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A super pan-genomic landscape of rice
Revealing why depression compromises antiviral immunity
Arc weakens synapses by modulating PSD phase separation
Female sterility mutation enables mechanized hybrid rice breeding



Center for Excellence in Molecular Cell Science
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SPRINGER NATURE

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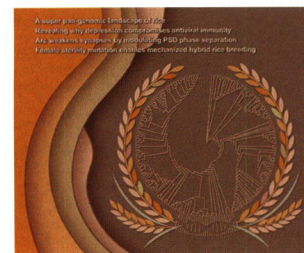
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RESEARCH HIGHLIGHTS

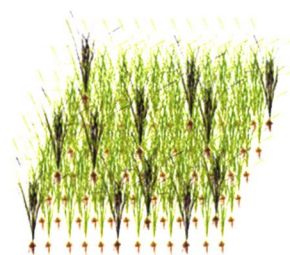
- 867 The rice pangenome branches out**
Kenneth M. Olsen
- 869 AHI1: linking depression and impaired antiviral immune response**
Weili Yang, Shihua Li, Xiao-Jiang Li
- 871 Inducing Z-DNA overcomes immune checkpoint blockade resistance**
Huipeng Jiao, Manolis Pasparakis
- 873 Profiling spatial gene activity in marmoset embryos**
Guangdun Peng, Patrick P. L. Tam
- 875 Structure-based evolutionary relationship between IscB and Cas9**
Ning Jia, Dinshaw J. Patel

ARTICLES

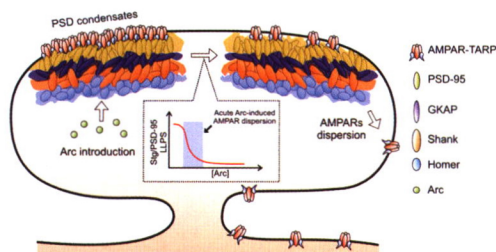
- 878 A super pan-genomic landscape of rice Open**
Liangang Shang, Xiaoxia Li, Huiying He, Qiaoling Yuan, Yanni Song, Zhaoran Wei, Hai Lin, Min Hu, Fengli Zhao, Chao Zhang, Yuhua Li, Hongsheng Gao, Tianyi Wang, Xiangpei Liu, Hong Zhang, Ya Zhang, Shuaimin Cao, Xiaoman Yu, Bintao Zhang, Yong Zhang, Yiqing Tan, Mao Qin, Cheng Ai, Yingxue Yang, Bin Zhang, Zhiqiang Hu, Hongru Wang, Yang Lv, Yuexing Wang, Jie Ma, Quan Wang, Hongwei Lu, Zhe Wu, Shanlin Liu, Zongyi Sun, Hongliang Zhang, Longbiao Guo, Zichao Li, Yongfeng Zhou, Jiayang Li, Zuofeng Zhu, Guosheng Xiong, Jue Ruan, Qian Qian
- 897 Depression compromises antiviral innate immunity via the AVP-AHI1-Tyk2 axis**
Hong-Guang Zhang, Bin Wang, Yong Yang, Xuan Liu, Junjie Wang, Ning Xin, Shifeng Li, Ying Miao, Qiuyu Wu, Tingting Guo, Yukang Yuan, Yibo Zuo, Xiangjie Chen, Tengfei Ren, Chunsheng Dong, Jun Wang, Hang Ruan, Miao Sun, Xingshun Xu, Hui Zheng
- 914 Arc weakens synapses by dispersing AMPA receptors from postsynaptic density via modulating PSD phase separation**
Xudong Chen, Bowen Jia, Yoichi Araki, Bian Liu, Fei Ye, Richard Huganir, Mingjie Zhang



Cover: The phylogenetic tree covering 251 different varieties of rice forms a badge together with the ear of rice consisting of rice grains in different colors. This badge pattern is the epitome of the work that rice super pan-genome from large natural populations captures genetic diversity and reveals genomic complexity. Meanwhile, it can be seen as a tribute to our ancestors who created the iconic terraces (illustrated in paper cut style) in rice farming civilization for their wisdom and diligence. Cover art is contributed by Zisong Ma. See page 878–896 by Liangang Shang et al. for details.



The lack of self-seed production from the restorer line *tfs1* (female-sterile line) enables direct harvesting of hybrid seeds from the field. See page 931–945 by Haoxuan Li et al. for details.



A model depicting that rise of Arc concentration can sensitively disperse AMPAR clustering via reversing phase separation of Stg and PSD-95. See page 914–930 by Xudong Chen et al. for details.

- 931 A spontaneous thermo-sensitive female sterility mutation in rice enables fully mechanized hybrid breeding**
Haoxuan Li, Chenjiang You, Manabu Yoshikawa, Xiaoyu Yang, Haiyong Gu, Chuanguo Li, Jie Cui, Xuemei Chen, Nenghui Ye, Jianhua Zhang, Guanqun Wang

LETTERS TO THE EDITOR

- 946 Structural basis for pore blockade of human voltage-gated calcium channel $\text{Ca}_v1.3$ by motion sickness drug cinnarizine**
Open
Xia Yao, Shuai Gao, Nieng Yan

- 949 Omicron-specific mRNA vaccine elicits potent immune responses in mice, hamsters, and nonhuman primates**
Yi Wu, Yanqiong Shen, Namei Wu, Xinghai Zhang, Shaohong Chen, Chang Yang, Junhui Zhou, Yan Wu, Da Chen, Li Wang, Chao Wang, Huajun Zhang, Nimuo Xia, Sandra Chiu, Yucai Wang
- 953 Retractosomes: small extracellular vesicles generated from broken-off retraction fibers**
Yizheng Wang, Shuaixin Gao, Yuheng Liu, Dongju Wang, Boqi Liu, Dong Jiang, Catherine C. L. Wong, Yang Chen, Li Yu

CORRECTION

- 957 Author Correction: Cryo-EM structure of the AVP–vasopressin receptor 2–Gs signaling complex**
Lei Wang, Jun Xu, Sheng Cao, Dapeng Sun, Heng Liu, Qiuyuan Lu, Zheng Liu, Yang Du and Cheng Zhang

ADVANCE ONLINE PUBLICATION

CRISPR FISHer enables high-sensitivity imaging of nonrepetitive DNA in living cells through phase separation-mediated signal amplification

Xin-Yuan Lyu, Yuan Deng, Xiao-Yan Huang, Zhen-Zhen Li, Guo-Qing Fang, Dong Yang, Feng-Liu Wang, Wang Kang, En-Zhi Shen and Chun-Qing Song

A novel protein RASON encoded by a lncRNA controls oncogenic RAS signaling in KRAS mutant cancers

Open
Rongjie Cheng, Fanying Li, Maolei Zhang, Xin Xia, Jianzhuang Wu, Xinya Gao, Huangkai Zhou, Zhi Zhang, Nunu Huang, Xuesong Yang, Yaliang Zhang, Shunli Shen, Tiebang Kang, Zexian Liu, Feizhe Xiao, Hongwei Yao, Jianbo Xu, Chao Yan and Nu Zhang

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Cell Discov. 2022 Jun 6;8(1):53. doi: 10.1038/s41421-022-00419-w.

Dynamic O-GlcNAcylation coordinates ferritinophagy and mitophagy to activate ferroptosis
Cell Discov. 2022 May 3;8(1):40. doi: 10.1038/s41421-022-00390-6.

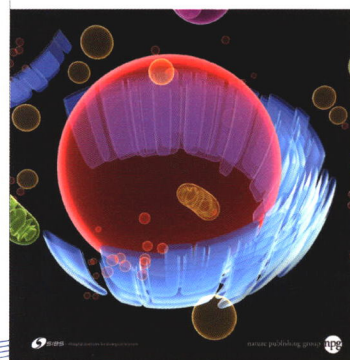
Treatment of SARS-CoV-2-induced pneumonia with NAD⁺ and NMN in two mouse models
Cell Discov. 2022 Apr 29;8(1):38. doi: 10.1038/s41421-022-00409-y.

CRISPR signal conductor 2.0 for redirecting cellular information flow
Cell Discov. 2022 Mar 15;8(1):26. doi: 10.1038/s41421-021-00371-1.

Cross-species metabolomic analysis identifies uridine as a potent regeneration promoting factor
Cell Discov. 2022 Feb 1;8(1):6. doi: 10.1038/s41421-021-00361-3.

Histone lysine methacrylation is a dynamic post-translational modification regulated by HAT1 and SIRT2
Cell Discov. 2021 Dec 28;7(1):122. doi: 10.1038/s41421-021-00344-4.

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