### 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 Ravanaz, 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
  Jaclyn F. Hechtman, Justyna Sadowska, Jason T. Huse, Laetitia Borsu, Rona Yaeger, Jinru 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 Truesdell, Jalna 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.