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AIDS

THE UNANSWERED QUESTIONS



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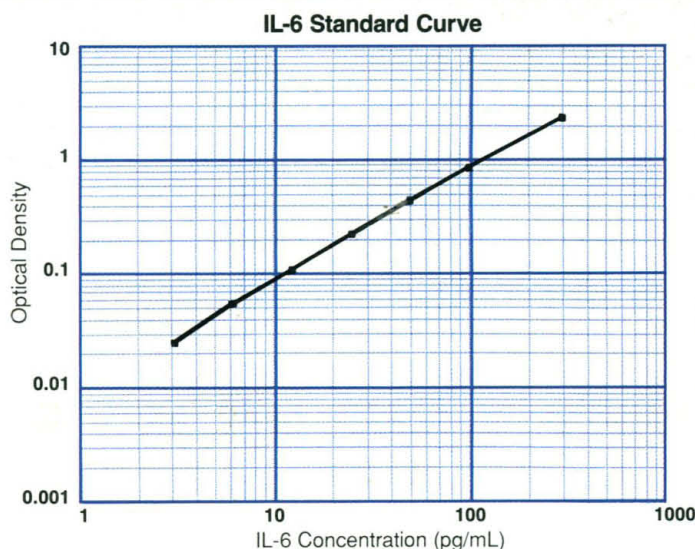
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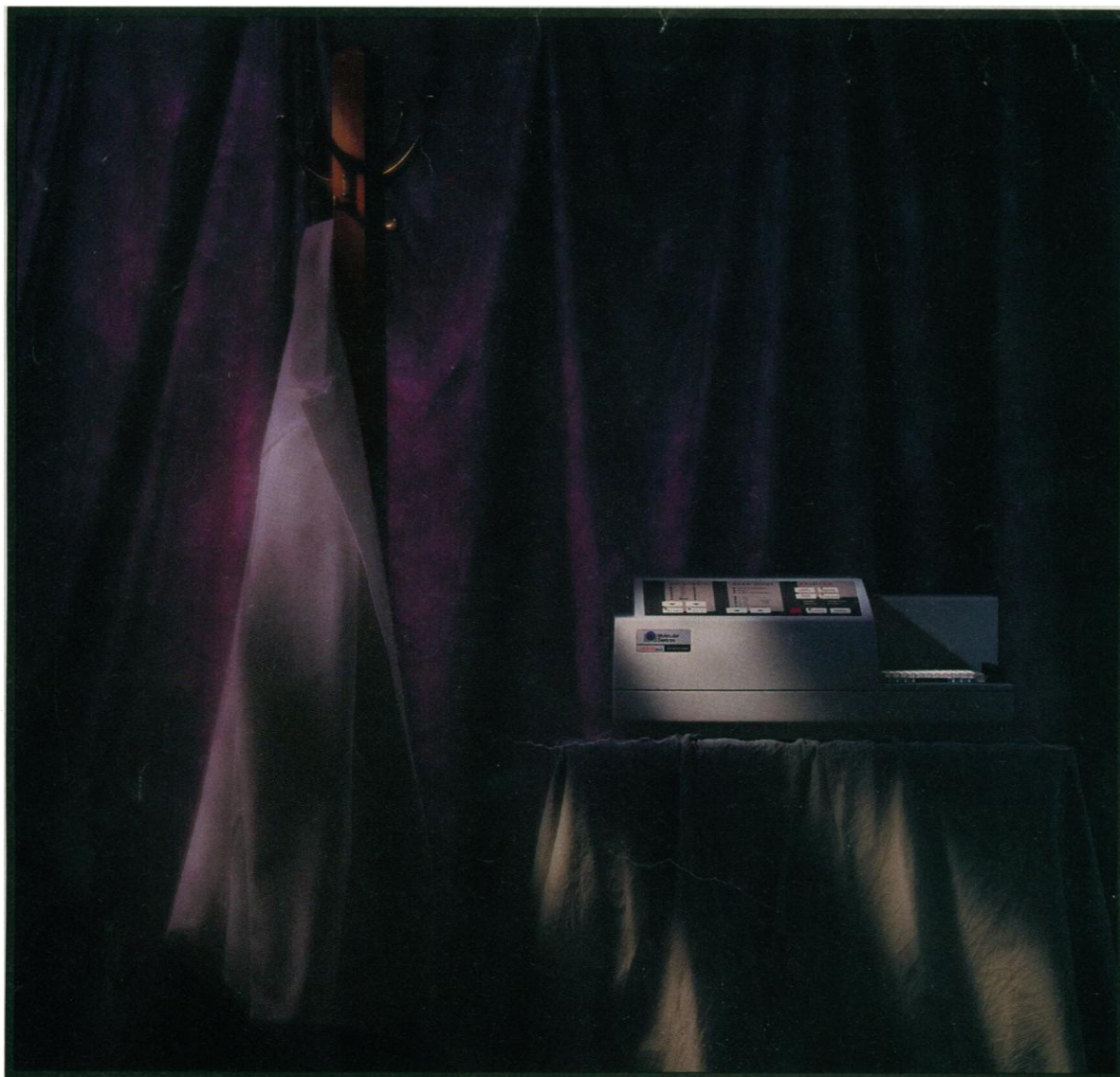


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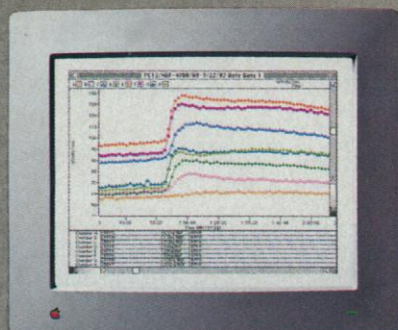
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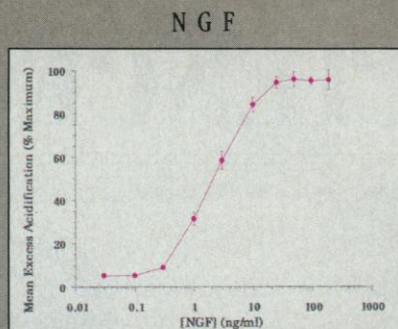
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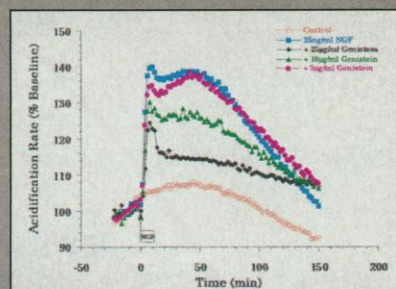




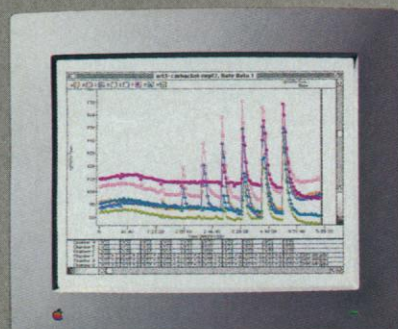
1 Real-time acidification rate data showing the response of PC12 cells to a 12 min. exposure to nerve growth factor (NGF:0.1-200ng/ml).



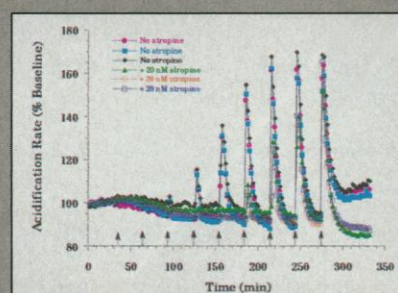
2 Dose response of PC12 cells to NGF. Each point represents data from 7 separate experiments. Calculated EC_{50} value for NGF was 1.9 ± 0.7 ng/ml ($\sim 152 \pm 50$ pM) ($\bar{x} \pm S.E.M.$).



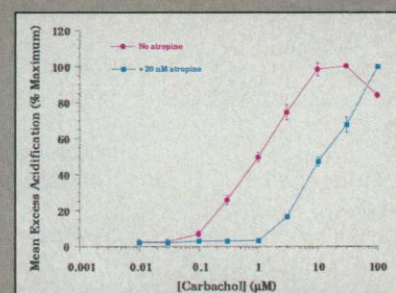
3 Dose dependent inhibition of NGF (25 ng/ml) response by tyrosine kinase inhibitor, genistein (3-25 μ g/ml).



1 Real-time acidification rate data demonstrating stimulation of CHO cells transfected with muscarinic M_1 receptor. Increasing doses of carbachol (10nM-100 μ M) given in presence and absence of 20nM atropine.



2 Data from previous figure expressed as percentages of the basal acidification rates before initial addition of carbachol.



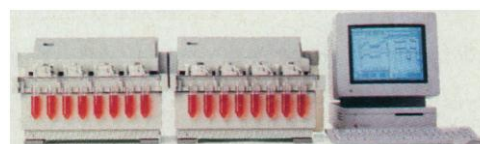
3 The EC_{50} values were calculated as 1.00 ± 0.08 μ M for carbachol alone and 11.50 ± 0.09 μ M for carbachol in the presence of 20nM atropine ($\bar{x} \pm S.E.M.$; $n = 3$).

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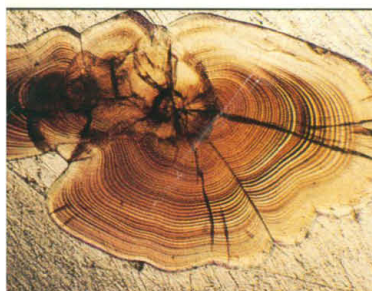
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Fish ear stones tell
age-old tales

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COVER

All of science is driven by unanswered questions, but in AIDS research 10 years after the discovery of the AIDS virus, there are many more questions than answers. In a special section beginning on page 1253, *Science* offers several different views of the

most important questions in the field, from a survey of 150 AIDS investigators to provocative views from leading researchers. The only remaining question is: When will there be answers?



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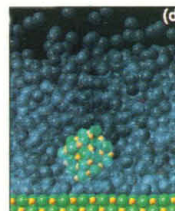
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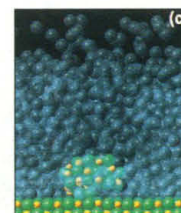
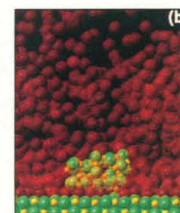
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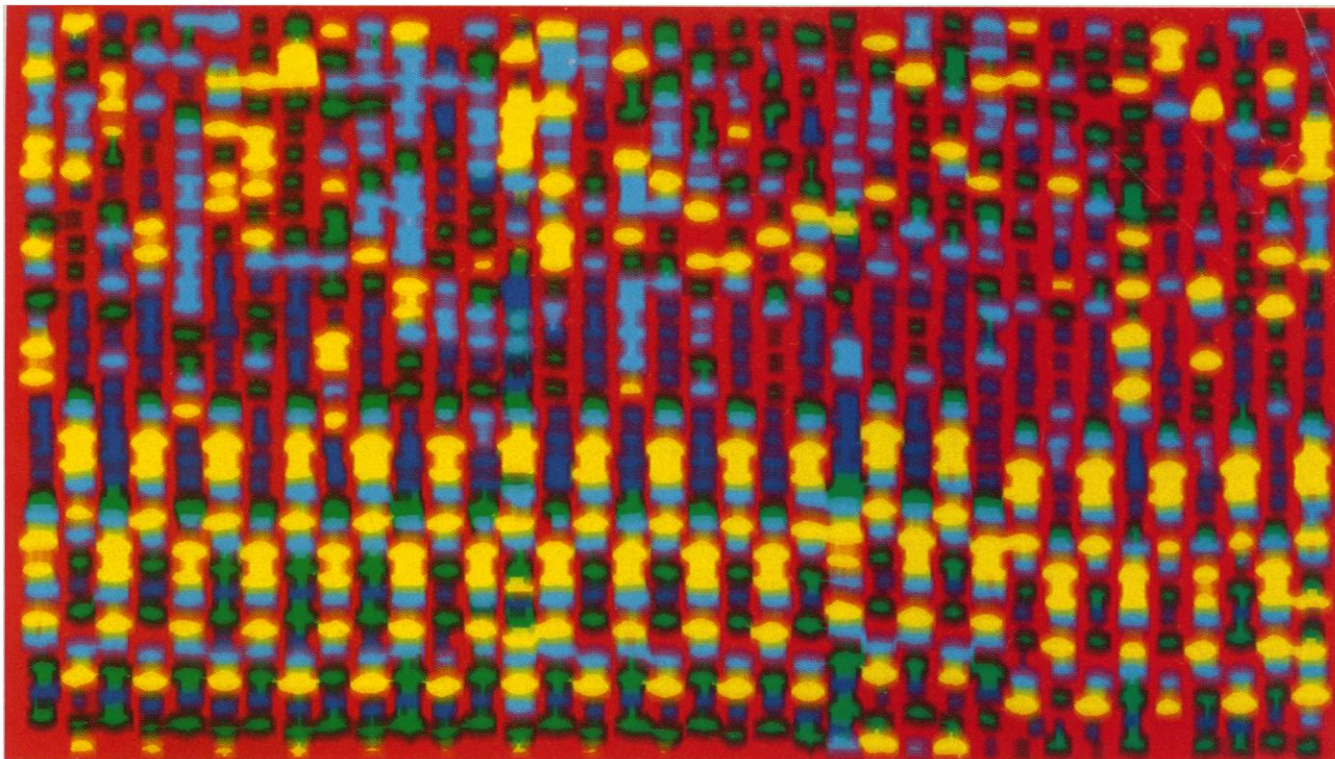


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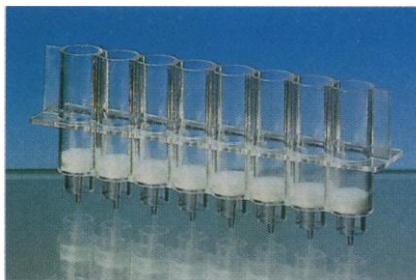
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Soft landings

One approach to growing materials is to consolidate nanoclusters after they are deposited onto a surface. Cheng and Landman (p. 1304) have performed numerical simulations of the dynamics of sodium chloride clusters during impact with adsorbed noble gas liquid films. For a liquid neon film, the cluster transfers energy efficiently to the cushion layer, leading to a controlled soft landing. For impacts on a higher density film of liquid argon, the cluster slows abruptly and is rapidly heated, with subsequent melting and fast cooling by argon evaporation. Such impact-processed clusters might be useful for the controlled growth of nanophase materials.

Polysaccharide synthesis

Chemical synthesis of peptides and oligonucleotides usually takes advantage of anchoring the polymer to a solid-phase support so that reactions take place at only one end of the chain. Danishefsky *et al.* (p. 1307) have developed solid-phase methods for oligosaccharide synthesis, which is complicated by the formation of stereospecific bonds at one to five hydroxyl sites. They eliminate several problems by using glycols to direct the chemistry. The glycol is epoxidized and can then glycosylate another glycol at the appropriate hydroxyl group. Failed couplings inactivate the double bond, so interior deletions are avoided.

New iron phase

Earth's liquid outer core and solid inner core are composed primarily of iron. Knowledge of

Polycrystalline films approach natural diamond

For electronic applications under extreme conditions, diamond offers numerous advantages, but single-crystal diamond films are expensive. Plano *et al.* (p. 1310) describe how chemical vapor deposition can be used to generate lower cost polycrystalline diamond films that overcome their main limitation as a semiconductor material, namely, their low carrier mobilities. Optimization of plasma deposition conditions have allowed high growth temperatures to be used and have increased grain size and decreased impurities. The carrier mobilities and lifetimes approach that of the highest quality single crystals of natural diamond.

the phase diagram of iron at high pressures and temperatures is key to understanding the composition and formation of the core. Saxena *et al.* (p. 1312) present experimental evidence for the existence of a previously unknown iron phase, which they call the β phase. Its structure is unknown, but it is stable at higher temperatures than the ϵ phase (hexagonal close-packed), which was generally thought to form the solid core.

Tumor suppressor identified

Von Hippel-Lindau (VHL) disease is a dominantly inherited syndrome that predisposes individuals to multiple cancers, including renal cell carcinomas. Previous work has shown that the VHL gene behaves as a tumor suppressor and maps to chromosome 3p25-p26. Latif *et al.* (p. 1317; see news story by Travis, p. 1233) have now identified the causative gene by characterizing deletions in VHL families that showed coincident transmission with the disease. The VHL gene, which also appears to be disrupted in some sporadic renal cell carcinomas, is evolutionarily conserved, and encodes an acidic repeat domain found in a trypanosome surface membrane protein.

Avoiding detection

Viruses are mainly detected by the immune system through major histocompatibility complex (MHC) class I antigens, which bind to viral peptides. Many viruses reduce MHC class I expression after infection, but Howcroft *et al.* (p. 1320) show that HIV-1 dramatically decreases expression by repressing class I promoter activity. Surprisingly, deletion constructs mapped this repressive effect to a transcriptional activator, the Tat protein. Cotransfection experiments showed that the spliced, or two-exon Tat protein, effects this repression but that the one-exon Tat protein has only a minor effect.

Runaway recombination

Defects in genetic recombination are thought to underlie ataxia-telangiectasia (A-T), an autosomal recessive disease characterized by neurological degeneration, immune dysfunction, and a high risk for cancer. Meyn (p. 1327) studied the rate of recombination in normal and A-T human fibroblasts. Although recombination rates were similar for both cell types for rearrangements between chromosomes, intrachromosomal recombination rates were 30 to 200 times higher for the

A-T cells. One possible cause for this increase could be defects in damage-sensitive checkpoints in the cell cycle that would allow DNA replication to finish before repair processes are completed.

Free at last

The lymphoid-specific transcription factor Elf-1 participates in the regulation of inducible gene expression during T cell activation, even though Elf-1 is present in similar quantities in both resting and activated T cells. Wang *et al.* (p. 1330) find that the Elf-1 protein may be sequestered or inactivated in resting T cells by interacting with unphosphorylated retinoblastoma protein (Rb). Upon T cell activation, Rb is phosphorylated and Elf-1 dissociates.

Getting the signal

The molecules that couple hormone and growth factor receptors on the cell surface to activation of the guanine nucleotide binding protein Ras are proving to be remarkably similar in all eukaryotes. Chardin *et al.* (p. 1338) have cloned the human homolog of the son of sevenless protein (hSos), a putative guanine nucleotide releasing factor (GRF) for Ras, from *Drosophila*. They found that hSos enhanced guanine nucleotide exchange by Ras and thus appears to function as an activator of Ras. The hSos protein could substitute for the Ras-GRF CDC25 in yeast and can bind to GRB2, a so-called adaptor protein that also binds to activated growth factor receptors. All of the molecular components required to transmit an extracellular signal to Ras have now been identified.



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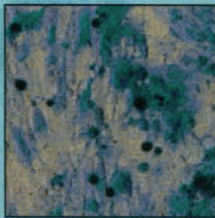


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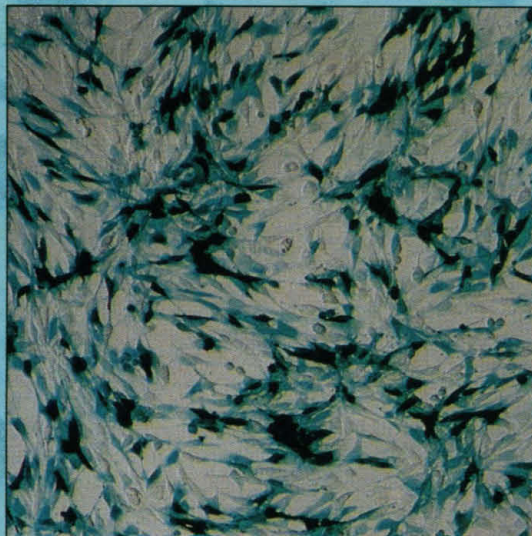
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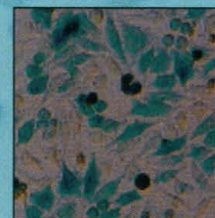
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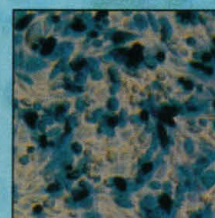
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Photographs depict β -galactosidase expression of cells transfected with pCMV β -gal DNA and LIPOFECTAMINE Reagent.

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1. Hawley-Nelson, P., Ciccarone, V., Gebeyehu, G., Jessee, J. and Felgner, P., (1993) *Focus* 15, in press.
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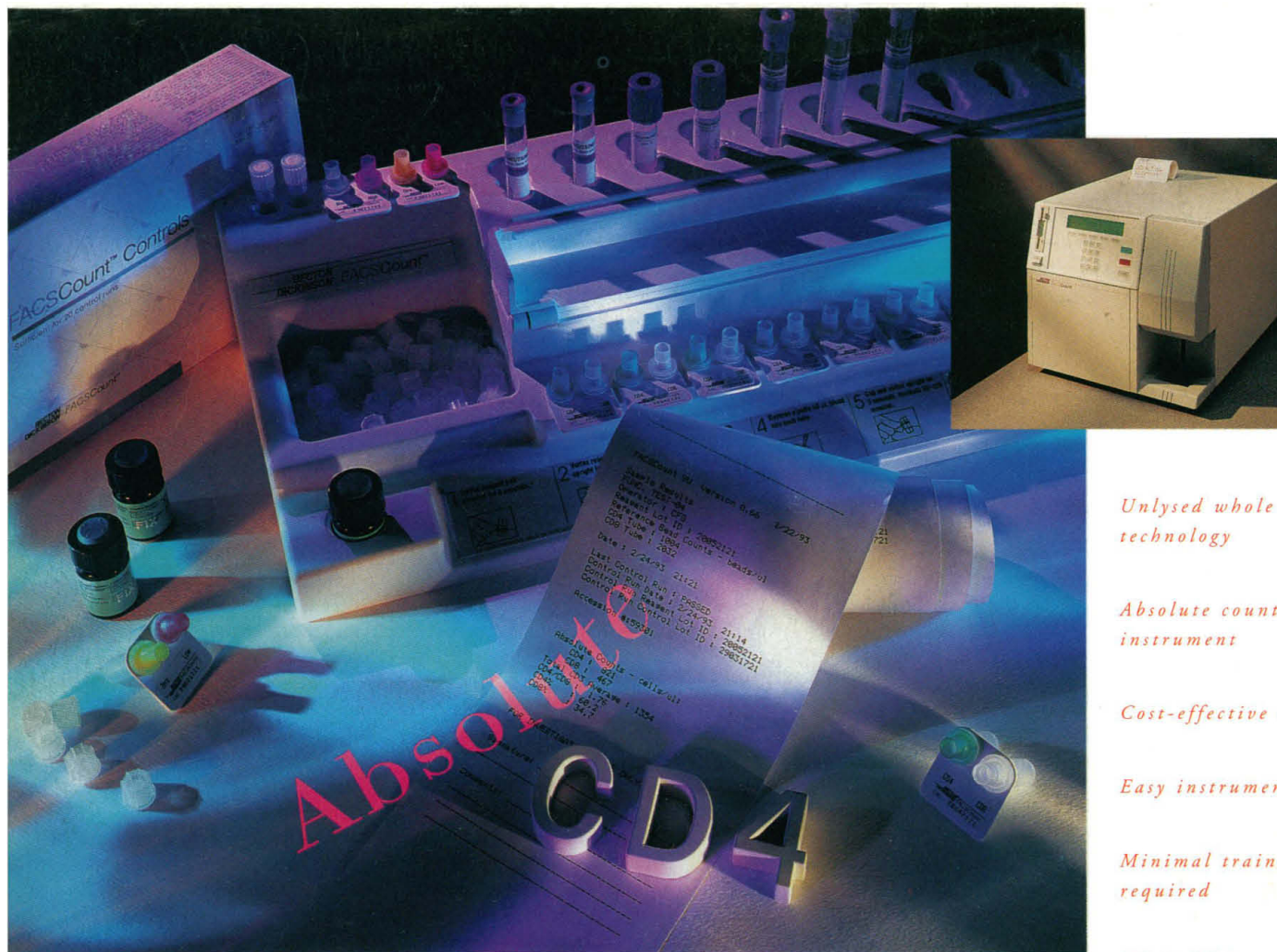
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Nikon asked inverted microscope users to make a wish list.

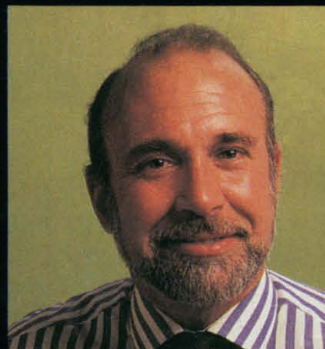
"Our microscope is used by a number of people. I wish the height of the eyepiece tube could be adjusted for each of us."



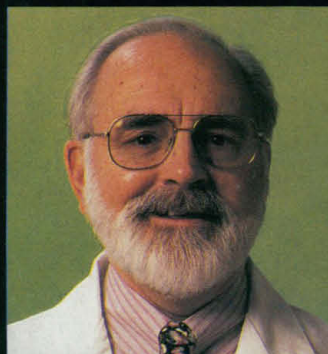
"With all the new fluorochromes available, I wish there was a quick-change multiple filter block system."



"High resolution DIC has become so much more important, I wish there was an easily set-up high N.A. DIC condenser."



"To accommodate extra-large chambers and micromanipulators, I wish there was more working distance above the stage."



"I just wish someone would design an inverted microscope with patch clamping, low-light fluorescence, IVF, confocal microscopy and other advanced applications in mind from the very start."



"I wish I had a microscope with more optical ports for simultaneous imaging."



"I'm at my microscope for hours at a time. I wish the controls were further forward so I could work more comfortably."



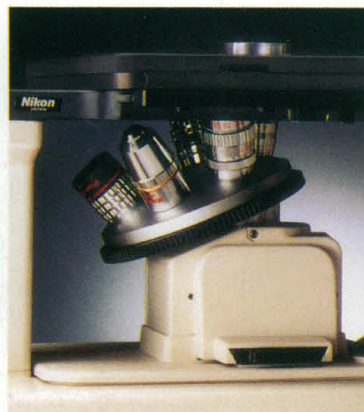
"For really critical specimen handling, my wish would be for a very low, very large, super stable stage."

Wish no more ...

Introducing the new Diaphot®. It's everything you've wished



Controls are far forward for maximum comfort.



Easy access to nosepiece and filters: dichroic cube fixed relative to focusing nosepiece.



0.85 and 1.4 N.A. DIC condensers for high resolution, high magnification microscopy.



Expanded space above stage and tiltable 100W illumination pillar for large specimens and micromanipulation.

Improving upon the world's finest inverted microscope was no easy task. But, after asking hundreds of users for their suggestions, we did it. The new Diaphot® features even brighter images, greater flexibility, more imaging capability, stability and comfort than its renowned predecessor, without sacrificing any of its legendary optical performance.

The new Diaphot is designed to suit applications from basic microscopy to UV fluorescence, *in-vitro* fertilization, laser scanning confocal microscopy, micromanipulation, patch clamping, microphotometry and high resolution video microscopy.

Important new CF® objectives with exceptional image sharpness, brightness and clarity have been added to the already extensive line of Nikon CF optics. Innovative optical design and advanced lens coating technology enhance transmission efficiency.

Comfort over long periods of operation was important to many people. We responded by placing the most frequently used controls of the new Diaphot within

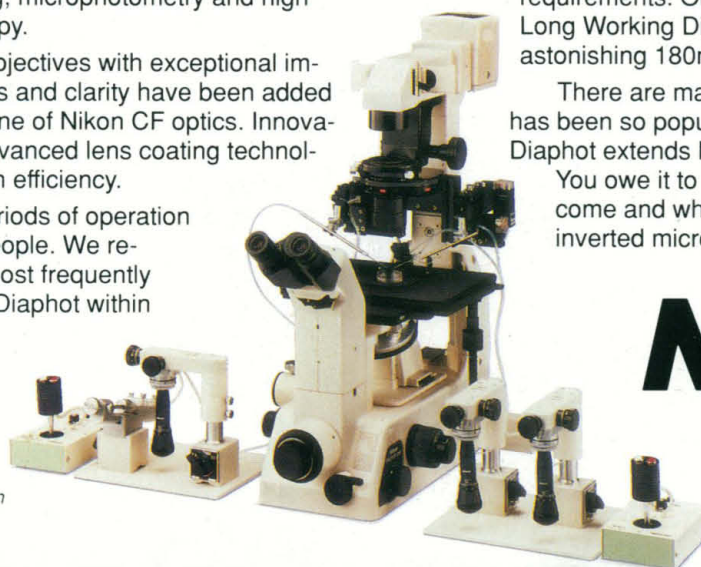


easy reach and by lowering the front and rear mounted stage for easier specimen handling and increased stability. In addition, the new Diaphot features an ergonomic height-adjustable eyepiece tube and 1 μ -sensitive coaxial focusing control with refocusing preset.

The new Diaphot also addresses users' needs for versatility and expandability. A new "systems condenser" lets you select working distance, DIC prisms and phase contrast to suit your individual microscopy requirements. Or choose the new SLWD (Super Long Working Distance) condenser for an astonishing 180mm of specimen clearance.

There are many reasons why the Diaphot has been so popular for so many years. The new Diaphot extends Nikon's tradition of excellence.

You owe it to yourself to see how far we've come and why the new Diaphot is the finest inverted microscope available today.



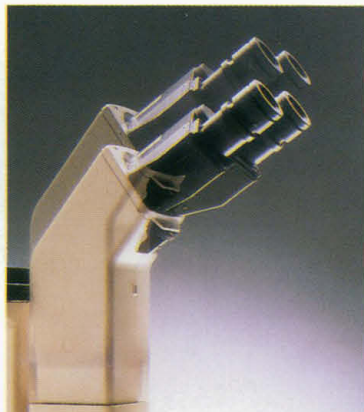
Designed with video microscopy, patch clamp and micromanipulation in mind. The DC power supply eliminates electronic noise.

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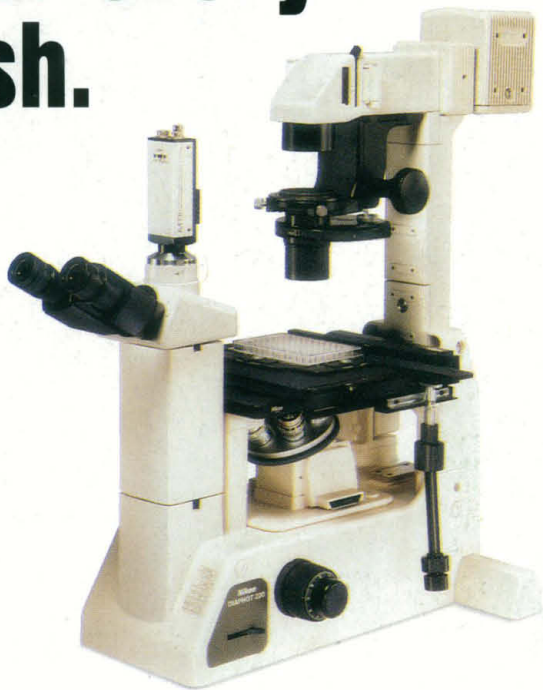
Multiple image ports include side, front, and optional trinocular eyepiece tube.



Height of BT1 eyepiece tube can be adjusted for user comfort.

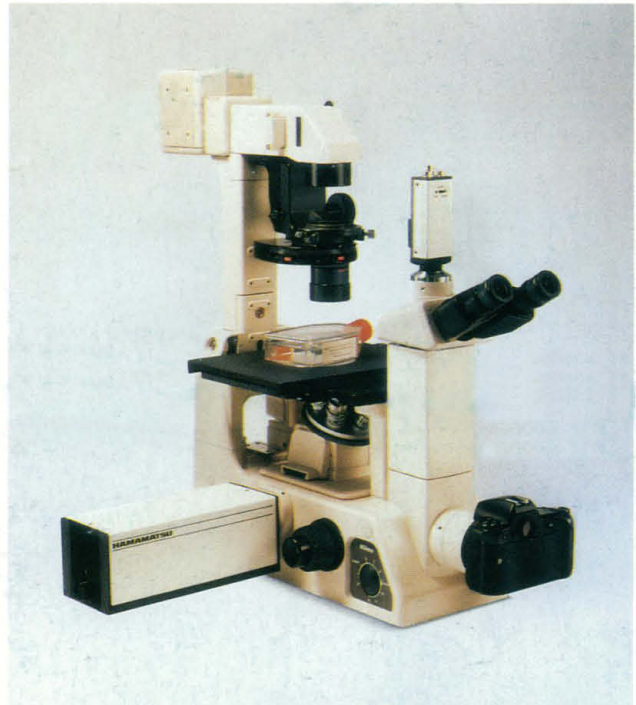
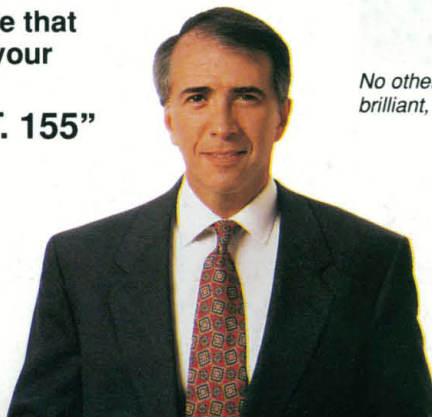


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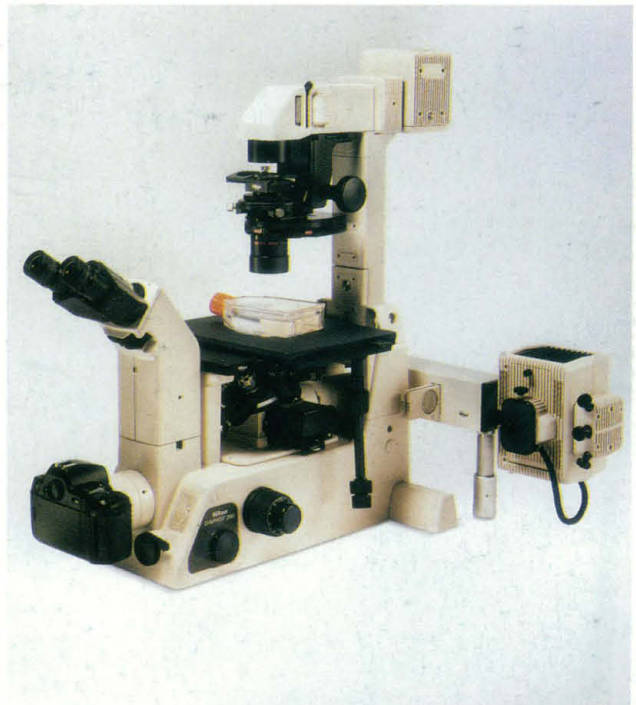


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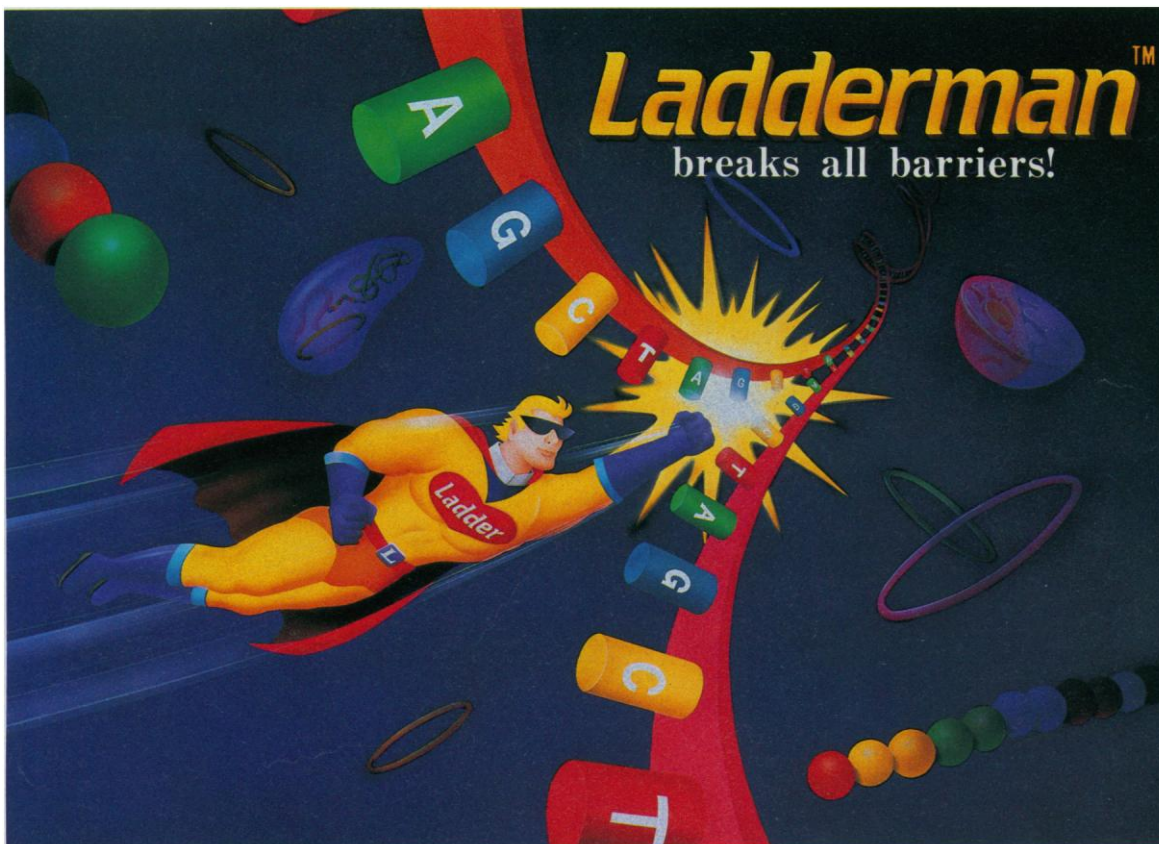


With multiple optical ports, Diaphot 300 can be configured for video, microphotometry and 35mm photomicrography simultaneously.



No other microscope can match the new Diaphot for exceptionally brilliant, high resolution epi-fluorescence over the broadest spectrum.

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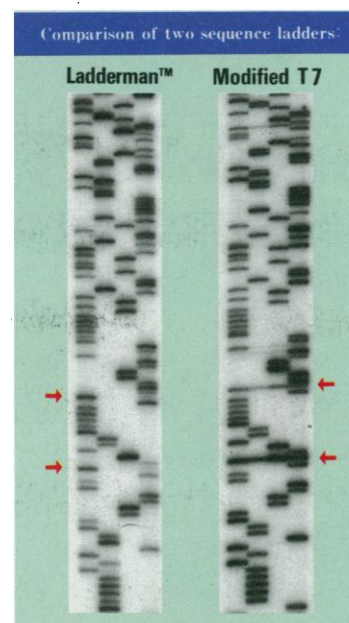


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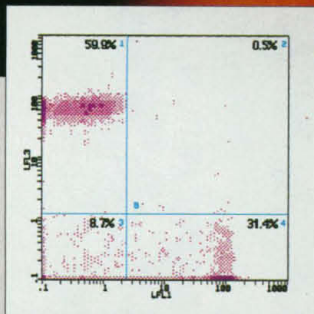
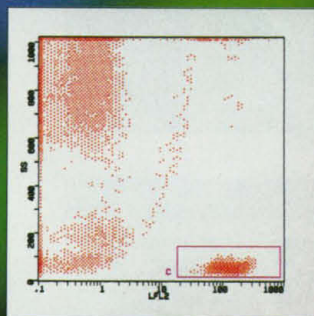
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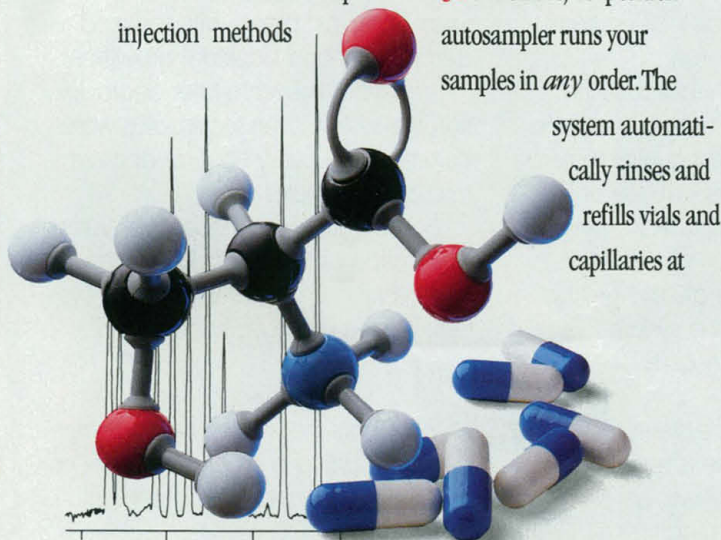
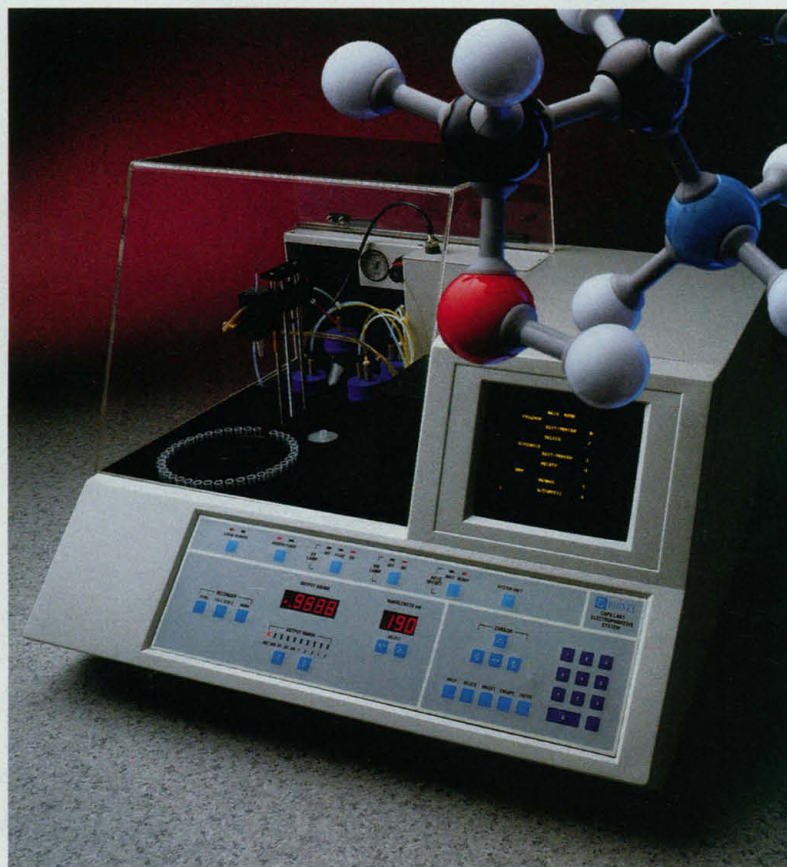
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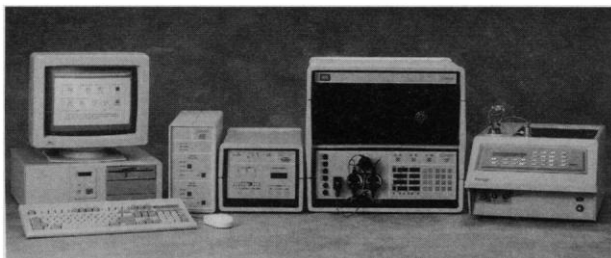
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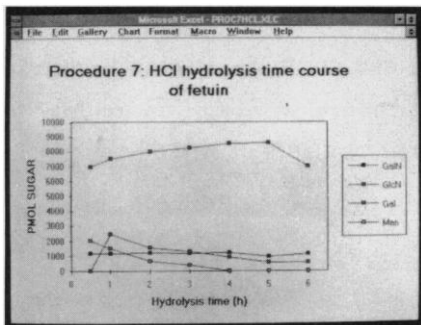
The system provides excellent resolution and sensitivity for detecting non-derivatized carbohydrates at picomole levels, and for resolving structures differing in only a single linkage position.

The GlycoStation can be fully automated, using software developed expressly for carbohydrate analysis, along with specialized

analytical procedures.

The system's self-teaching user's guide and intuitive, Windows™-driven software make it possible to teach yourself carbohydrate analysis in a few weeks. Fourteen steps take you from system setup and sample prep to monosaccharide/oligosaccharide analyses. Accurate and reliable data are virtually ensured.

The GlycoStation is expandable with a variety of detectors and columns, including the new CarboPac® MA1 column for separating reduced mono- and disaccharides (see next column).

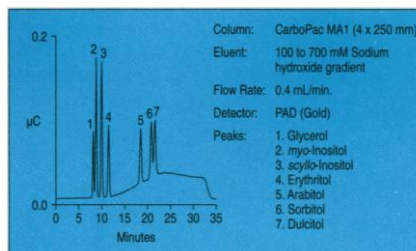


In addition to chromatography, the GlycoStation automatically produces easy-to-interpret graphs and charts.

A unique method for the separation of reduced carbohydrates

The CarboPac® MA1 is a high-performance anion-exchange column optimized for the sep-

reagents for the analysis of monosaccharides and polysaccharides.



The CarboPac MA1 is optimized for reduced mono- and disaccharides. These are typical sugar alcohols found in tissues and physiological fluids.

aration of reduced carbohydrates commonly found in physiological fluids, reduced glycoconjugate samples and food products.

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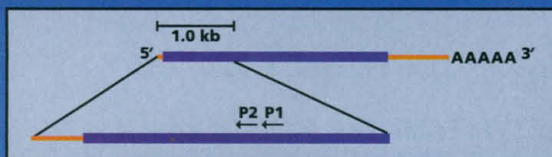
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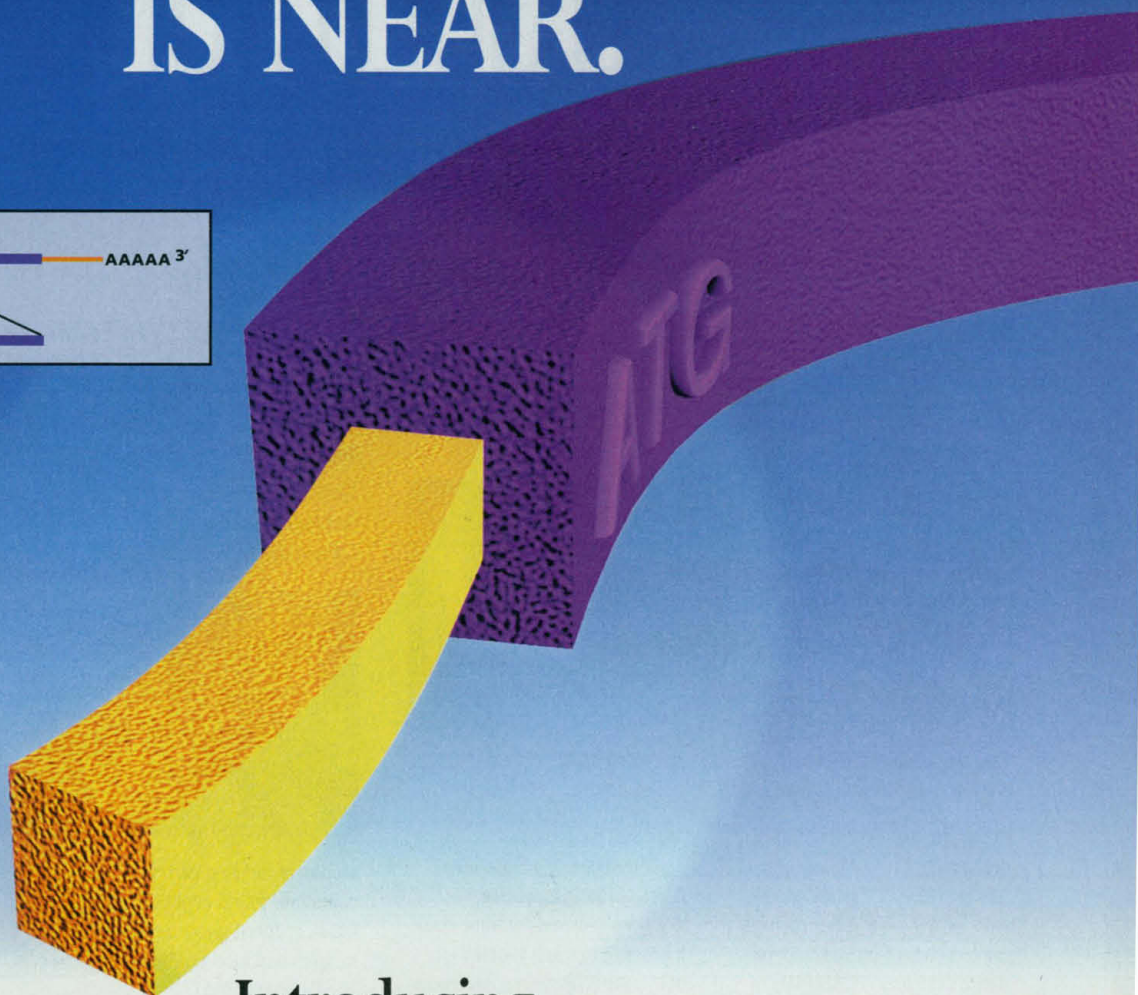
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Schematic diagram of the 4.7-kb human transferrin receptor gene transcript. The locations of the nested gene-specific primers, P1 and P2, are indicated.



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PCR amplification of the 5' ends of the human transferrin receptor (Lane 2: 0.87 kb. Lane 3: 0.45 kb) and tissue-type plasminogen activator genes (Lane 4: 0.40 kb).

AIDS

THE UNANSWERED QUESTIONS

If ever there were a pivotal year for AIDS research, 1993 could well be that year. It is precisely a decade since Luc Montagnier of the Pasteur Institute published the first report on the virus that is now known to cause AIDS, yet despite the high-powered arsenal of contemporary biology, there is nothing on the horizon remotely resembling a cure for AIDS. Nor is there anything like a workable vaccine.

In fact, as this special section on AIDS suggests, the more rapidly knowledge of the disease accumulates, the faster assumptions that seemed solid a year ago begin to crumble. And as solid ground disappears, scientific questions proliferate almost as rapidly as HIV itself. There are thousands of them—in vaccine work, drug research, pathogenesis studies, epidemiology, public health, and molecular biology.

To cut through that thicket of questions and focus on only the most important among them, *Science* has assembled a package combining news coverage with scientific papers. From the news team comes an overview of the crucial issues facing AIDS researchers as they were spelled out by 74 of the field's leaders in response to a survey devised by *Science*'s AIDS reporter Jon Cohen.

Offering greater depth on specific topics are six papers solicited by *Science* senior editor Barbara Jasny. In addition, the news team has previewed some of the most intriguing sessions that will be taking place at this year's International AIDS Conference in Berlin. We've also profiled four European AIDS researchers who our survey respondents told us were top contributors.

All told, we think this package offers the state of what is still very much an art: finding a way to stop HIV.

—John Benditt, Features Editor

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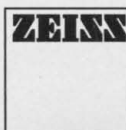
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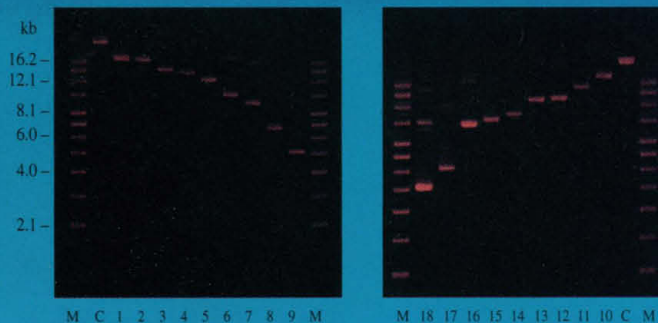
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Analysis of Deletion Subclones. A 17-kb human chromosomal DNA insert in pDELTA1 (pYA50) was subjected to the DELETION FACTORY System. Representative deletion subclones recovered from sucrose/tetracycline selection (*left panel*) and from streptomycin/kanamycin selection (*right panel*) were separated by electrophoresis. Lane numbers indicate individual clones; C = pYA50, and M = GIBCO BRL Supercoiled DNA Ladder.

I. Wang, G., Berg, C.M., Chen, J., Young, A.C., Blakesley, R.W., Lee, L.Y., and Berg, D.E. *Focus* 15, 47.

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H U M A N

MONOCLONAL

CLONE	BINDS TO AMINO ACID RESIDUES NO.	NEUTRALIZING
<i>Anti-gp41 anti-gp160 monoclonal antibodies</i>		
IAM 41-3D6	env 604-617	No
IAM 41-4D4	env 590-600	No
IAM 41-4B3	env 580-604	No
IAM 41-25C2	(a)	No
IAM 41-2F5	env 662-667	Yes
IAM 41-3H12	env 693-856	No
IAM 41-5F3	(a)	No
<i>Anti-gp120 anti-gp160 monoclonal antibodies</i>		
IAM 120-2G12	(b)	Yes
IAM 120-3D12	(b)	Yes
IAM 120-1B1	(b) (blocks CD4 binding)	Yes
IAM 120-2G6	(b)	No
<i>Anti-p24 monoclonal antibodies</i>		
IAM 24-37G12	structural epitope	No
IAM 24-3A6	p 24 122-149	No
IAM 24-1D7	unknown	No

(a) Binding site at residues 532-542, 556-566, or 683-693 of gp160; or structural epitope.
(b) Binds to carbohydrate moieties anchored between residues 001-201, 216-254, 285-302, 328-414, or 434-475 of gp160.
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$$\text{pk}_a = -\log k_a$$

$$E = mc^2$$

$$\frac{\partial^2 \psi}{\partial x^2} + \frac{\partial^2 \psi}{\partial y^2} + \frac{\partial^2 \psi}{\partial z^2} + \frac{8\pi^2 m}{h^2} (E - V) \psi = 0$$

$$E = 0.76 - \frac{0.059}{2} \log_{10} \frac{(1)(1)}{(10^{-3})^2}$$

$$a^n + b^n = c^n$$

$$X_i = kP_i$$

$$\int_0^{\infty} \frac{a dx}{a^2 + x^2} = \frac{\pi}{2}, \text{ if } a > 0; 0, \text{ if } a = 0; -\frac{\pi}{2}, \text{ if } a < 0$$

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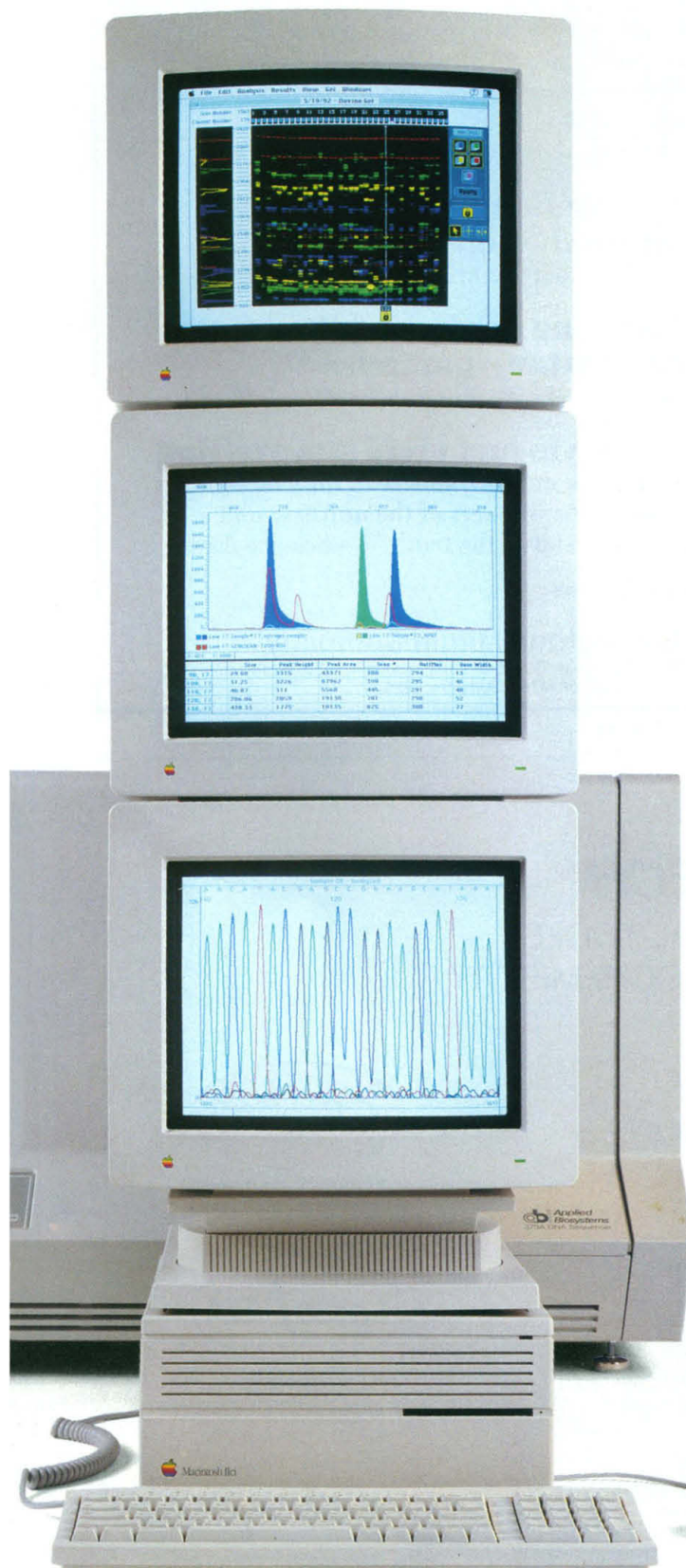
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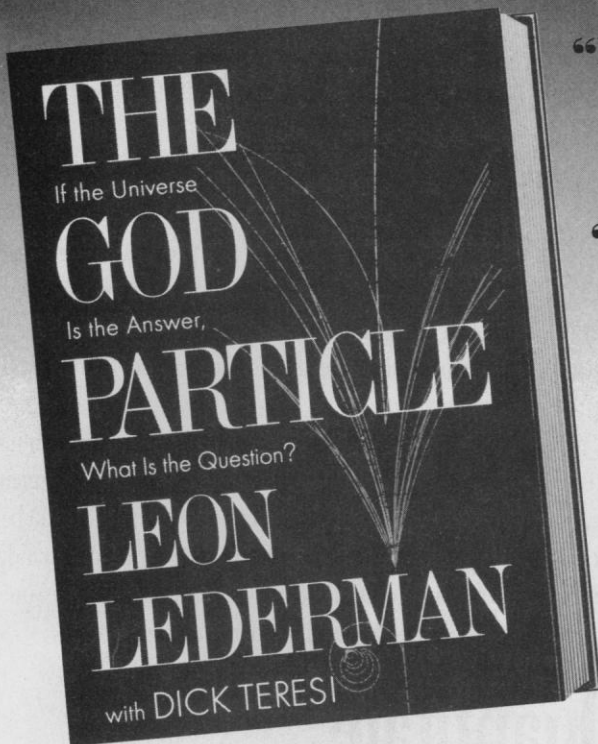
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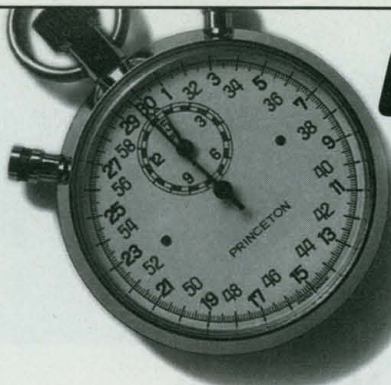
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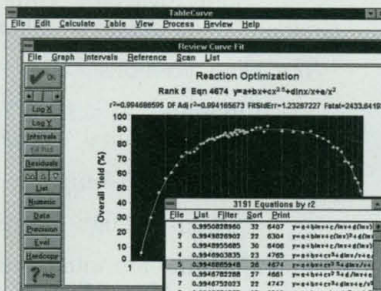


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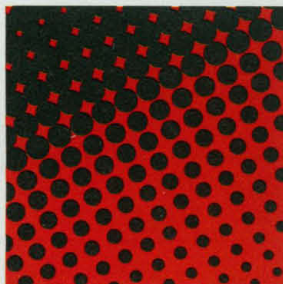
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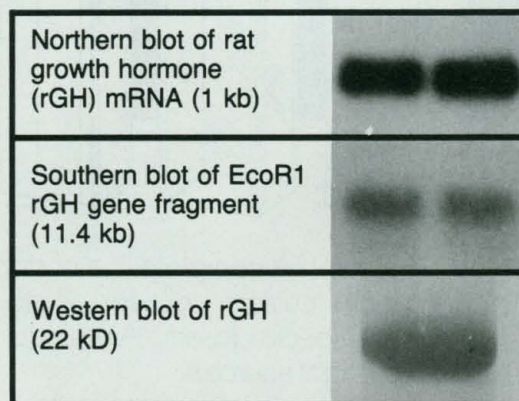
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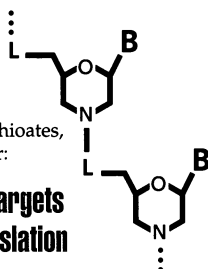
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Technical Report #2, ANTISENSE R&D, 3(1), following page 153.
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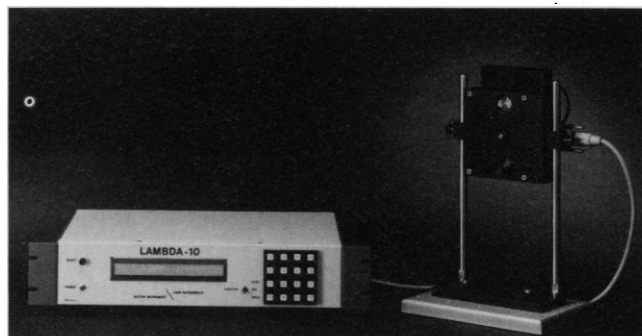
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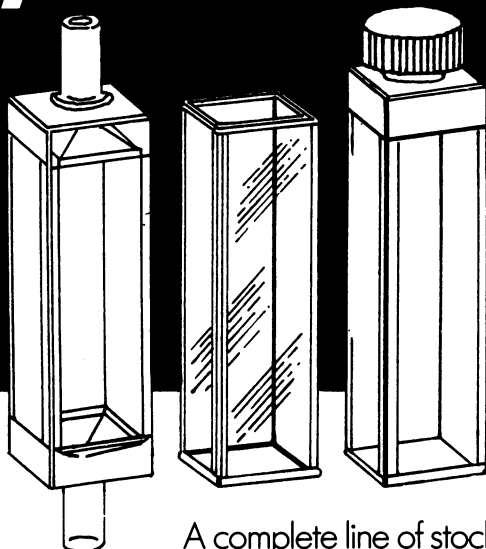
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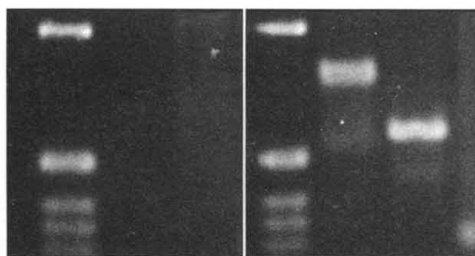
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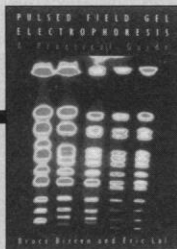
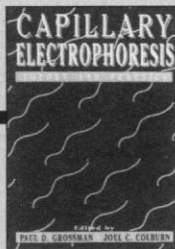
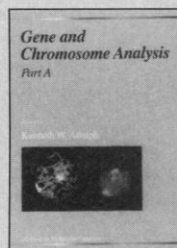
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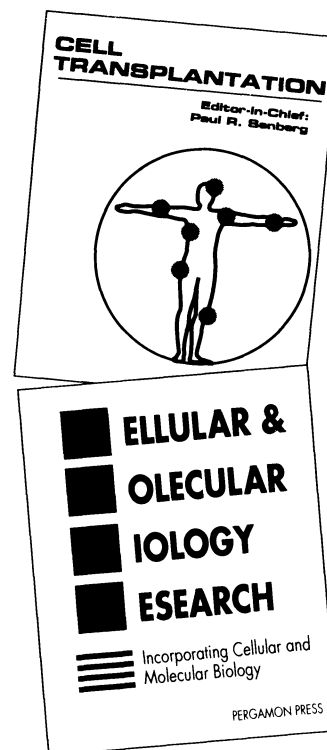
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