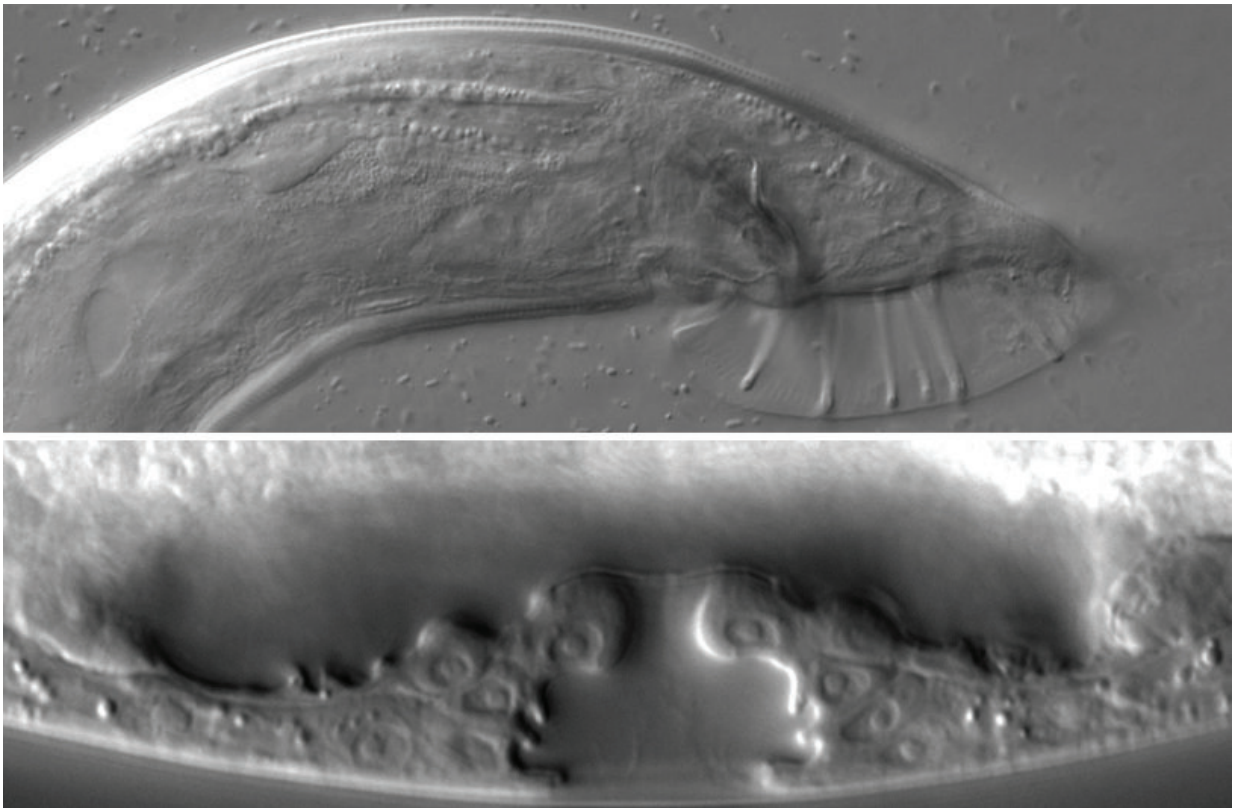


A key player in the maturation of sexual organs

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Top: the male tail, bottom: the developing vulva. Credit: Helge Grosshans, FMI

Puberty is a period of extensive changes of body morphology and function. Relatively little is known about what sets the whole process in motion. Thanks to studies in the tiny worm *C. elegans*, the group of Helge Großhans is getting closer to understanding how the onset of

puberty is genetically controlled. Recently, they uncovered a mechanism that initiates sexual organ maturation.

The *let-7* gene is considered a fundamental regulator of developmental timing in organisms as distinct as [worms](#) and mammals. It produces a small RNA, known as the *let-7* microRNA (miRNA), which can silence other genes. In the [model organism](#) *C. elegans*, *let-7* controls the transition from a juvenile to an adult animal. This transition, like human puberty, involves the formation of mature [sexual organs](#), in this case the vulval-uterine tract for the hermaphrodite and the tail in the male. (There are no female *C. elegans* worms—only hermaphrodites that can either self-fertilize or mate with the less abundant males.) The vital importance of *let-7* is highlighted by the fact that worms with a dysfunctional *let-7* die a dramatic death: they rupture through the vulva!

In a study published in *Life Science Alliance*, the group of Helge Großhans now presents new findings about how *let-7* controls worm puberty. Using precision genome engineering, the researchers specifically interfered with regulation of one or all targets of *let-7*. They found that the only relevant target of *let-7* for the formation of sexual organs is an mRNA called *lin-41*. This mRNA encodes an RNA binding protein that silences four other mRNAs. So basically, *let-7* inhibits *lin-41*, which therefore cannot inhibit the expression of the four genes anymore; those get expressed and promote the transition of the worm to adulthood.

"Although extensive studies of the *C. elegans* developmental timing pathway have helped to identify many of its players, we still don't understand well how these players function together. With this study, we are beginning to fill the gaps and to elucidate the [molecular mechanisms](#) underlying the transition from juvenile to adult—an important transition in the life of worms, and humans," says Großhans. "Intriguingly, timing defects in human puberty have been linked to genetic variations that

alter let-7 activity. It will therefore be interesting to study whether mammalian LIN41 also controls sexual organ maturation in mammals and possibly other puberty-related events."

Grosshans also notes that the study provides unexpected insight into another field, that of miRNA function. "miRNAs are often thought to act through a network activity in which they silence many targets modestly but coordinately. However, our work clearly demonstrates that the let-7 miRNA functions through only one target, lin-41. Although experiments comparable to what we have done here for let-7 are lacking for other miRNAs, there are other instances that seem to fit better to a model of only one or a few key miRNA targets. Hence, one might question whether the network activity model really describes the predominant mode of miRNA function in animals."

More information: Florian Aeschmann et al. let-7 coordinates the transition to adulthood through a single primary and four secondary targets. *Life Science Alliance*, April 2019 | Volume 2, No. 2 (published online March 25, 2019) [DOI: 10.26508/201900335](https://doi.org/10.26508/201900335) , [www.life-science-alliance.org/ ... content/2/2/e201900335](http://www.life-science-alliance.org/content/2/2/e201900335)

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