

# HMM for Bioinformatics

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27 lutego 2019

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

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- 2 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 synthesize 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.

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- 2 By the early 1900s it was shown that DNA was a long molecule consisting of four types of bases;
- 3 By the 1920s, nucleic acids were grouped into two classes called DNA and RNA, that differ slightly in their base composition.

# Short history of DNA and RNA

## 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?

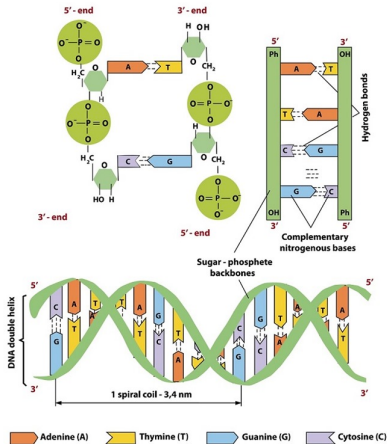
- 1 1950 – Erwin Chargraff discovered one-to-one ratio of the adynine-to-thymine and guanine-to-cytosine in DNA;
- 2 1951 – Maurice Wilkins and Rosalind Franklin obtained sharp x-ray images of DNA;
- 3 1953 – James Watson and Francis Crick determined the double helical structure of a DNA molecule.

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

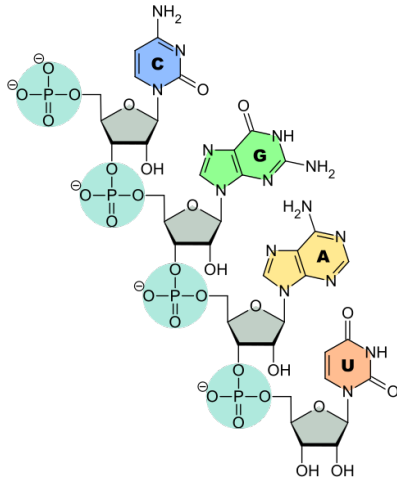


## DNA Structure

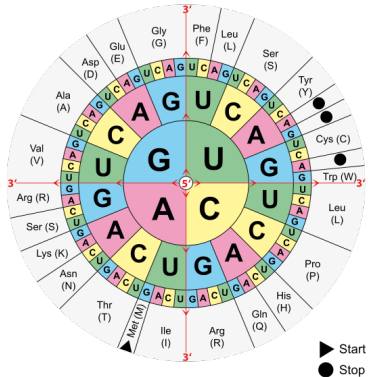


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# DNA and RNA



# The standard genetic code



# The Central Dogma of Genetics

