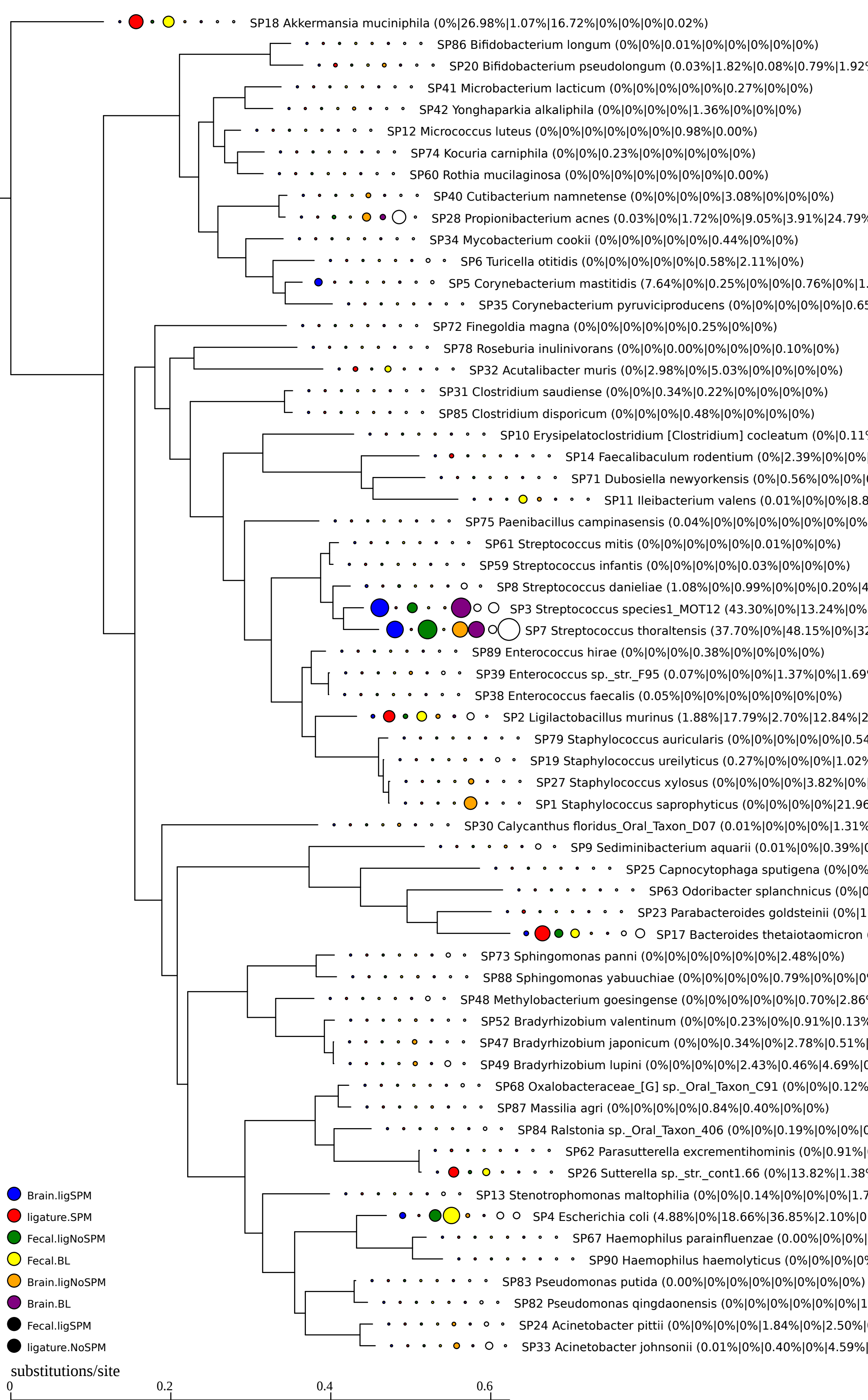
SP18 Akkermansia muciniphila (0%|26.98%|1.07%|16.72%|0%|0%|0%|0.02%)
SP86 Bifidobacterium longum (0%|0%|0.01%|0%|0%|0%|0%|0%)
SP20 Bifidobacterium pseudolongum (0.03%|1.82%|0.08%|0.79%|1.92
SP41 Microbacterium lacticum (0%|0%|0%|0%|0%|0.27%|0%|0%)
SP42 Yonghaparkia alkaliphila (0%|0%|0%|0%|1.36%|0%|0%|0%)
SP12 Micrococcus luteus (0%|0%|0%|0%|0%|0%|0.98%|0.00%)
SP74 Kocuria carniphila (0%|0%|0.23%|0%|0%|0%|0%|0%)
SP60 Rothia mucilaginosa (0%|0%|0%|0%|0%|0%|0%|0.00%)
SP40 Cutibacterium namnetense (0%|0%|0%|0%|3.08%|0%|0%|0%)
SP28 Propionibacterium acnes (0.03%|0%|1.72%|0%|9.05%|3.91%|24.79%
SP34 Mycobacterium cookii (0%|0%|0%|0%|0%|0.44%|0%|0%)
SP6 Turicella otitidis (0%|0%|0%|0%|0%|0.58%|2.11%|0%)
SP5 Corynebacterium mastitidis (7.64%|0%|0.25%|0%|0%|0.76%|0%|1.
SP35 Corynebacterium pyruviciproducens (0%|0%|0%|0%|0%|0.65
SP72 Finegoldia magna (0%|0%|0%|0%|0%|0.25%|0%|0%)
SP78 Roseburia inulinivorans (0%|0%|0.00%|0%|0%|0%|0.10%|0%)
SP32 Acutalibacter muris (0%|2.98%|0%|5.03%|0%|0%|0%|0%)
SP31 Clostridium saudiense (0%|0%|0.34%|0.22%|0%|0%|0%|0%)
SP85 Clostridium disporicum (0%|0%|0%|0.48%|0%|0%|0%|0%)
SP10 Erysipelatoclostridium [Clostridium] cocleatum (0%|0.11%
SP14 Faecalibaculum rodentium (0%|2.39%|0%|0%
SP71 Dubosiella newyorkensis (0%|0.56%|0%|0%|
SP11 Ileibacterium valens (0.01%|0%|0%|8.8
SP75 Paenibacillus campinasensis (0.04%|0%|0%|0%|0%|0%|0%|0%
SP61 Streptococcus mitis (0%|0%|0%|0%|0%|0.01%|0%|0%)
SP59 Streptococcus infantis (0%|0%|0%|0%|0.03%|0%|0%|0%)
SP8 Streptococcus danieliae (1.08%|0%|0.99%|0%|0%|0.20%|4
SP3 Streptococcus species1_MOT12 (43.30%|0%|13.24%|0%
SP7 Streptococcus thoraltensis (37.70%|0%|48.15%|0%|32
SP89 Enterococcus hirae (0%|0%|0%|0.38%|0%|0%|0%|0%)
SP39 Enterococcus sp._str._F95 (0.07%|0%|0%|0%|1.37%|0%|1.69%
SP38 Enterococcus faecalis (0.05%|0%|0%|0%|0%|0%|0%|0%)
SP2 Ligilactobacillus murinus (1.88%|17.79%|2.70%|12.84%|2
SP79 Staphylococcus auricularis (0%|0%|0%|0%|0%|0.54
SP19 Staphylococcus ureilyticus (0.27%|0%|0%|0%|1.02%
SP27 Staphylococcus xylosus (0%|0%|0%|0%|3.82%|0%
SP1 Staphylococcus saprophyticus (0%|0%|0%|0%|21.96
SP30 Calycanthus floridus_Oral_Taxon_D07 (0.01%|0%|0%|0%|1.31%
SP9 Sediminibacterium aquarii (0.01%|0%|0.39%|0
SP25 Capnocytophaga sputigena (0%|0%
SP63 Odoribacter splanchnicus (0%|0
SP23 Parabacteroides goldsteinii (0%|1
SP17 Bacteroides thetaiotaomicron
SP73 Sphingomonas panni (0%|0%|0%|0%|0%|0%|2.48%|0%)
SP88 Sphingomonas yabuuchiae (0%|0%|0%|0%|0.79%|0%|0%|0
SP48 Methylobacterium goesingense (0%|0%|0%|0%|0%|0.70%|2.86%
SP52 Bradyrhizobium valentinum (0%|0%|0.23%|0%|0.91%|0.13%
SP47 Bradyrhizobium japonicum (0%|0%|0.34%|0%|2.78%|0.51%|
SP49 Bradyrhizobium lupini (0%|0%|0%|0%|2.43%|0.46%|4.69%|0
SP68 Oxalobacteraceae_[G] sp._Oral_Taxon_C91 (0%|0%|0.12%
SP87 Massilia agri (0%|0%|0%|0%|0.84%|0.40%|0%|0%)
SP84 Ralstonia sp._Oral_Taxon_406 (0%|0%|0.19%|0%|0%|0
SP62 Parasutterella excrementihominis (0%|0.91%|
SP26 Sutterella sp._str._cont1.66 (0%|13.82%|1.38%
SP13 Stenotrophomonas maltophilia (0%|0%|0.14%|0%|0%|0%|1.7
SP4 Escherichia coli (4.88%|0%|18.66%|36.85%|2.10%|0
SP67 Haemophilus parainfluenzae (0.00%|0%|0%|
SP90 Haemophilus haemolyticus (0%|0%|0%|0%
SP83 Pseudomonas putida (0.00%|0%|0%|0%|0%|0%|0%|0%)
SP82 Pseudomonas qingdaonensis (0%|0%|0%|0%|0%|0%|1
SP24 Acinetobacter pittii (0%|0%|0%|0%|1.84%|0%|2.50%|
SP33 Acinetobacter johnsonii (0.01%|0%|0.40%|0%|4.59%|
Brain.ligSPM
ligature.SPM
Fecal.ligNoSPM
Fecal.BL
Brain.ligNoSPM
Brain.BL
Fecal.ligSPM
ligature.NoSPM
substitutions/site
0
0.2
0.4
0.6