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 "The I.M.A.G.E. Consortium: An Integrated Molecular Analysis of Genomes and their Expression", Lennon, G.G., Auffray, C., Polymeropoulos, M., and Soares, M. B. [1995] Genomics.

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SCIENCE (ISSN 0036-8075) is published weekly on Friday, except the last week in December, by the American Association for the Advancement of Science, 1200 New York Avenue, NW, Washington, DC 20005. Periodicals Mail postage (publication No. 484460) paid at Washington, NW, Washington, DC 20005. Periodicals Mail postage (publication No. 484460) paid at Washington, DC, and additional mailing offices. Copyright © 1997 by the American Association for the Advancement of Science. The title SCIENCE is a registered trademark of the AAAS. Domestic individual membership and subscription (51 issues): \$105 (\$58 allocated to subscription). Domestic institutional subscription (51 issues): \$260. Foreign postage extra: Mexico, Caribbean (surface mail) \$55; other countries (air assist delivery) \$90. First class, airmail, student, and emeritus rates on request. Canadian rates with GST available upon request, GST #1254 88122. IPM #1069624. Printed in the U.S.A.

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# THIS WEEK IN SCIENCE

#### edited by PHIL SZUROMI

#### **Faster together**

Diffusion of atoms and molecules on surfaces is important for catalytic processes. Although strongly bound systems, such as metals on metal surfaces, have been investigated for some time, the study of weakly bound systems, such as carbon monoxide on metals, has been more difficult, as measurement can more easily affect the molecular motion. Briner et al. (p. 257) performed scanning tunneling microscopy of carbon monoxide on copper surfaces and found that dimers and longer chains move faster, through single molecule jumps, than monomers; this difference is attributed to a reduction of vibrational entropy in the chains rather than to a difference in activation energy.

#### Pore alignment

Mesoporous materials, with typical pore sizes of a few nanometers, are complementary to zeolites, which have smaller pores and thus can only take up molecules up to a certain size. However, large-scale applications of mesoporous materials often require alignment of the pores over macroscopic length scales, which has been difficult to achieve. Tolbert et al. (p. 264) have devised a strategy whereby the precursor material, a liquid crystal, is aligned in a magnetic field; the alignment is preserved during the processing stages, resulting in a material with macroscopically aligned mesopores.

#### **Io-Jupiter feedback**

Io, the most volcanically active body known in the solar system and Jupiter's nearest large satellite, is intimately coupled to Jupiter's magnetosphere. Brown and Bouchez (p. 268; see the Perspective by McGrath, p. 237) observed sulfur ions (S<sup>+</sup>) in the Io plasma torus and sodium atoms in the neutral plasma cloud

#### **Recognized by their caps**

T cells recognize proteins and peptides through major histocompatibility complex (MHC) proteins, which bind peptides and "present" them to antigen receptors on T cells. However, T cells see more than just peptides—they can be activated by binding to certain lipids and glycolipids, which are presented by proteins that are distant cousins of those encoded in the MHC region. Moody *et al.* (p. 283) identified a glycolipid antigen from mycobacteria and did a detailed analysis of which parts of the glycolipid were critical for recognition. Recognition depended mainly on hydrophilic cap of the glycolipids; naturally occurring substitutions of the hydrophobic tail were well tolerated. This result is consistent with the recently solved structure of CD1 both sets of work support the concept that the deep CD1 binding groove is designed to accept a hydrophobic tail, which allows the carbohydrate cap to stick out and become a target for T cells.

associated with the jovian magnetosphere for 6 months. During this time, a volcanic eruption on Io created a large plume that increased the Na content of the neutral cloud, followed by a delayed increase in the S<sup>+</sup> content



of the lo torus. These excess  $S^+$  ions dissipated nonlinearly, which suggests that the jovian magnetosphere stabilizes itself with a nonlinear feedback mechanism against the sporadic volcanic outbursts of lo.

#### Trapped organics on Ganymede and Callisto

Jupiter's largest satellites, Ganymede and Callisto, show evidence for water ice on their surfaces that may have been trapped early in the evolution of our solar system. McCord *et al.* (p. 271), using near-infrared spectra from the Galileo orbiter, found evidence for  $CO_2$ ,  $SO_2$ , and some organics trapped within the water ice and hydrated minerals on the surfaces. Such components could be derived from a variety of sources, including interstellar ices, the jovian magnetosphere, and meteorites. Fitting the spectra with these sources will help define how and where carbon, hydrogen, and oxygen were distributed in the early solar system.



#### Pain relief

Prolonged pain can be extremely debilitating, but sensitivity to painful stimuli is critical for the avoidance of serious injury. Two reports look at the mechanisms that generate acute pain but that allow the avoidance of longterm pain, or extreme sensitivity, to what should be nonpainful stimuli (see the Perspective by Iadarola, p. 239). When noxious stimuli are applied, a peptide, known as substance P (SP), is released and binds to neurons possessing the SP receptor. The role of these neurons in acute and chronic pain perception has been unclear; Mantyh et al. (p 275) now show that these neurons can be specifically removed by applying SP coupled to a toxin. The removal of these neurons did not change the animals' responses to acute pain but made them far less sensitive to chronic pain. Peripheral nerve injury can cause an increased sensitivity to pain (neuropathic pain). Malmberg et al. (p. 279) found that this pain state requires the expression of a brain-specific form of protein kinase C (PKC $\gamma$ ). Mice genetically engineered to lack the PKC $\gamma$  protein had a normal sensitivity to acute pain but were almost totally lacking in neuropathic pain after injury.

#### Viral infection and angiogenesis

Continued analysis of proteins encoded by the Kaposi's sarcoma virus HHV-8 are providing insights into pathogenesis and potential as therapeutic agents against HIV. Boshoff et al. (p. 290) found that two viral chemokine-like proteins, vMIP-I and -II, promoted the formation of blood vessels, which may be important in the development of Kaposi's sarcoma. The vMIP-II can activate and attract certain white blood cells (eosinophils), primarily through the CCR3 receptor. HIV infection through this receptor is also the most sensitive to inhibition by vMIP-II.

# Unraveling actin during cell death

When a cell gets a signal to initiate the programmed cell death cascade, a series of proteases called caspases are activated by cleavage of one another. Other proteins can also serve as substrates for caspase cleavage, and Kothakota et al. (p. 294) show that gelsolin, a protein that severs actin chains in a calciumdependent fashion, is cut during apoptosis. This cleavage liberates an activated gelsolin fragment that depolymerizes actin into monomers in an unregulated fashion and which is responsible for the morphologic changes associated with apoptosis. Cells that had no gelsolin had delayed apoptosis and only limited morphologic changes. Thus, gelsolin seems to potentiate the cell death process, a role consistent with its recently noted down-regulation in many human tumors.

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# **MJ** RESEARCH NOTEBOOK



Volume VII...No. 2

A Bulletin of Technological Advance in Molecular Biology

Autumn 1997



DNA Engine with interchangeable Alpha<sup>™</sup> blocks. The one mounted on the cycler has a motorized Power Bonnet<sup>™</sup> heated lid; the other Alphas—both dual and single block models—have manual Hot Bonnet<sup>™</sup> heated lids.

### Twin Tower *In Situ* Alpha Fits DNA Engines & Tetrads

The versatility of the DNA Engine system is well illustrated by the Twin Tower<sup>M</sup> in situ Alpha. Two new methods of DNA amplifica-



w methods of DNA amplification—*in situ* PCR\* and PRINS have recently come to common usage, but they usually require a specialty thermal cycler that processes only glass slides.

But not with the DNA Engine or Tetrad. Instead, a "swappable" dual block, called the Twin Tower, can be put into either chassis, and each Alpha cycles 2 x 16 glass slides rapidly, accurately, and precisely.

## Accuracy & Data Export Needed for Diagnostics

Medicine is on the cusp of a new era when diagnosis of disease will be based increasingly upon the analysis of amplified DNA. But the thermal cyclers that actually do the amplification must be of specific and certifiable quality.

The College of American Pathologists and the NCCLS have chosen to focus upon two criteria for particular attention: accuracy and the recording of thermal data from every run. Each DNA Engine has NIST-traceable & field-verifiable accuracy. Thermal data may be continuously reported via serial or GPIB ports, or sent to a printer for hard copy, in order to record data.

# Tetrads Lead the Charge into Genomics

High-Capacity Yet Compact—Cycler Holds Four 96 or 384-well Plates

A recent development in biology is an unrelenting surge of interest in genomics. Already increasing knowledge of the human and other genomes has had major influence upon the biotech, pharmaceutical, and medical industries, and Wall Street seems to be betting on further advances. According to a recent series of articles in *Science* (275, 767-782, 7 Feb 97) the pharmaceutical industry alone has invested at least \$1 billion into genomics companies.

And what is the thermal cycler of choice among these companies? Why, the MJ RESEARCH • PCR is covered by patents owned by Hoffmann-La Roche. Inc. & F. Hoffmann-La Roche Ltd. Users should obtain license to perform the reaction.



Tetrad cycler with four swappable blocks

Tetrad. This thermal cycler has all the features of the DNA Engine—and it also offers 4 independent blocks in one compact chassis. It fits easily inside robots, it can be controlled manually from its keypad or digitally through its ports, and when fitted with Power Bonnets, it can be operated without manual intervention.

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# DNA Engine & Tetrad Set New Standards for Thermal Cyclers

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#### Intuitive Programming Especially Easy

WATERTOWN, Mass. — In late 1994, MJ RESEARCH introduced a revolutionary new line of thermal cyclers for PCR<sup>\*</sup> and cycle sequencing. Called the "DNA Engine<sup>TM</sup>" line, this new design builds upon experience the company had

gained manufacturing its pioneering PTC-100<sup>™</sup> and MiniCycler<sup>™</sup> instruments. Improvements included increased speed of ramping, higher precision and greater accuracy—as well as a new "swappable" block format. But perhaps the most significant advances were in the software, for these allowed easier user interface and better thermal control—as well as communication between cyclers and other digital devices.

The new software was the culmination of years of effort by engineers Joe Pacatte and Andrea Wolga. Building upon the intuitive concepts of PTC-100 software, they managed to create a powerful new system that allows for expansion, revision, and network communications via IEEE-488 or RS-232 ports-features that would impress any engineer. But they concentrated their efforts on the needs of users, creating a whole host of features to allow smoother functioning in the lab. These include the ability to store programs in individual folders, edit one program while another is running, choose from 3 different methods of thermal control, and calculate total run time from entered parameters. These and many other software features make DNA Engines a joy to use.

Other engineers did their jobs well too. In particular, the modular design of swappable blocks (called "Alphas") allows configuration for different vessels—96 or 384-well plates, 0.2 or 0.5ml tubes, for example. Last but not least, two available sizes of chassis—one holding 1 Alpha and the other 4—allows maximum flexibility in planning, purchase, and expansion. E-MAIL: SALES@MJR.COM • WEB: WWW.MJR.COM

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