

On the Cover: Plant cell growth is constrained by cell walls that glue neighboring cells together, yet the growth of these cells within a tissue can be highly variable. To understand this, Tauriello et al. (pp. 2342–2358) observed and analyzed dividing cells in live *Arabidopsis* (*Arabidopsis thaliana*) sepals. They tracked cell lineages at 6-hour intervals to extract growth curves for all lineages. These growth curves showed varying growth rates in space and time. The constant property was that all cell lineages reached the same maximum growth rate relative to their size. Thus, despite the observed variability, individual cell lineages follow similarly shaped growth curves. The cover image shows cell lineages on the abaxial epidermis of an *Arabidopsis* sepal grown for 66 h. Cell outlines are marked with blue dots. Each cell and all of its daughters in a single lineage have the same color. Cell lineages are superimposed on plasma membranes and nuclei (in gray). Cover image credits: Gerardo Tauriello (ETH Zürich), Heather M. Meyer (Cornell University), and Adrienne H.K. Roeder (Cornell University).

ON THE INSIDE

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COMMENTARY

A View to a Kill: Markers for Developmentally Regulated Cell Death in Plants. Alexis Maizel 2341

BREAKTHROUGH TECHNOLOGIES

^[OPEN]Variability and Constancy in Cellular Growth of *Arabidopsis* Sepals. Gerardo Tauriello, Heather M. Meyer, Richard S. Smith, Petros Koumoutsakos, and Adrienne H.K. Roeder

The growth of cells in the developing sepal is highly variable, but all cells exhibit a surprisingly common, asynchronous maximum growth rate relative to their size. 2342

^[OPEN]phenoVein—A Tool for Leaf Vein Segmentation and Analysis. Jonas Bühler, Louai Rishmawi, Daniel Pflugfelder, Gregor Huber, Hanno Scharr, Martin Hülskamp, Maarten Koornneef, Ulrich Schurr, and Siegfried Jahnke

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Ethylene: Traffic Controller on Hormonal Crossroads to Defense. Colette Broekgaarden, Lotte Caarls, Irene A. Vos, Corné M.J. Pieterse, and Saskia C.M. Van Wees

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Ethylene Control of Fruit Ripening: Revisiting the Complex Network of Transcriptional Regulation. Mingchun Liu, Julien Pirrello, Christian Chervin, Jean-Paul Roustan, and Mondher Bouzayen

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RESEARCH ARTICLES

BIOCHEMISTRY AND METABOLISM

[OPEN] An Intracellular Laccase Is Responsible for Epicatechin-Mediated Anthocyanin Degradation in Litchi Fruit Pericarp. Fang Fang, Xue-lian Zhang, Hong-hui Luo, Jia-jian Zhou, Yi-hui Gong, Wen-jun Li, Zhao-wan Shi, Quan He, Qing Wu, Lu Li, Lin-lin Jiang, Zhi-gao Cai, Michal Oren-Shamir, Zhao-qi Zhang, and Xue-qun Pang

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[OPEN] Four Isoforms of Arabidopsis 4-Coumarate:CoA Ligase Have Overlapping yet Distinct Roles in Phenylpropanoid Metabolism. Yi Li, Jeong Im Kim, Len Pysh, and Clint Chapple

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[OPEN] Highly Branched Xylan Made by IRREGULAR XYLEM14 and MUCILAGE-RELATED21 Links Mucilage to Arabidopsis Seeds. Cătălin Voiniciuc, Markus Günl, Maximilian Heinrich-Wilhelm Schmidt, and Björn Usadel

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[OPEN] The Arabidopsis Class III Peroxidase AtPRX71 Negatively Regulates Growth under Physiological Conditions and in Response to Cell Wall Damage. Sara Raggi, Alberto Ferrarini, Massimo Delledonne, Christophe Dunand, Philippe Ranocha, Giulia De Lorenzo, Felice Cervone, and Simone Ferrari

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[OPEN] RNA Silencing of Exocyst Genes in the Stigma Impairs the Acceptance of Compatible Pollen in Arabidopsis. Darya Safavian, Yara Zayed, Emily Indriolo, Laura Chapman, Abdalla Ahmed, and Daphne R. Goring

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Iron stress modulates auxin and ethylene pathways to regulate lateral root formation in Arabidopsis. 2608

[^{OPEN}] The Interplay between Sulfur and Iron Nutrition in Tomato. *Sabrina Zuchi, Mutsumi Watanabe, Hans-Michael Hubberten, Mariusz Bromke, Sonia Osorio, Alisdair R. Fernie, Silvia Celletti, Anna Rita Paolacci, Giulio Catarcione, Mario Ciaffi, Rainer Hoefgen, and Stefania Astolfi*

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[^{OPEN}] GmEXPB2, a Cell Wall β -Expansin, Affects Soybean Nodulation through Modifying Root Architecture and Promoting Nodule Formation and Development. *Xinxin Li, Jing Zhao, Zhiyuan Tan, Rensen Zeng, and Hong Liao*

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[^{OPEN}] Metabolite Profiles of Maize Leaves in Drought, Heat, and Combined Stress Field Trials Reveal the Relationship between Metabolism and Grain Yield. *Toshihiro Obata, Sandra Witt, Jan Liseč, Natalia Palacios-Rojas, Igor Florez-Sarasa, Salima Yousfi, Jose Luis Araus, Jill E. Cairns, and Alisdair R. Fernie*

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[^{OPEN}] Polyploidy Enhances F₁ Pollen Sterility Loci Interactions That Increase Meiosis Abnormalities and Pollen Sterility in Autotetraploid Rice. *Jinwen Wu, Muhammad Qasim Shahid, Lin Chen, Zhixiong Chen, Lan Wang, Xiangdong Liu, and Yonggen Lu*

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A phylogenetically conserved group of transcription factors form trimers to regulate root nodule development and early gene expression during the legume-Rhizobium symbiosis.

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MEMBRANES, TRANSPORT, AND BIOENERGETICS

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[^{OPEN}]Phosphorylation of the Light-Harvesting Complex II Isoform Lhcb2 Is Central to State Transitions. *Paolo Longoni, Damien Douchi, Federica Cariti, Geoffrey Fucile, and Michel Goldschmidt-Clermont*

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Leaf herbivore attack changes free and conjugated phenolic acids in maize roots and thereby triggers an avoidance response in a specialist root herbivore. 2884

[^{OPEN}]TaFROG Encodes a Pooideae Orphan Protein That Interacts with SnRK1 and Enhances Resistance to the Mycotoxigenic Fungus *Fusarium graminearum*. *Alexandre Perochon, Jia Jianguang, Amal Kahla, Chanemougasoundharam Arunachalam, Steven R. Scofield, Sarah Bowden, Emma Wallington, and Fiona M. Doohan*

A Pooideae orphan protein that enhances plant resistance against a toxigenic fungus physically interacts with a key signaling protein. 2895

The Rice Transcription Factor WRKY53 Suppresses Herbivore-Induced Defenses by Acting as a Negative Feedback Modulator of Mitogen-Activated Protein Kinase Activity. *Lingfei Hu, Meng Ye, Ran Li, Tongfang Zhang, Guoxin Zhou, Qi Wang, Jing Lu, and Yonggen Lou*

A transcription factor functions as a negative feedback modulator of two mitogen-activated protein kinases and thereby acts as an early suppressor of herbivore-induced defenses in rice.

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[OPEN]Short Hypocotyl in White Light1 Interacts with Elongated Hypocotyl5 (HY5) and Constitutive Photomorphogenic1 (COP1) and Promotes COP1-Mediated Degradation of HY5 during Arabidopsis Seedling Development. *Anjil Kumar Srivastava, Dhirodatta Senapati, Archana Srivastava, Moumita Chakraborty, Sreeramaiah N. Gangappa, and Sudip Chattopadhyay*

A unique regulator of proteasome-mediated degradation of photomorphogenesis-promoting factors affects Arabidopsis seedling development.

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[OPEN]OsMADS26 Negatively Regulates Resistance to Pathogens and Drought Tolerance in Rice. *Giang Ngan Khong, Pratap Kumar Pati, Frédérique Richaud, Boris Parizot, Przemyslaw Bidzinski, Chung Duc Mai, Martine Bès, Isabelle Bourrié, Donald Meynard, Tom Beeckman, Michael Gomez Selvaraj, Ishitani Manabu, Anna-Maria Genga, Christophe Brugidou, Vinh Nang Do, Emmanuel Guiderdoni, Jean-Benoit Morel, and Pascal Gantet*

A MADS-box transcription factor represses resistance to pathogenic microorganisms and water deficit, and its down-regulation results in improved biotic and abiotic stress tolerance of rice.

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[OPEN]PBL13 Is a Serine/Threonine Protein Kinase That Negatively Regulates Arabidopsis Immune Responses. *Zuh-Jyh Daniel Lin, Thomas W.H. Liebrand, Koste A. Yadeta, and Gitta Coaker*

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[OPEN]Glutathione Regulates 1-Aminocyclopropane-1-Carboxylate Synthase Transcription via WRKY33 and 1-Aminocyclopropane-1-Carboxylate Oxidase by Modulating Messenger RNA Stability to Induce Ethylene Synthesis during Stress. *Riddhi Datta, Deepak Kumar, Asma Sultana, Saptarshi Hazra, Dipto Bhattacharyya, and Sharmila Chattopadhyay*

Glutathione induces ethylene biosynthesis by enhancing ACC synthase transcription and increasing the mRNA stability of ACC oxidase during stress.

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SYSTEMS AND SYNTHETIC BIOLOGY

[OPEN]Evolution of Cis-Regulatory Elements and Regulatory Networks in Duplicated Genes of Arabidopsis. *Andrej A. Arsoovski, Julian Pradinuk, Xu Qiu Guo, Sishuo Wang, and Keith L. Adams*

Cis-regulatory elements and regulatory networks have diverged considerably in duplicated genes.

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Engineering Monolignol *p*-Coumarate Conjugates into Poplar and Arabidopsis Lignins. *Rebecca A. Smith, Eliana Gonzales-Vigil, Steven D. Karlen, Ji-Young Park, Fachuang Lu, Curtis G. Wilkerson, Lacey Samuels, John Ralph, and Shawn D. Mansfield*

Novel monolignol p-coumarate conjugates in eudicot lignin result from the introduction of a monocot acyltransferase.

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[OPEN] Unraveling the Light-Specific Metabolic and Regulatory Signatures of Rice through Combined in Silico Modeling and Multiomics Analysis. *Meiyappan Lakshmanan, Sun-Hyung Lim, Bijayalaxmi Mohanty, Jae Kwang Kim, Sun-Hwa Ha, and Dong-Yup Lee*

Combined in silico modeling and multiomics data analysis elucidate the transcriptional control of rice cellular metabolism upon light signaling.

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Changes in the Phosphoproteome and Metabolome Link Early Signaling Events to Rearrangement of Photosynthesis and Central Metabolism in Salinity and Oxidative Stress Response in Arabidopsis. *Yanmei Chen and Wolfgang Hoehenwarter*

Rearrangement of central metabolism and photosynthesis is linked to the dynamic phosphoproteome in abiotic stress signaling in Arabidopsis.

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