Science

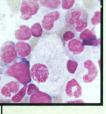
The **I**

he Human Genome Project is expeof DNA representing the functional history of the human species. How sequence is thought to specify the portion genes that encode proteins. Thus an impogenomics is to identify and localize these transcript mapping. When genes are expr converted into messenger RNA transcrip form of complementary DNAs (cDNAs). A human genes had been sampled as of 15

•1

600





GBA In Gaucher disease, the defective enzyme is unable to metabolize glucocerebrosides which accumulate in characteristic, distended phagocytic cells.

5

AD4 Brain scans of a healthy elderly person and a patient with Alzheimer's disease.

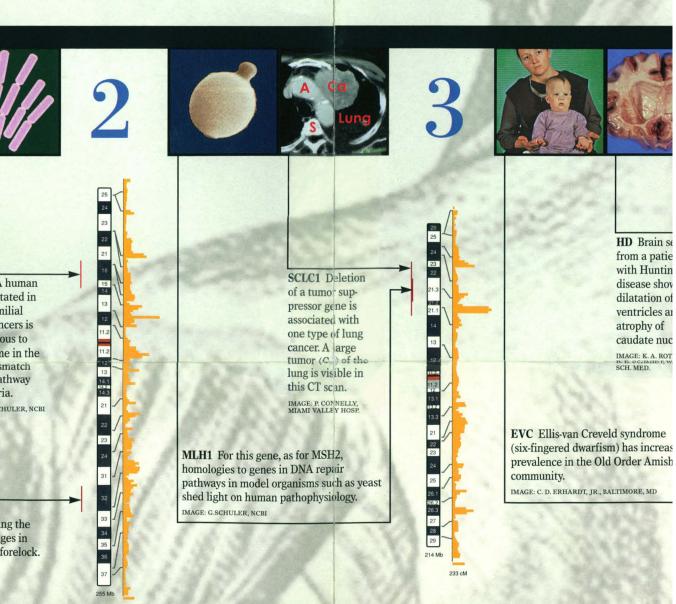
IMAGE: K. JOHNSON, BRIGHAM AND WOMEN'S HOSP.

MSH2 A human gene mutated in some familial colon cancers is homologous to an enzyme in the DNA mismatch repair pathway in bacteria.

PAX3 Portion of a pedigree of Waardenburg syndrome, indicating the occurrence of deafness and changes in pigmentation, including a white forelock. IMAGE: V. MCKUSICK, JOHNS HOPKINS HOSP.

Human Tran

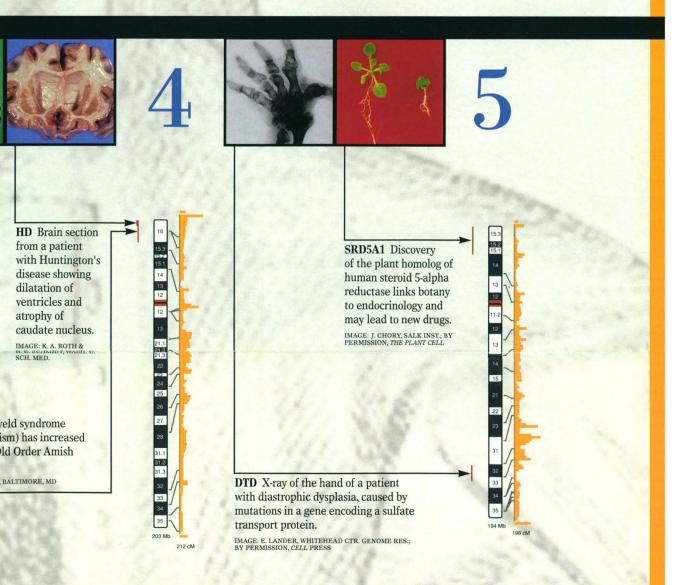
oject is expected to produce a sequence the functional blueprint and evolutionary pecies. However, only about 3% of this of the portions of our 50,000 to 100,000 hus an important part of basic and applied calize these genes in a process known as thes are expressed, their sequences are first A transcripts, which can be isolated in the s (cDNAs). Approximately half of all ed as of 15 June, 1996. A snall portion of each cDNA sequence is all that is need develop unique gene markers, known as sequence-tagged site which can be detected in chromosomal DNA by assays based polymerase chain reaction (PCR). To construct this transcript cDNA sequences from a master catalog of human genes were uted tomapping laboratories in North America, Europe, and These cDNAs were converted to STSs and their physical loca chromosomes determined on one of two radiation hybrid (RH or a yeast artificial chromosome (YAC) library containing hu genomic DNA. This mapping data was integrated relative to t



nscript Map

that is needed to e-tagged sites or STSs, ssays based on the s transcript map, genes were distriburope, and Japan. hysical locations on hybrid (RH) panels itaining human relative to the human genetic map and then cross-referenced to cytogenetic band maps of the chromosomes. (Further details are available in the accompanying article in the 25 October issue of *SCIENCE* and on the World Wide Web at http://www.ncbi.nlm.nih.gov/SCIENCE96/).

The histograms (
) reflect the distributions and densities of genes along the chromosomes. Because the individual genes (>16,000) are too numerous to represent, images have been chosen to illustrate the myriad aspects of human biology, pathology, and relationships with other organisms that can be revealed by analysis of genes and their protein products.



GBA In Gaucher disease, the defective enzyme is unable to metabolize glucocerebrosides which accumulate in characteristic, distended phagocytic cells.

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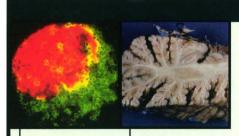


MSH2 A human gene mutated in some familial colon cancers is homologous to an enzyme in the DNA mismatch repair pathway in bacteria.

PAX3 Portion of a pedigree of Waardenburg syndrome, indicating the occurrence of deafness and changes in pigmentation, including a white forelock. IMAGE: V. MCKUSICK, JOHNS HOPKINS HOSP.







SCA1

Degeneration of the cerebellum leads to loss of muscle coordination in patients with spinocerebellar atrophy.

IMAGE: K. A. ROTH AND R. E. SCHMIDT, WASH. U. SCH. MED.

IDDM1 Juvenile onset diabetes, in which the body's own T lymphocytes infiltrate and destroy insulin-producing, pancreatic islet cells, is associated with the major histocompatibility complex. MAGE: A. COOKE & J. TODD, WELLCOME TRUST CTR. HUMAN GENET.



22



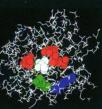
CFTR The gene encoding a chloride ion channel is defective in patients with cystic fibrosis.

IMAGE: Q. AL-AWQATI, COLUMBIA U., K. SUTLIFF, ADAPTED FROM SCIENCE

OBS The obese (Ob) mutation in the mouse provides a useful model system for studying human obesity.

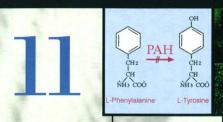
IMAGE: J. FRIEDMAN, ROCKEFELLER U.; REPRINTED FROM SCIENC







201 cM



human tated in nilial ncers is ous to ne in the smatch athway ria.

11.2

13

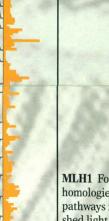
14.

14.2 14.3

21

277 cM

ng the ges in forelock.



SCIECT Determined of a tumo: suppressor gene is associated with one type of lung cancer. A arge tumor (C) of the lung is visible in this CT scan. **IMAGE: P. CONNELLY, MAME VALLEY HOSP.**

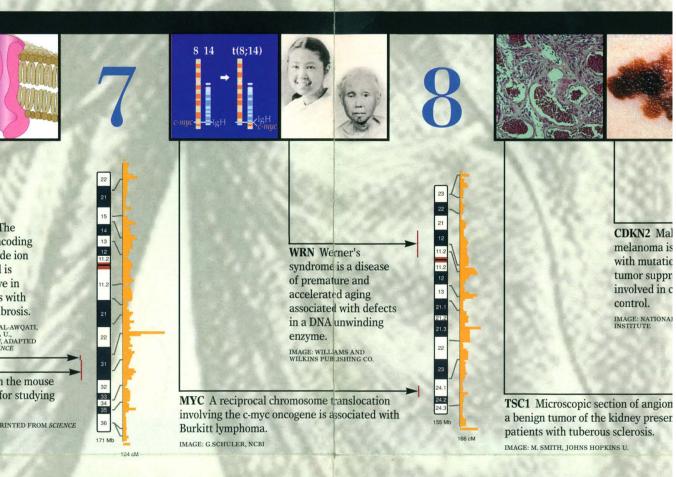
MLH1 For this gene, as for MSH2, homologies to genes in DNA repair pathways in model organisms such as yeast shed light on human pathophysiology. MAGE: G.SCHULER, NCBI

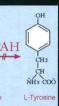


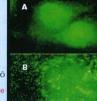
disease show dilatation of ventricles an atrophy of caudate nuc MAGE: K. A. ROT N. B. OCIMPIP, N.

EVC Ellis-van Creveld syndrome (six-fingered dwarfism) has increas prevalence in the Old Order Amish community.

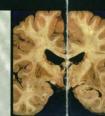
IMAGE: C. D. ERHARDT, JR., BALTIMORE, MD

















disease showing dilatation of ventricles and atrophy of caudate nucleus.

veld syndrome ism) has increased Ild Order Amish

BALTIMORE, MD

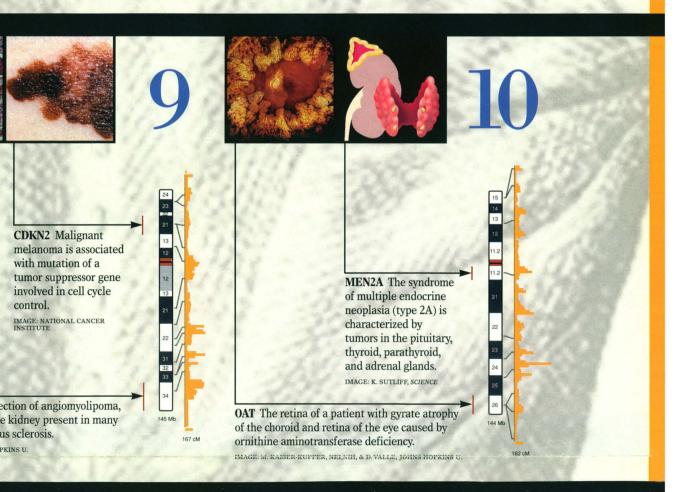


human steroid 5-alpha reductase links botany to endocrinology and may lead to new drugs. IMAGE: J. CHORY, SALK INST.; BY PERMISSION, THE PLANT CELL

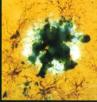
198 cM

DTD X-ray of the hand of a patient with diastrophic dysplasia, caused by mutations in a gene encoding a sulfate transport protein.

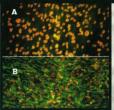
IMAGE: E. LANDER, WHITEHEAD CTR. GENOME RES.; BY PERMISSION, CELL PRESS











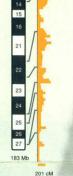




atrophy.

IMAGE: K. A. ROTH AND R. E. SCHMIDT, WASH. U. SCH. MED.

IDDM1 Juvenile onset diabetes, in which the body's own T lymphocytes infiltrate and destroy insulin-producing, pancreatic islet cells, is associated with the major histocompatibility complex. MAGE: A. COOKE & J. TODD, WELLCOME TRUST CTR. HUMAN GENET.

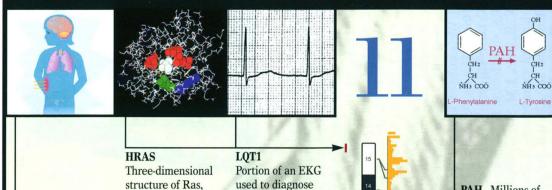


patients with cystic fibrosis.

IMAGE: Q. AL-AWQATI, COLUMBIA U., K. SUTLIFF, ADAPTED FROM SCIENCE

OBS The obese (Ob) mutation in the mouse provides a useful model system for studying human obesity.

IMAGE: J. FRIEDMAN, ROCKEFELLER U.; REPRINTED FROM SCIENC



structure of Ras, the product of an oncogene that is mutated in many human cancers. IMAGE: M. BOGUSKI, NCBI LQT1 Portion of an EKG used to diagnose long-QT syndrome, an inherited cardiac arrythmia associated with mutations in an ion channel protein.

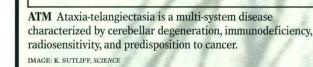


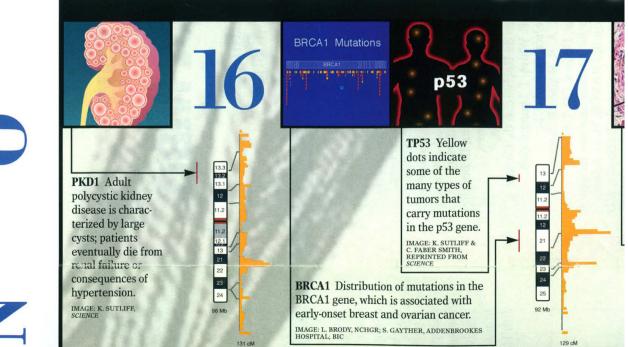
156 cM

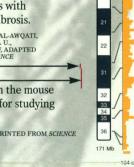
PAH Millions of newborns per year are screened for deficiency of phenylalanine hydroxylase to identify those susceptible to phenylketonuria.



T







R

accelerated aging associated with defects in a DNA unwinding enzyme. IMAGE: WILLIAMS AND WILKINS PUB JSHING CO.

MYC A reciprocal chromosome translocation involving the c-myc oncogene is associated with Burkitt lymphoma. IMAGE: G.SCHULER, NCBI

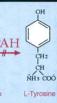


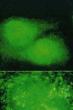
166 cM

control. IMAGE: NATIONAL

TSC1 Microscopic section of angion a benign tumor of the kidney preser patients with tuberous sclerosis. IMAGE: M. SMITH, JOHNS HOPKINS U.

Lesion









BRCA2 Mammogram showing breast cancer. IMAGE: P. CONNELLY, MIAMI VALLEY HOSP

ATP7B Ir Wilson's disease, toxic levels of copper accumula e and damage many tissues and organs, including the basal ganglia of the brain. IMAGE: K. A. FOTH & R. E. SCHMIDT, WASH. U. SCH. MED.

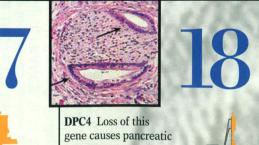
fillions of ns per year ened for icy of alanine lase to those ible to cetonuria. CHULER, NCBI

129 cM

Zellweger syndrome fibroblasts (A) but can be reconstituted by transfection with the PXR1 gene (B). IMAGE: N. BRAVERMAN, G. DODT, S. J. GOULD, & D. VALLE, JOHNS HOPKINS U.

PXR1 Peroxisomes

are not detected in



cancers to grow aggressively, as seen by tumor cells invading a nerve bundle. IMAGE: R. H. HRUBAN, JOHNS HOPKINS HOSP



85 Mb

124 cM

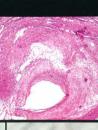
143 M

169 cM



117

IMAGE: M. A DE LA FLOR, REPRINTET FROM SCIENCE



APOE Atherosclerotic coronary artery disease is associated with the gene encoding apolipoprotein E, a ligand for the LDL receptor. IMAGE: M. BOGUSKI, NCBI







IMAGE: J. GOLDSTEIN & M.S. BROWN, U. TEXAS SW MED. CTR.; BY PERMISSION, EXCERPTA MEDICA INC.



13.3

control. IMAGE: NATIONAL CANCER INSTITUTE

ection of angiomyolipoma, e kidney present in many us sclerosis. PKINS U.



167 cM

of multiple endocrine neoplasia (type 2A) is characterized by tumors in the pituitary, thyroid, parathyroid, and adrenal glands. IMAGE: K. SUTLIFF, SCIENCE

OAT The retina of a patient with gyrate atrophy of the choroid and retina of the eye caused by ornithine aminotransferase deficiency.

IMAGE. M. KAISER-KUPFER, NEI,NIH; & D. VALLE, JOHNS HOFKINS U.









118 cM

13.

12

13.3

13.4

67 Mb

110 cM

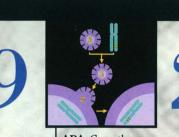
AD3 Neuritic plaques and neurofibrillary tangles are the major microscopic abnormalities in the brains of patients with Alzheimer's disease.



FBN1 Extracellular fibrillin-1 microfibrils (stained green) are absent in Marfan syndrome (A) compared to a control sample (B).

IMAGE: H. DIETZ, JOHNS HOPKINS U.; BY PERMISSION, ELSEVIER SCIENCE IRELAND LTD.





ADA Gene therapy has been attempted to treat severe combined immunodeficiency caused by a missing enzyme, adenosine deaminase.

IMAGE: NATIONAL CANCER INSTITUTE



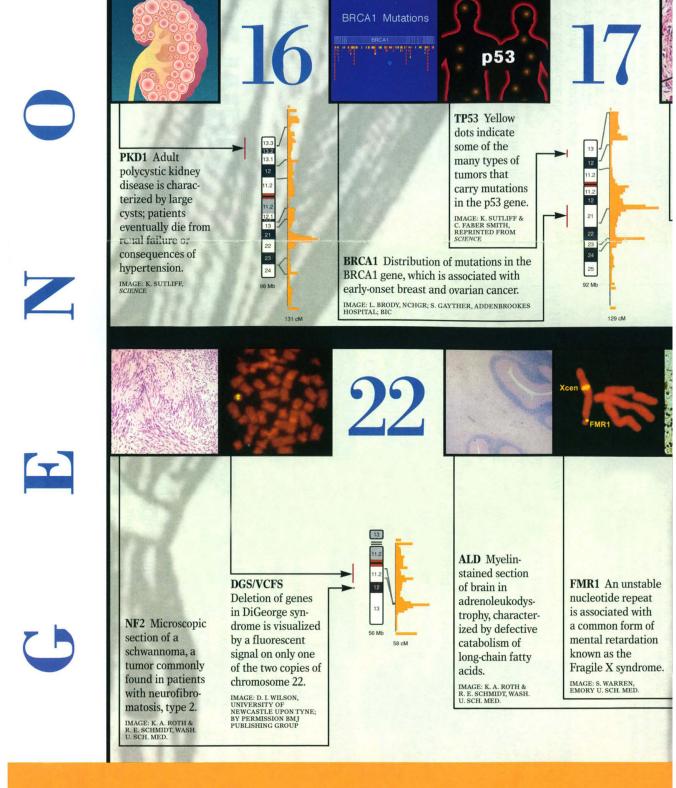




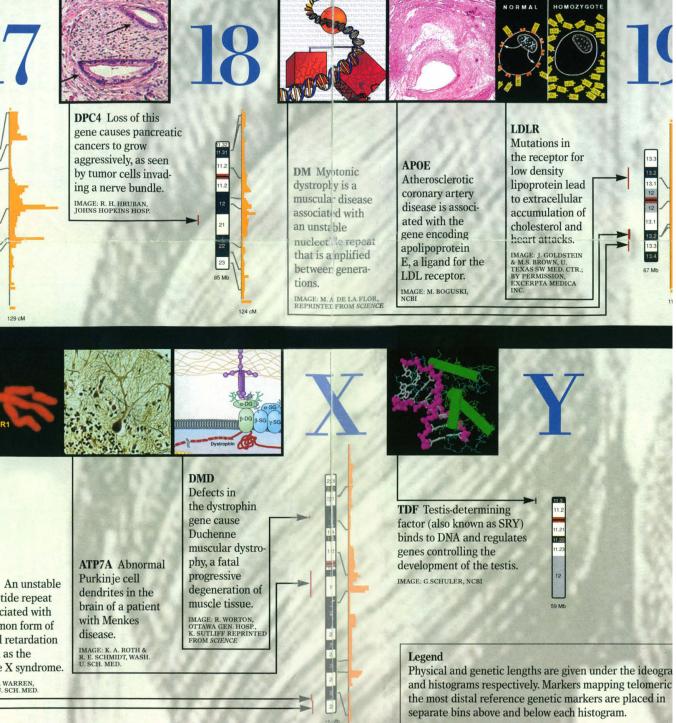
SOD1 Amyotrophic lateral sclerosis, known as Lou Gehrig's disease, is caused in some cases by a deficiency in the enzyme superoxide dismutase.



60 cM



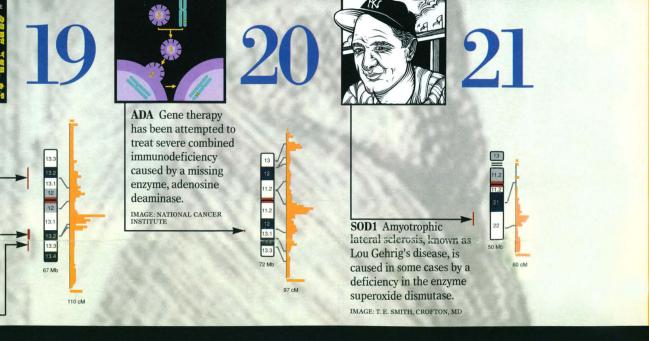
Gene Specific STSs and Matching cDNA Clones for Each Mapped Transcript • BAC and YAC Librari Radiation Hybrid Mapping Panels • I.M.A.G.E. Consortium (LLNL) cDNA Cl



- 10 genes per cM

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gram.

Credits: Science Coordinator: Barbara R. Jasny Authors: G. D. Schuler, M. S. Boguski, National Center for Biotechnology Information, Bethesda, MD, USA T. J. Hudson, L. Hui, J. Ma, A. B.Castle, X. Wu, J. Silva, H. C. Nusbaum, B. B. Birren, D. K. Slonim, S. Rozen, L. D. Stein, D. Page, E. S. Lander, Center for Genome Research, Whitehead Institute, Cambridge, MA, USA; E. A. Stewart, A. Aggarwal, E. Bajorek, S. Brady, A. Chu, N. Fang, D. Hadley, M. Harris, S. Hussain, A. Maratukulam, S. Perkins, M. Piercy, F. Qin, T. Reif, C. Sanders, X. She, W.-L. Sun, P. Tabar, S. Voyticky, C. Mader, K. B. McKusick, J.-B. Fan, S. Cowles, J. Quackenbush, D. Vollrath, R. M. Myers, D. R. Cox, Stanford Human Genome Center, Stanford, CA USA; A. Butler, C. Clee, T. Dibling, C. East, C. Edwards, C. Garrett, L. Green, P. Harrison, A. Hicks, E. Holloway, S. Ranby, A. MacGilvery, A. Mungall, A. Peck, T. Wilmer, C. Soderlund, K. Rice, I. Dunham, D. Bentley, P. Deloukas, The Sanger Center, Cambridge, UK; G.Gyapay, N. Chiannilkulchai, C. Fizames, S. Bentolila, S. Duprat, N. Vega-Czarny, D. Muselet, N. Drouot, J. Morissette, J. Beckmann, J. Weissenbach, Généthon, CNRS, Evry, FRANCE; J. Morissette, Centre de Recherche du Centre Hospitalier de l'Université Laval, Qúebec, CANADA; M. R. James, R. E. White, T. Thangarajah, C. Louis-Dit-Sully, P. J. R. Day, Wellcome Trust Centre for Human Genetics, Oxford, UK; P.N. Goodfellow, K. Schmitt, Cambridge University, Cambridge, UK; N. A. R. Walter, R. Berry, K. R. Iorio, J. M. Sikela, University of Colorado Health Sciences Center, Denver, CO, USA; M. H. Polymeropoulos, R. Torres, S. E. Ide, A. Dehejia, National Center for Human Genome Research, Bethesda, MD, USA; R. Houlgatte, C. Auffray, Genexpress, CNRS, Villejuif, FRANCE; M. D. Adams, C. Phillips, R. Brandon, M. Sandusky, J. C. Venter, The Institute for Genomic Research, Rockville, MD, USA; N. Seki, T. Nagase, K. Ishikawa, N. Nomura, Kazusa DNA Research Institute, Kisarazu, JAPAN; P. Rodriguez-Tome, The EMBL Outstation, The European Bioinformatics Institute, Cambridge, UK; T. C. Matise, The Rockefeller University, New York, NY, USA; W. Y. Lee, K. A. Swanson, J. R. Hudson Jr., Research Genetics, Huntsville, AL, USA; Reviewers: R. Kucherlapati, Albert Einstein College of Medicine, New York, NY, USA; V. McKusick, Johns Hopkins Hospital, Baltimore, MD, USA. Art Director, Amy Decker Henry; Design, Tracy Keaton-Drew; Production, Linda Hannigan.

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