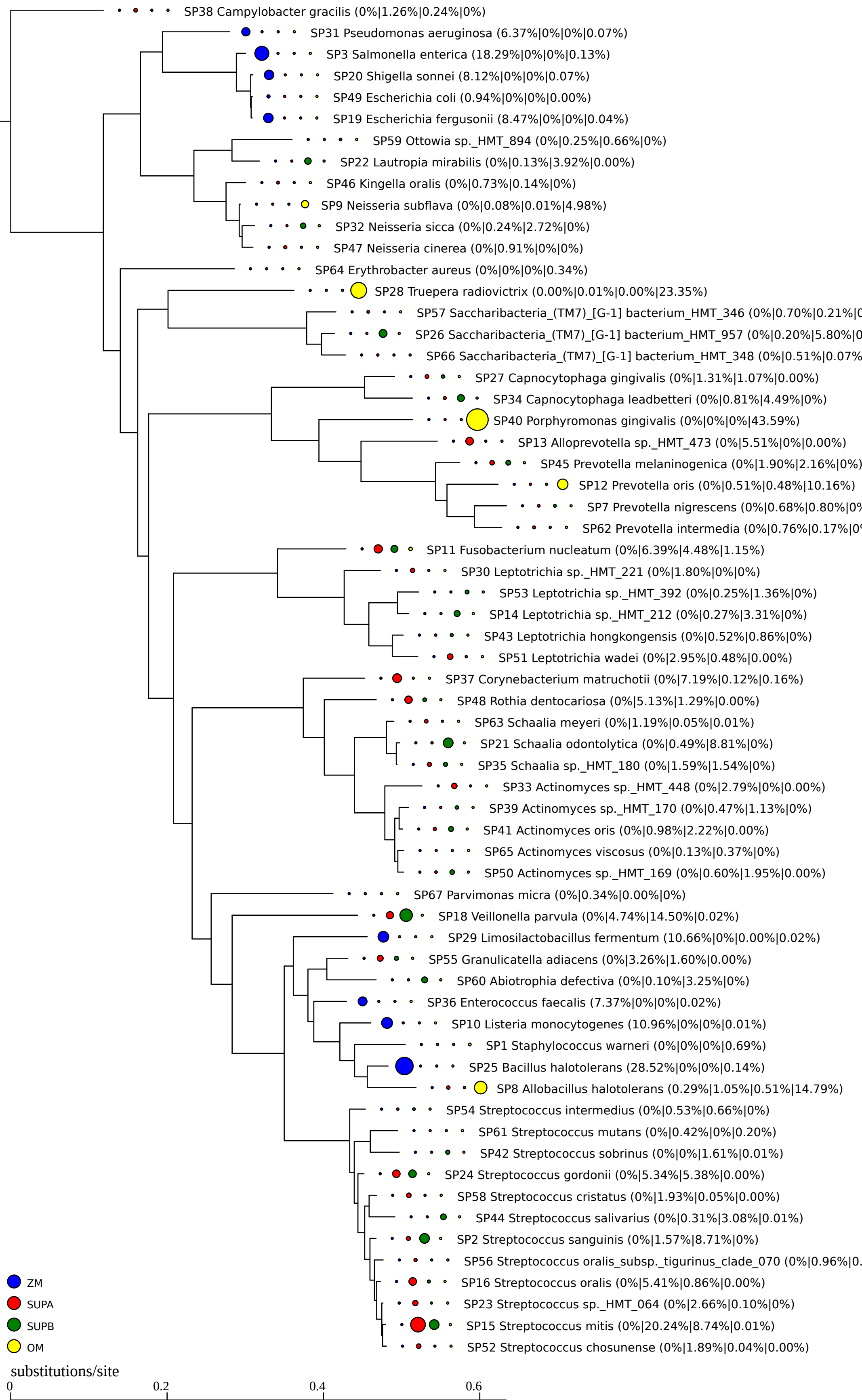
SP38 Campylobacter gracilis (0%|1.26%|0.24%|0%)
SP31 Pseudomonas aeruginosa (6.37%|0%|0%|0.07%)
SP3 Salmonella enterica (18.29%|0%|0%|0.13%)
SP20 Shigella sonnei (8.12%|0%|0%|0.07%)
SP49 Escherichia coli (0.94%|0%|0%|0.00%)
SP19 Escherichia fergusonii (8.47%|0%|0%|0.04%)
SP59 Ottowia sp._HMT_894 (0%|0.25%|0.66%|0%)
SP22 Lautropia mirabilis (0%|0.13%|3.92%|0.00%)
SP46 Kingella oralis (0%|0.73%|0.14%|0%)
SP9 Neisseria subflava (0%|0.08%|0.01%|4.98%)
SP32 Neisseria sicca (0%|0.24%|2.72%|0%)
SP47 Neisseria cinerea (0%|0.91%|0%|0%)
SP64 Erythrobacter aureus (0%|0%|0%|0.34%)
SP28 Truepera radiovictrix (0.00%|0.01%|0.00%|23.35%)
SP57 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0%|0.70%|0.21%|0
SP26 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0%|0.20%|5.80%|0
SP66 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0%|0.51%|0.07%
SP27 Capnocytophaga gingivalis (0%|1.31%|1.07%|0.00%)
SP34 Capnocytophaga leadbetteri (0%|0.81%|4.49%|0%)
SP40 Porphyromonas gingivalis (0%|0%|0%|43.59%)
SP13 Alloprevotella sp._HMT_473 (0%|5.51%|0%|0.00%)
SP45 Prevotella melaninogenica (0%|1.90%|2.16%|0%)
SP12 Prevotella oris (0%|0.51%|0.48%|10.16%)
SP7 Prevotella nigrescens (0%|0.68%|0.80%|0%
SP62 Prevotella intermedia (0%|0.76%|0.17%|0
SP11 Fusobacterium nucleatum (0%|6.39%|4.48%|1.15%)
SP30 Leptotrichia sp._HMT_221 (0%|1.80%|0%|0%)
SP53 Leptotrichia sp._HMT_392 (0%|0.25%|1.36%|0%)
SP14 Leptotrichia sp._HMT_212 (0%|0.27%|3.31%|0%)
SP43 Leptotrichia hongkongensis (0%|0.52%|0.86%|0%)
SP51 Leptotrichia wadei (0%|2.95%|0.48%|0.00%)
SP37 Corynebacterium matruchotii (0%|7.19%|0.12%|0.16%)
SP48 Rothia dentocariosa (0%|5.13%|1.29%|0.00%)
SP63 Schaalia meyeri (0%|1.19%|0.05%|0.01%)
SP21 Schaalia odontolytica (0%|0.49%|8.81%|0%)
SP35 Schaalia sp._HMT_180 (0%|1.59%|1.54%|0%)
SP33 Actinomyces sp._HMT_448 (0%|2.79%|0%|0.00%)
SP39 Actinomyces sp._HMT_170 (0%|0.47%|1.13%|0%)
SP41 Actinomyces oris (0%|0.98%|2.22%|0.00%)
SP65 Actinomyces viscosus (0%|0.13%|0.37%|0%)
SP50 Actinomyces sp._HMT_169 (0%|0.60%|1.95%|0.00%)
SP67 Parvimonas micra (0%|0.34%|0.00%|0%)
SP18 Veillonella parvula (0%|4.74%|14.50%|0.02%)
SP29 Limosilactobacillus fermentum (10.66%|0%|0.00%|0.02%)
SP55 Granulicatella adiacens (0%|3.26%|1.60%|0.00%)
SP60 Abiotrophia defectiva (0%|0.10%|3.25%|0%)
SP36 Enterococcus faecalis (7.37%|0%|0%|0.02%)
SP10 Listeria monocytogenes (10.96%|0%|0%|0.01%)
SP1 Staphylococcus warneri (0%|0%|0%|0.69%)
SP25 Bacillus halotolerans (28.52%|0%|0%|0.14%)
SP8 Allobacillus halotolerans (0.29%|1.05%|0.51%|14.79%)
SP54 Streptococcus intermedius (0%|0.53%|0.66%|0%)
SP61 Streptococcus mutans (0%|0.42%|0%|0.20%)
SP42 Streptococcus sobrinus (0%|0%|1.61%|0.01%)
SP24 Streptococcus gordonii (0%|5.34%|5.38%|0.00%)
SP58 Streptococcus cristatus (0%|1.93%|0.05%|0.00%)
SP44 Streptococcus salivarius (0%|0.31%|3.08%|0.01%)
SP2 Streptococcus sanguinis (0%|1.57%|8.71%|0%)
SP56 Streptococcus oralis_subsp._tigurinus_clade_070 (0%|0.96%|0.
SP16 Streptococcus oralis (0%|5.41%|0.86%|0.00%)
SP23 Streptococcus sp._HMT_064 (0%|2.66%|0.10%|0%)
SP15 Streptococcus mitis (0%|20.24%|8.74%|0.01%)
SP52 Streptococcus chosunense (0%|1.89%|0.04%|0.00%)
ZM
SUPA
SUPB
OM
substitutions/site
0
0.2
0.4
0.6