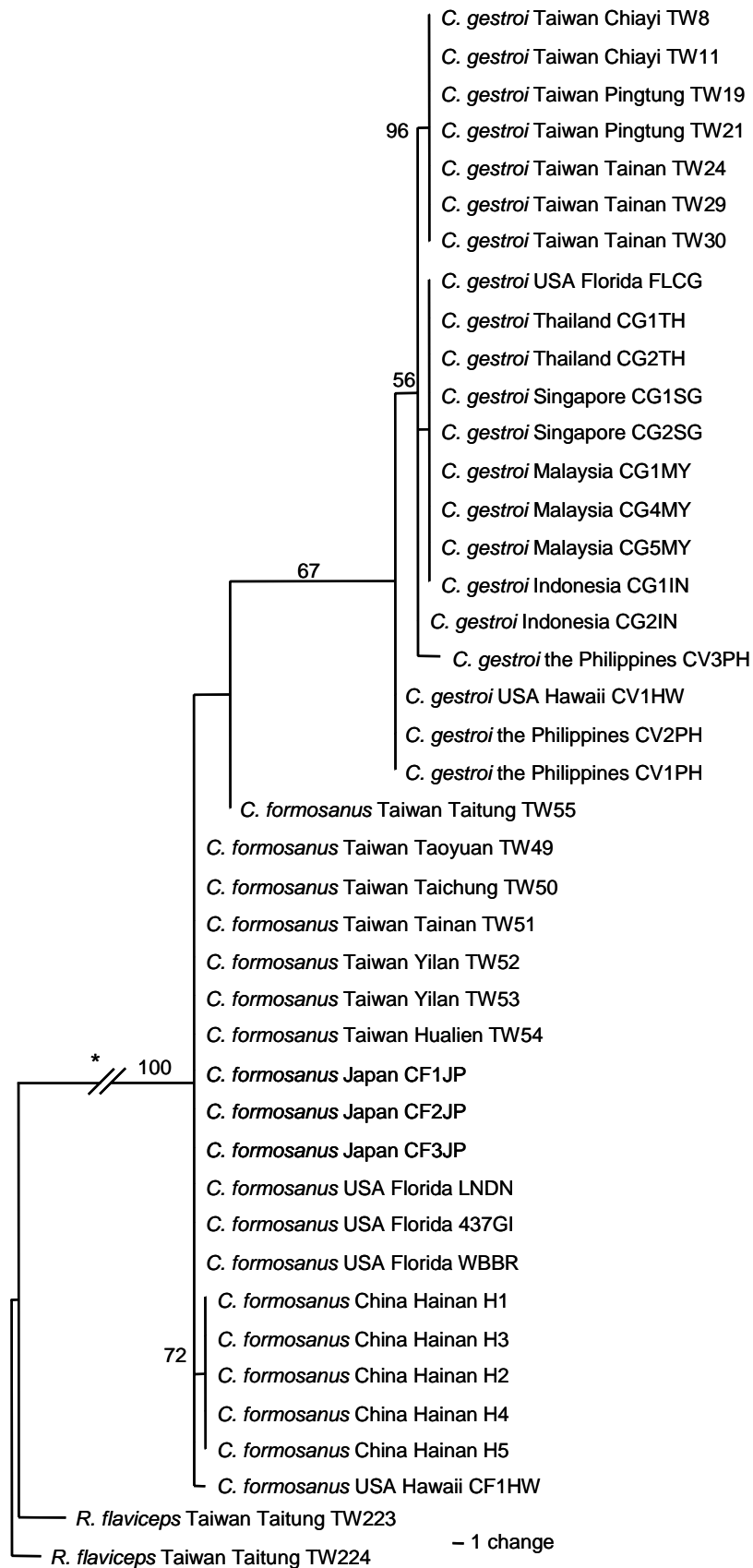


The 10001st Bayesian tree inferred from 12S rRNA gene sequences. Posterior probability values exceeding 50 are given on appropriate clades. *, > 20 changes.



The 10001st Bayesian tree inferred from 16S rRNA gene sequences. Posterior probability values exceeding 50 are given on appropriate clades. *, > 40 changes.