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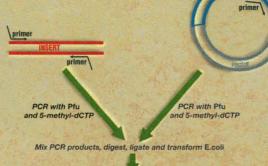
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Padgett, K.A., and Sorge, J.A. (1996) Gene 168: 31-35

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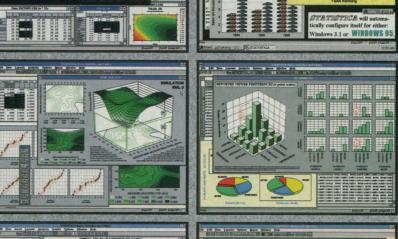
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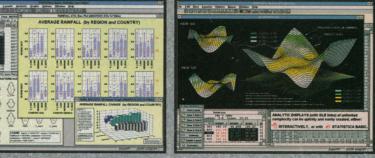


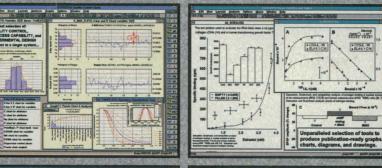
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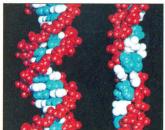
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COVER

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sciences. In the fall of 1996, Gordon Conferences will convene at Queens College, Oxford (cover); Pruhonice in the Czech Republic; and two sites in Japan. See page 826 for scheduling information. [Photo: Chris Andrews, Oxford Picture Library]



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792 & 795
Stretching DNA to the limit

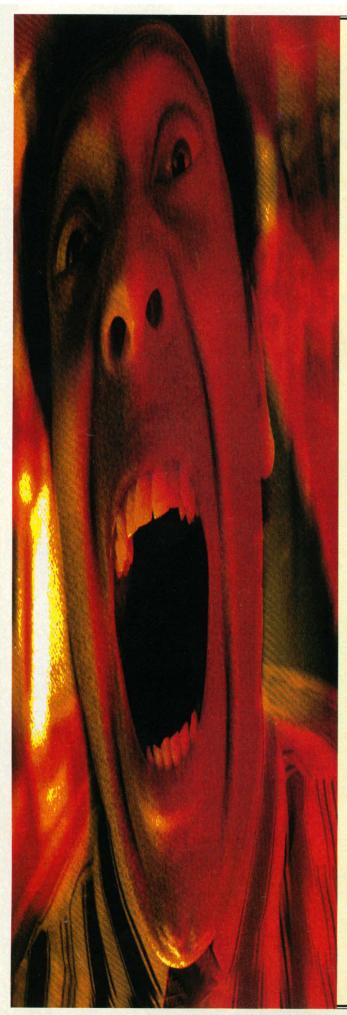
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1. Bernard, P. et al. (1994) Gene 148: 71-74.

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THIS WEEK IN SCIENCE

edited by DAVID VOSS

Icy breakup

Recently, several large ice shelves in Antarctica have broken up or diminished in size. These ice shelves border large ice sheets. Rott et al. (p. 788) analyze radar images from the ERS-1 satellite and show that the Larsen Ice Shelf, which covered an area of 4200 square kilometers, fractured and disintegrated within a few days. In a Perspective, Fahnestock (p. 775) amplifies some of the possible implications for understanding the origin and dynamics of the larger ice shelves.

Pulling hard on DNA

Laser tweezer methods make it possible to manipulate large molecules in solution. Attachment of latex beads to the ends of DNA strands allows tension to be applied to the strand so that its mechanical properties (extension, given applied force) can be determined. Two reports by Cluzel et al. (p. 792) and Smith et al. (p. 799) show that at relatively high forces (70 piconewtons), there is a transition where the DNA suddenly stretches by 70 percent. Smith et al. show that the force plateau matches that for singlestranded DNA, suggesting that nicks along the double-strand lead to unraveling of the base pairs. These results may be relevant to recombination—such a transition may reduce the energy needed for proteins such as RecA to stretch DNA.

Water on acid catalysts

Solid acid catalysts, such as aluminosilicates and silicoaluminophosphates, are workhorses of industrial and petrochemistry, yet the nature of acidic species

Under the reaper

In *Drosophila*, the *reaper* gene (*rpr*) appears to play a major role in mediating apoptosis, and sequence data suggest similarities with mammalian apoptotic pathways. Two reports focus on mechanistic aspects of *reaper* expression. White *et al.* (p. 805) generated transgenic flies that allowed control of expression of the *rpr* gene.

Overexpression of rpr in the fly retina resulted in compound eye ablation in a dosage-dependent manner. Pronk et al. (p. 808) induced expression of the RPR protein in Drosophila Schneider cells and showed that this in-





creased ceramide production. In both reports, the apoptotic effects could be blocked by protease inhibitors. This result suggests that an interleukin-1 β converting enzyme (ICE)-like protein plays a role in activating apoptosis.

generated at their surfaces is difficult to determine experimentally. Water is often used as a probe for acidity—does it form the H₃0+ cation, or is only hydrogen-bonded water present? Smith et al. (p. 795) used powder neutron scattering and infrared spectroscopy to study water bound to a microporous synthetic catalyst, HSAPO-34. that converts methanol to light alkenes. The ordered structure of the catalyst allowed both H₁0⁺ and hydrogen-bonded water to be identified at two different sites. In a Perspective, Sauer (p. 774) discusses these results in relation to recent quantum chemical calculations.

Making mast cells

Understanding the origins and development of mast cells, which play a key role in allergic and inflammatory reactions, is of relevance to both basic and clinical science. Rodewald *et al.* (p. 818) report the identification of a committed mast cell precursor. It can be distinguished both morphologically,

by the presence of cytoplasmic granules, and by its cell surface phenotype. It appears early in ontogeny, being found at day 15.5 of gestation in mice in the fetal blood; thus, commitment to the mast cell lineage can precede tissue emigration.

Insulin secretion

Diabetics are frequently treated with a group of drugs known as sulfonylureas. These compounds stimulate secretion of insulin from pancreatic β cells by inhibiting certain potassium channels. Eliasson *et al.* (p. 813) now show that in addition to the known indirect effects on plasma membrane channels, these drugs directly interfere with the cell's secretory machinery.

Cold comfort

To adapt to the cold, poikilothermic animals restore fluidity to cold-rigidified membranes by increasing the unsaturation of membrane phospholipids. Tiku *et al.* (p. 815) have cloned the enzyme, Δ^9 -desaturase from carp that incorporates the first degree of unsaturation into saturated fatty acids. They also examined the enzyme's regulation in carp during the exposure to cold. Cold induced an increase in Δ^9 -desaturase activity that was the result of increased transcription and the activation of latent desaturase.

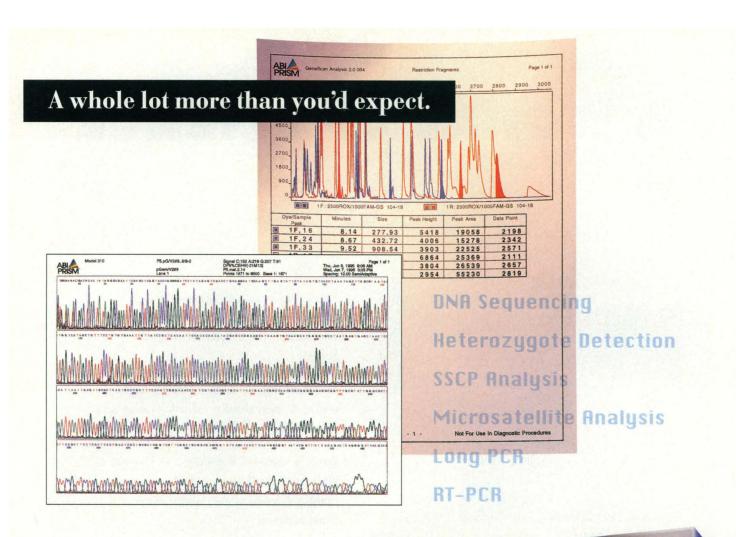
B cell activation

Human X-linked agammaglobulinemia is caused by a defective signaling pathway involving Bruton's tyrosine kinase (BTK) in B cells that results in a drastic loss of γ-globulin and antibody production. Several studies have suggested that BTK interacts with SRC family kinases. Rawlings et al. (p. 822) now show that SRC kinases transphosphorylate BTK at one site and that autophosphorylation of BTK at a second site results in activation. These same sites are also phosphorvlated when B cells are stimulated by immunoglobulin M. Certain mutations in BTK identified in patients map to the activation loop of BTK.

Following the flow

The hydrologic cycle of evaporation and precipitation on land provides nearly all of the fresh water used to sustain humanity and other life. Postel et al. (p. 785) provide an accounting of the use of this renewable fresh water by humans, including for agriculture, navigation, drinking, and other necessary activities. They conclude that humans appropriate 24 percent of evapotranspiration globally, and 54 percent of the runoff that is generally accessible (for example, flood waters are excluded).





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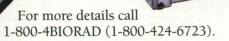
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References

- 1. Fodde, R., and Loosekoot, M., Human Mutation, 3, 83, (1994).
- 2. Fischer, S.G. and Lerman, L.S., Proc. Natl. Acad. Sci., U.S.A., 80, 1579 (1083)
- 3. Borresen, A. L., et al., Proc. Natl. Acad. Sci., U.S.A., 88, 8405 (1991).



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Winter 1996

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Fully Compatible With Earlier DNA Engine; Uses Same Interchangeable Alpha™ Blocks

WATERTOWN, Mass. – MJ RESEARCH proudly announces the introduction of an ultra-high-capacity model in its DNA Engine™ line of thermal cyclers. Called the PTC-225 DNA Engine Tetrad™, this speedy cycler has four fully-independent blocks, accurate and reliable Peltier-Joule heat pumps, and networking capabilities that make the cycler fully compatible with earlier DNA Engines—as well as with new automated systems.

In fact, the Tetrad cycler uses the same Alpha[™] sample-block/heat-pump assemblies that fit the earlier PTC-200 DNA Engine. These interchangeable blocks deliver the same thermal precision and NIST-traceable accuracy no matter what machine they are plugged into-and swapping an Alpha takes just ten seconds. Eight different Alphas are currently available, and they fit 0.5ml or 0.2ml tubes, 96-well or 192-well plates—or even combinations of vessels in dual blocks. Two newer blocks fit 384-well plates (see below) and microscope slides for in situ reactions. These different Alphas can be mixed or matched in a single Tetrad, for a total capacity of up to 1536 simultaneous reactions. No cycler made by any other manufacturer offers such versatility or throughput.

This instrument is the latest in the long line of Peltier thermal cyclers offered by MJ RESEARCH. Since 1988, this innovative manufacturer has pioneered development of Peltier instrumentation for laboratories, having introduced its first PTC-100 cycler that year. This venerable instrument was later followed by three models of the portable PTC-150 MiniCycler™ as well as five more models of PTC-100. MJ RESEARCH is also the company that blazed the trail to *in situ* amplification, and its line of PTC-200 DNA Engines sets a standard against which all other cyclers are now judged.

FOR MORE INFO, PLEASE CALL OR E-MAIL: SALES@MJR.COM



Manufacturer of Peltier-effect Thermal Cyclers 149 Grove St. • Watertown, MA 02172 U.S.A. (800) 729-2165 • Fax (617) 923-8080 Distributors Worldwide—Please Fax or E-mail for List



The PTC-225 Tetrad™ with four independent blocks, each with its own Hot Bonnet™ heated lid.

Automated Systems

Slim Cycler Works Well With Robots

The Tetrad was designed to integrate easily with robotic and automated systems, and its hardware and software features were carefully crafted to make integrations straightforward and reliable.

For example, an important consideration with robots is geometry. Thus, the Tetrad has a compact footprint (37 x 55cm), low height (25cm), and frontback airflow—features that facilitate easy fit into a robot without excessive occupation of the work envelope. Further, motorized Power Bonnet heated lids are available, and these open a full 115° to allow easy access to the block area. They operate automatically and use variable-ratio, dual-overhead cams to seat the heated lid firmly and evenly.

NETWORKING SOFTWARE

Control Can be Effected Through Keypad or Computer

Perhaps the most impressive feature of the DNA Engine line of thermal cyclers (the PTC-200 & PTC-225) is the sophisticated networking software that is exclusive to MJ RESEARCH. Not only does the software offer three methods of thermal control, advanced editing and filing features, and multi-tasking capability—it also allows up to fifteen cyclers to operate on a single, computerized network. Full control can be effected by a central computer through either a RS-232 or an IEEE-488 port, or alternatively, the individual blocks can be programmed or controlled through use of a keypad and the LCD/LED displays on the cycler itself.

"What About the 384-Well Format?" Ask Scientists in the Human Genome Community

The Quest for Colossal Capacity

Now that thermal-cycle sequencing of M13 templates seems to have become the sequencing



V-wells on 4.5mm centers (actual size) method of choice for the Human Genome Project (*Science* **267**, 783-4; *Nature* **375**, 93-4), investigators are faced with the engineering chore of scaling up equipment. Three billion bases in human DNA

need decoding, and the older standard format of disposable vessel—96-well plates—is generally too small for this sort of large-scale investigation.

Thus, a new 384-well format is in development. It shares the same basic V-well shape as the 0.2ml

96-well format, but the density has been multiplied four times by decreasing the well-to-well distance from 9mm to 4.5mm. This allows the use of the same multi-channel pipettors and automated dispensers/harvesters as with the 96-well format; the equipment merely accesses alternate wells in a back-and-forth fashion. MJ RESEARCH is working with others to establish standards for disposables, and although vessels (and the full utility of the system) are not yet available, 384-well Alphas for PTC-200 & 225 cyclers can be ordered. Reactions must now be conducted in 96-well plates; these vessels fit the 384 block adequately, but useful reaction volume is decreased to 20µl per well.

PCR is covered by patents owned by Hoffmann-La Roche, Inc. and F. Hoffmann-La Roche Ltd. Users should obtain license to perform the reaction.

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Access to Clones: Choose your clone based upon your homology searches or take advantage of the Genome Systems cDNA products and services.

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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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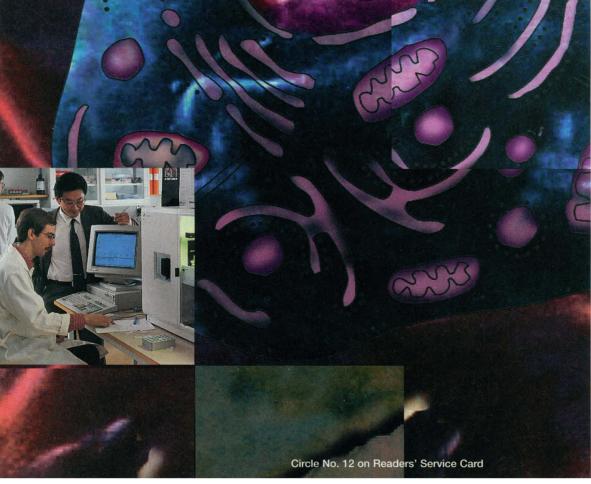
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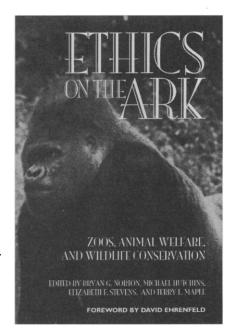
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