Xiong Jinsong



Areas of Research Plant disease resistance

Contact Information

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Research Interests

Plant disease resistance

Our research mainly focuses on functional genomics by the use of strawberry as model system. We are particularly interested in the following aspect of strawberry:

Disease resistance

Plants are sessile organisms that are constantly under the threaten of potential virulent microbes. Among these microbes, Botrytis cinerea, the cause of grey mold disease, is one of the most disastrous pathogenic fungus to strawberry. However, the defense mechanisms of strawberry to B. cinerea are largely unknown. We utilize the approaches of molecular biology, biochemistry and genetics to identification and characterization key genes that play roles in disease resistance, to dissect the molecular defense mechanisms against fungal pathogens in strawberry.

Education Background

Bachelor: Hunan University of Arts and Science

Master: Sun Yat-Sen University **Doctor:** Sun Yat-Sen University

Work experience

Postdoctoral research, Institute of Genetics and Developmental Biology, Chinese Academy of Sciences, 2010.07-2013.09

Lecturer, Nanjing Agricultural University, 2013.11-2019.11

Associate Professor, Nanjing Agricultural University, 2019.12-

Selected Publication

Xiong Jinsong*, Bai Yibo, Ma Chuangju, Zhu Hongyu, Zheng Dan, Cheng Zongming*. Molecular Cloning and Characterization of SQUAMOSA-Promoter Binding Protein-Like Gene FvSPL10 from Woodland Strawberry (Fragaria vesca). *Plants*, 2019, 8 (9), 342.

Liu Hui, **Xiong Jinsong**, Jiang Yueting, Wang Li, Cheng Zongming. Evolution of the R2R3-MYB gene family in six Rosaceae species and expression in woodland strawberry. *Journal of Integrative Agriculture*, 2019, 18 (12): 2753-2770.

Shao Gaoneng, Lu Zefu, **Xiong Jinsong**, Wang Bing, Jing Yanhui, Meng Xiangbing, Liu Guifu, Ma Haiyan, Liang Yan, Chen Fan, Wang Yonghong, Li Jiayang, Yu Hong. Tiller Bud Formation Regulators MOC1 and MOC3 Cooperatively Promote Tiller Bud Outgrowth by Activating FON1 Expression in Rice. *Molecular Plant*, 2019, 12 (8): 1090-1102.

Xiong Jinsong, Zheng Dan, Zhu Hongyu, Chen Jianqiu, Na Ran, Cheng Zongming. Genome-wide identification and expression analysis of the SPL gene family in woodland strawberry Fragaria vesca. *Genome*, 2018, 61 (9): 675-683.

Xiong Jinsong#, Zhu Hongyu#, Bai Yibo, Liu Hui, Cheng Zongming. RNA sequencing-based transcriptome analysis of mature strawberry fruit infected by necrotrophic fungal pathogen Botrytis cinerea. *Physiological and Molecular Plant Pathology*, 2018, 104: 77–85.

Cai Jie, Zhang Lanyue, Liu Wei, Tian Ye, **Xiong Jinsong**, Wang Yihan, Li Rujie, Li Haoming, Wen Jiangqi, Mysore Kirankumar S, Boller Thomas, Xie Zhiping, Staehelin Christian. Role of the Nod Factor Hydrolase MtNFH1 in Regulating Nod Factor Levels during Rhizobial Infection and in Mature Nodules of Medicago truncatula. *The Plant Cell*, 2018, 30: 397-414.

Song Xiaoguang#, Lu Zefu#, Yu Hong#, Shao Gaoneng, **Xiong Jinsong**, Meng Xiangbing, Jing Yanhui, Liu Guifu, Xiong Guosheng, Duan Jingbo, Yao Xuefeng, Liu Chunming, Li Hongqi, Wang Yonghong, Li Jiayang. IPA1 functions as a downstream

transcription factor repressed by D53 in strigolactone signaling in rice. *Cell Research*, 2017, 27(9): 1128–1141.

Liu Jinyi, Chen Nana, Cheng Zongming*, **Xiong Jinsong***. Genome-wide identification, annotation and expression profile analysis of SnRK2 gene family in grapevine. *Australian Journal of Grape and Wine Research*, 2016, 22: 478–488.

Xiong Jinsong*, Ding Jing, Li Yi*. Genome-editing technologies and their potential application in horticultural crop breeding. *Horticulture Research*, 2015, 2, 15019.

Wang Xiaolong, Zhong Yan, Cheng Zongming, **Xiong Jinsong***. Divergence of the bZIP Gene Family in Strawberry, Peach, and Apple Suggests Multiple Modes of Gene Evolution after Duplication. *International Journal of Genomics*, 2015, 2015, 536943.