

HMM for Bioinformatics

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- W.H. Majoros, “Methods for Computational Gene Prediction”, Cambridge University Press 2007;

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- T. Koski, “Hidden Markov Models for Bioinformatics”, Kluwer Academic Publishers 2001.

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- <http://smorfland.uni.wroc.pl/~blazej/HMM/>

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- ➋ To persons with a mathematical training, biology appears to be bewildering array of terminology and conventions;

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- ➋ All the biological statements that will appear during this lecture are "probably approximately correct".

Statement

All life on this planet depends on three types of molecule DNA, RNA and proteins,

Statement

DNA is a vast library describing how the cell works

Statement

RNA acts to transfer certain short pieces of this library to different places in the cell at which point those smaller volumes of information are used as a templates to synthetize proteins.

Statement

Proteins form enzymes that perform biochemical reactions, send signals to other cells, form the body's major component and otherwise perform the actual work of the cell.

Short history of DNA and RNA

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- ③ By the 1920s, nucleic acids were grouped into two classes called DNA and RNA, that differ slightly in their base composition.

fact 1.

Ironically, for a long time biologists paid a little attention to DNA.

Important date

1940 – Oswald Avery and colleagues proved that genes indeed reside on DNA.

What is the structure of DNA?

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What is the structure of DNA?

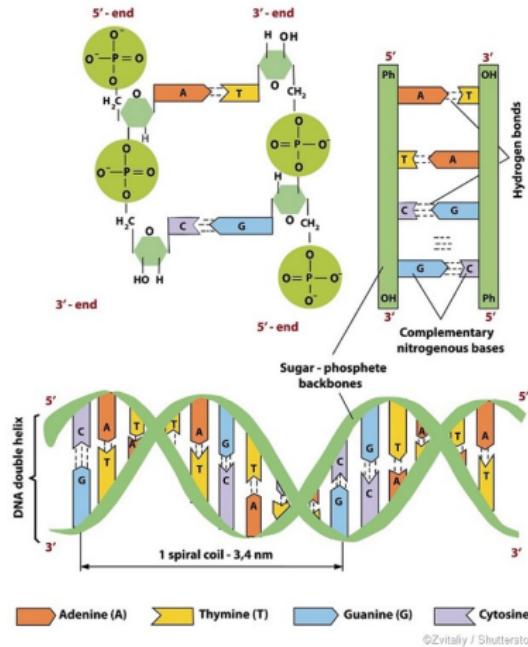
- ① 1950 – Erwin Chargaff discovered one-to-one ratio of the adenine-to-thymine and guanine-to-cytosine in DNA;
- ② 1951 – Maurice Wilkins and Rosalind Franklin obtained sharp x-ray images of DNA;
- ③ 1953 – James Watson and Francis Crick determined the double helical structure of a DNA molecule.

The Human Genome Project originally aimed to map the nucleotides contained in a human haploid reference genome (more than three billion). The "genome" of any given individual is unique; mapping the "human genome" involved sequencing a small number of individuals and then assembling these together to get a complete sequence for each chromosome. Therefore, the finished human genome is a mosaic, not representing any one individual.

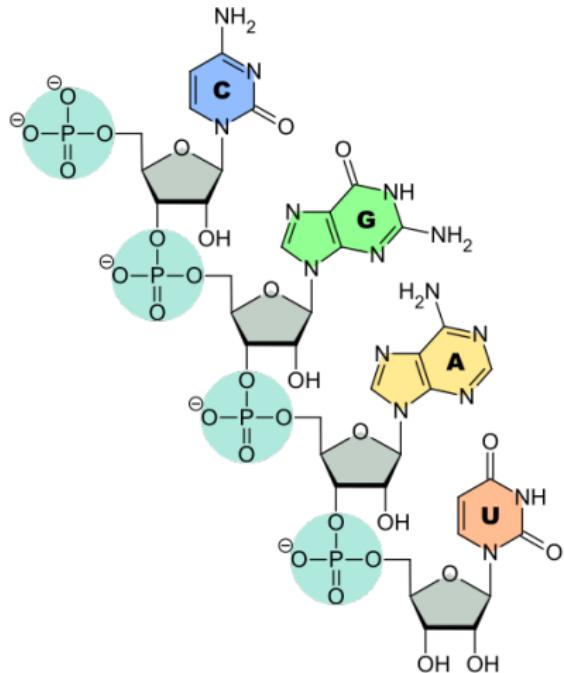
Rosetta Stone



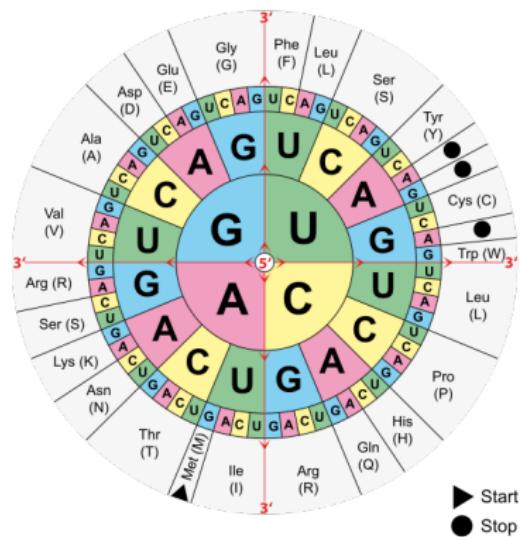
DNA Structure



DNA and RNA



The standard genetic code



The Central Dogma of Genetics

