

Pyrethrin Biosynthesis: From a Phytohormone to Specialized Metabolite

Dalmatian chrysanthemum (*Tanacetum cinerariifolium*) contains pyrethrins, which are highly effective natural insecticides that are nontoxic to most animals and biodegradable. Purification of these natural compounds, however, is more costly than production of their synthetic analogs, although the synthetic analogs are less biodegradable and can pose a danger for vertebrates. As we have become more environmentally conscious, the biosynthesis pathway of pyrethrins has attracted greater interest.

Pyrethrins are synthesized by the linking of two parts (Fig. 1). The first is an irregular monoterpene whose biosynthesis is well established (Xu et al., 2019), and the second is derived from the phytohormone jasmonate. The biosynthesis pathway of jasmonates and their role as defensive hormones are well established (Wasternack and Hause, 2013), but less is known about how pyrethrins are synthesized. Recent publications have already elucidated some of the steps, including the production of jasmolone from jasmonate by jasmone hydroxylase, but the enzyme responsible for the desaturation of the alkyl chain in jasmolone remains unknown (Li et al., 2018). This step is specifically important for the biosynthesis of pyrethrin I and II, which are the most abundant compounds and namesakes for this class of compounds.

In this issue of *Plant Physiology*, Li et al. (2019) investigated this desaturation reaction. Previous expression analysis identified 10 candidate enzymes, all belonging to the cytochrome P450 (CYP) superfamily (Xu et al., 2018). Most plants encode large numbers of CYPs in their genome, and these enzymes are known to catalyze a multitude of complex reactions that are essential for natural product biosynthesis (Hamberger and Bak, 2013). To identify the enzyme catalyzing jasmolone desaturation, candidates were transiently expressed together with *TcJMH*, the gene encoding jasmone hydroxylase, in *Nicotiana benthamiana*, with subsequent immersion in jasmone solution for substrate delivery. Only one CYP, which was henceforth named pyrethrolone synthase (TcPYS), was able to produce pyrethrolone. Further characterizations of TcPYS showed it to be highly substrate specific and localized to the endoplasmic reticulum membrane of trichomes on the ovaries of chrysanthemum flowers.

TcPYS, which is also called CYP82Q3 according to CYP nomenclature, is part of the CYP82 family. This CYP family is heavily involved in the biosynthesis of natural products from the flavonoid, alkaloid, terpene,

and coumarin classes. The majority of characterized CYP82 enzymes catalyze hydroxylation reactions, which makes the desaturation catalyzed by TcPYS unusual. Moreover, only a very limited number of CYPs outside of the CYP82 family are known to catalyze such a reaction (Zhang and Li, 2017; Guengerich, 2018). Therefore, TcPYS is also an interesting target for investigations into the reaction mechanism of this desaturation reaction that can potentially progress through two different reaction pathways.

This publication identified a mechanistically interesting step in pyrethrin biosynthesis, which could ultimately lead to a less expensive synthetic biology route for the production of these natural insecticides that are less ecotoxic than their synthetic analogs. However, for this to occur, the earlier biosynthetic steps from jasmonic acid, or potentially a precursor of jasmonic acid, must be identified.

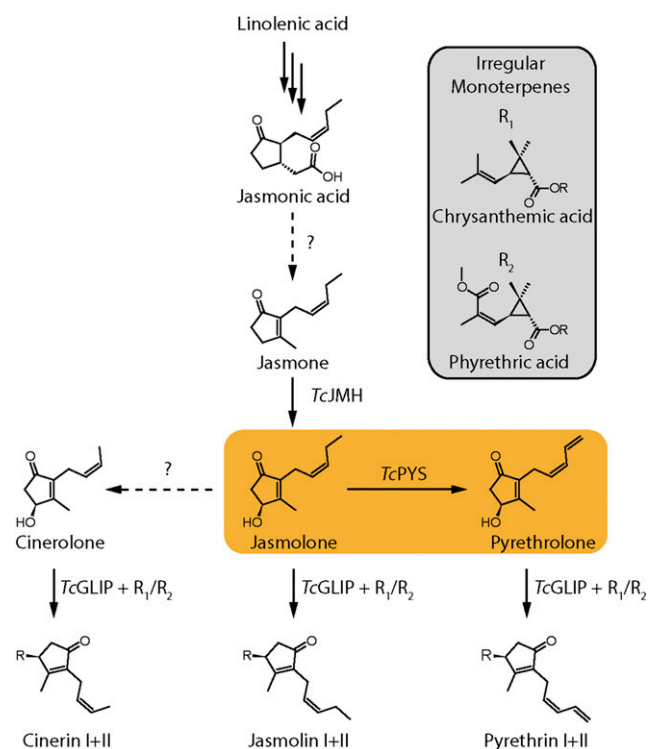


Figure 1. Core pyrethrin biosynthetic pathway. The enzyme(s) for jasmone and cinerolone biosynthesis are unknown, as indicated with question marks. TcJMH is the *T. cinerariifolium* jasmolone hydroxylase and TcGLIP is the GDSL lipase-like protein that links one of the irregular monoterpene parts of the molecule to cinerolone, jasmolone, or pyrethrolone. TcPYS is the pyrethrolone synthase investigated by Li et al. (2019), which, unusually for a CYP, catalyzes a desaturation reaction. (Adapted from Li et al., 2018.)

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This pathway is also intriguing as it has previously been shown that jasmonate hormones induce the production of pyrethrins through transcriptional regulation of pyrethrin biosynthetic genes, and jasmonate also serves as their biosynthetic precursor. This observation raises questions about the complicated regulation of a highly abundant natural product that is derived from a significantly less abundant phytohormone. How is the balance maintained?

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