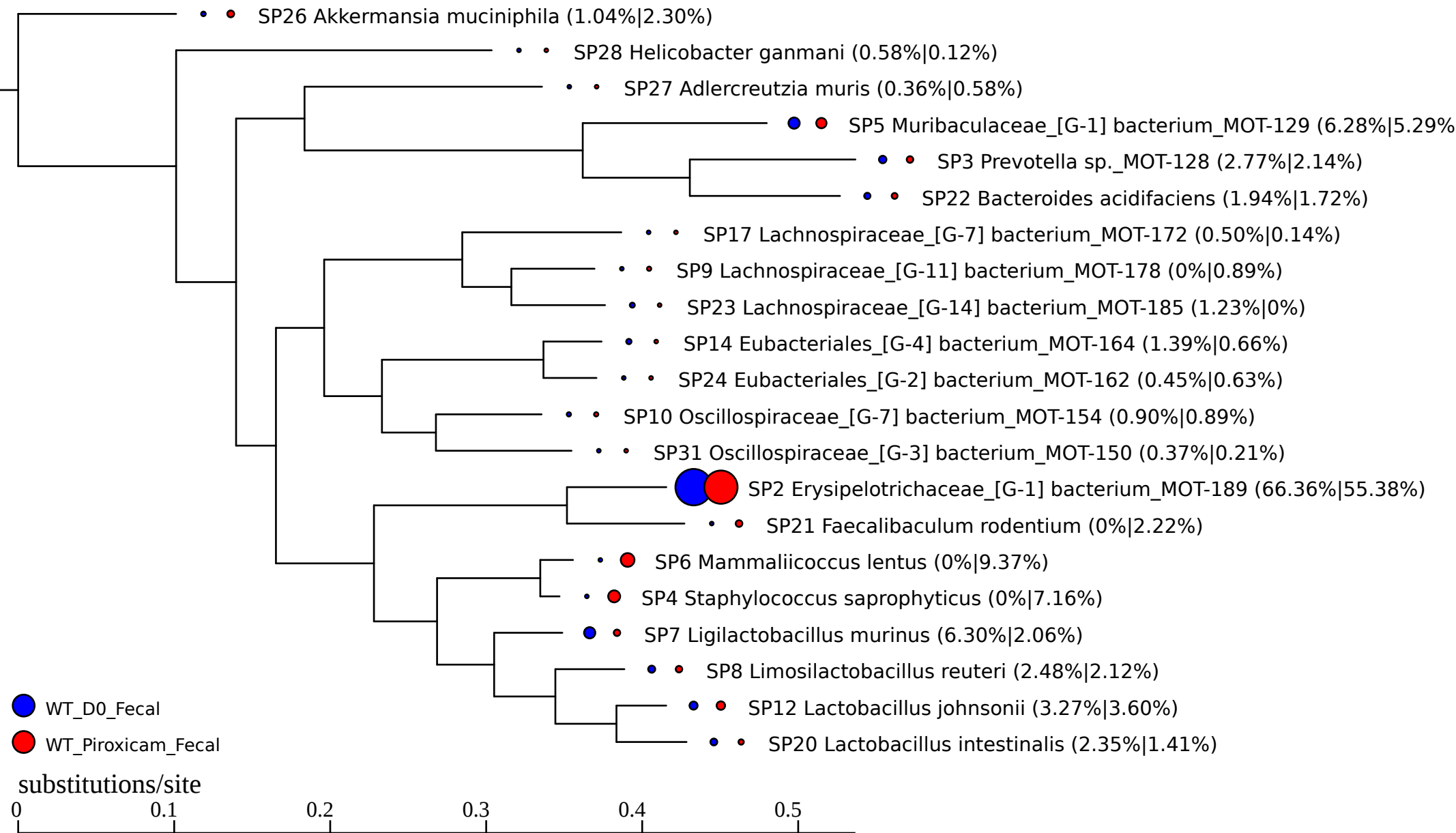
SP26 Akkermansia muciniphila (1.04%|2.30%)
SP28 Helicobacter ganmani (0.58%|0.12%)
SP27 Adlercreutzia muris (0.36%|0.58%)
SP5 Muribaculaceae_[G-1] bacterium_MOT-129 (6.28%|5.29%
SP3 Prevotella sp._MOT-128 (2.77%|2.14%)
SP22 Bacteroides acidifaciens (1.94%|1.72%)
SP17 Lachnospiraceae_[G-7] bacterium_MOT-172 (0.50%|0.14%)
SP9 Lachnospiraceae_[G-11] bacterium_MOT-178 (0%|0.89%)
SP23 Lachnospiraceae_[G-14] bacterium_MOT-185 (1.23%|0%)
SP14 Eubacteriales_[G-4] bacterium_MOT-164 (1.39%|0.66%)
SP24 Eubacteriales_[G-2] bacterium_MOT-162 (0.45%|0.63%)
SP10 Oscillospiraceae_[G-7] bacterium_MOT-154 (0.90%|0.89%)
SP31 Oscillospiraceae_[G-3] bacterium_MOT-150 (0.37%|0.21%)
SP2 Erysipelotrichaceae_[G-1] bacterium_MOT-189 (66.36%|55.38%)
SP21 Faecalibaculum rodentium (0%|2.22%)
SP6 Mammaliicoccus lentus (0%|9.37%)
SP4 Staphylococcus saprophyticus (0%|7.16%)
SP7 Ligilactobacillus murinus (6.30%|2.06%)
SP8 Limosilactobacillus reuteri (2.48%|2.12%)
SP12 Lactobacillus johnsonii (3.27%|3.60%)
SP20 Lactobacillus intestinalis (2.35%|1.41%)
WT_D0_Fecal
WT_Piroxicam_Fecal
substitutions/site
0
0.1
0.2
0.3
0.4
0.5