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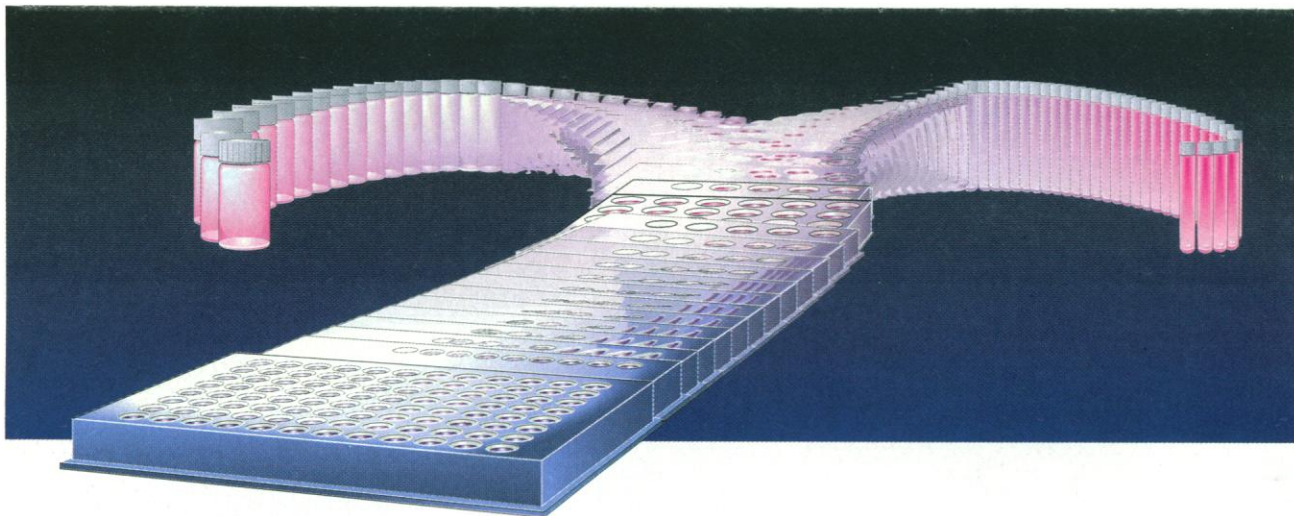
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therapy for hemophilia B



**110**  
Optical mapping  
of chromosomes

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As humans and mice look at each other through the eyes of the Human Genome Project, their similarities, which make the mouse a key model system, are more important than their differences. In this issue, a chart highlighting the similarity between the two genomes and showing the fusion of two mouse genome maps is

presented on page 67, along with an accompanying Article (page 57); a Policy Forum, Perspectives, and a News report indicate future challenges for the Genome Project. [Illustration: Susan Nowoslawski, Washington, DC]



## REPORTS

### Inclined-Field Structure, Morphology, and Pinning of the Vortex Lattice in Microtwinning YBa<sub>2</sub>Cu<sub>3</sub>O<sub>7</sub>

B. Keimer, F. Dogan, I. A. Aksay, R. W. Erwin, J. W. Lynn, M. Sarikaya

### Anatomy of the Photodissociation Region in the Orion Bar

A. G. G. M. Tielens, M. M. Meixner, P. P. van der Werf, J. Bregman, J. A. Tauber, J. Stutzki, D. Rank

### Protein Ladder Sequencing

B. T. Chait, R. Wang, R. C. Beavis, S. B. H. Kent

### Early Cambrian Ostracode Larvae with a Univalved Carapace

X.-g. Zhang and B. R. Pratt

### Deep and Bottom Water of the Weddell Sea's Western Rim

A. L. Gordon, B. A. Huber, H. H. Hellmer, A. Ffield

### Pressure-Induced Enhancement of $T_c$ Above 150 K in Hg-1223

M. Nuñez-Regueiro, J.-L. Tholence, E. V. Antipov, J.-J. Capponi, M. Marezio

### Protein Enhancement of Hammerhead Ribozyme Catalysis

Z. Tsuchihashi, M. Khosla, D. Herschlag

### Interaction of Mammalian Splicing Factor SF3a with U2 snRNP and Relation of Its 60-kD Subunit to Yeast PRP9

R. Brosi, K. Gröning, S.-E. Behrens, R. Lührmann, A. Krämer

### Correspondence Between a Mammalian Spliceosome Component and an Essential Yeast Splicing Factor

M. Bennett and R. Reed

### Interaction Between PRP11 and SPP91 Yeast Splicing Factors and Characterization of a PRP9-PRP11-SPP91 Complex

P. Legrain and C. Chapon

### Ordered Restriction Maps of *Saccharomyces cerevisiae* Chromosomes Constructed by Optical Mapping

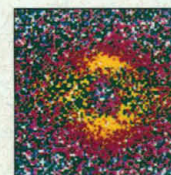
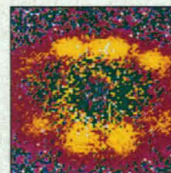
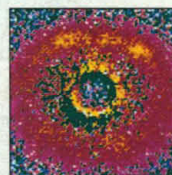
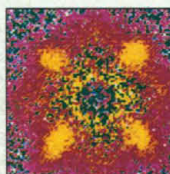
D. C. Schwartz, X. Li, L. I. Hernandez, S. P. Ramnarain, E. J. Huff, Y.-K. Wang

### Erythrocyte P Antigen: Cellular Receptor for B19 Parvovirus

K. E. Brown, S. M. Anderson, N. S. Young

### In Vivo Gene Therapy of Hemophilia B: Sustained Partial Correction in Factor IX-Deficient Dogs

M. A. Kay, S. Rothenberg, C. N. Landen, D. A. Bellinger, F. Leland, C. Toman, M. Finegold, A. R. Thompson, M. S. Read, K. M. Brinkhous, S. L. C. Woo



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Superconductor vortices revealed

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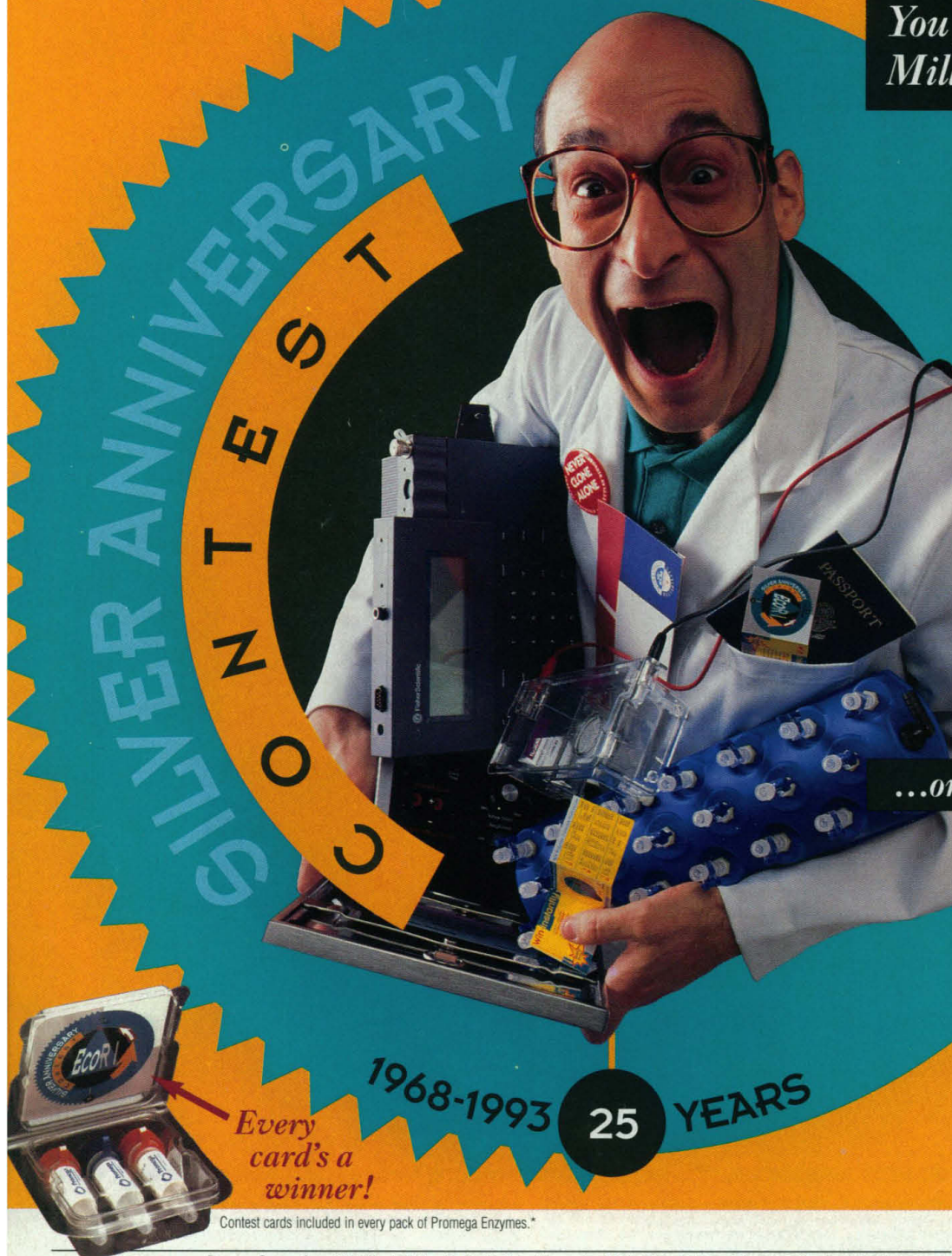
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## Superconductor vortices and twins

Neutron scattering can provide a sensitive measure of the vortex structure of a superconductor throughout its bulk structure. Keimer *et al.* (p. 83) studied a single crystal of  $\text{YBa}_2\text{Cu}_3\text{O}_7$  in a 0.5-tesla magnetic field that was varied over angles of  $0^\circ$  to  $80^\circ$  with respect to the crystal's *c* axis. They find that twin planes lock in the vortex lattice orientation up to a surprisingly high inclination angle of  $70^\circ$ ; for larger angles, a chain state is observed. Such strong flux pinning by twin planes should prove useful for maintaining high critical currents in applications.

## A difference approach to protein sequencing

Protein sequencing is almost synonymous with the Edman degradation, in which the sequence is determined by removing one residue at a time. Chait *et al.* (p. 89) have developed a method that has the promise of high sample throughput and low cost. The protein is subjected to numerous cycles of a rapid Edman degradation in which a terminating agent is used. This process generates a series of peptide fragments that differ from the next by one residue. The mass spectrum of this mixture reveals parent peaks whose mass differences yield the protein sequence. Posttranslationally modified and nonnatural amino acids are readily determined.

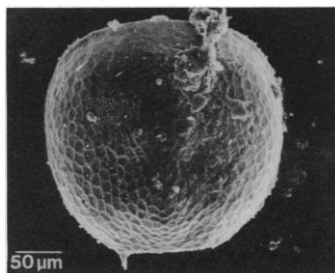
## Ostracode ontogeny

Most extant and fossil ostracodes, which are small crustaceans, are bivalved and have bivalved instars. Zhang and Pratt (p. 93), however, describe fossils from Lower Cambrian

## Lighting up the bar

Neutral atomic and molecular gas is one of the major components of the galactic interstellar medium but has been difficult to study because the neutral gas clouds have low surface brightness and emit and absorb primarily at infrared wavelengths. Tielens *et al.* (p. 86) conducted high-sensitivity, high-resolution observations of the Orion Bar, a prominent front of emission associated with nearby ionized gas features. By separately mapping emission from hydrogen, carbon monoxide, and polycyclic aromatic hydrocarbons, they could track the penetration of ionizing ultraviolet radiation into the neutral gas and show that the emission is due to external ionization and not, as had been proposed, a shock wave.

rocks in China indicating that some of the earliest ostracodes had univalved instars that later developed into bivalved instars and adults. The recognition of this transition during ontogeny



implies that a bivalved capacity evolved before the appearance of mineralized skeletons in the Cambrian.

## Higher temperature superconductors

One way to increase the transition temperature ( $T_c$ ) of copper oxide superconductors is to squeeze them. Nuñez-Reguero *et al.* (p. 97) found that the  $T_c$  of the homologous series  $\text{HgBa}_2\text{Ca}_{n-1}\text{Cu}_n\text{O}_{2n+2+\delta}$  increased continuously with applied pressure. At atmospheric pressure,  $T_c$  is 133 kelvin; it increased to 157 kelvin, the highest-yet observed for any material, at an applied pressure of 23.5 gigapascals. No saturation effects were observed; thus, modifying the solid in a way that mimics

the effects of compression might result yield higher  $T_c$ 's at ambient pressure. See also a News story by Amato (p. 31).

## More rapid ribozymes

To increase the specificity of ribozyme function, a long recognition sequence is required. Although they increase the binding of the substrate, paradoxically such long recognition sequences reduce the rate of product release and also reduce the specificity of the ribozyme. Tsuchihashi *et al.* (p. 99) report on the use of a derivative of the p7 nucleocapsid (NC) protein of HIV-1, a nonspecific RNA binding protein, in ribozyme reactions. The NC protein enhanced both the cleavage rate and the specificity of a hammerhead ribozyme possibly as a result of the strand exchange activity of NC.

## Joining splicing

Although yeasts utilize fewer proteins in their splicing machinery than mammals, multiple approaches have shown that elements of the splicing machinery have been conserved through evolution. The assembly of the prespliceosome in mammals requires the splicing

factor, SF3a. Brosi *et al.* (p. 102) show that SF3a is composed of three proteins, one is homologous to yeast PRP9, and all three are part of the U2 snRNP. Bennett and Reed (p. 105) isolated a mammalian spliceosome component (another of the SF3a proteins) and found that it is homologous to yeast PRP11 splicing factor. Legrain and Chapon (p. 108) show that PRP9 and PRP11 are also complexed in yeast.

## Restriction maps under the microscope

A direct optical approach has been developed for determining the physical map of restriction enzyme cutting sites from small amounts of genomic DNA. Schwartz *et al.* (p. 110) imaged large fluorescently stained DNA molecules (0.2 to 1.0 megabases) that were elongated under flow conditions and then fixed in agarose gel. Restriction enzyme cutting releases the flow-induced tension along the molecule. The tips of nascent ends curl up to form bright balls that are easily observed under the microscope. The fragments can then be sized by length or overall intensity. Maps for several yeast chromosomes were in good agreement with those determined from pulsed-field gel electrophoresis.

## Mouse map

The focus of this year's special genome issue is the mouse, one of the classic biological model systems (see Editorial, p. 11, for a synopsis). As for earlier issues on the human and *Drosophila* genomes and the X chromosome, a chart highlights areas of recent progress, intense scrutiny, and remaining gaps.

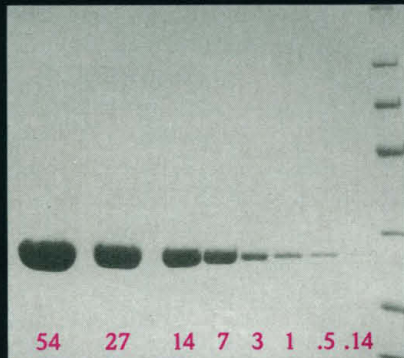


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SDS-PAGE of recombinant BamH I restriction endonuclease demonstrating purity of enzyme preparation from the cloned BamH I gene. Lanes represent a 2-fold dilution from 54 to 0.14  $\mu$ g BamH I per lane. Lane 9 contains molecular weight standards. Molecular weight of BamH I protein = 24,570 daltons. This preparation was used to grow BamH I crystals which led to solving its three-dimensional structure. (A. Aggarwal, personal communication)

### 62 Recombinant Restriction Endonucleases

|         |           |          |                    |
|---------|-----------|----------|--------------------|
| Aat II  | BspE I    | Hind III | Not I              |
| Acc I   | BspH I    | Hinf I   | Pst I              |
| Afl II  | Dde I     | HinP1 I  | Pvu II             |
| Afl III | Dpn I     | Hpa I    | Sac II             |
| Alu I   | Dpn II    | Hpa II   | Sal I              |
| Ase I   | Eag I     | Kas I    | Sau96 I            |
| Ava I   | EcoO109 I | Mbo II   | Sfi I              |
| Ava II  | EcoR I    | Msc I    | Sma I              |
| Avr II  | EcoR V    | Msp I    | Sph I              |
| BamH I  | Fok I     | Mwo I    | Sty I              |
| Ban I   | Fsp I     | Nae I    | Taq <sup>+</sup> I |
| Bbv I   | Hae II    | Nco I    | Xba I              |
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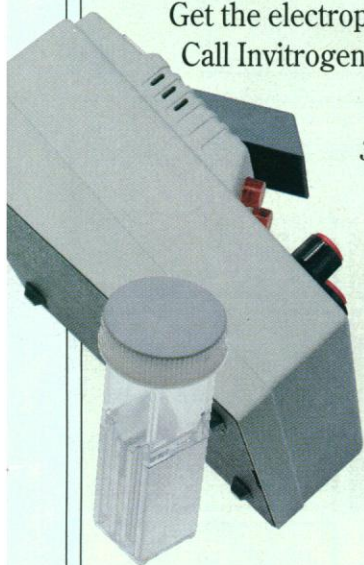
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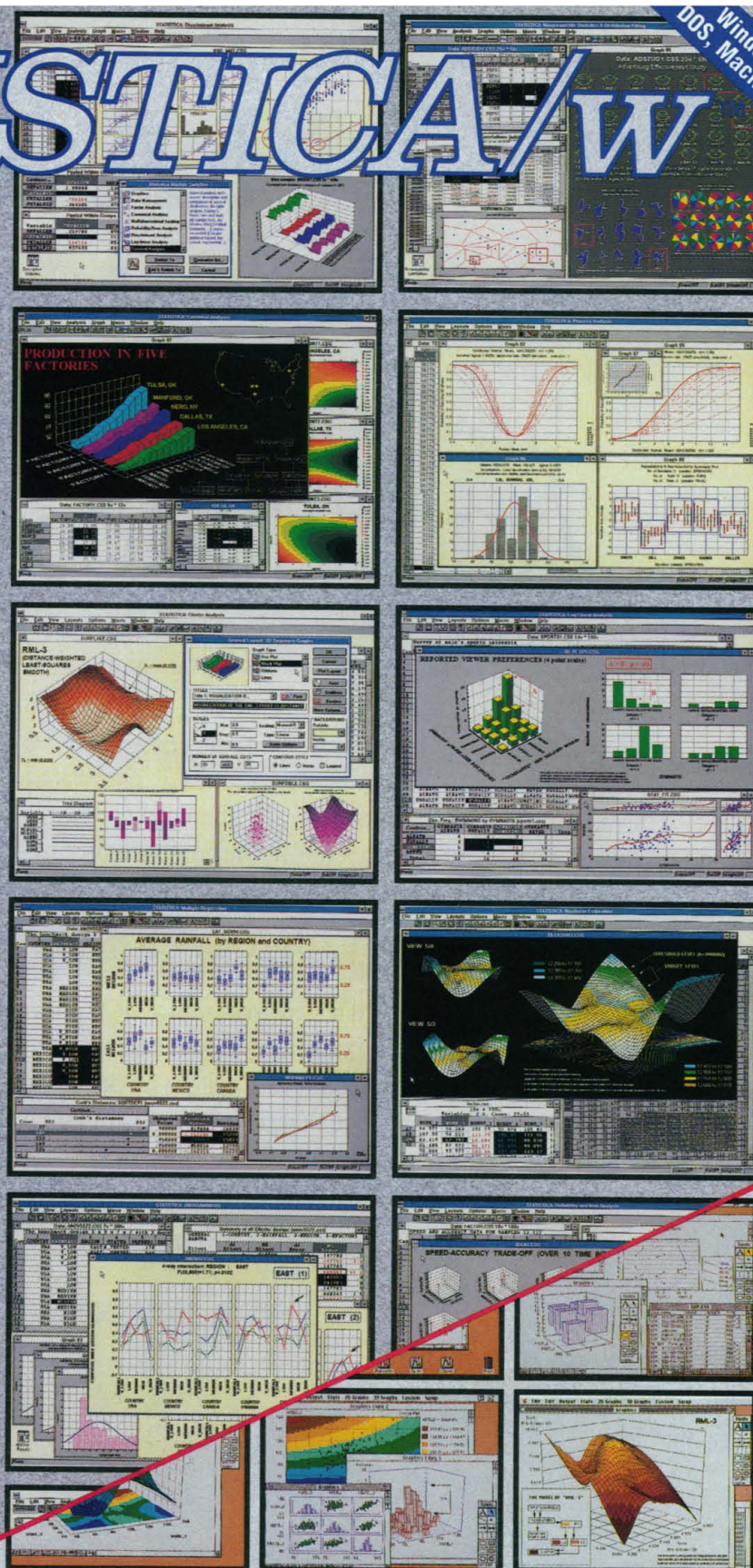


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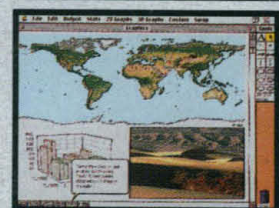
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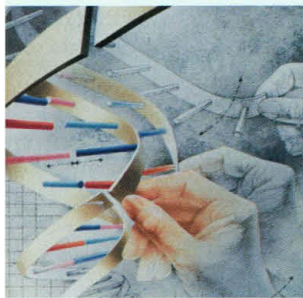


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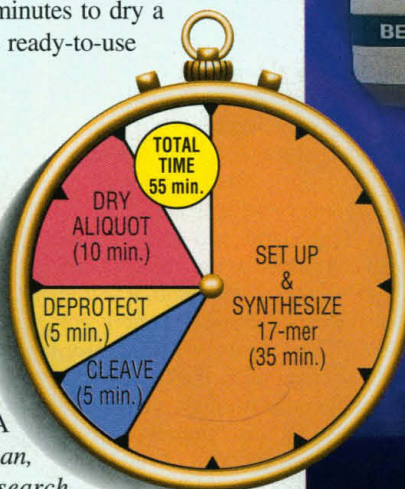
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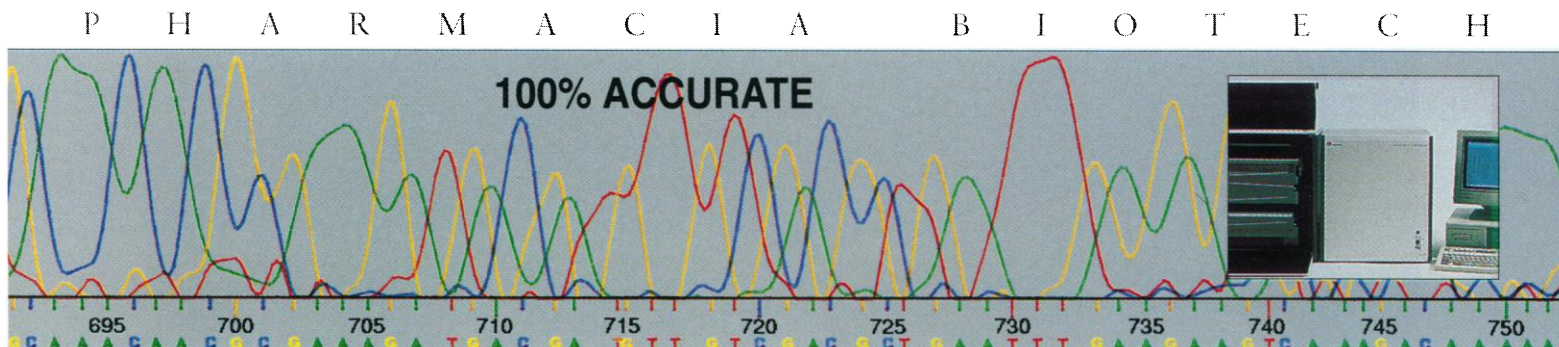
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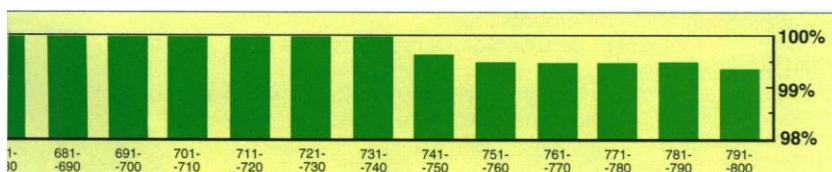
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1. Data supplied by M. Uhlen and T. Hultman from routine sequencing run at the Royal Institute of Technology, Stockholm, Sweden.
2. Comparison of three non-isotopic automated DNA sequence analysis systems. Poster presentation at the San Diego Conference on Nucleic acids, Nov. 20-22, 1991. Van Ranst, M., Fiten, P., Voet, M., Volckaert, G., Opdenakker, G.
3. Uniform scoring system for the assessment of DNA sequencing accuracy. *Meth. Mol. Cell. Biol.* 3 (1992) 243-245, Van Ranst, M., Fiten, P., Voet, M., Volckaert, G., Opdenakker, G.
4. Sequence length and error analysis of Sequenase and automated Taq cycle sequencing methods. *BioTechniques* 14 (1993) 442-447, Koop, B.F., Rowan, L., Chen, W.-Q., Deshpande, P., Lee, H., Hood, L.
5. An efficient low redundancy large scale DNA sequencing strategy: Primer walking on plasmid and cosmid DNA using T7 DNA polymerase and fluorescein-15'-dATP as internal label. Submitted for publication in *BioTechniques*, Voss, H., Wiemann, S., Zimmermann, J., Grothues, D., Sensen, C., Schwager, C., Stegemann, J., Erfle, H., Rupp, T., Sproat, B., Ansorge, W.

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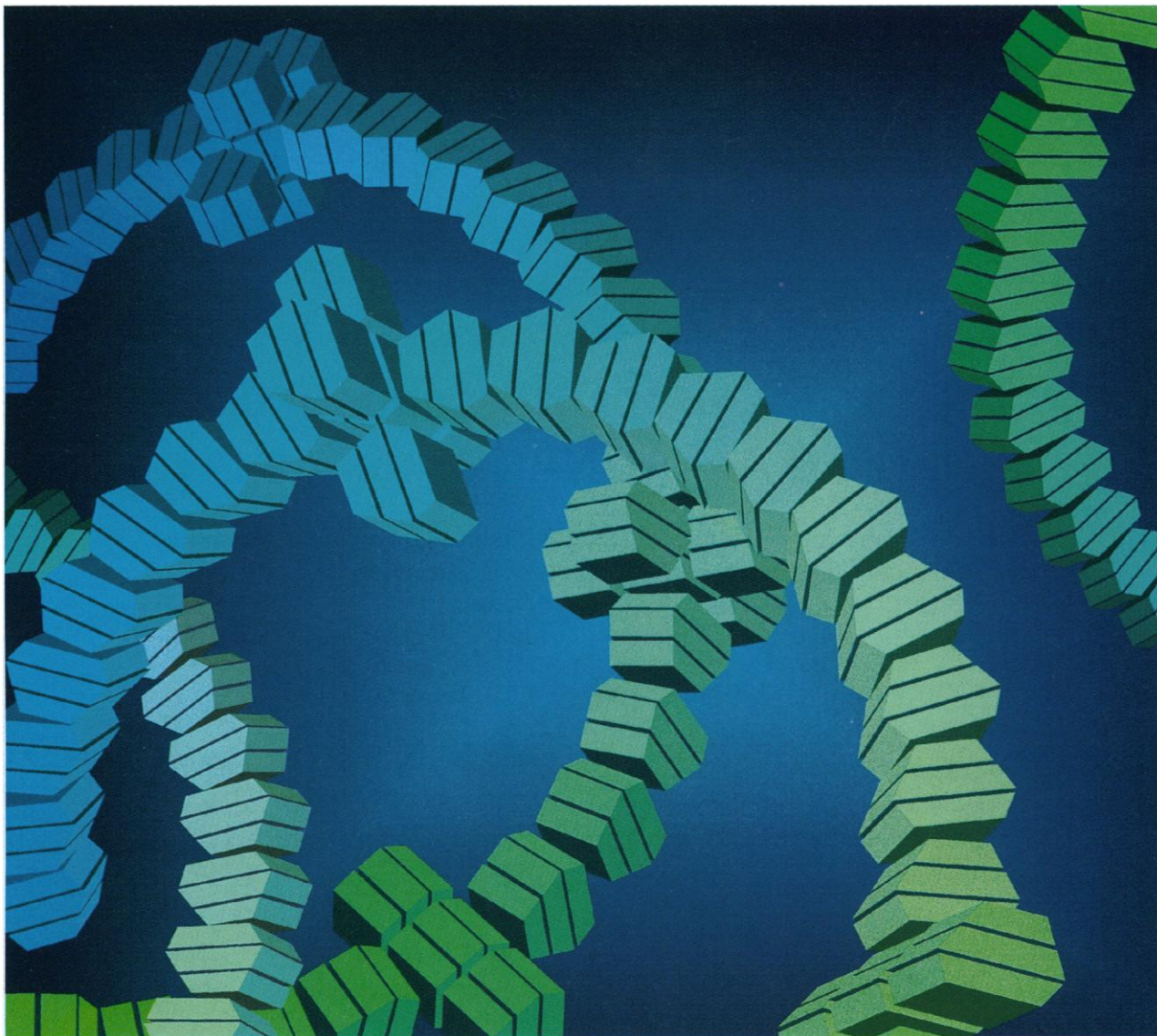
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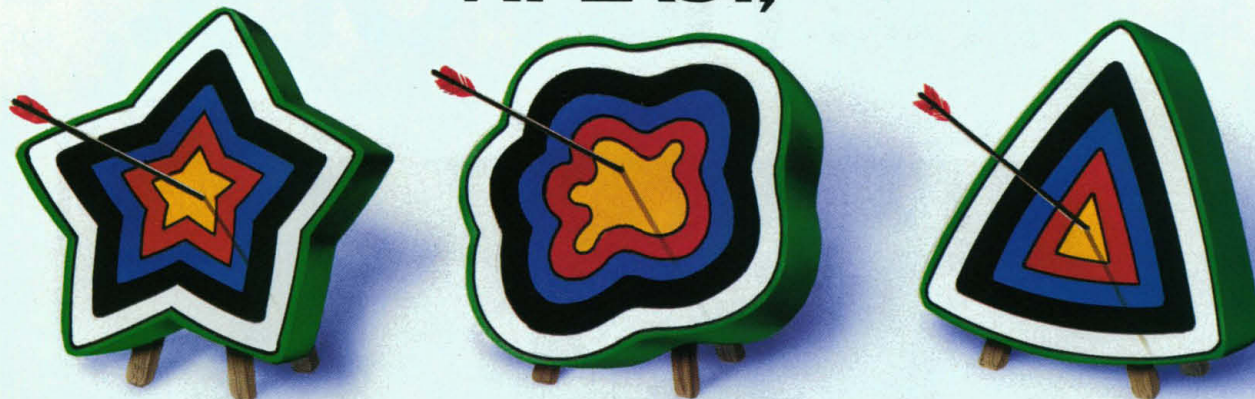
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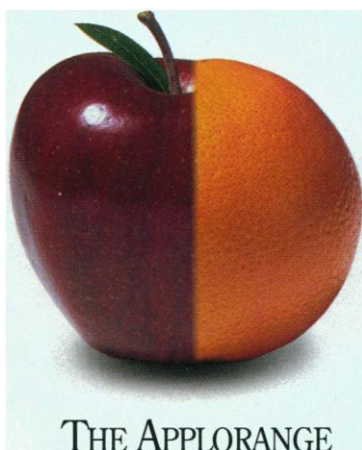
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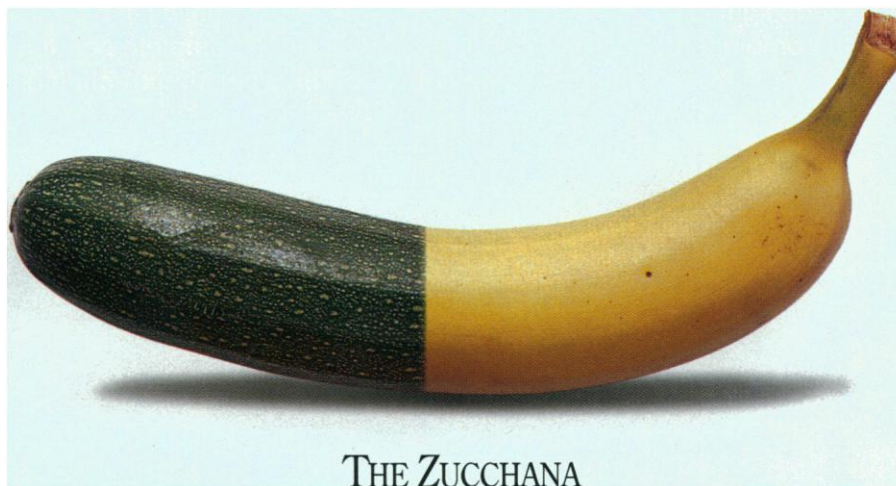




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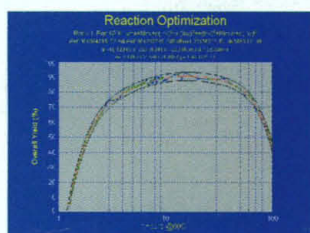
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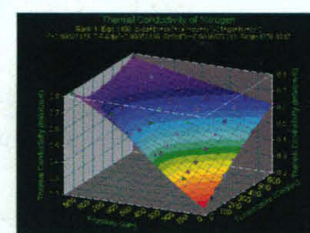
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| Hardware                  | 486 CPU/33MHz |
| No. of data points        | 100           |
| No. of equations          | 37,231        |
| Time to fit               | 168 seconds   |

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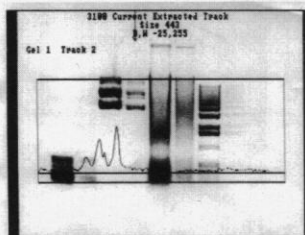


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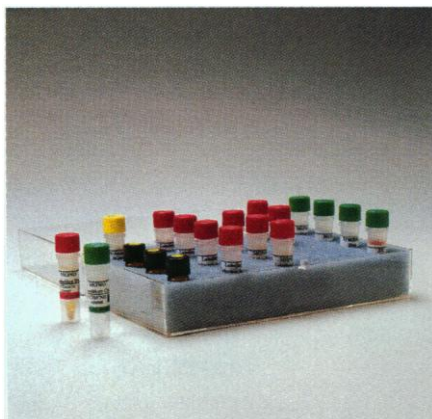


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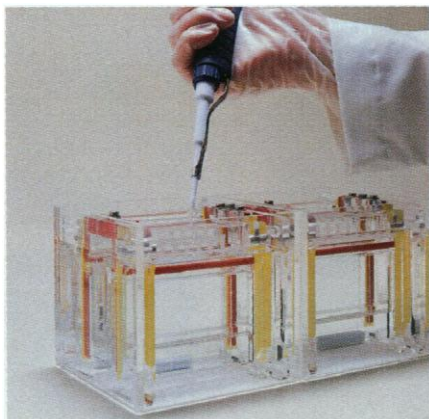
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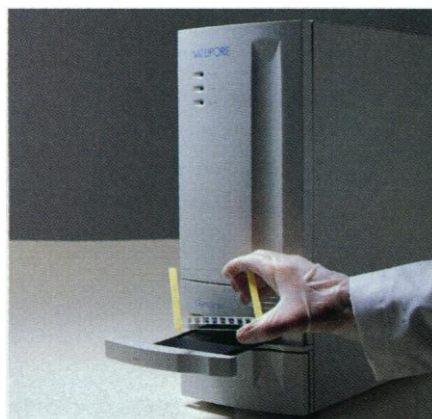
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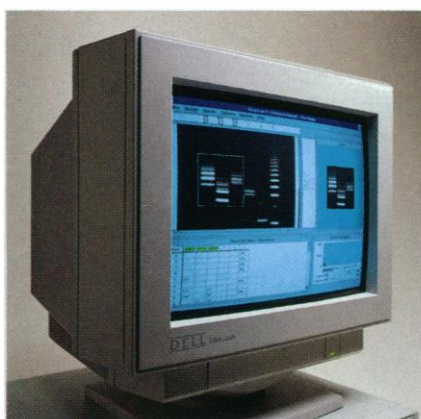
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MILLIPORE



## The Division of Cancer Etiology

## National Cancer Institute

Announces to the Scientific Community the Availability of the Following  
Resources/Services for Cancer Related Research As Noted Below:

### Biological Resources

■ Cell Culture Identification Service. Using Isozyme Analysis, Immunofluorescence and Karyotypic Analysis (Chromosome Banding).

**Contact:** Dr. Joseph Kaplan  
Children's Hospital of Michigan  
3901 Beaubien Boulevard  
Detroit, MI 48201  
(313) 745-5570

Citing Contract #N01-CP-33063

**Cost:** Reasonable; inquire with specific requests.

■ Goat Antisera Against: Avian, Bovine, Feline, Murine, and Primate Intact Viruses and Viral Proteins; Antibodies to Immunoglobulins for a number of species. Preimmune Sera available for some Virus Antisera.

**Contact:** Alice K. Robison, Ph.D.  
BCB Repository  
Quality Biotech, Inc.  
1667 Davis Street  
Camden, NJ 08104  
(609) 966-8000  
(609) 342-8078 FAX  
Citing Contract #N01-CP-15665

**Cost:** \$75.00/5 ml. (Antisera)  
25.00/5 ml. (Preimmune Sera)  
65.00/100 ml. (Immunoglobulins)  
(Frozen Material)

■ Viruses: Avian, Feline, Murine, and Primate Viruses Produced in vivo and in vitro.

**Contact:** Alice K. Robison, Ph.D.  
BCB Repository  
Quality Biotech, Inc.  
1667 Davis Street  
Camden, NJ 08104  
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Citing Contract #N01-CP-15665

**Cost:** Reasonable; inquire with specific requests.

■ Monoclonal Antibodies are available with specificities for synthetic peptides representing the amino acid sequences of the left end, right end and active site of the oncogene products of avian and mammalian retroviruses. Blocking peptides are also available, as are a limited number of cell lines producing the monoclonal antibodies.

**Contact:** Alice K. Robison, Ph.D.  
BCB Repository  
Quality Biotech, Inc.  
1667 Davis Street  
Camden, NJ 08104  
(609) 966-8000  
(609) 342-8078 FAX  
Citing Contract #N01-CP-15665

**Cost:** Peptides —\$25.00/mg.  
Ascites Fluid — 45.00/ml.  
Cell Culture —100.00/culture.  
(Plus Shipping and Handling)

■ Human sera from donors with various malignancies (including nasopharyngeal carcinoma), non-malignant disorders, and from normal individuals.

**Contact:** Program Director, Research  
Resources, Biological  
Carcinogenesis Branch,  
DCE, NCI, NIH  
Executive Plaza North, Room 540  
Bethesda, MD 20892  
(301) 496-1951  
(301) 496-2025 FAX

**Cost:** Shipping and handling charges only.

■ The Division of Cancer Etiology's Registry of Experimental Cancers announces the availability of 16 different study sets containing histologic slides of rodent tumors. The study sets, with accompanying syllabi, illustrate a variety of spontaneous and induced tumors, chiefly of rats, mice, and mastomys. Each set is available to cancer investigators worldwide, without charge, for up to two months. Requests or inquiries should be addressed to:

**Contact:** Registry of Experimental Cancers  
National Cancer Institute  
Building 41, Room D311  
NIH, Bethesda, MD 20892  
USA

### Environmental Cancer

■ NCI's Chemical Carcinogenesis Research Information System (CCRIS) is available online through the National Library of Medicine's Toxicology Data Network (TOXNET) system. Through an interagency agreement between NCI and NLM, the CCRIS database has been built and will be maintained and updated as one of TOXNET's sponsored databases in the broad areas of chemistry, toxicology, and hazardous waste information. The CCRIS database contains evaluated data and information on carcinogens, mutagens, tumor promoters, cocarcinogens, metabolites of carcinogens, and carcinogen inhibitors derived from published review articles, ongoing current awareness survey of primary literature, NCI/NTP's short- and long-term bioassay studies, the IARC Monographs on the Evaluation of Carcinogenic Risk of Chemicals to Man, and special studies and reports.

**Contact:** Dr. Thomas P. Cameron  
Office of the Director  
Division of Cancer Etiology  
National Cancer Institute  
Executive Plaza North, Room 712  
Bethesda, MD 20892  
(301) 496-1625

**Cost:** Inquire

■ The Special Assistant for Environmental Cancer, Office of the Director, announces the availability of a limited number of copies of the following publications, which have been prepared under contract to NCI:

Survey of Compounds Which Have Been Tested for Carcinogenic Activity, PHS-149, 1987-1988 and 1989-1990.

**Contact:** Dr. Thomas P. Cameron  
Office of the Director  
Division of Cancer Etiology  
National Cancer Institute  
Executive Plaza North, Room 712  
Bethesda, MD 20892  
(301) 496-1625

**Cost:** Inquire

■ The National Cancer Institute, along with the National Institute of Environmental Health Sciences, the Centers for Disease Control, and the Food and Drug Administration, has, for many years, supported a study by the Michigan Department of Public Health dealing with an accidental exposure to polybrominated biphenyls.

The Michigan Long Term PBB Study is a well-maintained longitudinal database on 4,000 participants from rural farms in Michigan. This group was exposed to polybrominated biphenyls through consumption of contaminated farm animals and food products. The cohort was enrolled and characterized in 1975-76, establishing a database containing demographic, health history, medical condition, reproductive history, blood and tissue analyses, and chemical/environmental exposure information. Major life events—birth, death, cancer and major illnesses have been confirmed and updated annually. The project is currently completing a detailed recharacterization of all cohort members and their children. This longitudinal database is available for collaborative research investigating biological and human health outcomes from halogenated biphenyl exposure.

**Contact:** Dr. Harold E. B. Humphrey  
Michigan Department of  
Public Health  
Division of Health Risk Assessment  
3423 North Logan, P.O. Box 30195  
Lansing, MI 48909  
(517) 335-8350

**Cost:** Free to qualified investigators.



## Epidemiology Resources

■ The Tumor Virus Epidemiology Repository (TVER) contains sera and other biological samples from more than 13,000 patients and controls obtained in 12 countries. The TVER was established primarily to support collaborative research on the role of Epstein-Barr virus (EBV) in Burkitt's lymphoma and related diseases. Sera characterized for human herpes virus 6 (HHV) antibodies are also available. The TVER collection is available for new collaborative studies and some independent research. The most extensive collections are serum samples from patients with Burkitt's lymphoma.

**Contact:** Dr. Paul H. Levine  
Viral Epidemiology Branch  
DCE, NCI, NIH  
Executive Plaza North, Room 434  
Bethesda, MD 20892  
(301) 496-8115

**Cost:** Free to Collaborating Investigators;  
Others: Dependent on Processing Time

■ The National Cancer Institute has available the Animal Morbidity/Mortality Survey of Colleges of Veterinary Medicine in North America (also known as the Veterinary Medical Data Program). This registry of veterinary medical information represents patient data on animals seen at collaborating veterinary teaching facilities; 3 million hospital episodes have been abstracted and computerized in a standardized record format. Disease information is coded using the scheme of the Standard Nomenclature of Veterinary Disease and Operations. The computer tapes will be made available upon request.

**Contact:** Dr. Howard M. Hayes  
Environmental Epidemiology Branch  
EPB, DCE, NCI, NIH  
Executive Plaza North, Room 443  
Bethesda, MD 20892  
(301) 496-1691

**Cost:** Inquire

■ Human fibroblast cultures from individuals at high risk of cancer, selected members of cancer-prone families, and some normal family members are available. The collection is historical with unknown viability and contamination status. Catalog of cell lines unavailable and followup on many individuals is unavailable. Information requests should include potential use of cultures.

**Contact:** Chief, Genetic Epidemiology Branch  
DCE, NCI, NIH  
Executive Plaza North, Room 439  
Bethesda, MD 20892  
(301) 496-4375

**Cost:** Free to collaborating investigators.  
Others: \$70/cell line.

■ The National Institute of Allergy and Infectious Diseases and the National Cancer Institute have developed a repository of biological specimens from homosexual men. The specimens were collected through cooperative agreements with five major U.S. universities for studies of the natural history of acquired immune deficiency syndrome (AIDS). Information about applying for collaborative use of these specimens is available from the NIAID Project Officer or the NCI Co-Project Officer.

**Contact:** Chief, Epidemiology Branch, AIDS  
Program, NIAID  
CDC Bldg., Room 240  
National Institutes of Health  
Bethesda, MD 20892

or to Chief  
Extramural Programs Branch, EPB,  
Division of Cancer Etiology, NCI  
Executive Plaza North, Room 535  
Bethesda, MD 20892

■ The Epidemiology and Biostatistics Program of the National Cancer Institute has developed the Observed versus Expected (O/E) software system which calculates: (1) the number of observed events (e.g. cancer cases or deaths) in a study group at risk; (2) the number of expected events in a study group based on the rate of occurrence in some standard or referent population; (3) the ratio of observed to expected events; and (4) the significance of this ratio. The system is user friendly and capable of executing a series of calculations by different variables such as age, time group, date of exposure, age at date of exposure, duration of exposure, year relative to entry and cause of event. The O/E System provides tables by race, sex and user defined variables, allows user defined latency intervals and accepts standard or user prepared rates. O/E is written in COBOL and is exportable to most mainframes.

**Contact:** Ruth Wolfson  
Epidemiology and Biostatistics  
Program, DCE, NCI, NIH  
Executive Plaza North, Room 443  
Bethesda, MD 20892  
(301) 496-1691

**Cost:** Free to investigators interested in  
epidemiologic research.

■ The Epidemiology and Biostatistics Program of the National Cancer Institute (NCI) has developed an Occupational Mortality Analysis software system which calculates Proportionate Mortality Ratios, Proportionate Cancer Mortality Ratios, or Mortality Odds Ratios using occupational information on the death certificates from 24 states for 1984-1989. The data were assembled through a collaborative effort involving the National Center for Health Statistics, the National Institute for

Occupational Safety and Health and NCI. The program is user friendly and allows analysis of data by (1) occupation, industry, or occupational/industry combinations; (2) age group; (3) states or geographic regions; (4) race groups, (black and white); (5) sex, and (6) underlying causes of death. The program is written in Wylbur Command Procedures and is exportable to most mainframes.

**Contact:** (To obtain the program and relate information):

Mustafa Dosemeci, Ph.D.  
Occupational Studies Section  
EBP, DCE, NCI, NIH  
Executive Plaza North, Room 418  
Bethesda, MD 20892  
Phone: (301) 496-9093  
Fax: (301) 402-1819

(For questions concerning the nature and source of the occupational mortality data):

Jeff Maurer, M.S.  
Mortality Statistics Branch  
Division of Vital Statistics, NCHS  
6525 Belcrest Rd., Room 840  
Hyattsville, MD 20782  
Phone: (301) 436-8884  
Fax: (301) 436-7066

**Cost:** Free to investigators interested in occupational epidemiologic research.

■ The Epidemiology and Biostatistics Program of the National Cancer Institute (NCI) has developed a computer-aided occupational and industrial code searching program (CODESEARCH) which allows the code assigner to select appropriate codes from existing classification systems for job or industrial titles from work histories of the study subjects. The program is user friendly and allows searches from four occupational Classification Systems (1977 Standard Occupational Classification Manual [SOC]; 1980 SOC; 1970 Bureau of Census Occupational Classification System [BOCOC]; and 1980 BOCOC, and four industrial classification systems (1972 Standard Industrial Classification System [SIC]; 1987 SIC; 1970 Bureau of Census Industrial Classification System [BOCIC]; and 1980 BOCIC). The program is written using PC-Clipper software and is exportable to most 486 PCs.

**Contact:** Mustafa Dosemeci, Ph.D.  
Occupational Studies Section  
EBP, DCE, NCI, NIH  
Executive Plaza North, Room 418  
Bethesda, MD 20892  
Phone: (301) 496-9093  
Fax: (301) 402-1819

**Cost:** Free to investigators interested in assignment of occupational and industrial titles from work history data

## Chemical Resources

■ Analytical support for the collection, separation, and elucidation of environmental carcinogens including combustion and smoking-related exposures. A contractor with experience in the development of analytical methods for the determination of constituents of cigarette smoke and cigarette smoke condensates, and of specialty instrumentation for inhalation toxicology is available to assist qualified investigators with particular interest in human and animal model exposure to environmental and sidestream smoke. An extensive chemical data base on smoke and smoke condensate components is available.

**Contact:** Harold E. Seifried, Ph.D.  
Chemical and Physical  
Carcinogenesis Branch, DCE, NCI  
Executive Plaza North, Room 700  
Bethesda, MD 20892  
(301) 496-5471  
(301) 496-1040 FAX

**Cost:** Inquire

■ Chemical Carcinogen Reference Standard Repository: Reference quantities of over 750 compounds are available. The newest additions are dilute aqueous standards of PAH deoxyguanosine-3'-monophosphates for Randerath <sup>32</sup>P post labelling assays. Other classes of available compounds are: fecapentaenes, food mutagens, polynuclear aromatic hydrocarbons (PAH); PAH metabolites, radiolabeled PAH metabolites, nitrogen heterocycles, nitrosamines/nitrosamides, aromatic amines, aromatic amine metabolites, azo/azoxy aromatics, inorganics, nitroaromatics, pesticides, pharmaceuticals, natural products, dyes, dioxins and chlorinated aliphatics. A number of radiolabeled PAH metabolites and nitrosamines are also available. Data Sheets provided with the compounds include chemical and physical properties, analytical data, hazards, storage, and handling information. Catalog available upon request.

**Contact:** Manager, NCI Chemical  
Carcinogen Repository  
Midwest Research Institute  
425 Volker Boulevard  
Kansas City, MO 64110  
(816) 753-7600, Ext. 523  
(816) 753-3664 FAX

Manager, NCI Radiolabeled  
Chemical Repository  
CHEMSYN Science Laboratories  
13605 W. 96th Terrace  
Lexena, KS 66215  
(913) 541-0525  
(913) 888-3582 FAX

**Cost:** Subject to chemical class code and quantity (see catalog) includes handling and shipping charges



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**SCIENCE**  
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**GENOME MAPS IV**

**SCIENCE COORDINATOR:** Barbara R. Jasny

**AUTHORS**

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Michael Wessel, Hillary Katz, Lincoln D. Stein,  
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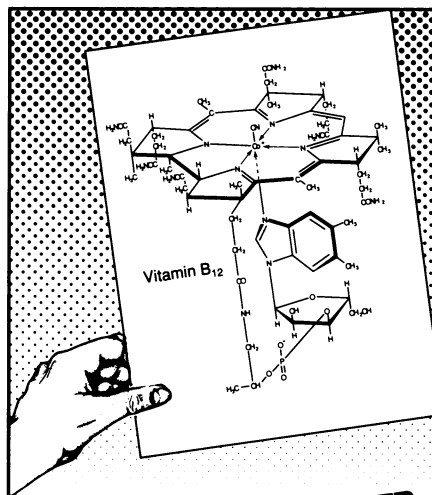
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