COSC 348: Computing for Bioinformatics

Lecture 1: Introduction

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http://www.cs.otago.ac.nz/cosc348/

Why Computing for Bioinformatics?

- Bioinformatics is an *applied* area of computing.
- We encounter problems that are unlike any in studying compilers or operating systems, computer architecture, networks, graphics, etc.
- The kind of problems Bioinformatics presents us with are fundamentally *algorithmic* problems.
- We have to understand the limitations of our answers, especially how well they scale with problem size.

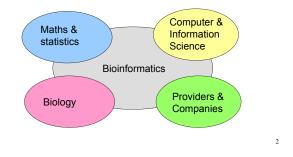
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• We have to start by understanding what the biological data and problem actually are.

Bioinformatics

• Bioinformatics is the field of science, in which biology, mathematics, statistics, computer science, and information technology merge into a single discipline.



Computing for Bioinformatics

- This course is designed for "tool-developers" rather than tool users.
 It is not about statistical analysis of data neither. This is the subject of STAT 435 at the Dept. of Math & Stats.
- Three major sub-disciplines in relation to computation:
 Algorithms to assess relationships between molecules;
 - Algorithms to analyse various types of molecular data;
 - Tools to enable efficient access and management of different types of data and information (= database and IR tools).

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Assessment

- Assessment consists of two programming assignments (worth respectively 14% and 14% of the final mark):
 http://www.cs.otago.ac.nz/cosc348/assignments.php
 - http://www.cs.otago.ac.nz/cosco46/assignments.php
- 12% for lab work (1% for each lab work starting at week 2):
 http://www.cs.otago.ac.nz/cosc348/labs.php
- a comprehensive closed book final examination (worth 60%):
 http://www.otago.ac.nz/library/exams/
- Lecture notes and other supplementary material for study:
 http://www.cs.otago.ac.nz/cosc348/lectures.php

Bioinformatics jobs (quotes from real ads)

 Job description: Design, development, implementation and testing of bioinformatics tools.

Typical requirements:

- Experience with the use of bioinformatics applications and tools for genomic analysis;
- Knowledge of necessary biological databases;
- Knowledge and experience with scripting and programming languages;
- Knowledge and well developed IT technical skills with emphasis on Unix, pipeline development and web based technologies.

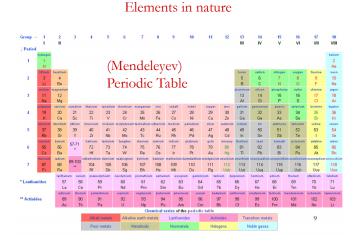
Bioinformatics companies and research in NZ NZ Genomics (provides genomics technology and bioinformatics

• AgResearch (analysis of bovine, sheep and plant genomes)

services to underpin research in a broad range of areas)

- PEBL (Pacific Edge Biotechnology), Dunedin (development of cancer diagnostic tests)
- University of Otago

 Database of RNA sequences / motifs, analysis of various gene data, etc.
- University of Auckland / Institute of Bioinformatics
 - New pharmaceuticals, bacterial genomes analysis, etc.



Introduction: 3 lectures

• Basic concepts

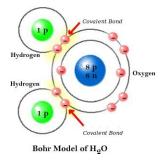
- Elements, atoms, molecules
- Cell, nucleus, chromosomes
- DNA, RNA, proteins
- Genetic code, genes, "junk" DNA
- Gene transcription regulation

• Big picture

- Individual development during life
- Genetic diseases
- Evolution

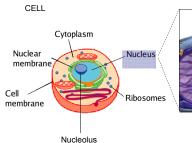
Elements, atoms, molecules

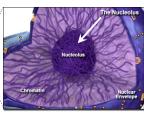
- Number assigned to each element is the number of protons/neutrons in the nucleus of its atom and the corresponding number of electrons in its orbit
- Atoms form molecules via chemical bonds
 - E.g. molecule of water
 H₂O, is made of two atoms of hydrogen and one atom of oxygen

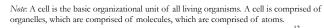


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Cell and nucleus







Biological molecules: COHN

- Amino acids: building blocks of proteins
 We are built by 20 amino
 - acids – Each has a different
 - residue group R
- Nucleotides: building blocks of DNA & RNA
 - 4 nucleotides Each has a different base
 - group

