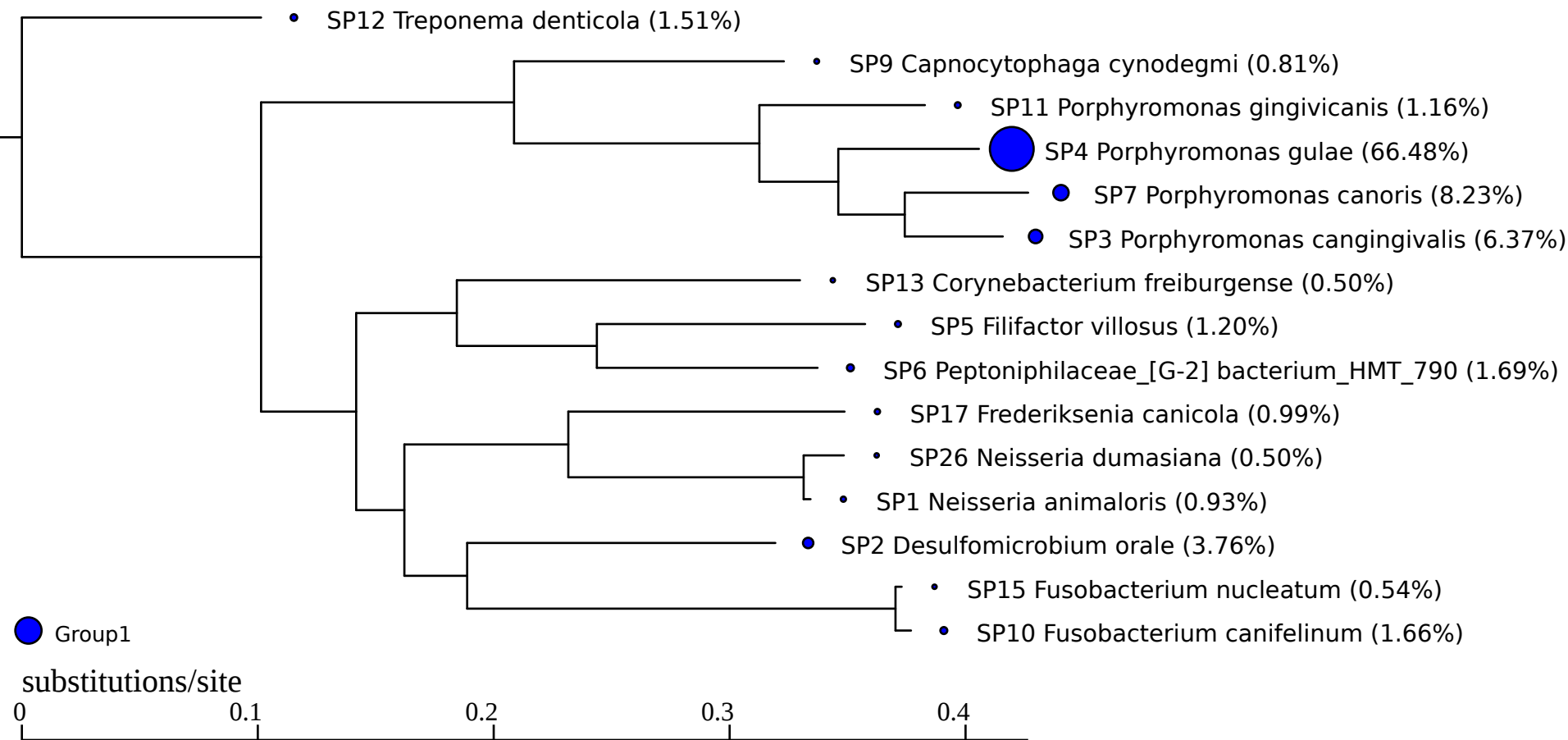
SP12 Treponema denticola (1.51%)
SP9 Capnocytophaga cynodegmi (0.81%)
SP11 Porphyromonas gingivicanis (1.16%)
SP4 Porphyromonas gulae (66.48%)
SP7 Porphyromonas canoris (8.23%)
SP3 Porphyromonas cangingivalis (6.37%)
SP13 Corynebacterium freiburgense (0.50%)
SP5 Filifactor villosus (1.20%)
SP6 Peptoniphilaceae_[G-2] bacterium_HMT_790 (1.69%)
SP17 Frederiksenia canicola (0.99%)
SP26 Neisseria dumasiana (0.50%)
SP1 Neisseria animaloris (0.93%)
SP2 Desulfomicrobium orale (3.76%)
SP15 Fusobacterium nucleatum (0.54%)
SP10 Fusobacterium canifelinum (1.66%)
Group1
substitutions/site
0
0.1
0.2
0.3
0.4