# Curriculum Vitae – Mei-Chun Cheng (Nina)

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**Research interests:** Plant stress and light biology; stress and light signal; translation mechanisms during stress and light perception, the regulation of stress-responsive genes and their roles in stress resistance.

### EDUCATION

06/2015	<b>Ph.D.</b> in Plant Biology, National Taiwan University, Taiwan Dissertation: The Arabidopsis ERF1 regulates abiotic-stress-responsive gene expression by binding to different cis-acting elements under different stress signaling
06/2008	<b>M.S.</b> in Plant Biology, National Taiwan University, Taiwan Thesis: Functional study of a drought-induced transcription factor, At2g20880, in Arabidopsis
06/2006	B.S. in Life Science, National Taiwan University, Taiwan

## SKILLS

- Advanced laboratory experience in molecular biology (plant transformation, gel electrophoresis, primer design, molecular cloning, PCR, RT-PCR, real-time RT-PCR, cDNA, N/S/Western blotting, GUS staining, Yeast one/two-hybrid screening, SDS-PAGE, protoplasts isolation and transfection, florescence microscopy, protein purification, Co-IP, ChIP assay, ubiquitination assay, EMSA, polysome profiling), genetics (crossing, mutant screening, map-based cloning) and genomics (gene expression profiling, knowledge of bioinformatic tools and statistical methods to integrate microarray data, promoter elements analysis, knock-out mutants characterization).
- Outstanding creativity and problem-solving capacities. Possess a great ability to combine relevant technical expertise and basic biology to challenge existing concepts. Superior time management, work well under pressure and on multiple projects. Supervised others.
- Thoroughly trained in writing and used to peer-review materials for publication. Good presentation skills. Excellent interpersonal and communication skills, very much at ease in international environments. Team player.
- Languages: Chinese (native), English (fluent), German (conversational).

### **RESEARCH EXPERIENCE**

- 02/20 present **Assistant Professor** in Department of Biochemical Science and Technology, National Taiwan University.
- 10/17 12/19 **Postdoctoral Fellow** in laboratory of Dr. Enamul Huq, Department of Molecular Biosciences, University of Texas at Austin. Regulation of PCH1/L degradation under dark by COP1 and its function in phyB dark reversion.
- 06/15 09/17 **Postdoctoral Fellow** in laboratory of Dr. Tsan-Piao Lin, Institute of Plant Biology, National Taiwan University. Identification and characterization of ERF1 interacting proteins. Discovered a mechanism that modulates the stability of ERF1 under light/dark cycle and regulates stress response.
- 08/11 06/15 **Ph.D. Student** in laboratory of Dr. Tsan-Piao Lin, Institute of Plant Biology, National Taiwan University. Dissertation: The Arabidopsis ERF1 regulates abiotic stress-responsive gene expression by binding to different cis-acting elements under different stress signaling. Studied the effect of GSH on the translational change.

- 07/08 07/11 **Research assistant** in laboratory of Dr. Tsan-Piao Lin, Institute of Plant Biology, National Taiwan University, Taiwan. Identified RGLG2 as an E3 ligase that mediates the ubiquitination of AtERF53 and negatively regulates abiotic stress response.
- 08/06 06/08 **M.S. Student** in laboratory of Dr. Tsan-Piao Lin, Institute of Plant Biology, National Taiwan University. Thesis: Functional study of a drought-induced transcription factor, At2g20880 (AtERF53), in Arabidopsis

## PUBLICATION

- 1. Cheng, M.C., Kathare, P.K., Paik, I., Huq, E. (2021) Phytochrome signaling network. Annual Review of Plant Biology (accepted contribution)
- Cheng, M.C., Enderle, B., Kathare, P.K., Hiltbrunner, A., Huq, E. (2020) PCH1 and PCHL directly interact with PIF1, promote its degradation and inhibit its transcriptional function during photomorphogenesis. Molecular Plant doi: https://doi.org/10.1016/j.molp.2020.02.003.
- 3. Cheng, M.C., Wang, Y.M., Kuo, W.C., Lin, T.P. (2017) UBC18 mediates ERF1 degradation under light-dark cycles. New Phytologist 213: 1156-1167.
- 4. Chen, H.Y., Hsieh, E.J., **Cheng, M.C.,** Chen, C.Y., Hwang, S.Y., Lin, T.P. (2016) ORA47 (octadecanoid-responsive AP2/ERF-domain transcription factor 47) regulates jasmonic acid and abscisic acid biosynthesis and signaling through binding to a novel cis-element. **New Phytologist** 211: 599-613.
- Cheng, M.C., Ko, K., Chang, W.L., Kuo, W.C., Chen, G.H., Lin, T.P. (2015). Increased level of glutathione contributes to stress tolerances and global translational change in Arabidopsis. Plant Journal 83: 926-939.
- Cheng, M.C., Liao, P.M., Kuo, W.W., Lin, T.P. (2013) The Arabidopsis ETHYLENE RESPONSE FACTOR1 regulates abiotic stress-responsive gene expression by binding to different cis-acting elements in response to different stress signals. Plant Physiology 162: 1566-1582.
- 7. Hsieh, E.J., Cheng, M.C., Lin, T.P. (2013) Functional characterization of an abiotic stress-inducible transcription factor AtERF53 in Arabidopsis thaliana. Plant Molecular Biology 82: 223-237.
- 8. **Cheng, M.C.,** Hsieh, E.J., Chen, J.H., Chen, H.Y., Tsan-Piao Lin, T.P. (2012). Arabidopsis RGLG2, functioning as a RING E3 ligase, interacts with AtERF53 and negatively regulates the plant drought stress response. **Plant Physiology** 158: 363-375.

#### AWARDS & HONORS

•	MOST Young Scholar Fellowship – Einstein project	2019	
•	MOST Postdoctoral Research Abroad Program	2016	
•	Ministry of Education Travel Award	2014	
National Taiwan University			
•	Departmental Scholarship	2013	
•	Departmental Scholarship	2012	
•	Dean Award	2008	

### **CONFERENCE ABSTRACT**

1. Cheng, M.C., Enderle, B., Kathare, P.K., Islam, R., Hiltbrunner, A., Huq, E. (2019) PCH1 and PCHL regulate light responses by interacting with PIF1 and mediate its degradation in Arabidopsis. American Society of Plant Biology, San Jose

- 2. Cheng, M.C., Ko, K., Chang, W.L., Kuo, W.C., Chen, G.H., Lin, T.P. (2016). Increased level of glutathione contributes to stress tolerances and global translational change in Arabidopsis. International Conference of Arabidopsis Research, Gyeonju. (Poster talk)
- **3.** Cheng, M.C., Hsieh, E.J., Lin, T.P. (2013) Functional study of Arabidopsis ERF1 in response to abiotic stress. Root system biology symposium, Taipei.
- 4. **Cheng, M.C.,** Hsieh, E.J., Chen, J.H., Chen, H.Y., Lin, T.P. (2012). Arabidopsis RGLG2, functioning as a RING E3 ligase, interacts with AtERF53 and negatively regulates the plant drought stress response. International Congress on Plant Molecular Biology, Jeju.
- 5. **Cheng, M.C.,** Hsieh, E.J., Chen, J.H., Chen, H.Y., Lin, T.P. (2012). Arabidopsis RGLG2, functioning as a RING E3 ligase, interacts with AtERF53 and negatively regulates the plant drought stress response. 2012 Annual meeting of society of experimental biology, Salzburg
- Cheng, M.C., Hsieh, E.J., Chen, J.H., Chen, H.Y., Lin, T.P. (2012). Functional study of a drought-induced transcription factor, At2g20880 (AtERF53), in Arabidopsis. International Symposium on the Kanagawa University – National Taiwan University, Kanagawa.

# INVITED TALKS

- 1. Plant biotechnology and green industry symposium. From data to publication: Stories of ERF transcription factors. Jul. 2016
- 2. National Sun Yat-Sen University. The Regulation of ETHYLENE RESPONSE FACTOR (ERF) in Abiotic Stress Response. Mar. 2016
- 4<sup>th</sup> Cross-strait conference of plant biology. The Arabidopsis ETHYLENE RESPONSE FACTOR1 regulates abiotic stress-responsive gene expression by binding to different cis-acting elements in response to different stress signals. Jun. 2013

# TEACHING EXPERIENCE

Summer School Invited Lecturer

- IPB summer school on Plant Molecular Biology (NTU) Summer 2014

**Teaching Assistant** 

- NTU: Proteomics Fall 2013
- NTU: Plant hormone analysis (Experiment) Spring 2012
- –NTU: Plant stress biology Fall 2007

# REFEREES

Prof. Tsan-Piao Lin Institute of Plant Biology, National Taiwan University tpl@ntu.edu.tw +886-2-33662537

Prof. Shih-Tong Jeng

Institute of Plant Biology, National Taiwan University stjeng@ntu.edu.tw +886-2-33662538

**Prof. Enamul Huq** Department of Molecular Biosciences, University of Texas at Austin huq@austin.utexas.edu