Supporting Information

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Fig. S1. Map with the current distribution of Rhacophoridae.



Fig. S2. Bayesian phylogenetic tree based on the nuclear dataset (Bayesian posterior probability ≥90% retained).



Fig. S3. Bayesian phylogenetic tree of the combined nuclear and mitochondrial dataset with posterior probabilities for branches (\geq 90% retained) and 95% credibility intervals for divergence time estimates with the full species name.

 Table S1. Samples, with sampling site, museum voucher nos., and GenBank accession nos. of corresponding sequences

Table S1

Table S2. Divergence times in million years before present and biogeographic reconstructions for nodes

Table S2

Table S3. Comparison of different temporal dispersal models using Lagrange, indicating that dispersal between the Indian subcontinent and Eurasia most likely occurred after 35 Ma, but before 25 Ma

Table S3

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