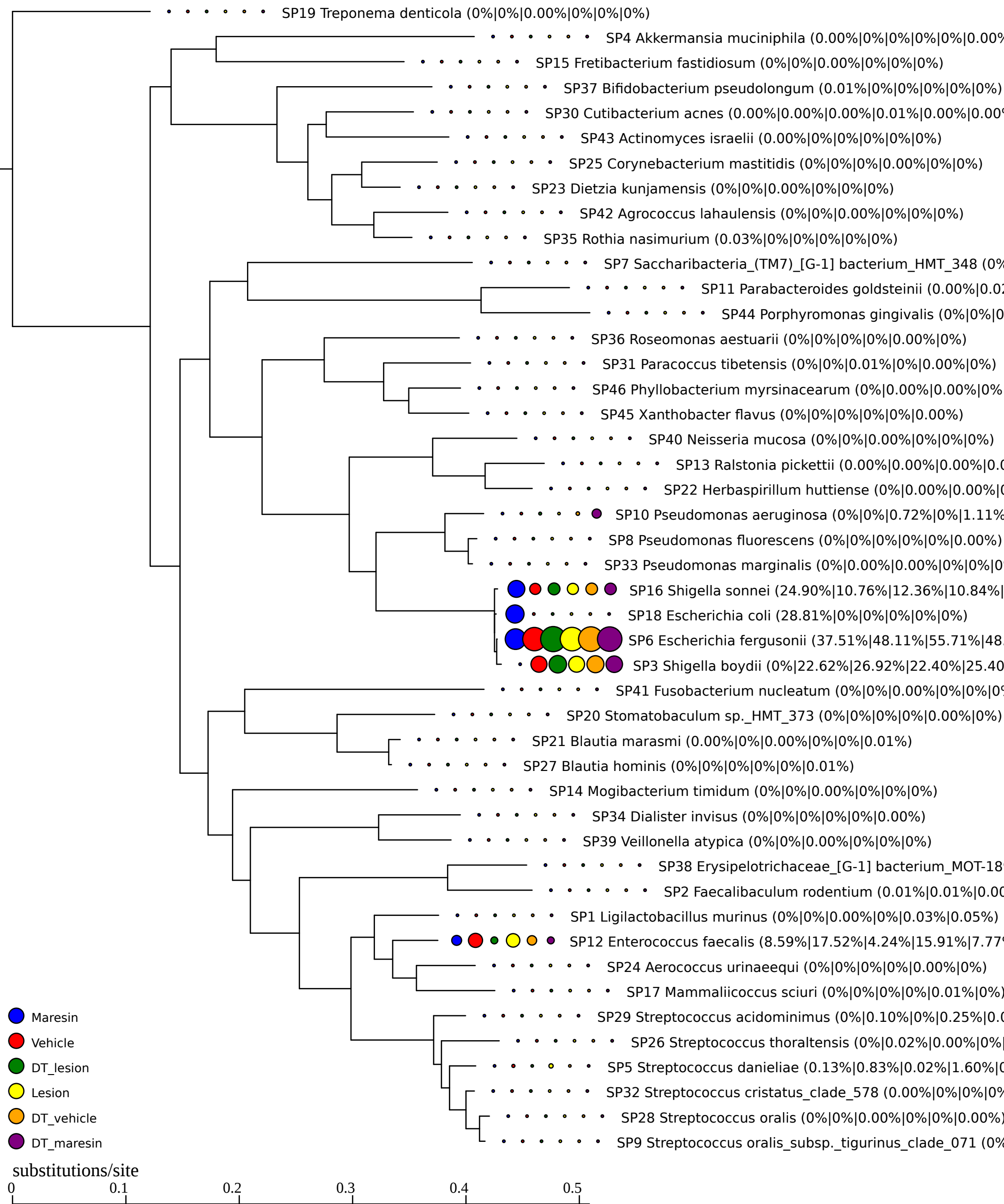
SP19 Treponema denticola (0%|0%|0.00%|0%|0%|0%)
SP4 Akkermansia muciniphila (0.00%|0%|0%|0%|0%|0.00%
SP15 Fretibacterium fastidiosum (0%|0%|0.00%|0%|0%|0%)
SP37 Bifidobacterium pseudolongum (0.01%|0%|0%|0%|0%|0%)
SP30 Cutibacterium acnes (0.00%|0.00%|0.00%|0.01%|0.00%|0.00
SP43 Actinomyces israelii (0.00%|0%|0%|0%|0%|0%)
SP25 Corynebacterium mastitidis (0%|0%|0%|0.00%|0%|0%)
SP23 Dietzia kunjamensis (0%|0%|0.00%|0%|0%|0%)
SP42 Agrococcus lahaulensis (0%|0%|0.00%|0%|0%|0%)
SP35 Rothia nasimurium (0.03%|0%|0%|0%|0%|0%)
SP7 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0%
SP11 Parabacteroides goldsteinii (0.00%|0.02
SP44 Porphyromonas gingivalis (0%|0%|0
SP36 Roseomonas aestuarii (0%|0%|0%|0%|0.00%|0%)
SP31 Paracoccus tibetensis (0%|0%|0.01%|0%|0.00%|0%)
SP46 Phyllobacterium myrsinacearum (0%|0.00%|0.00%|0%
SP45 Xanthobacter flavus (0%|0%|0%|0%|0%|0.00%)
SP40 Neisseria mucosa (0%|0%|0.00%|0%|0%|0%)
SP13 Ralstonia pickettii (0.00%|0.00%|0.00%|0.0
SP22 Herbaspirillum huttiense (0%|0.00%|0.00%|0
SP10 Pseudomonas aeruginosa (0%|0%|0.72%|0%|1.11%
SP8 Pseudomonas fluorescens (0%|0%|0%|0%|0%|0.00%)
SP33 Pseudomonas marginalis (0%|0.00%|0.00%|0%|0%|0
SP16 Shigella sonnei (24.90%|10.76%|12.36%|10.84%|
SP18 Escherichia coli (28.81%|0%|0%|0%|0%|0%)
SP6 Escherichia fergusonii (37.51%|48.11%|55.71%|48.
SP3 Shigella boydii (0%|22.62%|26.92%|22.40%|25.40
SP41 Fusobacterium nucleatum (0%|0%|0.00%|0%|0%|0%
SP20 Stomatobaculum sp._HMT_373 (0%|0%|0%|0%|0.00%|0%)
SP21 Blautia marasmi (0.00%|0%|0.00%|0%|0%|0.01%)
SP27 Blautia hominis (0%|0%|0%|0%|0%|0.01%)
SP14 Mogibacterium timidum (0%|0%|0.00%|0%|0%|0%)
SP34 Dialister invisus (0%|0%|0%|0%|0%|0.00%)
SP39 Veillonella atypica (0%|0%|0.00%|0%|0%|0%)
SP38 Erysipelotrichaceae_[G-1] bacterium_MOT-18
SP2 Faecalibaculum rodentium (0.01%|0.01%|0.00
SP1 Ligilactobacillus murinus (0%|0%|0.00%|0%|0.03%|0.05%)
SP12 Enterococcus faecalis (8.59%|17.52%|4.24%|15.91%|7.77
SP24 Aerococcus urinaeequi (0%|0%|0%|0%|0.00%|0%)
SP17 Mammaliicoccus sciuri (0%|0%|0%|0%|0.01%|0%)
SP29 Streptococcus acidominimus (0%|0.10%|0%|0.25%|0.0
SP26 Streptococcus thoraltensis (0%|0.02%|0.00%|0%|
SP5 Streptococcus danieliae (0.13%|0.83%|0.02%|1.60%|0
SP32 Streptococcus cristatus_clade_578 (0.00%|0%|0%|0%
SP28 Streptococcus oralis (0%|0%|0.00%|0%|0%|0.00%)
SP9 Streptococcus oralis_subsp._tigurinus_clade_071 (0%
Maresin
Vehicle
DT_lesion
Lesion
DT_vehicle
DT_maresin
substitutions/site
0
0.1
0.2
0.3
0.4
0.5