## Table of Contents

### November 2017 • Volume 15 • Number 11

#### Highlights of This Issue

1467

### CELL DEATH AND SURVIVAL

1469  Targeting AR Variant–Coactivator Interactions to Exploit Prostate Cancer Vulnerabilities

Fiorella Magani, Stephanie O. Peacock, Meghan A. Rice, Maria J. Martinez, Ann M. Greene, Pablo S. Magani, Rolando Lyles, Jonathan R. Weitz, and Kerry L. Burnstein

1481  Chemoradiotherapy Resistance in Colorectal Cancer Cells is Mediated by Wnt/β-catenin Signaling

Georg Emons, Melanie Spitzner, Sebastian Reinke, Jannke Möller, Noam Auslander, Frank Kramer, Yue Hu, Tim Beissbarth, Hendrik A. Wolff, Margret Rave-Frank, Elisabeth Hefmann, Jochen Gaedcke, B. Michael Ghadimi, Steven A. Johnsen, Thomas Ried, and Marian Grade

### CHROMATIN, EPIGENETICS, AND RNA REGULATION

1491  Dual Src Kinase/Pretubulin Inhibitor KX-01, Sensitizes ERα-negative Breast Cancers to Tamoxifen through ERα Reexpression

Muralidharan Anbalagan, Mei Sheng, Brian Fleischer, Yifang Zhang, Yuanjun Gao, Van Hoang, Margarite Matossian, Hope E. Burks, Matthew E. Burow, Bridgette M. Collins-Burow, David Hangauer, and Brian G. Rowan

### DNA DAMAGE AND REPAIR

1503  EGFR Mutations Compromise Hypoxia-Associated Radiation Resistance through Impaired Replication Fork–Associated DNA Damage Repair


### METABOLISM

1517  EWS/FLI is a Master Regulator of Metabolic Reprogramming in Ewing Sarcoma

Jason M. Tanner, Claire Bensard, Peng Wei, Nathan M. Krah, John C. Schell, Jamie Gardiner, Joshua Schiffman, Stephen L. Lessnick, and Jared Rutter

### GENOMICS

1531  CDK4/6 Inhibition on Glucose and Pancreatic Beta Cell Homeostasis in Young and Aged Rats

Aida I. Sacaan, Stephane Thibault, Miyoung Hong, Nagesha G. Kondegowda, Tim Nichols, Rosemary Li, Carolina Rosselot, Winston Evering, Rafael Fenuitria, Allison Vitsky, Thomas Brown, Martin Finkelstein, Adolfo García-Ocaña, Nasir Khan, Andrew F. Stewart, and Rupangi C. Varavada

1542  Comprehensive Genomic Profiling of Metastatic Squamous Cell Carcinoma of the Anal Canal

Van Morris, Xiayu Rao, Curtis Pickering, Wai Chin Foo, Asif Rashid, Karina Eterovic, Taebekim Kim, Ken Chen, Jing Wang, Kenna Shaw, and Cathy Eng

1551  Comprehensive Molecular Profiling of Olfactory Neuroblastoma Identifies Potentially Targetable FGFR3 Amplifications

Lorena Lazo de la Vega, Jonathan B. McHugh, Andi K. Cani, Komal Kunder, Frances M. Walocko, Chia-Jen Liu, Daniel H. Hovelson, Dan Robinson, Arul M. Chinnaiyan, Scott A. Tomlins, and Paul W. Harms

1558  Novel Aberrations Uncovered in Barrett’s Esophagus and Esophageal Adenocarcinoma Using Whole Transcriptome Sequencing


### ONCOGENES AND TUMOR SUPPRESSORS

1570  miR-432 Induces NRF2 Stabilization by Directly Targeting KEAP1

Burak Akdemir, Yasuaki Nakajima, Johji Inazawa, and Jun Inoue

1579  Nuclear Envelope Rupture Is Enhanced by Loss of p53 or Rb

Zhe Yang, John Maciejowski, and Titia de Lange

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SELECTIVE MET KINASE INHIBITION IN MET-DEPENDENT GLIOMA MODELS ALTERS GENE EXPRESSION AND INDUCES TUMOR PLASTICITY

EARLY-STAGE METASTASIS REQUIRES MDM2 AND NOT P53 GAIN OF FUNCTION

ADAM12 IS A NOVEL REGULATOR OF TUMOR ANGIOGENESIS VIA STAT3 SIGNALING
Roopali Roy, Adelle Dagher, Catherine Butterfield, and Marsha A. Moses

HEPARAN SULFATE GLYCOSAMINOGLYCANs IN GLIOBLASTOMA PROMOTE TUMOR INVASION

ABOUT THE COVER
The cover image depicts chromosomal copy alterations detected by next generation sequencing in olfactory neuroblastomas—rare, highly aggressive tumors of the sinonasal cavity. In this issue, Lazo de la Vega and colleagues (beginning on page 1551) demonstrate that olfactory neuroblastomas are characterized by recurrent chromosomal numerical alterations, including amplifications of the potential oncogenic driver FGFR3 in a subset. The finding suggests that FGFR3 may be a rational therapeutic target in olfactory neuroblastoma.