



乔 鑫

职 称：副教授，硕士生导师

邮 箱：qiaoxin@njau.edu.cn

联系电话：025-84396485

办公地址：生科楼 A6009

### 研究方向：

1. 果树基因组学与遗传育种（主要围绕梨树开展研究）
2. 生物信息学与组学大数据挖掘
3. 植物基因组进化规律研究，基因组多倍化和重复基因进化机制解析

### 教育经历：

2012. 09 – 2018. 12	南京农业大学，果树学，农学博士学位
2015. 10 – 2016. 10	美国佐治亚大学，国家公派联合培养博士
2008. 09 – 2012. 07	山西农业大学，园艺，农学学士学位

### 工作经历：

2019. 05 – 今	南京农业大学园艺学院，副教授，硕导
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### 代表性科研成果：

2020

Zeng W<sup>†</sup>, **Qiao Xin<sup>†</sup>**, Li Q, Liu C, Wu J, Yin H, Zhang S: Genome-wide identification and comparative analysis of the ADH gene family in Chinese white pear (*Pyrus bretschneideri*) and other Rosaceae species. **Genomics** 2020, 112(5):3484-3496.

Liu C<sup>†</sup>, **Qiao Xin<sup>†</sup>**, Li Q, Zeng W, Wei S, Wang X, Chen Y, Wu X, Wu J, Yin H et al: Genome-wide comparative analysis of the BAHD superfamily in seven Rosaceae species and expression analysis in pear (*Pyrus bretschneideri*). **BMC Plant Biology** 2020, 20(1):14.

## 2019

**Qiao Xin**, Li Q, Yin H, Qi K, Li L, Wang R, Zhang S\* and Paterson AH\*: Gene duplication and evolution in recurring polyploidization-diploidization cycles in plants. **Genome Biology** 2019, 20:38.  
Li Q<sup>†</sup>, **Qiao Xin<sup>†</sup>**, Yin H, Zhou Y, Dong H, Qi K, Li L, Zhang S\*: Unbiased subgenome evolution following recent whole-genome duplication in pear (*Pyrus bretschneideri*). **Horticulture Research** 2019, 6:34.

## 2018

**Qiao Xin**, Yin H, Li L, Wang R, Wu J, Wu J, Zhang S\*: Different Modes of Gene Duplication Show Divergent Evolutionary Patterns and Contribute Differently to the Expansion of Gene Families Involved in Important Fruit Traits in Pear (*Pyrus bretschneideri*). **Frontiers in Plant Science** 2018, 9(161).

## 2015

**Qiao Xin**, Li M, Li L, Yin H, Wu J, Zhang S\*: Genome-wide identification and comparative analysis of the heat shock transcription factor family in Chinese white pear (*Pyrus bretschneideri*) and five other Rosaceae species. **BMC Plant Biology** 2015, 15(1):12.

## 2014

乔鑫, 李梦, 殷豪, 李雷廷, 吴俊, 张绍铃\*: 果树全基因组测序研究进展. **园艺学报** 2014, 41(1):165-177.