

Mobile DNA element found in mosquito parasite has potential for infectious disease control

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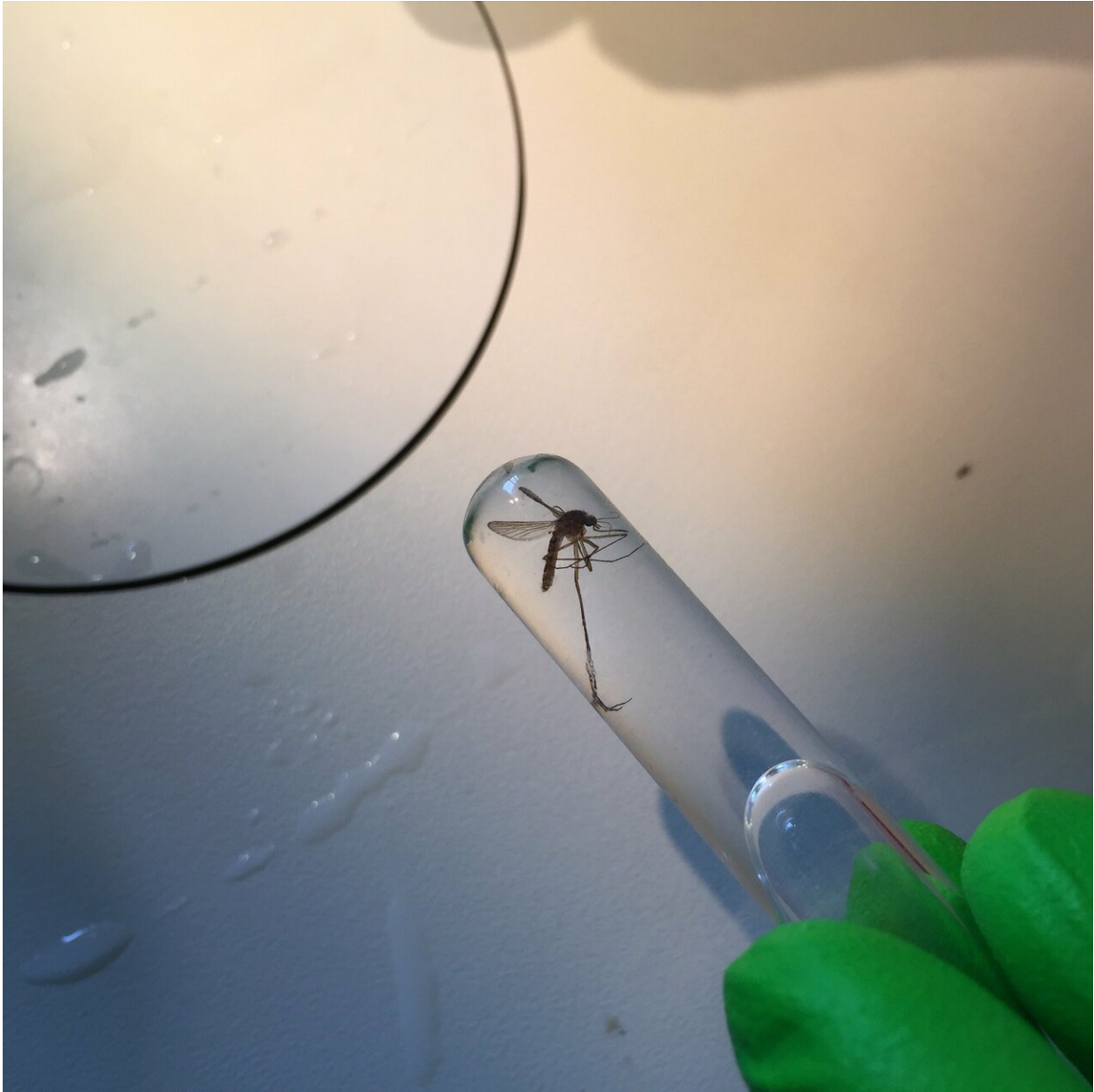
The Wolbachia bacterium (colorized purple). Credit: Dennis Kunkel, Sarah Bordenstein and Robert Brucker

Controlling mosquito-borne illnesses, such as Dengue or West Nile virus, has historically been difficult due to a lack of effective vaccines and concerns about the environmental impact of insecticides. Thus, scientists have turned to manipulating *Wolbachia*, a parasitic bacterium within mosquitoes, as a way to control the reproductive fitness of mosquito populations that transmit human disease.

In a new study in *Nature Communications*, an international team including scientists from the Marine Biological Laboratory (MBL) and the University of Chicago identified a new mobile DNA element in *Wolbachia*, which may contribute to improved control strategies for mosquito vectors of disease.

Led by former MBL scientists Julie Reveillaud of INRA, France, and Sarah Bordenstein of Vanderbilt University, the researchers reconstructed near-complete genomes of *Wolbachia* isolated from individual ovaries of four *Culex pipiens* mosquitoes. In the process, they identified a novel plasmid—a circular piece of DNA that can replicate independently of the chromosomes. Because a plasmid is a mobile DNA element, it can transfer from one cell to another and can have great implications for the fitness and evolution of a microbial species. Mobile genetic elements that can spread through different *Wolbachia* cells, and thus across a *Wolbachia* population, hold promise for controlling mosquito populations that may carry disease.

"Our data show that this novel plasmid is widespread across natural *Wolbachia* populations that infect *C. pipiens* mosquitoes throughout the world, which implies it has an essential role. The idea that it may enable transformation of the *Wolbachia* populations is simply very exciting," says the senior author of the study, A. Murat Eren (Meren), an assistant professor of medicine at the University of Chicago and an MBL Fellow.



Surface sterilizing a *Culex pipiens* mosquito sample before dissection. Credit: Julie Reveillaud

Wolbachia is transmitted from mother to offspring and can influence the reproductive behavior of its mosquito host. *Wolbachia* can modify sperm so if an infected male mates with an uninfected female, or with one that

carries a different strain of *Wolbachia*, embryos cannot develop. The mechanism behind this embryonic killing lies within another mobile genetic element in the *Wolbachia* genome.

"*Wolbachia* populations do not lend themselves to direct genetic modification. While a natural plasmid sounds very promising to bypass those limitations, we do not yet have any evidence to suggest the possibility or efficacy of any transformation through this plasmid," Meren says. "We just discovered its existence, and the time will tell how useful it is."

"Other exciting questions for which we do not have yet clear answers include whether there is an ancestral relationship between the plasmid and other mobile genetic elements of *Wolbachia*, and whether these interact or not," adds Reveillaud.

This collaboration began at MBL in 2011, when Meren and Reveillaud brought their computational and molecular evolution backgrounds together to explore questions in microbial ecology in the MBL's Bay Paul Center. (See Reveillaud's [blog post](#) about the collaboration.) Years later, they began examining *Wolbachia* genomic diversity by focusing on individual [mosquitoes](#) captured in the wild, rather than using laboratory mosquito strains. With the help of former MBL scientists Sarah Bordenstein and Seth Bordenstein of Vanderbilt University, who have expertise in *Wolbachia*, they began unraveling the intricacies of the *Wolbachia* genome.



Trap of *Culex pipiens* mosquitoes from France, provided by EID (Entente Interdépartementale Démoustication) in Montpellier, France. Credit: Julie Reveillaud

"*Wolbachia* is particularly difficult to study because, unlike free-living bacteria, it cannot be cultured in cell-free media. Through this

interdisciplinary collaboration, we were able to tease apart the genomes of the mosquito host, *Wolbachia*, and its mobile elements," says Sarah Bordenstein.

While fragments of this novel plasmid had appeared in previous *Wolbachia* sequencing studies, these fragments had never been assembled into a complete circular piece and its extrachromosomal nature was not recognized, most likely due to computational limitations. The team was able to overcome these limitations by employing a combination of cutting-edge strategies, such as genome-resolved metagenomics and long-read sequencing.

"The key bioinformatics platform that enabled us to study mosquito metagenomes and *Wolbachia* pangenomes in this study, and the connections between scientists with distinct skills that made this discovery possible, were developed at the MBL," Meren says. "Almost everyone in this paper is somehow linked to the MBL."

More information: Julie Reveillaud et al, The *Wolbachia* mobilome in *Culex pipiens* includes a putative plasmid, *Nature Communications* (2019). [DOI: 10.1038/s41467-019-08973-w](https://doi.org/10.1038/s41467-019-08973-w)

Provided by Marine Biological Laboratory

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