

## Computational methods in bioinformatics: Lecture 1

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## What is biology?

Ecosystem	Rain forest, desert, fresh water lake, digestive tract of an animal
Community	All species in an ecosystem
Population	All individuals of a single species
Organism	One single individual
Organ System	A specialised functional system of an organism, e.g. nervous system or immune system
Organ	A specialised structural system of an organism, e.g. brain or kidney
Tissue	A specialised substructure of an organ, e.g. nervous tissue, smooth muscle
Cell	A single cell, e.g. neuron, skin cell, stem cell, bacteria
Molecule	e.g. protein, DNA, RNA, sugar, fatty acid, metabolites, pharmaceutical drugs



## What is bioinformatics?

*"Research, development, or application of computational tools and approaches for expanding the use of biological, medical, behavioral or health data, including those to acquire, store, organize, archive, analyze, or visualize such data."*

*"Bioinformatics applies principles of information sciences and technologies to make the vast, diverse, and complex life sciences data more understandable and useful."*

Working definition by the NIH Biomedical Information Science and Technology Initiative Consortium, 2000

<http://www.bisti.nih.gov/docs/CompuBioDef.pdf>



## What is computational biology?

*"The development and application of data-analytical and theoretical methods, mathematical modeling and computational simulation techniques to the study of biological, behavioral, and social systems."*

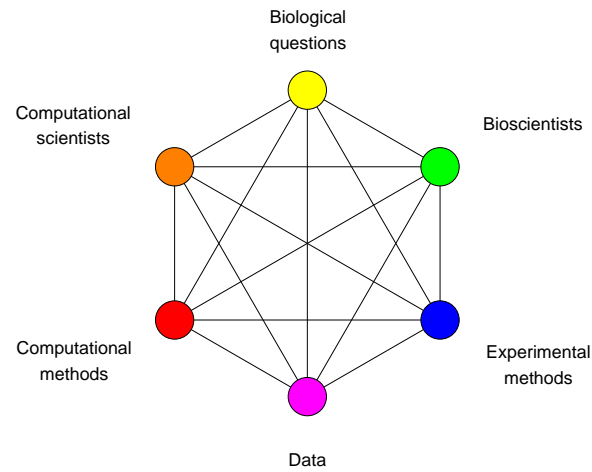
*"Computational biology uses mathematical and computational approaches to address theoretical and experimental questions in biology."*

Working definition by the NIH Biomedical Information Science and Technology Initiative Consortium, 2000

<http://www.bisti.nih.gov/docs/CompuBioDef.pdf>



# Addressing biological questions



# Sequences, structures and systems

## Sequences

- ▶ Nucleic acids (DNA and RNA) and proteins are (unbranched) polymers. Their composition can be described by the sequence of units (nucleotides or amino acid residues) in a chain.

## Structures

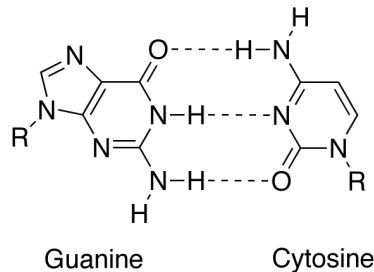
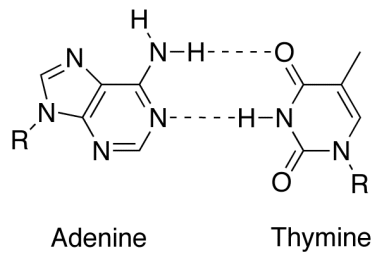
- ▶ Three-dimensional structures can give insights into the molecular basis of biological functions.

## Systems

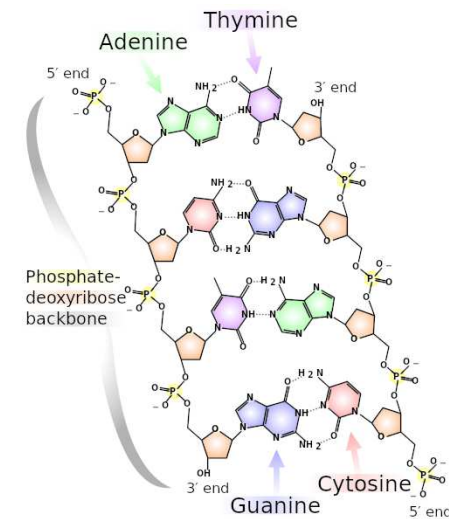
- ▶ Biological processes consist of the coordinated actions of molecules.



# Base pairing in DNA



# Structure of DNA



## Protein structure

### Primary structure

- ▶ sequence of amino acid residues linked in a chain

### Secondary structure

- ▶ locally, the main chain forms helices and strands

### Tertiary structure

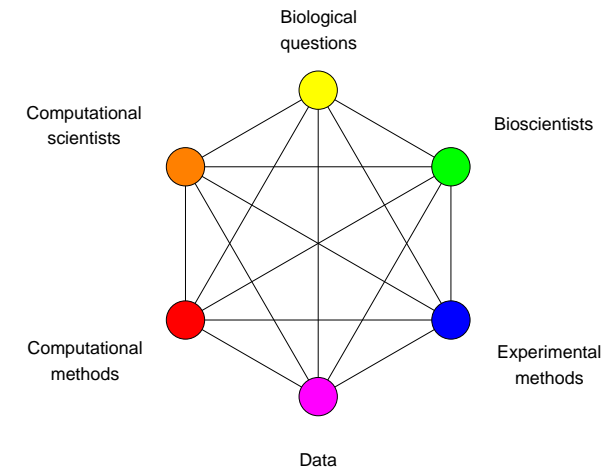
- ▶ the 3-D structure
- ▶ assembly and interaction of helices and sheets

### Quaternary structure

- ▶ assembly of subunits



## Addressing biological questions



## Biological sequences: some experimental methods

- ▶ DNA sequencing
- ▶ Protein sequencing
- ▶ Next-generation sequencing (NGS)



## Biological sequences: some questions

- ▶ How similar are a pair of sequences?
- ▶ Identify the corresponding units in a pair of homologous molecules that have undergone substitutions and insertions/deletions during their evolutionary history (*pairwise sequence alignment*).
- ▶ Given a new sequence, has anything similar (in whole or part) been seen before?
- ▶ Reconstruct a phylogenetic tree from the sequences of a set of homologous molecules.
- ▶ Given the sequences of many overlapping DNA fragments from a single organism, assemble them to reconstruct a full genome.
- ▶ Given the sequences of many DNA fragments from a mixture of organisms, identify the species present in the mixture.



## Biological structures: some experimental methods

Find the atomic structure of a macromolecule or complex

- ▶ X-ray crystallography
- ▶ Nuclear magnetic resonance (NMR) spectroscopy

Identify a low-resolution “envelope” enclosing a large macromolecular complex

- ▶ Cryo-electron microscopy
- ▶ Small-angle x-ray scattering



## Biological structures: some questions

- ▶ Can differences in the functions of two similar proteins be explained by differences in their structures?
- ▶ Can a drug be designed to fit into the active site of a target protein?
- ▶ Can the safety and efficacy of a potential therapeutic protein be predicted from its structure?
- ▶ Can the function of a protein be altered by changing its composition, and hence its structure?
- ▶ Can a protein's structure be predicted from its sequence?
  - ▶ the protein folding problem
- ▶ Given the structures of two proteins, will they associate with one another? If so, how will they fit together?
  - ▶ the protein docking problem



## Biological systems: some experimental methods

Which mRNA molecules are being expressed?

- ▶ Microarray gene expression
- ▶ RNA-Seq

Which proteins are being expressed?

- ▶ (2-D) gel electrophoresis
- ▶ Mass spectrometry

In which tissue(s) are particular genes expressed?

- ▶ *in situ* hybridization



## Biological systems: some questions

- ▶ Which genes/proteins are co-expressed (i.e. have similar expression profiles)?
- ▶ Which genes are expressed in tumour cells but not in healthy cells?
- ▶ If a gene is “knocked out”, will an organism survive, and how will the expression of other genes be affected?
- ▶ Can protein expression profiles identify proteins that could be targets for drug development?
- ▶ Can an individual's expression profile indicate whether they are likely to respond to a particular therapeutic treatment?
- ▶ How do biological networks respond to injury or to treatment with a therapeutic drug?

