Visualizing Gene Expression in Real-Time

Stephen W. Carmichael¹ Mayo Clinic stephen.carmichael@mayo.edu

Gene expression has been visualized for a few decades, but in static forms such as blots and gene chips. Susan Janicki, Toshiro Tsukamoto, Simone Salghetti, William Tansey, Ravi Sachidanandam, Kannanganattu Prasanth, Thomas Ried, Yaron Shav-Tal, Edouard Bertrand, Robert Singer, and David Spector have recently designed a cell line in which gene expression can be observed with stunningly accurate spatial and temporal resolution.² Gene expression is a cascade of events beginning with transcription of RNA from the DNA template and ending with translation into a protein sequence. Janicki et al. were able to visualize the entire process at the levels of DNA, RNA, and proteins in living cells!

As pointed out by the authors, several specific steps in the gene expression paradigm have been discovered and extensively studied in *vitro*. However, the dynamics of mRNA synthesis at a single specific transcription site has not been revealed. They developed a cell line that allows the investigation of how events of gene expression are coordinated spatially and temporally in vivo. They began by constructing a plasmid, containing a transcription unit, and stably integrating it into a human cancer cell line. With the addition of other plasmids, they had a cell line that produced a series of detectable markers at specific events

during gene expression. Transfected cells were maintained in a physiologic chamber on an inverted microscope and time-lapse imaging was acquired. Stacks of several images 0.5 µm apart were taken to assure that at least one image was focused on the structure of interest for each time point. Different wavelengths were used to excite the different reporter fluorochromes.

After extensive analysis, the interconnectedness of chromatin remodeling, transcription, mRNA processing, and messenger ribonucleoprotein (mRNP) export became apparent. Janicki et al. developed a cell system that allows the visualization of an inducible array of transcription units and their RNA and protein products in living cells. This allows the evaluation of dynamic changes in chromatin structure, RNA synthesis, and factor dissociation/association during the induction of transcription. The movie can be seen at http://spectorlab.cshl.edu. This study provided significant insight into the dynamic spatial and temporal changes that occur as chromatin transitions from a heterochromatic to a euchromatic state. This system has considerable potential to address a broad range of questions relating to gene expression, DNA replication, and chromatin stability!

References

- 1. The author gratefully acknowledges Dr. David Spector for reviewing this article.
- 2. Janicki, S.M., T. Tsukamoto, S.E. Salghetti, W.P. Tansey, R. Sachidanandam, K.V. Prasanth, T. Ried, Y. Sharon Shav-Tal, E. Bertrand, R.H. Singer, and D.L. Spector, From silencing to gene expression: Real-time analysis in single genes, Cell 116:683-698, 2004.

INDEX OF ARTICLES

Visualizing Gene Expression in Real-Time
Stephen W. Carmichael, Mayo Clinic
"Objects Worthy of Notice" Microscopical Anatomy of
Selected Plants Collected by The Lewis & Clark Expedition8
Harry A. Alden and Roland H. Cunningham, Smithsonian Center for
Materials Research and Education; Kevin Ryan and Paul T. Jantzen
Media Cybernetics, Inc.; and David R. Dobbins, Millersville University
A Novel GEMINI [®] STEM Detector System
Jack Vermeulen and Heiner Jaksch, Carl Zeiss SMT, Germany
Phase Measurement in Electron Microscopy
Using the Transport of Intensity Equation
Kazuo Ishizuka, HREM Research Inc. and Brendan Allman, IATIA Ltd.
High Resolution Light Microscopy of Live Cells
Vitaly Vodyanoy, Auburn University
Nondestructive, High-Resolution Materials Characterization
with the Confocal Raman-AFM
U. Schmidt, A. Jauss, W. Ibach, K. Weishaupt, and O. Hollricher
WITec GmbH. Ulm, Germany
Making Anaglyphs in Photoshop
Jerry Sedgewick, University of Minnesota
Ultra High Resolution SEM on Insulators and
Contaminating Samples40
Trisha Rice, Hillsboro, OR and Ralph Knowles FEI Eindhoven,
The Netherlands
Influence of Illumination Conditions on Temperature in Sample
Cell and the Output of a Quadrant Detector in an Optical
Tweezers System44
Yuqiang Jiang, ^{1,2,3} H. Guo, ² C. Liu, ² Z. Li,2 B. Cheng, ² D. Zhang, ²
and S. Jia; ¹ Shanxi University, Taiyuan, P. R. China
² Chinese Academy of Science, Beijing, P. R. China
³ Shanxi Datong University, Datong, P. R. China

Microwave Protocols for Plant and Animal Paraffin Microtechnique50
Denise Schichnes, Jeffrey A. Nemson, and Steven E. Ruzin, The University of California at Berkeley
Raman Microscopy as an Aid in Failure Analysis –
Examples From the Lab54
W. John Wolfgong, Raytheon, McKinney, TX
Virtual Electron Microscopy for Undergraduate/Graduate Classes
New and Interesting PITTCON 2005
Industry News
NetNotes
Index of Advertisers



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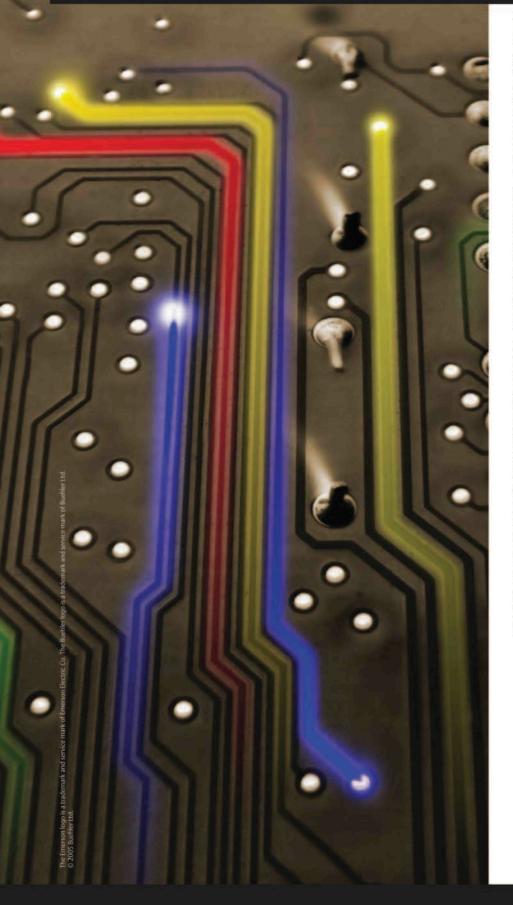
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Thomas E. Phillips, Contributing Editor

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