

Drought makes Borneo's trees flower at the same time

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The study species, *Shorea beccariana*. Credit: UZH

Tropical plants flower at supra-annual irregular intervals. In addition, mass flowering is typical for the tropical forests in Borneo and elsewhere, where hundreds of different plant timber species from the Dipterocarpaceae family flower synchronously. This phenomenon is all the more puzzling because both temperature and day length are relatively constant all year round due to geographical proximity to the equator. Up

to now it was supposed that several weeks of drought may trigger mass flowering in Borneo's forests. However, no empirical data and genetic analyses were available.

An international research team headed up by [evolutionary biologists](#) at the University of Zurich has now identified two [genes](#) responsible for the flowering of a tropical deciduous tree species *Shorea beccariana*. After drought periods, the two genes SbFT and SbSVP undergo dramatic transcriptional changes directly before flowering. The researchers can also confirm the flowering functions of these two genes using transgenic *Arabidopsis thaliana* plants.

85-meter canopy crane necessary for sample collection

The PhD student Masaki Kobayashi, his supervisor Professor Kentaro Shimizu and their Malaysian, Taiwanese and Japanese colleagues collected multiple buds from a single *Shorea beccariana* tree shortly before the start of flowering. "Given the fact that *Shorea* is a giant tree, having its crown at 40 meters of height, this sample collection was not easy at all", says Shimizu. Only with the help of an 85m high canopy crane were they able to collect samples at six different time points over a two-year period. Next, they analyzed the sample material using a next-generation sequencing procedure, which was initially developed for [human genome](#) and disease research. In this way, Kobayashi and Shimizu identified 98 genes that are associated with the flowering of the plant – including the genes SbFT and SbSVP, which showed transcriptional changes after a drought period and directly before flowering. The scientists then combined their genetic results with the meteorological data of the region. Kobayashi concludes that "Flowering in *Shorea beccariana* is triggered by a four-week drought in combination with elevated sucrose levels."



Community-level mass flowering (general flowering) in the Lambir Hills National Park in Borneo 2009 (Malaysia). Credit: UZH

Toward prediction of mass flowering

Climate change will affect the frequency of [drought](#) periods and is thus predicted to affect also the frequency of mass flowering. Environmental protection and restoration of the forests have so far been severely hindered by the irregularity of the mass flowering intervals, which are thus difficult to predict. It was never possible to know when the seeds needed could be collected. The genes that have been identified now indicate when mass flowering is about to happen. "Successively

monitoring of gene activity can help predict when mass flowering will take place", explains Kobayashi. This will make it possible to coordinate the collection of seeds and improve biodiversity and conservation programs substantially. Kentaro Shimizu and his colleagues will continue to explore these interactions in the newly created University Research Priority Program "Global Change and Biodiversity" at the University of Zurich.

More information: Masaki, J. et al. Mass flowering of tropical tree *Shorea beccariana* was preceded by expression changes in flowering and drought responsive genes, *Molecular Ecology*. May 8, 2013. [doi: 10.1111/mec.12344](https://doi.org/10.1111/mec.12344)

Provided by University of Zurich

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