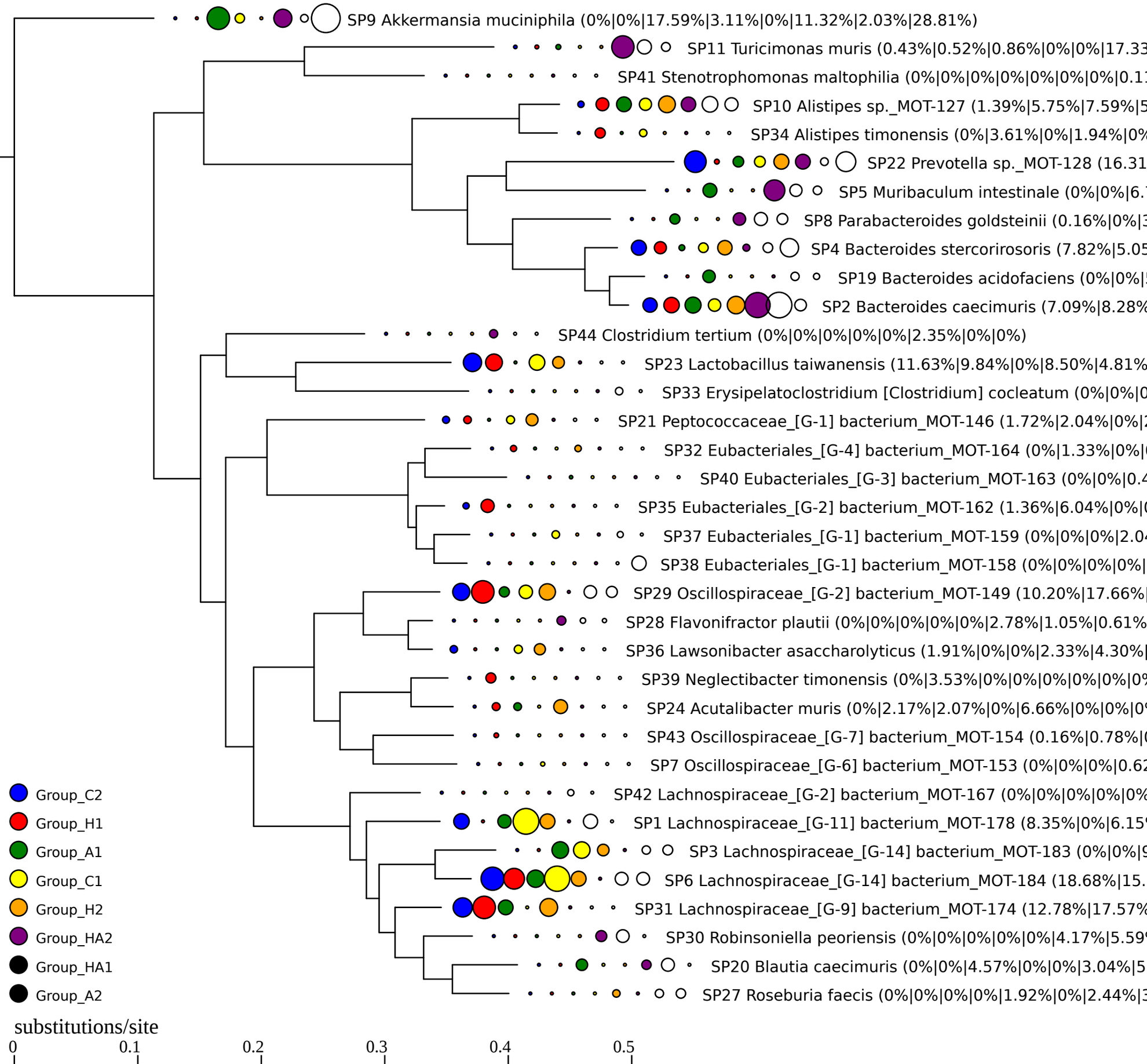
SP9 Akkermansia muciniphila (0%|0%|17.59%|3.11%|0%|11.32%|2.03%|28.81%)
SP11 Turicimonas muris (0.43%|0.52%|0.86%|0%|0%|17.33
SP41 Stenotrophomonas maltophilia (0%|0%|0%|0%|0%|0%|0%|0.1
SP10 Alistipes sp._MOT-127 (1.39%|5.75%|7.59%|5
SP34 Alistipes timonensis (0%|3.61%|0%|1.94%|0%
SP22 Prevotella sp._MOT-128 (16.31
SP5 Muribaculum intestinale (0%|0%|6.
SP8 Parabacteroides goldsteinii (0.16%|0%|3
SP4 Bacteroides stercorirosoris (7.82%|5.05
SP19 Bacteroides acidofaciens (0%|0%|
SP2 Bacteroides caecimuris (7.09%|8.28%
SP44 Clostridium tertium (0%|0%|0%|0%|0%|2.35%|0%|0%)
SP23 Lactobacillus taiwanensis (11.63%|9.84%|0%|8.50%|4.81%
SP33 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|0
SP21 Peptococcaceae_[G-1] bacterium_MOT-146 (1.72%|2.04%|0%|
SP32 Eubacteriales_[G-4] bacterium_MOT-164 (0%|1.33%|0%|
SP40 Eubacteriales_[G-3] bacterium_MOT-163 (0%|0%|0.4
SP35 Eubacteriales_[G-2] bacterium_MOT-162 (1.36%|6.04%|0%|
SP37 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|0%|2.04
SP38 Eubacteriales_[G-1] bacterium_MOT-158 (0%|0%|0%|0%|
SP29 Oscillospiraceae_[G-2] bacterium_MOT-149 (10.20%|17.66%
SP28 Flavonifractor plautii (0%|0%|0%|0%|0%|2.78%|1.05%|0.61%
SP36 Lawsonibacter asaccharolyticus (1.91%|0%|0%|2.33%|4.30%
SP39 Neglectibacter timonensis (0%|3.53%|0%|0%|0%|0%|0%|0%
SP24 Acutalibacter muris (0%|2.17%|2.07%|0%|6.66%|0%|0%|0
SP43 Oscillospiraceae_[G-7] bacterium_MOT-154 (0.16%|0.78%|
SP7 Oscillospiraceae_[G-6] bacterium_MOT-153 (0%|0%|0%|0.62
SP42 Lachnospiraceae_[G-2] bacterium_MOT-167 (0%|0%|0%|0%|0%
SP1 Lachnospiraceae_[G-11] bacterium_MOT-178 (8.35%|0%|6.15
SP3 Lachnospiraceae_[G-14] bacterium_MOT-183 (0%|0%|9
SP6 Lachnospiraceae_[G-14] bacterium_MOT-184 (18.68%|15.
SP31 Lachnospiraceae_[G-9] bacterium_MOT-174 (12.78%|17.57%
SP30 Robinsoniella peoriensis (0%|0%|0%|0%|0%|4.17%|5.59
SP20 Blautia caecimuris (0%|0%|4.57%|0%|0%|3.04%|5
SP27 Roseburia faecis (0%|0%|0%|0%|1.92%|0%|2.44%|3
Group_C2
Group_H1
Group_A1
Group_C1
Group_H2
Group_HA2
Group_HA1
Group_A2
substitutions/site
0
0.1
0.2
0.3
0.4
0.5