BREAKING ADVANCES

6375  Highlights from Recent Cancer Literature

REVIEWS

6377  A Genome-wide View of Microsatellite Instability: Old Stories of Cancer Mutations Revisited with New Sequencing Technologies
Tae-Min Kim and Peter J. Park

6383  Chimeric Antigen Receptor T-cell Therapy to Target Hematologic Malignancies
Saad Sirop Kenderian, Marco Ruella, Saar Gill, and Michael Kalos

PRIORITY REPORT

6390  Discrepancies in Cancer Genomic Sequencing Highlight Opportunities for Driver Mutation Discovery
Andrew M. Hudson, Tim Yates, Yaoyong Li, Eleanor W. Trotter, Shameem Fawdar, Phil Chapman, Paul Lorigan, Andrew Blankin, Crispin J. Miller, and John Brognard
Précis: These findings highlight major discrepancies in mutational profiles of identical cancer cell lines sequenced by two different institutes, with implications for identifying previously undiscovered driver mutations.

INTEGRATED SYSTEMS AND TECHNOLOGIES

6397  Mathematical Modeling of Tumor Growth and Metastatic Spreading: Validation in Tumor-Bearing Mice
Niklas Hartung, Séverine Molland, Dominique Barbolosi, Assia Benabdallah, Guillemette Chapuisat, Gerard Henry, Sarah Giacometti, Athanasios Iliadis, Joseph Ciccolini, Christian Faivre, and Florence Hubert
Précis: This work advances efforts to predict metastatic spreading during the earliest stages of cancer, at points that could help clinicians make the best decisions on treatment strategies.

6408  Direct Chemosensitivity Monitoring Ex Vivo on Undissociated Melanoma Tumor Tissue by Impedance Spectroscopy
Heinz-Georg Jahnke, Sarah Poenick, Jan Maschke, Michael Kendler, Jan C. Simon, and Andrea A. Robitzki
Précis: This study presents a novel and more accurate tissue-based method to determine chemotherapeutic drug sensitivity using small fragments of tumor tissue, addressing a need to personalize therapy for patients to improve treatment outcomes.

MICROENVIRONMENT AND IMMUNOLOGY

6419  CXM: A New Tool for Mapping Breast Cancer Risk in the Tumor Microenvironment
Michael J. Fister, Bradley T. Endres, Nathan Rudemiller, Allison B. Sarkis, Stephanie Santarriaga, Ishan Roy, Angela Lemke, Aron M. Geurs, Carol Moreno, Sophia Ran, Shiring-Wern Tsaih, Jeffery De Pons, Daniel F. Carlson, Wenfang Tan, Scott C. Fahrenkrug, Zelmira Lazarova, Jozef Lazur, Paula E. North, Peter S. LaViolette, Michael B. Dwinell, James D. Shull, and Howard J. Jacob
Précis: These results establish the utility of a novel model of breast cancer that can localize genetic variants that affect breast cancer risk through actions on the tumor microenvironment, rather than the tumor cell itself.

6430  Ag-Presenting CpG-Activated pDCs Prime Th17 Cells That Induce Tumor Regression
Leslie Guéry, Juan Dubrot, Carla Lippens, Dale Brighouse, Pauline Malinge, Magali Irla, Caroline Pot, Walter Reith, Jean-Marc Waldburger, and Stéphanie Hugues
Précis: This study identifies a new antigen-presenting strategy that may improve cancer immunotherapy involving Th17 cells.

6441  SA-4-1BBL and Monophosphoryl Lipid A Constitute an Efficacious Combination Adjuvant for Cancer Vaccines
Abhishek K. Srivastava, Gunes Dinc, Rajesh K. Sharma, Esma S. Yolcu, Hong Zhao, and Haval Shirwan
Précis: These results offer preclinical proof of concept for the use of a powerful new adjuvant system for tumor antigen-based cancer vaccines, with immediate implications for its clinical evaluation in the oncology clinic.
Host Deficiency in Caveolin-2 Inhibits Lung Carcinoma Tumor Growth by Impairing Tumor Angiogenesis
Yajun Liu, Sungchan Jang, Leike Xie, and Grzegorz Sowa

Précis: Loss of a protein that helps organize lipid rafts on the plasma membrane reduces cancerous cell growth, with possible implications for a generalized approach to cancer targeting.

Histone H1.3 Suppresses H19 Noncoding RNA Expression and Cell Growth of Ovarian Cancer Cells
Magdalena Medrzycki, Yunzhe Zhang, Weijia Zhang, Kaixiang Cao, Chenyi Pan, Nathalie Lailler, John F. McDonald, Eric E. Bouhassira, and Yuhong Fan

Précis: These results provide new information about the regulation of a noncoding RNA in ovarian cancer cells, advancing work in a timely new area of RNA physiology and cancer.

Oncogenic Properties of a Spermatogenic Meiotic Variant of Fer Kinase Expressed in Somatic Cells
Etai Yaffe, Elad Hikri, Yoav Elkis, Ortal Cohen, Ariela Segal, Adar Makovski, Alexander Varvak, Sally Shpungin, and Uri Nir

Précis: This provocative study reveals a molecular alteration in the mitochondria of cancer cells that may represent a common pathophysiological root, with possible implications for broad-based treatments.

BRG1/SMARCA4 Inactivation Promotes Non–Small Cell Lung Cancer Aggressiveness by Altering Chromatin Organization
Tess Orvis, Austin Hepperla, Vonn Walter, Shuie Song, Jeremy Simon, Joel Parker, Matthew D. Wilkerson, Nisarg Desai, Michael B. Major, D. Neil Hayes, Ian J. Davis, and Bernard Weissman

Précis: These results offer direct evidence of a tumor suppressor role for a core ATPase found in SWI/SNF chromatin regulatory complexes, the inactivation of which contributes to lung cancer aggressiveness by altering nucleosome positioning and expression at many cancer-associated genes.

Deubiquitination of γ-Tubulin by BAP1 Prevents Chromosome Instability in Breast Cancer Cells
Reihaner Zarrizi, Julien Albert Menard, Mattias Belting, and Ramin Massoumi

Précis: These findings illuminate a core mechanism preventing genomic instability, with implications for understanding malignant progression.

The Notch Pathway Inhibits TGFβ Signaling in Breast Cancer through HEYL-Mediated Crosstalk
Liangfeng Han, Adam Diehl, Nguyen K. Nguyen, Preethi Korangath, Weiwen Teo, Soonweng Cho, Scott Kominsky, David L. Huso, Lionel Feigenbaum, Alan Rein, Pedram Argani, Goran Landberg, Manfred Gessler, and Saraswati Sukumar

Précis: These findings identify a particular mechanism of TGFβ signaling as a key element in the development of drug resistance in breast cancer.

STAT1 Drives Tumor Progression in Serous Papillary Endometrial Cancer
Budiman Kharma, Tsukasa Baba, Noriomi Matsumura, Hyun Sook Kang, Junzo Hamanishi, Ryusuke Murakami, Melissa M. McConathy, Samuel Leung, Ken Yamaguchi, Yuko Hosoe, Yumiko Yoshioka, Susan K. Murphy, Masaki Mandai, David G. Hunstman, and Ikuo Konishi

Précis: This study identifies a molecular signature and root oncogenic driver of serous papillary endometrial cancer, a relatively rare and poorly characterized form of uterine cancer that arises in post-menopausal women, with implications for improving its prognosis and treatment.

IGF2 Preserves Osteosarcoma Cell Survival by Creating an Autophagic State of Dormancy That Protects Cells against Chemotherapeutic Stress

Précis: This study provides a mechanistic rationale for blunting IGF/insulin-mediated survival signals in osteosarcoma, a pediatric tumor notorious for its intrinsic therapeutic resistance, as a strategy to improve treatment outcomes.
CTBP2 Modulates the Androgen Receptor to Promote Prostate Cancer Progression
Ken-ichi Takayama, Takashi Suzuki, Tetsuya Fujimura, Tomohiko Urano, Satoru Takahashi, Yukio Homma, and Satoshi Inoue

**Précis:** A transcriptional corepressor linked to prostate cancer susceptibility is found here to be an androgen-regulated gene that modulates pro-cancerous downstream signals from the androgen receptor.

Genetic Evidence of a Precisely Tuned Dysregulation in the Hypoxia Signaling Pathway during Oncogenesis

**Précis:** These findings offer evidence in humans to validate the “continuum” model of tumor suppression, advancing work beyond the mouse in developing a successor to the classic “two-hit” model in the field.

In Vivo Disruption of an Rb-E2F-Ezh2 Signaling Loop Causes Bladder Cancer
Mirentxu Santos, Mónica Martinez-Fernández, Marta Dueñas, Ramón Garcia-Escudero, Begoña Alfaya, Felipe Villacampa, Cristina Saiz-Laderà, Clotilde Costa, Marta Oteo, José Duarte, Víctor Martínez, Mª José Gómez-Rodríguez, Mª Luisa Martín, Manoli Fernández, Patrick Viautour, Miguel A. Morcillo, Julien Sage, Daniel Castellano, José L. Rodríguez-Peralto, Federico de la Rosa, and Jesús M Paramio

**Précis:** This study addresses the gap in knowledge concerning the genetic and epigenetic underpinnings of bladder cancer development, which still remain relatively obscure.

Suppression of Deacetylase SIRT1 Mediates Tumor-Suppressive NOTCH Response and Offers a Novel Treatment Option in Metastatic Ewing Sarcoma

**Précis:** These findings offer a mechanistic rationale for the use of pharmacological inhibitors of a p53 deacetylase to treat cancers in which NOTCH acts a tumor suppressor.

High Serum Iron Is Associated with Increased Cancer Risk
Chi Pang Wen, June Han Lee, Ya-Ping Tai, Christopher Wen, Shuian Be Wu, Min Kuang Tsai, Dennis P.H. Hsieh, Hung-Che Chiang, Chao Agnes Hsiung, Chung Y. Hsu, and Xifeng Wu

**Précis:** This large cohort study reveals that high levels of iron in blood serum is a risk marker for a variety of adult cancer, most dramatically in conferring a 3-fold increased risk for liver cancer.

Microenvironment-Derived HGF Overcomes Genetically Determined Sensitivity to Anti-MET Drugs
Selma Pennacchietti, Manuela Cazzanti, Andrea Bertotti, William M. Rideout III, May Han, Jeno Gyuris, Timothy Perera, Paolo M. Comoglio, Livio Trusolino, and Paolo Michieli

**Précis:** This study offers preclinical proof of concept for the use of antibodies that neutralize hepatocyte growth factor along with MET-targeting agents as a more effective therapeutic strategy to treat MET-dependent tumors.

Targeting Cancer Stem–like Cells as an Approach to Defeating Cellular Heterogeneity in Ewing Sarcoma
Sandrine Cornaz-Buros, Nicolo Riggi, Claudio DeVito, Alexandre Sarre, Igor Letovanec, Paolo Provero, and Ivan Stamenkovic

**Précis:** These results suggest a broadly and immediately applicable approach to improve the treatment of solid tumors that are marked by extensive cellular heterogeneity, likely driven by the plastic nature of cancer stem-like cells, with immediate applications for clinical evaluation.

PREVENTION AND EPIDEMIOLOGY

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**TUMOR AND STEM CELL BIOLOGY**

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GPx2 Suppression of H$_2$O$_2$ Stress Links the Formation of Differentiated Tumor Mass to Metastatic Capacity in Colorectal Cancer

Benjamin L. Emmink, Jamila Laoukili, Anna P. Kipp, Jan Koster, Klaas M. Govaert, Szabolcs Fatrai, Andre Verheem, Ernst J.A. Steller, Regina Brigelius-Flohé, Connie R. Jimenez, Inne H.M. Borel Rinkes, and Onno Kranenburg

Précis: Results reveal an unexpected redox-controlled link between formation of a tumor mass and its capacity for metastasis.

RETRACTION

Retraction: Novel HSP90 Inhibitor NVP-HSP990 Targets Cell-Cycle Regulators to Ablate Olig 2-Positive Glioma Tumor–Initiating Cells

CORRECTIONS

Correction: Diffusion-Weighted Imaging in Cancer: Physical Foundations and Applications of Restriction Spectrum Imaging

Correction: A Novel Wnt Regulatory Axis in Endometrioid Endometrial Cancer

ABOUT THE COVER

CD8$^{+}$ T cells are critical for elimination of cancer. A major limitation of therapeutic cancer vaccines is their inability to activate and mobilize CD8$^{+}$ T cells for infiltration into tumor. A vaccine formulation containing SA-4-1BBL and MPL as a novel adjuvant system shows robust efficacy in activating and recruiting CD8$^{+}$ T cells into the tumor, with subsequent effective tumor destruction in preclinical models. For details, see article by Srivastava and colleagues on page 6441.