Phase I Expansion and Pharmacodynamic Study of the Oral MEK Inhibitor RO4987655 (CH4987655) in Selected Patients with Advanced Cancer with RAS–RAF Mutations

Acquired Resistance to Endocrine Treatments Is Associated with Tumor-Specific Molecular Changes in Patient-Derived Luminal Breast Cancer Xenografts
Paul Cotto, Ivan Bièche, Franck Assayag, Rania El Botty, Sophie Chatseau-Joubert, Aurélie Thuleau, Thomas Bagarre, Benoît Albaud, Audrey Rapinat, David Gentien, Pierre de la Grange, Vonick Sibut, Sophie Vacher, Rana Hatem, Jean-Luc Servely, Jean-Jacques Fontaine, Didier Decaudin, Jean-Yves Pierga, Sérgio Roman-Roman, and Elisabetta Marangoni

Multifactorial T-cell Hypofunction That Is Reversible Can Limit the Efficacy of Chimeric Antigen Receptor–Transduced Human T cells in Solid Tumors

Sonic Hedgehog Paracrine Signaling Activates Stromal Cells to Promote Perineural Invasion in Pancreatic Cancer
Xuqi Li, Zheng Wang, Qingyong Ma, Qinhong Xu, Han Liu, Wanxing Duan, Jianjun Lei, Jinguang Ma, Xiu Wang, Shifang Lv, Liang Han, Wei Li, Jian Guo, Kun Guo, Dong Zhang, Erxi Wu, and Keping Xie

Functional Kinomics Identifies Candidate Therapeutic Targets in Head and Neck Cancer
Russell Moser, Chang Xu, Michael Kao, James Annis, Luisa Angelica Lerma, Christopher M. Schaupp, Kay E. Gurley, In Sock Jang, Asel Biktasova, Wendell G. Yarbrough, Adam A. Margolin, Carla Grandori, Christopher J. Kemp, and Eduardo Mendez
See related commentary, p. 4173

Intraperitoneal Oxidative Stress in Rabbits with Papillomavirus-Associated Head and Neck Cancer Induces Tumoricidal Immune Response That Is Adoptively Transferable
Annette Rossmann, Robert Mandic, Jochen Heinis, Helmut Hoffken, Oliver Kissner, Ralf Kinscherf, Eberhard Weihe, and Michael Bette

Targeting Cancer Cells via the Reactive Oxygen Species-Mediated Unfolded Protein Response with a Novel Synthetic Polyphenol Conjugate
Soon Young Shin, Jong Min Lee, Mi So Lee, Dongsoo Koh, Hyeryoun Jung, Yoongho Lim, and Young Han Lee

Kidney Cancer Is Characterized by Aberrant Methylation of Tissue-Specific Enhancers That Are Prognostic for Overall Survival
Caroline Y. Hu, Davoud Mohtat, Yiting Yu, Yi-An Ko, Niraj Shenoy, Sanchari Bhattacharya, Maria C. Izquierdo, Ae Seo Deok Park, Orsolya Gírizc, Nishanth Vallunselia, Krishna Gundabolu, Kristin Ware, Tushar D. Bhaagat, Masako Suzuki, James Pullman, X. Shirley Liu, John M. Greally, Katalin Susztak, and Amit Verma

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Molecular Testing for Lymph Node Metastases as a Determinant of Colon Cancer Recurrence: Results from a Retrospective Multicenter Study
A Novel KLF4/LDHA Signaling Pathway Regulates Aerobic Glycolysis in and Progression of Pancreatic Cancer
Min Shi, Juju Cui, Jiawei Du, Daoyan Wei, Zhiliang Jia, Jun Zhang, Zhenggang Zhu, Yong Gao, and Keping Xie

Preoperative GNAS and KRAS Testing in the Diagnosis of Pancreatic Mucinous Cysts

Circulating CD4⁺ T Cells That Produce IL4 or IL17 When Stimulated by Melan-A but Not by NY-ESO-1 Have Negative Impacts on Survival of Patients with Stage IV Melanoma

PBX1 Is a Favorable Prognostic Biomarker as It Modulates 13-cis Retinoic Acid–Mediated Differentiation in Neuroblastoma
Nilay Shah, Jianjun Wang, Julia Selich-Anderson, Garrett Graham, Hasan Siddiqui, Xin Li, Javed Khan, and Jeffrey Toretsky

Correction: Interleukin-7 Mediates Selective Expansion of Tumor-Redirected Cytotoxic T Lymphocytes (CTLs) without Enhancement of Regulatory T-Cell Inhibition

Correction: A Phase I Trial of LY2510924, a CXCR4 Peptide Antagonist, in Patients with Advanced Cancer

Correction: Bim Polymorphisms: Influence on Function and Response to Treatment in Children with Acute Lymphoblastic Leukemia

ABOUT THE COVER
Integrative analysis of clear cell renal cell carcinoma shows predominant hypermethylation. A circos plot shows alterations in DNA methylation (hypermethylation in blue, outer ring), copy number variations (amplifications in red and deletions in green, middle ring), and expression (underexpressed in green, overexpressed in orange, innermost ring) in renal cell carcinoma when compared. For details, see the article by Hu and colleagues on page 4349 of this issue.