Insecticides play a central role in efforts to counter global impacts of mosquito- 
spread malaria and other diseases, which cause an estimated 750,000 deaths each 
year. These insect-specific chemicals, which cost more than $100 million to develop 
and bring to market, also are critical to controlling insect-driven damage on crops 
that poses a challenge to food security.

But in recent decades many insects have genetically adapted to become less 
sensitive to the potency of insecticides. In Africa, where long-lasting insecticide- 
treated bed nets and indoor spraying are major weapons in the fight against 
malaria, many species of mosquitoes across the continent have developed 
insecticide resistance that reduces the efficacy of these key interventions. In certain 
areas climate change is expected to exacerbate these problems.

University of California biologists have now developed a method that reverses 
insecticide resistance using CRISPR/Cas9 technology. A team including UC Santa 
Barbara researchers Craig Montell and Menglin Li, UC San Diego researchers 
Bhagyashree Kaduskar, Raja Kushwah and Professor Ethan Bier of UCSD’s Tata 
Institute for Genetics and Society (TIGS) used the genetic editing tool to replace an 
insecticide-resistant gene in fruit flies with the normal insecticide-susceptible form. 
Their achievement, described in Nature Communications, could significantly reduce 
the amount of insecticides used.
“This strategy could be used to reverse the resistance of mosquito disease vectors that spread devastating diseases that impact hundreds of millions of people each year,” said Craig Montell, a professor of molecular, cellular and developmental Biology at UC Santa Barbara.

“This technology also could be used to increase the proportion of a naturally occurring genetic variant in mosquitoes that renders them refractory to transmission or malarial parasites,” said Bier, a UCSD professor of cell and developmental biology and senior author of the paper.

The researchers used a modified type of gene-drive, a technology that employs CRISPR/Cas9 to cut genomes at targeted sites, to spread specific genes throughout a population. As one parent transmits genetic elements to their offspring, the Cas9 protein cuts the chromosome from the other parent at the corresponding site and the genetic information is copied into that location so that all offspring inherit the genetic trait. The new gene-drive includes an add-on that Bier and his colleagues previously engineered to bias the inheritance of simple genetic variants (also known as alleles) by also, at the same time, cutting an undesired genetic variant (e.g., insecticide resistant) and replacing it with the preferred variant (e.g., insecticide susceptible).

In the new study, the researchers used this “allelic drive” strategy to restore genetic susceptibility to insecticides, similar to insects in the wild prior to their having developed resistance. They focused on an insect protein known as the voltage-gated sodium channel (VGSC) which is a target for a widely used class of insecticides. Resistance to these insecticides, often called the knockdown resistance, or “kdr,” results from mutations in the vgsc gene that no longer permit the insecticide to bind to its VGSC protein target. The authors replaced a resistant kdr mutation with its normal natural counterpart that is susceptible to insecticides.

Starting with a population consisting of 83% kdr (resistant) alleles and 17% normal alleles (insecticide susceptible), the allelic drive system inverted that proportion to 13% resistant and 87% wild-type in 10 generations. Bier also notes that adaptions conferring insecticide resistance come with an evolutionary cost, making those insects less fit in a Darwinian sense. Thus, pairing the gene drive with the selective advantage of the more fit wild-type genetic variant results in a highly efficient and cooperative system, he says.
Treating fields with insecticides leads to the emergence of resistant pests and reduces the diversity of beneficial insects. Allelic-drives can help restore insecticide susceptibility and balance natural levels of insect populations.

**Photo Credit:** BIER LAB, UC SAN DIEGO

Similar allelic drive systems could be developed in other insects, including mosquitoes. This proof-of-principle adds a new method to pest- and vector-control toolboxes since it could be used in combination with other strategies to improve insecticide-based or parasite-reducing measures to drive down the spread of malaria.

“Through these allelic replacement strategies, it should be possible to achieve the same degree of pest control with far less application of insecticides,” said Bier. “It also should be possible to design self-eliminating versions of allelic drives that are programmed to act only transiently in a population to increase the relative frequency of a desired allele and then disappear. Such locally acting allelic drives could be reapplied as necessary to increase the abundance of a naturally occurring preferred trait with the ultimate endpoint being no GMO left in the environment.”

“An exciting possibility is to use allelic drives to introduce novel versions of the VGSC that are even more sensitive to insecticides than wild-type VGSCs,” suggested Montell. “This could potentially allow even lower levels of insecticides to be introduced into the environment to control pests and disease vectors.”

The study’s other authors are: Ankush Auradkar (UC San Diego), Annabel Guichard (UC San Diego and Tata Institute for Genetics and Society), Jared Bennett (UC Berkeley), Alison Henrique Ferreira Julio and John Marshall (UC Berkeley).

Funding for the research was provided by a gift from the Tata Trusts in India to TIGS-UC San Diego; National Institutes of Health (R01GM117321); a Paul G. Allen Frontiers Group Distinguished Investigator Award; the National Institute on Deafness and Other Communication Disorders (R01-DC007864); and the National Institute of Allergy and Infectious Diseases (AI165575).

*Competing interest note: Bier has equity interest in two companies he co-founded: Synbal Inc. and Agragene, Inc., which may potentially benefit from the research results. He also serves on Synbal’s board of directors and the scientific advisory board for both companies.*
About UC Santa Barbara

The University of California, Santa Barbara is a leading research institution that also provides a comprehensive liberal arts learning experience. Our academic community of faculty, students, and staff is characterized by a culture of interdisciplinary collaboration that is responsive to the needs of our multicultural and global society. All of this takes place within a living and learning environment like no other, as we draw inspiration from the beauty and resources of our extraordinary location at the edge of the Pacific Ocean.