BINF 630: Introduction to Bioinformatics

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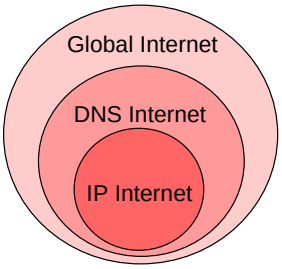
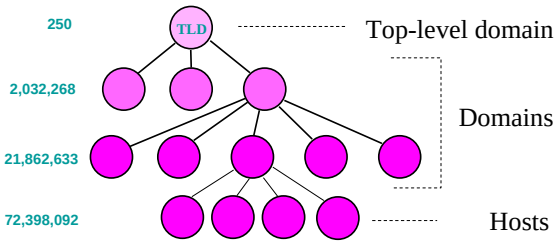
What is Internet



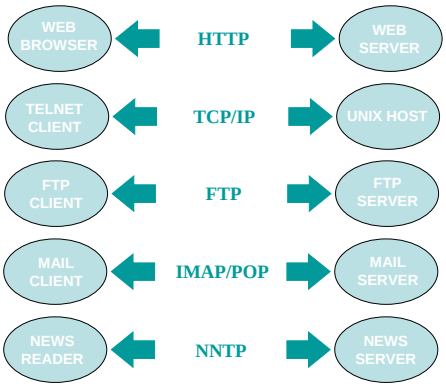
Domain Name System



Data from ISC
(January 2000)

Client - Server Model

Client - Server Model



Uniform Resource Locator (URL)

protocol://host.domain[:port]/path/filename

ftp - an anonymous FTP server (ftp://ftp.pdb.pdb.gov)
http - a World Wide Web server (http://mmlin4.pha.unc.edu/~cmb96)
telnet - a telnet session (telnet://nun.oit.unc.edu)

Network applications in science

- **Virtual Laboratory**
- **Virtual Library**
- **Virtual Conference**
- **Virtual Classroom**

Network collaboration

Real-time data sharing -- exchange of information between remote participants in the project

Resources sharing -- remote access to the instruments and computers

Resources integration -- simultaneous use of remote instruments and computers

Bioinformatics servers

Remote data access -- databse search, cross-links between the databases

Remote computing -- use of server's processing capabilities (sequence alignment, structure prediction, homology modeling)

Infospace navigation -- pointers to the available resources

Bioinformatics servers

Real-time

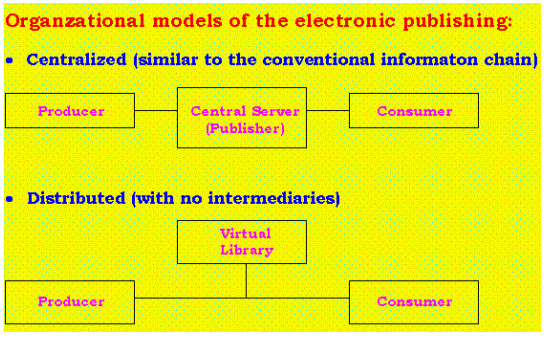
Asynchronous

Digital information cycle

Creation and capture
Storage and management
Rights management
Search and access
Distribution

Electronic publishing

Quality (peer review, retrospective evaluation)
Reliability (stability of serves, control over alterations, proper archiving and mirroring)



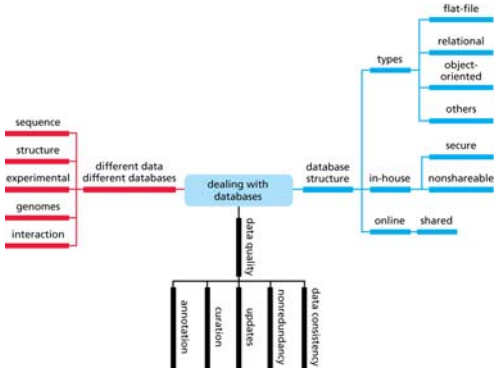
Hypertext Functionality in Scientific Literature

Active references
Forwarding references
Dynamic publishing

Ethical, Legal, and Economical Issues of Electronic Publishing

Intellectual property rights
Ownership of information
Information as a commodity

Databases in Bioinformatics



Molecular Databases

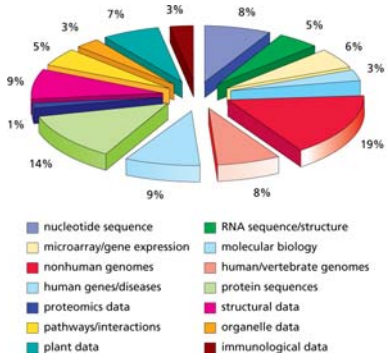
Nucleic acid sequences: GenBank, DNA Data Bank of Japan, EMBL Nucleotide Sequence Database
Nucleic acid structures: NDB - Nucleic Acid Database

Protein sequences: PIR - Protein Information Resource, SWISS-PROT
Protein structures: PDB - Protein Data Bank , NRL_3D

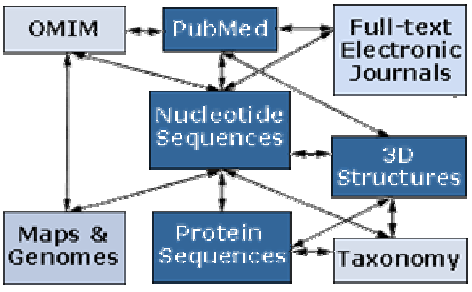
Physical properties: Biological Macromolecule Crystallization Database, BioMagResBank

Molecular images: SWISS-3DIMAGE, Molecules R US

Molecular Databases



NCBI integrated search and retrieval system
Entrez



NCBI Databases

- **nr** - All non-redundant GenBank CDS translations+PDB+SwissProt+PIR
- **month** - All new or revised GenBank CDS released in the last 30 days
- **swissprot** - the last major release of the SWISS-PROT protein sequence database (no updates)
- **yeast** - Yeast (Saccharomyces cerevisiae) protein sequences.
- **E. coli** - E. coli genomic CDS translations
- **pdb** - Sequences derived from the 3-dimensional structure Brookhaven Protein Data Bank
- **kabat** - Kabat's database of sequences of immunological interest

Database

Database

database	a collection of related structured information about entities
file	a collection of records
record	a set of fields
field	a single characteristic of an entity
character	a symbol used in data field

Example of a Genbank entry

```
LOCUS       VIBHALUXA    3141 bp    DNA             BCT       15-FEB-1996
DEFINITION  V.harveyi luciferase alpha and beta subunit (luxA and luxB) genes,
            complete cds.
ACCESSION   M10961 M13494
NID         g155174
KEYWORDS    luciferase.
SOURCE      Vibrio harveyi DNA.
  ORGANISM  Vibrio harveyi
            Eubacteria; Proteobacteria; gamma subdivision; Vibrionaceae;
            Vibrio.
REFERENCE   1  (bases 1 to 1838)
  AUTHORS   Cohn,D.H., Mileham,A.J., Simon,M.I., Nealson,K.H., Rausch,S.K.,
            Bonam,D. and Baldwin,T.O.
  TITLE     Nucleotide sequence of the luxA gene of Vibrio harveyi and the
            complete amino acid sequence of the alpha subunit of bacterial
            luciferase
  JOURNAL   J. Biol. Chem. 260 (10), 6139-6146 (1985)
  MEDLINE   85207595
REFERENCE   2  (bases 1745 to 3141)
  AUTHORS   Johnston,T.C., Thompson,R.B. and Baldwin,T.O.
  TITLE     Nucleotide sequence of the luxB gene of Vibrio harveyi and the
            complete amino acid sequence of the beta subunit of bacterial
            luciferase
  JOURNAL   J. Biol. Chem. 261 (11), 4805-4811 (1986)
  MEDLINE   86168191
```

Example of a Genbank entry

```
FEATURES             Location/Qualifiers
     gene            707..1774
                     /gene="luxA"
     CDS             707..1774
                     /gene="luxA"
                     /codon_start=1
                     /product="luciferase alpha subunit"
                     /db_xref="PID:g155175"
                     /transl_table=11
                     /translation="MKFGNFLLTYQPPELSQTEVMKRLVNLGKASEGCGFDTVWLLEH
                     HFTEFGLLGNPYVAAAHLLGATETLNVGTAAIVLPTAHPVRQAEDVNLLDQMSKGRFR
                     FGICRGLYDKDFRVFGTDMDNSRALMDCWYDLMKEGFNEGYIAADNEHIKFPKIQLNP
                     SAYTQGGAPVYVVAESASTTEWAAERGLPMILSWIINTHEKKAQLDLYNEVATEHGYD
                     VTKIDHCLSYITSVDHDSNRAKDICRNFLGHWYDSYVNATKIFDDSDQTKGYDFNKGQ
                     WRDFVLKGHKDTNRRIDYSYEINPVGTPEECIAIIQQDIDATGIDNICCGFEANGSEE
                     EIIASMKLFQSDVMPYLKEKQ"
BASE COUNT      883 a    665 c    741 g    852 t
ORIGIN      1 bp upstream of EcoRI site.
        1 gaattcacca tgacgacggg caaaaatagt ttgtgcactg tttatcactg gctgcagacc
       61 aagggcacac aaaacattgg cttgattgcg gcaagtctct cagctcgtgt cgcctatgaa
      121 gttatctctg atctggagct gtcttttctg attactgcgg ttggtgtggt gaacttgcgt
      181 gacacactag aaaaagcgct tggttttgat tacctcagtt tgcctatcga tgagctacca
          ....
```