

浙江省科学技术奖公示信息表（单位提名）

提名奖项：自然科学奖

成果名称	杂草基因组及其与作物协同进化机制研究
提名等级	一等奖
提名书 相关内容 (附表)	见附表
主要完成人	<p>樊龙江，排名 1，教授，浙江大学；</p> <p>陆永良，排名 2，研究员，中国水稻研究所；</p> <p>叶楚玉，排名 3，教授，浙江大学；</p> <p>郭龙彪，排名 4，研究员，中国水稻研究所；</p> <p>柏连阳，排名 5，教授，湖南省农业科学院；</p>
主要完成单位	<p>1. 浙江大学</p> <p>2. 中国水稻研究所</p> <p>3. 湖南省农业科学院</p> 
提名单位	浙江大学
提名意见	<p>农业杂草与作物之间就有着密切关系，研究杂草对作物生产具有重要意义。申报人聚焦稻田杂草稗草和杂草稻，在杂草基因组特征的解析，杂草与水稻互作机制的阐明，杂草起源演化规律的揭示等方面开展工作，取得了系列原创性成果。包括获得稗属高质量基因组，解析稗属多倍化基因组演化机制，阐明稗草拟态遗传机制，揭示稗草抑制水稻分子机制，弄清杂草稻起源方式。研究结果为杂草绿色防控、抑草水稻品种培育提供重要理论指导，推动了杂草遗传资源挖掘利用，也丰富了作物起源演化认知。相关研究结果在 Nature 子刊、PNAS、Genome Biology、Trends in Plant Science、Molecular Plant 等发表多篇论文，产生了广泛的国际影响。</p> <p>提名该成果为省自然科学奖一等奖。</p>

代表性论文专著目录

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2. Mao Lingfeng, Hiroshi Kawaide, Sho Miyazaki, Meihong Chen, Yoshiki Hirata, Honoka Kimura, Toshiya Higuchi, Miyu Teruya, Kaoru Fujiwara, Keisuke Tomita, Koji Miyamoto, Hisakazu Yamane, Ken-ichiro Hayashi, Hideaki Nojiri, Lei Jia, Jie Qiu, Chuyu Ye, Michael P. Timko, Longjiang Fan* and Kazunori Okada*. 2020. Genomic evidence for convergent evolution of gene clusters for momilactone biosynthesis in land plants. *PNAS*, doi:<https://doi.org/10.1073/pnas.1914373117>
3. Qiu Jie, Lei Jia, Dongya Wu, Xifang Weng, Lijuan Chen, Jian Sun, Meihong Chen, Lingfeng Mao, Bowen Jiang, Chuyu Ye, Guilherme Turra, Longbiao Guo, Guoyou Ye, Qianhao Zhu, Toshiyuki Imaizumi, Beng-Kah Song, Laura Scarabel, Aldo Merotto, Kenneth Olsen*, Longjiang Fan*. 2020. Diverse genetic mechanisms underlie worldwide convergent rice feralization. *Genome Biology*, 21:70-81
4. Ye Chu-Yu, Wei Tang, Dongya Wu, Lei Jia, Jie Qiu, Meihong Chen, Lingfeng Mao, Feng Lin, Haiming Xu, Xiaoyue Yu, Yongliang Lu, Yonghong Wang, Kenneth M Olsen, Michael P. Timko, Longjiang Fan*. 2019. Genomic evidence of human selection on Vavilovian mimicry. *Nature Ecology & Evolution*, 3:1474-1482
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7. Qiu Jie, Yongjun Zhou, Lingfeng Mao, Yongyi Yu, Chuyu Ye, Jianping Zhang, Weidi Wang, Fei Fu, Yunfei Wang, Feijian Qian, Ting Qi, Sanling Wu, Ya-Nan Cao, Yu Wang, Michael P. Timko, Song Ge, Longjiang Fan*, Yongliang Lu*. 2017. Genomic variation associated with local adaptation of weedy rice during de-domestication. *Nature Communications*, 8: 15323. doi:10.1038/ncomms15323
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