

Evolutionary discovery to rewrite textbooks

June 12 2019



Choanocyte cross section. Credit: The University of Queensland



Scientists at The University of Queensland have upended biologists' century-old understanding of the evolutionary history of animals.

Using <u>new technology</u> to investigate how multi-celled animals developed, their findings revealed a surprising truth.

Professor Bernie Degnan said the results contradicted years of tradition.

"We've found that the first multicellular animals probably weren't like the modern-day sponge <u>cells</u>, but were more like a collection of convertible cells," Professor Degnan said.

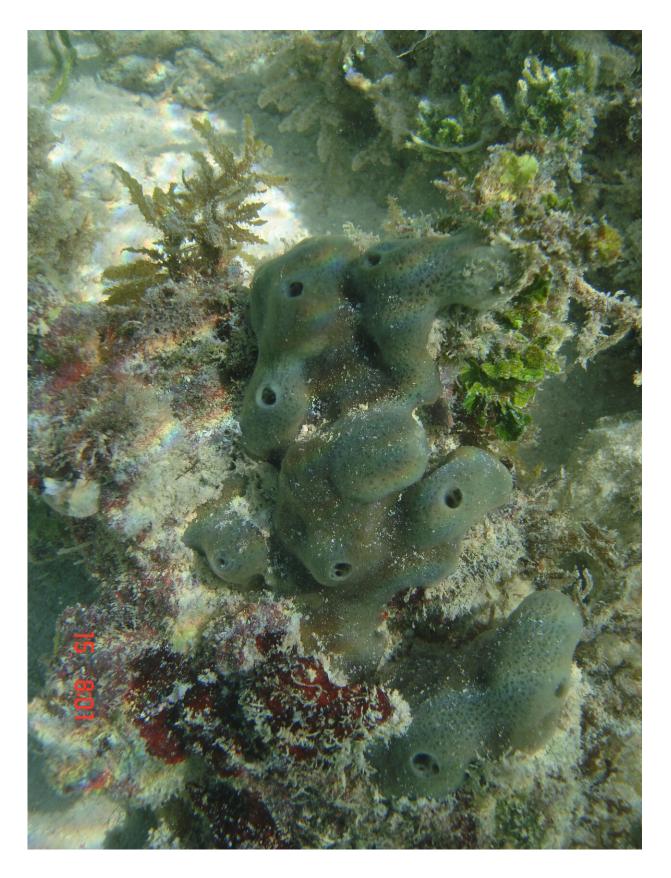
"The great-great-grandmother of all cells in the <u>animal kingdom</u>, so to speak, was probably quite similar to a stem cell.

"This is somewhat intuitive as, compared to plants and fungi, animals have many more cell types, used in very different ways—from neurons to muscles—and cell-flexibility has been critical to <u>animal evolution</u> from the start."

The findings disprove a long-standing idea: that multi-celled animals evolved from a single-celled ancestor resembling a modern sponge cell known as a choanocyte.

"Scattered throughout the history of evolution are major transitions, including the leap from a world of microscopic single-cells to a world of multi-celled animals," Professor Degnan said.







The adult sponge *Amphimedon queenslandica*. Credit: The University of Queensland

"With multicellularity came incredible complexity, creating the animal, plant, fungi and algae kingdoms we see today.

"These large organisms differ from the other more-than-99-per-cent of biodiversity that can only be seen under a microscope."

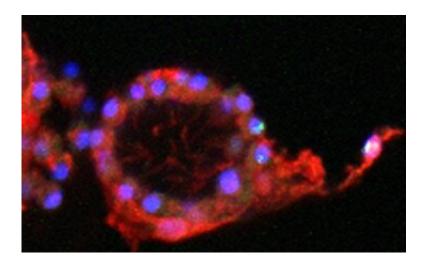
The team mapped individual cells, sequencing all of the genes expressed, allowing the researchers to compare similar types of cells over time.

Fellow senior author Associate Professor Sandie Degnan said this meant they could tease out the evolutionary history of individual <u>cell types</u>, by searching for the 'signatures' of each type.

"Biologists for decades believed the existing theory was a no-brainer, as sponge choanocytes look so much like single-celled choanoflagellates—the organism considered to be the closest living relatives of the animals," she said.

"But their transcriptome signatures simply don't match, meaning that these aren't the core building blocks of animal life that we originally thought they were.





Cells and flagella are labelled red and nuclei blue. Credit: The University of Queensland

"This technology has been used only for the last few years, but it's helped us finally address an age-old question, discovering something completely contrary to what anyone had ever proposed."

"We're taking a core theory of evolutionary biology and turning it on its head," she said.

"Now we have an opportunity to re-imagine the steps that gave rise to the first <u>animals</u>, the underlying rules that turned single cells into multicellular animal life."

Professor Degnan said he hoped the revelation would help us understand our own condition and our understanding of our own <u>stem cells</u> and cancer.

The study has been published in *Nature*.

More information: Pluripotency and the origin of animal



multicellularity, *Nature* (2019). <u>DOI: 10.1038/s41586-019-1290-4</u>, <u>www.nature.com/articles/s41586-019-1290-4</u>

Provided by University of Queensland

Citation: Evolutionary discovery to rewrite textbooks (2019, June 12) retrieved 25 April 2024 from https://phys.org/news/2019-06-evolutionary-discovery-rewrite-textbooks.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.