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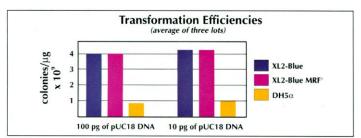
Stratagene has broken the 1 x 10 $^9$  barrier for competent *E.coli*! We did it by genetically modifying our Epicurian Coli<sup>®</sup> XL1-Blue and XL1-Blue MRF' cells. The result is our new XL2-Blue and XL2-Blue MRF' ultracompetent cells,\* which achieve transformation efficiencies as high as 5 x 10 $^9$  transformants/µg of plasmid control DNA. To maintain these ultrahigh efficiencies, we package the cells in convenient single-use aliquots that eliminate freeze/thaw cycles.

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\* Patent pending

\*\*The Polymerase Chain Reaction (PCR) process is covered by patents owned by Hoffmann-La Roche.

Use of the PCR process requires a license.



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Epicurian Coli®XL2-Blue Ultracompetent CellsCatalog # 200150Epicurian Coli®XL2-Blue MRF¹ Ultracompetent CellsCatalog # 200151Epicurian Coli®SURE® 2 Supercompetent CellsCatalog # 200152Epicurian Coli®XL1-Blue Subcloning-Grade Competent CellsCatalog # 200130



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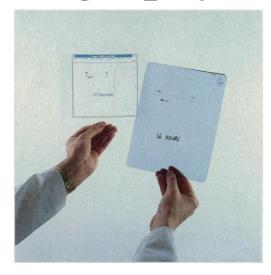
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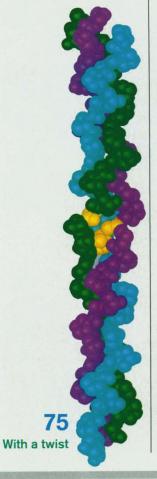
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# Science

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| NEWS & COMMENT  |               |
|---|---------------|
| Turning an Info-Glut Into a Library   | 20            |
| Bottleneck Keeps Existing Vaccine<br>Off the Market   | 22            |
| India Scrambles to Distribute Vaccine   | 23            |
| Euro-Russian Accord Begins With<br>30 Days of Solitude<br>Testing the Psychology of Would-Be Astronau | <b>24</b> its |
| cDNA Sequences: HGS Opens Its<br>Databanks—For a Price  | 25            |
| Europe's New X-ray Source Beams<br>With Confidence  | 26            |
| Grant Applications Pile Up at NIH   | 27            |
| RESEARCH NEWS   |               |
| Who Profits From Ecological Disaster? Between Extinctions, Evolutionary Stasis                        | 28<br>29      |
| Solar Farms May Reap Gamma Rays   | 30            |
| A Giant Licks Its Wounds  | 31            |
| Chemists Clean Up Synthesis With<br>One-Pot Reactions   | 32            |
| Crafting a Cage for Cholesterol   | 34            |

| urning an Info-Glut Into a Library   | 20        |
|--|-----------|
| Sottleneck Keeps Existing Vaccine Off the Market   | 22        |
| India Scrambles to Distribute Vaccine  | 23        |
| ouro-Russian Accord Begins With O Days of Solitude Testing the Psychology of Would-Be Astronau | <b>24</b> |
| DNA Sequences: HGS Opens Its<br>Databanks—For a Price  | 25        |
| Surope's New X-ray Source Beams<br>With Confidence   | 26        |
| Grant Applications Pile Up at NIH  | 27        |
| RESEARCH NEWS  |           |
| Who Profits From Ecological Disaster?<br>Between Extinctions, Evolutionary Stasis              | 28<br>29  |
| Solar Farms May Reap Gamma Rays  | 30        |
| A Giant Licks Its Wounds   | 31        |
|  |           |
| Chemists Clean Up Synthesis With One-Pot Reactions   | 32        |
|  | 32<br>34  |

|          | POLICY FORUMS   |  |  |  |  |  |
|----------|---|--|--|--|--|--|
| 20<br>22 | The Role of Biomedical Research in Health Care Reform M. W. Kirschner, E. Marincola, E. O. Teisberg |  |  |  |  |  |
| 22       | M. W. Klischher, E. Marincola, E. O. Teisberg   |  |  |  |  |  |
| 23       | The Paradox of Critical Mass for 51 Women in Science  |  |  |  |  |  |
| 24       | H. Etzkowitz, C. Kemelgor, M. Neuschatz, B. Uzzi<br>J. Alonzo                                       |  |  |  |  |  |
| its      | PERSPECTIVES  |  |  |  |  |  |
| 25       | Are Teflon "Ponytails" the Coming 55  |  |  |  |  |  |
| 26       | Fashion for Catalysts? J. A. Gladysz  |  |  |  |  |  |
| 27       | 14-3-3: Modulators of Signaling Proteins? D. Morrison   |  |  |  |  |  |
|          | ARTICLE   |  |  |  |  |  |
| 28       | ARTIOLE   |  |  |  |  |  |
| 29       | Physical and Biological Controls on Carbon Cycling in the Equatorial Pacific 58                     |  |  |  |  |  |
| 30       | J. W. Murray, R. T. Barber, M. R. Roman, M. P. Bacon, R. A. Feely                                   |  |  |  |  |  |
| 31       |   |  |  |  |  |  |
| 32       | RESEARCH ARTICLES   |  |  |  |  |  |
| 02       | A Strong Candidate for the Breast and Ovarian Cancer Susceptibility Gene BRCA1                      |  |  |  |  |  |
| 24       | V Mile I Swangan D Shattuck Fidens P A  |  |  |  |  |  |

| DEP  | AF        |
|--|-----------|
| THIS WEEK IN SCIENCE   | 9         |
| EDITORIAL Magnetic Energy Storage  | 11        |
| Smithsonian Science Exhibit: A. Molella; C. Wein<br>J. Lankford • Beta-Carotene and the Prevention<br>Cancer: J. R. Totter • HIV Prevention Program<br>D. R. Holtgrave and N. L. Qualls; M. M. Vilenchik | of<br>ns: |
| SCIENCESCOPE   | 19        |

| RESEARCH ARTICLES   |  |  |  |
|---|--|--|--|
| A Strong Candidate for the Breast and Ovarian Cancer Susceptibility Gene BRCA1 Y. Miki, J. Swensen, D. Shattuck-Eidens, P. A Futreal, K. Harshman, S. Tavtigian, Q. Liu C. Cochran, L. M. Bennett, W. Ding, R. Bell |  |  |  |
| IMENTS  |  |  |  |
| RANDOM SAMPLES 35   |  |  |  |
| QUARTERLY AUTHOR INDEX 82   |  |  |  |
| BOOK REVIEWS  Public Reactions to Nuclear Waste, reviewed by R. Cantor • Molecular Magnetism, P. Day • Design Paradigms, R. Mark • The Undivided Universe, D. M. Greenberger • Vignettes • Books Received           |  |  |  |

Frederick W. Alt Don L. Anderson Michael Ashburner Stephen J. Benkovic David E. Bloom Floyd E. Bloom Piet Borst Henry R. Bourne Michael S. Brown

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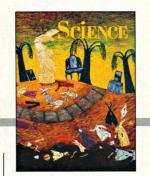
159

COVER

99

"Walking with the Ghosts of My Grandmothers" (1992), signature oil painting of artist Hollis Sigler's Breast Cancer Journal series. Through this series Sigler has tried to convey the emotional impact of a disease that affects one in nine women in the United States.

Two papers in this issue (see pages 66 and 120) report a major step forward in breast cancer research—the identification of the long-sought *BRCA1* gene, a critical determinant of hereditary breast cancer. [Photo: Michael Tropea]



J. Rosenthal, C. Hussey, T. Tran, M. McClure, C. Frye, T. Hattier, R. Phelps, A. Haugen-Strano, H. Katcher, K. Yakumo, Z. Gholami, D. Shaffer, S. Stone, S. Bayer, C. Wray, R. Bogden, P. Dayananth, J. Ward, P. Tonin, S. Narod, P. K. Bristow, F. H. Norris, L. Helvering, P. Morrison, P. Rosteck, M. Lai, J. C. Barrett, C. Lewis, S. Neuhausen, L. Cannon-Albright, D. Goldgar, R. Wiseman, A. Kamb, M. H. Skolnick

Facile Catalyst Separation Without Z 72
Water: Fluorous Biphase Hydroformylation
of Olefins

I. T. Horváth and J. Rábai

Crystal and Molecular Structure of a 75 Collagen-Like Peptide at 1.9 Å Resolution J. Bella, M. Eaton, B. Brodsky, H. M. Berman

#### **REPORTS**

Thorium-230 Ages of Corals and
Duration of the Last Interglacial Sea-Level
High Stand on Oahu, Hawaii
B. J. Szabo, K. R. Ludwig, D. R. Muhs, K. R.

B. J. Szabo, K. R. Ludwig, D. R. Muhs, K. R. Simmons

Interannual Variability of Temperature at a 96 Depth of 125 Meters in the North Atlantic Ocean S. Levitus, J. I. Antonov, T. P. Boyer

Atomic-Scale Dynamics of a Two-Dimensional Gas-Solid Interface S. J. Stranick, M. M. Kamna, P. S. Weiss

Direct Determination of Grain Boundary Atomic Structure in SrTiO<sub>3</sub>

M. M. McGibbon, N. D. Browning, M. F. Chisholm, A. J. McGibbon, S. J. Pennycook, V. Ravikumar, V. P. Dravid

Time-Resolved Detection of Structural
Changes During the Photocycle of
Spin-Labeled Bacteriorhodopsin

H.-J. Steinhoff, R. Mollaaghababa, C. Altenbach, K. Hideg, M. Krebs, H. G. Khorana, W. L. Hubbell

### Cystic Fibrosis Heterozygote Resistance to Cholera Toxin in the Cystic Fibrosis Mouse Model

S. E. Gabriel, K. N. Brigman, B. H. Koller, R. C. Boucher, M. J. Stutts

The Mobile Flavin of 4-OH Benzoate 110 Hydroxylase

D. L. Gatti, B. A. Palfey, M. S. Lah, B. Entsch, V. Massey, D. P. Ballou, M. L. Ludwig

RNA Editing: Transfer of Genetic 114
Information from gRNA to Precursor
mRNA in Vitro

S. D. Seiwert and K. Stuart

Identification of the ron Gene Product as the Receptor for the Human Macrophage Stimulating Protein

M.-H. Wang, C. Ronsin, M.-C. Gesnel, L. Coupey, A. Skeel, E. J. Leonard, R. Breathnach

BRCA1 Mutations in Primary Breast and Ovarian Carcinomas

P. A. Futreal, Q. Liu, D. Shattuck-Eidens, C. Cochran, K. Harshman, S. Tavtigian, L. M. Bennett, A. Haugen-Strano, J. Swensen, Y. Miki, K. Eddington, M. McCLure, C. Frye, J. Weaver-Feldhaus, W. Ding, Z. Gholami, P. Söderkvist, L. Terry, S. Jhanwar, A. Berchuck, J. D. Iglehart, J. Marks, D. G. Ballinger, J. C. Barrett, M. H. Skolnick, A. Kamb, R. Wiseman

Phosphate-Regulated Inactivation of the Kinase PHO80-PHO85 by the CDK Inhibitor PHO81

K. R. Schneider, R. L. Smith, E. K. O'Shea

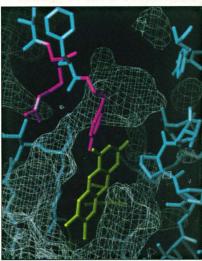
Interaction of the Protein Kinase Z 126
Raf-1 with 14-3-3 Proteins

H. Fu, K. Xia, D. C. Pallas, C. Cui, K. Conroy, R. P. Narsimhan, H. Mamon, R. J. Collier, T. M. Roberts

Association of the Protein Kinases c-Bcr and Bcr-Abl with Proteins of the 14-3-3 Family

G. W. Reuther, H. Fu, L. C. Cripe, R. J. Collier, A. M. Pendergast

110 Moving images



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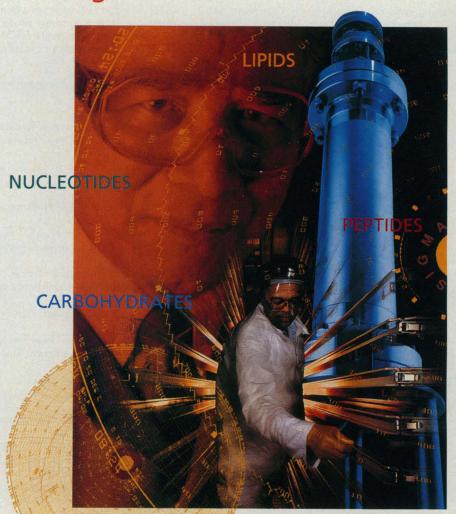
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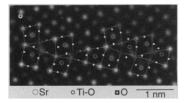
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#### THIS WEEK IN SCIENCE

edited by PHIL SZUROMI

#### Going with the grain

Although grain boundaries in crystals play an important role in process such as diffusion and electrical conduction, their atomic structure is difficult to determine. McGibbon *et al.* (p.



102) present a structure of a grain boundary in a SrTiO<sub>3</sub> bicrystal determined from a combination of high-resolution scanning transmission electron microscopy images, electron energy loss spectroscopy, and bond-valence sum calculations.

#### **High water mark**

Dates of exposed ancient coral reefs provide a key record of sea level changes, particularly for the timing of interglacials when sea level was highest. A key question has been the timing of the last interglacial. Szabo et al. (p. 93) present 29 ages from 17 samples of reefs formed during the last interglacial on Oahu, Hawaii, which imply that sea level peaked 1 to 2 meters higher than today about 131,000 years ago. This date is somewhat earlier than that inferred from marine core records.

#### Ocean oscillations

Understanding of the variability of temperature of the oceans at depths even a few meters below the surface has been poor. An early analysis of one station in the North Atlantic implied that temperature oscillated with a period of about 10 years and decreased overall during the past

#### Biphase catalysis without the water

Two-phase systems are often used in solution chemistry as a tool for maintaining product purity and recovering expensive catalysts. In phase transfer catalysis, a catalyst is dissolved in an aqueous phase and transported by a lipophilic cation into the organic phase. However, side reactions with water can occur, and many organic compounds dissolve poorly in water. Horváth and Rabái (p. 72; see the Perspective by Gladysz, p. 55) overcame these difficulties by replacing the aqueous phase with a fluorocarbon phase. Catalysts can be rendered soluble in the fluorous phase by attaching fluorocarbon pendant groups. They present applications in olefin hydroformylation and rhodium catalyst recovery.

several decades. Levitus *et al.* (p. 96) have gleaned data from other stations in the Atlantic to extend this analysis. The data imply that temperature trends in the subtropical and subarctic gyres were opposite and that decadal-scale oscillations in temperature extend throughout the North Atlantic Ocean.

#### **Breast cancer gene**

A strong candidate for BRCA1, a gene on chromosome 17q that confers susceptibility to earlyonset breast and ovarian cancer, has been isolated by Miki et al. (p. 66). The gene specifies a protein containing a zinc finger. Germline mutations of the BRCA1 candidate gene were found in members of 17q-linked families who are presumed to segregate BRCA1 susceptibility alleles. A study of sporadic breast and ovarian tumors by Futreal et al. (p. 120) found no evidence of somatic mutations in the BCRA1 candidate, suggesting that the gene may not play a critical role in these more common cancers.

#### **Triple helixes**

Collagen and collagen proteins have a number of unusual sequence features, such as the occurrence of the small glycine residue at every third position, that cause them to adopt their characteristic triple-helix structure. Such a regular structure has made crystallographic studies difficult, in that these molecules often disorder in the solid state, so that only fiber diffraction data can be obtained. Bella et al. (p. 75) have obtained a high-resolution x-ray structure of a 30-residue collagen-like peptide in which one glycine has been modified to alanine. Their analysis reveals an extensive hydrogen-bonding network of the peptide with water molecules. Similar mutations occur in a number of human diseases; a local untwisting of the triple helix occurs in the vicinity of the substitution.

#### **Increased resistance**

One in 20 Caucasians carry the mutation responsible for causing cystic fibrosis (CF). These carriers are healthy, whereas people who inherit two copies of the mutated gene are likely to die before reaching adulthood. Gabriel et al. (p. 107) used the CF mouse model to assess to what extent heterozygous mice possess an advantage that might explain the high incidence of the mutated gene in the human population. Mice carrying one

copy of the mutated gene were less sensitive to cholera toxin. Carriers of the gene might have a selective advantage of resistance to cholera or to other forms of secretory diarrhea.

#### **Editing guidelines**

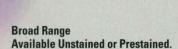
In the mitochondria of kinetoplastid protozoa, the sequence of a mature messenger RNA molecule is derived not only from the gene that encodes it, but also from a post-transcriptional editing process in which specific uridines are added or deleted. Seiwert and Stuart (p. 114) have developed an in vitro system for studying this editing process. Successful editing seems to depend upon adenosine triphosphate as well as upon base-pairing between the pre-mRNA and the guide RNA at the editing site.

#### More than cell cycle control

The function of cyclin-dependent kinases (CDKs) is not restricted to control of the cell cycle. The response of yeast to extracellular concentrations of phosphate is mediated by the CDK PHO85. In the presence of high concentrations of phosphate, an active complex of PHO85 and the cyclin PHO80 is formed, which leads to decreased expression of a secreted acid phosphatase. Schneider et al. (p. 122) show that in conditions where phosphate is limited, the PHO80-PHO85 complex is inhibited by PHO81, a protein with similarity to mammalian CDK inhibitors. Thus, similar mechanisms of regulation appear to modify the activities of CDKs that control either cell growth or responses to environmental stimuli.

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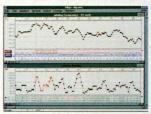
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# OLIGO 5.0 PCR SOFTWARE. WITH 22 NEW FEATURES, WHO KNOWS WHAT YOU MIGHT COME UP WITH?

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Now, NBI has ushered in a worthy successor: Oligo 5.0. And all of a sudden, everything else seems prehistoric.



So powerful, yet so easy. Just click on any oligo for instant analysis.

#### BETTER PCR PRIMERS, BY DESIGN.

If you're big on PCR, here's something you can sink your teeth into. Because with Oligo 5.0, you have the ultimate program for selecting *all* optimal PCR

primer pairs from any DNA/RNA sequence file. In seconds.

Its easy-to-use search function selects primers which are not only unique, highly specific, and crosscompatible, they're

also free of dimer, hairpin, and sequence repeats. Oligo 5.0 even searches for sequencing primers and hybridization probes. So you can design the best possible primers by simply lifting a finger.



What's more, it calculates optimal annealing temperature, PCR product composition, various Tm calculations, and extinction coefficients.

But perhaps most impressive is what we've added: Like LCR analysis, a GC clamp, primer Tm matching, highly accurate false priming checks, and nested primer selection.



All optimal PCR primer pairs are listed with complete analyses.

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Oligo 5.0 – for Windows or Mac – works so efficiently, so

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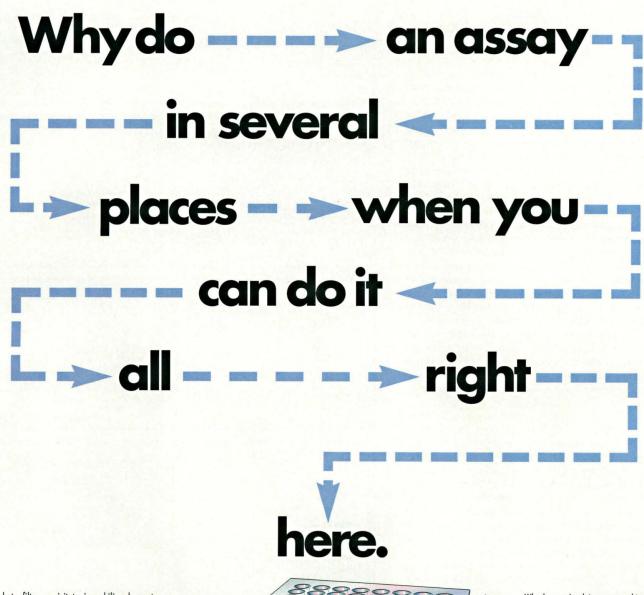
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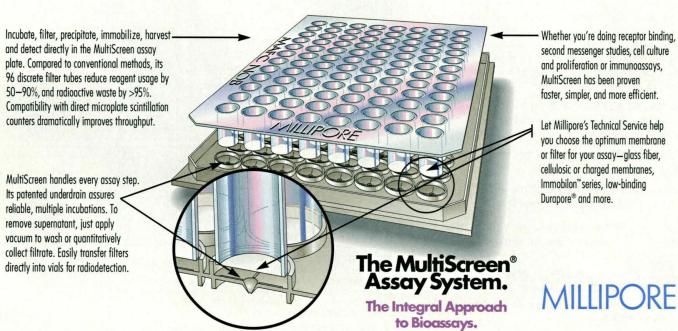
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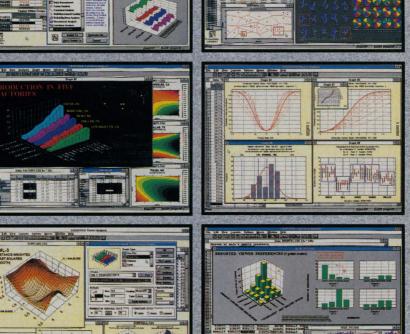


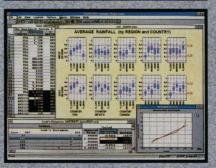
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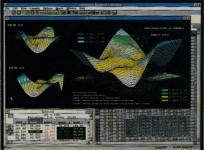
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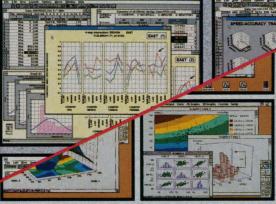
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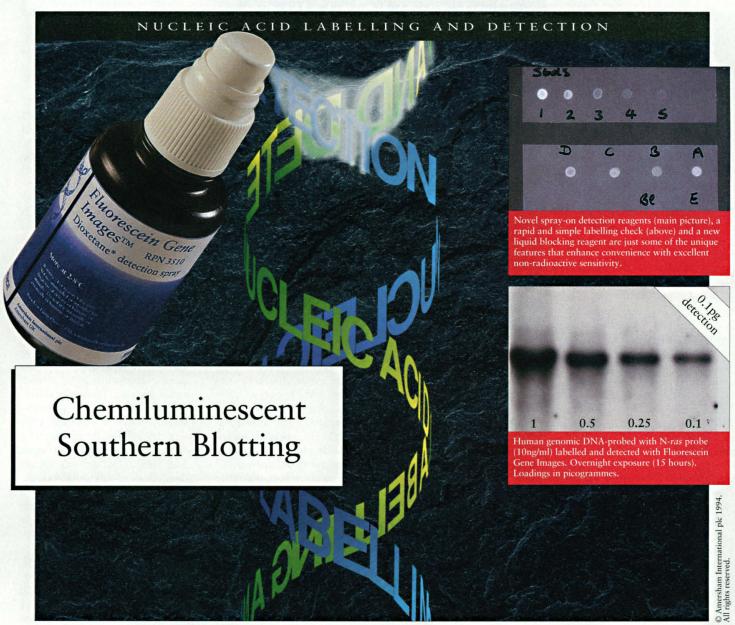






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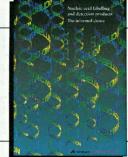


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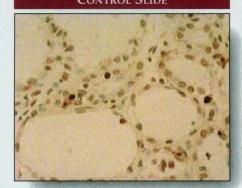
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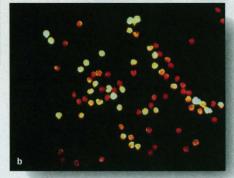
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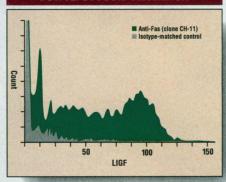
Apoptosis in 4-day post-weaning rat mammary gland (ApopTag<sup>™</sup> Control Slide), Apoptag<sup>™</sup> Peroxidase Kit and methyl green (X400)

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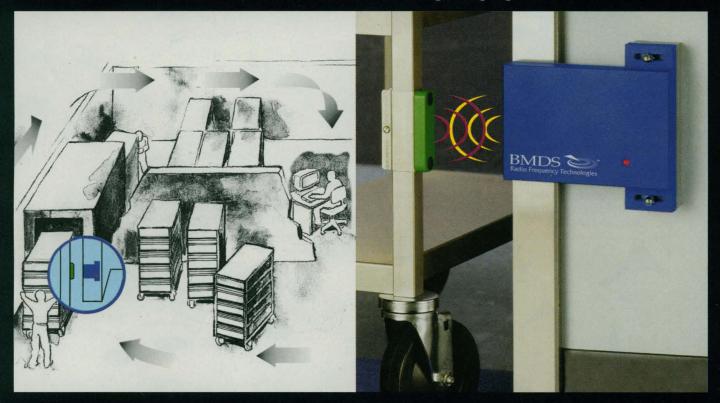
<sup>3</sup> X Li, W James, F Traganos & Z Darzynkiewicz, (1993) manuscript submitted.

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<sup>&</sup>lt;sup>2</sup> JFR Kerr, J Searle, BV Harmon & CJ Bishop, in: Potten, CS (ed) (1987) Perspectives in mammalian cell death.
Oxford U. Press, pp. 93-128. Z Zakeri, D Quaglino, T Latham & R Lockshin, (1993) FASEB Journal; 7:470-478; and manuscripts submitted.

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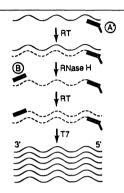
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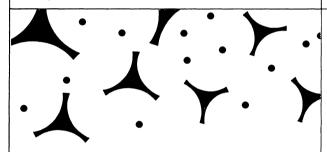
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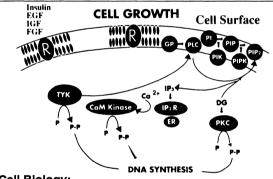
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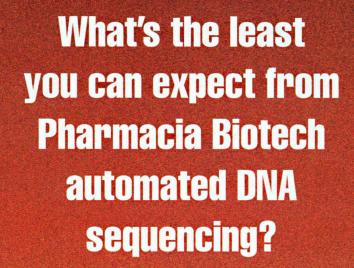
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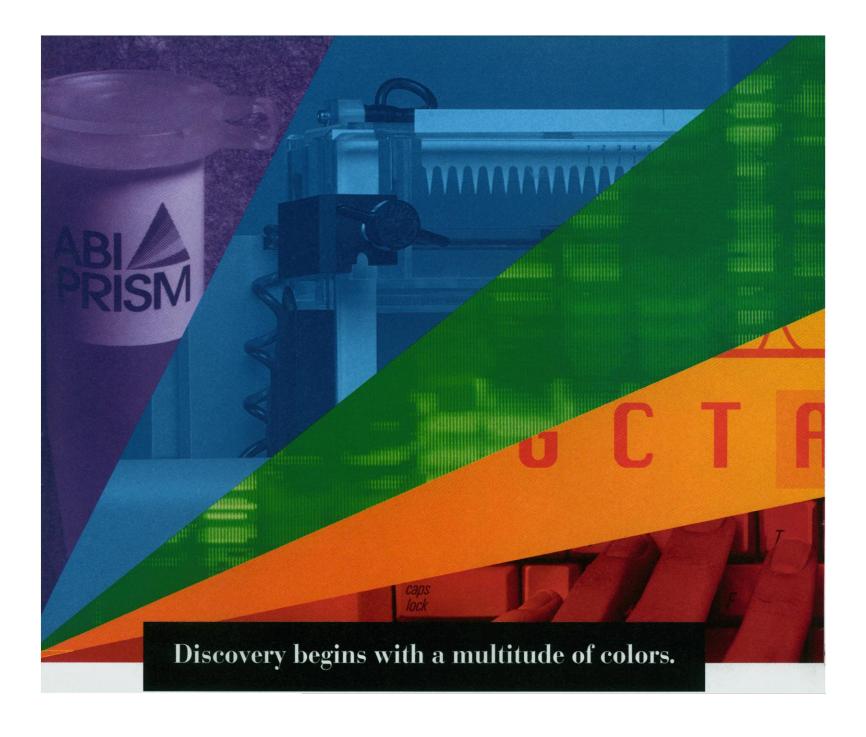
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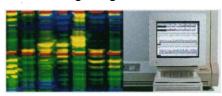
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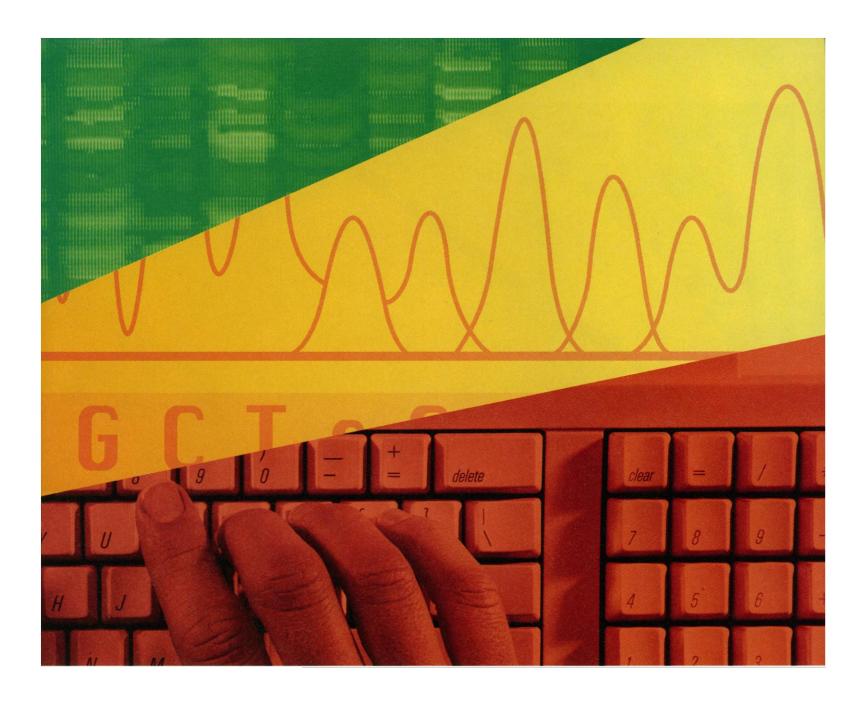
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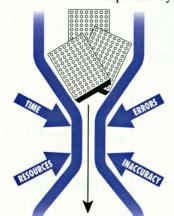
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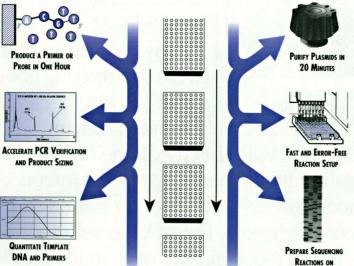
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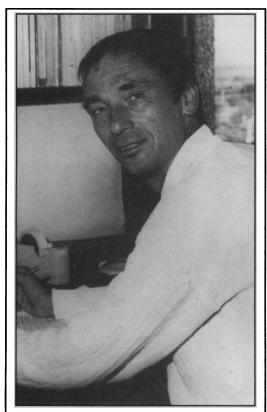
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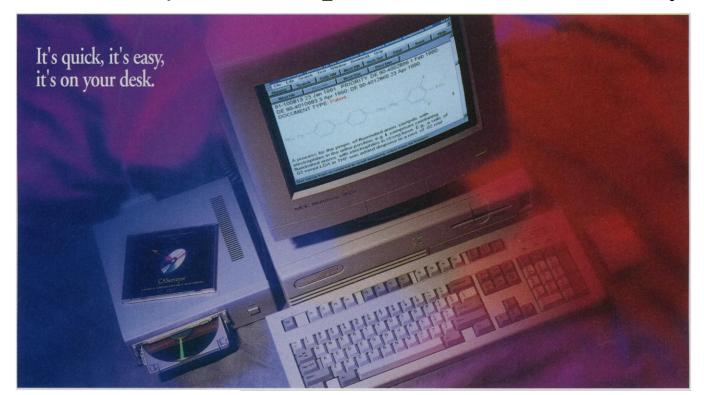
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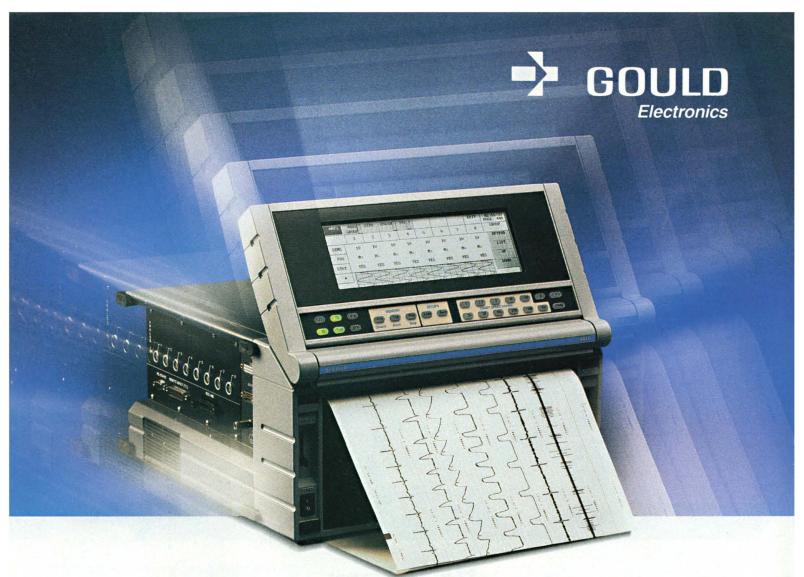
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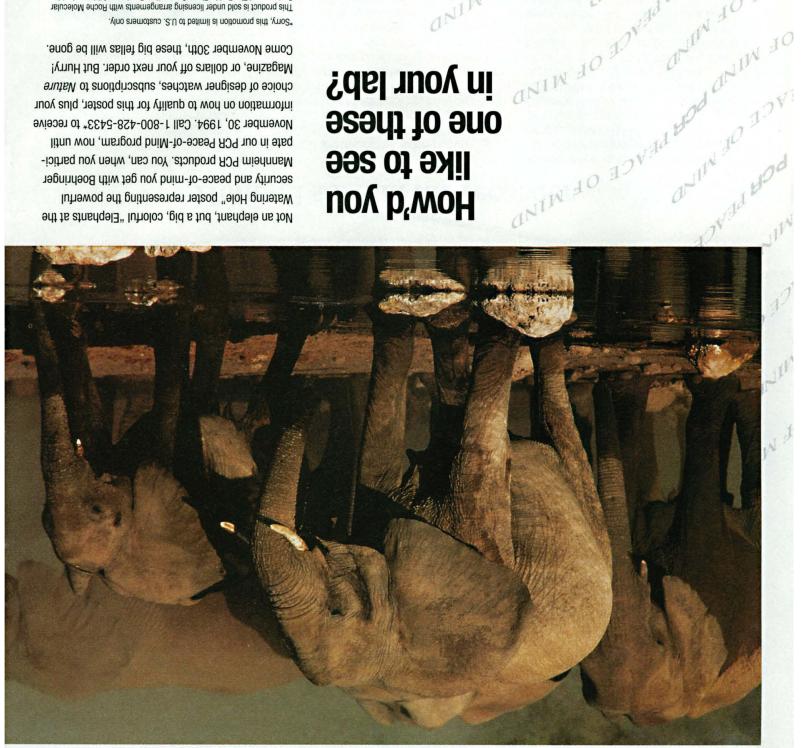
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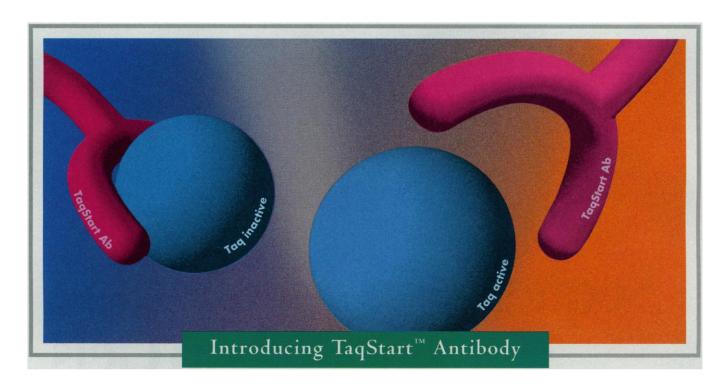
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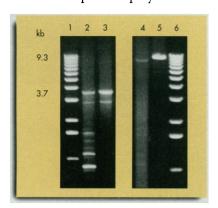
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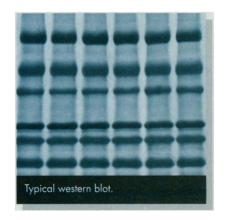
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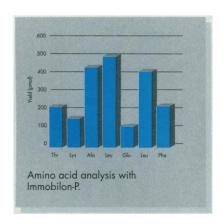
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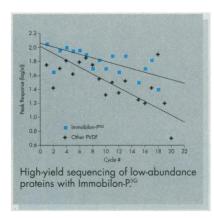
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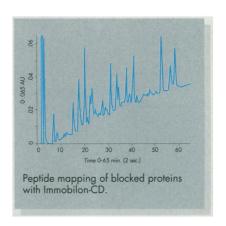




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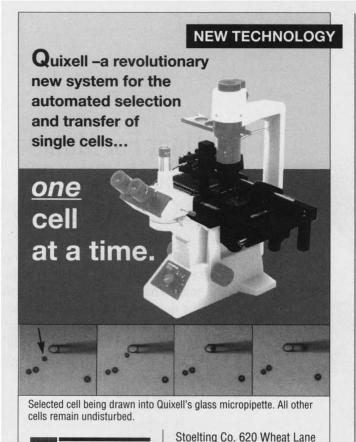
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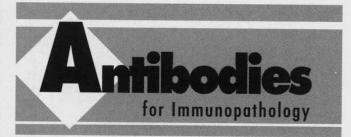
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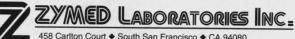
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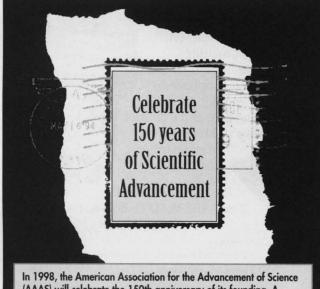
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- 1. Fodde, R., and Loosekoot, M., Human Mutation, 3, 83, (1994).
- Fischer, S.G. and Lerman, L.S., Proc. Natl. Acad. Sci., U.S.A., 80, 1579 (1983).
- 3. Borresen, A. L., et al., Proc. Natl. Acad. Sci., U.S.A., 88, 8405 (1991).



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