On the Cover: Weedy or red rice (*Oryza sativa*) infests cultivated rice fields worldwide, reducing yields and causing great economic losses. In recent years, use of herbicide resistant (HR) rice cultivars known as Clearfield rice has increased greatly in the United States, providing an option to control weedy rice using imidazolinone herbicides. However, the impact of HR cultivated rice on the evolutionary dynamics of U.S. weedy rice has not been explored. In this issue, Burgos et al. (pp. 1208–1220) characterize the morphology and population structure of weedy rice that has acquired the herbicide resistance trait, sampled from cultivated fields with a history of Clearfield use. Great morphological variation is seen in HR weedy rice, with some plants exhibiting reduced weediness in terms of reduced seed dormancy, later flowering, and reduced seed shattering. A significant portion of the local rice crop genome is introgressed into HR weedy rice, though crop-weed introgression occurred very rarely prior to Clearfield use. Whereas the majority of weedy rice in the United States prior to Clearfield use had straw-colored hulls, black-colored hulls predominate among contemporary HR weedy rice. Genotyping-by-sequencing data confirm that the evolution of HR weedy rice is a consequence of preferential hybridization between cultivated rice and historical black-hull weedy populations. The cover shows a cultivated rice field in Arkansas with at least two seasons of Clearfield use, which is being sampled for escaped weedy rice plants. The inset is a straw-hull weedy rice plant sampled from the field. Cover image credits: David Gealy, Nilda Burgos (photography) and Ana Caicedo (design).
Use of Multicopy Transposons Bearing Unfitness Genes in Weed Control: Four Example Scenarios.
Jonathan Gressel and Avraham A. Levy

Multicopy transposons could disseminate potentially lethal/unfit genes in weed populations, especially when coupled with positive selection to facilitate rapid dissemination.

RESEARCH ARTICLES

Dual Function of the Cytochrome P450 CYP76 Family from Arabidopsis thaliana in the Metabolism of Monoterpenols and Phenylurea Herbicides. René Höfer, Benoît Boachon, Hugues Renault, Carole Gavira, Laurence Miesch, Juliana Iglesias, Jean-François Ginkelinger, Lionel Allouche, Michel Miesch, Sebastien Grec, Romain Larbat, and Danièle Werck-Reichhart

Fast diversification and versatility of a subfamily of cytochrome P450 enzymes in Brassicaceae has been important in their metabolism of both monoterpenols and herbicides.


A modified native promoter, dual protein localization, and an evolved desensitized maize protein variant enables field tolerance in soybean to multiple herbicides.

Indaziflam Herbicidal Action: A Potent Cellulose Biosynthesis Inhibitor. Chad Brabham, Lei Lei, Ying Gu, Jossef Stork, Michael Barrett, and Seth DeBolt

Indaziflam is a cellulose biosynthesis inhibitor that has a different mechanism of action than previously characterized inhibitors and causes an increase in CELLULOSE SYNTHASE A particle density and reduced, but not paused, velocity at the plasma membrane focal plane.

De Novo Assembly and Characterization of the Transcriptome of the Parasitic Weed Dodder Identifies Genes Associated with Plant Parasitism. Aashish Ranjan, Yasunori Ichihashi, Moran Farhi, Kristina Zumstein, Brad Townsley, Rakefet David-Schwartz, and Neelima R. Sinha

Transcriptional dynamics during parasitism in the parasitic weed Cuscuta pentagona reveals increased expression of genes encoding transporters and stimulus response regulators, and a decrease in the expression of genes encoding photosynthetic proteins.


Genes encoding enolpyruvylshikimate phosphate synthase are tandemly arranged on chromosomes of field-evolved glyphosate-resistant Kochia scoparia.


Use of herbicide-resistant rice cultivars in the United States has led to the emergence of herbicide-resistant weedy rice formed predominantly by hybridization of cultivars with historical weeds characterized by black hulls and awns.

Continued on next page
A novel rice cytochrome P450 monooxygenase is involved in multiple-herbicide detoxification and could be useful in herbicide development and molecular breeding in crops.

De novo genome assembly and genomic resources of horseweed will be useful to understand the genetic and molecular bases of weediness.

Rapid vacuole sequestration, the glyphosate resistance mechanism observed in glyphosate-resistant horseweed, is associated with plasma membrane and tonoplast ATP-binding cassette transporters.

Mapping-by-sequencing identifies a naturally occurring variant of a kinesin that causes root hairs to branch.

Efficient gene editing in tomato in the first generation using the clustered regularly interspaced short palindromic repeats/CRISPR-associated9 system. The CRISPR/Cas9 system is highly efficient at generating targeted mutations in stable transgenic tomato plants, and homozygous deletions of a desired size can be created in the first generation.
RESEARCH ARTICLES

BIOCHEMISTRY AND METABOLISM

Pepper Mitochondrial FORMATE DEHYDROGENASE1 Regulates Cell Death and Defense Responses against Bacterial Pathogens.  
Du Seok Choi, Nak Hyun Kim, and Byung Kook Hwang

Formate dehydrogenase contributes to defense-related hormone regulation and defense gene activation, ultimately leading to hypersensitive cell death and defense response to bacterial pathogens.

Mitochondrial Alternative Oxidase Maintains Respiration and Preserves Photosynthetic Capacity during Moderate Drought in Nicotiana tabacum.  
Keshav Dahal, Jia Wang, Greg D. Martyn, Farkhunda Rahimy, and Greg C. Vanlerbergh

Alternative oxidase reduces feedback on the photosynthetic apparatus that can otherwise limit photosynthetic capacity during moderate drought.

Seed-Specific Expression of a Feedback-Insensitive Form of CYSTATHIONINE-γ-SYNTHASE in Arabidopsis Stimulates Metabolic and Transcriptomic Responses Associated with Desiccation Stress.  
Hagai Cohen, Hadasa Israeli, Ifat Matityahu, and Rachel Amir

Methionine metabolism in Arabidopsis seeds is strongly associated with stress-related metabolites and transcripts, suggesting that its content leads to desiccation stress during seed development.

CELL BIOLOGY

Heterodimeric Capping Protein from Arabidopsis Is a Membrane-Associated, Actin-Binding Protein.  
Jose C. Jimenez-Lopez, Xia Wang, Simeon O. Kotchoni, Shanjin Huang, Daniel B. Szymanski, and Christopher J. Staiger

A key regulator of the actin cytoskeleton is shown to be associated with endosomal membranes.

Peroxisomal Ubiquitin-Protein Ligases Peroxin2 and Peroxin10 Have Distinct But Synergistic Roles in Matrix Protein Import and Peroxin5 Retrotranslocation in Arabidopsis.  
Sarah E. Burkhart, Yun-Ting Kao, and Bonnie Bartel

Two Arabidopsis membrane peroxins contribute to peroxisomal matrix protein import via distinct roles in recycling and degrading a matrix receptor protein.

Association of the P6 Protein of Cauliflower mosaic virus with Plasmodesmata and Plasmodesmal Proteins.  
Andres Rodriguez, Carlos A. Angel, Lindy Lutz, Scott M. Leisner, Richard S. Nelson, and James E. Schoelz

Inclusion bodies of Cauliflower mosaic virus function in delivery of virions to the plasmodesma for transport to adjacent cells.

Chao Cai, Jessica L. Henty-Ridilla, Daniel B. Szymanski, and Christopher J. Staiger

A myosin triple knockout mutant has altered cytoskeletal organization and single filament turnover.

ECOPHYSIOLOGY AND SUSTAINABILITY

Transient Silencing of CHALCONE SYNTHASE during Fruit Ripening Modifies Tomato Epidermal Cells and Cuticle Properties.  
Laura España, José A. Heredia-Guerrero, José J. Reina-Pinto, Rafael Fernández-Muñoz, Antonio Heredia, and Eva Dominguez

Alteration of flavonoid accumulation during tomato fruit ripening affects cuticle wax accumulation, biomechanics, water permeability, and the molecular arrangement of cuticle components.
Capturing Arabidopsis Root Architecture Dynamics with ROOT-FIT Reveals Diversity in Responses to Salinity. Magdalena M. Julkowska, Huub C.J. Hoefsloot, Selena Mol, Richard Feron, Gert-Jan de Boer, Michel A. Haring, and Christa Testerink

Natural variation in the dynamic response to salt stress of the Arabidopsis root system can be divided between four distinct strategies.

Variation in Sulfur and Selenium Accumulation Is Controlled by Naturally Occurring Isoforms of the Key Sulfur Assimilation Enzyme ADENOSINE 5'-PHOSPHOSULFATE REDUCTASE2 across the Arabidopsis Species Range. Dai-Yin Chao, Patrycja Baraniecka, John Danku, Anna Koprivova, Brett Lahner, Hongbing Luo, Elena Yakubova, Brian Dilkes, Stanislav Kopriva, and David E. Salt

The ability of Arabidopsis to assimilate and accumulate sulfur varies widely across the species range and is controlled in part by diversity in a key reductase enzyme.

GENES, DEVELOPMENT, AND EVOLUTION

Neofunctionalization of Duplicated Tic40 Genes Caused a Gain-of-Function Variation Related to Male Fertility in Brassica oleracea Lineages. Xiaoling Dun, Wenhao Shen, Kaining Hu, Zhengfu Zhou, Shengqian Xia, Jing Wen, Bin Yi, Jinxiong Shen, Chaozhi Ma, Jinxing Tu, Tingdong Fu, and Ulf Lagercrantz

Divergent duplicated genes in the Brassica oleracea gain a novel function that could restore the fertility of a Brassica napus male-sterile mutant.

MEMBRANES, TRANSPORT, AND BIOENERGETICS

The Mitochondrial Protein Import Component, TRANSLOCASE OF THE INNER MEMBRANE17-1, Plays a Role in Defining the Timing of Germination in Arabidopsis. Yan Wang, Simon R. Lau, Aneta Ivanova, Olivier van Aken, Szymon Kabiszewski-Jakubiak, Vindya Uggalla, Margaretha van der Merwe, Owen Duncan, Reena Narisai, James Whelan, and Monika W. Murcha

A mitochondrial import protein determines the timing of mitochondrial biogenesis during germination.

Inducible Knockdown of MONOGALACTOSYLDIACYLGLYCEROL SYNTHASE1 Reveals Roles of Galactolipids in Organelle Differentiation in Arabidopsis Cotyledons. Sho Fujii, Koichi Kobayashi, Yuki Nakamura, and Hajime Wada

Biosynthesis of the major thylakoid membrane lipids coordinates the differentiation of chloroplasts with peroxisomes and mitochondria during the greening of Arabidopsis cotyledons.

Elucidating the Role of Transport Processes in Leaf Glucosinolate Distribution. Svend Roesen Madsen, Carl Erik Olsen, Hussam Hassan Sour-Eldin, and Barbara Ann Halkier

Within-leaf allocation of glucosinolates implicates an intracellular route from site of biosynthesis to peripheral leaf layers.

Grain setting defect1, Encoding a Remorin Protein, Affects the Grain Setting in Rice through Regulating Plasmodesmatal Conductance. Jinshan Gui, Chang Liu, Junhui Shen, and Laigeng Li

A remorin protein plays a role in regulating photoassimilate translocation by modulating plasmodesmata permeability in rice.


Silencing aquaporins of the xylem bundle sheath cells reduces leaf hydraulic conductance and affects the water permeability of adjacent mesophyll cells, suggesting a bundle sheath-mesophyll feed-forward hydraulic control.
A family of drugs reduces the growth response, the lipid modifications, and the pattern of gene expression triggered by inorganic phosphate starvation.


A zinc deficiency transcription factor moderates oxidative stress resistance.

An Arabidopsis receptor-like kinase negatively regulates abscisic acid signaling and promotes infection by filamentous (hemi)biotrophs.

An Arabidopsis phosphotyrosyl phosphatase activator is essential for assembly of a protein phosphatase and plays important roles in hormone signaling, salt stress response, and plant development.

Phytochrome-mediated phototropic enhancement depends on PINOID family kinases.

An orthologous pair of genes from barley and Arabidopsis is involved in long-chain fatty acid signaling that is required for germination of conidia from distantly related powdery mildews.

A drought stress-inducible receptor-like kinase is important for plant growth and grain yield in rice.

Two Arabidopsis genes with a role in thermotolerance are oriented in a head-to-head manner in the genome and the intergenic region between these genes functions as a heat-inducible bidirectional promoter.
SYSTEMS AND SYNTHETIC BIOLOGY

[OPEN] Systematic Phenotypic Screen of Arabidopsis Peroxisomal Mutants Identifies Proteins Involved in β-Oxidation. Gaëlle Cassin-Ross and Jianping Hu

Recently discovered Arabidopsis peroxisomal mutants identify proteins involved in β-oxidation. 1546


A cell type- and leaf tissue-specific model provides new insights into nitrogen metabolism in the maize leaf. 1659

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