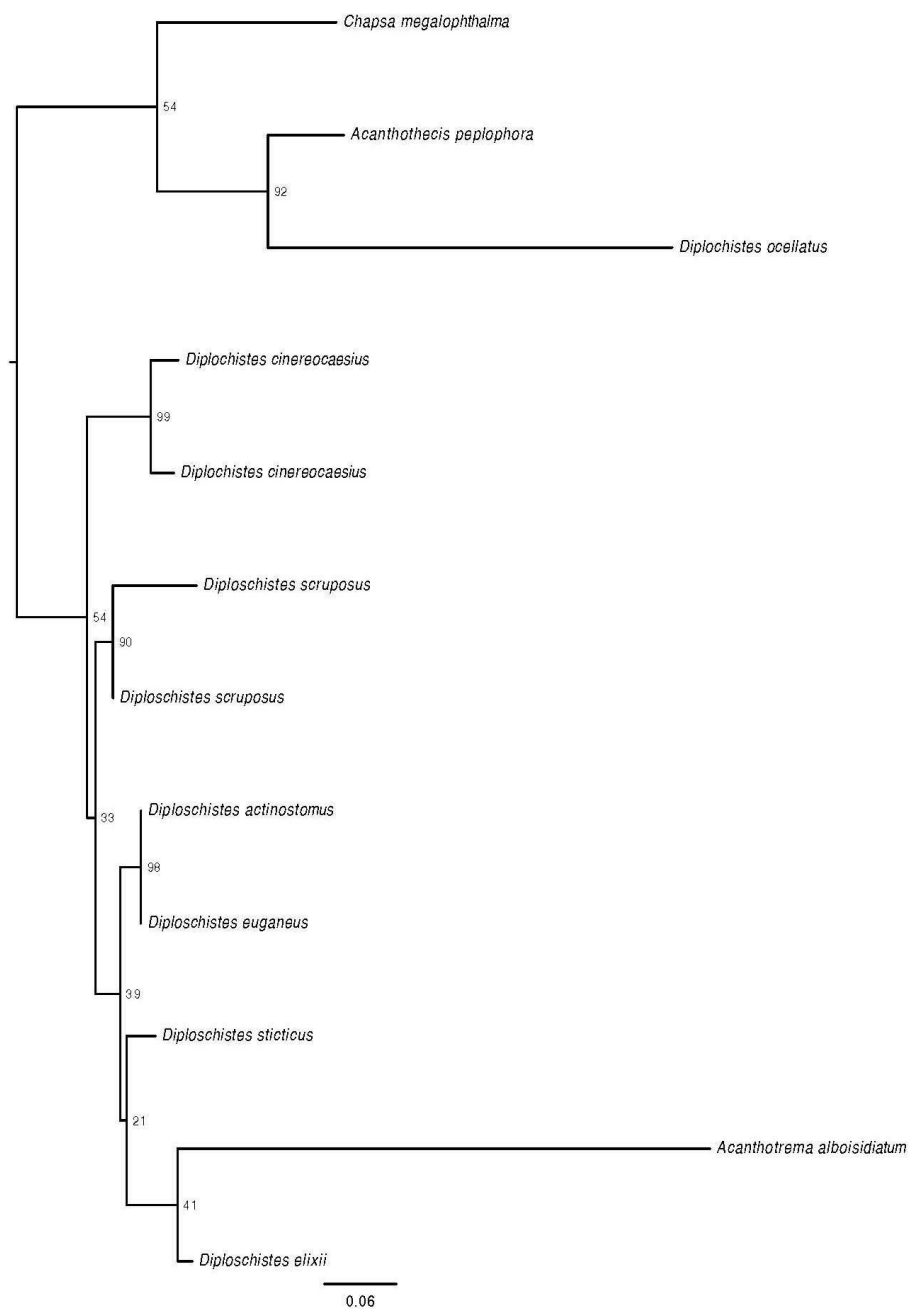


**Fig. S4.** Tree from a maximum-likelihood analysis of *RPB2* gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.



**Fig. S5.** Tree from a maximum-likelihood analysis of nuITS gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.