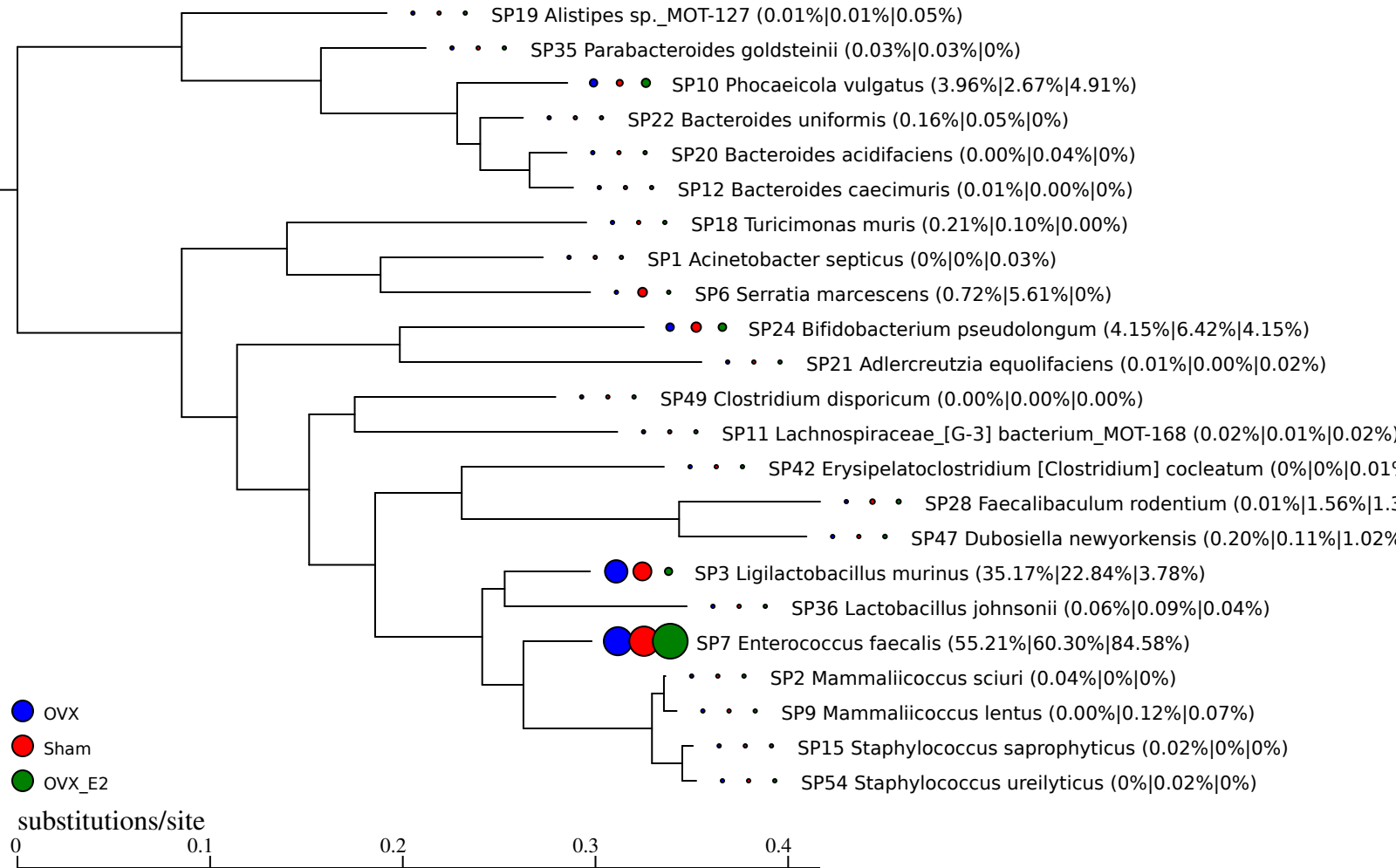
SP19 Alistipes sp._MOT-127 (0.01%|0.01%|0.05%)
SP35 Parabacteroides goldsteinii (0.03%|0.03%|0%)
SP10 Phocaeicola vulgatus (3.96%|2.67%|4.91%)
SP22 Bacteroides uniformis (0.16%|0.05%|0%)
SP20 Bacteroides acidifaciens (0.00%|0.04%|0%)
SP12 Bacteroides caecimuris (0.01%|0.00%|0%)
SP18 Turicimonas muris (0.21%|0.10%|0.00%)
SP1 Acinetobacter septicus (0%|0%|0.03%)
SP6 Serratia marcescens (0.72%|5.61%|0%)
SP24 Bifidobacterium pseudolongum (4.15%|6.42%|4.15%)
SP21 Adlercreutzia equolifaciens (0.01%|0.00%|0.02%)
SP49 Clostridium disporicum (0.00%|0.00%|0.00%)
SP11 Lachnospiraceae_[G-3] bacterium_MOT-168 (0.02%|0.01%|0.02%)
SP42 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|0.01
SP28 Faecalibaculum rodentium (0.01%|1.56%|1.3
SP47 Dubosiella newyorkensis (0.20%|0.11%|1.02%
SP3 Ligilactobacillus murinus (35.17%|22.84%|3.78%)
SP36 Lactobacillus johnsonii (0.06%|0.09%|0.04%)
SP7 Enterococcus faecalis (55.21%|60.30%|84.58%)
SP2 Mammaliicoccus sciuri (0.04%|0%|0%)
SP9 Mammaliicoccus lentus (0.00%|0.12%|0.07%)
SP15 Staphylococcus saprophyticus (0.02%|0%|0%)
SP54 Staphylococcus ureilyticus (0%|0.02%|0%)
OVX
Sham
OVX_E2
substitutions/site
0
0.1
0.2
0.3
0.4