

BCB 590
Practical Bioinformatics

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Introductions

- Name
- Lab
- Research Project
- What do you hope to learn in the next two weeks about bioinformatics?

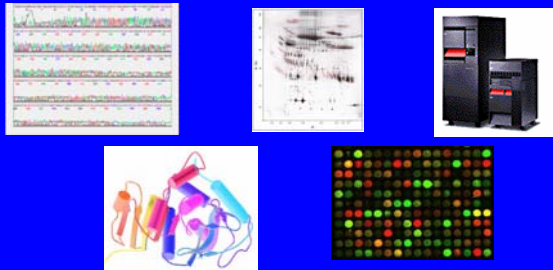
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Course Format

- Short introductory lecture
- Lab exercise
- Answer summary questions about the exercise and email answers to the instructor
- Final exam - lab practical
- Show up, do the labs, get an A

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What is Bioinformatics?
Computational Systems Biology?



Thanks to Drena Dobbs (ISU), Mark Gerstein (Yale) & Eric Green (NIH) for borrowed & modified PPTs

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What is Bioinformatics?
(& Computational Biology?)

Wikipedia:

- Bioinformatics & computational biology involve the use of techniques from mathematics, informatics, statistics & computer science (& engineering) to solve biological problems

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What is Bioinformatics?
(& Computational Biology?)

Gerstein:

- (Molecular) Bioinformatics is conceptualizing biology in terms of molecules & applying "informatics" techniques - derived from disciplines such as mathematics, computer science, and statistics - to organize and understand information associated with these molecules, on a large scale

Modified from Mark Gerstein

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What is Systems Biology?

Wikipedia:

- **Systems biology** - is a relatively new **biological study field** that focuses on the systematic study of complex interactions in biological systems, thus using a new perspective (integration instead of reduction) to study them

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What is the Information? Biological Sequences, Structures, Processes

<p>Central Dogma of Molecular Biology</p> <ul style="list-style-type: none"> • DNA sequence <ul style="list-style-type: none"> → RNA → Protein → Phenotype • Molecules <ul style="list-style-type: none"> • Sequence, Structure, Function • Processes <ul style="list-style-type: none"> • Mechanism, Specificity, Regulation 	<p>Central Paradigm for Bioinformatics</p> <ul style="list-style-type: none"> • Genomic (DNA) Sequence <ul style="list-style-type: none"> → mRNAs & other RNA sequences → Protein sequences → RNA & Protein Structures → RNA & Protein Functions → Phenotype • Large Amts of Information <ul style="list-style-type: none"> • Standardized • Statistical
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Modified from Mark Gerstein (idea from D. Braker, Stanford; graphics from S. Strahl)

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
Explosion of "Omes" & "Omics!" Genome, Transcriptome, Proteome



• **Genome** - the complete collection of DNA (genes and "non-genes") of an organism



• **Transcriptome** - the complete collection of RNAs (mRNAs & others) expressed in an organism



• **Proteome** - the complete collection of proteins expressed in an organism

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Even MORE Omes:

Glycome
Metabolome
Cytome
Diseasome!!!

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Genome = Constant (MOSTLY) Transcriptome & Proteome = Variable

- **Genome** - the complete collection of DNA (genes and "non-genes") of an organism
- **Transcriptome** - the complete collection of RNAs (mRNAs & others) expressed in an organism*
- **Proteome** - the complete collection of proteins expressed in an organism*

* Note: Although the DNA is "identical" in all cells of an organism, the sets of RNAs or proteins expressed in different cells & tissues of a single organism vary greatly -- and depend on variables such as environmental conditions, age, developmental stage, disease state, etc.

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Molecular Biology Information: DNA & RNA Sequences

Functions:

- Genetic material
- Information transfer (mRNA)
- Protein synthesis (tRNA/mRNA)
- Catalytic & regulatory activities (some very new!)

DNA sequence:

```
atggcaattaaatctggatcaatggtttggctctat
geacaacacgtgacatggaattgtaggtatnaa
atgcttatatgtgaantatgattcaacacggtcag
aaagatggttaacttagtggatgglanaacatccg
GcaaacTaaactgggggcaactcgtttgatcgtcttaactgatgaaa
ctgctcgtaaacatcacctcaggcggcaaaaagtt
```

RNA sequence has "U" instead of "T"

Information:

- 4 letter alphabet (DNA nucleotides: AGCT)
- ~ 1,000 base pairs in a small gene
- ~ 3 X 10⁹ bp in a genome (human)

- Where are the genes?
- Which DNA sequences encode mRNA?
- Which DNA sequences are "junk"?
- Which RNA sequences encode protein?

Modified from Mark Gerstein

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Molecular Biology Information: Protein Sequences

Functions: *Most cellular functions are performed or facilitated by proteins*

- Biocatalysis
- Cofactor transport/storage
- Mechanical motion/support
- Immune protection
- Regulation of growth and differentiation

Information:

- 20 letter alphabet (amino acids)
 - ACDEFGHIKLMNPQRSTVWY (but not BIQUXZ)
- ~ 300 aa in an average protein (in bacteria)
- ~ 3×10^6 known protein sequences

Protein sequences:

```

d1dhfa_
LNCIVAVSQNMIGIKNGDLPWPLRNEFRYFQRMIT
d8df_
LNSIVAVCQNMIGIKDGNLPWPLRNEYKYFQRMIS
            
```

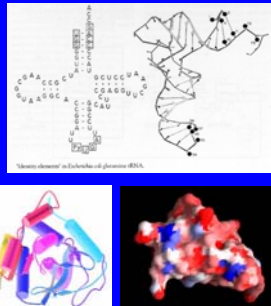
- What *is* this protein?
- Which amino acids are most important -- for folding, activity, interaction with other proteins?
- Which sequence variations are harmful (or beneficial)?

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Molecular Biology Information: Macromolecular Structures

DNA/RNA/Protein Structures

- How does a protein (or RNA) sequence fold into an active 3-D structure?
- Can we predict structure from sequence?
- Can we predict function from structure (or perhaps, from sequence alone?)

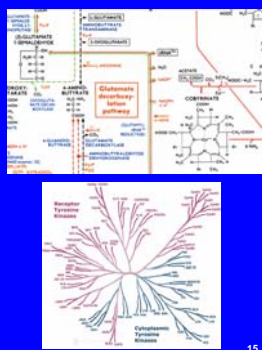


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Molecular Biology Information: Biological Processes & Systems

Functional Genomics

- How do patterns of gene expression determine *phenotype*?
- Which genes and proteins are required for differentiation during *normal development*? Are involved in disease?
- How do *proteins, signaling molecules and metabolites* interact in biological networks?
- Which genes and pathways have been most highly conserved during evolution? & *Why*?




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Gene Expression Data: Transcriptome (& Proteome)

MicroArray Data

Yeast Expression Data:

- Levels for all 6,000 genes!
- Experiments to investigate how genes respond to changes in environment or how patterns of expression change in normal vs cancerous tissue



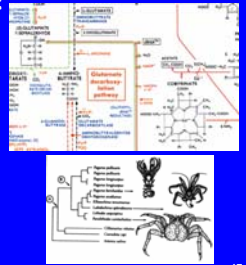
ISU's Biotechnology Facilities include state-of-the-art Microarray & Proteomics instrumentation

Modified from Mark Gerstein (courtesy of J Hager) 16

Molecular Biology Information: Integrating Data

Understanding the function of genomes requires integration of many diverse and complex types of information:

- Metabolic pathways
- Regulatory networks
- Whole organism physiology
- Evolution, phylogeny
- Environment, ecology
- Literature (MEDLINE)

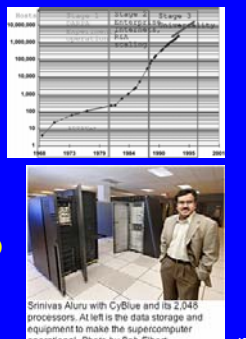


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Storing & Analyzing Large-scale Information: Exponential Growth of Data Matched by Development of Computer Technology

CPU vs Disk & Net

- Both the increase in computer speed and the ability to store large amounts of information on computers have been crucial
- Improved computing resources have been a driving force in Bioinformatics



ISU's supercomputer "CyBlue" is (WAS...) among 100 most powerful in the world

Modified from Mark Gerstein (Internet picture extracted from B. Karpis, Stanford) 18

Resources for Bioinformatics & Computational Biology

- [Wikipedia: Bioinformatics](#)
- [Wikipedia: Systems Biology](#)
- [NCBI - National Center for Biotechnology Information](#)
- [ISCB - International Society for Computational Biology](#)
- [JCB - Jena Center for Bioinformatics](#)
- [UBC - Bioinformatics Links Directory](#)

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(a few) ISU Resources & Experts

ISU Research Centers & Graduate Training Programs:

- [BCB - Bioinformatics & Computational Biology](#)
- [Baker Center - Bioinformatics & Biological Statistics](#)
- [CIAG - Center for Integrated Animal Genomics](#)
- [CILD - Computational Intelligence, Learning & Discovery](#)

ISU Facilities:

- [Biotech - Instrumentation Facilities](#)
- [CIAG - Center for Integrated Animal Genomics](#)
- [PSI - Plant Sciences Institute](#)
- [PSI Centers](#)

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Things we won't cover

- Comparative and functional genomics
- Gene prediction
- Microarray analysis
- Proteomics
- Phylogenetics
- Multiple sequence alignment
- Networks
- And more ...

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Topics for our course

- Biological databases
- Making sense of DNA and protein sequences
- BLAST
- RNA structure prediction
- Protein structure and function
- Protein structure prediction
- Protein structure visualization

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Our focus

- Effective use of some of the most commonly used bioinformatics tools and databases
- How to interpret your results
- How to find the information you need
- How to learn to use new tools

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- <http://dobbslab.gdcb.iastate.edu/BCB590>

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