Bioinformatics 2

Introduction

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Lecture 1

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- Course Overview & Assessment
- Introduction to Bioinformatics
- Careers and PhD options
- Proteomics guest lecture

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About me...

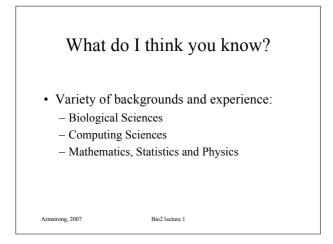
- Started in Biology (behaviour genetics)
- Got interested in databases (anatomy)
- Commercial and Academic Experience
- · 'wet lab' and bioinformatics projects
- Office in FH, Lab in HRB

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The class (2007) • M.Sc. Classes: • Quantitative Genetics and Genome Analysis (assignment 1 and term paper) • Bioinformatics 2 (assignment 1 and exam)

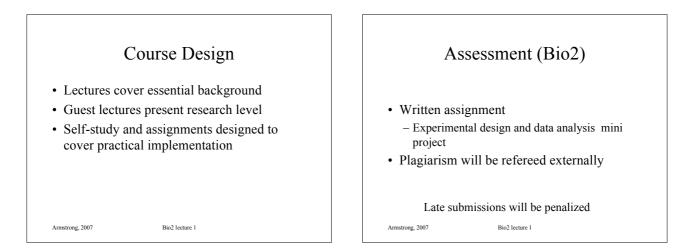
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Course Outcomes

- Know the core algorithms in bioinformatics
- Experience in using and/or implementing simple solutions
- Appreciate the current 'state of the art'
 - what has been solved?what are the key limitations?
- Be familiar with the available resources

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Bioinformatics?

- What is Bioinformatics?
- What does Bioinformatics do for CS?
- What does Bioinformatics do for Biology?

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- What guest bioinformatics lecture would you like?
- Discuss in groups for 10 min.

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What is BioInformatics?

- · Sequence analysis and genome building
- Molecular Structure prediction
- Evolution, phylogeny and linkage
- Automated data collection and analysis
- Simulations
- · Biological databases and resources

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BioInf and CS

- Provides CS with new challenges with clear medical significance.
- Complex and large datasets sometimes very noisy with hidden structures.
- Can biological solutions be used to inspire new computational tools and methods?

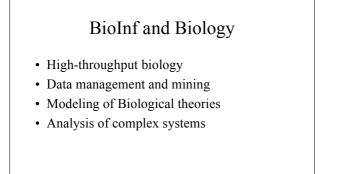
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BioInf and Biology

- High-throughput biology:
 - around 1989, the sequence of a 1.8kb gene would be a PhD project
 - by 1993, the same project was an undergraduate project
 - in 2000 we generated 40kb sequence per week in a non-genomics lab.

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bioinformatics

- <u>www.bbsrc.ac.uk</u> oasis database
- Bioinformatics.oxfordjournals.org
- <u>www.biomedcentral.org/bmcbioinformatics</u>

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• www.nature.com/msb

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- Database integration
- Data provenance

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- Evolutionary and genetic computation
- Gene expression databases
- High performance data structures for semistructured data (Vectorised XML)

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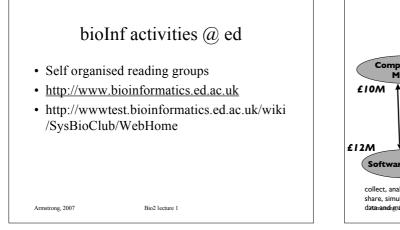
- Machine learning
- Microarray data analysis
- Natural language and bio-text mining
- Neural computation, visualisation and simulation

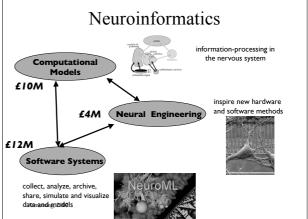
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- Protein complex modeling
- Systems Biology

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Career Options

- Academic Routes
 - Get Ph.D, do Postdoctoral Research lectureship and independent group
 - M.Sc. RA becomes semi independent usually linked to one or more academic groups. Career structure is less defined but improving. RAs can do Ph.D. part-time.

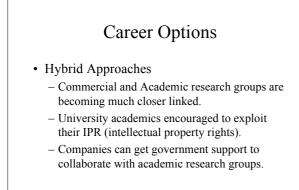
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Career Options

- Commercial Sector
 - Big Pharma Accept PhD and MSc entry. Normally assigned to projects and work within defined teams. Defined career structure (group leaders, project managers etc)
 - Spin-out/Small biotech Accept PhD and MSc entry. More freedom and variety. A degree of 'maintenance' work is to be expected.

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Ph.D.

- Assuming a start date of October 2007
- 'prize' studentships advertised on jobs.ac.uk, nature, science etc starting NOW!
 - Many linked to nationality/residency (Check details carefully).
- UK 'quota' studentships vary with department but contact/apply early.

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Ph.D.

- US studentships take longer but are better paid and have extra training/coursework
 - require an entry exam
 - again, deadlines are very soon for '07

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