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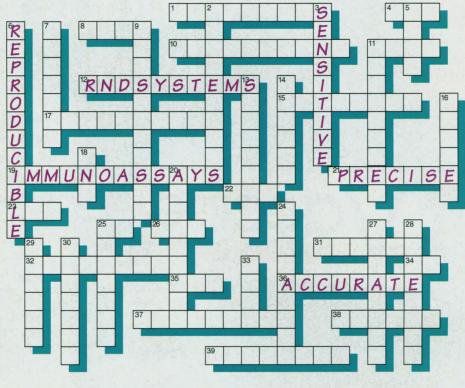
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ACROSS

- 1 1 of 4 specific families of CAMS 4 Abbrev. for biological response modifiers
- 8 Abbrev, for human antimouse antibody
- 10 R&D Systems' line of immunoassays for the measurement of cytokines
- 11 Basic structural and functional unit of all organisms 12 R&D Systems URL -
- http://www. .com 15 Glycoprotein substance produced by B lymphoid
- cells in response to stimulation with an immunogen 17 Proteolytic complex with a
- mass of about 1.000.000 19 Tests that measure antigen
- or antibody

within close specified limits 22 Abbrev. for cell-selective protein that promotes adhesion of cells

21 Successively reproducible

- 23 Abbrev. for gram negative endotoxin
- 25 Site on DNA lacking either purines or pyrimidines
- 26 Abbrev. for a detection enzyme 31 Obtained from an affinity
- column 32 "The Ice Man Cometh" for
- cells 34 Abbrev, for cluster of
- differentiation
- 35 Abbrev. for superoxide dismutase
- 36 In exact conformity to fact 37 Cross-linked molecules

38 Discrete portion of a molecule 39 Region of an antigen that combines with an MHC class II molecule

DOWN

- 2 Cell migration from the interior of small vessels into tissue spaces
- 3 A high level of discrimination; with 21 and 36 across, descriptive of Quantikine kits
- 5 Evidence ignored by
- Simpson jury 6 Capable of replication
- 7 An antigenic determinant
- 9 An end-to-end union or join-
- ing together of blood vessels 11 What R&D Systems is your source for
- 13 Substance acted upon by enzymes

- 14 The weight of a single hydrogen atom or a member of an outlaw gang
- 16 Winner who shared 1984 Nobel prize with Milstein
- Dawkins' selfish entity 20 A molecule that serves as a
- homing device 24 Having a single binding site
- 25 A substance with which an antibody molecule or T cell
- receptor may bind
- 27 A defining example 28 Complementary binding site
- 29 "M" of ECM
 - 30 A specimen of known content used together with an unknown in order that the two may be compared
 - 33 Abbrev. for an anticoagulant that binds divalent cations

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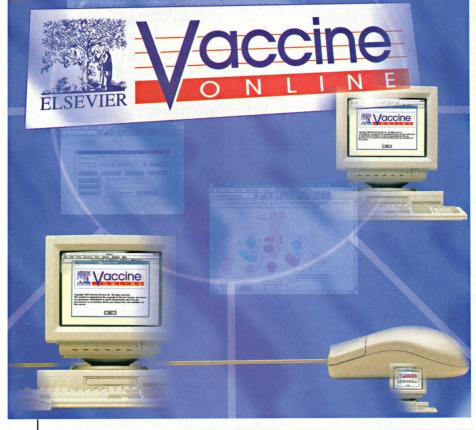
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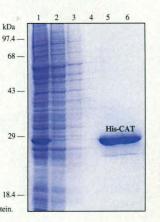
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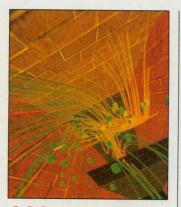
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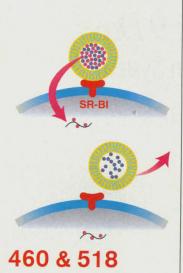
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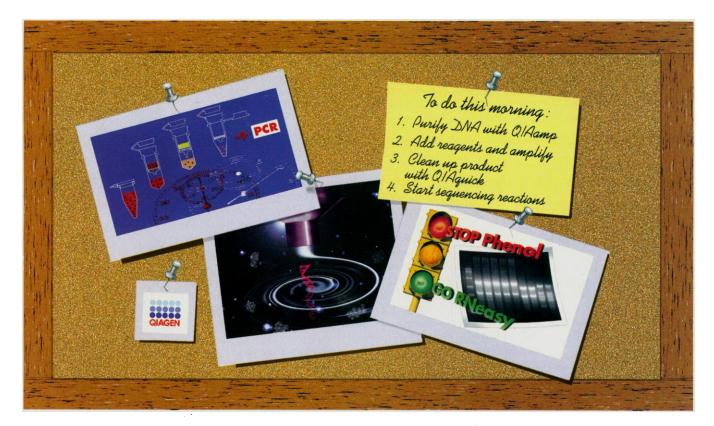
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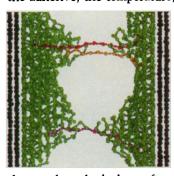
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This Week in Science

edited by PHIL SZUROMI

Torn apart

What happens at the molecular level when two surfaces, held together by an adhesive film, are pulled apart? The answer depends on the type and state of the adhesive, the temperature,



the speed at which the surfaces are pulled apart, and the strength of bonding between the surfaces and the adhesive. Entangled adhesive molecules resist rupture, leading to a buildup of energy that is dissipated differently, depending on the state of the polymer. Baljon and Robbins (p. 482) performed molecular dynamics calculations to follow individual polymer changes during this process, and have identified three mechanisms of energy dissipation during rupture that have parallels to macroscopic phenomena observed in commercial adhesives.

Random magnetism

Materials that contain chain structures can approximate onedimensional systems, whose properties, such as the magnetic coupling between electrons, can be guite different from that seen in three-dimensional systems. Nguyen et al. (p. 489) describe a compound, Sr₃CuPt_{0.5}Ir_{0.5}O₆, that is a solid solution of ferromagnetic Sr₃CuPtO₆ and antiferromagnetic Sr₃CuIrO₆. This intermediate compound forms a random quantum spin chain paramagnet—the coupling of spins along the chain through

Linking atmospheric oxygen and phosphorus

Oxygen levels in Earth's atmosphere evidently have been maintained at levels that allow life to persist for billions of years. The controls that essentially buffer the system have been uncertain. Van Cappellen and Ingall (p. 493; see the Perspective by Kump and Mackenzie, p. 459) suggest that the key is the phosphorus cycle; phosphorus is a key nutrient driving oceanic production and respiration, is a minor component of the atmosphere, and its burial is extremely sensitive to the oxidation state of the oceans. Analysis of phosphorus burial in the past shows that when waters are low in oxygen, less phosphorus is buried.

exchange interactions can be randomly ferromagnetic or antiferromagnetic.

Enzyme clocks for kingdom divergence

Most mutations in proteins occur as random events over time, so, in principle, one can estimate when divergence occurred from a common ancestor by analyzing sequences of the same protein from different organisms. In practice, proteins mutate at different rates, and functionally similar proteins may in fact have different origins. Doolittle et al. (p. 470; see the news story by Morrell, p. 448) analyzed divergence times for 57 different enzymes from organisms from all of the major biological groups. The analysis of this large data set indicates that eukaryotes and eubacteria had a common ancestor about 2 billion years ago, or about twice as long ago as the plant-animal divergence.

-

HIV dynamics

During the first few weeks of HIV infection, the concentration of virus in plasma rises but then drops off. Phillips (p. 497) has modeled the course of early HIV infection and suggests that the drop-off is not an HIVspecific immune response but is actually due to population dynamics in HIV replication. The supply of uninfected cells is quickly depleted, thus decreasing virus production.

Ancient enzymes

Although the DNA sequence of an ancestral gene may be predictable, the enzymatic function of the protein product may not be. Chandrasekharan et al. (p. 502) used extant chymase gene sequences to predict the ancestral form of the enzyme, and then expressed the enzyme and characterized its kinetics for converting angiotensin I to angiotensin II. The ancestral form is very specific and efficient for this reaction, and some modern chymases actually have less substrate specificity than this ancestral form.

Hepatitis G virus

In some hepatitis cases, the causative agent is unknown, which suggests that some hepatitis viruses are still unidentified. Linnen *et al.* (p. 505) identified an RNA virus, hepatitis G virus (HGV), by analyzing immunoreactive complementary DNA clones from a patient who was also infected with hepatitis C virus. Although the pathogenesis of HGV remains to be determined, it is associated with both acute and chronic hepatitis and is present in the blooddonor population.

-

Overactive enzyme

About 20 to 25% of patients with familial amyotrophic lateral sclerosis (FALS) have mutations in the gene encoding copper-zinc superoxide dismutase (CuZnSOD). Wiedau-Pazos et al. (p. 515; see the news story by Marx, p. 446) show that the mutant CuZnSOD is more active than wild type in catalyzing oxidation of a model substrate, and that this enhanced activity can be inhibited by copper chelators. In a cell culture model of FALS, copper chelators inhibited the cell death induced by mutant CuZnSOD. Aberrant oxidative reactions catalyzed by mutant CuZnSOD may underlie the neuropathologic changes in FALS, and copper chelators may have therapeutic value for some FALS patients.

Apoptosis links

Programmed cell death (PCD), which typically occurs through apoptosis, is a normal function in multicellular organisms. Disregulated PCD, however, can cause numerous disorders, including neurodegenerative diseases and cancer. Several studies indicate that Ca²⁺ plays a role in apoptosis; Vito et al. (p. 521) developed a selection system which allowed them to identify two genes, ALG-2 and ALG-3, that mediate apoptosis. The ALG-2 gene encodes a Ca2+-binding protein required for several apoptotic pathways, and the ALG-3 gene is homologous to the Alzheimer's disease gene STM2.

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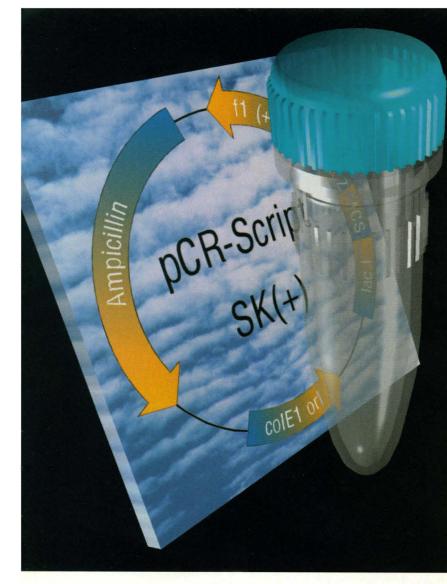
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Bauer, J., et al. (1992) Stratagies 5:2-64
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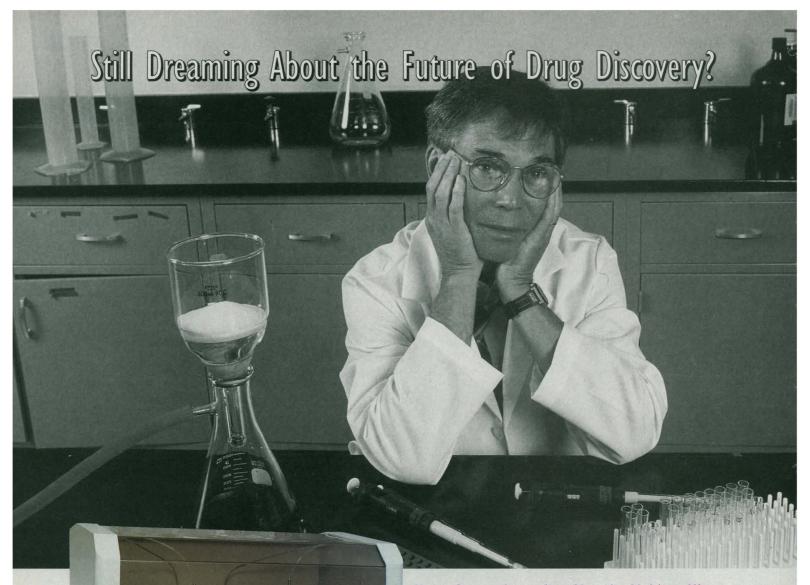
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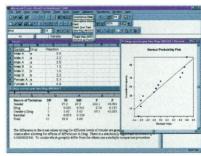
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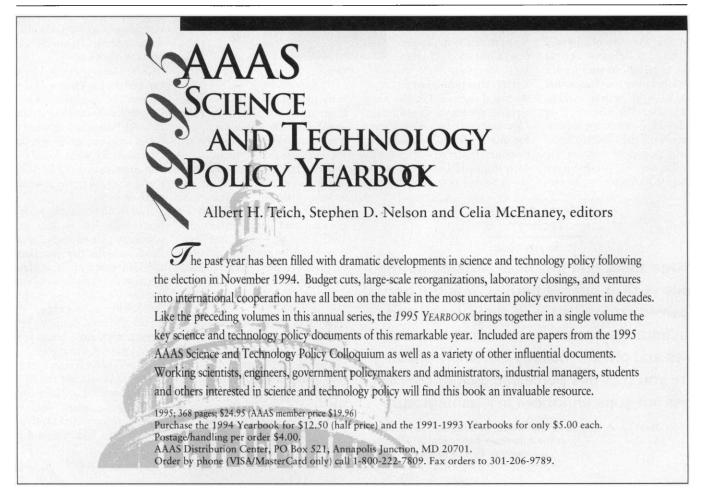
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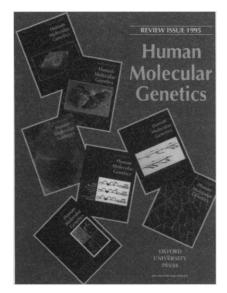
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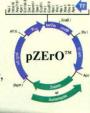
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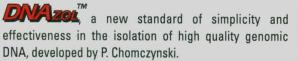
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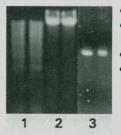
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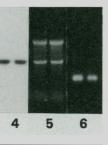




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