

## Studies on regulation of the plasma membrane H<sup>+</sup>-ATPase activity with respect to blue light- dependent stomatal opening

山内, 翔太

<https://hdl.handle.net/2324/1806836>

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出版情報：九州大学, 2016, 博士（理学）, 課程博士  
バージョン：  
権利関係：やむを得ない事由により本文ファイル非公開（3）

氏 名 : 山内 翔太

論 文 名 : Studies on regulation of the plasma membrane H<sup>+</sup>-ATPase activity with respect to blue light-dependent stomatal opening (青色光依存の気孔開口を指標とした細胞膜 H<sup>+</sup>-ATPase の活性制御に関する研究)

区 分 : 甲

### 論 文 内 容 の 要 旨

The plasma membrane H<sup>+</sup>-ATPase acts as a primary transporter in fungi and plants, and drives a large number of secondary transporters by enhancing the negative membrane potential and pH gradient across the plasma membrane. The H<sup>+</sup>-ATPase is responsible for various physiological responses, including cell elongation, phloem loading, and stomatal opening. Stomatal opening is one of the suitable responses to investigate the activity of H<sup>+</sup>-ATPase. Functions of H<sup>+</sup>-ATPase have been analyzed mainly by biochemical technique, but genetic evidence for the functions are scarcely available because all 11 H<sup>+</sup>-ATPase isoforms in *Arabidopsis* are expressed in stomatal guard cells. The functional redundancy of the H<sup>+</sup>-ATPase makes it difficult to identify the specific role of the H<sup>+</sup>-ATPase in plants.

In this study, I performed genetic screening by thermography and isolated *blus2* (*blue light signaling2*), in which opening responses of stomata were reduced. Whole genome re-sequencing demonstrated that the responsible gene of *blus2* is *AHA1* (*Arabidopsis H<sup>+</sup>-ATPase1*) that encodes one of the isoform of H<sup>+</sup>-ATPase. I thus renamed *blus2* as *ahal-10*. By contrast, the mutation in *AHA2* (*aha2*) and *AHA5* (*aha5*) did not affect stomatal opening in response to blue light, although the transcripts of two isoforms were highly expressed in guard cells. I showed that the amount of AHA1 protein was larger than those of AHA2 and AHA5 in guard cells. From these results, I conclude that AHA1 plays a major role in blue light-dependent stomatal opening.

Phosphorylation of the penultimate threonine in the C-terminus and subsequent 14-3-3 protein binding are essential for the activation of H<sup>+</sup>-ATPase. However, no genetic evidence for this was obtained. Using *ahal-9* mutant that lacks AHA1, I determined the activity of H<sup>+</sup>-ATPase with respect to stomatal response. To do this, I transformed *ahal-9* with non-phosphorylatable form of the penultimate threonine in AHA1 (Threonine 948 to Ala; T948A). The transformant having T948A did not recover the stomatal phenotype of *ahal-9*. 14-3-3 protein were not bound to the H<sup>+</sup>-ATPase in response to blue light in the transformant. The results provide genetic evidence that phosphorylation of the penultimate threonine is essential for H<sup>+</sup>-ATPase activation.