

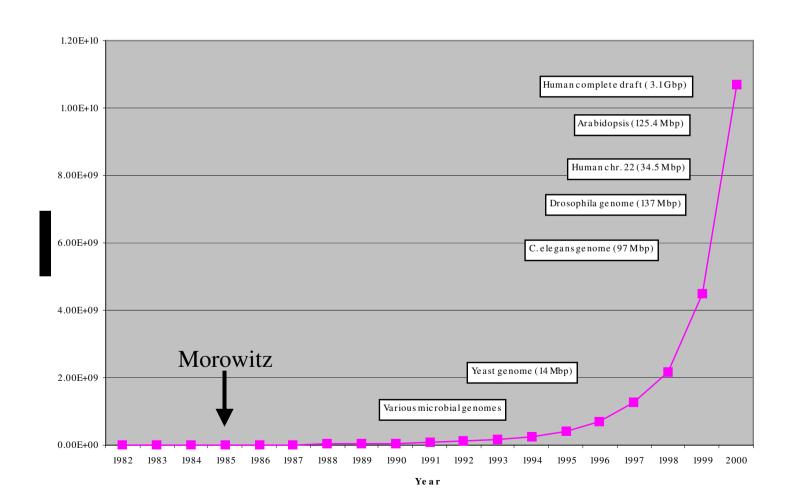
Wisdom world-wide?

Portals to bio-medical information viewed from a user perspective

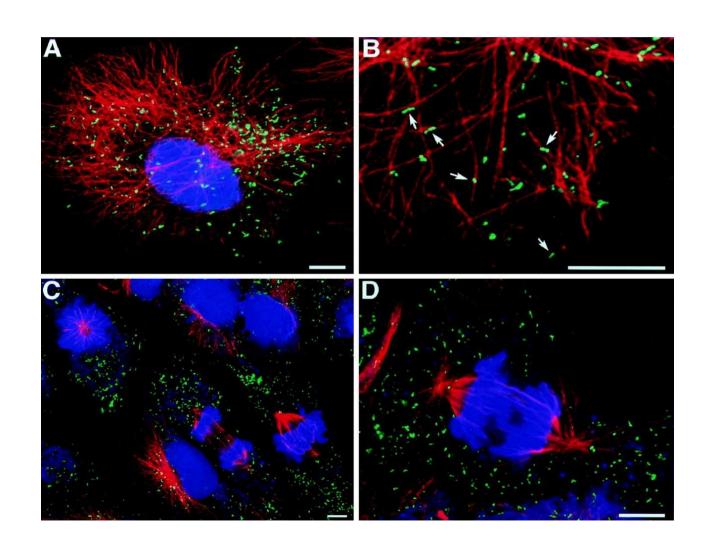
Les Grivell European Molecular Biology Organization "Biological research has reached a point where new generalizations and higher order biological laws are being approached, but may be obscured by the simple mass of data"

> Harold Morowitz, 1985 Report to the U.S. National Academy of Sciences

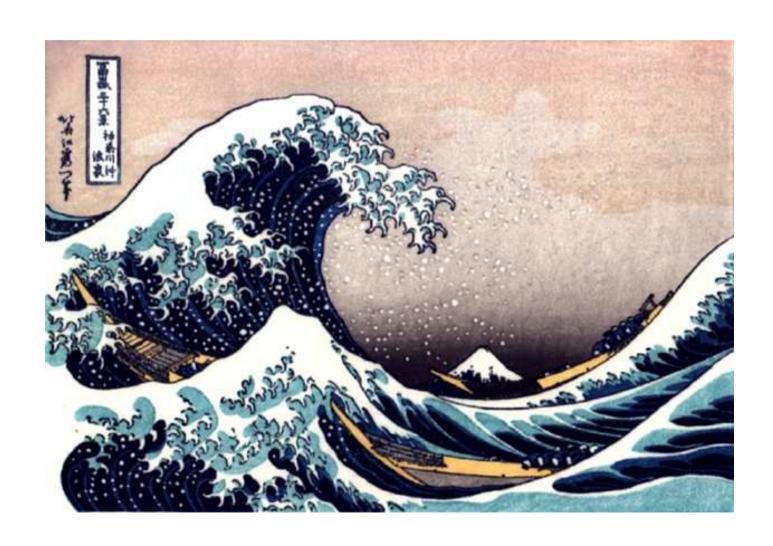
One part of the information explosion



New technologies also generate digital information



The problem



...and the user perspective



Biological information: current reality

- Knowledge scattered across the literature in many thousands of non-computer readable journal articles
- Hundreds of different databases, many in flat-file format
- Non-uniform or lack of external identifiers
- Lack of interoperability at the level of syntax and semantics

What do most users want?

- Intuitive interfaces
- To navigate smoothly from one resource to another
- Immediate availability of authoritative information; free of charge at the point-of-use
- Information that can be integrated / manipulated / visualized /output in another form

A sample scenario.....



Friedreich ataxia is a disease characterized by neurodegeneration and cardiomyopathy. The gene responsible for the disease is frataxin. It encodes a mitochondrial protein required for iron homeostasis and is evolutionarily conserved from yeast to man.

- Critically assess the experimental evidence that frataxin functions in iron-sulphur cluster synthesis.
- Review the effectiveness of anti-oxidants as treatment for the disease.
- Assemble as far as possible all other known information on the frataxin gene and the action of its protein

Now choose a portal ...



* [TOP] [Part 1] [Part 2] [Part 3] [All] [Journals] []

Part 1: Molecular Biology Search and Analysis

- 3D-ALI Database relating Proteins Structures and Sequences (see Documentation) at EMBL-Heidelberg W3
- AA Analysis 🏃 Protein Identification in SwissProt and PIR using Amino Acid Composition at EMBL-Heidelberg W3
- AA CompIdent Protein Identification in SwissProt using Amino Acid Composition at ExPASy, Switzerland W3
- AA CompSim Compare the Amino Acid Composition of a SwissProt Entry with Other Entries at ExPASy, Switzerland W3
- AA Sequence Analysis Multiple Analysis of a Native or Modified Amino Acid Sequence at Rockefeller U. W3
- AbCheck Test an Antibody Sequence Against the Kabat Database at <u>U. College-London</u>, UK W3
- ALIGN Optimal Global Alignment of Two Sequences with No Short-cuts at EERIE-Nimes, France and Align at MIPS, Gern
- AllAll Relationships in a Set of Related Peptides at ETH-Zürich, Switzerland W3
- AllAllDB^{*}
 Character Search the All-Against-All Database of SwissProt at ETH-Zürich, Switzerland W3
- AMAS Analysis of Protein Multiple Sequence Alignments at U. Oxford, UK W3
- ASC☆ Analytic Surface Calculation of PDB Protein Structures at EMBL-Heidelberg W3
- ATLAS

 ✓ Search DNA and Protein Sequence Databases at MIPS, Germany W3
- BCM Search Launcher at Baylor College of Medicine W3
- BEAUTY BLAST Enhanced Alignment Utility (see Documentation) at Baylor College of Medicine W3
- BERLIN RNA Databank of 5S rRNA and 5S rRNA Gene Sequences at <u>CAOS/CAMM</u>, The Netherlands W3
- BIOACCELERATOR at Weizmann Institute of Science, Israel W3
- bioCUSI Multiple Access to Biological Servers at OCMS-Oxford W3
- Biological Units Expanded PDB Entries with Full Biological Units at Brookhaven National Laboratory FTP
- Biologist's Search Palette Collection of Search Engines for Biological Databases at Oxford U., UK W3
- BLAST Basic Local Alignment Research Tool at NCBI, BLAST (hiv, vector, etc) at Stanford U., BLAST at GenomeNet, Ja France BLAST (with Entrez and SRS Links) at Baylor College of Medicine and BLAST at EPFL, Switzerland W3
- BLASTPAT BLAST-based Pattern Database Search (see Documentation) at Baylor College of Medicine W3
- BLITZ Fast comparison of protein sequences against SwissProt (see Documentation) at EBI, UK W3
- BLOCKS Database of Highly Conserved Regions in Proteins W3 and BLOCKS-WAIS Gopher at Fred Hutchinson Cancer
- BioMagResBank A Database of NMR-Derived Protein Structures at BIMAS-NIH W3
- BioSCAN Biological Sequence Comparative Analysis at U. North Caroline W3
- <u>BMCD</u> Biological Macromolecule Crystallization Database at <u>CARB</u> W3

There's plenty of choice!



* [TOP] [Part 1] [Part 2] [Part 3] [All] [Journals]

Part 2: Bibliographic, Text and WWW Searches

- AA-Index Amino Acid Index Database at GenomeNet, Japan and NAKAI at CAOS/CAMM, The Netherlands W3
- AGRICOLA Bibliographic References (Plant Genome Subset) at USDA-NAL Gopher
- Annual Reviews
 → Bibliographic Searches at LMMB-National Cancer Institute W3
- ArchiePlexForm Search Files Available by Anonymous FTP at LeRC-NASA W3
- ATCG Anderson's Timesaving Comparative Guides for Restriction Enzymes, Modifying Enzymes, Clone Libraries and Fi
- Beckman-Bioresearch Information on Protein and DNA Applications, Meetings and Product Information at Beckman I
- BIBLIO Molbio Journal Searches at Genethon, France Gopher and Biology Journal Contents at CERN, Switzerland W3
- BIBSYS Interuniversitary Bibliographic System in Norway W3
- BIOBIB Biocomputing Bibliography at ICGEB-Trieste, Italy Gopher and CompAlgo at CERN, Switzerland W3
- BioCatalog Catalog of Molecular Biology Programs at EBI, Bio-catalogue at Genethon, France and BIOCAT at CAOS/
- Bio-Email List of Biomedical E-Mail Servers and Bio-Email-Search at Washington U. W3
- Bioline Bio-Publications at Base de Dados Tropical, Brazil ₩3
- BioMOO The Biologists' Virtual Meeting Place at Weizmann Institute of Science, Israel W3
- Bionet Journal Articles and Bionet News Gopher
- BIOSCI Archives at EMBnet-UK Gopher and BIOSCI at CERN, Switzerland W3
- BioSpace Find People, Companies and Resources in Biotech Industry at Synergistic Designs W3
- Biotech Company WWW Registry Martin Leach's List of Commercial Net sites/Email addresses for Molecular Biolog.
- Biotechnology Abstracts DataBase at Stack-Serpukhov, Russia Gopher
- Biotechnology Bibliographies at BIC-U. Maryland/College Park W3

Although some may be difficult to find ...

Not Found

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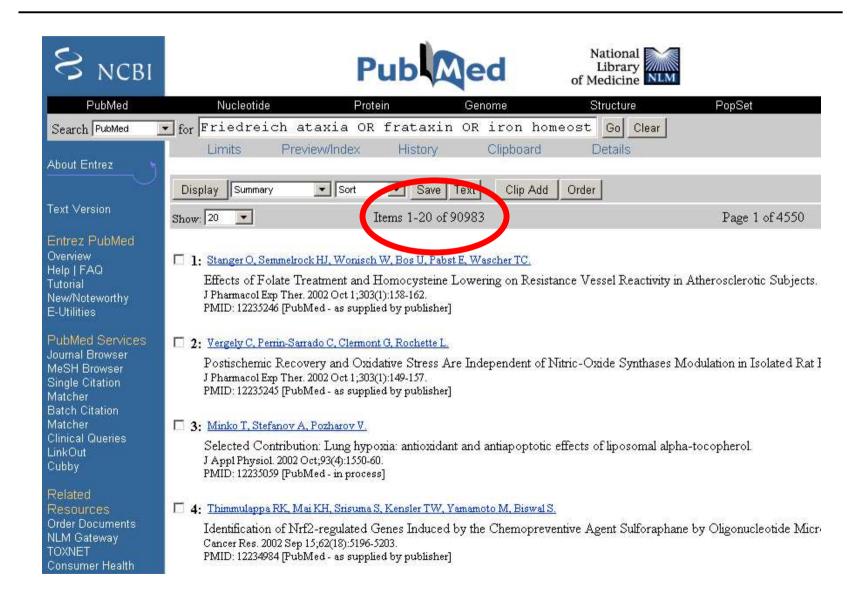
Apachell.3.26 Server at fire2.scl.genome.ad.jp Port 80

(23% web-links invalid after 1 yr; rising to 53% after 6 years Lawrence et al., 2001)

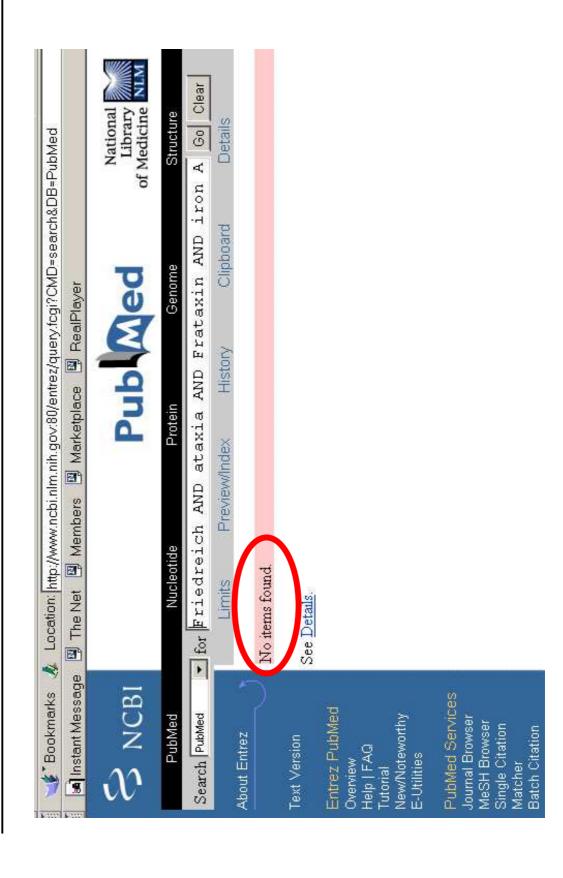
Try again ...

Comprehensive Yeast Genome Database **♦** home **√** Index ? help Gene / ORF Yeast - Full text search Go Reset Search: homology, div. Your Query: Frataxin AND iron AND antioxidant ... DB PPI / Complex Search Ascomycota Please note: this search-engine is not capable of logical operations as "AND" or "OR" View/Get chromosomes, seq. Tables misc., RNAs Catalogues Pathways

Another attempt



and another



And again?...













Success?



Google Search

Web Images Groups Directory

Searched the web for Friedreich ataxia Frataxin iron homeostasis yeast iron sulphur antioxidant

Antioxidants - Wholesale Pricing on Proancynol

www.healthwisdom.com Free shipping with purchase over \$50. Safe online ordering

Friedreich Ataxia

... mitochondrial iron homeostasis ... JM (2001) Antioxidant ... mitochondrial iron-sulphur ... Normal

serum iron ... with Friedreich's ataxia ... in yeast ... the frataxin ...

Description: An in depth look at this disorder. Includes a summary, diagnosing, molecular genetic testing, clinical...

Category: Health > Conditions and Diseases > ... > Friedreich Ataxia

www.geneclinics.org/profiles/friedreich/ - 85k - Cached - Similar pages

por Idebenone and reduced cardiac hypertrophy in Friedreich's ataxia

File Format: PDF/Adobe Acrobat - View as HTML

... acting as an antioxidant ... 8 and of yeast ... of cell iron homeostasis ... DISCUSSION Friedreich's

ataxia ... 20 Frataxin deficient cells ... deficiency of iron ...

www.ataxie.com/new/heart.pdf - Similar pages

[pnr]Clinical, biochemical and molecular genetic correlations in ...

File Format: PDF/Adobe Acrobat

... November 1999 Friedreich's ... of progressive ataxia ... of iron sulphur ... lipid soluble antioxidant ... homologue

of frataxin ... function, iron homeostasis ... the YFH1 yeast ...

ww2.mcgill.ca/biology/undergra/c524a/9-275.pdf - Similar pages

PDFIPIL: S0166-2236(00)01584-8

File Format: PDF/Adobe Acrobat - View as HTML

... Disruption of the yeast ... There is increased iron ... in the gene for frataxin ... type identical

to Friedreich's ataxia ... abnormalities in copper homeostasis ...

neuro.med.comell.edu/NYH-CMC/bealenergetics.pdf - Similar pages



A next generation, scalable information service for the life sciences that will interlink factual and image data repositories with the research literature

EU Quality of Life research infrastructure: platform under construction



E-B oSci The current E-BioSci partnership



- Distributed network of information resources
- Europe-based; world-wide role (Act locally think globally!)



The E-BioSci platform

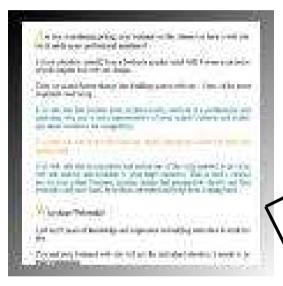
- Free access to the academic research community
- Set of *distributed* resources (literature, sequence- and image-databases)
- Integrated access to bibliographic abstracts, full text, factual and image databases
- Full-text search implemented via conceptual fingerprinting
 - Cross-repository
 - Cross-language
 - Implementation of effective 2-way links between databases and literature via gene symbol recognition





Conceptual fingerprints





Full text document

Linkage of index terms to the saurus

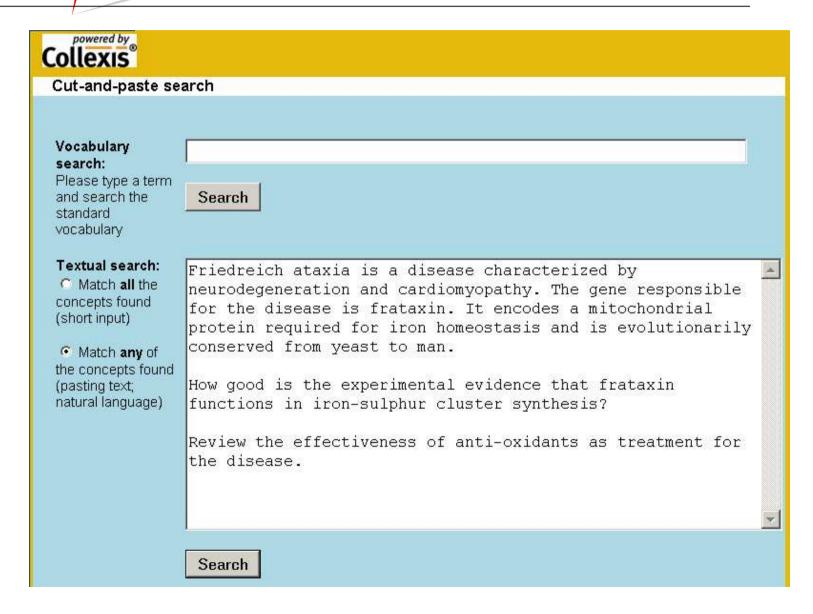
C19881 0.99 C92992 0.67 C02002 0.66 C99229 0.44 C00392 0.33 C93939 0.21

Fingerprint database

- •1 CFP = 400 bytes
- •Abstraction: 250.000 pages/PC/day
- •Matching: 500.000 CFP's: 40 millisec.



E-B OSci Our literature search once more....





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Refining the search



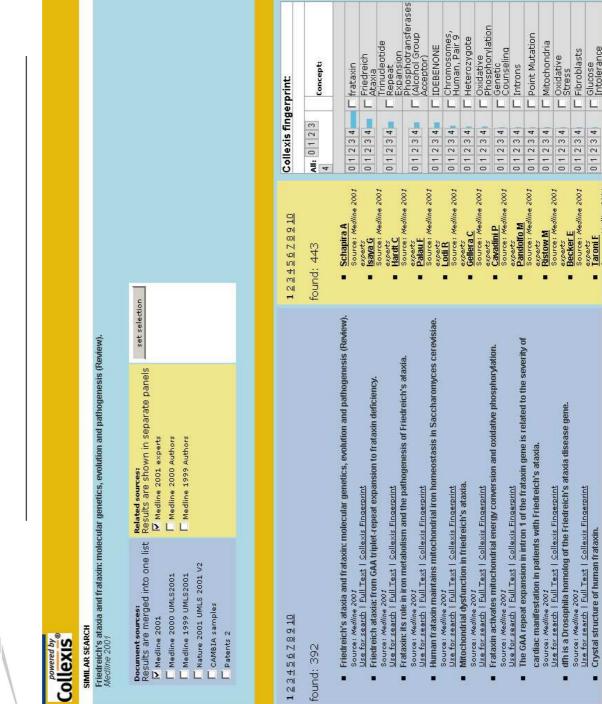
100 frataxin 74 Friedreich Ataxia 64 Trinucleotide Repeat Expansion 68 Phosphotransferases (Alcohol 69 Group Acceptor) 88 Chromosomes, Human, Pair 9 81 Heterozygote 27 Introns 27 Point Mutation 26 IDEBENONE 25 Oxidative Stress 23 Fibroblasts 22 Sequence Homology, Amino Acid 20 Alleles 19 Oxidative Phosphorylation 17 Phenotype 15 Amino Acid Sequence 14 Molecular Sequence Data

14 Iron 14 Mutation

Concept:	frataxin	Friedreich Ataxia	Trinucleotide Repeat Expansion	Phosphotransferases (Alcohol Group Acceptor)	Chromosomes, Human, Pair 9	Heterozygote	Introns	Point Mutation	IDEBENONE	Mitochondria	Oxidative Stress	Fibroblasts	Sequence Homology, Amino Acid	Alleles	Oxidative Phosphorylation	Phenotype	Amino Acid Sequence	Molecular Seguence Data	Iron	
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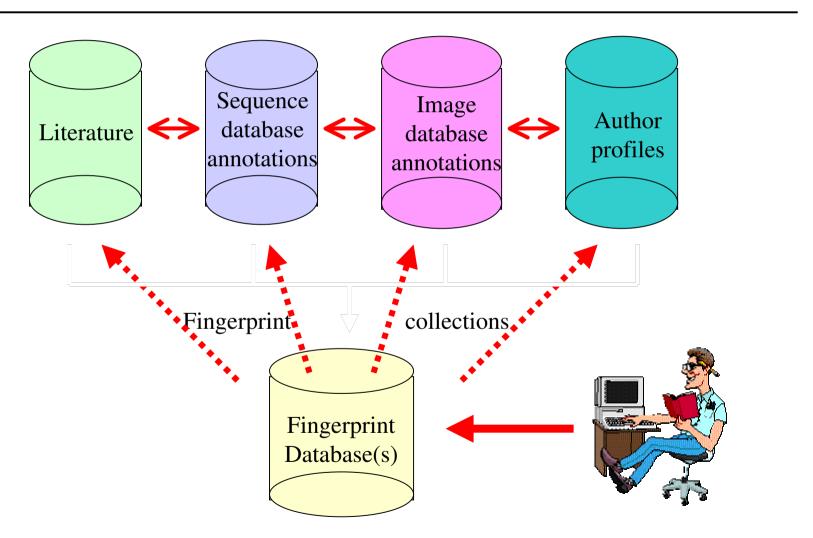
The result



Crystal structure of human frataxin.



E-Bosci Interconnecting searchable resources





The prototype platform:



Currently under construction, pre-testing, evaluation

See http://www.e-biosci.org for further information



Acknowledgements

- Frank Gannon, Executive Director EMBO
- ... and many others who contributed ideas to the concept of E-BioSci
- The E-BioSci partners
- European Commission (contract no QLRI-2001-30266)





