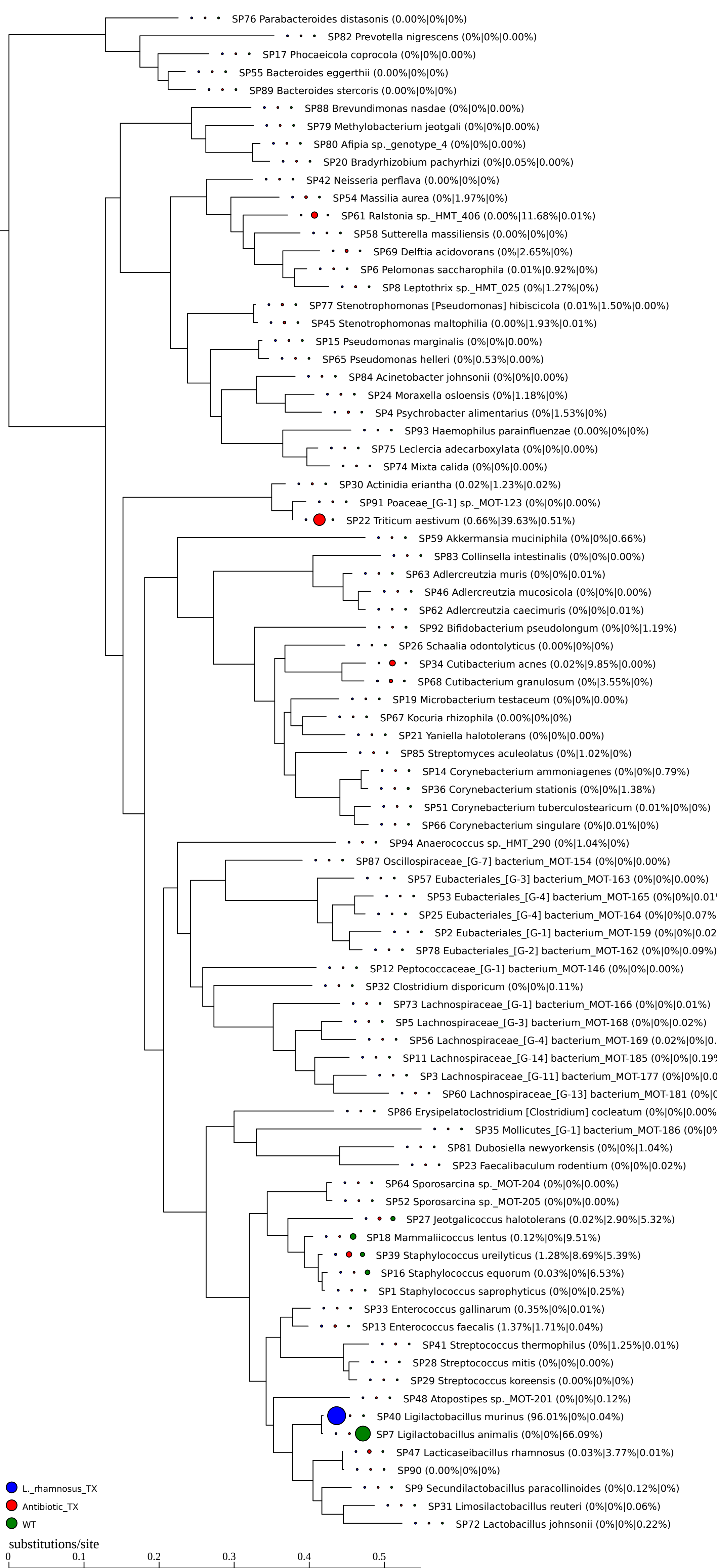
SP76 Parabacteroides distasonis (0.00%|0%|0%)
SP82 Prevotella nigrescens (0%|0%|0.00%)
SP17 Phocaeicola coprocola (0%|0%|0.00%)
SP55 Bacteroides eggerthii (0.00%|0%|0%)
SP89 Bacteroides stercoris (0.00%|0%|0%)
SP88 Brevundimonas nasdae (0%|0%|0.00%)
SP79 Methylobacterium jeotgali (0%|0%|0.00%)
SP80 Afipia sp._genotype_4 (0%|0%|0.00%)
SP20 Bradyrhizobium pachyrhizi (0%|0.05%|0.00%)
SP42 Neisseria perflava (0.00%|0%|0%)
SP54 Massilia aurea (0%|1.97%|0%)
SP61 Ralstonia sp._HMT_406 (0.00%|11.68%|0.01%)
SP58 Sutterella massiliensis (0.00%|0%|0%)
SP69 Delftia acidovorans (0%|2.65%|0%)
SP6 Pelomonas saccharophila (0.01%|0.92%|0%)
SP8 Leptothrix sp._HMT_025 (0%|1.27%|0%)
SP77 Stenotrophomonas [Pseudomonas] hibiscicola (0.01%|1.50%|0.00%)
SP45 Stenotrophomonas maltophilia (0.00%|1.93%|0.01%)
SP15 Pseudomonas marginalis (0%|0%|0.00%)
SP65 Pseudomonas helleri (0%|0.53%|0.00%)
SP84 Acinetobacter johnsonii (0%|0%|0.00%)
SP24 Moraxella osloensis (0%|1.18%|0%)
SP4 Psychrobacter alimentarius (0%|1.53%|0%)
SP93 Haemophilus parainfluenzae (0.00%|0%|0%)
SP75 Leclercia adecarboxylata (0%|0%|0.00%)
SP74 Mixta calida (0%|0%|0.00%)
SP30 Actinidia eriantha (0.02%|1.23%|0.02%)
SP91 Poaceae_[G-1] sp._MOT-123 (0%|0%|0.00%)
SP22 Triticum aestivum (0.66%|39.63%|0.51%)
SP59 Akkermansia muciniphila (0%|0%|0.66%)
SP83 Collinsella intestinalis (0%|0%|0.00%)
SP63 Adlercreutzia muris (0%|0%|0.01%)
SP46 Adlercreutzia mucosicola (0%|0%|0.00%)
SP62 Adlercreutzia caecimuris (0%|0%|0.01%)
SP92 Bifidobacterium pseudolongum (0%|0%|1.19%)
SP26 Schaalia odontolyticus (0.00%|0%|0%)
SP34 Cutibacterium acnes (0.02%|9.85%|0.00%)
SP68 Cutibacterium granulosum (0%|3.55%|0%)
SP19 Microbacterium testaceum (0%|0%|0.00%)
SP67 Kocuria rhizophila (0.00%|0%|0%)
SP21 Yaniella halotolerans (0%|0%|0.00%)
SP85 Streptomyces aculeolatus (0%|1.02%|0%)
SP14 Corynebacterium ammoniagenes (0%|0%|0.79%)
SP36 Corynebacterium stationis (0%|0%|1.38%)
SP51 Corynebacterium tuberculostearicum (0.01%|0%|0%)
SP66 Corynebacterium singulare (0%|0.01%|0%)
SP94 Anaerococcus sp._HMT_290 (0%|1.04%|0%)
SP87 Oscillospiraceae_[G-7] bacterium_MOT-154 (0%|0%|0.00%)
SP57 Eubacteriales_[G-3] bacterium_MOT-163 (0%|0%|0.00%)
SP53 Eubacteriales_[G-4] bacterium_MOT-165 (0%|0%|0.01%
SP25 Eubacteriales_[G-4] bacterium_MOT-164 (0%|0%|0.07%
SP2 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|0.02
SP78 Eubacteriales_[G-2] bacterium_MOT-162 (0%|0%|0.09%)
SP12 Peptococcaceae_[G-1] bacterium_MOT-146 (0%|0%|0.00%)
SP32 Clostridium disporicum (0%|0%|0.11%)
SP73 Lachnospiraceae_[G-1] bacterium_MOT-166 (0%|0%|0.01%)
SP5 Lachnospiraceae_[G-3] bacterium_MOT-168 (0%|0%|0.02%)
SP56 Lachnospiraceae_[G-4] bacterium_MOT-169 (0.02%|0%|0.
SP11 Lachnospiraceae_[G-14] bacterium_MOT-185 (0%|0%|0.19%
SP3 Lachnospiraceae_[G-11] bacterium_MOT-177 (0%|0%|0.0
SP60 Lachnospiraceae_[G-13] bacterium_MOT-181 (0%|0
SP86 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|0.00%
SP35 Mollicutes_[G-1] bacterium_MOT-186 (0%|0%
SP81 Dubosiella newyorkensis (0%|0%|1.04%)
SP23 Faecalibaculum rodentium (0%|0%|0.02%)
SP64 Sporosarcina sp._MOT-204 (0%|0%|0.00%)
SP52 Sporosarcina sp._MOT-205 (0%|0%|0.00%)
SP27 Jeotgalicoccus halotolerans (0.02%|2.90%|5.32%)
SP18 Mammaliicoccus lentus (0.12%|0%|9.51%)
SP39 Staphylococcus ureilyticus (1.28%|8.69%|5.39%)
SP16 Staphylococcus equorum (0.03%|0%|6.53%)
SP1 Staphylococcus saprophyticus (0%|0%|0.25%)
SP33 Enterococcus gallinarum (0.35%|0%|0.01%)
SP13 Enterococcus faecalis (1.37%|1.71%|0.04%)
SP41 Streptococcus thermophilus (0%|1.25%|0.01%)
SP28 Streptococcus mitis (0%|0%|0.00%)
SP29 Streptococcus koreensis (0.00%|0%|0%)
SP48 Atopostipes sp._MOT-201 (0%|0%|0.12%)
SP40 Ligilactobacillus murinus (96.01%|0%|0.04%)
SP7 Ligilactobacillus animalis (0%|0%|66.09%)
SP47 Lacticaseibacillus rhamnosus (0.03%|3.77%|0.01%)
SP90 (0.00%|0%|0%)
SP9 Secundilactobacillus paracollinoides (0%|0.12%|0%)
SP31 Limosilactobacillus reuteri (0%|0%|0.06%)
SP72 Lactobacillus johnsonii (0%|0%|0.22%)
L._rhamnosus_TX
Antibiotic_TX
WT
substitutions/site
0
0.1
0.2
0.3
0.4
0.5