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Correspondences

- 1961 **The main restorer *Rf3* of maize S type cytoplasmic male sterility encodes a PPR protein that functions in reduction of the transcripts of *orf355***
Xiner Qin, Shike Tian, Wenliang Zhang, Qi Zheng, Hao Wang, Yang Feng, Yanan Lin, Jihua Tang, Yi Wang, Jianbing Yan, Mingqiu Dai, Yonglian Zheng, and Bing Yue
- 1965 **WheatOmics: A platform combining multiple omics data to accelerate functional genomics studies in wheat**
Shengwei Ma, Meng Wang, Jianhui Wu, Weilong Guo, Yongming Chen, Guangwei Li, Yanpeng Wang, Weiming Shi, Guangmin Xia, Daolin Fu, Zhensheng Kang, and Fei Ni

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- 1969 **Toward haplotype studies in polyploid plants to assist breeding**
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- 1973 **Phosphoinositides: Emerging players in plant salinity stress tolerance**
Sergey Shabala and Min Yu
- 1976 **Ubiquitination and PARYlation cross-talk about immunity**
Marco Trujillo
- 1979 **Fine-tuning ROS homeostasis by ROD1 is a battleground between rice and *Magnaporthe oryzae***
Yoji Kawano
- 1982 **TMK: A crucial piece of the acid growth puzzle**
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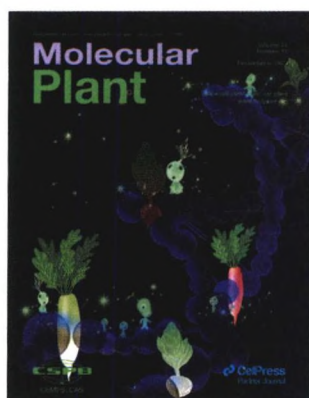
Resource Article

- 1985 **A single-cell morpho-transcriptomic map of brassinosteroid action in the *Arabidopsis* root**
Moritz Graeff, Surbhi Rana, Jos R. Wendrich, Julien Dorier, Thomas Eekhout, Ana Cecilia Aliaga Fandino, Nicolas Guex, George W. Bassel, Bert De Rybel, and Christian S. Hardtke

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- 2000 **Dynamic changes of phosphatidylinositol and phosphatidylinositol 4-phosphate levels modulate H⁺-ATPase and Na⁺/H⁺ antiporter activities to maintain ion homeostasis in *Arabidopsis* under salt stress**
Yongqing Yang, Xiuli Han, Liang Ma, Yujiao Wu, Xiao Liu, Haiqi Fu, Guoyong Liu, Xiaoguang Lei, and Yan Guo
- 2015 **An angiosperm NLR Atlas reveals that NLR gene reduction is associated with ecological specialization and signal transduction component deletion**
Yang Liu, Zhen Zeng, Yan-Mei Zhang, Qian Li, Xing-Mei Jiang, Zhen Jiang, Ji-Hong Tang, Dijun Chen, Qiang Wang, Jian-Qun Chen, and Zhu-Qing Shao
- 2032 **Pan-genome of *Raphanus* highlights genetic variation and introgression among domesticated, wild, and weedy radishes**
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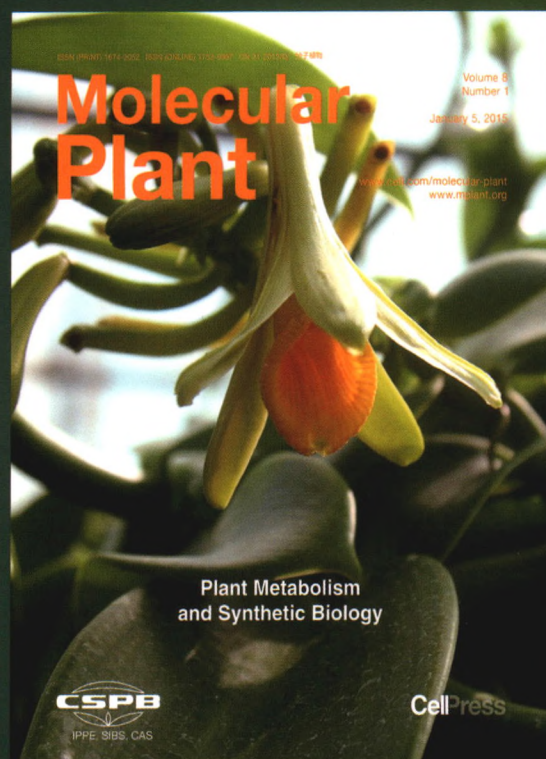
- 2056 Cytoplasmic and nuclear genome variations of rice hybrids and their parents inform the trajectory and strategy of hybrid rice breeding**
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- 2072 The BZR1-EDS1 module regulates plant growth-defense coordination**
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- 2088 Coordinated regulation of plant immunity by poly(ADP-ribosylation) and K63-linked ubiquitination**
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On The Cover

Radish pan-genome, which represents a genus-level pan-genome of *Raphanus*, provides new insights into genome evolution underlying post-polyploid diploidization and lays the foundation for genetic improvement of radish crops, biological control of weeds, and conservation of wild germplasms. The radish pan-genome was constructed through *de novo* genome assemblies of eleven accessions covering most of typical sub-species and varieties of domesticated, wild and weedy radishes from East Asia, South Asia, Europe and America. The cover image illustrates that the pan-genome is made up of the genomes of diverse radish accessions. The elves symbolize the chromosome-associated proteins that play important roles in DNA replication, DNA repair, chromatin modifications, and genome integrity maintenance. Image by: Jing Chen.

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