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Angiogenesis, Metastasis, and the Cellular Microenvironment

Imatinib Mesylate (STI-571) Reduces Bcr-Abl-Mediated Vascular Endothelial Growth Factor Secretion in Chronic Myelogenous Leukemia89
John M.L. Ebos, Jennifer Tran, Zubin Master, Daniel Dumont, Junia V. Melo, Elisabeth Buchdunger, and Robert S. Kerbel

A Fucosylated Chondroitin Sulfate From Echinoderm Modulates *in Vitro* Fibroblast Growth Factor 2-Dependent Angiogenesis96
Jacqueline Tapon-Bretaudière, Delphine Chabut, Maximiliano Zierer, Sabine Matou, Dominique Helley, Andrée Bros, Paulo A.S. Mourão, and Anne-Marie Fischer

Cancer Genes and Genomics

Damage Recovery Pathways in *Saccharomyces cerevisiae* Revealed by Genomic Phenotyping and Interactome Mapping103
Thomas J. Begley, Ari S. Rosenbach, Trey Ideker, and Leona D. Samson

Prostate Stem Cell Antigen Is a Marker of Late Intermediate Prostate Epithelial Cells113
Chau P. Tran, Connie Lin, Joyce Yamashiro, and Robert E. Reiter

Cell Cycle, Cell Death, and Senescence

Apoptosis of Mortal Human Fibroblasts Transformed by the Bovine Papillomavirus E5 Oncoprotein122
Ying Zhang, John M. Lehman, and Lisa M. Petti

DNA Damage and Cellular Stress Responses

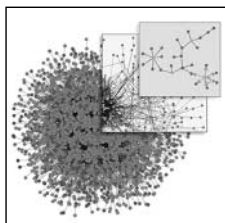
Role of Reactive Oxygen Species in Cells Overexpressing Manganese Superoxide Dismutase: Mechanism for Induction of Radioresistance137
Yasunari Takada, Misao Hachiya, Sang-Hee Park, Yoshiaki Osawa, Toshihiko Ozawa, and Makoto Akashi

Signaling and Regulation

Extracellular Signal-Regulated Protein Kinase Signaling Is Uncoupled From Initial Differentiation of Central Nervous System Stem Cells to Neurons147
Maria Enarsson, Anna Erlandsson, Helena Larsson, and Karin Forsberg-Nilsson

Regulation of Lck and Fyn Tyrosine Kinase Activities by Transmembrane Protein Tyrosine Phosphatase Leukocyte Common Antigen-Related Molecule155
Kazutake Tsujikawa, Tomoko Ichijo, Kazuki Moriyama, Noriko Tadotsu, Kazuhiro Sakamoto, Naoki Sakane, So-ichiro Fukada, Tatsuhiko Furukawa, Haruo Saito, and Hiroshi Yamamoto

Information for Authors i



On the Cover

Screening mutant phenotypes against a network of 12,232 protein-protein (blue lines) and protein-DNA (red arrows) interactions. *Large inset*, filtered network containing proteins that contribute to viable recovery (green nodes) after agent treatment and their immediate neighbors. *Small inset*, a concentrated subnetwork. (Red and gray nodes, no phenotype and untested proteins.) For details, see article by Begley et al. on p. 103 this issue

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