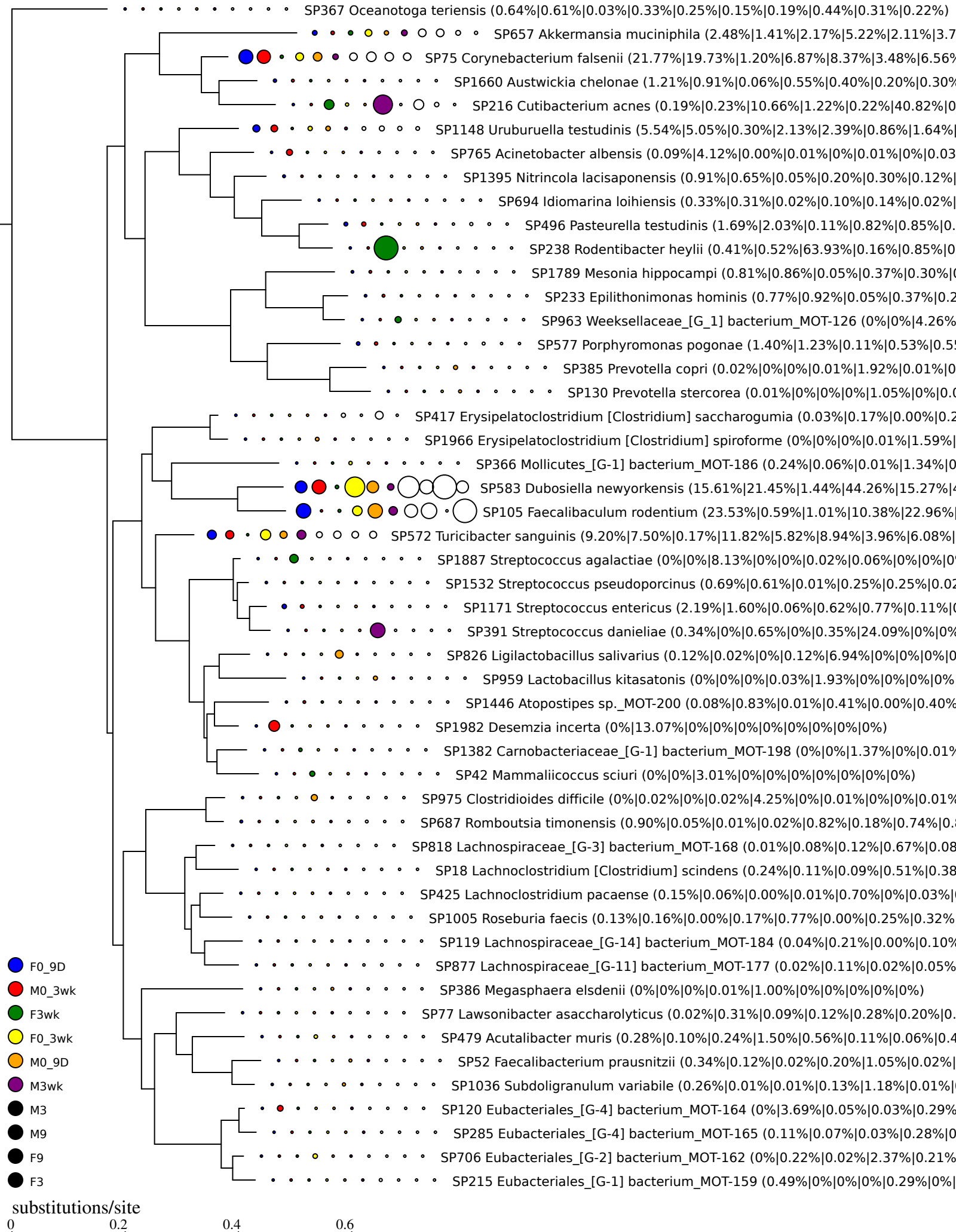
SP367 Oceanotoga teriensis (0.64%|0.61%|0.03%|0.33%|0.25%|0.15%|0.19%|0.44%|0.31%|0.22%)
SP657 Akkermansia muciniphila (2.48%|1.41%|2.17%|5.22%|2.11%|3.7
SP75 Corynebacterium falsenii (21.77%|19.73%|1.20%|6.87%|8.37%|3.48%|6.56%
SP1660 Austwickia chelonae (1.21%|0.91%|0.06%|0.55%|0.40%|0.20%|0.30%
SP216 Cutibacterium acnes (0.19%|0.23%|10.66%|1.22%|0.22%|40.82%|0
SP1148 Uruburuella testudinis (5.54%|5.05%|0.30%|2.13%|2.39%|0.86%|1.64%|
SP765 Acinetobacter albensis (0.09%|4.12%|0.00%|0.01%|0%|0.01%|0%|0.03
SP1395 Nitrincola lacisaponensis (0.91%|0.65%|0.05%|0.20%|0.30%|0.12%|
SP694 Idiomarina loihiensis (0.33%|0.31%|0.02%|0.10%|0.14%|0.02%|
SP496 Pasteurella testudinis (1.69%|2.03%|0.11%|0.82%|0.85%|0.
SP238 Rodentibacter heylii (0.41%|0.52%|63.93%|0.16%|0.85%|0
SP1789 Mesonia hippocampi (0.81%|0.86%|0.05%|0.37%|0.30%|
SP233 Epilithonimonas hominis (0.77%|0.92%|0.05%|0.37%|0.2
SP963 Weeksellaceae_[G_1] bacterium_MOT-126 (0%|0%|4.26%
SP577 Porphyromonas pogonae (1.40%|1.23%|0.11%|0.53%|0.5
SP385 Prevotella copri (0.02%|0%|0%|0.01%|1.92%|0.01%|0
SP130 Prevotella stercorea (0.01%|0%|0%|0%|1.05%|0%|0.
SP417 Erysipelatoclostridium [Clostridium] saccharogumia (0.03%|0.17%|0.00%|0.2
SP1966 Erysipelatoclostridium [Clostridium] spiroforme (0%|0%|0%|0.01%|1.59%|
SP366 Mollicutes_[G-1] bacterium_MOT-186 (0.24%|0.06%|0.01%|1.34%|0
SP583 Dubosiella newyorkensis (15.61%|21.45%|1.44%|44.26%|15.27%|4
SP105 Faecalibaculum rodentium (23.53%|0.59%|1.01%|10.38%|22.96%|
SP572 Turicibacter sanguinis (9.20%|7.50%|0.17%|11.82%|5.82%|8.94%|3.96%|6.08%|
SP1887 Streptococcus agalactiae (0%|0%|8.13%|0%|0%|0.02%|0.06%|0%|0%|0
SP1532 Streptococcus pseudoporcinus (0.69%|0.61%|0.01%|0.25%|0.25%|0.02
SP1171 Streptococcus entericus (2.19%|1.60%|0.06%|0.62%|0.77%|0.11%|
SP391 Streptococcus danieliae (0.34%|0%|0.65%|0%|0.35%|24.09%|0%|0%
SP826 Ligilactobacillus salivarius (0.12%|0.02%|0%|0.12%|6.94%|0%|0%|0%|0
SP959 Lactobacillus kitasatonis (0%|0%|0%|0.03%|1.93%|0%|0%|0%|0%
SP1446 Atopostipes sp._MOT-200 (0.08%|0.83%|0.01%|0.41%|0.00%|0.40%
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%|0.01%
SP42 Mammaliicoccus sciuri (0%|0%|3.01%|0%|0%|0%|0%|0%|0%|0%)
SP975 Clostridioides difficile (0%|0.02%|0%|0.02%|4.25%|0%|0.01%|0%|0%|0.01%
SP687 Romboutsia timonensis (0.90%|0.05%|0.01%|0.02%|0.82%|0.18%|0.74%|0.
SP818 Lachnospiraceae_[G-3] bacterium_MOT-168 (0.01%|0.08%|0.12%|0.67%|0.08
SP18 Lachnoclostridium [Clostridium] scindens (0.24%|0.11%|0.09%|0.51%|0.38
SP425 Lachnoclostridium pacaense (0.15%|0.06%|0.00%|0.01%|0.70%|0%|0.03%|
SP1005 Roseburia faecis (0.13%|0.16%|0.00%|0.17%|0.77%|0.00%|0.25%|0.32%
SP119 Lachnospiraceae_[G-14] bacterium_MOT-184 (0.04%|0.21%|0.00%|0.10%
SP877 Lachnospiraceae_[G-11] bacterium_MOT-177 (0.02%|0.11%|0.02%|0.05%
SP386 Megasphaera elsdenii (0%|0%|0%|0.01%|1.00%|0%|0%|0%|0%|0%)
SP77 Lawsonibacter asaccharolyticus (0.02%|0.31%|0.09%|0.12%|0.28%|0.20%|0.
SP479 Acutalibacter muris (0.28%|0.10%|0.24%|1.50%|0.56%|0.11%|0.06%|0.4
SP52 Faecalibacterium prausnitzii (0.34%|0.12%|0.02%|0.20%|1.05%|0.02%|
SP1036 Subdoligranulum variabile (0.26%|0.01%|0.01%|0.13%|1.18%|0.01%|
SP120 Eubacteriales_[G-4] bacterium_MOT-164 (0%|3.69%|0.05%|0.03%|0.29%
SP285 Eubacteriales_[G-4] bacterium_MOT-165 (0.11%|0.07%|0.03%|0.28%|0
SP706 Eubacteriales_[G-2] bacterium_MOT-162 (0%|0.22%|0.02%|2.37%|0.21%
SP215 Eubacteriales_[G-1] bacterium_MOT-159 (0.49%|0%|0%|0%|0.29%|0%|
F0_9D
M0_3wk
F3wk
F0_3wk
M0_9D
M3wk
M3
M9
F9
F3
substitutions/site
0
0.2
0.4
0.6