# Table of Contents

## Highlights of This Issue 955

### CHROMATIN, GENE, AND RNA REGULATION

957 A YAP/TAZ-Regulated Molecular Signature Is Associated with Oral Squamous Cell Carcinoma
- Samantha E. Hiemer, Liye Zhang, Vinay K. Kartha, Trevor S. Packer, Munirah Almershed, Vikki Noonan, Maria Kukuruzinska, Manish V. Bais, Stefano Monti, and Xaralabos Varelas

969 A Protein Interaction between β-Catenin and Dnmt1 Regulates Wnt Signaling and DNA Methylation in Colorectal Cancer Cells
- Jing Song, Zhanwen Du, Mate Ravaasz, Bohan Dong, Zhenghe Wang, and Rob M. Ewing

### DNA DAMAGE AND REPAIR

982 EphB1 Suppression in Acute Myelogenous Leukemia: Regulating the DNA Damage Control System

### GENOMICS

993 Homologous Mutation to Human BRAF V600E Is Common in Naturally Occurring Canine Bladder Cancer—Evidence for a Relevant Model System and Urine-Based Diagnostic Test
- Brennan Decker, Heidi G. Parker, Deepika Dhawan, Erika M. Kwon, Eric Karlins, Brian W. Davis, José A. Ramos-Vara, Patty L. Bonney, Elizabeth A. McNiel, Deborah W. Knapp, and Elaine A. Ostrander

### ONCOGENES AND TUMOR SUPPRESSORS

1003 AKT1 E17K in Colorectal Carcinoma Is Associated with BRAF V600E but Not MSI-H Status: A Clinicopathologic Comparison to PIK3CA Helical and Kinase Domain Mutants
- Jacyln F. Hechtman, Justyna Sadowska, Jason T. Huse, Laetitia Borsu, Rona Yaeger, Jinrut Shia, Efsevia Vakiani, Marc Ladanyi, and Maria E. Arcila

1009 Anti-miR-21 Suppresses Hepatocellular Carcinoma Growth via Broad Transcriptional Network Deregulation

1022 COX-2 Elevates Oncogenic miR-526b in Breast Cancer by EP4 Activation
- Mousumi Majumder, Erin Landman, Ling Liu, David Hess, and Peeyush K. Lala

### SIGNAL TRANSDUCTION

1034 The Role of Nitric Oxide Synthase Uncoupling in Tumor Progression

1044 Endophilin A2 Promotes TNBC Cell Invasion and Tumor Metastasis
- Tomas Baldassarre, Kathleen Watt, Peter Truedsell, Jaina Meens, Mark M. Schneider, Sandip K. Sengupta, and Andrew W. Craig

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ABOUT THE COVER

The cover shows a heat map of oral squamous cell carcinoma (OSCC) tumor expression data from "The Cancer Genome Atlas (TCGA)" projected onto the YAP/TAZ-activated transcriptional signature identified from microarray studies performed by Hiemer and colleagues (beginning on page 957). Hierarchical clustering identified two tumor-associated sub-clusters which were enriched in either cell-cycle genes or genes responding to cancer-related signaling pathways.
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13 (6)


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