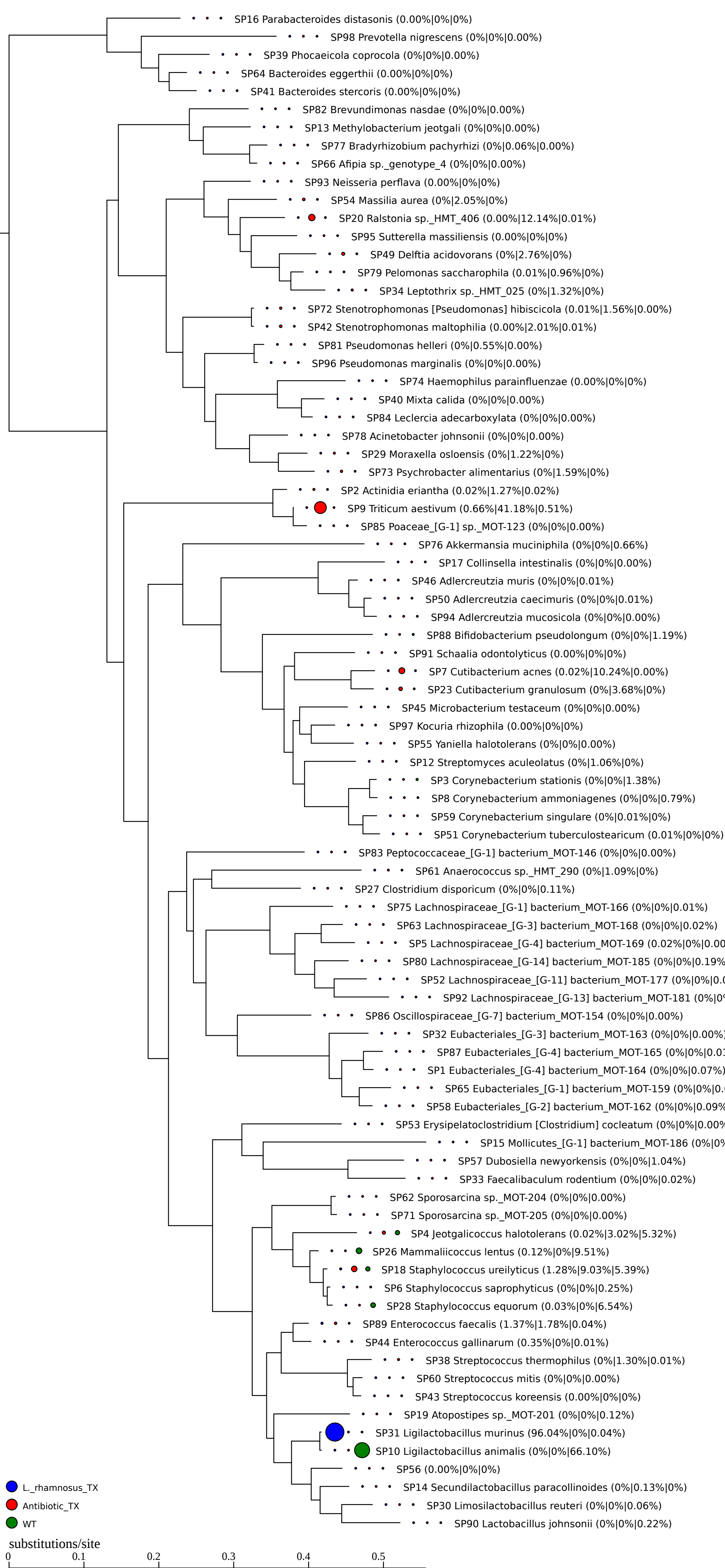
SP16 Parabacteroides distasonis (0.00%|0%|0%)
SP98 Prevotella nigrescens (0%|0%|0.00%)
SP39 Phocaeicola coprocola (0%|0%|0.00%)
SP64 Bacteroides eggerthii (0.00%|0%|0%)
SP41 Bacteroides stercoris (0.00%|0%|0%)
SP82 Brevundimonas nasdae (0%|0%|0.00%)
SP13 Methylobacterium jeotgali (0%|0%|0.00%)
SP77 Bradyrhizobium pachyrhizi (0%|0.06%|0.00%)
SP66 Afipia sp._genotype_4 (0%|0%|0.00%)
SP93 Neisseria perflava (0.00%|0%|0%)
SP54 Massilia aurea (0%|2.05%|0%)
SP20 Ralstonia sp._HMT_406 (0.00%|12.14%|0.01%)
SP95 Sutterella massiliensis (0.00%|0%|0%)
SP49 Delftia acidovorans (0%|2.76%|0%)
SP79 Pelomonas saccharophila (0.01%|0.96%|0%)
SP34 Leptothrix sp._HMT_025 (0%|1.32%|0%)
SP72 Stenotrophomonas [Pseudomonas] hibiscicola (0.01%|1.56%|0.00%)
SP42 Stenotrophomonas maltophilia (0.00%|2.01%|0.01%)
SP81 Pseudomonas helleri (0%|0.55%|0.00%)
SP96 Pseudomonas marginalis (0%|0%|0.00%)
SP74 Haemophilus parainfluenzae (0.00%|0%|0%)
SP40 Mixta calida (0%|0%|0.00%)
SP84 Leclercia adecarboxylata (0%|0%|0.00%)
SP78 Acinetobacter johnsonii (0%|0%|0.00%)
SP29 Moraxella osloensis (0%|1.22%|0%)
SP73 Psychrobacter alimentarius (0%|1.59%|0%)
SP2 Actinidia eriantha (0.02%|1.27%|0.02%)
SP9 Triticum aestivum (0.66%|41.18%|0.51%)
SP85 Poaceae_[G-1] sp._MOT-123 (0%|0%|0.00%)
SP76 Akkermansia muciniphila (0%|0%|0.66%)
SP17 Collinsella intestinalis (0%|0%|0.00%)
SP46 Adlercreutzia muris (0%|0%|0.01%)
SP50 Adlercreutzia caecimuris (0%|0%|0.01%)
SP94 Adlercreutzia mucosicola (0%|0%|0.00%)
SP88 Bifidobacterium pseudolongum (0%|0%|1.19%)
SP91 Schaalia odontolyticus (0.00%|0%|0%)
SP7 Cutibacterium acnes (0.02%|10.24%|0.00%)
SP23 Cutibacterium granulosum (0%|3.68%|0%)
SP45 Microbacterium testaceum (0%|0%|0.00%)
SP97 Kocuria rhizophila (0.00%|0%|0%)
SP55 Yaniella halotolerans (0%|0%|0.00%)
SP12 Streptomyces aculeolatus (0%|1.06%|0%)
SP3 Corynebacterium stationis (0%|0%|1.38%)
SP8 Corynebacterium ammoniagenes (0%|0%|0.79%)
SP59 Corynebacterium singulare (0%|0.01%|0%)
SP51 Corynebacterium tuberculostearicum (0.01%|0%|0%)
SP83 Peptococcaceae_[G-1] bacterium_MOT-146 (0%|0%|0.00%)
SP61 Anaerococcus sp._HMT_290 (0%|1.09%|0%)
SP27 Clostridium disporicum (0%|0%|0.11%)
SP75 Lachnospiraceae_[G-1] bacterium_MOT-166 (0%|0%|0.01%)
SP63 Lachnospiraceae_[G-3] bacterium_MOT-168 (0%|0%|0.02%)
SP5 Lachnospiraceae_[G-4] bacterium_MOT-169 (0.02%|0%|0.00
SP80 Lachnospiraceae_[G-14] bacterium_MOT-185 (0%|0%|0.19%
SP52 Lachnospiraceae_[G-11] bacterium_MOT-177 (0%|0%|0.0
SP92 Lachnospiraceae_[G-13] bacterium_MOT-181 (0%|0%
SP86 Oscillospiraceae_[G-7] bacterium_MOT-154 (0%|0%|0.00%)
SP32 Eubacteriales_[G-3] bacterium_MOT-163 (0%|0%|0.00%)
SP87 Eubacteriales_[G-4] bacterium_MOT-165 (0%|0%|0.01
SP1 Eubacteriales_[G-4] bacterium_MOT-164 (0%|0%|0.07%)
SP65 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|0.
SP58 Eubacteriales_[G-2] bacterium_MOT-162 (0%|0%|0.09%
SP53 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|0.00%
SP15 Mollicutes_[G-1] bacterium_MOT-186 (0%|0%
SP57 Dubosiella newyorkensis (0%|0%|1.04%)
SP33 Faecalibaculum rodentium (0%|0%|0.02%)
SP62 Sporosarcina sp._MOT-204 (0%|0%|0.00%)
SP71 Sporosarcina sp._MOT-205 (0%|0%|0.00%)
SP4 Jeotgalicoccus halotolerans (0.02%|3.02%|5.32%)
SP26 Mammaliicoccus lentus (0.12%|0%|9.51%)
SP18 Staphylococcus ureilyticus (1.28%|9.03%|5.39%)
SP6 Staphylococcus saprophyticus (0%|0%|0.25%)
SP28 Staphylococcus equorum (0.03%|0%|6.54%)
SP89 Enterococcus faecalis (1.37%|1.78%|0.04%)
SP44 Enterococcus gallinarum (0.35%|0%|0.01%)
SP38 Streptococcus thermophilus (0%|1.30%|0.01%)
SP60 Streptococcus mitis (0%|0%|0.00%)
SP43 Streptococcus koreensis (0.00%|0%|0%)
SP19 Atopostipes sp._MOT-201 (0%|0%|0.12%)
SP31 Ligilactobacillus murinus (96.04%|0%|0.04%)
SP10 Ligilactobacillus animalis (0%|0%|66.10%)
SP56 (0.00%|0%|0%)
SP14 Secundilactobacillus paracollinoides (0%|0.13%|0%)
SP30 Limosilactobacillus reuteri (0%|0%|0.06%)
SP90 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