



Bioinformatics Panel Presentation

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My Background

- **Direct bioinformatics research group of 6 people in Artificial Intelligence Center at SRI International**
- **Stanford Computer Science Ph.D., 1989**
- **1.5 year post-doc at National Institutes of Health**
- **At SRI from 1991-present**
- **Vice President at DoubleTwist Inc. from 1997-1999**
- **Consulting assistant professor of Medicine at Stanford, 1994-present**

Introduction



- **What are**
 - Bioinformatics
 - Computational Biology
 - Biomedical computing
- **(Computer-Science * j) + (Biology * k)**
- **$j = 1 - k$**
- **$0 > j < 1$**
- **Education, research results, journals, funding sources, conferences, collaborators**

$$j = .9 ; k = .1 ;$$

- **Computer scientist who**

- Performs computer science research in the context of biological problems
- Designs computational paradigms based on biological systems
- Earned Ph.D. in Computer science
- Publishes in computer science journals only
- Funded by NSF Computer Science

- May or may not ever actually solve a biological problem
- May or may not have biologist collaborators





$j = .1 ; k = .9 ;$

- **Biologist who**
 - Applies existing bioinformatics software to solve biological problems
 - Earned Ph.D. in biological sciences
 - Programs in Perl and SQL
 - Publishes in biology journals
 - Funded by NSF Biology, NIH, DOE

 - Might have taken a few computer science classes
 - May have developed some programming proficiency in other languages
- **My terminology: Computational biologist, not bioinformatics researcher**

$$j = .3 ; k = .7 ;$$

- **Interdisciplinary researcher who**

- Develops a biological database or its supporting software, develops software for genome analysis or visualization
- Develops sophisticated software to solve challenging biological problems
- Earned Ph.D. in biological sciences, M.S. in computer science
- Publishes in a mix of bioinformatics and biology journals
- Funded by NIH, NSF, DOE


$$j = .7 ; k = .3 ;$$

- **Interdisciplinary researcher who**
 - Develops novel bioinformatics algorithms, ontologies
 - Uses state of the art computer science, or performs computer science research, to solve biological problems
 - Earned Ph.D. in computer science, B.A. in biology
 - Publishes in a mix of bioinformatics, computer science, and biology journals
 - Collaborates with biologists
 - Funded by NSF, NIH, DOE

- Can't find a job



Typical Mistakes Made by Computer Scientists New to Bioinformatics

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- **Develop a beautifully engineered program that uses an elegant algorithm to rapidly solve the wrong problem**
- **Underestimate the importance of content**
- **Discovery = Algorithms + Databases**



How Changes in Computer Science Education Can Help Bioinformatics

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- **Most natural scientists have little understanding of computer science**
 - Computer science is programming
 - Cannot appreciate the value that computer scientists bring to bioinformatics
 - Complexity of software engineering

Database Education

- **Science in the 21st Century is information intensive**
- **Over 300 databases in bioinformatics**
- **The database area of bioinformatics is where practice falls farthest behind the state of the art**
 - Few bioinformaticians trained in databases, knowledge representation, ontologies, formal languages
 - Little use of commercial DB technology until recently
 - Considerable design flaws in many DBs
 - Elementary mistakes made over and over
 - Dependency of databases on history
 - Database expertise vs mathematics expertise
- **All natural scientists should be educated in the area that spans databases, AI knowledge representation and ontologies, formal grammars, data models**

My Research



- **Symbolic systems biology**
- **Encode biological theories in declarative form**
- **Knowledge base describing *E. coli* genome, proteome, metabolic pathways**
- **Algorithms and ontologies for metabolic pathways**
- **Algorithm for predicting the metabolic pathways of an organism from its genome**
- **Research in integrating knowledge bases and databases**

