

New species may arise from rapid mitochondrial evolution

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Credit: Oregon State University

Genetic research at Oregon State University has shed new light on how isolated populations of the same species evolve toward reproductive incompatibility and thus become separate species.

Scientists sequenced the entire genome of a Pacific tidepool crustacean, *Tigriopus californicus*, a model [species](#) for differentiation based on geographic separation—an early stage of one species becoming multiple species.

They examined the co-evolution of mitochondrial and [nuclear genes](#). Mitochondria act as a cell's power plant, generating adenosine triphosphate, or ATP—a source of chemical energy.

As in all animals, most of a *T. californicus* cell's [genes](#) are in its nucleus but some are in the mitochondria.

"The mitochondria organelle contains a small chromosome with only 37 genes, but these genes are absolutely essential for metabolism," said the study's corresponding author, Felipe Barreto, assistant professor of integrative biology in OSU's College of Science. "In order for ATP to be produced properly in a cell, a few hundred other genes encoded in the nucleus must interact directly with the 37 [mitochondrial genes](#). Mutations in the mitochondrial genes may cause these interactions to be subpar and thus cause reductions in metabolic performance."

T. californicus populations along the Pacific coast of North America have mitochondrial genes that differ widely from one [population](#) to the next—there are lots of mutations relative to each other.

"As a result, hybrid offspring between populations suffer from lowered fitness in the form of lower fecundity, slow development and lower ATP production as determined by several previous experiments," Barreto said.

Barreto and collaborators from the University of California, San Diego, the University of Southern California and the University of North Carolina used molecular statistical models to screen the genomes of eight populations in order to detect which genes might be incompatible between populations.

"Those genes may therefore be [candidate genes](#) for understanding how different populations become incompatible and possibly eventually

become different species," he said.

More information: Felipe S. Barreto et al, Genomic signatures of mitonuclear coevolution across populations of *Tigriopus californicus*, *Nature Ecology & Evolution* (2018). [DOI: 10.1038/s41559-018-0588-1](https://doi.org/10.1038/s41559-018-0588-1)

Provided by Oregon State University

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