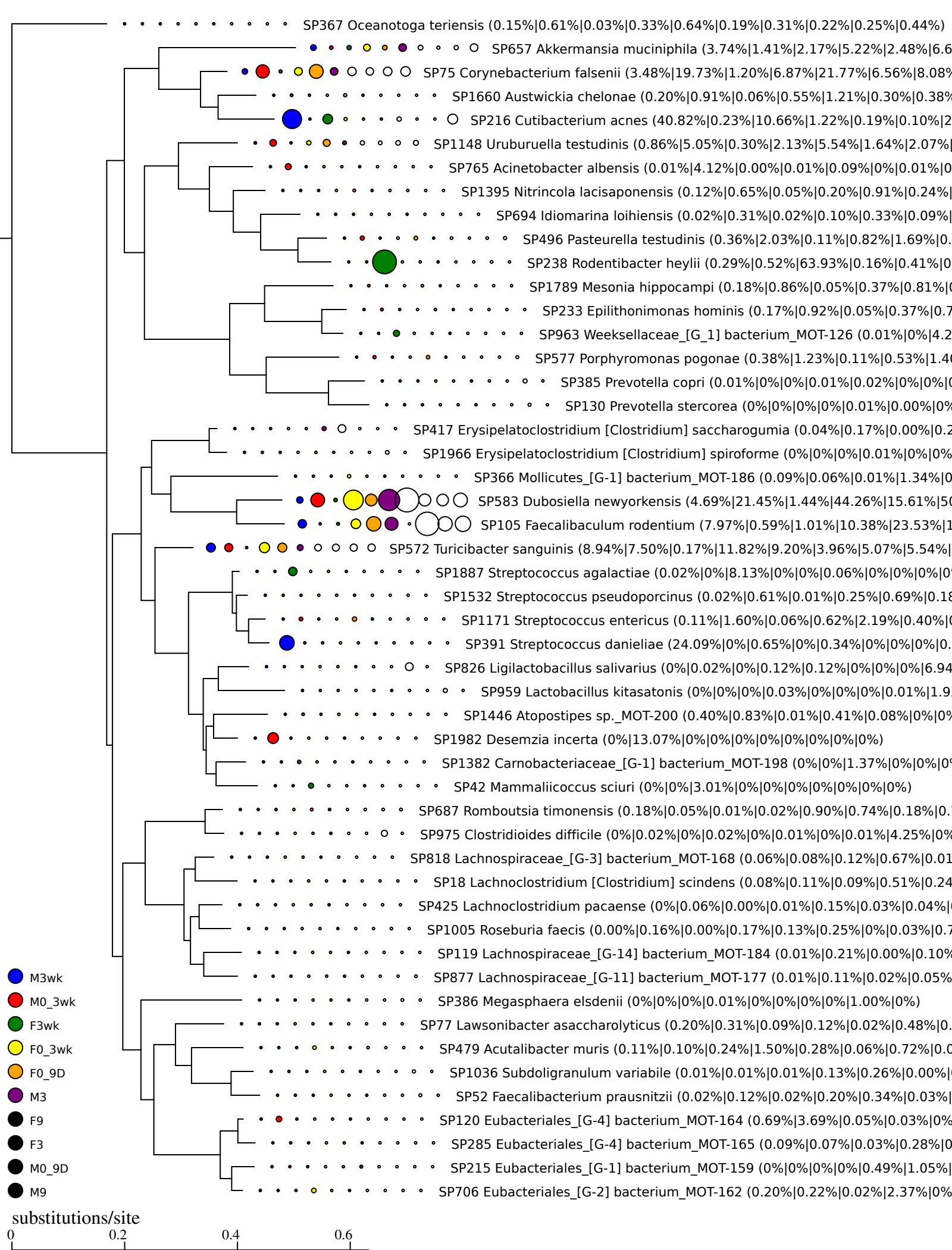
SP367 Oceanotoga teriensis (0.15%|0.61%|0.03%|0.33%|0.64%|0.19%|0.31%|0.22%|0.25%|0.44%)
SP657 Akkermansia muciniphila (3.74%|1.41%|2.17%|5.22%|2.48%|6.6
SP75 Corynebacterium falsenii (3.48%|19.73%|1.20%|6.87%|21.77%|6.56%|8.08%
SP1660 Austwickia chelonae (0.20%|0.91%|0.06%|0.55%|1.21%|0.30%|0.38%
SP216 Cutibacterium acnes (40.82%|0.23%|10.66%|1.22%|0.19%|0.10%|2
SP1148 Uruburuella testudinis (0.86%|5.05%|0.30%|2.13%|5.54%|1.64%|2.07%|
SP765 Acinetobacter albensis (0.01%|4.12%|0.00%|0.01%|0.09%|0%|0.01%|0
SP1395 Nitrincola lacisaponensis (0.12%|0.65%|0.05%|0.20%|0.91%|0.24%|
SP694 Idiomarina loihiensis (0.02%|0.31%|0.02%|0.10%|0.33%|0.09%|
SP496 Pasteurella testudinis (0.36%|2.03%|0.11%|0.82%|1.69%|0.
SP238 Rodentibacter heylii (0.29%|0.52%|63.93%|0.16%|0.41%|0
SP1789 Mesonia hippocampi (0.18%|0.86%|0.05%|0.37%|0.81%|0
SP233 Epilithonimonas hominis (0.17%|0.92%|0.05%|0.37%|0.7
SP963 Weeksellaceae_[G_1] bacterium_MOT-126 (0.01%|0%|4.2
SP577 Porphyromonas pogonae (0.38%|1.23%|0.11%|0.53%|1.4
SP385 Prevotella copri (0.01%|0%|0%|0.01%|0.02%|0%|0%|0
SP130 Prevotella stercorea (0%|0%|0%|0%|0.01%|0.00%|0%
SP417 Erysipelatoclostridium [Clostridium] saccharogumia (0.04%|0.17%|0.00%|0.2
SP1966 Erysipelatoclostridium [Clostridium] spiroforme (0%|0%|0%|0.01%|0%|0%
SP366 Mollicutes_[G-1] bacterium_MOT-186 (0.09%|0.06%|0.01%|1.34%|0
SP583 Dubosiella newyorkensis (4.69%|21.45%|1.44%|44.26%|15.61%|50
SP105 Faecalibaculum rodentium (7.97%|0.59%|1.01%|10.38%|23.53%|1
SP572 Turicibacter sanguinis (8.94%|7.50%|0.17%|11.82%|9.20%|3.96%|5.07%|5.54%|
SP1887 Streptococcus agalactiae (0.02%|0%|8.13%|0%|0%|0.06%|0%|0%|0%|0
SP1532 Streptococcus pseudoporcinus (0.02%|0.61%|0.01%|0.25%|0.69%|0.18
SP1171 Streptococcus entericus (0.11%|1.60%|0.06%|0.62%|2.19%|0.40%|0
SP391 Streptococcus danieliae (24.09%|0%|0.65%|0%|0.34%|0%|0%|0%|0.
SP826 Ligilactobacillus salivarius (0%|0.02%|0%|0.12%|0.12%|0%|0%|0%|6.94
SP959 Lactobacillus kitasatonis (0%|0%|0%|0.03%|0%|0%|0%|0.01%|1.9
SP1446 Atopostipes sp._MOT-200 (0.40%|0.83%|0.01%|0.41%|0.08%|0%|0%
SP1982 Desemzia incerta (0%|13.07%|0%|0%|0%|0%|0%|0%|0%|0%)
SP1382 Carnobacteriaceae_[G-1] bacterium_MOT-198 (0%|0%|1.37%|0%|0%|0
SP42 Mammaliicoccus sciuri (0%|0%|3.01%|0%|0%|0%|0%|0%|0%|0%)
SP687 Romboutsia timonensis (0.18%|0.05%|0.01%|0.02%|0.90%|0.74%|0.18%|0.
SP975 Clostridioides difficile (0%|0.02%|0%|0.02%|0%|0.01%|0%|0.01%|4.25%|0%
SP818 Lachnospiraceae_[G-3] bacterium_MOT-168 (0.06%|0.08%|0.12%|0.67%|0.01
SP18 Lachnoclostridium [Clostridium] scindens (0.08%|0.11%|0.09%|0.51%|0.24
SP425 Lachnoclostridium pacaense (0%|0.06%|0.00%|0.01%|0.15%|0.03%|0.04%|
SP1005 Roseburia faecis (0.00%|0.16%|0.00%|0.17%|0.13%|0.25%|0%|0.03%|0.7
SP119 Lachnospiraceae_[G-14] bacterium_MOT-184 (0.01%|0.21%|0.00%|0.10%
SP877 Lachnospiraceae_[G-11] bacterium_MOT-177 (0.01%|0.11%|0.02%|0.05%
SP386 Megasphaera elsdenii (0%|0%|0%|0.01%|0%|0%|0%|0%|1.00%|0%)
SP77 Lawsonibacter asaccharolyticus (0.20%|0.31%|0.09%|0.12%|0.02%|0.48%|0.
SP479 Acutalibacter muris (0.11%|0.10%|0.24%|1.50%|0.28%|0.06%|0.72%|0.0
SP1036 Subdoligranulum variabile (0.01%|0.01%|0.01%|0.13%|0.26%|0.00%|
SP52 Faecalibacterium prausnitzii (0.02%|0.12%|0.02%|0.20%|0.34%|0.03%|
SP120 Eubacteriales_[G-4] bacterium_MOT-164 (0.69%|3.69%|0.05%|0.03%|0%
SP285 Eubacteriales_[G-4] bacterium_MOT-165 (0.09%|0.07%|0.03%|0.28%|0
SP215 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|0%|0%|0.49%|1.05%|
SP706 Eubacteriales_[G-2] bacterium_MOT-162 (0.20%|0.22%|0.02%|2.37%|0%
M3wk
M0_3wk
F3wk
F0_3wk
F0_9D
M3
F9
F3
M0_9D
M9
substitutions/site
0
0.2
0.4
0.6