

Contents lists available at ScienceDirect

Data in Brief



Data Article

CPC-ETC1 chimeric protein localization data in *Arabidopsis* root epidermis

R. Tominaga-Wada*, T. Wada

Graduate School of Biosphere Sciences, Hiroshima University, 1-4-4 Kagamiyama, Higashi-Hiroshima 739-8528, Japan

ARTICLE INFO

Article history: Received 9 January 2018 Accepted 17 April 2018 Available online 25 April 2018

ABSTRACT

Intercellular movement of transcription factor proteins is essential for plant development. The R3 type MYB transcription factor protein, CAPRICE (CPC), moves from non-hair cells to root-hair cells where it promotes root hair formation in *Arabidopsis* root epidermis. In contrast, the CPC homolog of ENHANCER OF TRY AND CPC1 (ETC1) cannot move in root epidermal cells. In this work, we present protein localization data of CPC-ETC1 chimeric proteins. Localization of CPC-ETC1-GFP fusion proteins of chimera1 and chimera2 transgenic plants was observed using confocal laser scanning microscope. Insertion of ETC1-specific amino acids into CPC somewhat prevents normal protein localization of CPC in root epidermal cells. Cell-to-cell movement of chimera1 and chimera2 proteins from non-hair cells to root-hair cells was interfered. Nuclear localization was also inhibited, especially in chimera1.

© 2018 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

Specifications table

Subject area	Biology
More specific subject area	Plant Sciences
Type of data	Figure

DOI of original article: https://doi.org/10.1016/j.ydbio.2018.01.002 * Corresponding author.

Corresponding author.

https://doi.org/10.1016/j.dib.2018.04.055

E-mail address: rtomi@hiroshima-u.ac.jp (R. Tominaga-Wada).

^{2352-3409/© 2018} The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

How data was acquired	Confocal laser scanning microscope (Zeiss LSM-510 Meta)
Data format	Raw
Experimental factors	-
Experimental features	-
Data source location	Higashi-Hiroshima, Japan
Data accessibility	Data are presented in this article
Related research article	Effect of amino acid substitution of CAPRICE on cell-to-cell movement
	ability in Arabidopsis root epidermis, Developmental Biology, in press.

Value of the data

- The data provide information about the protein localization and cell-to-cell movement properties
 of CPC-ETC1 chimeric proteins in Arabidopsis root epidermal cells.
- This study shows the importance of precise amino acid sequence of CPC in proper cell-to-cell movement ability in *Arabidopsis* root epidermal cells.
- The cell-to-cell movement data of chimera proteins in *Arabidopsis* root epidermis helps to understand the functions of R3-type MYB transcription factors.

1. Data

Fig. 1 shows the localization of CPC-ETC1 chimera-GFP fusion proteins in *Arabidopsis* root epidermis. The level of GFP fluorescence was slightly lower in root hair cells than in non-hair cells of all transgenic plants of Chimera 1#2, Chimera 2#2, and Chimera 2#3 in this study. Clear nucleus localization of the GFP fusion protein was not observed in Chimera 1#2 transgenic epidermal cells.

2. Experimental design, materials and methods

2.1. Plant material and growth conditions

This study utilized previously reported transgenic *Arabidopsis thaliana* (L.) Heynh. lines CPC-ETC1 Chimera 1 #2, Chimera 2 #2, and Chimera 2 #3 [1] of the ecotype Columbia (Col-0). Seeds were surface-sterilized and sown on 1.5% agar plates as described previously [2]. The plates with sawn seeds were kept at 4 °C for 2 days and then incubated at 22 °C under constant white light (50–100 μ mol m⁻² s⁻¹). For each transgenic line, five-day-old seedlings were examined for the GFP fused chimeric protein localization.

2.2. Gene constructs

Gene constructs for CPC-ETC1 chimeric proteins were generated in the *CPCp:CPC:2xGFP* backbone [3] by TaKaRa (TaKaRa, Japan). To create the Chimera 1 construct, *ETC1*-specific DNA sequence corresponding to the NT amino acid sequence was inserted into the *CPC* coding region between the 11th (D) and 12th (K) position of the CPC amino acid sequence in *CPCp:CPC:2xGFP* [1]. To create the Chimera 2 construct, *ETC1*-specific DNA sequence corresponding to the HLKTNPTIV amino acid sequence was inserted into the *CPC* coding region between the 21st (K) and 22nd (A) position of the CPC amino acid sequence in *CPCp:CPC:2xGFP* [1].

2.3. Transgenic plants

The floral dip method was used for the plant transformation in this study [4], and the transgenic plants were selected on $0.5 \times$ Murashige and Skoog's agar plates containing 50 mg/L kanamycin. The homozygous transgenic lines were selected for kanamycin resistance.



Fig. 1. Distribution of GFP fluorescence in the transgenic *Arabidopsis* plants expressing Chimera 1 and Chimera 2 constructs. Homozygous transgenic lines of Chimera 1#2, Chimera 2#2, and Chimera 2#3 are shown. Confocal laser scanning microscope images showing GFP (green) and propidium iodide (red) fluorescence in the root epidermis of five-day-old seedlings. Asterisks indicate the root hair cell files. Scale bars: 100 µm.

2.4. Microscopy

For each transgenic line of Chimera 1#2, Chimera 2#2, and Chimera 2#3, five-day-old seedling roots were analyzed for GFP fluorescence. The transgenic GFP fusion lines were stained with $5 \mu g/mL$ propidium iodide for 30 s and then washed with water. Confocal images were obtained with a Zeiss LSM-510 Meta confocal laser scanning microscope using 488-nm laser lines for GFP excitation. Image processing was performed with Photoshop version 7.0 (Adobe Systems, CA, USA).

Acknowledgments

We wish to thank T. Kurata, R. Sano, and T. Ishida for useful comments, and M. Onishi, Y. Nukumizu, and M. Iwata for providing technical support. This work was supported by JSPS KAKENHI (grant numbers 15K14656 and 16K07644).

Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at http://dx.doi. org/10.1016/j.dib.2018.04.055.

Transparency document. Supporting information

Supplementary data associated with this article can be found in the online version at http://dx.doi. org/10.1016/j.dib.2018.04.055.

References

- R. Tominaga-Wada, T. Wada, Effect of amino acid substitution of CAPRICE on cell-to-cell movement ability in Arabidopsis root epidermis, Dev. Biol. 435 (2018) 1–5.
- [2] K. Okada, Y. Shimura, Reversible root tip rotation in Arabidopsis seedlings induced by obstacle-touching stimulus, Science 250 (1990) 274–276.

- [3] T. Wada, T. Kurata, R. Tominaga, Y. Koshino-Kimura, T. Tachibana, K. Goto, M.D. Marks, Y. Shimura, K. Okada, Role of a positive regulator of root hair development, CAPRICE, in Arabidopsis root epidermal cell differentiation, Development 129 (2002) 5409–5419.
- [4] S.J. Clough, A.F. Bent, Floral dip: a simplified method for Agrobacterium-mediated transformation of Arabidopsis thaliana, Plant J. 16 (1998) 735-743.