

Study sheds light on the genetic origins of the two sexes

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Female Eudorina colony with 32 large egg cells, some of which are pairing with small male sperm cells. On the upper right is a male sperm packet which has found the female colony and is about to release its sperm inside. Credit: Hiroko Kawai-Toyooka, University of Tokyo.



A new study published in the journal *Communications Biology* has shed light on the earliest stages in the evolution of male-female differentiation and sex chromosomes—and found the genetic origins of the two sexes to be unexpectedly modest.

James Umen, Ph.D., member, Enterprise Rent-a-Car Institute for Renewable Fuels and Joseph Varner Distinguished Investigator at the Donald Danforth Plant Science Center was part of a research team led by Dr. Hisayoshi Nozaki at the University of Tokyo who have been investigating the evolution of male and female sexes in a group of freshwater photosynthetic protists called volvocine green algae, a group that is well-known to scientists for capturing early stages in the evolution of sexes and multicellularity. Previous studies in animals and plants identified a general trend of expansion and differentiation between male and female sex chromosomes, often leading to large genetic differences between them; but these studies could not capture the earliest stages of evolution where distinct sperm and egg cell types first evolved from a simpler ancestral mating system with equal-sized gametes, known as isogamy.

The research team focused on two especially informative and closelyrelated multicellular volvocine species from the genera Yamagishiella and Eudorina which bracket the transition from isogamy to male/female sexes. While 32-celled Yamagishiella and Eudorina colonies look very similar to each other, the former is isogamous while the latter produces small male gametes and large female gametes. The team used highthroughput genome sequencing of the chromosomal regions that specify mating type in Yamagishiella and male-female differentiation in Eudorina, and then compared these regions.

While evolutionary theory predicted an expansion and/or increased genetic complexity of the sex determining <u>region</u> associated with the <u>evolution</u> of sexes in Eudorina, the results of the study showed the



opposite, with Eudorina having the most diminutive and genetically least complex sex-determining region found to date found among all volvocine species. In essence, the major difference between males and females in Eudorina could be reduced to the presence or absence of a single gene called MID that resides in a tiny chromosomal region.

"This new study punches a hole in the idea that increased genetic complexity of sex chromosomes accompanied the origin of sexes," said Umen. "Moreover, the work also has practical implications since it expands our understanding of how to identify mating types and sexes in new species of algae that we might want to breed as crops for improved traits relating to biofuel or biotechnology applications."

More information: Takashi Hamaji et al, Anisogamy evolved with a reduced sex-determining region in volvocine green algae, *Communications Biology* (2018). DOI: 10.1038/s42003-018-0019-5

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