

CONTENTS

Journal of **Cellular
Biochemistry**

VOLUME 111 • NUMBER 5

CONTENTS i

FEATURES v

VIEW POINTS

WNT Unrelated Activities in Commercially Available Preparations of Recombinant WNT3a

Lukas Cajanek, Linda Adlerz, Vitezslav Bryja, and Ernest Arenas 1077

PUBLISHED ONLINE 21 JULY 2010

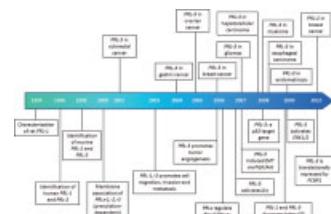


PAGE 1080

Revisiting Ancient mtDNA Equid Sequences From Pompeii

Susan M.R. Gurney 1080

PUBLISHED ONLINE 29 OCTOBER 2010



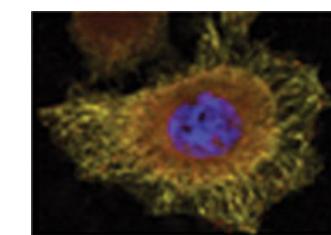
PAGE 1088

PROSPECTS

Activation of Signal Pathways and the Resistance to Anti-EGFR Treatment in Colorectal Cancer

Jiezhong Chen, Xu-Feng Huang, and Andrew Katsifis 1082

PUBLISHED ONLINE 4 NOVEMBER 2010

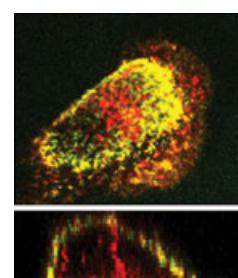


PAGE 1103

PRL-3 Phosphatase and Cancer Metastasis

Abdul Qader O. Al-Aidaroos and Qi Zeng 1087

PUBLISHED ONLINE 4 NOVEMBER 2010



PAGE 1129

ARTICLES

A New Nuclear Protease With Cathepsin L Properties Is Present in HeLa and Caco-2 Cells

Marcia Puchi, Jenaro García-Huidobro, Candy Cordova, Rodrigo Aguilar, Estefanie Dufey, Maria Imschenetzky, Paula Bustos, and Violeta Morin 1099

PUBLISHED ONLINE 19 MAY 2010

Fyn Promotes Proliferation, Differentiation, Survival and Function of Osteoclast Lineage Cells

Hyun-Ju Kim, Julia T. Warren, Shin-Yoon Kim, Jean C. Chappel, Carl J. DeSelms, F. Patrick Ross, Wei Zou, and Steven L. Teitelbaum 1107

PUBLISHED ONLINE 17 AUGUST 2010

VEGF Stimulation Enhances Livin Protein Synthesis Through mTOR Signaling

Biao Yan, Men Kong, Shi Chen, and Yi-han Chen 1114

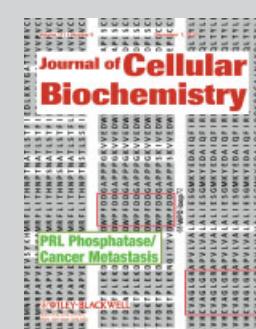
PUBLISHED ONLINE 17 AUGUST 2010

Recycling and LFA-1-Dependent Trafficking of ICAM-1 to the Immunological Synapse

Jae-Hyeok Jo, Min-Sung Kwon, Hyang-Ok Choi, Hyun-Mee Oh, Hyang-Jin Kim, and Chang-Duk Jun 1125

PUBLISHED ONLINE 2 AUGUST 2010

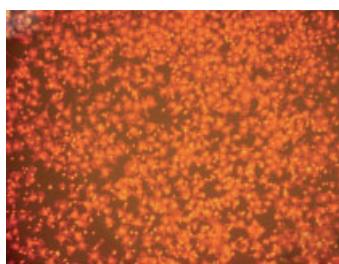
Cover: The cover shows amino acid sequence alignment of known and predicted PRL-3 homologues across different species. Key functional motifs of PRLs are boxed and labeled. Amino acids common to all species are shaded. See Prospect in this issue by Al-Aidaroos and Zeng, pages 1087–1098.



(continued on next page)

CONTENTS

CONTINUED



PAGE 1144

Osteoblasts Are a Major Source of Inflammatory Cytokines in the Tumor Microenvironment of Bone Metastatic Breast Cancer

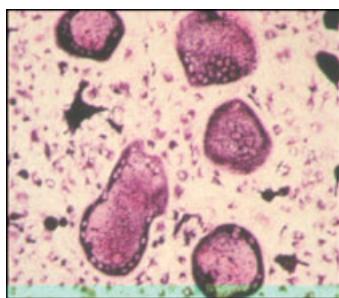
Karen M. Bussard, David J. Venzon, and Andrea M. Mastro 1138

PUBLISHED ONLINE 3 AUGUST 2010

Demethyl Fruticulin A (SCO-1) Causes Apoptosis by Inducing Reactive Oxygen Species in Mitochondria

Massimiliano Monticone, Angela Bisio, Antonio Daga, Paolo Giannoni, Walter Giaretti, Massimo Maffei, Ulrich Pfeffer, Francesco Romeo, Rodolfo Quarto, Giovanni Romussi, Giorgio Corte, and Patrizio Castagnola 1149

PUBLISHED ONLINE 3 AUGUST 2010



PAGE 1181

Lysyl Oxidase Propeptide Sensitizes Pancreatic and Breast Cancer Cells to Doxorubicin-Induced Apoptosis

Chengyin Min, Yingshe Zhao, Mathilde Romagnoli, Philip C. Trackman, Gail E. Sonenschein, and Kathrin H. Kirsch 1160

PUBLISHED ONLINE 17 AUGUST 2010

c-Cbl Regulates Glioma Invasion Through Matrix Metalloproteinase 2

Hojin Lee and Alexander Y. Tsygankov 1169

PUBLISHED ONLINE 17 AUGUST 2010

Microarray Profile of Gene Expression During Osteoclast Differentiation in Modelled Microgravity

Yuvaraj Sambandam, Jeremy J. Blanchard, Giffin Daughtridge, Robert J. Kolb, Srinivasan Shanmugarajan, Subramanya N. M. Pandruvada, Ted A. Bateman, and Sakamuri V. Reddy 1179

PUBLISHED ONLINE 17 AUGUST 2010



PAGE 1195

Capacitation Suppression by Mouse Seminal Vesicle Autoantigen Involves a Decrease in Plasma Membrane Ca^{2+} -ATPase (PMCA)-Mediated Intracellular Calcium

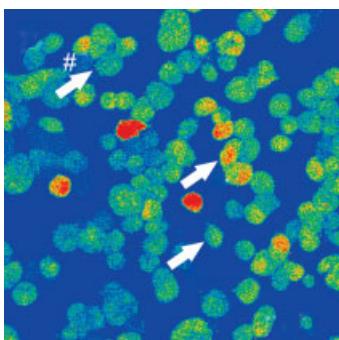
Shing-Hwa Lu, Yuan-Kuei Yen, Thai-Yen Ling, Kur-Ta Cheng, Jye-An Shu, Heng-Kien Au, and Yen-Hua Huang 1188

PUBLISHED ONLINE 17 AUGUST 2010

Hedgehog Signaling and Osteogenic Differentiation in Multipotent Bone Marrow Stromal Cells Are Inhibited by Oxidative Stress

Woo-Kyun Kim, Vicente Meliton, Noam Bourquard, Theodore J. Hahn, and Farhad Parhami 1199

PUBLISHED ONLINE 17 AUGUST 2010



PAGE 1225

Both FGF23 and Extracellular Phosphate Activate Raf/MEK/ERK Pathway via FGF Receptors in HEK293 Cells

Miwa Yamazaki, Keiichi Ozono, Tomoko Okada, Kanako Tachikawa, Hiroki Kondou, Yasuhisa Ohata, and Toshimi Michigami 1210

PUBLISHED ONLINE 17 AUGUST 2010

Thapsigargin-Induced Ca^{2+} Increase Inhibits TGF β 1-mediated Smad2 Transcriptional Responses via Ca^{2+} /Calmodulin-Dependent Protein Kinase II

Ming Ming, Ivan Manzini, Weidong Le, Kerstin Kriegelstein, and Björn Spittau 1222

PUBLISHED ONLINE 17 AUGUST 2010

(continued on next page)

CONTENTS CONTINUED

Role of Lysyl Oxidase Propeptide in Secretion and Enzyme Activity

Jessica L. Grimsby, Hector A. Lucero, Philip C. Trackman, Katya Ravid, and Herbert M. Kagan 1231

PUBLISHED ONLINE 17 AUGUST 2010



PAGE 1238

Characterization of E2F3a Function in HepG2 Liver Cancer Cells

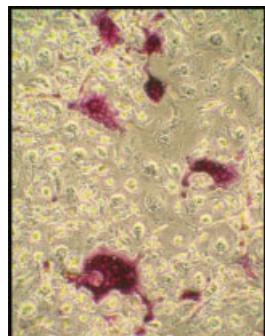
Wei Li, Guo-Xin Ni, Ping Zhang, Zheng-Xi Zhang, Wei Li, and Qiang Wu 1244

PUBLISHED ONLINE 27 AUGUST 2010

Glypican 3 Binds to GLUT1 and Decreases Glucose Transport Activity in Hepatocellular Carcinoma Cells

Hye-Sim Cho, Jung-Mo Ahn, Ho-Jae Han, and Je-Yoel Cho 1252

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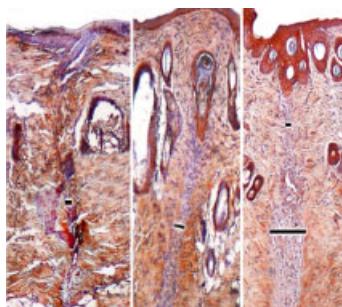


PAGE 1263

Compressive Mechanical Force Augments Osteoclastogenesis by Bone Marrow Macrophages Through Activation of c-Fms-Mediated Signaling

Eui-Sic Cho, Keun-Soo Lee, Young-Ok Son, Yong-Suk Jang, Seung-Youp Lee, So-Yeong Kwak, Yeon-Mi Yang, Seung-Moon Park, and Jeong-Chae Lee 1260

PUBLISHED ONLINE 27 AUGUST 2010



PAGE 1276

Angiogenesis and Inflammation Signaling Are Targets of Beer Polyphenols on Vascular Cells

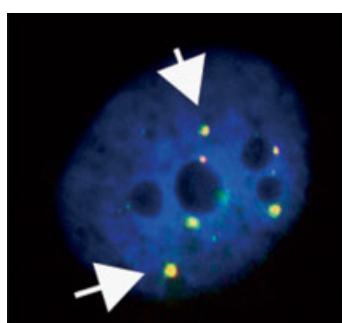
Rita Negrão, Raquel Costa, Delfim Duarte, Tiago Taveira Gomes, Mário Mendanha, Liane Moura, Luísa Vasques, Isabel Azevedo, and Raquel Soares 1270

PUBLISHED ONLINE 27 AUGUST 2010

p53 and p21(Waf1) Are Recruited to Distinct PML-Containing Nuclear Foci in Irradiated and Nutlin-3a-Treated U2OS Cells

Hong Shen and Carl G. Maki 1280

PUBLISHED ONLINE 27 JULY 2010



PAGE 1286

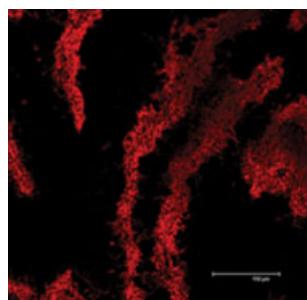
Modulation of GEF-H1 Induced Signaling by Heparanase in Brain Metastatic Melanoma Cells

Lon D. Ridgway, Michael D. Wetzel, and Dario Marchetti 1299

PUBLISHED ONLINE 27 AUGUST 2010

(continued on next page)

CONTENTS CONTINUED



PAGE 1326

SIRT1 Modulates High-Mobility Group Box 1-Induced Osteoclastogenic Cytokines in Human Periodontal Ligament Cells

Young-Suk Kim, Young-Man Lee, Jae-Sang Park, Suk-Keun Lee,
and Eun-Cheol Kim

1310

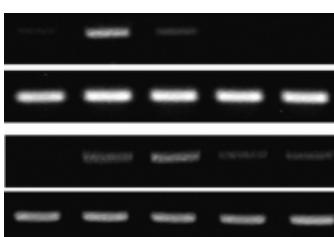
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Nitric Oxide Involved in the IL-1-Induced Inhibition of Fructose Intestinal Transport

Alberto García-Barrios, Natalia Guillén, Sonia Gascón, 1,2 Jesús Osada,
Carmen Mª Vazquez, José Luis Miguel-Carrasco, and Mª Jesús Rodríguez-Yoldi

1321

PUBLISHED ONLINE 27 AUGUST 2010



PAGE 1326

Identification of a Functional Hepatocyte Nuclear Factor 4 Binding Site in the Neutral Ceramidase Promoter

H. R. Maltesen, Jesper T. Troelsen, and Jørgen Olsen

1330

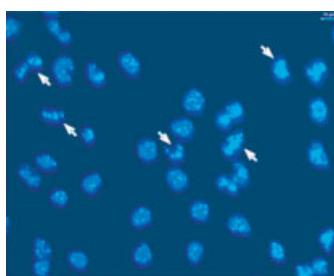
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Differential Regulation by Fucoidan of IFN- γ -Induced NO Production in Glial cells and Macrophages

Hang Do, Nam-Sung Kang, Suhkneung Pyo, Timothy R. Billiar, and
Eun-Hwa Sohn

1337

PUBLISHED ONLINE 9 SEPTEMBER 2010



PAGE 1341

FGF2 Promotes Msx2 Stimulated PC-1 Expression Via Frs2/MAPK Signaling

Yan Li, Jin Liu, Mark Hudson, Sungsu Kim, and Nan E. Hatch

1346

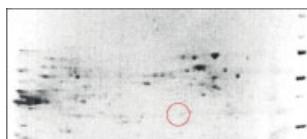
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Identification of p10 as a Neurotoxic Product Generated From the Proteolytic Cleavage of the Neuronal Cdk5 Activator

Jenny Chew, Minghui Jessica Chen, Alan Yiu-Wah Lee, Zhao Feng Peng,
Kevin Wai Yin Chong, Lisheng He, Boon Huat Bay, Jian Ming Jeremy Ng,
Robert Z. Qi, and Nam Sang Cheung

1359

PUBLISHED ONLINE 9 SEPTEMBER 2010



PAGE 1362

Prohibitin Is a Cholesterol-Sensitive Regulator of Cell Cycle Transit

Pei Dong, Jessica Flores, Kristine Pelton, and Keith R. Solomon

1367

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ERRATUM

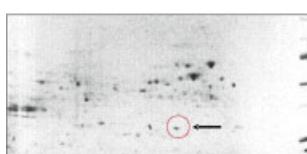
Erratum to "The Cellular Control of DNA Double-Strand Breaks"

[J Cell Biochem 2006, 99:1463–1475]

Shaun P. Scott and Tej K. Pandita

1375

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PAGE 1370