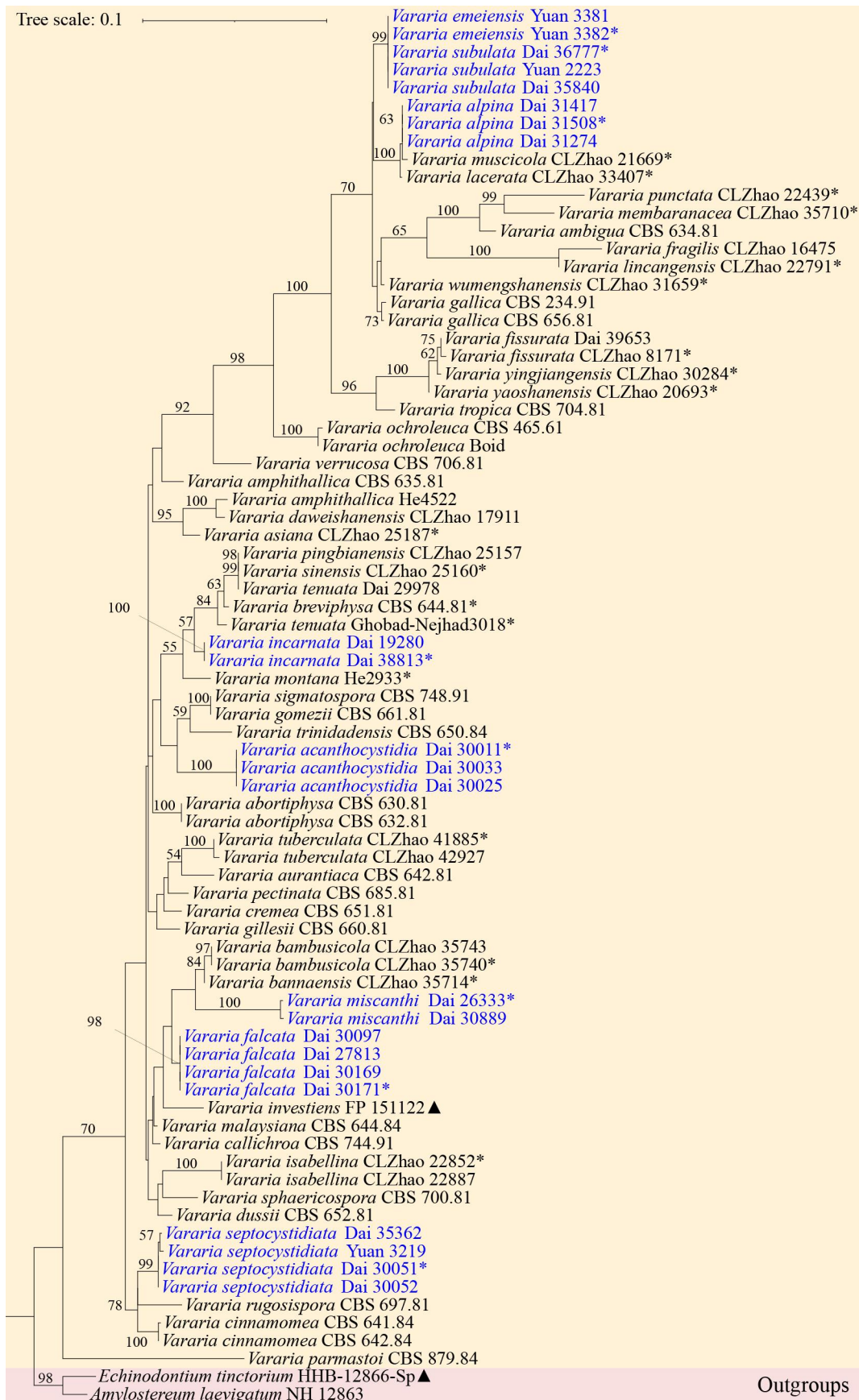




Supplementary Fig. S9 The phylogeny of *Vararia* was reconstructed using Maximum Likelihood analysis of combined ITS+LSU sequences (only the ML tree is presented). Branch nodes are annotated with Bootstrap support ($\geq 50\%$). Black triangle (▲) = type species; blue = new species.