Phylogenomic reclassification of the world's most venomous spiders (Mygalomorphae, Atracinae), with implications for venom evolution

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Supplemental Information

Supplemental Text. Summary of prior hexathelid phylogeny and classification studies.

Although our overall results contrast with earlier morphology-only studies of mygalomorph relationships, several points of agreement can be found with more recent molecular studies. Raven [12, 27] re-limited the family Hexathelidae, and identified one morphological synapomorphy (possession of numerous labial cuspules). Raven [12] discussed at length the uncomfortable placement of Atrax (then the only described atracine), including hypothesized morphological homoplasy with more derived mygalomorph taxa. Goloboff [22] conducted a formal cladistic analysis of morphology, and recovered a paraphyletic Hexathelidae as an early-diverging Avicularioidea lineage, but noted that near parsimonious trees included hexathelids as monophyletic. Using ribosomal DNA sequences, Hedin & Bond [23] found a paraphyletic Hexathelidae, with taxa intermixed with non-diplurine diplurids at the base of Avicularioidea. Sampled atracines were not recovered with the single sampled actinopodid. Using elongation factor-1 gamma sequences, Ayoub et al. [17] recovered a Hadronyche plus Missulena clade, separate from the hexatheline Bymainiella. Using multiple genes alone, or molecules plus morphology, Bond et al. [13] recovered Atrax + Hadronyche with Missulena + Actinopus (F Actinopodidae). Again, hexathelids were not monophyletic, with two hexathelines intermixed with early-diverging non-diplurine diplurids. Opatova and Arnedo [18] used the Bond et al. [13] multigenic matrixes, adding data for multiple Macrothele specimens. These authors found four separate hexathelid lineages (i.e., hexathelid nonmonophyly), a monophyletic *Macrothele*, and an atracine plus actinopodid sister relationship. Using a 6-gene dataset, Wheeler et al. [20] recovered a core hexathelid lineage (including the important South American taxa Mediothele and Scotinoecus as sister taxa), separate from a well-supported atracine plus actinopodid clade. Finally, Hamilton et al. [19] used anchored hybrid enrichment (AHE) sequence capture data to show hexathelid non-monophyly, and an atracinae plus actinopodid relationship (see FIG 1C).

Supplemental Table 1. Excel file with voucher, identification and UCE information.

Supplemental Figure 1. ASTRAL phylogenies.

