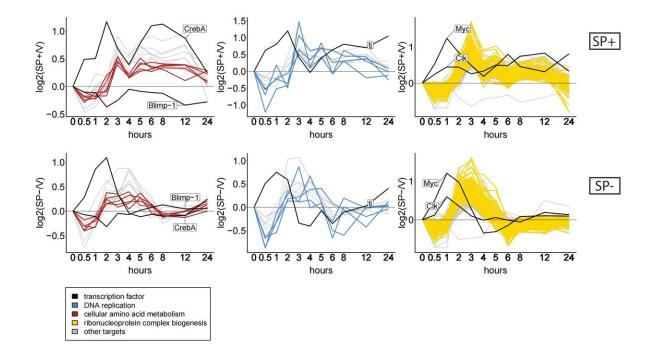


Mating causes 'jet lag' in female fruit flies, changing behavior





Within 4 hours after mating, expression changes are seen in features with roles in cell growth and protein biosynthesis. Log₂ fold changes of mated (SP+ or SP-) versus virgin (V) females for transcription factors and predicted targets, for features in group 1. Predicted targets are colored based on significant GO term enrichment (q-value Proceedings of the National Academy of Sciences (2023). DOI: 10.1073/pnas.2214883120

An innovative technique from Cornell University researchers finds



seminal fluid protein transferred from male to female fruit flies during mating changes the expression of genes related to the fly's circadian clock.

The finding, published in the *Proceedings of the National Academy of Sciences*, could help explain how this protein, called <u>sex peptide</u>, alters the female's behavior.

Post-mating, sex peptide has been shown to elicit increased egg-laying, aggression, activity and feeding, while reducing sleep and interest in mating in previously unmated females.

"Flies like to eat at certain times of day," said Mariana Wolfner, professor of molecular biology and genetics and one of the paper's senior authors. "They sleep at certain times, and the <u>circadian clock</u> machinery controls when flies are likely to do these things.

"What we're seeing," she said, "is that these very same behaviors—such as sleeping and eating—are changed after mating by the sex peptide. One way it might do that is by basically shifting the whole clock of the fly."

The surprising findings were made possible by examining transcriptomes—RNA sequencing that reveals <u>gene expression</u>, or which genes are turned on and off—at many different time points, providing high-resolution data that illuminates the order in which changes occur.

In the first four hours after mating, researchers found changes in expression of genes involved in the female fly's metabolism and the circadian clock. It is unknown what triggered these initial changes, but pheromones or seminal fluid proteins other than sex peptide are possible candidates. The initial effects were short term; they didn't last without sex peptide and seemed to prime the system.



In a second phase, four hours after <u>mating</u>, the researchers discovered that sex peptide caused changes to genes that regulate <u>circadian rhythms</u> and genes that are regulated by circadian clock pathways.

The study opens the door for future work that explores such questions as: how long the effect lasts and whether these effects occur in other organisms, given that circadian clock <u>genes</u> are highly conserved in many life forms.

More information: Sofie Y. N. Delbare et al, Time series transcriptome analysis implicates the circadian clock in the Drosophila melanogaster female's response to sex peptide, *Proceedings of the National Academy of Sciences* (2023). DOI: 10.1073/pnas.2214883120

Provided by Cornell University

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