The Research Grants Council of Hong Kong NSFC/RGC Joint Research Scheme ______Joint Completion Report___

(Please attach a copy of the completion report submitted to the NSFC by the Mainland researcher)

Part A: The Project and Investigator(s)

1. Project Title

Mechanistic Analysis of Palmitate Transferase DHHC11 that Regulates Multiple Developmental and Cellular Processes 棕榈酰转移酶 DHHC11 调控植物生长发育的分子及细胞机理研究

2. Investigator(s) and Academic Department/Units Involved

	Hong Kong Team	Mainland Team
Name of Principal Investigator (with title)	Professor JIANG Liwen	Professor ZHANG Yan
Post	Professor	Professor
Unit / Department / Institution	School of Life Sciences, CUHK	College of Life Sciences, Shandong Agricultural Univ. State Key Lab of Crop Biology
Contact Information	ljiang@cuhk.edu.hk	yzhang@sdau.edu.cn
Co-investigator(s) (with title and institution)		Researcher, Sha Li Researcher, Liang-Zi Zhou Researcher, Xin-Ying Zhao Researcher, Fu-Rong Ge (Shandong Agricultural University)

3. **Project Duration**

	Original	Revised	Date of RGC/ Institution Approval (must be quoted)
Project Start date	1-1-2013		
Project Completion date	31-12-2016		
Duration (in month)	48		
Deadline for Submission of Completion Report	31-12-2017		

Part B: The Completion Report

5. Project Objectives

- 5.1 Objectives as per original application
 - 1. Characterizing DHHC11-dependent developmental processes and abiotic stress responses.
 - 2. Analyzing subcellular localization of DHHC11 and identifying critical residues or motifs determining its subcellular localization.
 - 3. Identifying DHHC11-regulated vacuolar trafficking routes for tonoplast proteins.
 - 4. Identifying interacting proteins of DHHC11 and building an interacting network centred on DHHC11.
- 5.2 Revised Objectives

Date of approval from the RGC:

NSFC/RGC 8 (Revised 10/15)

Reasons for the change:

1. 2. 3.

6. Research Outcome

Major findings and research outcome *(maximum 1 page; please make reference to Part C where necessary)*

Post-translational modifications, such as phosphorylation and lipidation, play important roles in the subcellular localization, interaction, and function of substrate proteins. Protein S-acylation, or palmitoylation is an evolutionarily conserved post-translation modification that catalyzes the reversible addition of palmitate group to substrate cysteine residues. A large number of proteins in eukaryotes have been demonstrated to be palmitoylation substrates. However, functional analysis on the enzymes catalyzing this process, i.e. protein S-acyltransferases (PATs), was scarce. There was only one plant PAT gene with demonstrated function. The purpose of this project is to understand the function of plant PATs in growth, development, as well as responses to environmental cues. Detailed research findings are listed as follows.

1. We have reported functional analysis of *PAT10* (*DHHC11*) in Arabidopsis. *PAT10* is constitutively expressed. Functional loss of *PAT10* resulted in pleiotropic developmental defects, including reduced vegetative and reproductive growth, sporophytic and gametophytic male defects, and compromised ability of pistils to support pollen tube growth. Both cell expansion and cell division were reduced in *pat10*. Using stable transgenic plants for fluorescent colabeling, we showed that PAT10 localizes at vacuolar membrane but not at Golgi or any post-Golgi vesicles. In addition to the developmental defects, mutants of *PAT10* were hypersensitive to salt stresses. We further identified a subfamily of calcineurin B–like proteins (CBLs) as putative substrates of PAT10. Localization of CBL2, CBL3, and CBL6 at the tonoplast depended on functional PAT10. Our results demonstrate that PAT10-mediated protein palmitoylation at the tonoplast is critical for development and salt tolerance in Arabidopsis. Results were published on the Plant Cell (2013).

Related research:

With the support of this grant to graduate students, we have also performed other related researches. We thus consider these publications as the research output of this grant. Selected related research findings are listed as follows.

2. PAT14 plays an evolutionarily conserved role in SA-mediated leaf senescence. We reported the characterization of PAT14, a Golgi-localized PAT. Functional loss of *PAT14* resulted in precocious leaf senescence after floral transition. Transcriptomic data indicated that PAT14 mediates SA biosynthesis and signaling. Indeed, genetic interference of the NPR1-mediated SA signaling abolished the precocious leaf senescence induced by *PAT14* loss-of-function. Interestingly, we have isolated the maize and wheat homologs of Arabidopsis PAT14 and found out that they have conserved function. Results were published on Scientific Reports (2016).

3. Prenylation, the post-translational attachment of prenyl groups to substrate proteins, can affect their distribution and interactomes. Prenylation of ROPs by PLP mediates their membrane association and stability during root hair growth. Prenylation is a post-translational modification that anchors soluble proteins to membranes. Small GTPase ROPs are substrates of prenylation. We reported that PLP-mediated prenylation, together with GDI and palmitoylation, controls ROP signaling in root hairs. Results were published on Plant Journal (2016).

4. Protein palmitoylation, which is critical for membrane association and subcellular targeting of many signaling proteins, is catalyzed mainly by protein S-acyl transferases. By combining pharmacological and genetic approaches and using root hairs as a model, we show that protein palmitoylation, regulated by protein S-acyl transferases at different endomembrane compartments such as the Golgi and the vacuole, is critical for the polar

growth of root hairs in Arabidopsis. Inhibition of protein palmitoylation by 2-BP disturbed key intracellular activities in root hairs. Although some of these effects are likely indirect, the cytological data reported here will contribute to a deep understanding of protein palmitoylation during tip growth in plants. Results were published on BMC Plant Biology (2015).

5. Characterization of MON1 function in pollen development in Arabidopsis. Programmed cell death (PCD)-triggered degradation of plant tapetum is essential for microspore development and pollen coat formation; however, little is known about the cellular mechanism regulating tapetal PCD. Here, we demonstrate that Rab7-mediated vacuolar transport of tapetum degradation-related cysteine proteases is crucial for tapetal PCD and pollen development in Arabidopsis, with the following evidence: (1) The monensin sensitivity1 (mon1) mutants, which are defective in Rab7 activation, showed impaired male fertility due to a combined defect in both tapetum and male gametophyte development. (2) In anthers, MON1 showed preferential high-level expression in tapetal cell layers and pollen. (3) The mon1 mutants exhibited delayed tapetum degeneration and tapetal PCD, resulting in abnormal pollen coat formation and decreased male fertility. (4) MON1/CCZ1-mediated Rab7 activation was indispensable for vacuolar trafficking of tapetum degradation-related cysteine proteases, supporting that PCD-triggered tapetum degeneration requires Rab7-mediated vacuolar trafficking of these cysteine proteases. (5) MON1 mutations also resulted in defective pollen germination and tube growth. Taken together, tapetal PCD and pollen development require successful MON1/CCZ1-mediated vacuolar transport in Arabidopsis. Results were published on the Plant Physiology (2017).

Potential for further development of the research and the proposed course of action (maximum half a page)

The research projects have been successfully addressed. No further development of research is proposed.

7. The Layman's Summary

(describe <u>in layman's language</u> the nature, significance and value of the research project, in no more than 200 words)

Palmitoylation regulates enzyme activity, protein stability, subcellular localization, and intracellular sorting. Many plant proteins are palmitoylated. However, little is known about protein S-acyl transferases (PATs), which catalyze palmitoylation. Here, we report that the tonoplast-localized PAT10 (DHHC11) is critical for development and salt tolerance in Arabidopsis thaliana. We have identified several calcineurin B–like proteins as the substrates for PAT10 palmitoylation, by using the cell and molecular biology tools. We propose that PAT10-mediated palmitoylation is critical for vacuolar function by regulating membrane association or the activities of tonoplast proteins.

Part C: Research Output

8. Peer-reviewed journal publication(s) arising <u>directly</u> from this research project

(Please attach a copy of each publication and/or the letter of acceptance if not yet submitted in the previous progress report(s). All listed publications must acknowledge RGC's funding support by quoting the specific grant reference.)

The Latest S	tatus of Publications	Author(s)	Title and Journal/ Book	Submitted	Attached	Acknowl	Accessi
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Year of publicati on	Year of Acceptance (For paper	Under Review	Under Preparation	(bold the authors belonging to the project teams and	and other necessary	to RGC (indicate the year	report (Yes	edged the support of this	ble from the institutio
	accepted but not yet published)		(optional)	denote the corresponding author with an asterisk*)	specified)	ending of the relevant progress report)		Joint Research Scheme	nal repositor y (Yes or No)
2014				,	mechanisms for ER and Golgi membrane proteins. <i>Trends in</i>	2014	Yes	Yes	No
2014				Q, Gao C, Ding Y, Zeng YL, Ueda T, Nakano A	Activation of the Rab7 GTPase by the MON1-CCZ1	2014	Yes	Yes	No
2014				-	network-located AP1 gamma adaptins mediate		Yes	Yes	No
2015				Zhuang X, Cui Y, Fu X, He Y, Zhao Q, Zeng Y, Shen J, Luo	component FREE1 in		Yes	Yes	No

2015	Zhuang V	Endocytic	No	Yes	Yes	No
	Cui Y, Gao C and *Jiang L	pathways crosstalk in plants. <i>Current Opinion</i> <i>in Plant Biology</i> 28:39-47.				
2015	Chung KP, Li B, Lai CM, Lam SK, Wang X, Cui Y, Gao C, Luo M, Wong KB, Schekman R, and * Jiang L	function of protein ER export in Arabidopsis thaliana. Proceedings of the National Academy of Sciences of the United States of America 112(46):14360-5.	No	Yes	Yes	No
2015	Feng, Zhao,	Protein palmitoylation is critical for the polar growth of root hairs in Arabidopsis. <i>BMC</i> <i>Plant Biology</i> 15;50	No	Yes	Yes (NSF C: 31261 16049 0)	No
2016	Zhuang X, Wang J and	Biogenesis of Plant Prevacuolar Multivesicular Bodies. <i>Molecular</i> <i>Plant</i> 9(6):774-86.	No	Yes	Yes	No
2016		Using Fluorescent Protein Fusions to Study Protein Subcellular Localization and Dynamics in Plant Cells. <i>Methods in</i> <i>Molecular Biology</i> 1474: 113-123	No	Yes	Yes	No

2016		Song, Wang,	Precocious leaf senescence by functional loss of PROTEIN S-ACYLTRANSF ERASE14 involves the NPR1-dependent salicylic acid signaling. <i>Scientific Report</i> 6, 20309	No	Yes	Yes (NSF C: 31261 16049 0)	No
2016			PLURIPETALA mediates ROP2 localization and stability in parallel to SCN1 but synergistically with TIP1 in root hairs. <i>The Plant</i> <i>Journal</i> 86, 413-426	No	Yes	Yes (NSF C: 31261 16049 0)	No
2017			MONENSIN SENSITIVITY 1 (MON1)/CALCIU M CAFFEINE ZINC SENSITIVITY 1 (CCZ1)-mediated Rab7 activation regulates tapetal programmed cell death and pollen development. Plant Physiology 173(1):206-218.	No	Yes	Yes	No

9. Recognized international conference(s) in which paper(s) related to this research project was/were delivered (*Please attach a copy of each delivered paper. All listed papers must acknowledge RGC's funding support by quoting the specific grant reference.*)

Month/Year/	Title	Conference Name	Submitted	Attached	Acknowledged	Accessible
Place			to RGC	to this	the support of	from the
			(indicate the			institutional
			year ending	(Yes or No)	Research	repository
			of the		Scheme	(Yes or No)
			relevant		(Yes or No)	
			progress report)			

June 2013 Suzhou, China	function of COPII	Cold Spring Harbor Asia - Plant Cell and Developmental Biology	2014	Yes	Yes	No
June 2013 Suzhou, China	Endomembran	Cold Spring Harbor Asia - Plant Cell and Developmental Biology	2014	Yes	Yes	No
June 2013 Suzhou, China	Golgi Retention Mechanisms of Endomembran e Membrane Proteins (EMPs) in Eukaryotes		2014	Yes	Yes	No
Apr 2015 Shenzhen, China	MON1-CCZ1	CSCB The Biennial Conference and the 15th Congress of the Chinese Society for Cell Biology	No	Yes	Yes	No
Apr 2015 Shenzhen, China	Mon1 Repressor Mutant		No	Yes	Yes	No
Aug 2016 France	European Network for Plant Endomembran	Prevacuolar compartments contribute to the vacuole biogenesis in Arabidopsis root cortex and epidermis	No	Yes	Yes	No

10. Student(s) trained (*Please attach a copy of the title page of the thesis.*)

Name	Degree registered for	Date of registration	Date of thesis
			submission/
			graduation

NSFC/RGC 8 (Revised 10/15)

Zhuang Xiaohong	PhD	1/8/2008	1/8/2013
Shen Jinbo	PhD	1/8/2009	1/8/2013
Cui Yong	PhD	1/8/2011	1/9/2014
Lin Youshun	MPhil	1/8/2012	31/12/2014
Joanne WOO	MPhil	1/8/2012	1/6/2015
Lai Ching Man	MPhil	1/8/2012	1/3/2015
Wang Xiangfeng	PhD	3/8/2009	1/6/2015
Wang Juan	PhD	1/8/2010	1/12/2015
Zeng Yonglun	PhD	12/7/2010	1/1/2016

11. Other impact (e.g. award of patents or prizes, collaboration with other research institutions, *technology transfer, etc.*)