

# Why on earth would you ever use anything else?

# The answer is... you wouldn't.

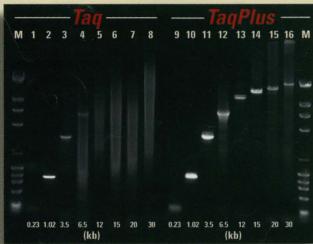
Because Stratagene's new TaqPlus™ DNA polymerase† does it all!

#### **ROBUST PERFORMANCE**

TaqPlus DNA polymerase boosts PCR yield. For long and short templates. For everyday PCR. Use it instead of Taq, and you'll understand the meaning of robust PCR!

#### **LONG PCR**

The robust nature of *TaqPlus* DNA polymerase makes it ideal for long PCR. It amplifies templates up to 35 kb long. So this potent polymerase is all you need.



#### **TOUGH TEMPLATES**

TaqPlus DNA polymerase reduces mismatch pausing. So more extension reactions reach completion in each cycle of PCR. Armed with this greater efficiency, TaqPlus DNA polymerase can even tame tough templates to bring out the bands.

Circle No. 44 on Readers' Service Card Electronic Marketplace: http://www.aaas.org TaqPlus<sup>™</sup> DNA Polymerase is an optimized blend of Stratagene's highest quality Taq DNA polymerase<sup>†</sup> and Taq Extender<sup>™</sup> PCR additive, the most powerful PCR-enhancement reagent available. <sup>1,2</sup>

## You'll never go back to Taq.

REFERENCES

1. Nielson, K.B., Scott, B., Bauer, J.C., and Kretz, K. (1994) Strategies 7: 64-65.

2. Nielson, K.B., Schoettlin, W., Bauer, J.C., and Mathur, E. (1994) Strategies 7: 27.

Nelson, K.B., Schoettlin, W., Bauer, J.C., and Mathu http://doi.org/10.1007/19.

POUNDLES.

Thurbase of this enzyme is accompanied by a license to use it in the polymerase chain reaction (PCRI process in conjunction with an authorized thermal cycler. This product is sold under licensing arrangements with Roche Molecular Systems, Inc., F. Hoffmann-La Roche Ltd. and The Perkin-Elme Corporation.

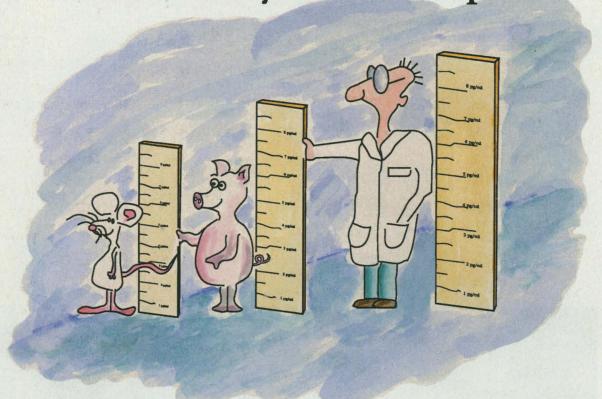
TaqPlus<sup>18</sup> polymerase 100 U Catalog #600203

INTERNET MAIL
tech\_services@stratagene.cor
USA
Corporate Headquarters
(800) 424-5444
GERMANY
Stratagene GmbH

UNITED KINGDOM Stratagene Ltd. Telephone: (0223) 420958 SWITZERLAND Stratagene (MI) 3641106



Human, Mouse & Pig Cytokine ELISA kits that really measure up.



**Superior Performance** 

Endogen ELISA kits are developed and manufactured to ensure the highest possible degree of sensitivity and precision in cytokine quantitation. The performance of every ELISA kit is documented

through a battery of validation studies to ensure that your cytokine measurements are accurate.

#### Reliable

Every Endogen kit lot meets stringent Quality Control specifications guaranteeing consistent, accurate results time after time.

#### **Small Sample Size**

We understand how precious your samples are. Only 50 µl of sample per well is required in our ELISAs.

#### Flexible

Each Endogen ELISA kit utilizes a consistent assay procedure for serum, plasma and culture supernatants, enabling you to test different sample types in the same assay. Depending on the number of samples to

be run, you choose whether to use the entire stripwell plate, or run two assays on separate occasions.

Endogen ELISA kits are fast, easy and reliable. Just collect your samples and go... start to finish.

#### **Technical Support**

Technical Service Specialists are standing by to answer your questions or fax datasheets on any Endogen ELISA kit or research reagent.

CALL TODAY!

#### ENDOGEN CYTOKINE ELISA KITS **Human ELISAs** • hIL-1a • hIL-6 · hIFNy • hIL-1B • hIL-8 • hIFNa • hIL-2 • hIL-10 • hTNFa · hGM-CSF · hIL-4 Pig ELISAs • pTNFa Mouse ELISAs • mIL-1a • mIL-4 · mGM-CSF • mIFNy • mIL-1B • mIL-5 • mIL-2 • mIL-6 • mTNFa • mIL-3 • mIL-10

Circle No. 13 on Readers' Service Card

640 MEMORIAL DRIVE, CAMBRIDGE, MA 02139 USA 617.225.0055 FAX 617.225.2040 For research use only. Not for use in diagnostic or therapeutic procedures.

Head Office for Europe: Biozol Diagnostica TEL 49+(89) 3192053

Biozol Diagnostica In Japan: Funakoshi Co., Ltd.
TEL 49+(89) 3192053 TEL 81+(3) 56841622
FAX 49+(89) 3193611 FAX 81+(3) 56841633

**ENDOGEN** 800-487-4885 You have this idea.

Could be great, could be a dead end. You're not sure yet because you can't see it.

So you knock out a 3-D model on your Power Macintosh." Render it. Rotate it. Shade it.

Run your model through a simulation. Ouch.



For more information by fax, call 800-503-3855; by mail, 800-732-3131, ext. 240. Or visit us on the Internet at http://www.apple.com.

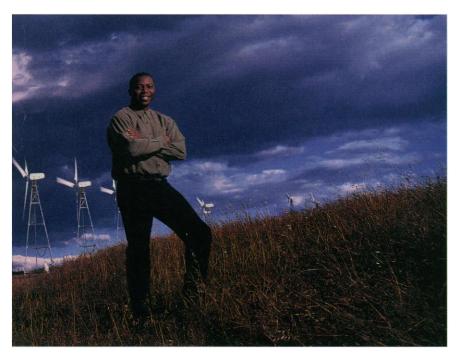
Revert to saved.

Modify your model. Run another simulation. Push it.

That's when you discover that it's only taken 30 minutes to figure out

that it isn't a dead end after all.

And it occurs to you:



It's how powerful the computer makes you.

## It's not how powerful the computer is.



Introducing the Power Macintosh 9500.

The power to be your best.



ISSN 0036-8075 8 SEPTEMBER 1995 VOLUME 269 NUMBER 5229

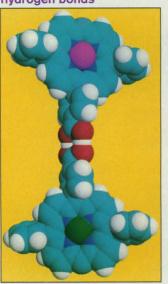
# Science





1332 Science and ethics in Africa

1409 Firing electrons through hydrogen bonds



#### **NEWS** Staging Ethical AIDS Trials in Africa 1332 A Model Collaboration Built to Last 1333 Testing AIDS Interventions: When Is the 1334 Price Too High? Japan Expands Graduate, Postdoc Slots 1335 1336 First Light From a Space Laser A Science Teaching System Honed in a 1337 Two-Room School Computer Processing Gives Imaging a 1338 Sharper View Chemists React to New Compounds 1339 in Chicago

#### PERSPECTIVES

| First-Principles Quantum Mechanics<br>E. Wimmer  |      |
|--|------|
| Digging into Caveolae<br>R. G. Parton and K. Simons  | 1398 |
| Tumor Suppression and Transcription<br>Elongation: The Dire Consequences of<br>Changing Partners  A. Krumm and M. Groudine | 1400 |

Computational Materials Design with

#### COMPUTERS '95: FLUID DYNAMICS

| FLUID DYNAMICS   |              |
|--|--------------|
| NEWS Indexing the Internet The Web-Crawler Wars  | 1354<br>1355 |
| On-Line Archives Let Biologists<br>Interrogate the Genome<br>The Visible Man Steps Out               | 1356<br>1358 |
| Off-the-Shelf Chips Conquer the<br>Heights of Computing<br>Swimming Against a<br>Supercomputing Tide | 1359<br>1360 |
| Mathematicians Open the Black Box of Turbulence  | 1361         |

#### RESEARCH ARTICLE

Inhibition of Transcription Elongation Inhibition of Transcription Elongation Inhibition Inhibition

#### REPORTS |

Macroscopic Evidence of the Effect of
Interfacial Slippage on Adhesion
B.-m. Zhang Newby, M. K. Chaudhury, H. R.
Brown

#### **DEPARTMENTS**

1397

| THIS WEEK IN SCIENCE EDITORIAL Wired or Wary?                                       | 1321<br>1323 | Plasma Physics and Fusion Research: D. Montgomery • Restoring Prince William Sound: R. B. Spies • Successful Grants Management: J. E. Dowling |                   |
|---|--------------|---|-------------------|
| LETTERS   | 1325         | SCIENCESCOPE  | 1331              |
| The Discipline of Epidemiology: W. V.   | Willett, S.  | RANDOM SAMPLES  | 1341              |
| Greenland, B. MacMahon, D. Trichop  |              | BOOK REVIEWS  | 1455              |
| Rothman, D. Thomas, M. Thun, N. Trichopoulos; J. Rapp; R. W. Miller; A              | . J. Saah;   | Dark Sun, reviewed by H. A. Bethe • Vignette ment at the Smithsonian • Books Received   | • Judge-          |
| G. B. Gori • Glass Ceiling: Bump, Bump<br>Picciotto and M. Hatch; S. Emerson and P. |              | PRODUCTS & MATERIALS  | 1458              |
| i icciotto and ivi. i lateli, o. Emerson and i .                                    | D. Coley     |   | The second second |

#### Board of Reviewing Editors

Frederick W. Alt Don L. Anderson Michael Ashburner Stephen J. Benkovic Alan Bernstein David E. Bloom Piet Borst Henry R. Bourne Michael S. Brown James J. Bull Kathryn Calame C. Thomas Caskey Dennis W. Choi David Clapham Adrienne E. Clarke John M. Coffin F. Fleming Crim Paul J. Crutzen James E. Dahlberg Robert Desimone Paul T. Englund Richard G. Fairbanks Douglas T. Fearon Harry A. Fozzard Klaus Friedrich Roger I. M. Glass Stephen P. Goff Peter N. Goodfellow Corey S. Goodman Peter Gruss Ira Herskowitz Tomas Hökfelt Susan D. Iversen

Eric F. Johnson Stephen M. Kosslyn Michael LaBarbera Nicole Le Douarin Charles S. Levings III Alexander Levitzki Harvey F. Lodish Richard Losick Reinhard Lührmann Diane Mathis Anthony R. Means Shigetada Nakanishi Kim Nasmyth Roger A. Nicoll Staffan Normark Stuart L. Pimm Yeshayau Pocker Dennis A. Powers Ralph S. Quatrano Martin Raff V. Ramanathan Douglas C. Rees T. M. Rice David C. Rubie Erkki Ruoslahti Gottfried Schatz Jozef Schell Ronald H. Schwartz Terrence J. Sejnowski Ellen Solomon Thomas A. Steitz Michael P. Stryker Robert T. N. Tijan Emil R. Unanue Geerat J. Vermeij Bert Vogelstein Arthur Weiss Zena Werb George M. Whitesides Owen N. Witte William A. Wulf

#### COVER

Simulation of highly turbulent compressible convection showing the vorticity (localized spin) of fluid motions concentrated into an evolving network of vortex tubes and rings near the upper surface. Such high-resolution fluid modeling of solar convection is now possible with

parallel computer systems. See page 1370 and the special section on computational fluid dynamics beginning on page 1353. [Simulation: D. Porter and P. Woodward on Cray T3D at Pittsburgh Supercomputing Center. Imaging: N. Brummell and D. Porter]



| Computer Scientists Rethink Their Discipline's Foundations   | 1363        |
|--|-------------|
| ARTICLES Keplerian Complexity: Numerical Simulations of Accretion Disk Transport J. F. Hawley        | 1365        |
| Turbulent Dynamics in the Solar<br>Convection Zone<br>N. Brummell, F. Cattaneo, J. Toomre            | 1370        |
| Modeling Ocean Circulation A. J. Semtner   | 1379        |
| Numerical Models of Caldera-Scale<br>Volcanic Eruptions on Earth, Venus, and Ma<br>S. Werner Kieffer | 1385<br>ars |

Direct Evaluation of Electronic Coupling 1409 Mediated by Hydrogen Bonds: Implications for Biological Electron Transfer P. J. F. de Rege, S. A. Williams, M. J. Therien

Synchrony and Causal Relations Between
Permian-Triassic Boundary Crises and
Siberian Flood Volcanism

P. R. Renne, Z. Zichao, M. A. Richards, M. T. Black, A. R. Basu

Short- and Intermediate-Range 1416
Structural Ordering in Glassy Boron Oxide
R. F. Youngman, S. T. Haubrich, I. W. Zwanziger

R. E. Youngman, S. T. Haubrich, J. W. Zwanziger, M. T. Janicke, B. F. Chmelka

Branched Threadlike Micelles in an 1420 Aqueous Solution of a Trimeric Surfactant D. Danino, Y. Talmon, H. Levy, G. Beinert, R. Zana

Reciprocal Stimulation of GTP Hydrolysis by Two Directly Interacting GTPases T. Powers and P. Walter

TRAF2-Mediated Activation of NF-κB by TNF Receptor 2 and CD40 M. Rothe, V. Sarma, V. M. Dixit, D. V. Goeddel Inducible Gene Targeting in Mice 1427 R. Kühn, F. Schwenk, M. Aguet, K. Rajewsky

The Chromodomain Protein Swi6: A Key
Component at Fission Yeast Centromeres

K. Ekwall, J.-P. Javerzat, A. Lorentz, H. Schmidt, G. Cranston, R. Allshire

A Simple Genetic Basis for a Complex
Psychological Trait in Laboratory Mice

J. Flint, R. Corley, J. C. DeFries, D. W. Fulker, J. A. Gray, S. Miller, A. C. Collins

Separation of Caveolae from Associated I 1435
Microdomains of GPI-Anchored Proteins
J. E. Schnitzer, D. P. McIntosh, A. M. Dvorak,
J. Liu, P. Oh

Elongin (SIII): A Multisubunit

Regulator of Elongation by RNA Polymerase II

T. Aso, W. S. Lane, J. W. Conaway, R. C. Conaway

Binding of the von Hippel–Lindau

Tumor Suppressor Protein to Elongin B and C

A. Kibel, O. Iliopoulos, J. A. DeCaprio, W. G. Kaelin Jr.

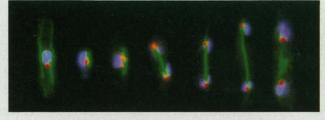
Decreased Muscarinic Receptor Binding 1446 in the Arcuate Nucleus in Sudden Infant Death Syndrome

H. C. Kinney, J. J. Filiano, L. A. Sleeper, F. Mandell, M. Valdes-Dapena, W. F. White

#### **TECHNICAL COMMENTS**

Possible Exceptions to Rules for α-Helix Termination by Glycine

E. L. Altschuler and M. Lades; G. Rose, R. Aurora, R. Srinivasan



Indicates accompanying feature

1429 Yeast Swi6 protein localized at centromeres

#### AAAS Board of Directors

Francisco J. Ayala Retiring President, Chairman Rita R. Colwell President Jane Lubchenco President-elect

William A. Lester Jr. Simon A. Levin Michael J. Novacek Anna C. Roosevelt Alan Schriesheim Jean E. Taylor Chang-Lin Tien Nancy S. Wexler

William T. Golden Treasurer Richard S. Nicholson Executive Officer ■ SCIENCE (ISSN 0036-8075) is published weekly on Friday, except the last week in December, by the American Association for the Advancement of Science, 1333 H Street, NW, Washington, DC 20005. Second-class postage (publication No. 484460) paid at Washington, DC, and additional mailing offices. Copyright © 1995 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): \$97 (\$50 allocated to subscription). Domestic institutional subscription (51 issues): \$228. Foreign postage extra Mexico, Caribbean (surface mail) \$53; other countries (air assist delivery) \$93. First class, airmail, student and emeritus rates on request. Canadian rates with GST available upon request, GST #1254 88122. Printed in the U.S.A.

Change of address: allow 6 weeks, giving old and new addresses and 11-digit account number. Postmaster: Send change of address to Science, P.O. Box 1811, Danbury, CT 06813–1811. Single copy sales: \$7.00 per issue prepaid includes surface postage; bulk rates on request. Authorization to photocopy material for internal or personal use under circumstances not falling within the fair use provisions of the Copyright Act is granted by AAAS to libraries and other users registered with the Copyright Clearance Center (CCC) Transactional Reporting Service, provided that \$3.00 per article is paid directly to CCC, 27 Congress Street, Salem, MA 01970. The identification code for Science is 0036-8075/83 \$3.00. Science is indexed in the Reader's Guide to Periodical Literature and in several procedure and indexes.

# NEW PRODUCTS for Soluble Protein Expression in E. coli.

Kpn I Sma I Sma I Xba I Aoc I Sa I I Pst I

#### End Insolubility

The ThioFusion™ Expression System from Invitrogen eliminates protein insolubility and the problems associated with it.

· No more difficult, time-consuming solubilization

 No more refolding of proteins that form inclusion bodies

With the unique E. coli protein thioredoxin as a fusion partner, formerly insoluble proteins are expressed in the soluble fraction. The ThioFusion™ Expression Kit contains the expression vector pTrxFus plus control vectors, essential media components, osmotic shock buffers, sequencing primers, and a comprehensive manual. Call for more information today!

ThioFusion™ Expression System

Catalog No. K350-01

Specific Detection

thioredoxin fusion protein with

the Anti-Thio™ Antibody. Anti-Thio™

is a monoclonal antibody that is

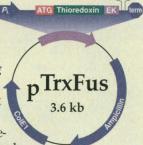
highly specific for thioredoxin. The antibody is ideal for

detection in western blots

and ELISAs. Call for more information.

Anti-Thio™ Antibody Catalog No. R920-25

Track expression of your



#### Simple Purification

Now you can purify your thioredoxin fusion proteins quickly and easily with ThioBond™ resin. This unique resin is specially formulated to specifically bind the active site of thioredoxin. Thioredoxin fusion proteins

stick, E. coli proteins don't. The result is pure thioredoxin fusion protein in less than 3 hours.

Call us today for more information.

ThioBond™ Resin (10 ml)

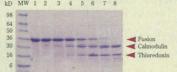
Catalog No. R350-10

#### Efficient Cleavage

The most efficient way to cleave your thioredoxin fusion protein is EnterokinaseMax™ (EKMax™). EKMax™ a recombinant preparation of the catalytic subunit of enterokinase, has many advantages over non-recombinant preparations including:

- EKMax<sup>™</sup> is free of contaminants
- EKMax<sup>™</sup> cleaves your fusion protein cleanly and efficiently

Digestion of Thioredoxin-Calmodulin Fusion with EKMax<sup>18</sup>



Lane 2: 0.004 units Lane 3: 0.007 units Lane 4: 0.015 units

Lane 6: 0.06 units Lane 7: 0.12 units

SDS-PAGE gel of a thioredoxin-calmodulin fusion protein digested with EKMax™ using the units shown on the right.

Don't waste your time and money on non-recombinant enzymes. Try EKMax™ today!

EKMax™ (250 units)

Catalog No. E180-01

Invitrogen BV
De Schelp 12, 9351 NV Leek
The Netherlands Tel: +31 (0) 594 515175 Fax: +31 (0) 594 515312

Australia 03 562 6888 Hong Kong 886 2 381 0844 Singapore 65 779 1919 or 29 29 783

From all other countries, please contact our European headquarters at +31 (0) 5945 15175

#### TOLL FREE TELEPHONE NUMBERS

#### ||| Invitrogen

The Gene Expression Folks. 1-800-955-6288

3985 B Sorrento Valley Blvd. San Diego, California 92121 Telephone (619) 597-6200 Fax (619) 597-6210

Circle No. 25 on Readers' Service Card

#### THIS WEEK IN SCIENCE

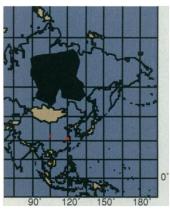
edited by PHIL SZUROMI

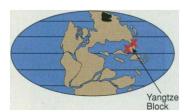
#### Giving them the slip

What makes adhesive tape stick better to some surfaces and not to others? Peeling is normally thought to be the way in which an adhesive is separated from its substrate, so adhesion strength should be proportional to surface free energies. Newby et al. (p. 1407) found that a transparent tape peeled off of a fluorocarbon surface less readily than for a corresponding hydrocarbon surface, despite the lower surface free energy of the fluorocarbon surface. These surfaces were silanes bearing different functional groups that were grafted onto silicon substrates. They also observed a fingered pattern in the region of adhesive failure and suggest that interfacial slippage at the adhesive interface may govern adhesive strength.

#### Volcanic extinction?

The most severe extinction in Earth's history was at the Per-





mian-Triassic boundary, 250 million years ago. One of the most massive episodes of volca-

#### **Tumor suppressor gives pause**

Patients with von Hippel–Lindau (VHL) disease are predisposed to a variety of cancers, including kidney cancer, and have been shown to carry germ-line mutations in the VHL gene. The wild-type VHL protein is thought to function as a tumor suppressor, but its exact cellular role has remained elusive. Duan et al. (p. 1402), Aso et al. (p. 1439), and Kibel et al. (p. 1444; see the Perspective by Krumm and Groudine, p. 1400) show that wild-type VHL inhibits the activity of a transcription elongation factor, Elongin, which normally functions to prevent pausing by RNA polymerase II during transcription. VHL mediates this inhibition by binding to Elongin's two regulatory subunits. The binding site on VHL includes a region that is frequently mutated in kidney cancer, suggesting that this interaction is important for its tumor suppressor function.

nic activity in Earth's history also occurred then, the Siberian flood basalts. Renne *et al.* (p. 1413) now provide a tantalizing chronologic link between these two events. They used argon-40–argon-39 dating to show that the age of volcanic rocks from southern China that mark the paleontologic boundary is indistinguishable (errors of a few hundred thousand years) from ages obtained in the same laboratory for the inception of the massive flood volcanism.

#### **Fully connected**

Surfactants can form a variety of aggregates (micelles) depending on their concentration. The viscosity of dimeric or trimeric surfactants, with two or three chains, increases very rapidly at low surfactant concentrations. and this has been attributed to branching of the micelles leading to a three-dimensional network. Such branching has also been predicted by calculations, but direct experimental evidence for branched micelles has not so far been obtained. Danino et al. (p. 1420) present transmission electron micrographs which show that trimeric surfactants can form such a branched network.

#### **Interacting GAPs**

Guanosine triphosphatases (GTPases) can act as molecular switches in various intracellular reactions. Two GTPases in Escherichia coli are thought to control protein targeting to the plasma membrane. Powers and Walter (p. 1422) changed the nucleotide specificity of each of the proteins so that instead of hydrolyzing GTP, they could hydrolyze xanthosine triphosphate. They then showed that the two GTPases interact with one another such that each protein stimulates the GTPase activity of the other proteinthus they act as mutual GTPase activating proteins, or GAPs. This control of reaction cycles by GTPases physically and temporally links both the activity and the inactivation of a participant in a reaction cycle.

#### **TRAF** traffic

The activation of members of the tumor necrosis factor (TNF) superfamily of receptors initiates an inflammatory response. TRAF (TNF receptor–associated factor) proteins are putative effectors involved in TNF signaling. Rothe *et al.* (p. 1424) examined the requirement for several TRAF proteins in the

TNF-R2 and the CD40 signaling pathways that activate nuclear factor  $\kappa B$  (NF- $\kappa B$ ). They find that TRAF2 is a common mediator of both TNF-R2 and CD40 signaling.

#### A matter of timing

Gene inactivation in mice has been targeted to particular cell types with the Cre/loxP recombination system of bacteriophage P1. Kühn et al. (p. 1427) have now used the inducible promoter of the mouse Mx1 to control the expression of a Cre recombinase transgene. The Mx1 gene, which is activated in response to viral infections, can be activated at a particular time in development in differentiated cells by application of interferon (IFN)  $\alpha$ , IFN- $\beta$ , or an IFN inducer. They used this method to delete the DNA polymerase  $\beta$  gene in 8-week-old mice. Deletion was nearly complete in the liver and nearly complete in lymphocytes a few days after injections.

#### **Critical for centromeres**

In the fission yeast Schizosaccharomyces pombe, centromeres, telomeres, and silent mating-type loci are transcriptionally silent. The swi6+ gene encodes a chromodomain protein that represses transcription at these structures which make up heterochromatin in this yeast and that is required for proper chromosome segregation during cell division. Ekwall et al. (p. 1429) have now localized the Swi6 protein to centromeres, telomeres, and silent mating loci. They also show that mutations in swi6 cause the centromeres to lag on the spindle at anaphase, which results in chromosome loss.

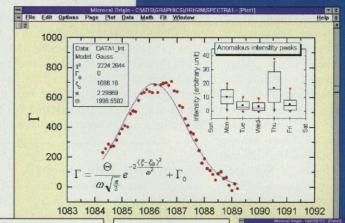


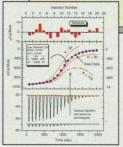
Fall 1995.0

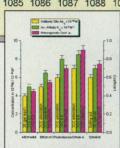
## Technical Graphics Data Analysis

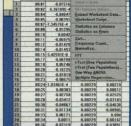
Origin was the first technical graphics and data analysis software developed for the MS-Windows™ environment. Today, Origin is recognized as a powerful, fast, and easy to use software that is ideally suited to meet the needs of scientists and engineers. Discover why thousands agree that Origin is the ultimate solution!

- Input data from keyboard, use functions or import from a wide range of data formats.
- Customize 100+ graph types for text font, line style, symbol, color, axis type, tick labels and more; Save and import templates.
- Analyze data with a wide selection of statistical and mathematical functions, including non-linear curve fitting.
- Automate and batch process tasks.
- Create publication quality graphics with Windows supported printers, plotters, and slide makers.

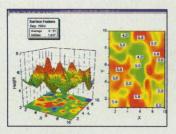




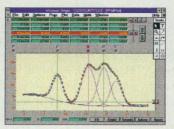




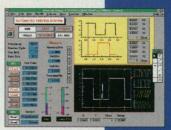
## and the rest of the Family



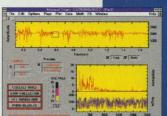
3D and Contour Module Generate 3D surface, scatter, and contour plots. Mix your XYZ plots and mesh surfaces with contour map and labels. Quickly rotate the viewing angle, and more!



Peak Fitting Module
Provides peak analysis for
chromatography and
spectroscopy. Fits up to 100
peaks, and generates
detailed reports in Origin's
superb graphics.



Data Acquisition System
A family of modules provides complete data acquisition support with quick, powerful, functional building blocks for plug-in boards, IEEE-488, and RS-232 devices.



Other Add-On Modules

- User Interface Module
- File Utilities Module
- Real Time Module
- Axon pClamp Module

Microcal

Microcal Software Inc. One Roundhouse Plaza Northampton, MA 01060 USA Fax (413) 585-0126 email address: info@microcal.com Call for Free Demo 1-800-969-7720



#### INTRODUCING

Change... Growth... Adaptation... Our vision of serving the future needs of life sciences means transforming the way we do business.







combines the strengths of each division to create a balance of skills, disciplines and customer support that

will set the standard into the next century.

Today, Sigma BioSciences stands as your single source for life science research and manufacturing products.

Our vision has built Sigma BioSciences - a union of our Cell Culture, Immunochemical and Molecular Biology divisions. Sigma BioSciences



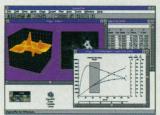
We are committed to the success of our customers through science, technology and service.

Count it! Measure it! Stat it! Fit it! Plot it! We've got it!



#### Introducing the latest generation of the Jandel software family

Jandel Scientific Software has long been the first name in integrated software solutions for scientists and engineers, with a full range of products that complement each other. Our powerful SigmaLink™ technology lets you share the same worksheet and graph page in SigmaPlot®, SigmaStat®, and SigmaScar™. Jandel Scientific – the leading family of scientific and engineering software since 1982.



#### SigmaPlot FLEXIBLE SCIENTIFIC GRAPHING

From nonlinear regressions to multiple axes per graph, more than 75,000 researchers depend on award-winning SigmaPlot's flexibility and power to analyze and graph their technical data the way they need to.

For Mac 70 For Win 71



#### SigmaGel GEL ANALYSIS SOFTWARE

Gel analysis-spot, lane, molecular weight, and DNA-on your lab or office PC.Works with inexpensive hand or desktop TWAIN-compatible scanners and imaging devices. Quality and speed previously available only in higher priced systems.

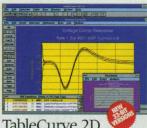
Circle No. 75



### SigmaStat ADVISORY STATISTICAL SOFTWARE

Advises on the test to use, checks for violations in underlying assumptions, runs the test, and explains the results. Handles messy data. Like having a professional statistical consultant at your fingertips.

Circle No. 72



## TableCurve 2D AUTOMATED CURVE FITTING TableCurve 3D AUTOMATED SURFACE FITTING

Equation generator and curve/surface fitter. Use built-in equation generator to select and rank best-fit equations or fit theoretical models with user-defined functions.

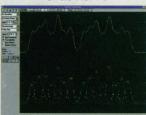
Circle No. 76



## SigmaScan IMAGE MEASUREMENT SigmaScan Pro IMAGE ANALYSIS

Easy, accurate and low-cost. SigmaScan for manual image measurements or SigmaScan Pro for automated measurements, counting, and image analysis.

Circle No. 73



#### PeakFit automated nonlinear peak analysis

Spectroscopists, chromatographers, and other researchers use PeakFit to separate peaks automatically. PeakFit contains the most complete set of nonlinear fitting functions available.

Circle No. 77



Windows® 95 Compatible

## SigmaSuite INTEGRATED GRAPHING, STATISTICS & IMAGE MEASUREMENT

The power of SigmaPlot, SigmaStat, and SigmaScan at a Suite price! Measure images, run statistics, and create graphs using SigmaLink for seamless data sharing in all three products.

Circle No. 74



#### SigmaPlot Graph Library VOLUME ONE

Create graphs quickly and easily by choosing from a visual catalog of 150 professionally designed graphs.

Circle No. 78

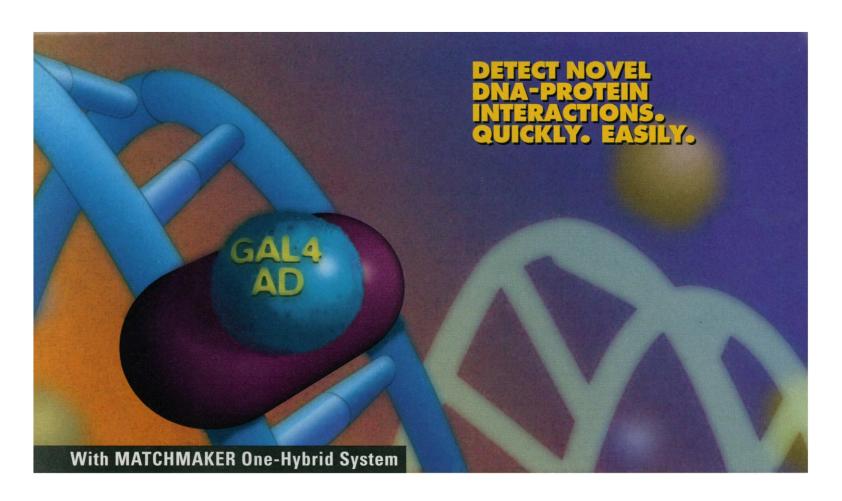
#### TO GET IT CALL 1-800-4-JANDEL (1-800-452-6335)

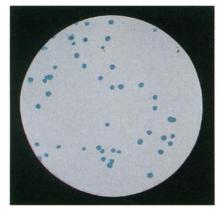
FOR MORE INFORMATION PLEASE CALL, FAX OR E-MAIL: Jandel Scientific: 2591 Kerner Blvd., San Rafael, CA 94901

Jandel Scientific: 2591 Kerner Blvd., San Rafael, CA 94901 415-453-6700 Fax 415-453-7769 Internet: sales@jandel.com In Europe: Jandel Scientific, Schimmelbuschstrasse 25, 40699 Erkrath, Germany 02104/9540 02104/95410 (FAX).

International Dealers: Australia 2 958 2688, Brazil 11 453 5588, Canada 519 767 1061, Chile 56 2 635 2422, Hong Kong 852 739 2001, India 40 31 3368, Israel 972 349 0823, Japan 3 3590 2311, 3 5689 8000, Korea 822 886 3983, Kuwait 965 242 1851, Malaysia 603 777 2597, Mexico 525 513 0399 South Africa, 12 660 2884, Taiwan 2 788 6777, 2 705 1590 All company and/or product names are trademarks of their respective companies. @1995 Jandel Corporation.







Results of a  $\beta$ -galactosidase filter assay on a positive clone resulting from a MATCHMAKER one-hybrid library screening. Plasmids isolated from the 12 large colonies that grew on SD/–His agar plates were transformed into the p53-lacZ reporter strain, plated on SD/–His/–Leu/+45 mM 3-AT, and incubated at 30°C for 4 days. A colony-lift  $\beta$ -galactosidase filter assay was performed according to the MATCHMAKER One-Hybrid protocol.

| Product                         | Size | Cat. #  |
|---------------------------------|------|---------|
| MATCHMAKER<br>One-Hybrid System | each | K1603-1 |

CLONTECH's MATCHMAKER One-Hybrid System is a new *in vivo* yeast genetic assay for studying DNA-protein interactions. The one-hybrid system allows you to rapidly identify novel DNA-binding proteins from cDNA or genomic libraries and immediately access the genes for the identified proteins. Maximal sensitivity is achieved because DNA-binding protein interactions are detected while proteins are in their native configurations. Use this sensitive assay to map DNA-binding domains of known transcription factors. All pre-made MATCHMAKER cDNA and genomic libraries are compatible with the new one-hybrid system. Only from CLONTECH. For more information or to place an order call 1-800-662-CLON or contact your local distributor.

ON'T EXPERIMENT WITH ANYTHING ELSE.



1020 East Meadow Circle, Palo Alto, California 94303 USA
FAX: 800/424-1350 415/424-1064 • TEL: 800/662-CLON 415/424-8222
e-mail: tech@CLONTECH.com orders@CLONTECH.com • internet:http://www.clontech.com
@1995 CLONTECH Laboratories, Inc.

## **Novel Gene DNA Sequencing**

# with the genomyxLR™

## Advanced Performance at an affordable price



#### Advanced "air impingement" temperature control

Proprietary air distribution technology maintains a constant and completely uniform temperature over a 62 x 33 cm gel during electrophoresis, eliminating smiling and other electrophoretic distortions. Brilliant band resolution and long read length assure maximum data per gel (Panel A).

#### Unsurpassed resolving power and accuracy

Programmable, microprocessor control and decoupling of voltage and temperature provide high resolution of data from different templates. Elevated temperature runs eliminate problems and data loss associated with sequencing GCrich templates (Panel B). Achieve 100% accuracy at up to 650 bases with new 200 µm spacer set.

Outstanding performance in the following applications...

#### Adjunct to automated sequencing Prevents data loss and assures full-length reads by supplementing automated systems when sequencing difficult templates,

prescreening clones or closing gaps.

#### The tool of choice for Molecular Systematics

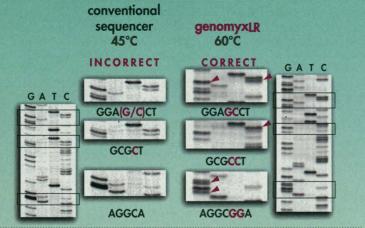
Achieve dramatically improved throughput and maximum data per gel for rapid completion of phylogenetic projects.

Long-read autorads are ideal for archival data storage.

#### Upgrade of conventional manual sequencing

A technically advanced, yet affordable standardized method for minimizing effort and maximizing throughput in dedicated manual sequencing labs.

### Eliminate compressions and data loss...



45°C sample run on conventional manual sequencing equipment under ambient conditions (6 hr, 60 W, 6% acrylamide, aluminum plate backing). 60°C sample run on **genomyx<sub>LR</sub>™ DNA Sequencer** (5 hr, 100 W, 6% HR-1000<sup>™</sup> gel). Sequencing data derived from mouse cardiotrophin cDNA (Genentech, Inc.)

#### Call today to discuss how the genomyxLR can help you with your sequencing project.

The **genomyxLR** is a programmable DNA sequencer incorporating proprietary **air-impingement technology**. This system includes a built-in power supply and all the equipment required to immediately begin running long-read gels. A complete line of **LR-OPTIMIZED**<sup>TM</sup> kits and reagents is available to ensure optimal performance with the **genomyxLR** System.



For additional information regarding the **genomyxLR** Sequencing System and our complete line of **LR-OPTIMIZED** products, please call us at **1-800-9GENOME**. **genomyx@aol.com** 

### **Differential Display**

# with the genomyxLR™

#### For a level of performance unachievable by conventional electrophoresis



#### Resolution

Dramatically enhanced resolution of all cDNA fragments across a broad size spectrum assures complete spatial separation of all bands for accurate pattern determination and band recovery (Panel C).

#### Read length

The only electrophoresis system capable of resolving cDNA fragments as large as 2-3 kb. Assures maximum coding information for efficient gene identification.

#### Registration

Completely even cDNA fragment migration across entire gel assures accurate pattern determination over the entire width of gel.

#### Recovery

Precise excision of single bands from gel is facilitated by high resolution electrophoresis, allowing accurate, quantitative recovery of unique cDNA fragments for reamplification and sequencing with the **genomyxlr**.

Panel A: Long read, high resolution Differential Display data provided by Drs. Lidia Averboukh and Arthur Pardee, Division of Cell Growth & Regulation, Dana Farber Cancer Institute. Samples run on genomyxLR (2.5 hr, 100 W, 3 kV, 50°C, HR-1000™ 4.5% Denaturing gel). Data represent display patterns under different RT-PCR conditions using total cellular RNA from LNCaP-FGC prostate carcinoma cell line. All reactions performed with the GenHunter RNAimage™ Differential Display Kit.

Panel B: Long read, high resolution Differential Display patterns of total cellular RNA from HeLa and transformed rat fibroblast cells using GenHunter RNAimage Differential Display Kit. Samples run on **genomyxLR** (2.5 hr, 100 W, 2.7 kV, 50°C, HR-1000 6% Denaturing gel). Data provided by GENOMYX Corporation.

#### Call today to discuss your Differential Display requirements with one of our scientists.

The **genomyxlr** is a programmable DNA sequencer incorporating proprietary **air-impingement technology**. This system includes a built-in power supply and all the equipment required to immediately begin running long read gels. A complete line of **LR-OPTIMIZED**<sup>TM</sup> kits and reagents is available to ensure optimal performance with the **genomyxlr** System.



For additional information regarding the **genomyxLR** DNA Sequencer and our complete line of LR-OPTIMIZED products, please call us at 1-800-9GENOME.

# Is combinatorial chemistry just multiplying your compounds, or compounding your problems?

# Project Library can help manage and make sense of your data.

Project Library helps you quickly store and access the structural and biological data on the thousands—even millions—of molecules you create with combinatorial chemistry. Affordable and easy to use, Project Library is a desktop software application uniquely equipped to help you perform critical tasks, such as:

- Complete or selective enumeration of libraries into sub-generics or specifics, by criteria such as molecular weight, Rgroup, Rgroup tag or identifier, and substructure.
- Elucidation of active structures from deconvolution, spatially addressable, or encoding experiments.
- Structure and data exchange with other scientific applications for analysis and reporting.

The true value of combinatorial chemistry, after all, lies in how well and how quickly you interpret the data. And Project Library offers you an efficient and cost-effective way to do just that. For more information, please call today:

United States: 1-800-225-9423 Europe: 41-61-4812180 United Kingdom: 01-276-681777 Japan: 81-6-302-9895

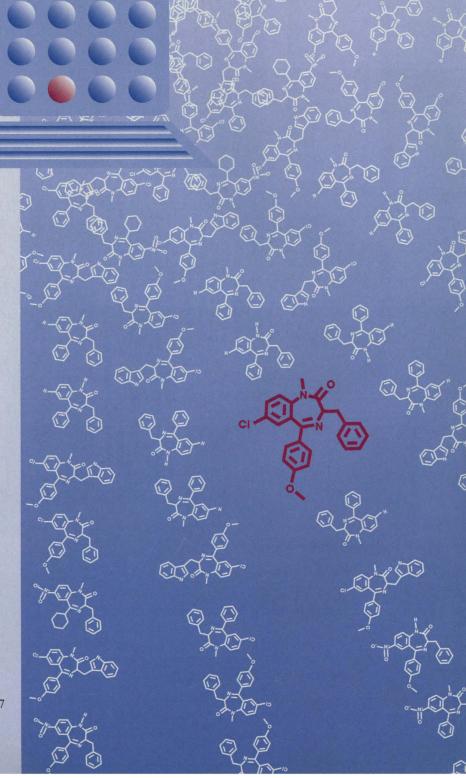
#### MDL

Information Systems, Inc.

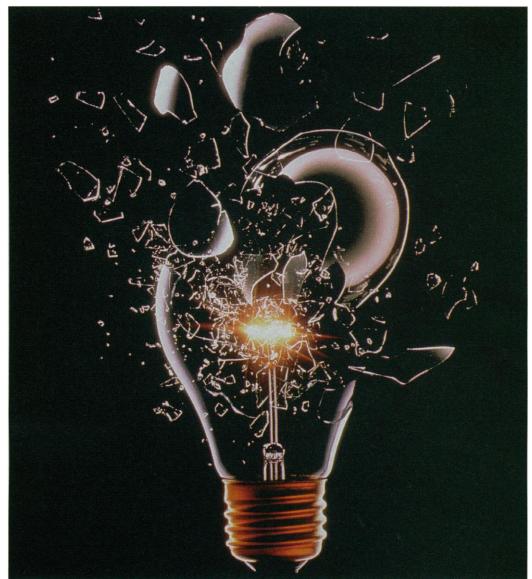
14600 Catalina Street, San Leandro, CA 94577 Fax: 510-483-4738

MDL and Project Library are trademarks of MDL Information Systems, Inc.

Circle No. 55 on Readers' Service Card



### **Break the Information Barrier**



#### Work smarter. Work faster. Work more efficiently.

nce you go beyond the confines of conventional information retrieval, amazing things happen. Suddenly, you can find your way through scientific and technical data with unbelievable agility. The journal articles, conference papers, and patent information you need appear before your eyes. And industry news from all over the world is as close as your computer.

You save time. You save money. And best of all, it's easy. With Knight-Ridder Information's advanced product line, everything you need is right at your fingertips.

- Flexible searching online, CD-ROM, and the Internet
- Convenient document delivery
- Unmatched range of scientific and technical data
- Value-based pricing
- Over 20 years of online expertise

So don't let the information barrier hold you back. Break through it with Knight-Ridder Information's tools. And watch your research take off.

Call us at 800-334-2564, 415-254-8800, 215-241-0131, or http://www.dialog.com

#### Featuring DIALOG®

Also:

DataStar®

 ${\rm KR} \; {\rm ProBase^{\rm SM}}$ 

DIALOG DirectsM

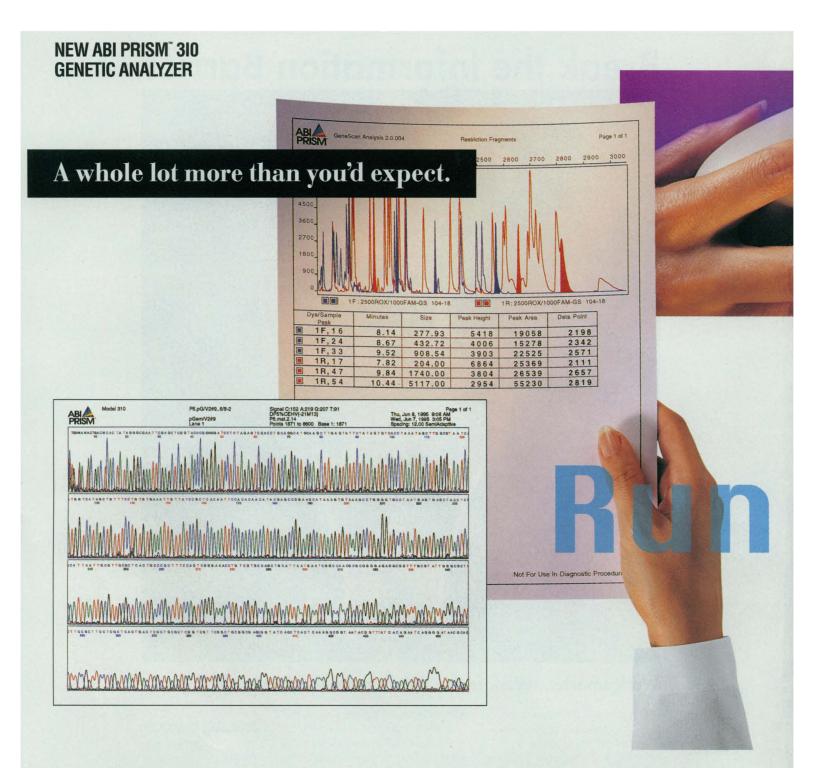
KR SourceOne<sup>SM</sup>

CustomDIALOG™

KR ScienceBasesM

KR Information OnDisc™

KNIGHT-RIDDER INFORMATION





The new ABI PRISM" 310 Genetic Analyzer is changing a lot of expecta-

tions about automated genetic analysis.

To begin with, it gives you a lot more versatility. The 310 system automates all the applications you now perform on acrylamide or agarose—and a whole lot more—without limitations.

At the same time, the 310 Genetic Analyzer makes genetic analysis a lot less work. You'll never have to pour another slab gel. Instead of conventional gels, the 310 system uses revolutionary new polymers that are injected into a capillary before each run. Not only do these polymers make automation possi-

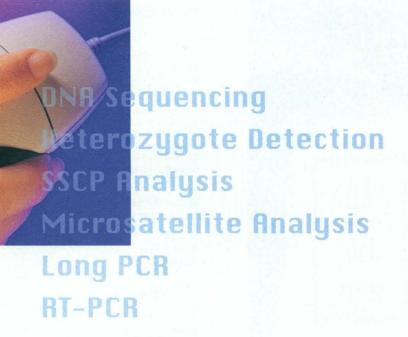
ble, their uniform consistency eliminates guesswork and helps to ensure reproducibility.

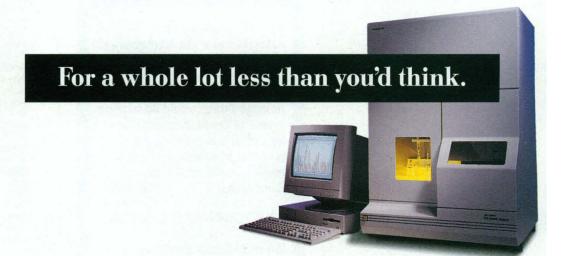
You don't even have to pipet your samples into the gel. Just perform your

#### ABI PRISM 310 GENETIC ANALYZER

| Application         | Performance                            |
|---------------------|--|
| Cycle Sequencing    | 425 nt, 98.5% base calling accuracy    |
| Microsatellites     | 2 nt resolution, 0.5% sizing precision |
| SSCP                | 0.5% pattern matching precision        |
| Quantitative RT-PCR | 4.0% peak height precision             |

A lot more, for a lot less. You can automate virtually any DNA electrophoresis application on the ABI PRISM 310 system. Here are just a few examples.





reactions, and place your labeled samples in the sample tray. The 310 system automatically injects your samples in whatever order you specify. You can even add samples during a run.

Sample preparation is a lot less trouble, too. ABI PRISM kits for fluorescent PCR and DNA sequencing reduce pipetting steps and variability. And there's no radioactivity or ethidium bromide.

Most importantly, the 310 Genetic Analyzer makes it a whole lot easier to get the results you want. In fact, the results are so accurate, you can transfer data directly to specialized programs for genotyping, mutation detection, and sequence assembly.

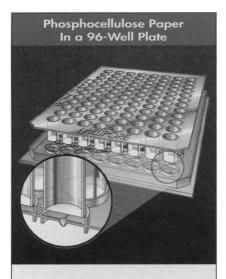
But if you expect the 310 Genetic Analyzer to cost a lot, you'll be surprised to find the price is well within the budget of individual laboratories. The ABI PRISM 310 system costs about half as much as the ABI PRISM 377 system, our highthroughput DNA sequencing system.

Circle No. 37 on Readers' Service Card Electronic Marketplace: http://www.aaas.org

The ABI PRISM 310 Genetic Analyzer. A whole lot more, for a whole lot less. To request information, call 1-800-345-5224. Outside the U.S. and Canada, contact your local Perkin-Elmer representative.

Europe Langen, Germany Tel: 49 6103 708 301 Fax: 49 6103 708 310 Japan Tokyo, Japan Tel: (0473) 80-8500 Fax: (0473) 80-8505 Latin America Mexico City, Mexico Tel: 52-5-651-7077 Fax: 52-5-593-6223 Australia Melbourne, Australia Tel: (03) 9212-8585 Fax: (03) 9212-8502

The Polymerase Chain Reaction (PCR) process is covered by U.S. patents owned by Hoffmann-La Roche, Inc., and F.Hoffmann-La Roche Ltd. Perkin-Elmer is a registered trademark and ABI PRISM is a trademark of The Perkin-Elmer Corporation.



### Increase Throughput For Kinase Assays

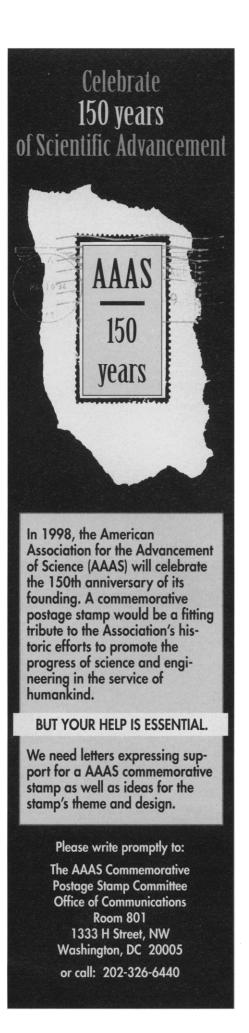
Signal transduction assays are an important tool in drug discovery. The Millipore MultiScreen Assay System uniquely offers a 96-well filter plate containing phosphocellulose paper and protocols to perform peptide substrate kinase assays. No more cutting and transferring the paper! Every step, from the phosphorylation reaction to direct scintillation counting, can be carried out in one plate.

The MultiScreen Assay System can also be used with other filter formats, including glass fiber or low protein binding Durapore® (PVDF) membrane both ideally suited for TCA precipitated substrates.

Call or fax for more information. U.S. and Canada, call Technical Services: 1-800-MILLIPORE (645-5476); in Japan, call: (03) 3474-9111; in Europe, fax: +33.88.38.91.95.

#### MILLIPORE

MILLIPORE LAB CATALOG ON INTERNET: ACCESS URL MENU AND TYPE: http://www.millipore.com



#### **Boehringer Mannheim** Worldwide Representatives

**Australia** Tel.: (02) 899 7999

Austria

Tel.: (0222) 277 87 DW 420 to 423

**Belgium** Tel.: (02) 247 4930

**Brazil** Tel.: 55 11 66 3565

**Canada** Tel.: (514) 686 7050

Central/Eastern Europe Tel.: (0222) 277 87

**China** Tel.: (852) 2485 7596

**Czech Republic** Tel.: (0324) 45 54, 58 71-2

**Denmark** Tel.: 49 13 83 42

Finland Tel.: (90) 429 2342

France Tel.: 76 76 30 86

**Germany** Tel.: 0621 759 8568

**Greece** Tel.: (01) 64 96 683-6

Hong Kong, see China India

Tel.: (22) 837 0794

Indonesia, see Singapore Ireland Tel.: 1 800 409 041

**Israel** Tel.: (03) 493111

**Italy** Tel.: (02) 270 96209

**Japan** Tel.: (03) 3432 3155

Malaysia, see Singapore Mexico Tel.: (5) 227 8967

Netherlands Tel.: (036) 539 4911

New Zealand Tel.: (09) 276 4157

Norway Tel.: 22 07 65 00

Philippines, see Singapore

**Poland** Tel./Fax: (022) 37 42 35 Tel./Fax: (026) 62 63 03

Portugal Tel.: (01) 417 1717

**Singapore** Tel.: 65 479 4111

South Africa Tel: (011) 886 2400

South Korea Tel.: 02 569 6902 (5 lines) 02 562 9947

**Spain** Tel.: (93) 201 4411

**Sweden** Tel.: 08 98 8150

Switzerland Tel.: 042 65 4242

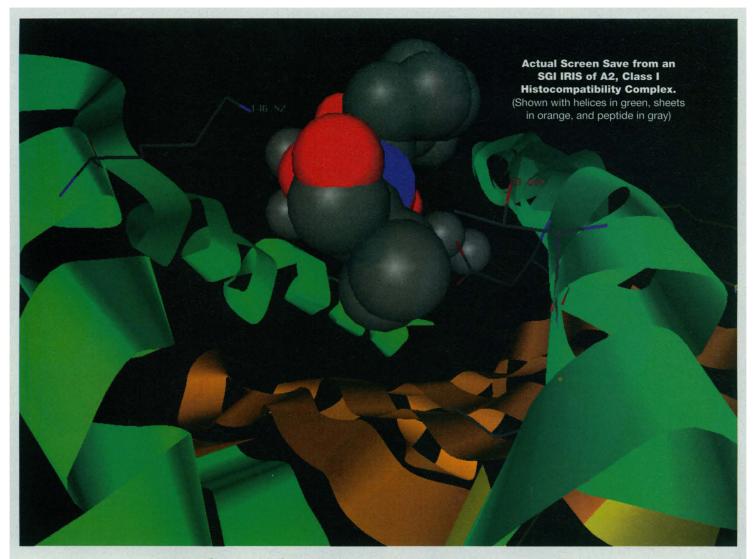
**Taiwan** Tel.: (02) 736 7125

Thailand, see Singapore

**Turkey** Tel.: 1 349 8176, -77, -78, -79

United Kingdom Tel.: 0800 521578

USA Tel.: 800 428 5433



## Bridging the Sequence-Structure Gap

INNOVATIVE UNDERSTANDING OF PROTEINS

The explosion of the field of X-ray crystallography has resulted in the amazing capacity to view the position and interrelation of every atom in a

Visualize your protein, directly from a known crystallized structure, or by homology modeling (using the SegMod accessory package)

large array of proteins. How can you use this structural information to inform and inspire your

experiments, to understand how proteins function?

*Introducing LOOK*: A unique software tool specifically designed to bring structural insight to molecular biology. LOOK integrates the study of

sequence with structure and function, through powerful representation and prediction.

Visualize and analyze your experimental results utilizing the structure of your protein. Design your future experi-

SeyWith

A lice was a lamining and a superior instance product and instance and ins

ments based on structural knowledge.

LOOK enables you to communicate those results with persuasive, insightful graphics ready for publications or slide presentations.

Additionally, the LOOK program includes a database of structures and

sequences
customized for your
interests. LOOK also
gives you a database of
all literature for your
system, that lets you see
what is known about any

given residue or region in your protein, simply by clicking on it. These databases

give you the information you need for your research – all

in one place.
For additional

information about LOOK for your Silicon

SLGYLRSYYNQSEAGH
RIALR
RIALR
Butter of the control of the control

Graphics®, DEC Alpha® or SUN® workstation call toll free 1-800-229-7382 (USA only), 415-473-3030 (worldwide), or email info@mag.com today.

LOOK
THE MOLECULAR
APPLICATIONS GROUP

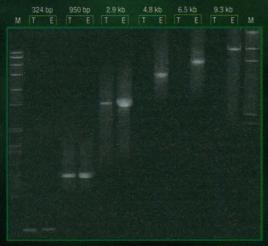
Title: Residues in pockets B and P of HLA-B27 are critical in the presentation of an influenza A virus nucleoprotein peptide and influence the stability of peptide – MHC complexes.

Author: Carreno BM: Whater CC; Taurog JD; Hansen TH; Biddison WE.

Expand Beyond the Limits of *Taq* DNA Polymerase

#### Better yields and fidelity than Tag

Achieve 2-fold higher yields and up to 3-fold greater fidelity with the Expand<sup>™</sup> PCR System from Boehringer Mannheim. The result is stronger bands than *Taq* DNA polymerase\* can offer. From even smaller amounts of short and long templates. With fewer misincorporations.



E= amplification with Expand High Fidelity; T= amplification with Taq; M= Molecular Weight Marker.

#### Perform PCR not possible with Tag

Two Expand blends combine the high processivity of our *Taq* DNA polymerase\* with the proofreading accuracy of our exclusive *Pwo* DNA polymerase\*, delivering better results than *Taq* DNA polymerase.

#### Expand High Fidelity

- increases yields and fidelity for improved PCR cloning experiments—even from difficult templates
- exhibits a greater tolerance for [Mg<sup>2+</sup>] variations, ensuring consistent PCR results.

#### Expand Long Template

- amplifies human genomic DNA of up to 27 kb
- amplifies λDNA of up to 40 kb.

#### Don't let Tag limit your research possibilities

Expand places premium performance within striking distance. Order today to begin to see striking results.

Expand™ is a trademark of Boehringer Mannheim.

\* Purchase of these products is accompanied by a limited license to use them in the Polymerase Chain Reaction (PCR) process for life science research in conjunction with a thermal cycler whose use in the automated performance of the PCR process is covered by the up-front license fee, either by payment to Perkin-Elmer or as purchased, i.e., an authorized thermal cycler.

© 1995 Boehringer Mannheim. All Rights Reserved.

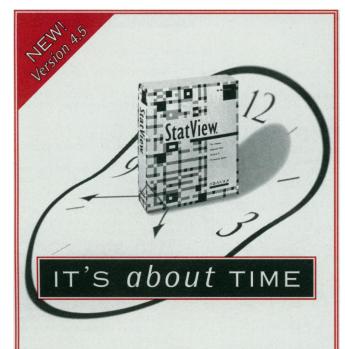
## EXPAND"

PREMIUM PCR PERFORMANCE

Striking Results — In the wild, a thin line separates predator and prey. When the falcon crosses that line, it does so with lightning speed and accuracy. Striking with a precision that delivers uncompromising results.



Biochemicals



Saving time – your time.

Since 1985, StatView's been the best selling data analysis software on the Macintosh. Why? Because StatView's easy. Easy to learn. Easy to use.

In less time than it takes to learn most data analysis programs, you get your results with StatView.

StatView is designed to work with you. The software's flexible and intuitive interface allows you to analyze data the way you want to.

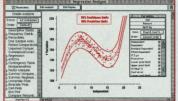
StatView features fully integrated statistics, data mangement, graphing and presentation tools – all the tools you need to get your work done.

And with StatView, your time is never wasted.

StatView's innovative templates allow you to save all your work simply by saving the document. You can open that template at any time to resume your analysis. Or apply it to any other data set.

Whether you need to analyze data on a daily basis or only every couple of months, StatView's unprecedented ease of use will save you time and effort.

Isn't it about time you used StatView?



Creating an analysis is as easy as clicking your mouse.

FOR MORE INFORMATION CALL
1 - 8 0 0 - 6 6 6 - S T A T



THE SIGNIFICANT DIFFERENCE"

1918 Bonita Avenue, Berkeley, CA 94704-1014 Phone: 510-540-1949, Fax: 510-540-0260 Internet: info@abacus.com



We have localized, international and student versions of StatView and license, academic and quantity discounts. Call for more information.

Circle No. 34 on Readers' Service Card Electronic Marketplace: http://www.aaas.org

# SCIENCE ENTERS CYBERSPACE!

Announcing the SCIENCE Electronic Marketplace: A new on-line product information service for readers.

To access the SCIENCE Electronic Marketplace go to:

World Wide Web: http://www.aaas.org Gopher: gopher.aaas.org

Choose SCIENCE from the menu, then select SCIENCE Electronic Marketplace

Participating advertisers have the words: **Electronic Market Place: http://www.aaas.org** printed beneath their advertisement. Look for them in this issue.

**Abacus** 

Academic Press, Inc.

**Advanced ChemTech** 

Advanced Gene Computing Technologies, Inc Apple Computer, Inc.

Ambion, Inc.

**Boehringer Mannheim Biochemicals** 

**Bio Image** 

**Bio Soft** 

**BTX** 

**Cell Systems** 

Genomyx

**Jandel** 

Jolley Consulting and Research, Inc.

**Molecular Applications Group** 

Leica

NCCS

Niles and Associates, Inc.

**Perkin Elmer** 

Personal Bibliographic Software, Inc.

**Pharmacia Biotech** 

Progen

**Research Software Design** 

Stratagene

**Scientific Endeavors** 

**Sigma Chemical Company** 

**Southwest Scientific Resources** 



A few great reasons why you should know more about ISIS.

t's no surprise that leading companies like these count on ISIS to help them discover and deliver new products. You see, ISIS, the Integrated Scientific Information System from MDL, has been setting the industry-standard for years.

What can ISIS do for *your* company? Our unique software systems can give everyone in your organization the power to quickly integrate, analyze, manage, and communicate *all kinds* of important chemical and biological information. And they can do it whether they're across the hall or across the ocean.

That's right. Chemical structures, property data, 3D models, synthetic reactions, and biological data from relational databases, can all be pulled together instantly. Decision making becomes easier because you have all the right information exactly when you need it.

The bottom line? The computers on your researchers' desktops become part of an information management strategy specifically designed to increase productivity and put more candidates in the discovery pipeline.

Best of all, ISIS comes from a company you can rely on—MDL Information Systems, Inc. We're proud of the fact that for almost two decades, we've been helping organizations of all sizes get the most from their intellectual assets—with software, databases, services, education, support, and more.

If you've got questions about what our time-tested software can do for *you*, find out what over 16,000 scientists in more than 200 companies have already discovered—that when it comes to information management, the answer is ISIS.



## FREE! Important Information

Call now and we'll send you important information with no obligation. Fact-filled case studies document how ISIS has helped companies like yours meet their business objectives. Industry leaders already rely on ISIS. Shouldn't you find out why?

#### Call 800-635-0064 x1326

Information Request Number: PK-ISIS-2



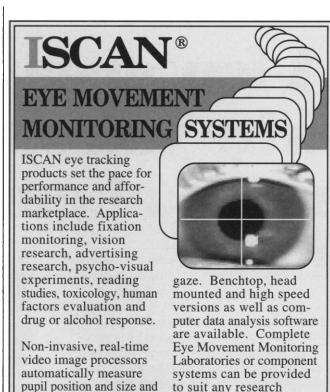
Information Systems, Inc.

14600 Catalina Street, San Leandro, CA 94577 (510) 895-1313 Fax: (510) 483-4738 Internet: isis@mdli.com World Wide Web: http://www.mdli.com

©1995 by MDL Information Systems, Inc. ISIS and MDL are trademarks of MDL Information Systems, Inc.



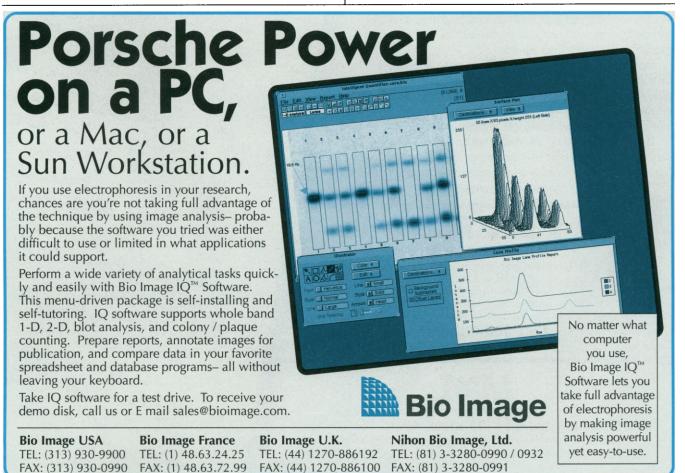
Circle No. 5 on Readers' Service Card



to suit any research compute subject's point of application. ISCAN, Inc. P.O. Box 382076

Cambridge, MA 02238-2076 USA Tel: 617-273-4455 Fax: 617-273-0076

Circle No. 11 on Readers' Service Card



# Hnnouncing...

## NO ANNUAL FEE and a Special 9.9%<sup>†</sup> Annual Percentage Rate (APR) on Cash Advances and **Balance Transfers for AAAS Members.**



**Call Now!** 1-800-847-7378, ext. 400

Please use this priority code when calling: NTMG

tThe Annual Percentage Rate (APR) for purchases is 16.4%, which may vary. The APR is 9.9% on cash advances through your first six statement closing dates commencing on the month after your account is opened and may vary thereafter (applies to both new and outstanding cash advance balances). The current indexed rate for cash advances is 16.4%, which may vary. The transaction fee for the purchase of wire transfers, noney for advances, to the property tickets, and casino gaming chips is 2% of each such purchase; \$2 minimum, \$25 maximum. Transaction fee for two specifically designated Premium Access Checks\* or Preferred Access Checks\* will be waived. The information about the cost of the card described in this application is accurate as of 1/95. The information may have changed after that date. To find out what may have changed, call 1.90.943.7378

\*\*Gash advances cannot be used to pay off or pay down another MBNA\* account.

MBNA America,\*\* Premium Access Checks and Preferred Access Checks are federally registered service marks of MBNA America Bank, N.A. MasterCard\* is a federally registered service mark of MasterCard International, Inc., used pursuant to license. Visa\* is a federally registered service mark of Visa U.S.A., Inc., used pursuant to license. © 1994 MBNA America Bank, N.A.

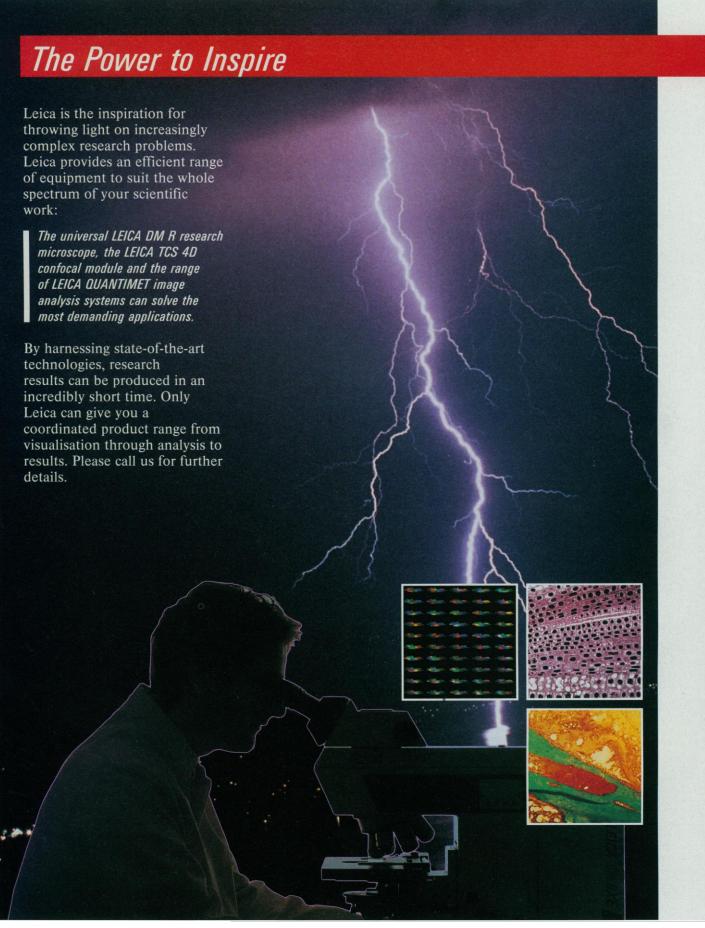
Introducing the AAAS Visa® Gold card, with NO ANNUAL FEE. Many AAAS members have already discovered the benefits of the AAAS Visa Gold card. Now, through this special offer, you can join your colleagues in carrying the only credit card endorsed by the AAAS.

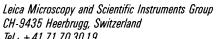
The American Association for the Advancement of Science is pleased to present a special introductory 9.9% APR on cash advances.\* As a part of your welcome package you will receive two Premium Access Checks that you can use to pay off or pay down other credit cards and installment loans.

Extra benefits include:

- A 16.4% variable APR on purchases
- Higher lines of credit ... up to \$50,000
- 30 minute credit line increases

Now you can carry the AAAS card with these and many other unsurpassed benefits and pay no annual fee.





Tel.: +41 71 70 30 19 Fax: +41 71 72 78 25

M&SI 1/94 e

Circle No. 17 on Readers' Service Card Electronic Marketplace: http://www.aaas.org



#### PERSONAL BIBLIOGRAPHIC DATABASES...

These cost more:

ENDNOTE • DMS 4 CITE • PRO-CITE REFERENCE MANAGER • REF-11

This does more:

# PAPYRUS

- Manages up to 2 million reference citations. Stores up to 16,000 characters and 100 keywords per reference.
- Dozens of predefined output formats, plus the ability to easily design your own.
- 100% compatible with WordPerfect\*, Microsoft Word\*, Ami Pro\*, WordStar, XyWrite, Signature, ChiWriter, TEX.
   \*including Windows™ versions
- Can also be used with virtually all other word processors.
- · Fast, powerful search capabilities.
- Able to import references from national databases, CD-ROM files, monthly diskette services, other bibliography programs, or almost any other database or text file.
- Allows an unlimited number of Notecards for each reference.
- · Powerful new user interface.
- Fully compatible with Windows.™

for IBM-PC and compatibles
also available for VAX-VMS

#### Research

SOFTWARE DESIGN

2718 SW Kelly Street, Suite 181 Portland, OR 97201 (503) 796-1368 FAX: 503-241-4260 Complete System \$99

Full money-back guarantee on purchase of Complete System.

#### Demo System \$25

Demo price credited toward subsequent Complete System purchase.

Outside North America, add \$20 shipping charge – U.S. funds, on a U.S. bank.

Circle No. 49 on Readers' Service Card Electronic Marketplace: http://www.aaas.org

#### Our Users Are Redefining the Readlengths You Can Achieve with DNA Sequencers!

# 1080 BASES\*

Congratulations to Dr. Hilary Morrison for winning the first Long ReadIR™ Championship Challenge. Dr. Morrison submitted a double-stranded DNA sample that *autosequenced\** to 1080 bases with 98.5% accuracy. To find out how LI-COR **infrared** DNA sequencers can increase the productivity of your lab, call for information today!

1-800-645-4267

Toll Free U.S. & Canada

LI-COR, Inc. 4421 Superior Street Lincoln, NE 68504 USA LI-COR®

Phone: 402-467-0700 FAX: 402-467-0819

\*Dr. Morrison is with Mitchell Sogin's Molecular Evolution Group at the Marine Biological Laboratory, Woods Hole, MA. Readlengths for the contest were determined using L1-COR's Base ImagIR<sup>TM</sup> software to automatically call bases with no editing or operator involvement.

# Custom Peptides for the Incurably Suspicious



If you're the kind of researcher who questions everything that could affect your results, Genosys is your kind of peptide supplier.

We encourage healthy skepticism, but you'll find there's never a doubt about a Genosys peptide. That's because every one of them is 100% satisfaction guaranteed. Every one is quality-certified by mass spectral analysis and HPLC. We even take extra steps to make sure your peptide will be as soluble (and easy to work with) as possible.

Choose the purity level and quantity that best suits your research, and your budget. Pick your modification: MAPS, biotin, N-acetylation, C-amidation, phosphotyrosine, D-amino acids, or KLH / BSA conjugation. No-hassle antisera services are also available.

But don't take our word for it (as if you would.) Call today to discuss your requirements. Then place an order, and validate our claims by the scientific method.

# GENOSYS

North America The Woodlands, TX U.S.A.

Phone: (800) 853-3385 Fax: (713) 363-2212 eMail: genosys@main.com Europe Cambridge, UK Phone: (+44) (0) 1223 425622

Fax: (+44) (0) 1223 425966 eMail: genosys@genosys.co.uk

VISA

Visit Our World Wide Web Site at http://www.genosys.com

- Q---

@ 1995 Génosys Bicsechnológies, Inc. 195-250-B&S

Circle No. 35 on Readers' Service Card

## Building Your VIRTUAL Library



#### New Electronic References from Academic

Base price allows for up to five users!



Recombinant DNA and Related Methodology

18 Selected Core Volumes with Cumulative Indexes for MIE Volumes 1-244

Includes Recombinant DNA, Parts A-I, DNA Structure, Parts A & B, Three Guide to's, and Related Volumes

John N. Abelson and Melvin I. Simon

Praise for the Print Version

"Incomparably useful."

-ANALYTICAL BIOCHEMISTRY

In 1995, the critically acclaimed Methods in Enzymology series celebrates its fortieth birthday. In honor of this occasion, Academic Press is proud to present Methods in Enzymology on CD-ROM, CD-1, focusing on recombinant DNA and related methodology. A perfect complement to the print series, **Methods in Enzymology on CD-ROM, CD-1**, combines indexing with high-quality, informative full texts to create a unique information resource that brings single-search access to two database collections. The CD-ROM runs on Windows, Macintosh, and UNIX machines.

#### CONTENTS

Includes these full-text volumes:

METHODS IN ENZYMOLOGY, VOLUMES 68, 100, 101, 152, 153, 154, 155, 182, 183, 185, 194, 208, 211, 212, 216, 217, 218, and 224

Includes these index volumes:

Cumulative Contributor Index and Table of Contents for Volumes 1-244 Cumulative Subject Index for Volumes 81-194

The CD-ROM uses EBT's Dynatext software (Electronic Book Technologies Inc., Providence, Rhode Island), and was developed by Lightbinders of San Francisco, California.

March 1995, \$999.00 \$750.00/ISBN: 0-12-000100-4\*

\*Price reduction valid through October 31, 1995.



When you order direct

Order from your local bookseller

or directly from



Order Fulfillment Dept. DM17915 6277 Sea Harbor Drive, Orlando, FL 32887 24-28 Oval Road, London NW1 7DX, U.K.

In the U.S. and Canada CALL TOLL FREE: 1-800-321-5068

FAX: 1-800-336-7377 e-mail: ap@acad.com In Europe, CALL 0181-300-3322

Call for a FREE electronic products catalog!

The Academic Press Dictionary of Science and Technology was recognized as one of American Libraries' Outstanding References of 1993!

## nary of Science and Technology CD-ROM

Version 1.0

Praise for the Print Version

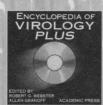
"Has a place on all reference shelves... Outstanding... Easy to use... The Academic Press Dictionary of Science and Technology captures the terminology of today and gives modern definitions... Highly recommended.

-BOOKLIST/REFERENCE BOOKS BULLETIN

Immediate information from the convenience of your desktop will soon be at hand, as the award-winning Academic Press Dictionary of Science **Technology** will be available this August on CD-ROM. This multiplatform CD-ROM enhances the same features that made the popular Academic Press Dictionary of Science and Technology comprehensive, informative, and easy-to-use, and now offers a quick, user-friendly environment from the convenience of your desktop.

This exciting new product will benefit high school and college students; science writers; science literates; libraries, including high school, college, research, legal, general, and public; corporate information specialists; and anyone who runs across scientific or technical terms while using a word processor.

August 1995, \$79.95/ISBN: 0-12-200401-9



## **Encyclopedia** of

R.G. Webster and A. Granoff

Praise for the Print Version

"I consider this encyclopedia to be a valuable addition to my own virology library. I believe the readership which will most benefit by this work includes basic scientists in virology and microbiology. If clinicians, students, and others need access to this information, they will find it on the shelves of libraries which choose to house this compilation, and I hope that many libraries will make that choice."

-HARLEY A. ROTBART, University of Colorado School of Medicine

The print version of the Encyclopedia of Virology, published in 1994, is one of Academic Press' finest achievements, receiving excellents reviews and recognition in the field. The CD-ROM version will contain many enhanced features, including structural photographs and diagrams cross-referenced to the text, a new section of high-resolution, color coded images and animations, plus advanced searching facilities including a thesaurus, annotation functions, and a journal recording device to record path through information searches.

Stand-Alone: \$475.00 (tentative)/ISBN: 0-12-000103-9 Networkable: \$675.00 (tentative)/ISBN: 0-12-000102-0 November 1995

Prices subject to change without notice. @1995 by Academic Press, Inc. All Rights Reserved. 20095-LK/KJ/TR/GEN