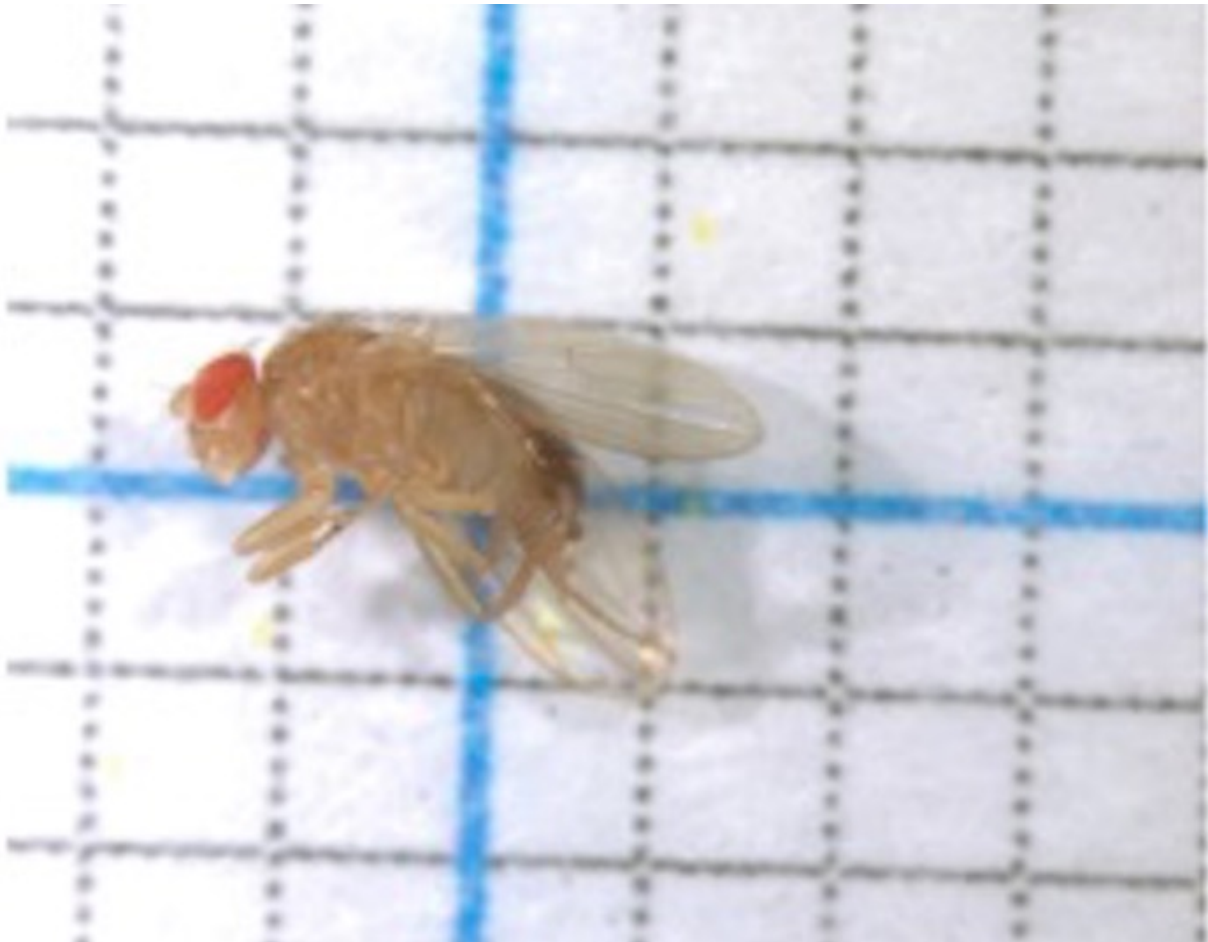


# Scientists engineer animals with ancient genes to test causes of evolution

January 13 2017

---



A transgenic fruit fly engineered to carry the alcohol dehydrogenase gene as it existed about 4 million years ago. Thousands of these 'ancestralized' flies were bred and studied for their ability to metabolize alcohol and to survive on an alcohol-rich food source. Credit: Kathleen Gordon

Scientists at the University of Chicago have created the first genetically modified animals containing reconstructed ancient genes, which they used to test the evolutionary effects of genetic changes that happened in the deep past on the animals' biology and fitness.

The research, published early online in *Nature Ecology & Evolution* on Jan. 13, is a major step forward for efforts to study the genetic basis of adaptation and evolution. The specific findings, involving the fruit fly's ability to break down alcohol in rotting fruit, overturn a widely-held hypothesis about the molecular causes of one of evolutionary biology's classic cases of adaptation.

"One of the major goals of modern evolutionary biology is to identify the genes that caused species to adapt to new environments, but it's been hard to do that directly, because we've had no way to test the effects of ancient genes on animal biology," said Mo Siddiq, a graduate student in the Department of Ecology and Evolution at the University of Chicago, one of the study's lead scientists.

"We realized we could overcome this problem by combining two recently developed methods—statistical reconstruction of ancient gene sequences and engineering of transgenic animals," he said.

Until recently, most studies of molecular adaptation have analyzed gene sequences to identify "signatures of selection"—patterns suggesting that a gene changed so quickly during its evolution that selection is likely to have been the cause. The evidence from this approach is only circumstantial, however, because genes can evolve quickly for many reasons, such as chance, fluctuations in population size, or selection for functions unrelated to the environmental conditions to which the organism is thought to have adapted.

Siddiq and his advisor, Joe Thornton, PhD, professor of ecology and

evolution and human genetics at the University of Chicago, wanted to directly test the effects of a gene's evolution on adaptation. Thornton has pioneered methods for reconstructing ancestral genes—statistically determining their sequences from large databases of present-day sequences, then synthesizing them and experimentally studying their molecular properties in the laboratory. This strategy has yielded major insights into the mechanisms by which biochemical functions evolve.

Thornton and Siddiq reasoned that by combining ancestral gene reconstruction with techniques for engineering transgenic animals, they could study how genetic changes that occurred in the deep past affected whole organisms—their development, physiology, and even their fitness.

"This strategy of engineering 'ancestralized animals' could be applied to many evolutionary questions," Thornton said. "For the first test case, we chose a classic example of adaptation—how fruit flies evolved the ability to survive the high alcohol concentrations found in rotting fruit. We found that the accepted wisdom about the molecular causes of the flies' evolution is simply wrong."

The fruitfly *Drosophila melanogaster* is one of the most studied organisms in genetics and evolution. In the wild, *D. melanogaster* lives in alcohol-rich rotting fruit, tolerating far higher alcohol concentrations than its closest relatives, which live on other food sources. Twenty-five years ago at the University of Chicago, biologists Martin Kreitman and John McDonald invented a new statistical method for finding signatures of selection, which remains to this day one of the most widely used methods in molecular evolution. They demonstrated it on the alcohol dehydrogenase (*Adh*) gene—the gene for the enzyme that breaks down alcohol inside cells—from this group of flies. *Adh* had a strong signature of selection, and it was already known that *D. melanogaster* flies break down alcohol faster than their relatives. So, the idea that the *Adh* enzyme was the cause of the fruit fly's adaptation to ethanol became

the first accepted case of a specific gene that mediated adaptive evolution of a species.

Siddiq and Thornton realized that this hypothesis could be tested directly using the new technologies. Siddiq first inferred the sequences of ancient Adh genes from just before and just after *D. melanogaster* evolved its ethanol tolerance, some two to four million years ago. He synthesized these genes biochemically, expressed them, and used biochemical methods to measure their ability to break down alcohol in a test tube. The results were surprising: the [genetic changes](#) that occurred during the evolution of *D. melanogaster* had no detectable effect on the protein's function.

Working with collaborators David Loehlin at the University of Wisconsin and Kristi Montooth at the University of Nebraska, Siddiq then created and characterized transgenic flies containing the reconstructed ancestral forms of Adh. They bred thousands of these "ancestralized" flies, tested how quickly they could break down alcohol, and how well the larvae and adult flies survived when raised on food with high alcohol content. Surprisingly, the transgenic flies carrying the more recent Adh were no better at metabolizing alcohol than flies carrying the more ancient form of Adh. Even more strikingly, they were no better able to grow or survive on increasing alcohol concentrations. Thus, none of the predictions of the classic version of the story were fulfilled. There is no doubt that *D. melanogaster* did adapt to high-alcohol food sources during its evolution, but not because of changes in the Adh enzyme.

"The Adh story was accepted because the ecology, physiology, and the statistical signature of selection all pointed in the same direction. But three lines of circumstantial evidence don't make an airtight case," Thornton said. "That's why we wanted to test the hypothesis directly, now that we finally have the means to do so."

Siddiq and Thornton hope that the strategy of making ancestralized transgenics will become the gold standard in the field to decisively determine the historical changes in genes to their changes on organisms' biology and fitness.

For his part, Kreitman, who is still a professor of ecology and evolution at UChicago, has been supportive of the new research, helping advise Siddiq on the project and sharing his vast knowledge about molecular [evolution](#) and *Drosophila* genetics.

"From the beginning, Marty was excited about our experiments, and he was just as supportive when our results overturned well-known conclusions based on his past work," Siddiq said. "I think that's extremely inspiring."

**More information:** Mohammad A. Siddiq et al, Experimental test and refutation of a classic case of molecular adaptation in *Drosophila melanogaster*, *Nature Ecology & Evolution* (2017). [DOI: 10.1038/s41559-016-0025](#)

Provided by University of Chicago Medical Center

Citation: Scientists engineer animals with ancient genes to test causes of evolution (2017, January 13) retrieved 20 June 2024 from <https://phys.org/news/2017-01-scientists-animals-ancient-genes-evolution.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.
---