

CCR Translations

- Expanding the Use of Retinoids in Acute Myeloid Leukemia: Spotlight on Bexarotene.**
□□ *Commentary on Tsai et al., p. 5619*
Suzan McNamara and Wilson H. Miller, Jr.5311

Molecular Pathways

- Oncogenic NOTCH1 Control of MYC and PI3K: Challenges and Opportunities for Anti-NOTCH1 Therapy in T-Cell Acute Lymphoblastic Leukemias and Lymphomas.**
Teresa Palomero and Adolfo Ferrando5314
- Targeting the MDM2-p53 Interaction for Cancer Therapy.** Sanjeev Shangary and Shaomeng Wang.....5318

Report from the FDA

- Tasigna for Chronic and Accelerated Phase Philadelphia Chromosome–Positive Chronic Myelogenous Leukemia Resistant to or Intolerant of Imatinib.** Maitreyee Hazarika, Xiaoping Jiang, Qi Liu, Shwu-Luan Lee, Roshni Ramchandani, Christine Garnett, Micheal S. Orr, Rajeshwari Sridhara, Brian Booth, John K. Leighton, William Timmer, Ravi Harapanhalli, Ramzi Dagher, Robert Justice, and Richard Pazdur5325

Human Cancer Biology

- Expanding the Roles for Pregnane X Receptor in Cancer: Proliferation and Drug Resistance in Ovarian Cancer.** Divya Gupta, Madhukumar Venkatesh, Hongwei Wang, Sean Kim, Michael Sinz, Gary L. Goldberg, Kathleen Whitney, Clifford Longley, and Sridhar Mani5332
- β2-Microglobulin Signaling Blockade Inhibited Androgen Receptor Axis and Caused Apoptosis in Human Prostate Cancer Cells.** Wen-Chin Huang, Jonathan J. Havel, Haiyen E. Zhou, Wei Ping Qian, Hui-Wen Lue, Chia-Yi Chu, Takeo Nomura, and Leland W.K. Chung5341
- Expression and Gene Amplification of Actinin-4 in Invasive Ductal Carcinoma of the Pancreas.** Satoru Kikuchi, Kazufumi Honda, Hitoshi Tsuda, Nobuyoshi Hiraoka, Issei Imoto, Tomoo Kosuge, Tomoko Umaki, Kaoru Onozato, Miki Shitashige, Umio Yamaguchi, Masaya Ono, Akihiko Tsuchida, Tatsuya Aoki, Johji Inazawa, Setsuo Hirohashi, and Tesshi Yamada5348
- Epithelial and Stromal Cathepsin K and CXCL14 Expression in Breast Tumor Progression.** Celina G. Kleer, Noga Bloushtain-Qimron, Yu-Hui Chen, Daniel Carrasco, Min Hu, Jun Yao, Stine-Kathrein Kraeft, Laura C. Collins, Michael S. Sabel, Pedram Argani, Rebecca Gelman, Stuart J. Schnitt, Ian E. Krop, and Kornelia Polyak5357
- Induction of Receptor for Advanced Glycation End Products by EBV Latent Membrane Protein 1 and Its Correlation with Angiogenesis and Cervical Lymph Node Metastasis in Nasopharyngeal Carcinoma.** Akira Tsuji, Naohiro Wakisaka, Satoru Kondo, Shigeyuki Muro, Mitsuru Furukawa, and Tomokazu Yoshizaki5368
- Overexpression of the Zinc Uptake Transporter hZIP1 Inhibits Nuclear Factor-κB and Reduces the Malignant Potential of Prostate Cancer Cells *In vitro* and *In vivo*.** Konstantin Golovine, Peter Makhov, Robert G. Uzzo, Tavis Shaw, David Kunkle, and Vladimir M. Kolenko5376

Cancer Therapy: Preclinical

- Vorinostat and Sorafenib Synergistically Kill Tumor Cells via FLIP Suppression and CD95 Activation.** Guo Zhang, Margaret A. Park, Clint Mitchell, Hossein Hamed, Mohamed Rahmani, Aditi Pandya Martin, David T. Curiel, Adly Yacoub, Martin Graf, Ray Lee, John D. Roberts, Paul B. Fisher, Steven Grant, and Paul Dent5385
- Oncostatin M Induces Bone Loss and Sensitizes Rat Osteosarcoma to the Antitumor Effect of Midostaurin *In vivo*.** Bénédicte Brounais, Céline Chipoy, Kanji Mori, Céline Charrier, Séverine Battaglia, Paul Pilet, Carl D. Richards, Dominique Heymann, Françoise Rédini, and Frédéric Blanchard5400

Postradiation Sensitization of the Histone Deacetylase Inhibitor Valproic Acid. Prakash Chinnaiyan, David Cerna, William E. Burgan, Katie Beam, Eli S. Williams, Kevin Camphausen, and Philip J. Tofilon	5410
Inhibition of the p53 E3 Ligase HDM-2 Induces Apoptosis and DNA Damage-Independent p53 Phosphorylation in Mantle Cell Lymphoma. Richard J. Jones, Qing Chen, Peter M. Voorhees, Ken H. Young, Nathalie Bruey-Sedano, Dajun Yang, and Robert Z. Orlowski	5416
Matuzumab Short-Term Therapy in Experimental Pancreatic Cancer: Prolonged Antitumor Activity in Combination with Gemcitabine. Axel Kleespies, Ivan Ischenko, Martin E. Eichhorn, Hendrik Seeliger, Christiane Amendt, Ole Mantell, Karl-Walter Jauch, and Christiane J. Bruns.....	5426
Targeting Aurora Kinase with MK-0457 Inhibits Ovarian Cancer Growth. Yvonne G. Lin, Anand Immaneni, William M. Merritt, Lingeowda S. Mangala, Seung Wook Kim, Mian M.K. Shahzad, Yvonne T.M. Tsang, Guillermo N. Armaiz-Pena, Chunhua Lu, Aparna A. Kamat, Liz Y. Han, Whitney A. Spannuth, Alpa M. Nick, Charles N. Landen, Jr., Kwong K. Wong, Michael J. Gray, Robert L. Coleman, Diane C. Bodurka, William R. Brinkley, and Anil K. Sood.....	5437
Inhibition of Glioblastoma Growth in a Highly Invasive Nude Mouse Model Can Be Achieved by Targeting Epidermal Growth Factor Receptor but not Vascular Endothelial Growth Factor Receptor-2. Tobias Martens, Yvonne Laabs, Hauke S. Günther, Dirk Kemming, Zhenping Zhu, Larry Witte, Christian Hagel, Manfred Westphal, and Katrin Lamszus.....	5447
Multi-Kinase Inhibitor E7080 Suppresses Lymph Node and Lung Metastases of Human Mammary Breast Tumor MDA-MB-231 via Inhibition of Vascular Endothelial Growth Factor-Receptor (VEGF-R) 2 and VEGF-R3 Kinase. Junji Matsui, Yasuhiro Funahashi, Toshimitsu Uenaka, Tatsuo Watanabe, Akihiko Tsuruoka, and Makoto Asada	5459
Combined Vascular Endothelial Growth Factor Receptor/Epidermal Growth Factor Receptor Blockade with Chemotherapy for Treatment of Local, Uterine, and Metastatic Soft Tissue Sarcoma. Wenhong Ren, Borys Korchin, Guy Lahat, Caimiao Wei, Svetlana Bolshakov, Theresa Nguyen, William Merritt, Adam Dicker, Alexander Lazar, Anil Sood, Raphael E. Pollock, and Dina Lev	5466
Thymidine Selectively Enhances Growth Suppressive Effects of Camptothecin/Irinotecan in MSI+ Cells and Tumors Containing a Mutation of MRE11. Rene Rodriguez, Lasse Tengbjerg Hansen, Geraldine Phear, Jennifer Scorch, Mogens Spang-Thomsen, Angela Cox, Thomas Helleday, and Mark Meuth	5476
Effective CpG Immunotherapy of Breast Carcinoma Prevents but Fails to Eradicate Established Brain Metastasis. Zhengming Xiong, Soheila Gharagozlou, Isabelita Vengco, Wei Chen, and John R. Ohlfest.....	5484
TMTP1, a Novel Tumor-Homing Peptide Specifically Targeting Metastasis. Wanhua Yang, Danfeng Luo, Shixuan Wang, Rui Wang, Rui Chen, Yan Liu, Tao Zhu, Xiangyi Ma, Ronghua Liu, Gang Xu, Li Meng, Yunping Lu, Jianfeng Zhou, and Ding Ma.....	5494
Matrilysin (MMP-7) Is a Novel Broadly Expressed Tumor Antigen Recognized by Antigen-Specific T Cells. Yuko Yokoyama, Frank Grünebach, Susanne M. Schmidt, Annkristin Heine, Maik Häntschel, Stefan Stevanovic, Hans-Georg Rammensee, and Peter Brossart.....	5503
Eradication of Ovarian Tumor Xenografts by Locoregional Administration of Targeted Immunotherapy. Michelandrea De Cesare, Claudia Calcaterra, Graziella Pratesi, Laura Gatti, Franco Zunino, Sylvie Ménard, and Andrea Balsari	5512
Experimental Therapy of Hepatoma with Artemisinin and Its Derivatives: <i>In vitro</i> and <i>In vivo</i> Activity, Chemosensitization, and Mechanisms of Action. Junmei Hou, Disong Wang, Ruiwen Zhang, and Hui Wang.....	5519

Imaging, Diagnosis, Prognosis

Genetic Clustering of Clear Cell Renal Cell Carcinoma Based on Array-Comparative Genomic Hybridization: Its Association with DNA Methylation Alteration and Patient Outcome. Eri Arai, Saori Ushijima, Hitoshi Tsuda, Hiroyuki Fujimoto, Fumie Hosoda, Tatsuhiko Shibata, Tadashi Kondo, Issei Imoto, Johji Inazawa, Setsuo Hirohashi, and Yae Kanai.....	5531
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Whole Chromosome Alterations Predict Survival in High-Risk Neuroblastoma without MYCN Amplification. Sven Bilke, Qing-Rong Chen, Jun S. Wei, and Javed Khan	5540
Magnetic Resonance Imaging–Measured Blood Flow Change after Antiangiogenic Therapy with PTK787/ZK 222584 Correlates with Clinical Outcome in Metastatic Renal Cell Carcinoma. Cedric de Bazelaire, David C. Alsop, Daniel George, Ivan Pedrosa, Yongyu Wang, M. Dror Michaelson, and Neil M. Rofsky	5548
Association of an Extracellular Matrix Gene Cluster with Breast Cancer Prognosis and Endocrine Therapy Response. Jozien Helleman, Maurice P.H.M. Jansen, Kirsten Ruigrok-Ritstier, Iris L. van Staveren, Maxime P. Look, Marion E. Meijer-van Gelder, Anieta M. Sieuwerts, Jan G.M. Klijn, Stefan Sleijfer, John A. Foekens, and Els M.J.J. Berns	5555
A Multigene Assay Is Prognostic of Survival in Patients with Early-Stage Lung Adenocarcinoma. Dan J. Raz, M. Roshni Ray, Jae Y. Kim, Biao He, Miquel Taron, Marcin Skrzypski, Mark Segal, David R. Gandara, Rafael Rosell, and David M. Jablons	5565
Molecular Diagnosis of Sentinel Lymph Node Metastases in Cervical Cancer Using Squamous Cell Carcinoma Antigen. Song-Hua Yuan, Xue-Fang Liang, Wei-Hua Jia, Jian-Ling Huang, Mei Wei, Ling Deng, Li-Zhi Liang, Xiao-Yun Wang, and Yi-Xin Zeng	5571
Combination of Quantitative IMP3 and Tumor Stage: A New System to Predict Metastasis for Patients with Localized Renal Cell Carcinomas. Zhong Jiang, Peigou G. Chu, Bruce A. Woda, Qin Liu, K.C. Balaji, Kenneth L. Rock, and Chin-Lee Wu	5579

Cancer Therapy: Clinical

Gemtuzumab Ozogamicin as Part of Reduced-Intensity Conditioning for Allogeneic Hematopoietic Cell Transplantation in Patients with Relapsed Acute Myeloid Leukemia. Martin Bornhäuser, Thomas Illmer, Uta Oelschlaegel, Johannes Schetelig, Rainer Ordemann, Markus Schaich, Mathias Hänel, Ulrich Schuler, Christian Thiede, Alexander Kiani, Uwe Platzbecker, and Gerhard Ehninger	5585
ABCB1 MDR1 Polymorphisms and Progression-Free Survival among Women with Ovarian Cancer following Paclitaxel/Carboplatin Chemotherapy. Sharon E. Johnatty, Jonathan Beesley, Jim Paul, Sian Fereday, Amanda B. Spurdle, Penelope M. Webb, Karen Byth, Sharon Marsh, Howard McLeod, AOCs Study Group, Paul R. Harnett, Robert Brown, Anna de Fazio, and Georgia Chenevix-Trench	5594
A Phase I Study of EKB-569 in Combination with Capecitabine in Patients with Advanced Colorectal Cancer. Dan Laheru, Gary Croghan, Ronald Bukowski, Michelle Rudek, Wells Messersmith, Charles Erlichman, Robert Pelley, Antonio Jimeno, Ross Donehower, Joseph Boni, Richat Abbas, Patricia Martins, Charles Zacharchuk, and Manuel Hidalgo	5602
Treatment of Metastatic Melanoma Using Interleukin-2 Alone or in Conjunction with Vaccines. Franz O. Smith, Stephanie G. Downey, Jacob A. Klapper, James C. Yang, Richard M. Sherry, Richard E. Royal, Udai S. Kammula, Marybeth S. Hughes, Nicholas P. Restifo, Catherine L. Levy, Donald E. White, Seth M. Steinberg, and Steven A. Rosenberg	5610
A Phase I Study of Bexarotene, a Retinoic X Receptor Agonist, in Non-M3 Acute Myeloid Leukemia. Donald E. Tsai, Selina M. Luger, Charalambos Andreadis, Dan T. Vogl, Allison Kemner, Melissa Potuzak, Ami Goradia, Alison W. Loren, Alexander E. Perl, Stephen J. Schuster, David L. Porter, Edward A. Stadtmauer, Steven C. Goldstein, James E. Thompson, Cezary Swider, Adam Bagg, Anthony R. Mato, and Martin Carroll	5619
CpG Oligodeoxynucleotides Alter Lymphocyte and Dendritic Cell Trafficking in Humans. W. Nicholas Haining, Jeffrey Davies, Holger Kanzler, Linda Drury, Thomas Brenn, John Evans, Jill Angelosanto, Steven Rivoli, Kate Russell, Suzanne George, Paul Sims, Donna Neuberger, Xiaochun Li, Jeffrey Kutok, Jeffrey Morgan, Patrick Wen, George Demetri, Robert L. Coffman, and Lee M. Nadler	5626

Cancer Prevention and Susceptibility

Effect of Genetic Variability within 8q24 on Aggressiveness Patterns at Diagnosis and Familial Status of Prostate Cancer. Olivier Cussenot, Abdel-Rahmène Azzouzi, Gladys Bantsimba-Malanda, Cecile Gaffory, Philippe Mangin, Luc Cormier, Georges Fournier, Antoine Valeri, Lionel Jouffe, Morgan Roupert, Gaëlle Fromont, Mathilde Sibony, Eva Comperat, and Géraldine Cancel-Tassin	5635
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Letters to the Editor

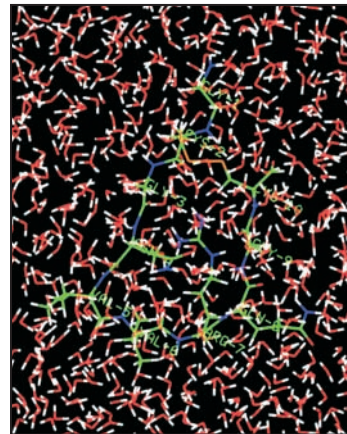
IMP3: A New and Important Biomarker of Systemic Malignancies. Shailendra Kapoor.....5640

Reply to the Letter to the Editor from S. Kapoor.

Zhong Jiang.....5640

About the Cover

Molecular dynamics simulation of TMTP1. TMTP1 peptide is solved in a water cubic periodic box. Ten amino acid residues form a cyclic peptide via a disulfide bond between the two cysteines that respectively lie at the head and the tail. The line view of TMTP1 peptide with side chains and the internal hydrogen bonds. For further details, please see Luo et al on page 5494 of this issue.



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