

Newly discovered molecular network regulates vitamin C biosynthesis in kiwifruit

June 9 2022, by Zhang Nannan



The circle bar charts showing the AsA content (log2) of 48 different Actinidia species. Credit: WBG



Actinidia (kiwifruit) is rich in ascorbic acid (AsA) and Vitamin C (Vc) and is known as the "King of Vc," which makes it very popular with consumers. Interestingly, the fruit AsA content of different kiwifruit species varies up to 300- to 400-fold. However, the molecular regulation mechanisms of AsA synthesis in kiwifruit are still poorly understood.

Supervised by Prof. Zhong Caihong and Associate Prof. Li Dawei, Ph.D. student Liu Xiaoying from the Wuhan Botanical Garden of the Chinese Academy of Sciences first carried out systematic measurements of fruit AsA content in 48 <u>kiwifruit</u> species and found that the differences in AsA between kiwifruit species were mainly due to the extent of AsA accumulation in the early fruit development.

In addition, RNA-Seq, transient transformation, transgenic and <u>gene</u> <u>editing</u> were performed on kiwifruit with a high AsA content (six to tenfold that of commercial kiwifruit) to uncover AceGGP3, a key gene for AsA synthesis in Actinidia eriantha (A. eriantha).

They identified a 1R-subtype myeloblastosis (MYB)-type transcription factor, AceMYBS1 by transcriptome analysis and RT-qPCR. Overexpression editing of AceMYBS1 in A. eriantha showed that AceMYBS1 promoted AsA accumulation.

Subsequently, yeast 1-hybrid, dual-luciferase <u>assay</u> and electrophoretic mobility shift assay revealed that AceMYBS1 bound directly to the AceGGP3 promoter and activated transcription of the AceGGP3 gene, increasing AsA accumulation.





Circle heatmap showing the expression profifile (log2FPKM) of orthologous genes in the L-galactose pathway in the kiwifruit. Credit: WBG

Further, they uncovered the downstream protein AceGBF3 (a G-box binding factor) that interacted with AceMYBS1 using yeast 2-hybrid library screening assay. Additionally, yeast-2 hybrid, bimolecular luminescence complementation assay, pull-down, bimolecular fluorescence complementation assay and transgenic assay were carried out to demonstrate that AceMYBS1 interacted with AceGBF3 in vitro and in vivo.

Moreover, the interaction between AceMYBS1 and AceGBF3 can activate the transcription of AceGGP3 and synergistically promote the AsA synthesis of A. eriantha.



Previous studies have suggested that hormones could affect AsA synthesis. The newly-discovered AceGBF3-AceMYBS1-AceGGP3 molecular regulatory network by Zhong's team can also be inhibited by <u>abscisic acid</u>, which is a foundation that will provide new clues for future elucidation of hormonal regulation of Vitamin C synthesis.

The study has been published in *New Phytologist*, titled "Kiwifruit MYBS1-like and GBF3 transcription factors influence L-<u>ascorbic acid</u> biosynthesis by activating transcription of GDP-L-galactose phosphorylase 3."

More information: Xiaoying Liu et al, Kiwifruit MYBS1-like and GBF3 transcription factors influence l -ascorbic acid biosynthesis by activating transcription of GDP-L-galactose phosphorylase 3, *New Phytologist* (2022). DOI: 10.1111/nph.18097

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