

Male-killing bugs hold key to butterflies' curious color changes

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An African monarch butterfly. Credit: Claude Martin

It is a spectacular butterfly breed with an intriguing back story—now scientists have revealed how male-killing bacteria are helping to create a dazzling hybrid of the African monarch.

Researchers have shed new light on the complex reproductive process which dictates that only female offspring of the *Danaus chrysippus* survive—all in close proximity to Kenya's capital, Nairobi.

Previous research had shown that all female butterflies in this region are infected with a bacteria, called *Spiroplasma*, and after the females mate with immigrant males from other subspecies, the bacteria kills all their sons.

It was not clear, however, how this influenced the female butterflies' color patterns. The international study, led by the University of Edinburgh, analyzed the entire DNA of the bacteria and the female butterflies' [chromosomes](#).

The findings reveal that the bacteria ensures the survival of one particular color pattern gene, one that is always passed from mother to daughter. However, this same gene causes the daughter to resemble her father, rather than her mother.

Scientists say that by forcing its way into [daughters](#) only, the color gene genetically hitchhikes with the bacteria to evade death.

"This infection sets up a strange situation in this butterfly where populations in East Africa are largely female and they rely on immigrant males for their propagation," said Professor Richard ffrench-Constant, of the Centre for Ecology and Conservation on the University of Exeter's Penryn Campus in Cornwall.

"These males have no sons and all their daughters appear identical to them."

Despite 50 years of study, scientists have, until now, been puzzled why this female-only population displays unique and dazzling color

patterns—that change seasonally.

Normally, each subspecies has a recognizable color pattern that warns predators that they are toxic. New and unusual color patterns are thought to make them more vulnerable to predators.

Previous studies also found that the females' color chromosome had become attached to the female sex chromosome, called the W chromosome.

By sequencing the DNA, researchers proved that this new fused chromosome, called the neo-W, alters color patterns and spreads rapidly through the population, hand-in-hand with the male-killing [bacteria](#).

There was, however, one puzzle for the scientists still to solve: if the females all carried the same color gene, why was the population so variable?

The study found that these [genes](#) in the daughters allow themselves to be overridden by the genes coming from the fathers.

Seasonal fluctuations in wind patterns are thought to affect which subspecies of male immigrants end up in this region, leading to changes in the females' color patterns.

Even though they always resemble their father, the infected hybrid daughters, unable to produce sons, represent a genetic dead end for fathers, whose color pattern genes only survive for one generation before being wiped out.

Dr. Simon Martin, of the School of Biological Sciences at the University of Edinburgh, said: "The relatively fast emergence and spread of a new chromosome, combined with the short life cycle of the butterfly, allows

us to study how the microbe is altering the evolution of the butterfly, almost in real-time.

"We are continually discovering new ways in which microbes manipulate their hosts, and male-killing is just one example of this.

"It makes you wonder to what extent the evolution of other organisms—even humans—is affected by such unseen forces."

The study, published in the journal *PLOS Biology*, was carried out in partnership with the Universities of Exeter, Cambridge and the Mpala Research Centre in Kenya.

More information: Simon H. Martin et al. Whole-chromosome hitchhiking driven by a male-killing endosymbiont, (2019). [DOI: 10.1101/703421](https://doi.org/10.1101/703421) , [journals.plos.org/plosbiology/...journal.pbio.3000610](https://journals.plos.org/plosbiology/article?id=10.1371/journal.pbio.3000610)

Provided by University of Exeter

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