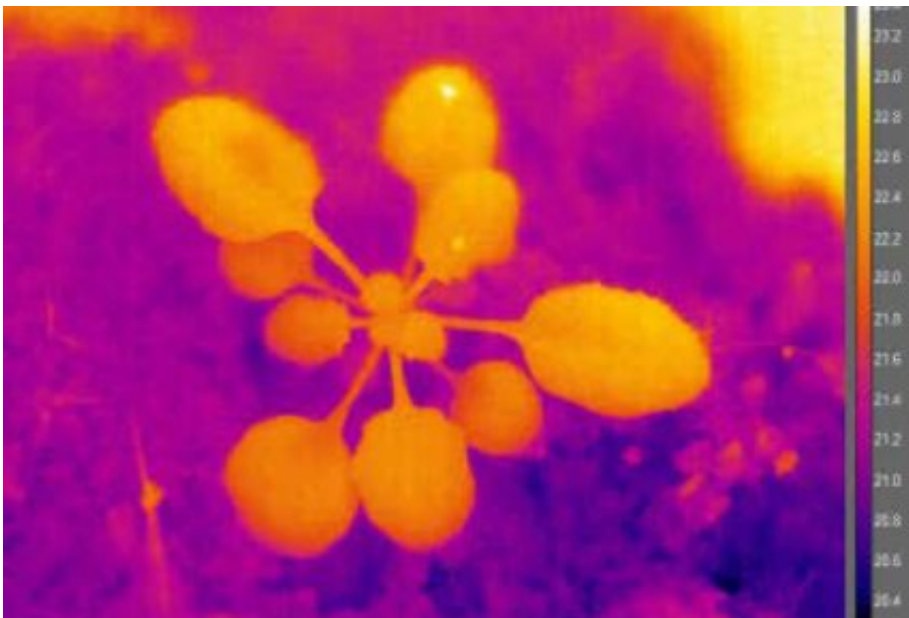


How plants adjust their body plan to cope with high temperature stress

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A thermal image of a plant that is able to keep its leaves cooler than its surroundings through perspiration and airflow. Stretching under high temperature leads to further cooling. Credit: Utrecht University

Biologists from Utrecht University have described a new molecular mechanism that allows plants to optimize their growth under suboptimal high-temperature conditions. The study offers promising leads for the development of climate warming-tolerant crops, which maintain a high yield under stressful high ambient temperatures, according to research leader Martijn van Zanten. They publish their findings November 25th

in *PNAS*.

Many crops suffer from high temperatures and each degree Celsius [temperature](#) increase can lead to up to 10 percent crop loss. This is a major problem in the context of current global warming and increasing demands for food. However, many [plant species](#) can adjust the shape of their stems and leaves, making them more resilient to high temperatures. This process is called thermomorphogenesis and leads to an 'open body plan,' allowing efficient evaporation, reduction of direct heat radiation from the sun and enables dissipation of the heat by improved air circulation around leaves. Thermomorphogenesis thus facilitates cooling and enables the plant to maintain optimal growth and production under sub-optimal high ambient temperatures. But many modern agricultural crops have lost that capacity.

Thermotolerant crops

The development of thermotolerant crops requires detailed knowledge of how plants perceive high temperatures and how they translate this signal into growth adjustments. "How plants detect temperature and which molecular factors contribute to thermomorphogenesis is, however, still poorly understood," explains last author Martijn van Zanten of Utrecht University, who led the international team of researchers from Utrecht, the United Kingdom, Italy, Sweden, Czech Republic, Australia and Wageningen. "We discovered an unknown molecular mechanism by which plants control thermomorphogenesis, especially in young seedlings, the most sensitive period in the plant's life when it comes to ambient temperature. These new pieces of the puzzle can make the development of future thermotolerant crops more efficient."

In the journal *PNAS*, the researchers demonstrate November 25th that the enzyme histone deacetylase 9 (HDA9) plays a key role in thermomorphogenesis in the widely used model plant *Arabidopsis*

thaliana. At increasing temperatures, the abundance of the enzyme rises, which results in the removal of epigenetic modifications of DNA-bound histone proteins that have an inhibiting effect on the synthesis of the well-known plant growth hormone auxin. As a result, auxin levels increase and the plant adjusts its stature.

Plant growth

"This new mechanism is scientifically very interesting, because it shows that a histone deacetylase 9 has an indirect positive effect on transcription, while histone deacetylases are generally accepted as suppressors of this process," says Van Zanten. "In addition, we show that HDA9 works independently of the only known temperature sensor, phytochrome B, which is also a light sensor. In doing so, we are exposing a new temperature signaling route in plants."

According to the researchers, this finding offers interesting possibilities for application. "We show that mutants in HDA9 are disturbed in the ability to adjust their body plan to high temperatures, but can still react normally to light signals from neighboring [plants](#) in dense vegetation. With this knowledge at hand, we now can uncouple light-controlled plant growth from temperature-controlled adjustments," says Van Zanten. This research therefore offers interesting starting points for the development of climate-warming-tolerant [crops](#), without compromising on other desirable properties.

More information: Lennard C. van der Woude et al., "Histone deacetylase 9 stimulates auxin-dependent thermomorphogenesis in *Arabidopsis thaliana* by mediating H2A.Z depletion," *PNAS* (2019). www.pnas.org/cgi/doi/10.1073/pnas.1911694116

Provided by Utrecht University

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