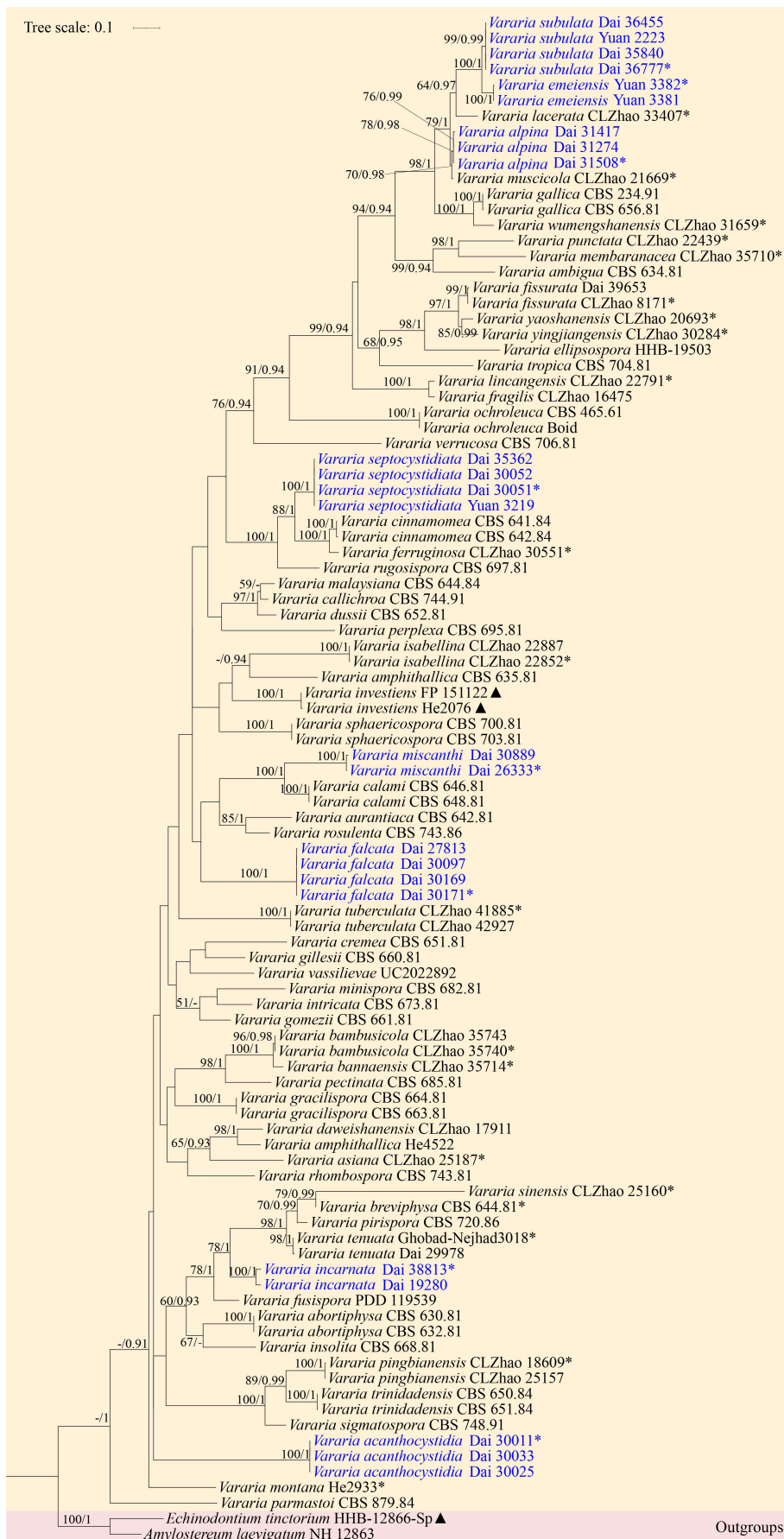




Supplementary Fig. S7 The phylogeny of *Vararia* was reconstructed using Maximum Likelihood analysis of ITS sequences alone. Bayesian analysis ran for 150 million generations (ASDSF < 0.05, ESS > 200). Branch nodes are annotated with Bootstrap support ($\geq 50\%$). Black triangle (▲) = type species; blue = new species.