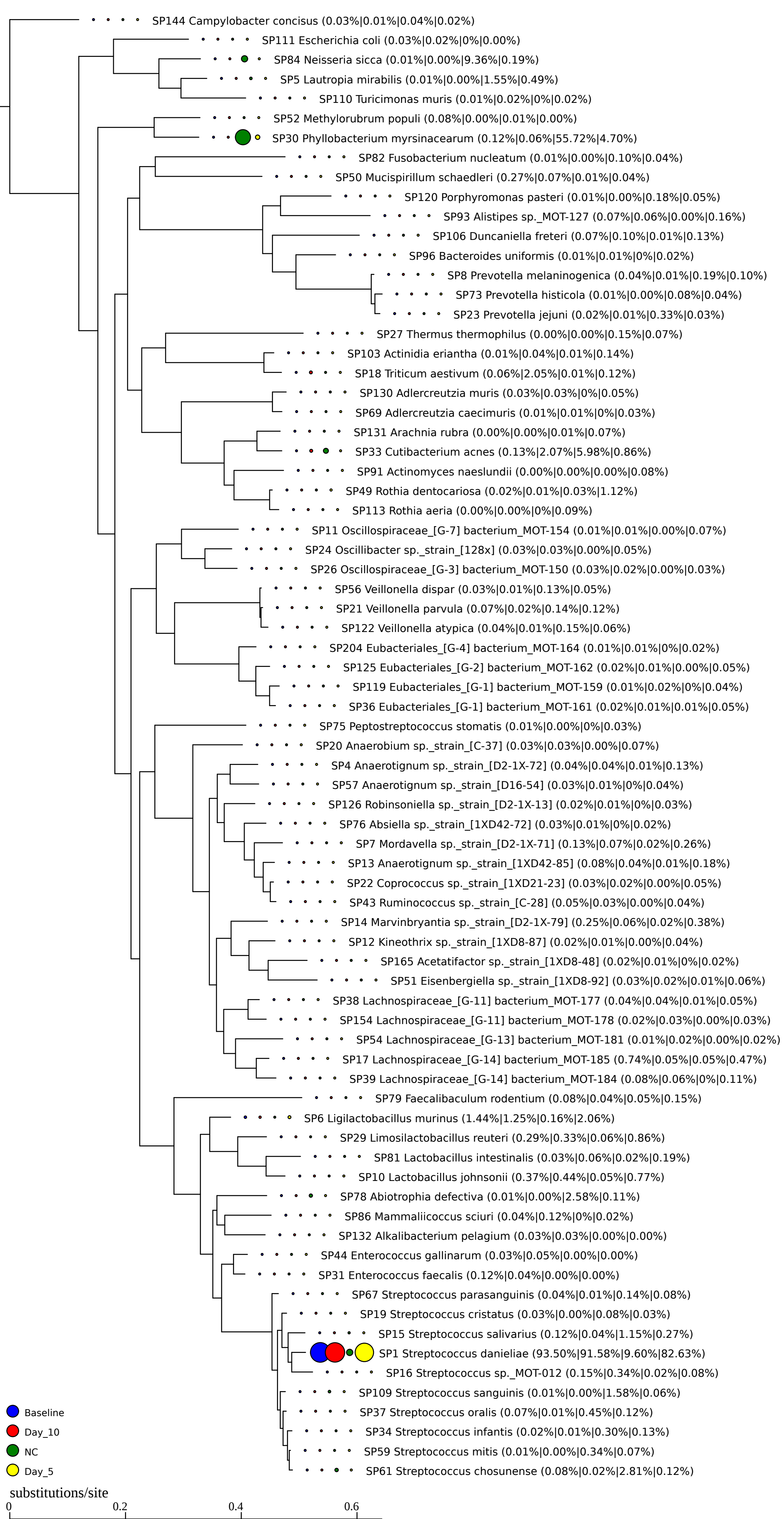
SP144 Campylobacter concisus (0.03%|0.01%|0.04%|0.02%)
SP111 Escherichia coli (0.03%|0.02%|0%|0.00%)
SP84 Neisseria sicca (0.01%|0.00%|9.36%|0.19%)
SP5 Lautropia mirabilis (0.01%|0.00%|1.55%|0.49%)
SP110 Turicimonas muris (0.01%|0.02%|0%|0.02%)
SP52 Methylorubrum populi (0.08%|0.00%|0.01%|0.00%)
SP30 Phyllobacterium myrsinacearum (0.12%|0.06%|55.72%|4.70%)
SP82 Fusobacterium nucleatum (0.01%|0.00%|0.10%|0.04%)
SP50 Mucispirillum schaedleri (0.27%|0.07%|0.01%|0.04%)
SP120 Porphyromonas pasteri (0.01%|0.00%|0.18%|0.05%)
SP93 Alistipes sp._MOT-127 (0.07%|0.06%|0.00%|0.16%)
SP106 Duncaniella freteri (0.07%|0.10%|0.01%|0.13%)
SP96 Bacteroides uniformis (0.01%|0.01%|0%|0.02%)
SP8 Prevotella melaninogenica (0.04%|0.01%|0.19%|0.10%)
SP73 Prevotella histicola (0.01%|0.00%|0.08%|0.04%)
SP23 Prevotella jejuni (0.02%|0.01%|0.33%|0.03%)
SP27 Thermus thermophilus (0.00%|0.00%|0.15%|0.07%)
SP103 Actinidia eriantha (0.01%|0.04%|0.01%|0.14%)
SP18 Triticum aestivum (0.06%|2.05%|0.01%|0.12%)
SP130 Adlercreutzia muris (0.03%|0.03%|0%|0.05%)
SP69 Adlercreutzia caecimuris (0.01%|0.01%|0%|0.03%)
SP131 Arachnia rubra (0.00%|0.00%|0.01%|0.07%)
SP33 Cutibacterium acnes (0.13%|2.07%|5.98%|0.86%)
SP91 Actinomyces naeslundii (0.00%|0.00%|0.00%|0.08%)
SP49 Rothia dentocariosa (0.02%|0.01%|0.03%|1.12%)
SP113 Rothia aeria (0.00%|0.00%|0%|0.09%)
SP11 Oscillospiraceae_[G-7] bacterium_MOT-154 (0.01%|0.01%|0.00%|0.07%)
SP24 Oscillibacter sp._strain_[128x] (0.03%|0.03%|0.00%|0.05%)
SP26 Oscillospiraceae_[G-3] bacterium_MOT-150 (0.03%|0.02%|0.00%|0.03%)
SP56 Veillonella dispar (0.03%|0.01%|0.13%|0.05%)
SP21 Veillonella parvula (0.07%|0.02%|0.14%|0.12%)
SP122 Veillonella atypica (0.04%|0.01%|0.15%|0.06%)
SP204 Eubacteriales_[G-4] bacterium_MOT-164 (0.01%|0.01%|0%|0.02%)
SP125 Eubacteriales_[G-2] bacterium_MOT-162 (0.02%|0.01%|0.00%|0.05%)
SP119 Eubacteriales_[G-1] bacterium_MOT-159 (0.01%|0.02%|0%|0.04%)
SP36 Eubacteriales_[G-1] bacterium_MOT-161 (0.02%|0.01%|0.01%|0.05%)
SP75 Peptostreptococcus stomatis (0.01%|0.00%|0%|0.03%)
SP20 Anaerobium sp._strain_[C-37] (0.03%|0.03%|0.00%|0.07%)
SP4 Anaerotignum sp._strain_[D2-1X-72] (0.04%|0.04%|0.01%|0.13%)
SP57 Anaerotignum sp._strain_[D16-54] (0.03%|0.01%|0%|0.04%)
SP126 Robinsoniella sp._strain_[D2-1X-13] (0.02%|0.01%|0%|0.03%)
SP76 Absiella sp._strain_[1XD42-72] (0.03%|0.01%|0%|0.02%)
SP7 Mordavella sp._strain_[D2-1X-71] (0.13%|0.07%|0.02%|0.26%)
SP13 Anaerotignum sp._strain_[1XD42-85] (0.08%|0.04%|0.01%|0.18%)
SP22 Coprococcus sp._strain_[1XD21-23] (0.03%|0.02%|0.00%|0.05%)
SP43 Ruminococcus sp._strain_[C-28] (0.05%|0.03%|0.00%|0.04%)
SP14 Marvinbryantia sp._strain_[D2-1X-79] (0.25%|0.06%|0.02%|0.38%)
SP12 Kineothrix sp._strain_[1XD8-87] (0.02%|0.01%|0.00%|0.04%)
SP165 Acetatifactor sp._strain_[1XD8-48] (0.02%|0.01%|0%|0.02%)
SP51 Eisenbergiella sp._strain_[1XD8-92] (0.03%|0.02%|0.01%|0.06%)
SP38 Lachnospiraceae_[G-11] bacterium_MOT-177 (0.04%|0.04%|0.01%|0.05%)
SP154 Lachnospiraceae_[G-11] bacterium_MOT-178 (0.02%|0.03%|0.00%|0.03%)
SP54 Lachnospiraceae_[G-13] bacterium_MOT-181 (0.01%|0.02%|0.00%|0.02%)
SP17 Lachnospiraceae_[G-14] bacterium_MOT-185 (0.74%|0.05%|0.05%|0.47%)
SP39 Lachnospiraceae_[G-14] bacterium_MOT-184 (0.08%|0.06%|0%|0.11%)
SP79 Faecalibaculum rodentium (0.08%|0.04%|0.05%|0.15%)
SP6 Ligilactobacillus murinus (1.44%|1.25%|0.16%|2.06%)
SP29 Limosilactobacillus reuteri (0.29%|0.33%|0.06%|0.86%)
SP81 Lactobacillus intestinalis (0.03%|0.06%|0.02%|0.19%)
SP10 Lactobacillus johnsonii (0.37%|0.44%|0.05%|0.77%)
SP78 Abiotrophia defectiva (0.01%|0.00%|2.58%|0.11%)
SP86 Mammaliicoccus sciuri (0.04%|0.12%|0%|0.02%)
SP132 Alkalibacterium pelagium (0.03%|0.03%|0.00%|0.00%)
SP44 Enterococcus gallinarum (0.03%|0.05%|0.00%|0.00%)
SP31 Enterococcus faecalis (0.12%|0.04%|0.00%|0.00%)
SP67 Streptococcus parasanguinis (0.04%|0.01%|0.14%|0.08%)
SP19 Streptococcus cristatus (0.03%|0.00%|0.08%|0.03%)
SP15 Streptococcus salivarius (0.12%|0.04%|1.15%|0.27%)
SP1 Streptococcus danieliae (93.50%|91.58%|9.60%|82.63%)
SP16 Streptococcus sp._MOT-012 (0.15%|0.34%|0.02%|0.08%)
SP109 Streptococcus sanguinis (0.01%|0.00%|1.58%|0.06%)
SP37 Streptococcus oralis (0.07%|0.01%|0.45%|0.12%)
SP34 Streptococcus infantis (0.02%|0.01%|0.30%|0.13%)
SP59 Streptococcus mitis (0.01%|0.00%|0.34%|0.07%)
SP61 Streptococcus chosunense (0.08%|0.02%|2.81%|0.12%)
Baseline
Day_10
NC
Day_5
substitutions/site
0
0.2
0.4
0.6