

How plants pass down genetic memories

August 28 2023



Arabidopsis thaliana is a plant species widely used to make fundamental biological discoveries. With the help of this versatile test subject, CSHL scientists have now dug up the secrets of a process that helps control inheritance. Credit: Martienssen lab/Cold Spring Harbor Laboratory

When organisms pass their genes on to future generations, they include more than the code spelled out in DNA. Some also pass along chemical markers that instruct cells how to use that code. The passage of these markers to future generations is known as epigenetic inheritance. It's



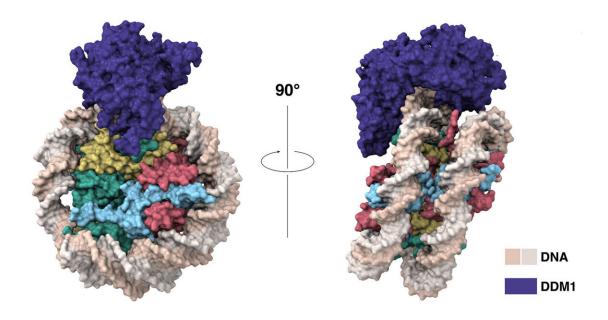
particularly common in plants. So, significant findings here may have implications for agriculture, food supplies, and the environment.

Cold Spring Harbor Laboratory (CSHL) Professors and HHMI Investigators Rob Martienssen and Leemor Joshua-Tor have been researching how <u>plants</u> pass along the markers that keep transposons inactive. Transposons are also known as jumping genes. When switched on, they can move around and disrupt other genes. To silence them and protect the genome, cells add regulatory marks to specific DNA sites. This process is called methylation. The research has been published in the journal *Cell*.

Martienssen and Joshua-Tor have now shown how protein DDM1 makes way for the enzyme that places these marks on new DNA strands. <u>Plant</u> <u>cells</u> need DDM1 because their DNA is tightly packaged. To keep their genomes compact and orderly, cells wrap their DNA around packing proteins called histones. "But that blocks access to the DNA for all sorts of important enzymes," Martienssen explains. Before methylation can occur, "you have to remove or slide the histones out of the way."

Martienssen and former CSHL colleague Eric Richards first <u>discovered</u> DDM1 30 years ago. Since then, researchers have learned it slides DNA along its packing proteins to expose sites needing methylation. Martienssen likens the movement to a yo-yo gliding along a string. The histones "can move up and down the DNA, exposing parts of the DNA at a time, but never falling off," he explains.





This cartoon model illustrates, for the first time, where and how the DDM1 protein (purple) grips onto DNA (beige) during cell division. Credit: Joshua-Tor lab/Cryo-EM Facility/Cold Spring Harbor Laboratory

Through genetic and biochemical experiments, Martienssen pinpointed the exact histones DDM1 displaces. Joshua-Tor used cryo-<u>electron</u> <u>microscopy</u> to capture detailed images of the enzyme interacting with DNA and associated packing proteins. They were able to see how DDM1 grabs onto particular histones to remodel packaged DNA. "An unexpected bond that ties DDM1 together turned out to correspond to the first mutation found all those years ago," Joshua-Tor says.

The experiments also revealed how DDM1's affinity for certain <u>histones</u> preserves epigenetic controls across generations. The team showed that a histone found only in pollen is resistant to DDM1 and acts as a placeholder during cell division. "It remembers where the histone was



during plant development and retains that memory into the next generation," Martienssen says.

Plants may not be alone here. Humans also depend on DDM1-like proteins to maintain DNA methylation. The new discovery may help explain how those proteins keep our genomes functional and intact.

More information: Robert A. Martienssen, Chromatin remodeling of histone H3 variants by DDM1 underlies epigenetic inheritance of DNA methylation, *Cell* (2023). DOI: 10.1016/j.cell.2023.08.001. www.cell.com/cell/fulltext/S0092-8674(23)00855-3

Provided by Cold Spring Harbor Laboratory

Citation: How plants pass down genetic memories (2023, August 28) retrieved 28 April 2024 from <u>https://phys.org/news/2023-08-genetic-memories.html</u>

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