

Some prairie vole brains are better wired for sexual fidelity

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A vole female with her pups and no male are shown. Researchers at The University of Texas at Austin have found that natural selection drives some male prairie voles to be fully monogamous and others to seek more partners. The surprising contrasts in the animals' brains result from differences in their DNA. Credit: Aubrey Kelly/Cornell University



Researchers at The University of Texas at Austin have found that natural selection drives some male prairie voles to be fully monogamous and others to seek more partners. The surprising contrasts in the animals' brains result from differences in their DNA.

The study, published this week in *Science*, compares faithful males with their less devoted counterparts and finds genetic differences in their brains. Just as people can be introverted or extroverted, <u>prairie voles</u> can be more or less prone to sexual fidelity because of these genetic differences.

Natural selection, the researchers conclude, does not appear to favor one set of genetic characteristics over the other, but instead allows for the coexistence of the seemingly contrasting traits, creating evolutionary space for both monogamous and nonmonogamous voles.

"This brain variation isn't just there by chance. It isn't random," says Steven Phelps, associate professor of integrative biology and the lead investigator on the study. "It's actually something that selection has kept around for a very long time. When it comes to <u>social behavior</u>, maybe there isn't a normal brain."

Prairie voles are small Midwestern rodents known for their monogamy. Males and females form pair-bonds and raise their young together as a couple. However, this does not stop some males from wandering off to make time with other females.

These dalliances come with both costs and benefits. Unfaithful males sire more offspring outside of their own nests. But while they are away, their mates may have encounters of their own. The result is that the offspring raised by an unfaithful male may not be his own. In contrast, a male that stays at home ensures that all of the pups in the nest belong to him.









A female vole and her pups are with the male vole in the background. Researchers at The University of Texas at Austin have found that natural selection drives some male prairie voles to be fully monogamous and others to seek more partners. The surprising contrasts in the animals' brains result from differences in their DNA. Credit: Aubrey Kelly/Cornell University

Phelps and his colleagues looked at the genetics of voles that stray and the voles that stay. Since neither of the two mating strategies is inherently more successful, both types persist in the population. The researchers found that the two types of male vole have <u>genetic</u> <u>differences</u>. These differences were found in parts of the DNA that turn genes on and off in specific brain regions, in this case a brain region important for spatial memory. The researchers speculate that faithful males are better at keeping tabs on a mate or are less likely to return to a place where they've encountered conflict.

"All of animal neuroscience is kind of predicated on minimizing genetic variation," says Phelps. "When we did this simple thing of going out in nature and looking at the brain, we were shocked at what we found."





A pair of voles, one with an ear tag that is used as a unique identifier for the vole. Credit: Aubrey Kelly/Cornell University

The findings support the notion that nature is favoring both brain types, thus increasing the diversity of the voles.

"I wouldn't be surprised if there were many genes whose variants have been kept around by selection in a similar way," says Phelps. "We may find this to be a common pattern in social behavior, including <u>personality</u> <u>differences</u>, in lots of species."

More information: "Sexual fidelity trade-offs promote regulatory variation in the prairie vole brain," by M. Okhovat et al. <u>www.sciencemag.org/lookup/doi/ ... 1126/science.aac5791</u>



Provided by University of Texas at Austin

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