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

SIGNAL TRANSDUCTION

1034 The Role of Nitric Oxide Synthase Uncoupling in Tumor Progression

1044 Endophilin A2 Promotes TNBC Cell Invasion and Tumor Metastasis
Tomas Baldasserre, Kathleen Watt, Peter Truesdell, Jalna Meens, Mark M. Schneider, Sandip K. Sengupta, and Andrew W. Craig
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.