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Education

2003-2010 Ph.D Shanghai Institute of Plant Physiology and Ecology, Chinese Academy of Sciences.

1999-2003 B.S Nanjing Agriculture University, China

Experience

2011.6-Present

Plant Systems Group, Oak Ridge National Lab

Postdoc Researcher

- Genomic studies in Crassulacean Acid Metabolism species.
- Characterization of small proteins in response to plant-microbe interaction.

2010.7-2011.6

Institute of Subtropical Forestry, Chinese Academy of Forestry

Research Assistant

- Functional analysis of ABC genes in *Camellia japonica*.

Publication

Chu-Yu Ye, Ting Li, **Hengfu Yin**, David J. Weston, Gerald A Tuskan, Timothy Tschaplinski, and Xiaohan Yang. Non-family genes will play important roles in evolutionary and functional genomics research. (in review)

Hengfu Yin, Peng Gao, Chengwu Liu, Jun Yang, Zhongchi Liu, Xiaohu Hu, and Da Luo. Control of stem development via PS synthesis genes in rice.(in review)

Wang, L., **Yin, H.**, Qian, Q., Yang, J., Huang, C., Hu, X. and Luo, D. (2009). NECK LEAF 1, a GATA type transcription factor, modulates organogenesis by regulating the expression of multiple regulatory genes during reproductive development in rice. *Cell Res.*

Luo, A.,* Qian, Q.,* **Yin, H.**,* Liu, X., Yin, C., Lan, Y., Tang, J., Tang, Z., Cao, S., Wang, X. et al. (2006). EUI1, encoding a putative cytochrome P450 monooxygenase, regulates internode elongation by modulating gibberellin responses in rice. *Plant Cell Physiol* 47, 181-91.(*equal contribution)