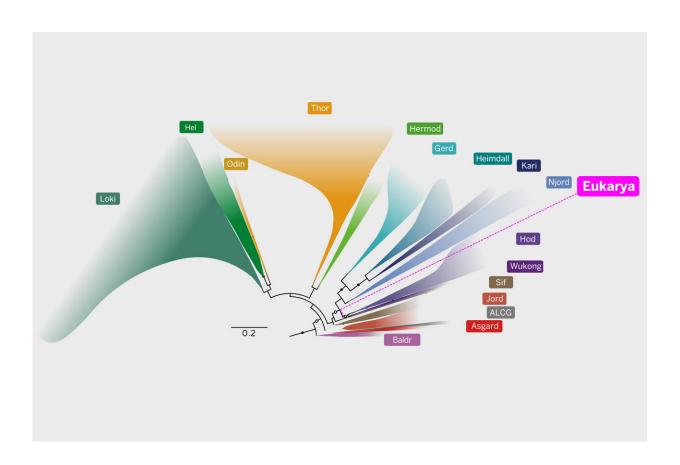


New clues about origin of complex life trace roots to common ancestor

June 14 2023, by Marc Airhart



According to this latest study, all complex life forms (a.k.a. eukaryotes) trace their roots back to a common ancestor among a group of microbes called the Asgard archaea. Credit: University of Texas at Austin

TThor, the legendary Norse god from the mythological city of Asgard, is



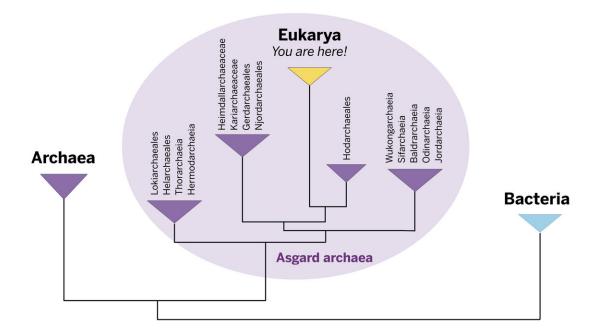
not alone. According to groundbreaking research published in the journal *Nature*, we humans—along with eagles, starfish, daisies and every complex organism on Earth—are, in a sense, Asgardians.

Analyzing the genomes of hundreds of different microbes called archaea, researchers at The University of Texas at Austin and other institutions have discovered that eukaryotes—complex life forms with nuclei in their cells, including all the world's plants, animals, insects and fungi—trace their roots to a common Asgard archaean ancestor.

That means eukaryotes are, in the parlance of evolutionary biologists, a "well-nested clade" within Asgard archaea, similar to how birds are one of several groups within a larger group called dinosaurs, sharing a common ancestor. The team has found that all eukaryotes share a common ancestor among the Asgards.

No fossils of eukaryotes have been found from farther back than about 2 billion years ago, suggesting that before that, only various types of microbes existed.





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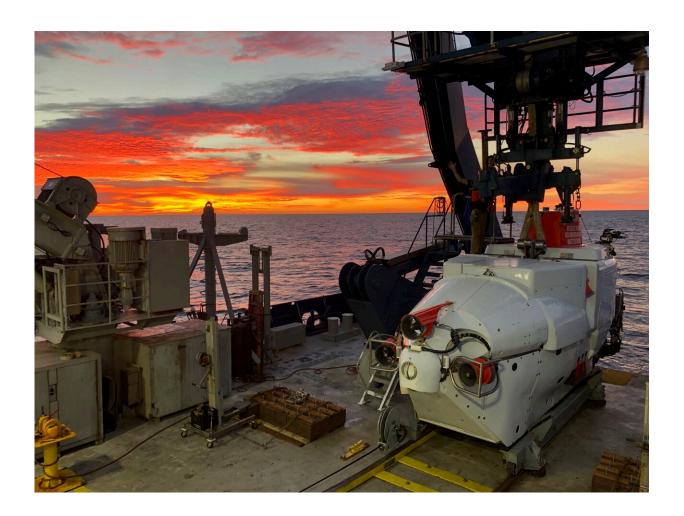
"So, what events led microbes to evolve into eukaryotes?" said Brett Baker, UT Austin associate professor of integrative biology and <u>marine science</u>. "That's a big question. Having this <u>common ancestor</u> is a big step in understanding that."

Led by Thijs Ettema of Wageningen University in the Netherlands, the research team identified the closest microbial relative to all complex life forms on the tree of life as a newly described order called the Hodarchaeales (or Hods for short). The Hods, found in marine sediments, are one of several subgroups within the larger group of Asgard archaea.

The Asgard archaea evolved more than 2 billion years ago, and their



descendants are still living. Some have been discovered in deep sea sediments and hot springs around the world, but so far only two strains have been successfully grown in the lab. To identify them, scientists collect their genetic material from the environment and then piece together their genomes. Based on genetic similarities with other organisms that can be grown in the lab and studied, the scientists can infer metabolism and other features of the Asgards.



Some of the microbes analyzed for this study were collected using the Alvin deep-sea submersible, seen here on a collection trip in the Guaymas Basin in November 2018. Credit: Brett Baker



"Imagine a <u>time machine</u>, not to explore the realms of dinosaurs or <u>ancient civilizations</u>, but to journey deep into the potential metabolic reactions that could have sparked the dawn of complex life," said Valerie De Anda, a researcher in Baker's lab. "Instead of fossils or <u>ancient artifacts</u>, we look at the genetic blueprints of modern microbes to reconstruct their past."

The researchers expanded the known Asgard genomic diversity, adding more than 50 undescribed Asgard genomes as input for their modeling. Their analysis indicates that the ancestor of all modern Asgards appears to have been living in hot environments, consuming CO₂ and chemicals to live. Meanwhile, Hods, which are more closely related to eukaryotes, are metabolically more similar to us, eating carbon and living in cooler environments.

"This is really exciting because we are looking for the first time at the molecular blueprints of the <u>ancestor</u> that gave rise to the first eukaryotic cells," De Anda said.

In Norse mythology, Hod (also spelled Höd, Höðr or Hoder) is a god, the blind son of Odin and Frigg, who is tricked into killing his own brother Baldr.

"I keep joking in my talks that 'We are all Asgardian," Baker said. "Now that's probably going to be on my tombstone."

More information: Laura Eme et al, Inference and reconstruction of the heimdallarchaeial ancestry of eukaryotes, *Nature* (2023). <u>DOI:</u> 10.1038/s41586-023-06186-2

Provided by University of Texas at Austin



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