

## Commentary

- Heterogeneity of Drug Target Expression Among Metastatic Lesions: Lessons from a Breast Cancer Autopsy Program.** Patricia S. Steeg .....3643

## CCR Practice of Translational Oncology

- Is There a Role for Circulating Tumor Cells in the Management of Breast Cancer?**  
Daniel F. Hayes and Jeffrey Smerage .....3646

## Molecular Pathways

- Advances in the Development of Cancer Therapeutics Directed against the RAS-Mitogen-Activated Protein Kinase Pathway.** Judith S. Sebolt-Leopold.....3651

## CCR Focus

- From the Editor.** Susan E. Bates .....3657

- Oncologic Phase 0 Trials Incorporating Clinical Pharmacodynamics: from Concept to Patient.** James H. Doroshow and Ralph E. Parchment.....3658

- The Development of Phase I Cancer Trial Methodologies: the Use of Pharmacokinetic and Pharmacodynamic End Points Sets the Scene for Phase 0 Cancer Clinical Trials.**  
A. Hilary Calvert and Ruth Plummer .....3664

- Leveraging Exploratory Investigational New Drug Studies to Accelerate Drug Development.**  
David Jacobson-Kram and George Mills .....3670

- Designing Phase 0 Cancer Clinical Trials.** Anthony J. Murgo, Shivaani Kummar, Larry Rubinstein, Martin Gutierrez, Jerry Collins, Robert Kinders, Ralph E. Parchment, Jiuping Ji, Seth M. Steinberg, Sherry X. Yang, Melinda Hollingshead, Alice Chen, Lee Helman, Robert Wilttrout, Joseph E. Tomaszewski, and James H. Doroshow .....3675

- Phase 0 Trials: An Industry Perspective.** Helen Eliopoulos, Vincent Giranda, Robert Carr, Rita Tiehen, Terri Leahy, and Gary Gordon .....3683

- Patient Perspectives on Phase 0 Clinical Trials.** Martin Gutierrez and Deborah Collyar.....3689

- The Ethics of Phase 0 Oncology Trials.** Emily Abdoler, Holly Taylor, and David Wendler.....3692

## Review

- Tissue-Based Research in Kidney Cancer: Current Challenges and Future Directions.**  
Sabina Signoretti, Gennady Bratslavsky, Frederick M. Waldman, Victor E. Reuter, John Haaga, Maria Merino, George V. Thomas, Michael R. Pins, Towia Libermann, John Gillespie, Joseph E. Tomaszewski, Carolyn C. Compton, Andrew Hruszkewycz, W. Marston Linehan, and Michael B. Atkins .....3699

## Human Cancer Biology

- T Regulatory Type 1 Cells in Squamous Cell Carcinoma of the Head and Neck: Mechanisms of Suppression and Expansion in Advanced Disease.** Christoph Bergmann, Laura Strauss, Yun Wang, Miroslaw J. Szczepanski, Stephan Lang, Jonas T. Johnson, and Theresa L. Whiteside .....3706

<b>Somatic Mutations of JAK1 and JAK3 in Acute Leukemias and Solid Cancers.</b> Eun Goo Jeong, Min Sung Kim, Hyo Kyung Nam, Chang Ki Min, Seok Lee, Yeun Jun Chung, Nam Jin Yoo, and Sug Hyung Lee .....	<b>3716</b>
<b>CIP2A Is Overexpressed in Gastric Cancer and Its Depletion Leads to Impaired Clonogenicity, Senescence, or Differentiation of Tumor Cells.</b> Wenjuan Li, Zheng Ge, Cheng Liu, Zhifang Liu, Magnus Björkholm, Jihui Jia, and Dawei Xu .....	<b>3722</b>
<b>Soluble ErbB3 Levels in Bone Marrow and Plasma of Men with Prostate Cancer.</b> Sue-Hwa Lin, Yu-Chen Lee, Michel B. Choueiri, Sijin Wen, Paul Mathew, Xiangcang Ye, Kim-Anh Do, Nora M. Navone, Jeri Kim, Shi-Ming Tu, Li-Yuan Yu-Lee, and Christopher J. Logothetis.....	<b>3729</b>
<b>A Metastasis Modifier Locus on Human Chromosome 8p in Uveal Melanoma Identified by Integrative Genomic Analysis.</b> Michael D. Onken, Lori A. Worley, and J. William Harbour.....	<b>3737</b>
<b>Association of p16 Homozygous Deletions with Clinicopathologic Characteristics and EGFR/KRAS/p53 Mutations in Lung Adenocarcinoma.</b> Reika Iwakawa, Takashi Kohno, Yoichi Anami, Masayuki Noguchi, Kenji Suzuki, Yoshihiro Matsuno, Kazuhiko Mishima, Ryo Nishikawa, Fumio Tashiro, and Jun Yokota .....	<b>3746</b>
<b>ssDNA-Binding Protein 2 Is Frequently Hypermethylated and Suppresses Cell Growth in Human Prostate Cancer.</b> Jun-Wei Liu, Jatin K. Nagpal, Wenye Sun, Juna Lee, Myoung Sook Kim, Kimberly L. Ostrow, Shaoyu Zhou, Carmen Jeronimo, Rui Henrique, Wim Van Criekinge, Chu So Moon, Joseph A. Califano, Barry Trink, and David Sidransky.....	<b>3754</b>
<b>BRCA1 and BRCA2 Mutation Prevalence and Clinical Characteristics of a Population-Based Series of Ovarian Cancer Cases from Denmark.</b> Marie Soegaard, Susanne Kruger Kjaer, Mark Cox, Eva Wozniak, Estrid Høgdall, Claus Høgdall, Jan Blaakaer, Ian J. Jacobs, Simon A. Gayther, and Susan J. Ramus .....	<b>3761</b>
<b>Down-Regulation of Hedgehog-Interacting Protein through Genetic and Epigenetic Alterations in Human Hepatocellular Carcinoma.</b> Motohisa Tada, Fumihiko Kanai, Yasuo Tanaka, Keisuke Tateishi, Miki Ohta, Yoshinari Asaoka, Motoko Seto, Ryosuke Muroyama, Kenichi Fukai, Fumio Imazeki, Takao Kawabe, Osamu Yokosuka, and Masao Omata .....	<b>3768</b>

## Imaging, Diagnosis, Prognosis

<b>Effect of Interleukin-10 Gene Polymorphisms on Clinical Outcome of Patients with Aggressive Non-Hodgkin's Lymphoma: An Exploratory Study.</b> Dieter Kube, Thanh-Duc Hua, Frederike von Bonin, Nils Schoof, Samira Zeynalova, Marita Klöss, Daniela Gocht, Bernd Potthoff, Mladen Tzvetkov, Jürgen Brockmöller, Markus Löffler, Michael Pfreundschuh, and Lorenz Trümper .....	<b>3777</b>
<b>Improved Prediction of Disease Relapse after Radical Prostatectomy through a Panel of Preoperative Blood-Based Biomarkers.</b> Shahrokh F. Shariat, Jose A. Karam, Jochen Walz, Claus G. Roehrborn, Francesco Montorsi, Vitaly Margulis, Fred Saad, Kevin M. Slawin, and Pierre I. Karakiewicz .....	<b>3785</b>
<b>Local Recurrence in Mismatch Repair–Proficient Colon Cancer Predicted by an Infiltrative Tumor Border and Lack of CD8<sup>+</sup> Tumor-Infiltrating Lymphocytes.</b> Inti Zlobec, Luigi M. Terracciano, and Alessandro Lugli .....	<b>3792</b>
<b>RHAMM, p21 Combined Phenotype Identifies Microsatellite Instability-High Colorectal Cancers with a Highly Adverse Prognosis.</b> Inti Zlobec, Kristi Baker, Luigi M. Terracciano, and Alessandro Lugli .....	<b>3798</b>
<b>Heterogeneous Ribonucleoprotein K and Thymidine Phosphorylase Are Independent Prognostic and Therapeutic Markers for Nasopharyngeal Carcinoma.</b> Lih-Chyang Chen, Chuen Hsueh, Ngan-Ming Tsang, Ying Liang, Kai-Ping Chang, Sheng-Po Hao, Jau-Song Yu, and Yu-Sun Chang .....	<b>3807</b>
<b>The Relative Distribution of Membranous and Cytoplasmic Met Is a Prognostic Indicator in Stage I and II Colon Cancer.</b> Fiona Ginty, Sudeshna Adak, Ali Can, Michael Gerdes, Melinda Larsen, Harvey Cline, Robert Filkins, Zhengyu Pang, Qing Li, and Michael C. Montalto .....	<b>3814</b>

<b>Association of <i>Megalyn</i> Genetic Polymorphisms with Prostate Cancer Risk and Prognosis.</b> Sarah K. Holt, Danielle M. Karyadi, Erika M. Kwon, Janet L. Stanford, Peter S. Nelson, and Elaine A. Ostrander .....	3823
<b>Imaging Immune Response <i>In vivo</i>: Cytolytic Action of Genetically Altered T Cells Directed to Glioblastoma Multiforme.</b> Jelena Lazovic, Michael C. Jensen, Evette Ferkassian, Brenda Aguilar, Andrew Raubitschek, and Russell E. Jacobs .....	3832
<b>Affibody Molecules for <i>In vivo</i> Characterization of HER2-Positive Tumors by Near-Infrared Imaging.</b> Sang Bong Lee, Moinuddin Hassan, Robert Fisher, Oleg Chertov, Victor Chernomordik, Gabriela Kramer-Marek, Amir Gandjbakhche, and Jacek Capala.....	3840
<b>Cytokeratin 10 and Cytokeratin 19: Predictive Markers for Poor Prognosis in Hepatocellular Carcinoma Patients after Curative Resection.</b> Xin-Rong Yang, Yang Xu, Guo-Ming Shi, Jia Fan, Jian Zhou, Yuan Ji, Hui-Chuan Sun, Shuang-Jian Qiu, Bing Yu, Qiang Gao, Yi-Zhou He, Weng-Zhen Qin, Rong-Xin Chen, Guo-Huan Yang, Bing Wu, Qing Lu, Zhi-Quan Wu, and Zhao-You Tang.....	3850

## Cancer Therapy: Clinical

<b>Are There Any Ethnic Differences in Molecular Predictors of Erlotinib Efficacy in Advanced Non-Small Cell Lung Cancer?</b> Myung-Ju Ahn, Byeong-Bae Park, Jin Seok Ahn, Sang We Kim, Heung-Tae Kim, Jong Seog Lee, Jin Hyung Kang, Jae Yong Cho, Hong Suk Song, Se Hoon Park, Chang Hak Sohn, Sang Won Shin, Jin Hyuck Choi, Chang-Seok Ki, Chan Keum Park, Alison J. Holmes, Pasi A. Jänne, and Keunchil Park.....	3860
<b>A Phase II Pharmacodynamic Study of Erlotinib in Patients with Advanced Non-Small Cell Lung Cancer Previously Treated with Platinum-Based Chemotherapy.</b> Enriqueta Felip, Federico Rojo, Martin Reck, Astrid Heller, Barbara Klughammer, Gemma Sala, Susana Cedres, Sergio Peralta, Heiko Maacke, Dorothee Foerzler, Marta Parera, Joachim Möcks, Cristina Saura, Ulrich Gatzemeier, and José Baselga .....	3867
<b>Interaction of Cisplatin with the Human Organic Cation Transporter 2.</b> Kelly K. Filipiski, Walter J. Loos, Jaap Verweij, and Alex Sparreboom.....	3875
<b>Dasatinib Cellular Uptake and Efflux in Chronic Myeloid Leukemia Cells: Therapeutic Implications.</b> Devendra K. Hiwase, Verity Saunders, Duncan Hewett, Amity Frede, Stephanie Zrim, Phuong Dang, Laura Eadie, L. Bik To, Junia Melo, Sharad Kumar, Timothy P. Hughes, and Deborah L. White .....	3881
<b>Phase I Study of GTI-2040, an Antisense to Ribonucleotide Reductase, in Combination with High-Dose Cytarabine in Patients with Acute Myeloid Leukemia.</b> Rebecca B. Klisovic, William Blum, Xiaohui Wei, Shujun Liu, Zhongfa Liu, Zhiliang Xie, Tamara Vukosavljevic, Cheryl Kefauver, LeNguyen Huynh, Jiuxia Pang, James A. Zwiebel, Steven Devine, John C. Byrd, Michael R. Grever, Kenneth Chan, and Guido Marcucci .....	3889
<b>Concerted Potent Humoral Immune Responses to Autoantigens Are Associated with Tumor Destruction and Favorable Clinical Outcomes without Autoimmunity.</b> Taylor Sittler, Jun Zhou, Joosang Park, Noah K. Yuen, Stefanie Sarantopoulos, Joseph Mollick, Ravi Salgia, Anita Giobbie-Hurder, Glenn Dranoff, and F. Stephen Hodi.....	3896
<b>Phase II Study of Dasatinib in Philadelphia Chromosome-Negative Acute and Chronic Myeloid Diseases, Including Systemic Mastocytosis.</b> Srdan Verstovsek, Ayalew Tefferi, Jorge Cortes, Susan O'Brien, Guillermo Garcia-Manero, Animesh Pardanani, Cem Akin, Stefan Faderl, Taghi Manshouri, Deborah Thomas, and Hagop Kantarjian.....	3906

## Cancer Therapy: Preclinical

<b>Selective Inhibition of BRCA2-Deficient Mammary Tumor Cell Growth by AZD2281 and Cisplatin.</b> Bastiaan Evers, Rinske Drost, Eva Schut, Michiel de Bruin, Eline van der Burg, Patrick W.B. Derksen, Henne Holstege, Xiaoling Liu, Ellen van Drunen, H. Berna Beverloo, Graeme C.M. Smith, Niall M.B. Martin, Alan Lau, Mark J. O'Connor, and Jos Jonkers .....	3916
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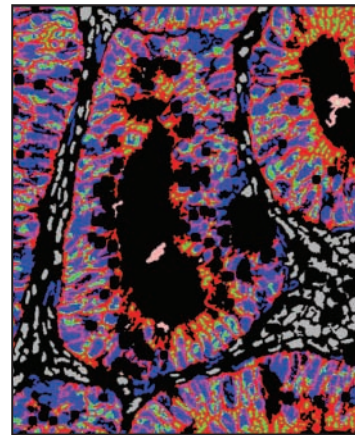
<b>Tumor Vascular Maturation and Improved Drug Delivery Induced by Methylselenocysteine Leads to Therapeutic Synergy with Anticancer Drugs.</b> Arup Bhattacharya, Mukund Seshadri, Steven D. Oven, Károly Tóth, Mary M. Vaughan, and Youcef M. Rustum.....	<b>3926</b>
<b>Intratumoral Coadministration of Hyaluronidase Enzyme and Oncolytic Adenoviruses Enhances Virus Potency in Metastatic Tumor Models.</b> Shanthi Ganesh, Melissa Gonzalez-Edick, Douglas Gibbons, Melinda Van Roey, and Karin Jooss.....	<b>3933</b>
<b>Oncogenic and Angiogenic Growth Factors Accumulate during Routine Storage of Apheresis Platelet Concentrates.</b> Julie Kanter, Samina Y. Khan, Marguerite Kelher, Lia Gore, and Christopher C. Silliman.....	<b>3942</b>
<b>RCAS/SCL-TVA Animal Model Allows Targeted Delivery of Polyoma Middle T Oncogene to Vascular Endothelial Progenitors <i>In vivo</i> and Results in Hemangioma Development.</b> Justin Sausville, Alfredo A. Molinolo, Xiangfei Cheng, Jon Frampton, Naoko Takebe, J. Silvio Gutkind, and Ricardo A. Feldman.....	<b>3948</b>
<b>Suppression of proHB-EGF Carboxy-Terminal Fragment Nuclear Translocation: A New Molecular Target Therapy for Gastric Cancer.</b> Takaya Shimura, Hiromi Kataoka, Naotaka Ogasawara, Eiji Kubota, Makoto Sasaki, Satoshi Tanida, and Takashi Joh.....	<b>3956</b>
<b>Transforming Growth Factor-<math>\beta</math> Receptor Blockade Augments the Effectiveness of Adoptive T-Cell Therapy of Established Solid Cancers.</b> Africa Wallace, Veena Kapoor, Jing Sun, Paul Mrass, Wolfgang Weninger, Daniel F. Heitjan, Carl June, Larry R. Kaiser, Leona E. Ling, and Steven M. Albelda.....	<b>3966</b>
<b>Intraoperative <math>^{186}\text{Re}</math>-Liposome Radionuclide Therapy in a Head and Neck Squamous Cell Carcinoma Xenograft Positive Surgical Margin Model.</b> Sean X. Wang, Ande Bao, Stephanie J. Herrera, William T. Phillips, Beth Goins, Cristina Santoyo, Frank R. Miller, and Randal A. Otto.....	<b>3975</b>
<b>DNA-Dependent Protein Kinase Is a Therapeutic Target and an Indicator of Poor Prognosis in B-Cell Chronic Lymphocytic Leukemia.</b> Elaine Willmore, Sarah L. Elliott, Tryfonia Mainou-Fowler, Geoffrey P. Summerfield, Graham H. Jackson, Fran O'Neill, Christopher Lowe, Anthony Carter, Robert Harris, Andrew R. Pettitt, Celine Cano-Soumillac, Roger J. Griffin, Ian G. Cowell, Caroline A. Austin, and Barbara W. Durkacz.....	<b>3984</b>
<b>PTEN Loss Does Not Predict for Response to RAD001 (Everolimus) in a Glioblastoma Orthotopic Xenograft Test Panel.</b> Lin Yang, Michelle J. Clarke, Brett L. Carlson, Ann C. Mladek, Mark A. Schroeder, Paul Decker, Wenting Wu, Gaspar J. Kitange, Patrick T. Grogan, Jennie M. Goble, Joon Uhm, Evanthia Galanis, Caterina Giannini, Heidi A. Lane, C. David James, and Jann N. Sarkaria.....	<b>3993</b>
<b>Increase in Brain Tumor Permeability in Glioma-Bearing Rats with Nitric Oxide Donors.</b> Dali Yin, Xiao Wang, Bindu M. Konda, John M. Ong, Jinwei Hu, Manuel R. Sacapano, MinHee K. Ko, Andres J. Espinoza, Dwain K. Irvin, Yan Shu, and Keith L. Black.....	<b>4002</b>

## Cancer Prevention and Susceptibility

<b>MDM2 Promoter Polymorphism and Pancreatic Cancer Risk and Prognosis.</b> Kofi Asomaning, Amy E. Reid, Wei Zhou, Rebecca S. Heist, Rihong Zhai, Li Su, Eunice L. Kwak, Lawrence Blaszczkowski, Andrew X. Zhu, David P. Ryan, David C. Christiani, and Geoffrey Liu.....	<b>4010</b>
<b>Phosphodiesterase 11A (PDE11A) and Genetic Predisposition to Adrenocortical Tumors.</b> Rossella Libé, Amato Fratticci, Joel Coste, Frédérique Tissier, Anelia Horvath, Bruno Ragazzon, Fernande Rene-Corail, Lionel Groussin, Xavier Bertagna, Marie Laure Raffin-Sanson, Constantine A. Stratakis, and Jérôme Bertherat.....	<b>4016</b>

## About the Cover

A colon cancer tissue microarray containing 583 stage I-IV patients was stained for pan-cadherin, DAPI, and Met protein. Using novel automated curvature and intensity-based image analysis, epithelial membrane (*red*), cytoplasm (*green*) and nuclei (*blue*) were segmented. Stromal nuclei are shown in gray. The probability distribution of Met protein within each compartment was determined. A higher distribution of Met in the cytoplasm relative to the membrane was a significant predictor of survival in stage I and II patients. For further details, please see Ginty and colleagues on page 3814 in this issue.



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