Yong Wang

Title: Associate Professor

Address: State Key Laboratory of Crop Stress Biology for Arid Areas, College of Agronomy, Northwest A&F University, Yangling 712100, Shaanxi, China.

Tel: +86-29-87082854

Email: wangyong2114@163.com

• Education Background:

2011 D.S.C. Nanjing Agricultural University, Nanjing, China;2006 B.A. Nanjing Agricultural University, Nanjing, China.



• Working Experiences:

2018-present Associate professor of the College of Agronomy, Northwest A&F University;2011-2017 Lecturer of the College of Agronomy, Northwest A&F University.

• Research Interests:

Biosynthesis and regulation of wheat cuticular wax. Molecular biology.

• Professional Activities:

Published articles as the first author or corresponding author in the Plant Physiology, Plant Journal, Journal of Experimental Botany, Plant and Cell Physiology, and Frontiers in Plant Science. As a reviewer of the Journal of Experimental Botany, Agriculture, Advances in Applied Physiology, and Frontiers in Microbiology.

Biosynthesis and drought resistance of cuticular wax alkanes in wheat

Yong Wang¹

⁽⁵Northwest A&F University)

Abstract: **(Objective)** The aim of the work was to identify and functionally characterize a key alkane biosynthesis gene TaCER1-6A from wheat. [Method] Overexpression and CRISPR/Cas9-mediated gene editing of TaCER1-6A were carried out in wheat, and the wax content, cuticle permeability and drought tolerance of TaCER1-6A transgenic lines and wild-type (WT) were further analyzed by gas chromatography-mass spectrometry (GC-MS), gas chromatography-flame ionization detection (GC-FID), chlorophyll leaching, and water loss assays. The biological function of TaCER1-6A was characterized in detail. And dual-luciferase (LUC), yeast one-hybrid (Y1H), and β -glucuronidase (GUS) activation assays were used to confirm the interactions of R2R3-type MYB transcription factors TaMYB96-2D/5D and the promoter of TaCER1-6A. [Result] The CRISPR/Cas9-mediated knockout mutation in TaCER1-6A greatly reduced the contents of C27, C29, C31, and C33 alkanes in wheat leaves, while TaCER1-6A overexpression significantly increased the contents of C27, C29, C31, and C33 alkanes in wheat leaves, suggesting that TaCER1-6A is specifically involved in the biosynthesis of C27-C33 alkanes on wheat leaf surfaces. TaCER1-6A knockout lines exhibited increased cuticle permeability and reduced drought tolerance, whereas TaCER1-6A overexpression lines displayed reduced cuticle permeability and enhanced drought tolerance. TaCER1-6A was highly expressed in flag leaf blades and seedling leaf blades and could respond to abiotic stresses and abscisic acid (ABA). TaCER1-6A was located in the endoplasmic reticulum (ER), which is the subcellular compartment responsible for wax biosynthesis. A total of three haplotypes (HapI/II/III) of TaCER1-6A were identified in 43 wheat accessions, and HapI was the dominant haplotype (95%) in these wheat varieties. Additionally, we identified two R2R3-MYB transcription factors TaMYB96-2D and TaMYB96-5D that bound directly to the conserved motif (CAACCA) in promoters of the TaCER1-6A. **Conclusion** TaCER1-6A is required for C27–C33 alkanes biosynthesis and improves drought resistance in wheat, suggesting that TaCER1-6A is a promising tool aiming at generating wheat cultivars with more alkanes contents and improved drought tolerance via molecular breeding and transgenic strategies.

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