

Fig. S1. Tree from a maximum-likelihood analysis of mtSSU gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.

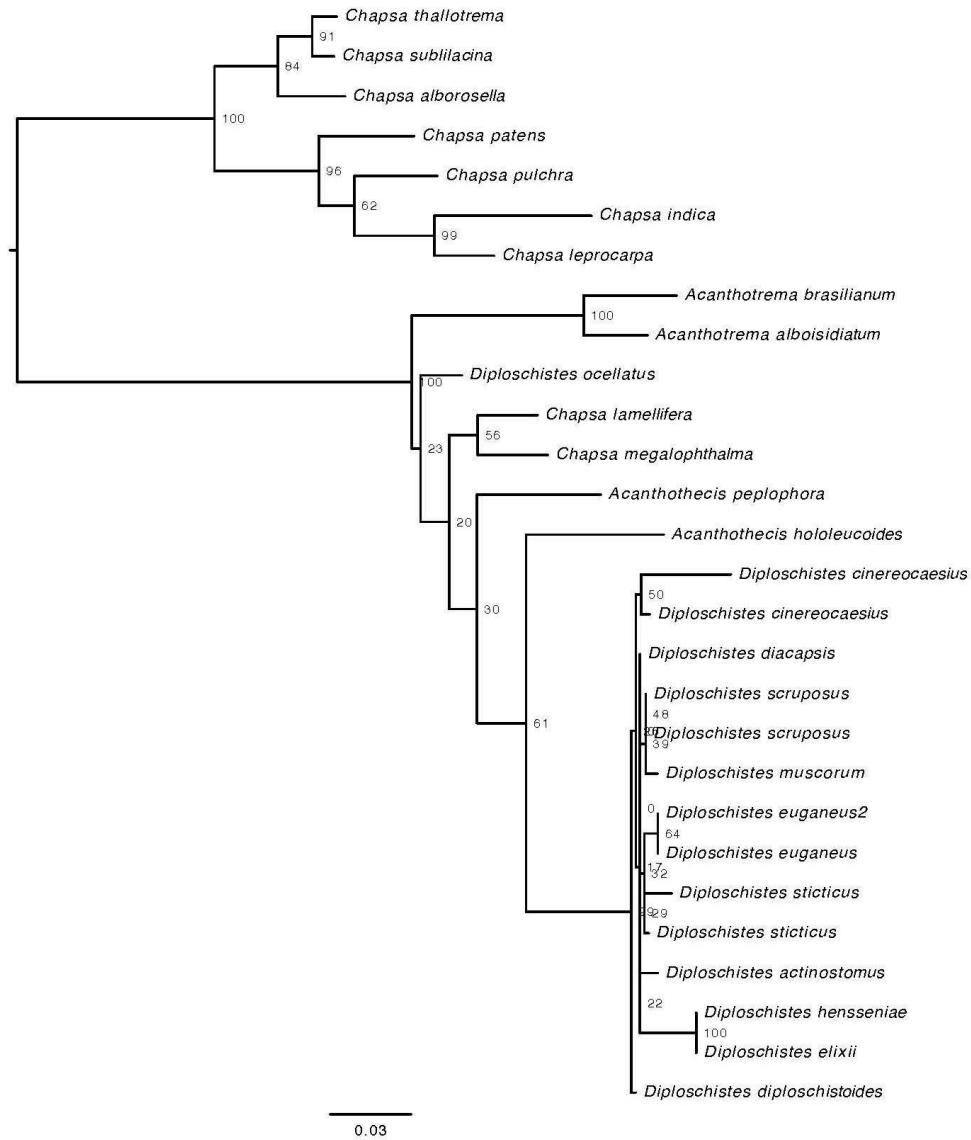


Fig. S2. Tree from a maximum-likelihood analysis of nuLSU gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.

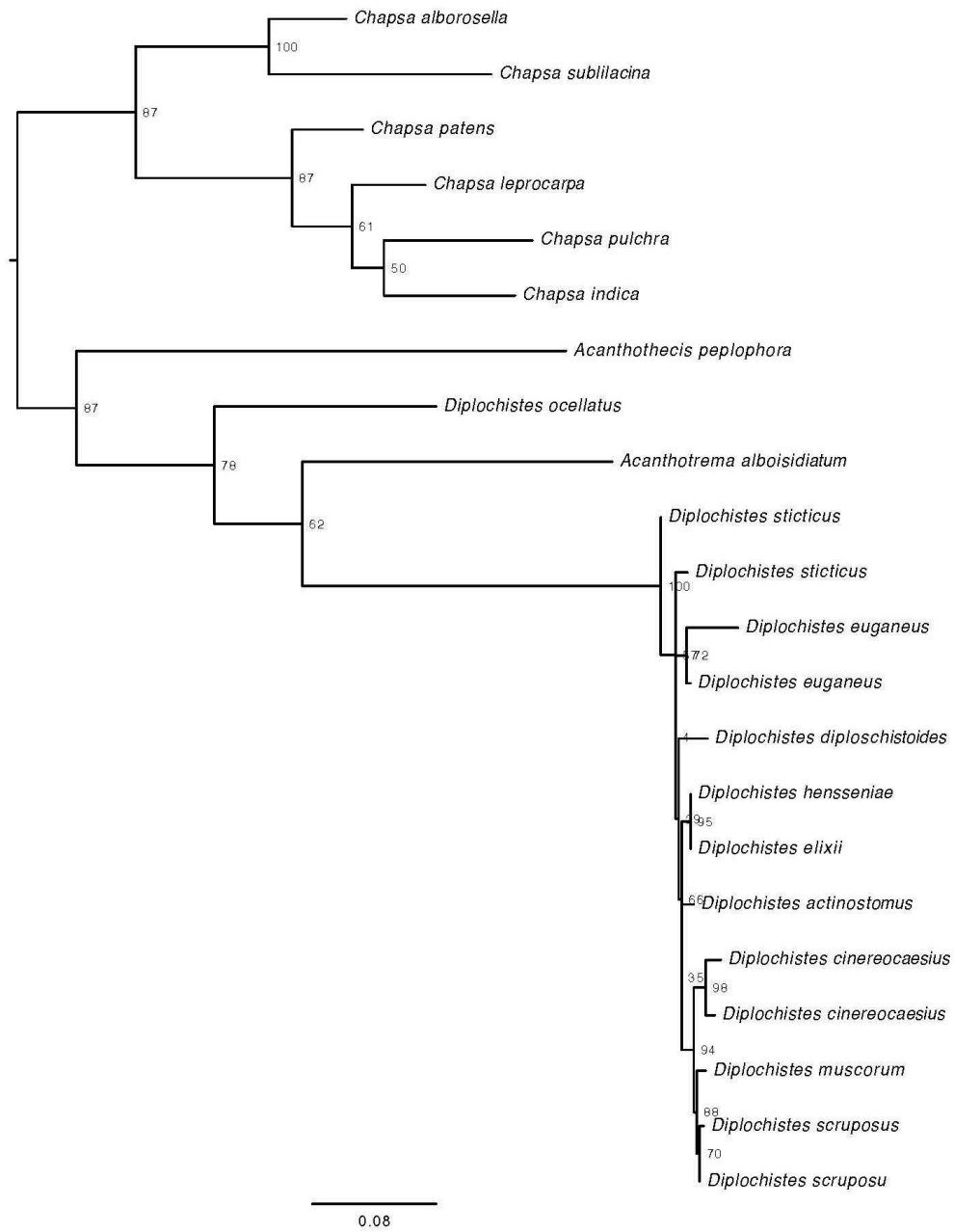


Fig. S3. Tree from a maximum-likelihood analysis of *RPBI* gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.

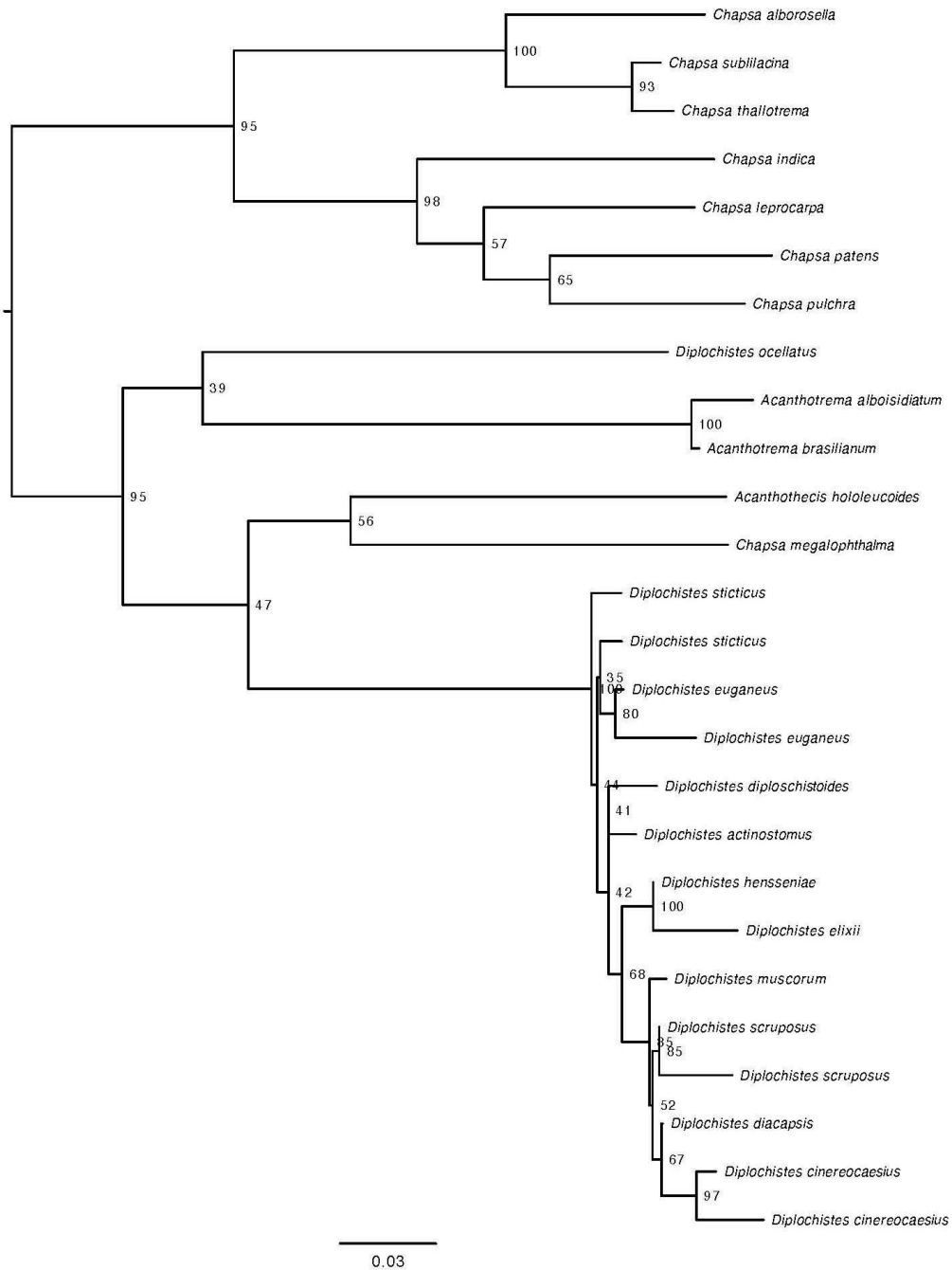


Fig. S4. Tree from a maximum-likelihood analysis of *RPB2* gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.

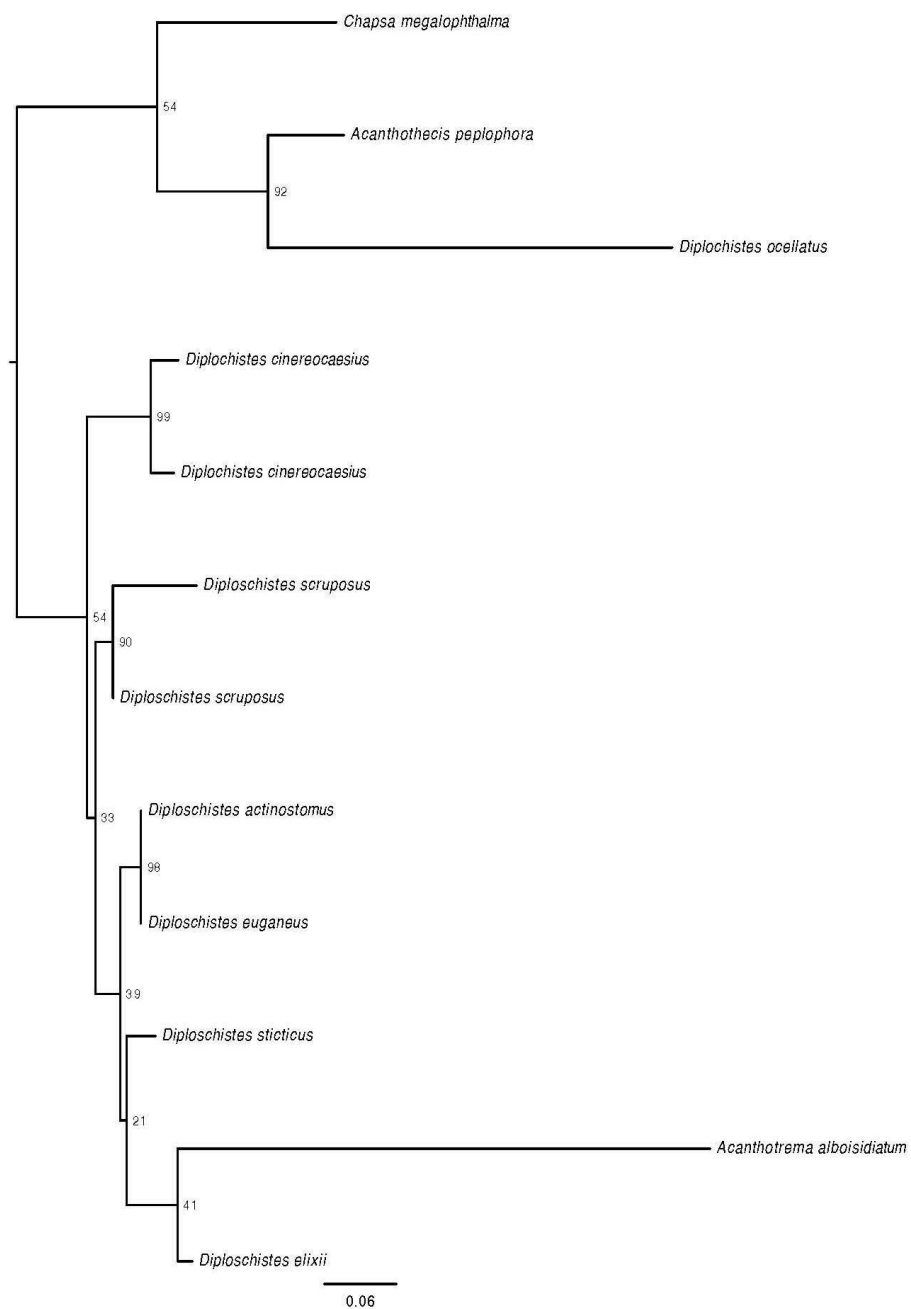


Fig. S5. Tree from a maximum-likelihood analysis of nuITS gene of *Diploschistes* and their relatives. The numbers at the nodes indicate their bootstrap support.