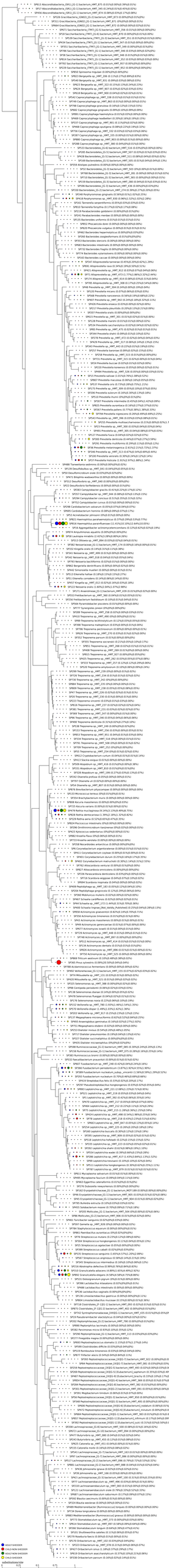
SP013 Abscondibacteria_(SR1_[G-1]) bacterium_HRT_343 (0.0%|0.0|0.0%|0.0%)
SP017 Abscondibacteria_(SR1_[G-1]) bacterium_HRT_343 (0.2%|0.0|0.0%|0.0%)
SP036 Abscondibacteria_(SR1_[G-1]) bacterium_HRT_343 (0.0%|0.0|0.0%|0.0%)
SP128 Gracilibacteria_(GN02_[G-2]) bacterium_HRT_873 (0.0%|0.0|0.0%|0.0%)
SP131 Gracilibacteria_(GN02_[G-2]) bacterium_HRT_873 (0.0%|0.0|0.0%|0.0%)
SP064 Parcubacteria_(OD1_[G-1]) bacterium_HRT_952 (0.0%|0.0|0.0%|0.0%)
SP125 Parcubacteria_(OD1_[G-1]) bacterium_HRT_952 (0.0%|0.0|0.0%|0.0%)
SP135 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_350 (0.0%|0.0|0.0%|0.0%)
SP139 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_351 (0.0%|0.0|0.0%|0.0%)
SP140 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_351 (0.0%|0.0|0.0%|0.0%)
SP151 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_349 (0.0%|0.0|0.0%|0.0%)
SP156 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_349 (0.0%|0.0|0.0%|0.0%)
SP158 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_348 (0.0%|0.0|0.0%|0.0%)
SP159 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_348 (0.0%|0.0|0.0%|0.0%)
SP160 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_348 (0.0%|0.0|0.0%|0.0%)
SP192 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_357 (0.0%|0.0|0.0%|0.0%)
SP193 Saccharibacteria_(TM7_[G-1]) bacterium_HRT_357 (0.0%|0.0|0.0%|0.0%)
SP051 Spirosoma_ningbense (0.0%|0.0|0.0%|0.0%)
SP013 Bergeyella sp._HRT_208 (0.1%|0.1|0.0%|0.0%)
SP040 Bergeyella sp._HRT_931 (0.0%|0.0|0.0%|0.0%)
SP053 Bergeyella sp._HRT_322 (0.1%|0.1|0.0%|0.0%)
SP054 Bergeyella sp._HRT_903 (0.0%|0.0|0.0%|0.0%)
SP063 Bergeyella sp._HRT_385 (0.0%|0.0|0.0%|0.0%)
SP042 Capnocytophaga sp._HRT_338 (0.0%|0.0|0.0%|0.0%)
SP049 Capnocytophaga sp._HRT_863 (0.0%|0.0|0.0%|0.0%)
SP058 Capnocytophaga granulosa (0.0%|0.0|0.0%|0.0%)
SP062 Capnocytophaga gingivalis (0.0%|0.0|0.0%|0.0%)
SP081 Capnocytophaga haemolytica (0.0%|0.0|0.0%|0.0%)
SP084 Capnocytophaga leadbetteri (0.0%|0.0|0.0%|0.0%)
SP103 Capnocytophaga sp._HRT_351 (0.1%|0.0|0.0%|0.0%)
SP060 Capnocytophaga ochracea (0.0%|0.0|0.0%|0.0%)
SP156 Capnocytophaga sp._HRT_332 (0.0%|0.0|0.0%|0.0%)
SP080 Capnocytophaga sp._HRT_195 (0.0%|0.0|0.0%|0.0%)
SP086 Capnocytophaga sp._HRT_903 (0.0%|0.0|0.0%|0.0%)
SP087 Capnocytophaga sp._HRT_380 (0.0%|0.0|0.0%|0.0%)
SP013 Sporobacter_(O.40) bacterium_HRT_516 (0.0%|0.0|0.0%|0.0%)
SP035 Bacteroides_(O.S) bacterium_HRT_511 (0.0%|0.0|0.0%|0.0%)
SP028 Bacteroides_(O.S) bacterium_HRT_511 (0.0%|0.0|0.0%|0.0%)
SP018 Bacteroides_(O.S) bacterium_HRT_511 (0.0%|0.0|0.0%|0.0%)
SP019 Alistipes putredinis (0.0%|0.0|0.0%|0.0%)
SP12 Bacteroides_(O.S) bacterium_HRT_291 (0.0%|0.0|0.0%|0.0%)
SP08 Bacteroides_(O.S) bacterium_HRT_291 (0.0%|0.0|0.0%|0.0%)
SP09 Bacteroides_(O.S) bacterium_HRT_291 (0.0%|0.0|0.0%|0.0%)
SP15 Bacteroides_(O.S) bacterium_HRT_280 (0.0%|0.0|0.0%|0.0%)
SP16 Bacteroides_(O.S) bacterium_HRT_280 (0.0%|0.0|0.0%|0.0%)
SP20 Bacteroides_(O.S) bacterium_HRT_274 (0.3%|0.2|0.2%|0.0%)
SP26 Porphyromonas gingivalis (0.5%|0.5|0.2%|0.2%)
SP418 Porphyromonas_sp._HRT_930 (0.9%|0.1|52%|0.2%)
SP21 Tannerella sarpentiformis (0.0%|0.0|0.0%|0.0%)
SP03 Tannerella forsythia (0.1%|0.0|0.0%|0.0%)
SP219 Parabacteroides goldsteinii (0.0%|0.0|0.0%|0.0%)
SP241 Parabacteroides merdae (0.0%|0.0|0.0%|0.0%)
SP128 Bacteroides uniformis (0.0%|0.0|0.0%|0.0%)
SP062 Prevotella direi (0.0%|0.0|0.0%|0.0%)
SP025 Prevotella corporis (0.0%|0.0|0.0%|0.0%)
SP042 Bacteroides hepatocytophilus (0.0%|0.0|0.0%|0.0%)
SP036 Bacteroides ovatus/xylanisolvens (0.0%|0.0|0.0%|0.0%)
SP053 Bacteroides vitrocis (0.0%|0.0|0.0%|0.0%)
SP065 Bacteroides thetaiotaomicron (0.0%|0.0|0.0%|0.0%)
SP072 Bacteroides fragilis (0.0%|0.0|0.0%|0.0%)
SP074 Bacteroides caccae/ovatus (0.0%|0.0|0.0%|0.0%)
SP075 Bacteroides caccae (0.0%|0.0|0.0%|0.0%)
SP147 Alistipes onderdonkii/senegalensis (0.2%|0.2|0.2%|0.2%)
SP061 Alistipes finegoldii