

Study reveals maintenance of male-related genes after loss of males in stick insects

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Traits are often lost during evolution, either because they are no longer beneficial or because they are too costly to maintain. When this happens, it is generally believed that the genes underlying the trait will eventually

degrade as well, making it difficult if not impossible for the trait to re-emerge. Yet there are numerous examples in nature of once-lost traits re-emerging in descendent lineages.

According to Giobbe Forni, a Research Fellow at the University of Bologna, "Mapping the presence and absence of traits onto a [species](#) tree suggests that some traits may have been lost in the lineages leading to extant species and then subsequently reinstated. Wings in stick insects are considered one of the more iconic instances of this evolutionary process."

This implies that the genes underlying these traits may be preserved, in some cases for millions of years. Unfortunately, research on the molecular basis of such reemergence is sparse, leaving the underlying mechanisms responsible for such preservation largely open to speculation until now.

In a [new study published](#) in *Genome Biology and Evolution*, Forni and his colleagues shed light on another complex trait that has been lost in some stick insects—the production of males.

Loss of the ability to produce males results in populations of only females, which reproduce by parthenogenesis, a form of asexual reproduction. The study reveals that genes that are highly connected in [regulatory networks](#) and involved in multiple biological processes may be maintained long after a trait is lost, providing a potential avenue for trait re-emergence over long evolutionary time scales.

In the new study, Forni and his co-authors Barbara Mantovani, Alexander S. Mikheyev, and Andrea Luchetti performed a comparative analysis of three species of stick insects in the genus *Bacillus*. While *Bacillus grandii maretimi* populations are composed of males and females that reproduce sexually, *Bacillus atticus* comprises populations

with only females that reproduce by parthenogenesis.

A third species, *Bacillus rossius*, includes both sexual and parthenogenetic populations. By studying the fates of genes involved in male reproduction in these three species, the authors sought to investigate the extent to which genes are preserved after trait loss and the potential mechanisms driving this preservation.

The researchers first identified gene networks whose expression was correlated with either male or female reproduction in the sexual species *B. marettimi* and then evaluated the same genes in *B. atticus* and *B. rossius*. Surprisingly, male-related genes exhibited no signs of weakened selection or accelerated evolution compared to female-related genes in the parthenogenetic species. Furthermore, male-related patterns of gene expression were partially preserved across both parthenogenetic species.

Delving deeper, the researchers found that genes in female-related networks were primarily expressed in female reproductive tissues, while those in male-related networks were expressed in male and female reproductive tissues, including both sexual and parthenogenetic females. This suggests that male-related genes may also play roles in female reproduction.

The involvement of a gene in multiple biological processes is known as pleiotropy, and this phenomenon may explain the preservation of male-related genes in these parthenogenetic stick insects, as previously hypothesized.

Moreover, the authors found that genes that were highly connected to many other genes in the network were more likely to be expressed in the reproductive tissues of parthenogens, suggesting that a gene's network connectivity may also influence its gene preservation after trait loss.

Taken together, these findings indicate "that the molecular ground plan of the once-lost male reproductive process may persist due to pleiotropic effects on other traits," explains Forni. "Different [genes](#) may undertake different trajectories of preservation and decay depending on the level of pleiotropy within the gene regulatory network."

This study not only sheds light on genetic architecture persistence after trait loss but also offers a potential glimpse into the emergence of rare males and cryptic sex (i.e., episodic generation of males and sexual reproduction), which have been observed in an increasing number of lineages that were thought to have lost the ability to produce males long ago. This opens up new potential avenues for research, with implications that may reach far beyond stick insects.

"Looking at how widespread genetic preservation after trait loss is on a larger scale remains fundamental. Although the *Bacillus* species complex offers a nice framework to address these issues, it would be useful to analyze a larger species complex where multiple transitions between reproductive strategies has occurred," notes Forni.

"While it is often necessary to rely on model species to discover and dissect [biological processes](#), it is even more important to test our hypotheses in a wider context. This will be possible only if we dedicate more effort to observing and analyzing the amazing diversity of organisms and their intricate adaptations."

More information: Giobbe Forni et al, Parthenogenetic Stick Insects Exhibit Signatures of Preservation in the Molecular Architecture of Male Reproduction, *Genome Biology and Evolution* (2024). [DOI: 10.1093/gbe/evae073](https://doi.org/10.1093/gbe/evae073)

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