



***Tisaniba*, a new genus of marpissoid jumping spiders from Borneo (Araneae: Salticidae)**

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Abstract

Six new species of marpissoid jumping spiders from Sarawak, Borneo, are described in the new genus *Tisaniba* Zhang & Maddison. They are the type species, *T. mulu* Zhang & Maddison **sp. nov.**, as well as the species *T. bijibijan* Zhang & Maddison **sp. nov.**, *T. dik* Zhang & Maddison **sp. nov.**, *T. kubah* Zhang & Maddison **sp. nov.**, *T. selan* Zhang & Maddison **sp. nov.**, and *T. selasi* Zhang & Maddison **sp. nov.** The spiders are small and brown to black, living in leaf litter in the tropical forest. Phylogenetic analyses based on 28s and 16sND1 genes indicate that they are a distinctive group within the marpissoids. Diagnostic illustrations and photographs of living spiders are provided for all species.

Key words: Araneae, Salticidae, Marpissoida, new genus, new species, jumping spider, molecular phylogeny, Borneo

Introduction

The jumping spider clade Marpissoida (Maddison & Hedin, 2003; Bodner & Maddison, 2012) is known to include two major subclades, the smaller Ballinae of the Old World (Benjamin, 2004), and a large, primarily New World, clade including the Dendryphantinae, Marpissinae, and some smaller groups (e.g., synagelines, *Attidops* Banks, *Itata* Peckham & Peckham; see Bodner & Maddison 2012). Among the several hundred marpissoid species, most are tree trunk or foliage dwellers (Maddison, unpublished), with only a handful of genera having ground dwelling species (e.g. *Leikung* Benjamin, *Marpissa* C. L. Koch, *Synageles* Simon, *Terralonus* Maddison). We here report the discovery of a third major subclade, a group of small leaf-litter dwelling salticids from Borneo, the new genus *Tisaniba*.

Material and methods

Photographs of living specimens were taken with a Pentax Optio 33WR digital camera. For the macro capability, a small lens was glued to it. Photographs of preserved specimens were taken under a Leica MZ16 dissecting microscope with Leica Application Suite version 3.1.0. Preserved specimens were examined under both dissecting microscopes and a compound microscope with reflected light. Drawings were made with a drawing tube on a Nikon ME600L compound microscope.

Terminology is standard for Araneae. All measurements are given in millimeters. Descriptions of color pattern are based on the alcohol-preserved specimens. Carapace length was measured from the base of the anterior median eyes not including the lenses to the rear margin of the carapace medially; abdomen length to the end of the anal tubercle. Specimens are deposited in the Spencer Entomological Collection at the Beaty Biodiversity Museum, University of British Columbia (UBC-SEM). Specimen identifiers are given in two ways, one (UBC SEM ARxxxxx) is the museum's number, the other (SWK12-xxx) is a field number derived from photographs.

Phylogenetic trees inferred are shown in Fig. 73 for the broad sample dataset and Fig. 74 for the marpissoid dataset. The broad phylogeny, using only two gene regions, has a few clear deep misplacements (e.g. *Cesonia*, *Cheiracanthium*), but otherwise is in concordance with previous results with more genes and taxa (e.g., Bodner & Maddison, 2012).

Tisaniba is placed within the Marpissoida, as sister group to the Ballinae in all 20 of the broad-sample search replicates (Fig. 73). With the more restricted marpissoid data set, *Tisaniba* continued to fall within the marpissoida, but the bootstrap analysis was unable to place *Tisaniba* clearly with either the Ballinae or the other marpissoids. This and the pattern of branch lengths (Fig. 73) suggests that *Tisaniba* branches deep within the Marpissoida, and can be considered a third major group of Marpissoida, alongside the Ballinae and the large primarily New World clade including the Marpissinae and Dendryphantinae.

This placement of *Tisaniba* is in accord with morphology. The embolic spiral and its orientation, as well as the furrow across the tegulum, make the male palp closely resemble ballines and some other marpissoids such as *Attidops*. Indeed, with *Attidops*, ballines and *Tisaniba* all branching deep in the marpissoids, their common form of palp could be credibly argued as the ancestral form for marpissoids.

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References

- Benjamin, S.P. (2004) Taxonomic revision and phylogenetic hypothesis for the jumping spider subfamily Ballinae (Araneae, Salticidae). *Zoological Journal of the Linnean Society*, 142, 1–82.
<http://dx.doi.org/10.1111/j.1096-3642.2004.00123.x>
- Bodner, M.R. & Maddison, W.P. (2012) The biogeography and age of salticid spider radiations (Araneae: Salticidae). *Molecular Phylogenetics and Evolution*, 65, 213–240.
<http://dx.doi.org/10.1016/j.ympev.2012.06.005>
- Guindon, S. & Gascuel, O. (2003) A simple, fast and accurate method to estimate large phylogenies by maximum-likelihood. *Systematic Biology*, 52, 696–704.
<http://dx.doi.org/10.1080/10635150390235520>
- Katoh, K., Kuma, K., Toh, H. & Miyata, T. (2005) MAFFT version 5: improvement in accuracy of multiple sequence alignment. *Nucleic Acids Research*, 33, 511–518.
<http://dx.doi.org/10.1093/nar/gki198>
- Katoh, K., Misawa, K., Kuma, K. & Miyata, T. (2002) MAFFT: a novel method for rapid multiple sequence alignment based on fast Fourier transform. *Nucleic Acids Research*, 30, 3059–3066.
<http://dx.doi.org/10.1093/nar/gkf436>
- Logunov, D.V. & Azarkina, G.N. (2008) Two new genera and species of Euophryinae (Aranei: Salticidae) from SE Asia. *Arthropoda Selecta*, 17, 111–115.
- Maddison, W.P. & Hedin, M.C. (2003) Jumping spider phylogeny (Araneae: Salticidae). *Invertebrate Systematics*, 17, 529–549.
<http://dx.doi.org/10.1071/is02044>
- Platnick, N.I. (2012) The world spider catalog, version 12.5. American Museum of Natural History. Available from: <http://research.amnh.org/iz/spiders/catalog> (accessed 2 July 2014)
- Posada, D. (2008) jModelTest: Phylogenetic model averaging. *Molecular Biology and Evolution*, 25, 1253–1256.
<http://dx.doi.org/10.1093/molbev/msn083>
- Zhang, J.X. & Maddison, W.P. (2013) Molecular phylogeny, divergence times and biogeography of spiders of the subfamily Euophryinae (Araneae: Salticidae). *Molecular Phylogenetics and Evolution*, 68, 81–92.
<http://dx.doi.org/10.1016/j.ympev.2013.03.017>
- Zwickl, D.J. (2006) *Genetic Algorithm Approaches for the Phylogenetic Analysis of Large Biological Sequence Datasets under the Maximum Likelihood Criterion*. PhD Dissertation. The University of Texas at Austin. [total page unknown]