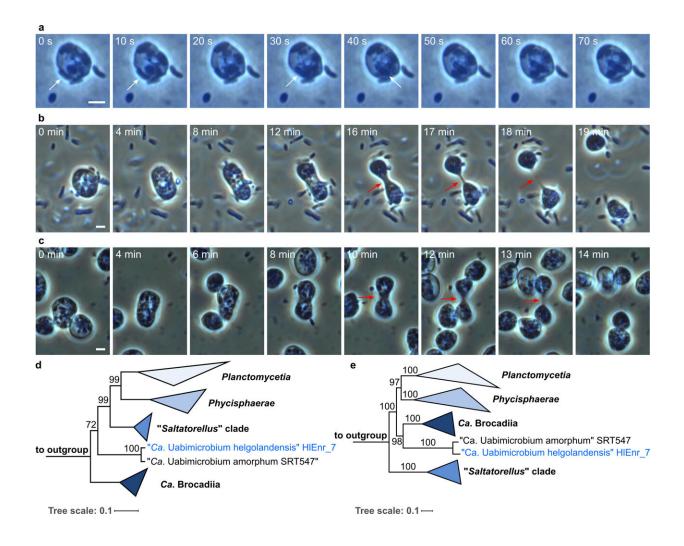


Research team discovers 'impossible' unicellular organism

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Overview of the cell biology and phylogeny of "Ca. U. amorphum" SRT547 as well as the novel isolate "Ca. U. helgolandensis" HIEnr_7. Phagocytosis-like uptake of surrounding prey bacteria by "Ca. U. amorphum" (large cell) (a); white arrows indicate the prey bacterium being internalized. Credit: *mBio* (2024). DOI: 10.1128/mbio.02044-24



The origin of eukaryotes is considered one of the greatest enigmas in biology: according to current doctrine, two prokaryotes, a so-called Asgard archaeon and a bacterium, are believed to have merged. The bacterium is said to have developed into a mitochondrion. Thanks to its mitochondrion, this eukaryotic ancestor had enough energy available to develop into the more complex cells known today.

A defining feature of such complex <u>eukaryotes</u> is endocytosis—the ability to eat other cells. Until now, however, it was deemed energetically impossible for a prokaryotic cell to be capable of endocytosis, as the energy source of the mitochondrion was considered indispensable for this.

However, a research team at the University of Jena has discovered prokaryotic bacteria that can eat other cells. The team <u>reports</u> on this surprising discovery in the scientific journal *mBio*.

Planctomycetes: 'Impossible' prokaryotes

For more than 10 years, Prof. Dr. Christian Jogler and his team have been pursuing a different approach to explaining eukaryogenesis. The researcher is focusing on the prokaryotic group of planctomycetes, very unusual bacteria that are considered by some scientists to be potential ancestors of eukaryotes due to their unusual cell biology.

"The idea of a 'fusion' of two different prokaryotes into one eukaryote seems incomprehensible to me from a cell biology perspective," explains the microbiologist, who conducts research in the Balance of the Microverse Cluster of Excellence at the University of Jena. "No one has ever observed anything like this before and such a hybrid would probably not be viable due to the different membrane structures and



molecular machineries," adds Jogler.

Microbe hunters

In 2014, Prof. Dr. Jogler's team found novel planctomycetes in Heiligendamm on the Baltic Sea, which provided arguments for the planctomycetal origin of eukaryotes. "These bacteria change their shape, they 'walk' over surfaces and flow around each other."

A team led by Takashi Shiratori from Japan also found and described comparable planctomycetes in 2019. These microorganisms even ate other bacteria and thus seemed to disprove the doctrine that prokaryotes could not be capable of endocytosis for energetic reasons.

"To be honest, I didn't believe Dr. Shiratori at first," says Jogler. He and his team therefore initially tried to disprove the hypothesis of prokaryotic endocytosis. However, after a year of intensive research, the researchers no longer have any doubts about the accuracy of Shiratori's results.

In the paper they have now published, they present a comparable organism that they have isolated and characterized from the North Sea: Uabimicrobium helgolandensis. These prokaryotes also feed on other bacteria and these organisms should therefore not even exist. The researchers have given this group of unusual planctomycetes the fitting name "bacteria of prey."

"By sequencing the genome of Uabimicrobium helgolandensis, we were also able to develop new hypotheses on the molecular mechanism of the absorption of prey <u>bacteria</u>," explains Jogler.

He sees the predatory planctomycetes as the microbial Archaeopteryx, a bridge organism between the <u>prokaryotes</u> and eukaryotes, and is



convinced that the planctomycetes played a role in eukaryogenesis, perhaps even in the origin of life itself.

More information: Carmen E. Wurzbacher et al, "Candidatus Uabimicrobium helgolandensis"—a planctomycetal bacterium with phagocytosis-like prey cell engulfment, surface-dependent motility, and cell division, *mBio* (2024). DOI: 10.1128/mbio.02044-24

Provided by Friedrich Schiller University of Jena

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