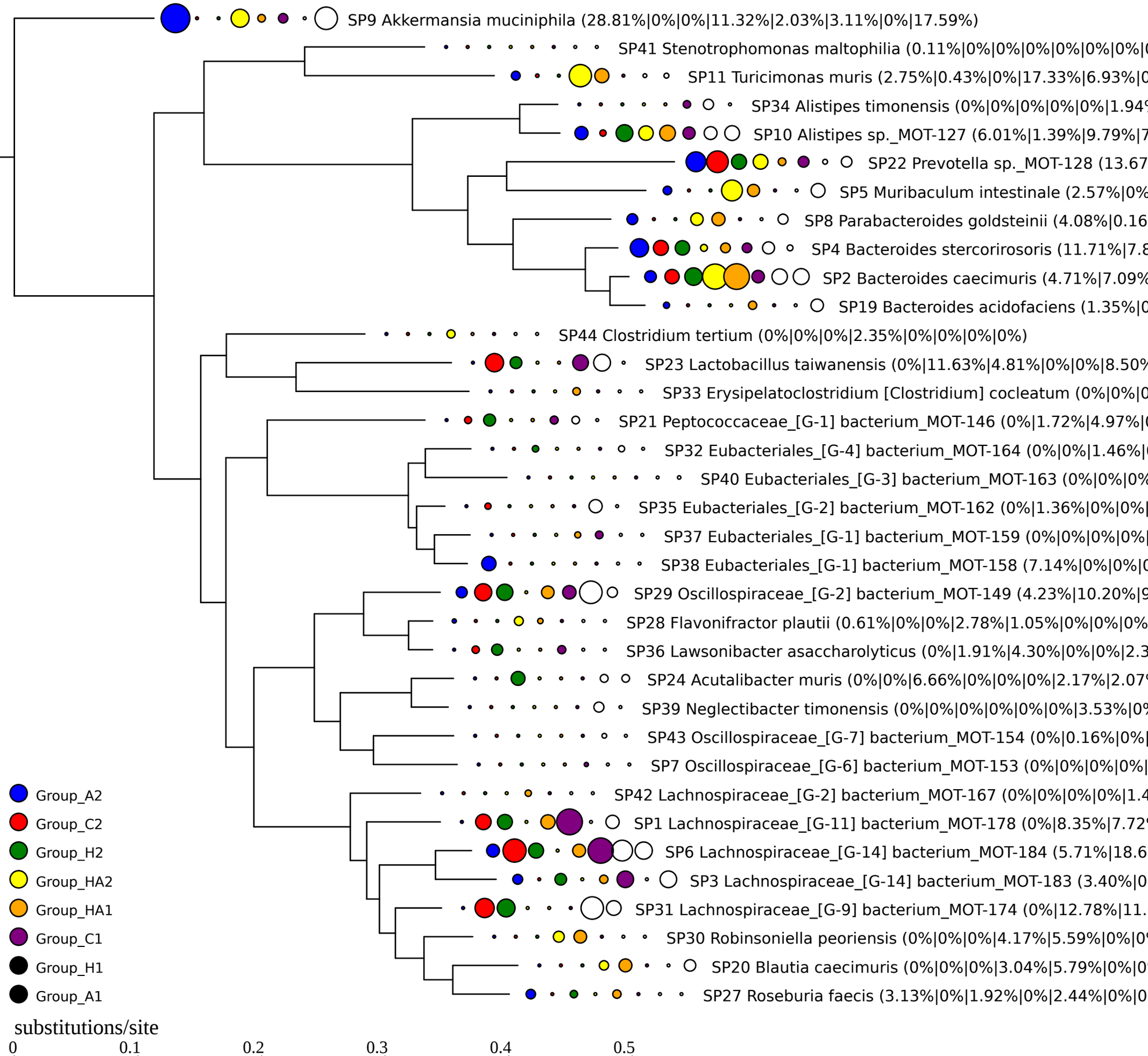
SP9 Akkermansia muciniphila (28.81%|0%|0%|11.32%|2.03%|3.11%|0%|17.59%)
SP41 Stenotrophomonas maltophilia (0.11%|0%|0%|0%|0%|0%|0%|
SP11 Turicimonas muris (2.75%|0.43%|0%|17.33%|6.93%|
SP34 Alistipes timonensis (0%|0%|0%|0%|0%|1.94
SP10 Alistipes sp._MOT-127 (6.01%|1.39%|9.79%|
SP22 Prevotella sp._MOT-128 (13.67
SP5 Muribaculum intestinale (2.57%|0%
SP8 Parabacteroides goldsteinii (4.08%|0.16
SP4 Bacteroides stercorirosoris (11.71%|7.8
SP2 Bacteroides caecimuris (4.71%|7.09%
SP19 Bacteroides acidofaciens (1.35%|
SP44 Clostridium tertium (0%|0%|0%|2.35%|0%|0%|0%|0%)
SP23 Lactobacillus taiwanensis (0%|11.63%|4.81%|0%|0%|8.50%
SP33 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|
SP21 Peptococcaceae_[G-1] bacterium_MOT-146 (0%|1.72%|4.97%|
SP32 Eubacteriales_[G-4] bacterium_MOT-164 (0%|0%|1.46%|
SP40 Eubacteriales_[G-3] bacterium_MOT-163 (0%|0%|0%
SP35 Eubacteriales_[G-2] bacterium_MOT-162 (0%|1.36%|0%|0%|
SP37 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|0%|0%|
SP38 Eubacteriales_[G-1] bacterium_MOT-158 (7.14%|0%|0%|
SP29 Oscillospiraceae_[G-2] bacterium_MOT-149 (4.23%|10.20%|9
SP28 Flavonifractor plautii (0.61%|0%|0%|2.78%|1.05%|0%|0%|0%
SP36 Lawsonibacter asaccharolyticus (0%|1.91%|4.30%|0%|0%|2.3
SP24 Acutalibacter muris (0%|0%|6.66%|0%|0%|0%|2.17%|2.07
SP39 Neglectibacter timonensis (0%|0%|0%|0%|0%|0%|3.53%|0
SP43 Oscillospiraceae_[G-7] bacterium_MOT-154 (0%|0.16%|0%|
SP7 Oscillospiraceae_[G-6] bacterium_MOT-153 (0%|0%|0%|0%|
SP42 Lachnospiraceae_[G-2] bacterium_MOT-167 (0%|0%|0%|0%|1.4
SP1 Lachnospiraceae_[G-11] bacterium_MOT-178 (0%|8.35%|7.72
SP6 Lachnospiraceae_[G-14] bacterium_MOT-184 (5.71%|18.6
SP3 Lachnospiraceae_[G-14] bacterium_MOT-183 (3.40%|0
SP31 Lachnospiraceae_[G-9] bacterium_MOT-174 (0%|12.78%|11.
SP30 Robinsoniella peoriensis (0%|0%|0%|4.17%|5.59%|0%|0
SP20 Blautia caecimuris (0%|0%|0%|3.04%|5.79%|0%|0
SP27 Roseburia faecis (3.13%|0%|1.92%|0%|2.44%|0%|0
Group_A2
Group_C2
Group_H2
Group_HA2
Group_HA1
Group_C1
Group_H1
Group_A1
substitutions/site
0
0.1
0.2
0.3
0.4
0.5