

New insights from a lost world

Unlocking the potential of museum collections using historical specimens

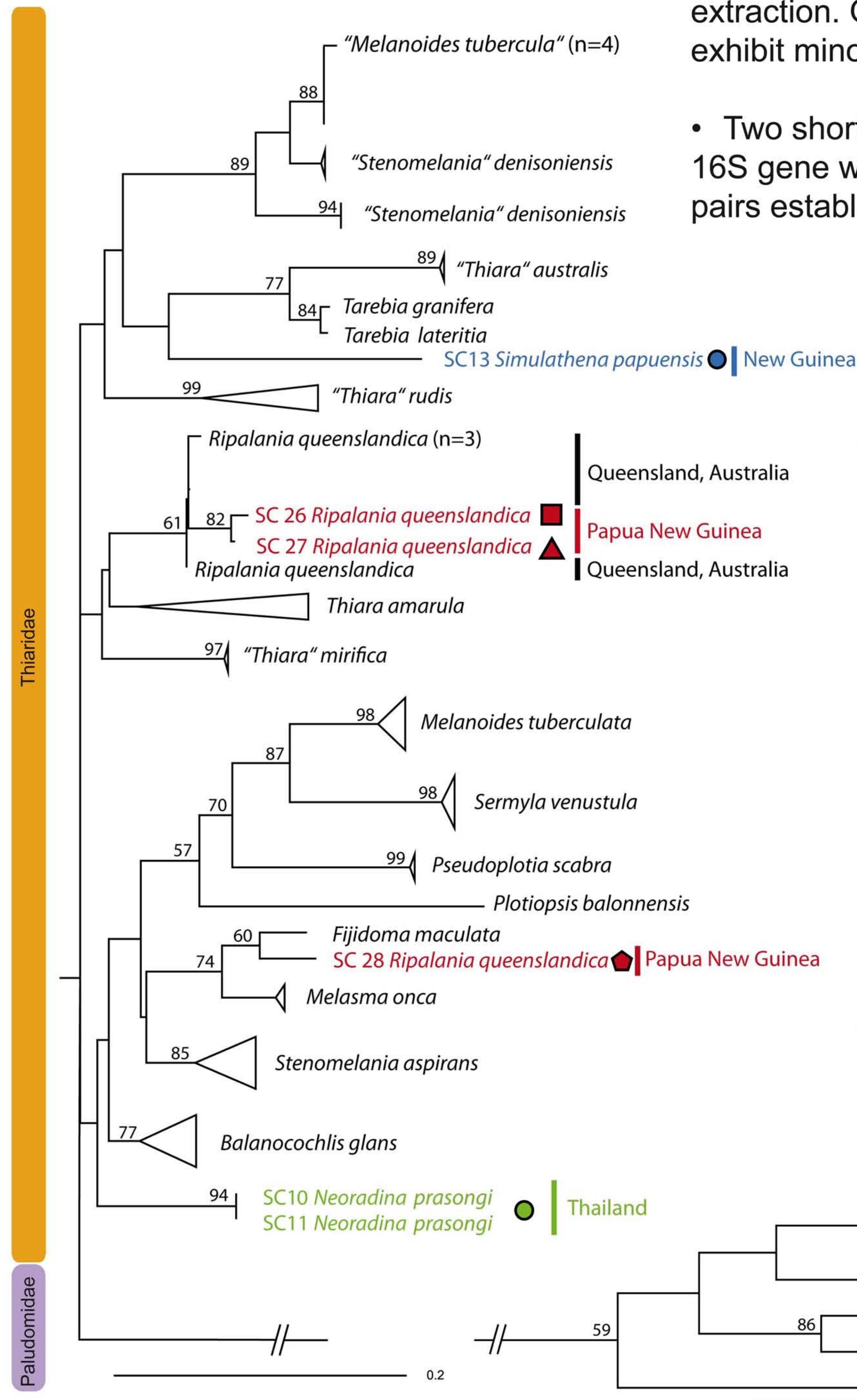
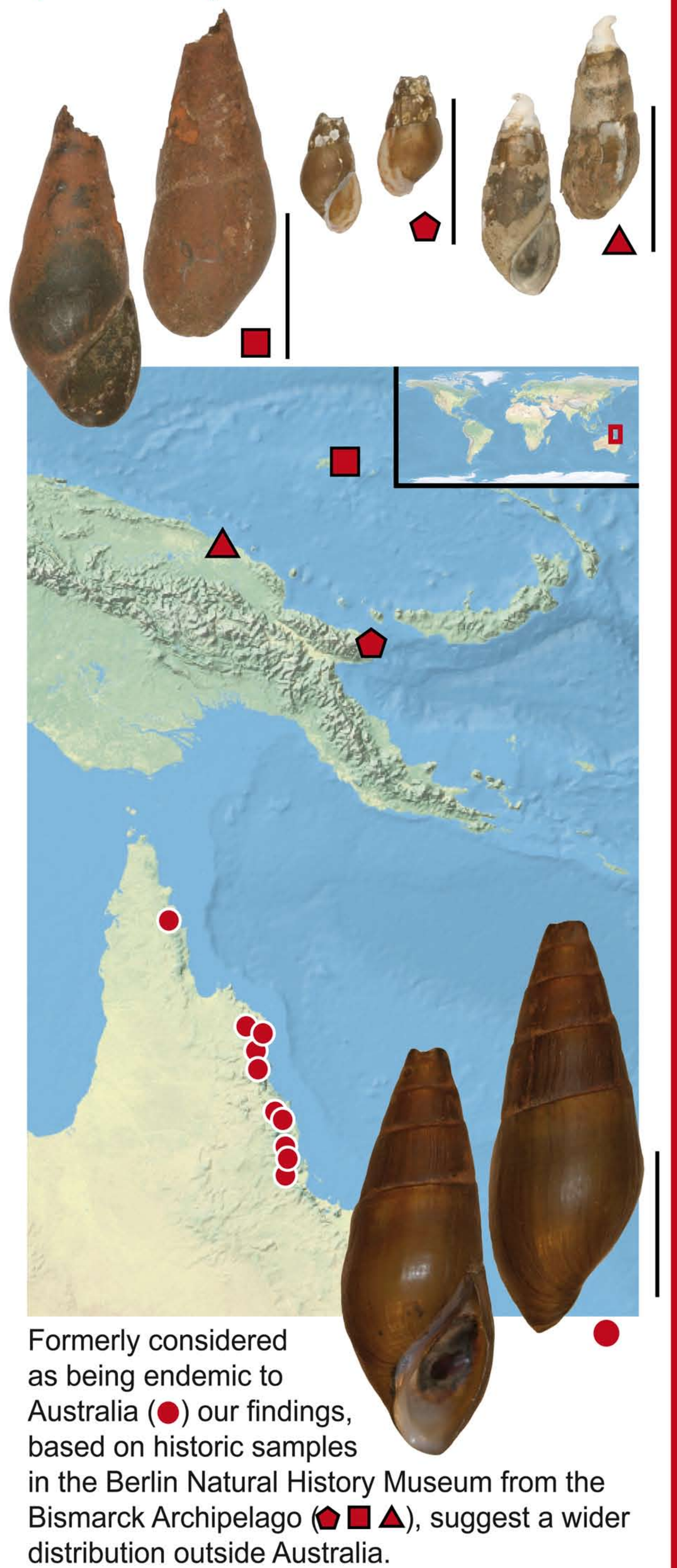
Matthias Glaubrecht¹, Alex Greenwood², Jana Ebersbach¹, Annabell Szymansky¹, Karin Hönig², Benedikt Wiggering¹, France Gimnich¹

¹Museum für Naturkunde Berlin, Leibniz Institute for Research in Evolution and Biodiversity at the Humboldt University, Invalidenstraße 43, 10115 Berlin, Germany;

²Leibniz Institute for Zoo and Wildlife Research in the Forschungsverbund Berlin e.V., Alfred-Kowalke-Straße 17, 10315 Berlin Germany
matthias.glaubrecht@mfn-berlin.de; benedikt.wiggering@mfn-berlin.de; france.gimnich@mfn-berlin.de

We tested a non-destructive ancient DNA extraction protocol for historical mollusks specimens, sampled during the last century, in order to explore for use and tap malacological museum collections, as e.g. from the Berlin Natural History Museum. The procedure was considered suitable for various historic tissue and shell samples from limnic thiarid and paludomid gastropods, for which here four case studies from SE Asia and Australia underline the potential of this method.

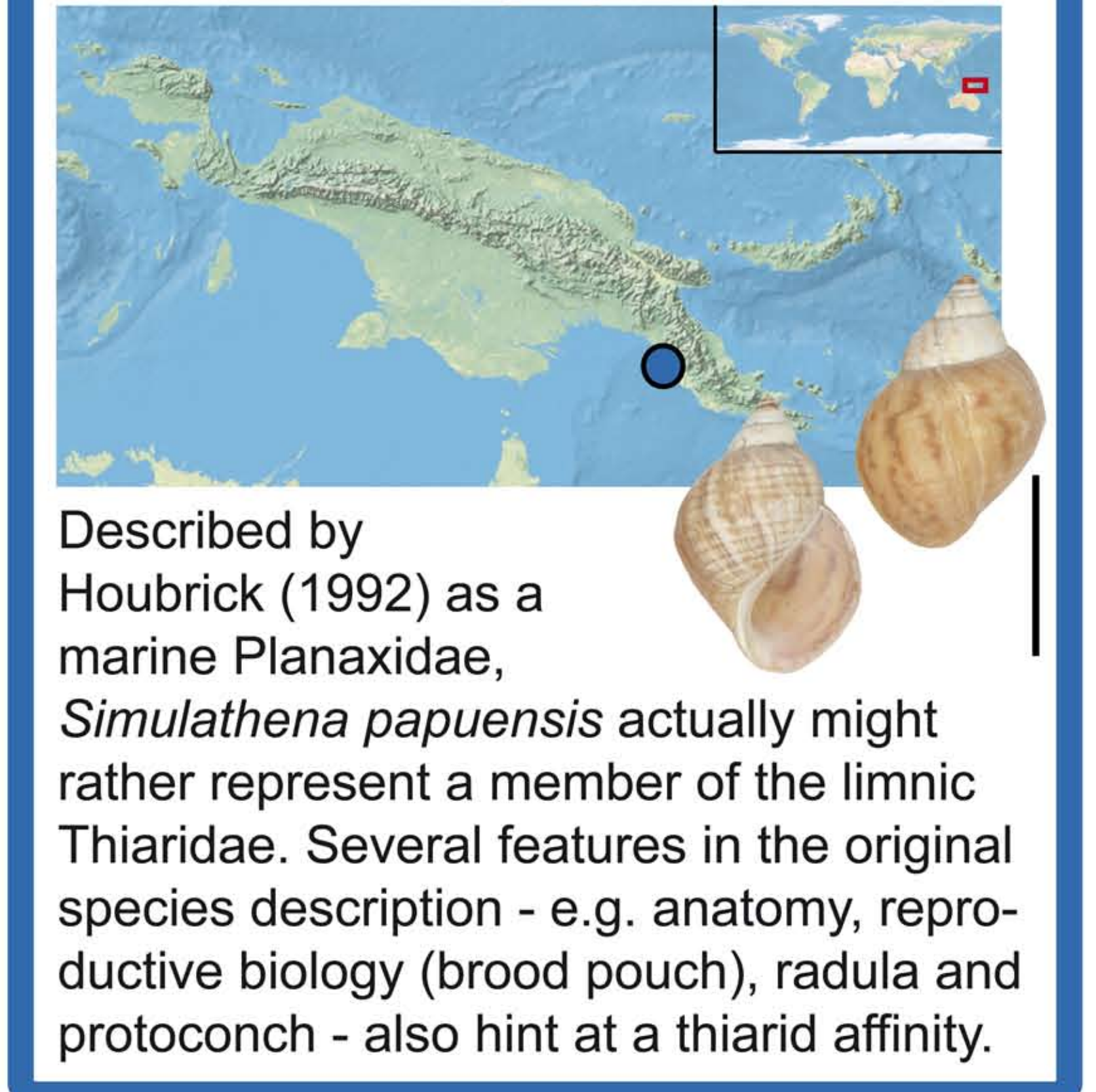
Ripalania queenslandica



- Entire shells or fragments, operculum and foot tissue from ethanol samples were used for DNA extraction. Only the periostracum in few shells exhibit minor alterations.

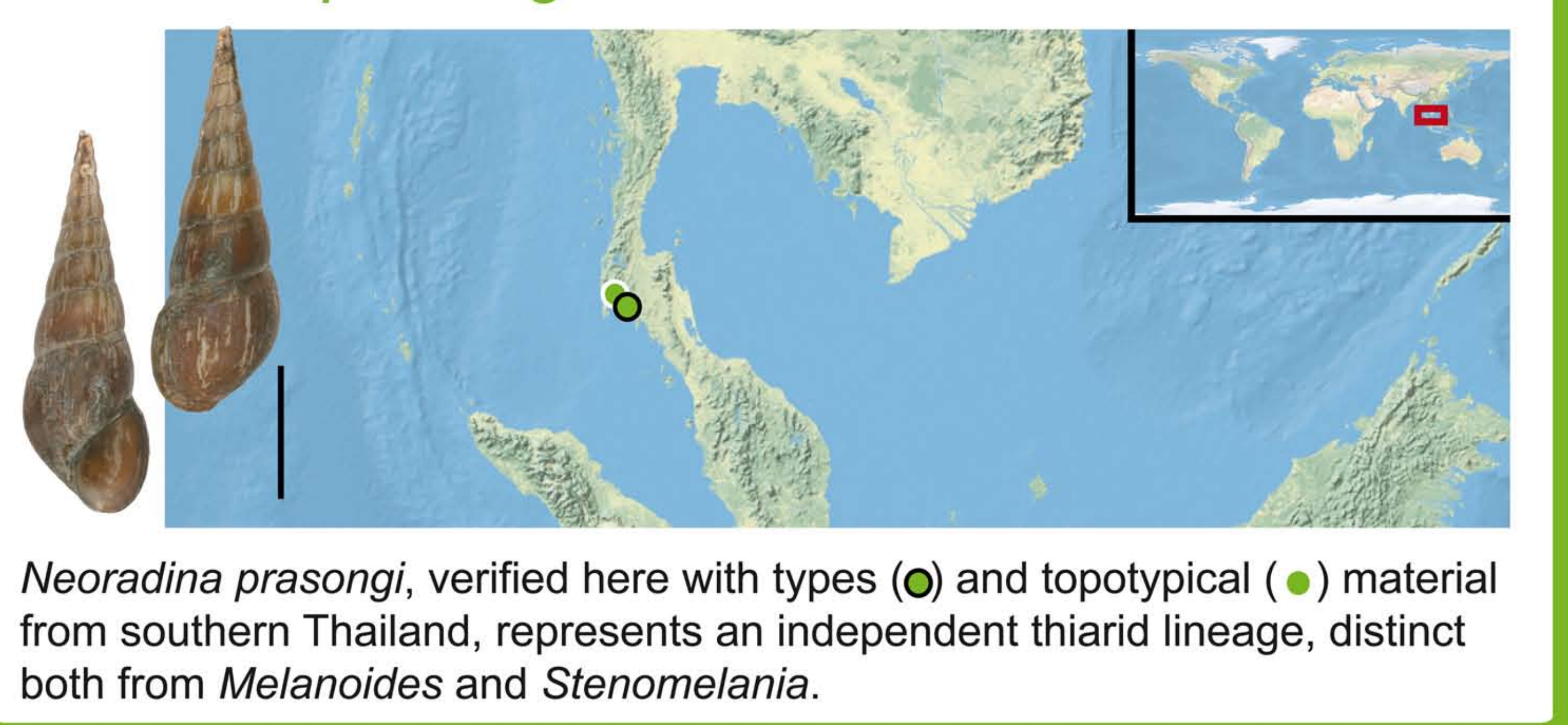
- Two short fragments (~100bp/~200bp) of the 16S gene were amplified via PCR using primer pairs established for Thiariidae.

Simulathena papuensis



▲ Maximum Likelihood Tree of thiarid and paludomid Cerithioidea, based on 16S mtDNA. Results of Bootstrap analysis are given (>50%). Colors indicate historic DNA samples of the respective case studies.

Neoradina prasongi



Paludomus sp.

