



Editorial: Lipid Metabolism in Development and Environmental Stress Tolerance for Engineering Agronomic Traits

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Keywords: lipid biosynthesis and turnover, membrane and storage lipids, development, biotic and abiotic stress, genetic engineering

Editorial on the Research Topic

OPEN ACCESS

Edited and reviewed by:

Xue-Rong Zhou, Agriculture and Food, Commonwealth Scientific and Industrial Research Organisation (CSIRO), Australia

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Specialty section:

This article was submitted to Plant Metabolism and Chemodiversity, a section of the journal Frontiers in Plant Science

> Received: 12 July 2021 Accepted: 26 July 2021 Published: 24 August 2021

Citation:

Du Z-Y, Hoffmann-Benning S, Wang S, Yin L, Zienkiewicz A and Zienkiewicz K (2021) Editorial: Lipid Metabolism in Development and Environmental Stress Tolerance for Engineering Agronomic Traits. Front. Plant Sci. 12:739786. doi: 10.3389/fpls.2021.739786 Lipid Metabolism in Development and Environmental Stress Tolerance for Engineering Agronomic Traits

Lipids are the essential building blocks of cellular membranes, and they play fundamental roles in numerous biological activities such as photosynthesis, protection, environmental and cellular communication, and storage of carbon and energy. Lipid metabolism is a dynamic and complicated process that includes lipid biosynthesis, transport, accumulation, turnover, and excretion, acting to regulate plant development and tolerance to various environmental stresses. Understanding how lipid metabolism regulates the development and growth in response to adverse conditions raises many interesting questions. Communication between membrane and storage lipids has been observed during the progress, for example, membrane degradation accompanied by lipid droplet (LD) accumulation. Recently, autophagy has been connected with lipid metabolism in plants and algae under stress conditions (Zienkiewicz and Zienkiewicz). Because of the vital role of lipid metabolism in photosynthetic organisms, exploring lipid processes is very important for engineering crops to obtain better agronomic traits including productivity, nutrition, stress tolerance and resilience.

This Research Topic includes 10 original review and research articles, with a special focus on lipid metabolism in crops. The review by Zienkiewicz and Zienkiewicz summarizes the most recent advances on LD degradation, which is a key process for the release of triacylglycerol (TAG). As the major reservoir of cellular carbon and energy, TAG is packed in LD, a dynamic organelle at the center of cellular metabolism. The review also compares the LD degradation in plants and algae (**Figure 1A**). Looking at lipid metabolism in the context of membrane composition and transport, Salvaing et al. report the function of the Arabidopsis ALA10, a phospholipid flippase of the P₄ type-ATPase family, in reducing the 18:2 desaturation of phosphatidylcholine in the endoplasmic reticulum (ER), while Liao et al. show the roles of two rice acyl-CoA-binding proteins, OsACBP4 (plasma membrane and ER) and OsACBP5 (apoplast), in the intra- and extracellular transport of acyl-CoA esters, respectively (**Figure 1B**). Other contributions directly address the role of lipid



metabolism during development, EgMIXTA1, an R2R3 MYB-type transcription factor, promotes the cuticular wax formation in Eustoma grandiflorum (lisianthus) leaves (Wang et al.); together with the specificities of extraplastidial acyltransferase enzymes, the olive OeFAD2-2 and OeFAD2-5 regulate the oleic/linoleic acid ratio in the mesocarp of olive cultivars (Figure 1C) (Hernández et al.). The Research Topic also highlights the significance of lipid metabolism in environmental stress response and tolerance (Figure 1D). In Arabidopsis, expression of CsKCS6, a gene coding the 3-Ketoacyl-CoA synthase (KCS) that catalyzes the biosynthesis of very-long-chain fatty acid (VLCFA) in Citrus sinensis (Newhall navel orange), reduces water loss and ion leakage and increases tolerance to drought and salt stress (Guo et al.); Arabidopsis phospholipid:diacylglycerol acyltransferase 1 (AtPDAT1) promotes the activities of acyl-CoA:lysophosphatidylcholine acyltransferase (LPCAT) and acylCoA:lysophosphatidylethanolamine acyltransferase (LPEAT) and improves resilience to heat and cold exposure (Demski et al.). In *Vaccinium corymbosum* L., cv. Duke (Duke blueberry), intermittent warming treatment of the fruit regulates its lipid metabolism such as increases in phosphatidylcholine, linoleic acid, and oleic acid, which enhances the low-temperature tolerance in the fruit (Dai et al.). In the roots of maize, integrated lipidomic and transcriptomic analyses reveal the membrane lipid remodeling and gene regulations under saline-alkaline (Xu et al.) or cold stress (Zhao et al.), which are applicable for engineering the crop for better performance under these stress conditions.

The collection of articles in this Research Topic demonstrates the significance of lipid metabolism in various biological processes and the findings will contribute to engineering agronomic traits of crops for the increasing demand for food and bioproducts.

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

ACKNOWLEDGMENTS

The editors would like to thank all reviewers who evaluated manuscripts and contributors for this Research Topic. Research of the Topic Editors was supported by the funds from CTAHR (2303491) and OVCR, UHM (Z-YD), Natural Science Foundation of Shaanxi Province, China (2021JM-604) (LY), NSF (1841251) and USDA-NIMSS (NC-1203) (SH-B).

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