

# Largest-ever study of spider genetics shows orb weaver spiders do not share common origins

17 July 2014



Close-up of the cephalothorax on *Eriophora* sp. (possibly *E. heroine* or *E. pustulosa*. Credit: JJ Harrison/Wikipedia/CC BY-SA 3.0

For decades, the story of spider evolution went like this: As insects became more and more diverse, with some species taking to the skies, spiders evolved new hunting strategies, including the ability to weave orb-shaped webs to trap their prey.

From that single origin, the story goes, orb-weaver [spiders](#) diverged along different evolutionary paths, leading to today, where several species weave similar – though not identical – webs.

It's a good story, but there's just one problem – Harvard scientists now know it's not true.

The largest-ever phylogenetic study of spiders, conducted by postdoctoral student Rosa Fernández, Gonzalo Giribet, Alexander Agassiz Professor of Zoology, and Gustavo Hormiga, a professor at George Washington University, shows

that, contrary to long-held popular opinion, the two groups of spiders that weave orb-shaped webs do not share a single origin. The study is described in a July 17 paper published in *Current Biology*.

"This study examines two different groups of orb-weaver spiders, as well as several other species," Giribet said. "Using thousands of genes, we did a comparative phylogenetic analysis, and what we now know is there is not a single origin for the orb-weaver spiders.

"There are two possible explanations for this," he continued. "One is that the orb web evolved far back in the lineage of the two groups, but has been lost in some groups. The other option is that the orb web evolved independently in these two groups. We still haven't resolved that question yet – we need to sample many more of these intermediate groups before we can say which option is correct."

The belief that orb-weaver spiders shared a common origin, Giribet said, came largely from earlier morphological studies.

Even as new genetic tools became more commonplace in the last two decades, the single origin theory held sway, in part, because early phylogenetic studies relied on just a handful of genes to draw a picture of the spider evolutionary tree.

"Some early analyses pointed out that spiders with orb webs didn't form a group – they appeared in different places along the tree," Giribet said. "But the genes that were being used weren't enough to elucidate the evolution of a very diverse group like spiders, so most people dismissed many of those results."

In recent years, however, sequencing technology

has dropped dramatically in cost, meaning researchers who once were able to study only a handful of genes can now examine the entire genome of a particular organism.

"The technology has changed what we are able to do in terms of the questions we can ask and the questions we can answer," Giribet continued. "Even just five years ago, we were spending thousands of dollars to sequence 3,000 genes. Today, we're spending just a few hundreds of dollars to sequence millions, which is almost an entire genome.

In the case of Giribet and Fernández, the technology allowed them to sequence genes from 14 different spiders, creating the largest genomic data set for the study of spiders.

"This paper is at the forefront of how these large data sets are being analyzed, and how we are now constructing phylogenies using molecular data," Giribet said. "We can now test all possible pitfalls of phylogenetic interference to make sure our results are as accurate as possible."

Though his hunch, Giribet said, is that the two orb-weaver groups evolved independently, he's now designing a study that will examine the genetics of as many as 150 spider species to test that hypothesis.

While the two groups create very similar webs, he said, their strategies for using them, how they manufacture silk, and even the silk used to construct them are very different. While one group passively uses their web to catch prey, some species in the other use it like a net, holding it between their legs, and trapping insects with it.

"Their behaviors are very different, the types of silk are very different – it's my bet that these would be two separate evolutions of a similar structure of web," Giribet said. "But we need to further test it, and we are now looking to expand this study to include as many as 150 different spiders."

Provided by Harvard University

APA citation: Largest-ever study of spider genetics shows orb weaver spiders do not share common

origins (2014, July 17) retrieved 25 September 2021 from <https://phys.org/news/2014-07-largest-ever-spider-genetics-orb-weaver.html>

*This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.*