

# Scientists solve reptile mysteries with landmark study on the evolution of turtles

November 24 2014

---



Turtles -- like this alligator snapping turtle *Macrochelys temminckii* -- are a diverse group of animals with a hotly contested evolutionary history. Researchers from the California Academy of Sciences used a next-generation genetic sequencing technique to reconstruct a robust turtle 'tree of life' and fill-in existing knowledge gaps. Credit: The California Academy of Sciences

A team of scientists, including researchers from the California Academy of Sciences, has reconstructed a detailed "tree of life" for turtles. The specifics of how turtles are related—to one another, to other reptiles,

and even to dinosaurs—have been hotly debated for decades. Next generation sequencing technologies in Academy labs have generated unprecedented amounts of genetic information for a thrilling new look at turtles' evolutionary history. These high-tech lab methods revolutionize the way scientists explore species origins and evolutionary relationships, and provide a strong foundation for future looks into Earth's fossil record.

Research results, appearing in *Molecular Phylogenetics and Evolution*, describe how a new genetic sequencing technique called Ultra Conserved Elements (UCE) reveal [turtles](#)' closest relatives across the animal kingdom. The new genetic tree uses an enormous amount of data to refute the notion that turtles are most closely related to lizards and snakes. Instead, authors place turtles in the newly named group "*Archelosauria*" with their closest relatives: birds, crocodiles, and dinosaurs. Scientists suspect the new group will be the largest group of vertebrates to ever receive a new scientific name.

The UCE technique used in high-tech labs allowed scientists to move beyond years of speculation and place the *Archelosauria* group in its rightful place on the reptile tree of life. UCE has been available since 2012, yet scientists are just beginning to tap its potential for generating enormous amounts of genetic data across vertebrates.

"Calling this is an exciting new era of sequencing technology is an understatement," says Brian Simison, PhD, Director of the Academy's Center for Comparative Genomics (CCG) that analyzed the study's massive amount of data. The CCG is a state-of-the-art facility comprised of a sequencing lab, frozen DNA collection, and computing resources that serves as the Academy's core genetic center. Established in the summer of 2008, the CCG continues to refine Academy research—including new turtle findings—on a global, evolutionary scale.

"In the space of just five years, reasonably affordable studies using DNA sequencing have advanced from using only a handful of genetic markers to more than 2,000—an unbelievable amount of DNA," adds Simison. "New techniques like UCE dramatically improve our ability to help resolve decades-long evolutionary mysteries, giving us a clear picture of how animals like turtles evolved on our constantly-changing planet."



The new study uses an enormous amount of data to refute the notion that turtles are most closely related to lizards and snakes. Instead, authors place turtles in the newly named group "*Archelosauria*" with their closest relatives: birds, crocodiles, and dinosaurs. Scientists suspect the new group will be the largest group of vertebrates to ever receive a new scientific name. Credit: The California Academy of Sciences

Major findings also resolve an evolutionary mystery surrounding softshell turtles—a bizarre group of scale-less turtles with snorkel-like

snouts. Until now, studies linked softshell turtles with a smaller semi-aquatic group called mud turtles, despite the fact that softshells appear in the fossil record long before their mud-loving counterparts. The Academy's study places softshells in a league of their own on the evolutionary tree, quite far removed from any turtle relatives. Their long independent history helps explain their striking looks as well as their ancient presence in the fossil record.

Study coauthor James Parham, PhD—Academy Research Associate, Assistant Professor of Geological Sciences at Cal State Fullerton, and renowned turtle expert—says cutting-edge testing techniques bring a new level of clarity to more than two decades of his turtle research. With large amounts of data backing up each evolutionary branch on the turtle tree of life, scientists are able to compare their evolution not only across species, but also across each continent's corresponding fossil records.

"I have been working on the [evolutionary relationships](#) of turtles for over 20 years using a variety of methods," says Parham. "Fossils are essential for showing us what extinct turtles looked like, but also in letting us know when and where they lived in the past."

Parham notes that studying turtle fossils—particularly the physical features of their bones—hasn't always painted an accurate evolutionary picture of turtle relationships across continents and through time. "The turtle tree of life based on fossil turtle anatomy didn't match up with the timing of their appearance in the fossil record, as well as their geography," Parham says. "But the tree of life generated at the Academy's CCG is consistent with time and space patterns we've gathered from the [fossil record](#). These new testing techniques help reconcile the information from DNA and fossils, making us confident that we've found the right tree."

Provided by California Academy of Sciences

Citation: Scientists solve reptile mysteries with landmark study on the evolution of turtles (2014, November 24) retrieved 2 May 2024 from <https://phys.org/news/2014-11-scientists-reptile-mysteries-landmark-evolution.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.