FORTHCOMING PAPERS

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Phillip J. McCown, Keith A. Corbino, Shira Stav, Madeline E. Sherlock, and Ronald R. Breaker

Computational analysis of nascent peptides that induce ribosome stalling and their proteomic distribution in *Saccharomyces cerevisiae*
Renana Sabi and Tamir Tuller

Reports
DROSHA targets its own transcript to modulate alternative splicing
Dooyoung Lee, Jin-Wu Nam, and Chanseok Shin

Structure of an SMG8–SMG9 complex identifies a G-domain heterodimer in the NMD effector proteins
Liang Li, Mahesh Lingaraju, Claire Basquin, Jérome Basquin, and Elena Conti

Articles
A genetic screen implicates a CWC16/Yju2/CCDC130 protein and SMU1 in alternative splicing in *Arabidopsis thaliana*
Tatsuo Kanno, Wen-Dar Lin, Jason L. Fu, Antonius J.M. Matzke, and Marjori Matzke

LncATLAS database for subcellular localization of long noncoding RNAs
David Mas-Ponte, Joana Carlevaro-Fita, Emilio Palumbo, Toni Hermoso Pulido, Roderic Guigo, and Rory Johnson

Topological and thermodynamic factors that influence the evolution of small networks of catalytic RNA species
Jessica A.M. Yeates, Philippe Nghe, and Niles Lehman

Characterization of the mammalian DEAD-box protein DDX5 reveals functional conservation with *S. cerevisiae* ortholog Dbp2 in transcriptional control and glucose metabolism
Zheng Xing, Siwen Wang, and Elizabeth J. Tran