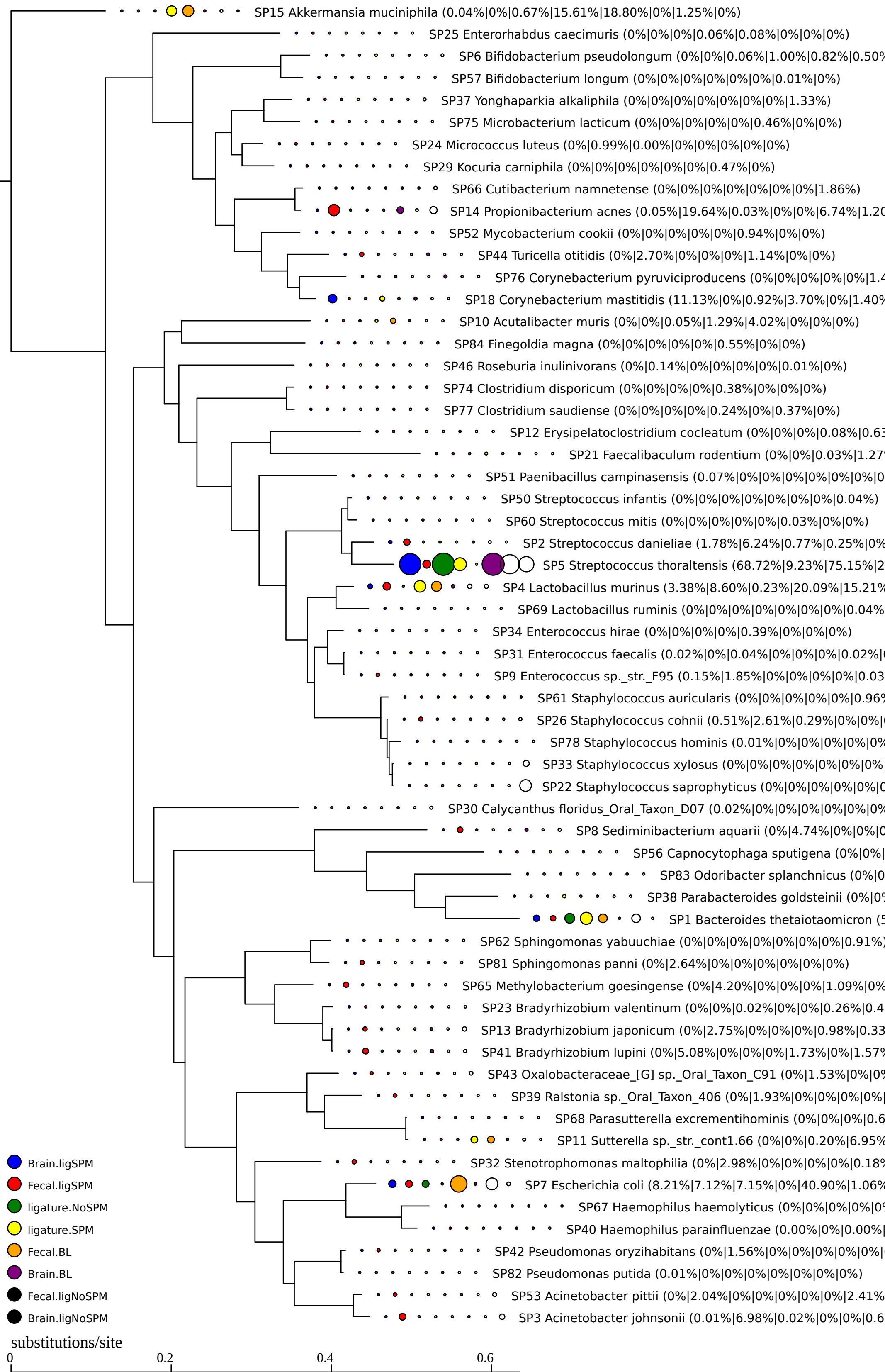
SP15 Akkermansia muciniphila (0.04%|0%|0.67%|15.61%|18.80%|0%|1.25%|0%)
SP25 Enterorhabdus caecimuris (0%|0%|0%|0.06%|0.08%|0%|0%|0%)
SP6 Bifidobacterium pseudolongum (0%|0%|0.06%|1.00%|0.82%|0.50%
SP57 Bifidobacterium longum (0%|0%|0%|0%|0%|0%|0.01%|0%)
SP37 Yonghaparkia alkaliphila (0%|0%|0%|0%|0%|0%|0%|1.33%)
SP75 Microbacterium lacticum (0%|0%|0%|0%|0%|0.46%|0%|0%)
SP24 Micrococcus luteus (0%|0.99%|0.00%|0%|0%|0%|0%|0%)
SP29 Kocuria carniphila (0%|0%|0%|0%|0%|0%|0.47%|0%)
SP66 Cutibacterium namnetense (0%|0%|0%|0%|0%|0%|0%|1.86%)
SP14 Propionibacterium acnes (0.05%|19.64%|0.03%|0%|0%|6.74%|1.20
SP52 Mycobacterium cookii (0%|0%|0%|0%|0%|0.94%|0%|0%)
SP44 Turicella otitidis (0%|2.70%|0%|0%|0%|1.14%|0%|0%)
SP76 Corynebacterium pyruviciproducens (0%|0%|0%|0%|0%|1.4
SP18 Corynebacterium mastitidis (11.13%|0%|0.92%|3.70%|0%|1.40%
SP10 Acutalibacter muris (0%|0%|0.05%|1.29%|4.02%|0%|0%|0%)
SP84 Finegoldia magna (0%|0%|0%|0%|0%|0.55%|0%|0%)
SP46 Roseburia inulinivorans (0%|0.14%|0%|0%|0%|0%|0.01%|0%)
SP74 Clostridium disporicum (0%|0%|0%|0%|0.38%|0%|0%|0%)
SP77 Clostridium saudiense (0%|0%|0%|0%|0.24%|0%|0.37%|0%)
SP12 Erysipelatoclostridium cocleatum (0%|0%|0%|0.08%|0.63
SP21 Faecalibaculum rodentium (0%|0%|0.03%|1.27
SP51 Paenibacillus campinasensis (0.07%|0%|0%|0%|0%|0%|0%|0
SP50 Streptococcus infantis (0%|0%|0%|0%|0%|0%|0%|0.04%)
SP60 Streptococcus mitis (0%|0%|0%|0%|0%|0.03%|0%|0%)
SP2 Streptococcus danieliae (1.78%|6.24%|0.77%|0.25%|0%
SP5 Streptococcus thoraltensis (68.72%|9.23%|75.15%|2
SP4 Lactobacillus murinus (3.38%|8.60%|0.23%|20.09%|15.21%
SP69 Lactobacillus ruminis (0%|0%|0%|0%|0%|0%|0%|0.04%
SP34 Enterococcus hirae (0%|0%|0%|0%|0.39%|0%|0%|0%)
SP31 Enterococcus faecalis (0.02%|0%|0.04%|0%|0%|0%|0.02%|
SP9 Enterococcus sp._str._F95 (0.15%|1.85%|0%|0%|0%|0%|0.03
SP61 Staphylococcus auricularis (0%|0%|0%|0%|0%|0.96%
SP26 Staphylococcus cohnii (0.51%|2.61%|0.29%|0%|0%|
SP78 Staphylococcus hominis (0.01%|0%|0%|0%|0%|0%
SP33 Staphylococcus xylosus (0%|0%|0%|0%|0%|0%|0%|
SP22 Staphylococcus saprophyticus (0%|0%|0%|0%|0%|0
SP30 Calycanthus floridus_Oral_Taxon_D07 (0.02%|0%|0%|0%|0%|0%|0%
SP8 Sediminibacterium aquarii (0%|4.74%|0%|0%|0
SP56 Capnocytophaga sputigena (0%|0%|
SP83 Odoribacter splanchnicus (0%|0
SP38 Parabacteroides goldsteinii (0%|0%
SP1 Bacteroides thetaiotaomicron (5
SP62 Sphingomonas yabuuchiae (0%|0%|0%|0%|0%|0%|0%|0.91%)
SP81 Sphingomonas panni (0%|2.64%|0%|0%|0%|0%|0%|0%)
SP65 Methylobacterium goesingense (0%|4.20%|0%|0%|0%|1.09%|0%
SP23 Bradyrhizobium valentinum (0%|0%|0.02%|0%|0%|0.26%|0.4
SP13 Bradyrhizobium japonicum (0%|2.75%|0%|0%|0%|0.98%|0.33
SP41 Bradyrhizobium lupini (0%|5.08%|0%|0%|0%|1.73%|0%|1.57%
SP43 Oxalobacteraceae_[G] sp._Oral_Taxon_C91 (0%|1.53%|0%|0%
SP39 Ralstonia sp._Oral_Taxon_406 (0%|1.93%|0%|0%|0%|0%|
SP68 Parasutterella excrementihominis (0%|0%|0%|0.6
SP11 Sutterella sp._str._cont1.66 (0%|0%|0.20%|6.95%
SP32 Stenotrophomonas maltophilia (0%|2.98%|0%|0%|0%|0%|0.18%
SP7 Escherichia coli (8.21%|7.12%|7.15%|0%|40.90%|1.06%
SP67 Haemophilus haemolyticus (0%|0%|0%|0%|0%
SP40 Haemophilus parainfluenzae (0.00%|0%|0.00%|
SP42 Pseudomonas oryzihabitans (0%|1.56%|0%|0%|0%|0%|0%|
SP82 Pseudomonas putida (0.01%|0%|0%|0%|0%|0%|0%|0%)
SP53 Acinetobacter pittii (0%|2.04%|0%|0%|0%|0%|0%|2.41%
SP3 Acinetobacter johnsonii (0.01%|6.98%|0.02%|0%|0%|0.6
Brain.ligSPM
Fecal.ligSPM
ligature.NoSPM
ligature.SPM
Fecal.BL
Brain.BL
Fecal.ligNoSPM
Brain.ligNoSPM
substitutions/site
0
0.2
0.4
0.6