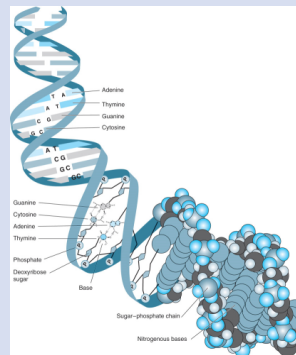
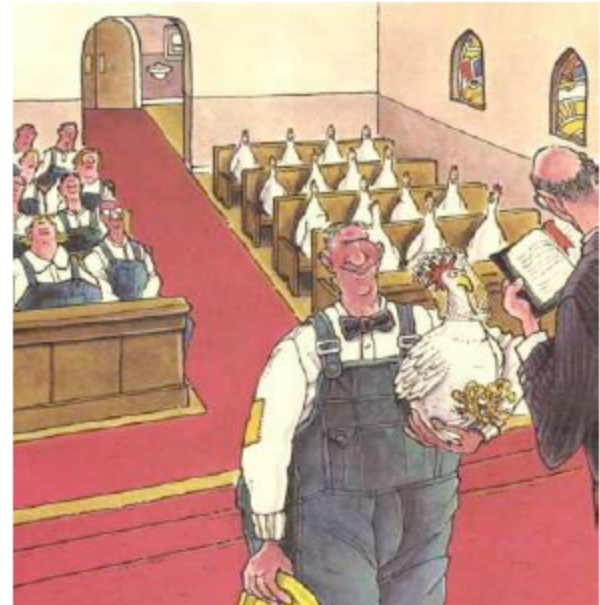


Bioinformatics



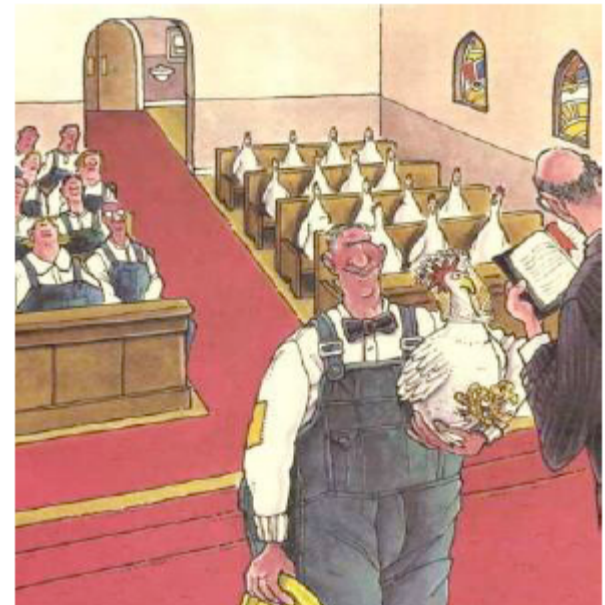
Introduction

- What is bioinformatics?
Interdisciplinary subject between:
 - Computer science
 - Biology
 - Medicine
 - Other life sciences
- A marriage between biology and computers



Introduction

- What is bioinformatics?
biological and medical data on:
 - Nucliec acids:
 - RNA
 - DNA
 - Genes
 - Proteins
 - Bioinformatics is generally defined as the analysis, prediction, and modeling of biological and medical data with the help of computers



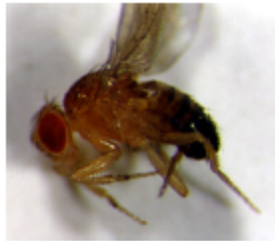
Bioinformatics introduction: continued

- Bioinformatics is the combination of biology and computing.
- DNA sequencing technologies have created massive amounts of information that can only be efficiently analyzed with computers.
- So far 70 species sequenced
 - Human, rat chimpanzee, chicken, and many others.
- As the information becomes ever so larger and more complex, more computational tools are needed to sort through the data.
 - Bioinformatics to the rescue!!!

Biological and medical data:

- Data storage and retrieval
- analysis of data
- Information extraction from data
- Gene functions
- Pathways
- Protein function
- Sequence and structure analysis





Drosophila



C. elegans



Rat



Human



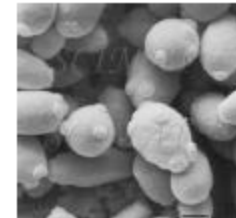
Mouse



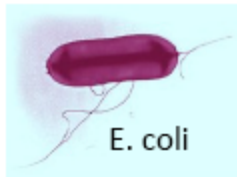
Rice



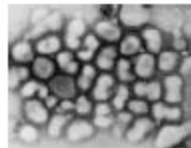
Mosquito



Yeast



E. coli



H. influenza



Arabidopsis

Species and Genomes

History

- Molecular biology began in 1600s
- Robert Hooke (1635-1703) discovered organisms are made up of **cells**



- Robert Hooke

Genetics

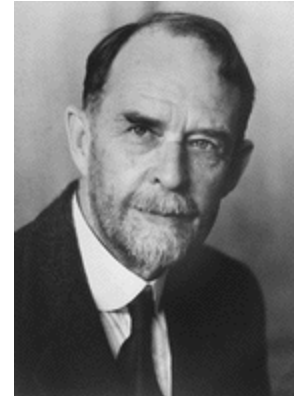
- **1865** **Gregor Mendel** discover the basic rules of **heredity** of garden pea.
 - An individual organism has two alternative heredity units for a given trait (**dominant trait** v.s. **recessive trait**)



Mendel:
The Father of **Genetics**

Genes and RNA

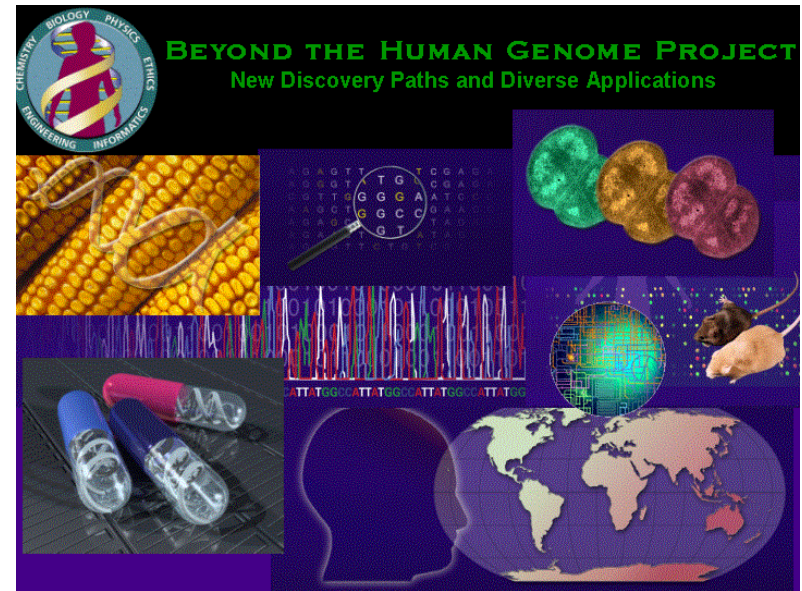
- **1911** – Thomas Hunt Morgan discovers **genes** on chromosomes are the discrete units of heredity
- **1911** Pheobus Aaron Theodore Lerene discovers **RNA**
- **1941** – George Beadle and Edward Tatum identify that genes make **proteins**



Thomas
Morgan

Genome

- **1986** Human **Genome** Initiative announced
- **1990** The 15 year Human Genome project is launched by congress
- **April 2003** Human Genome Project Completed. Mouse genome is sequenced.
- **April 2004** Rat genome sequenced.



The three main objects



DNA

Gene

Genome



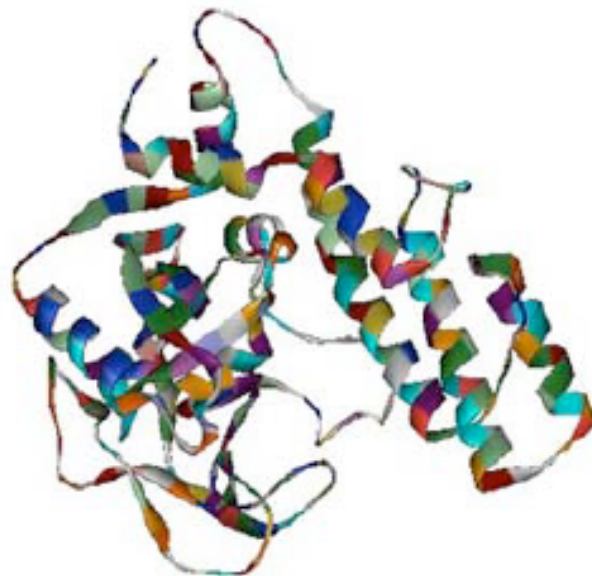
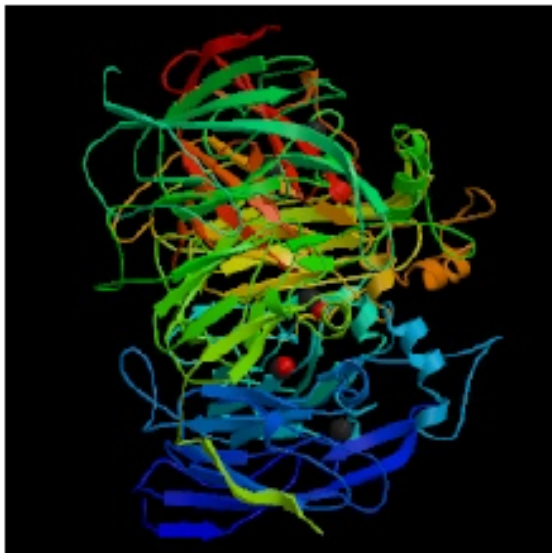
RNA

messenger



Protein

proteome

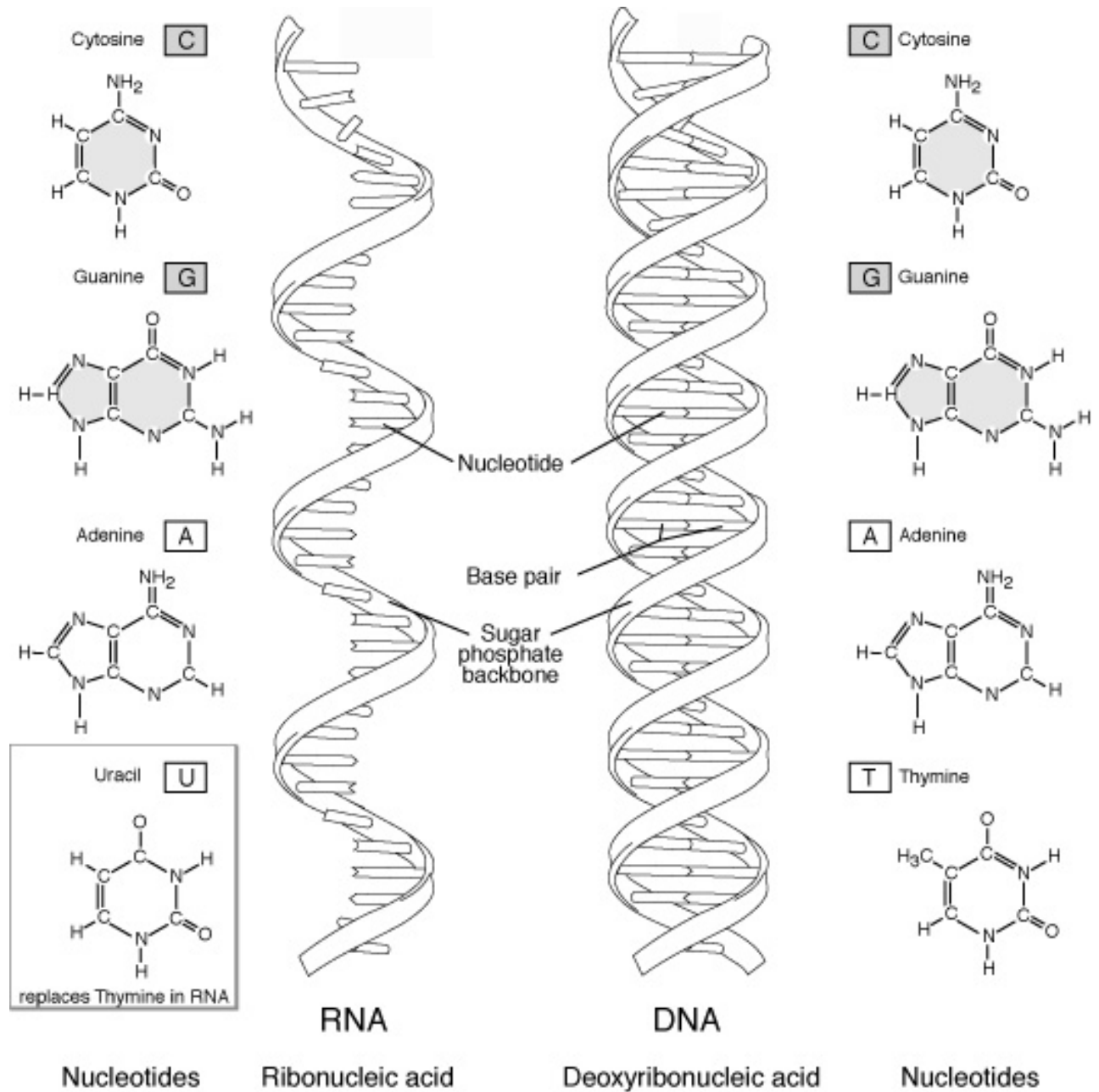
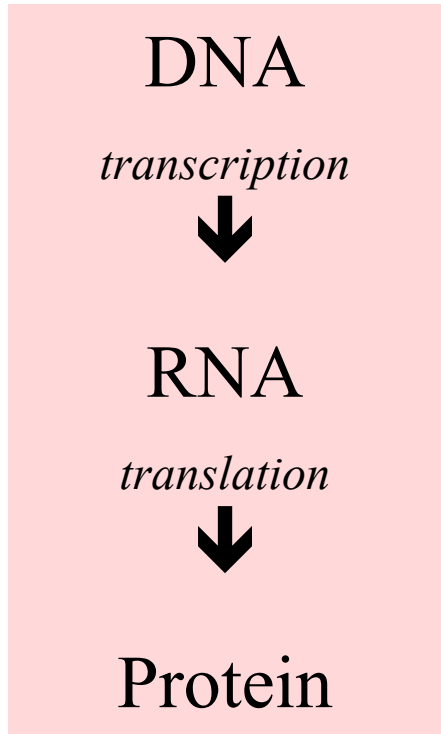


Protein 3D structure

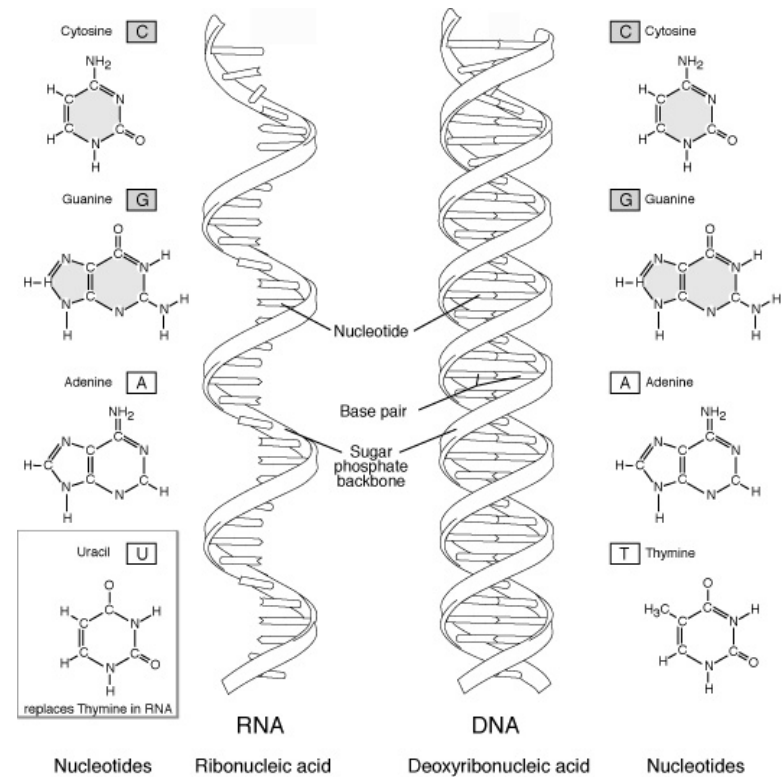


- DNA (**Deoxyribonucleic acid**) is arranged into 23 chromosome pairs in the nucleus of each cell.
- DNA: 4 bases:
 - A, C, G, T
- Just like human text, genomic sequences represent a language written in A, T, C, G

-DNA
-RNA
-Protein



-DNA
-RNA
-Protein



Protein: is a chain of amino acid

- there are around 20 amino acids known in human body

A protein is usually composed of 50 to 400+ amino acids.

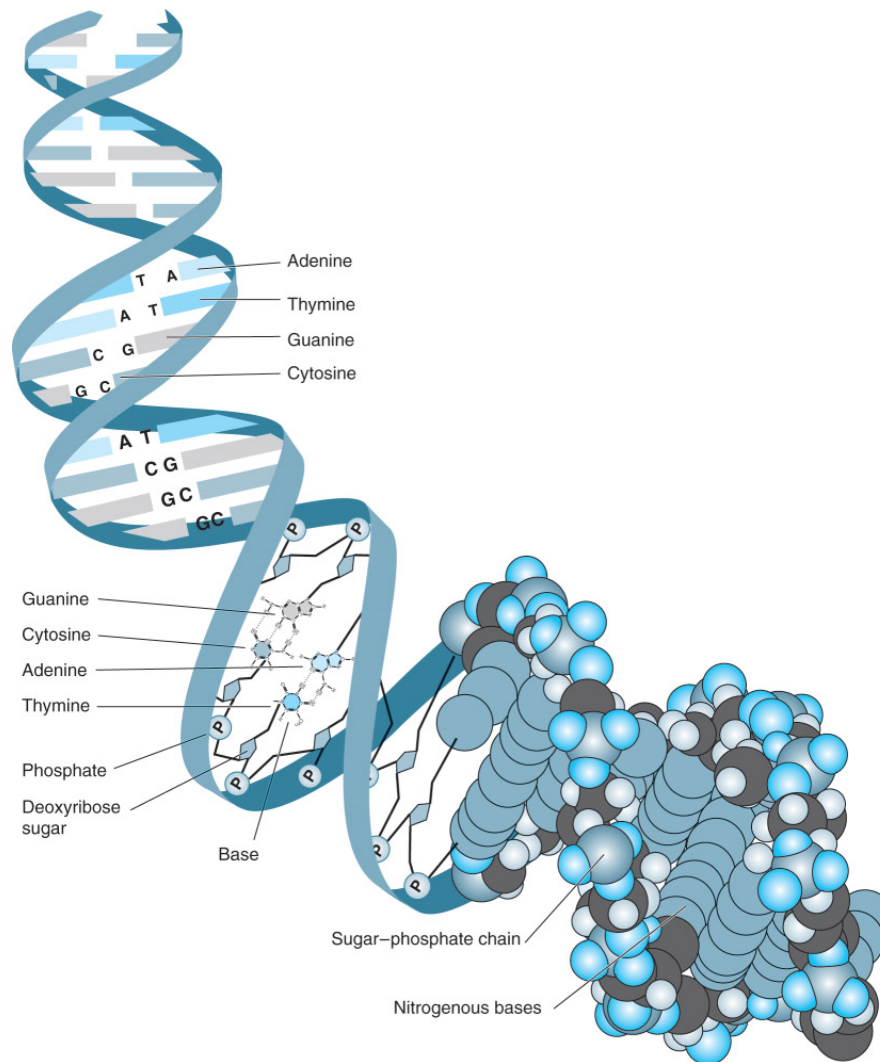
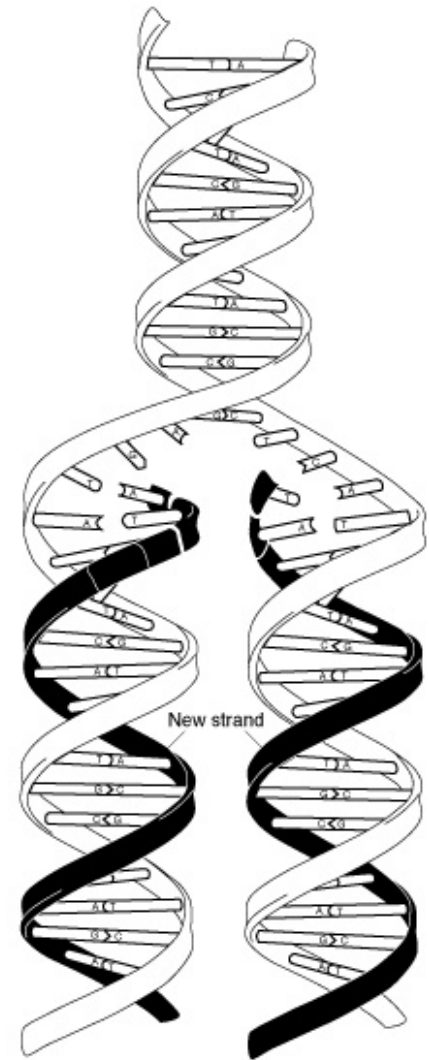
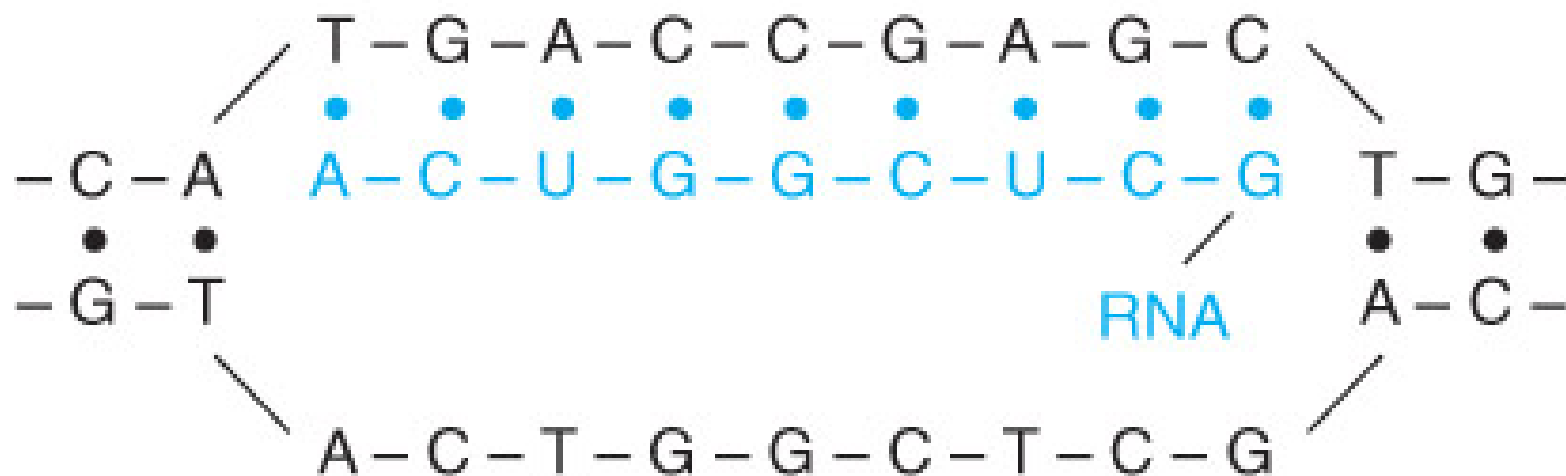
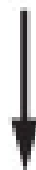
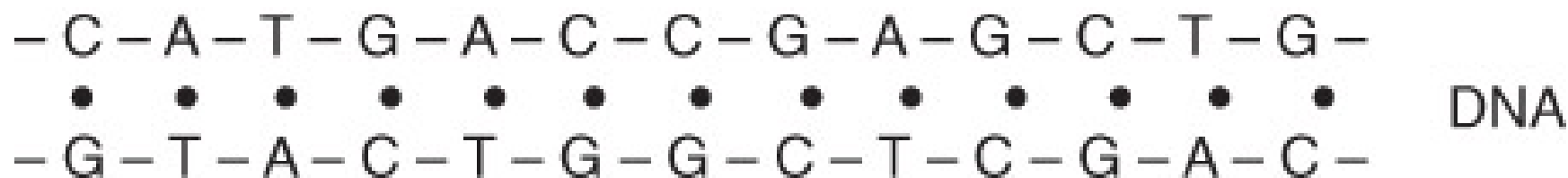


Figure 2.5: A DNA molecule is made up of two strands of nucleotides

DNA Replication

- Prior to cell division, all the genetic instructions must be “**copied**” so that each new cell will have a complete set
- DNA **polymerase** is the enzyme that copies DNA

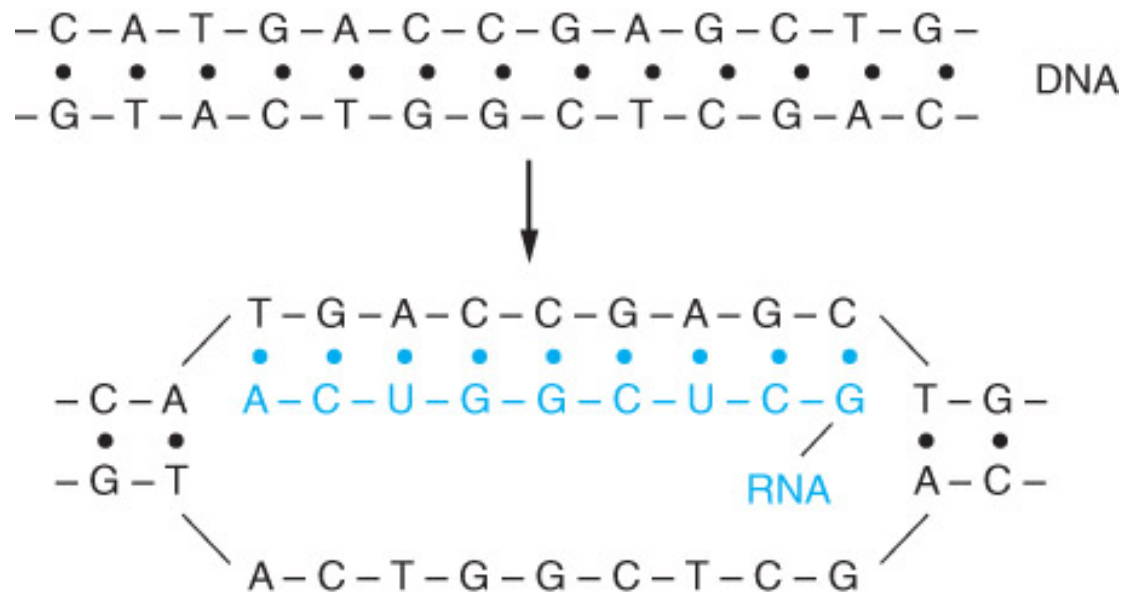




DNA: C G A T

RNA: C G A U

==> notice: RNA is single strand.



- DNA: C G A T
- RNA: C G A U
- Proteins: 20 letters from the English alphabet
(*all letters except for B, J, O, U, X, and Z*).

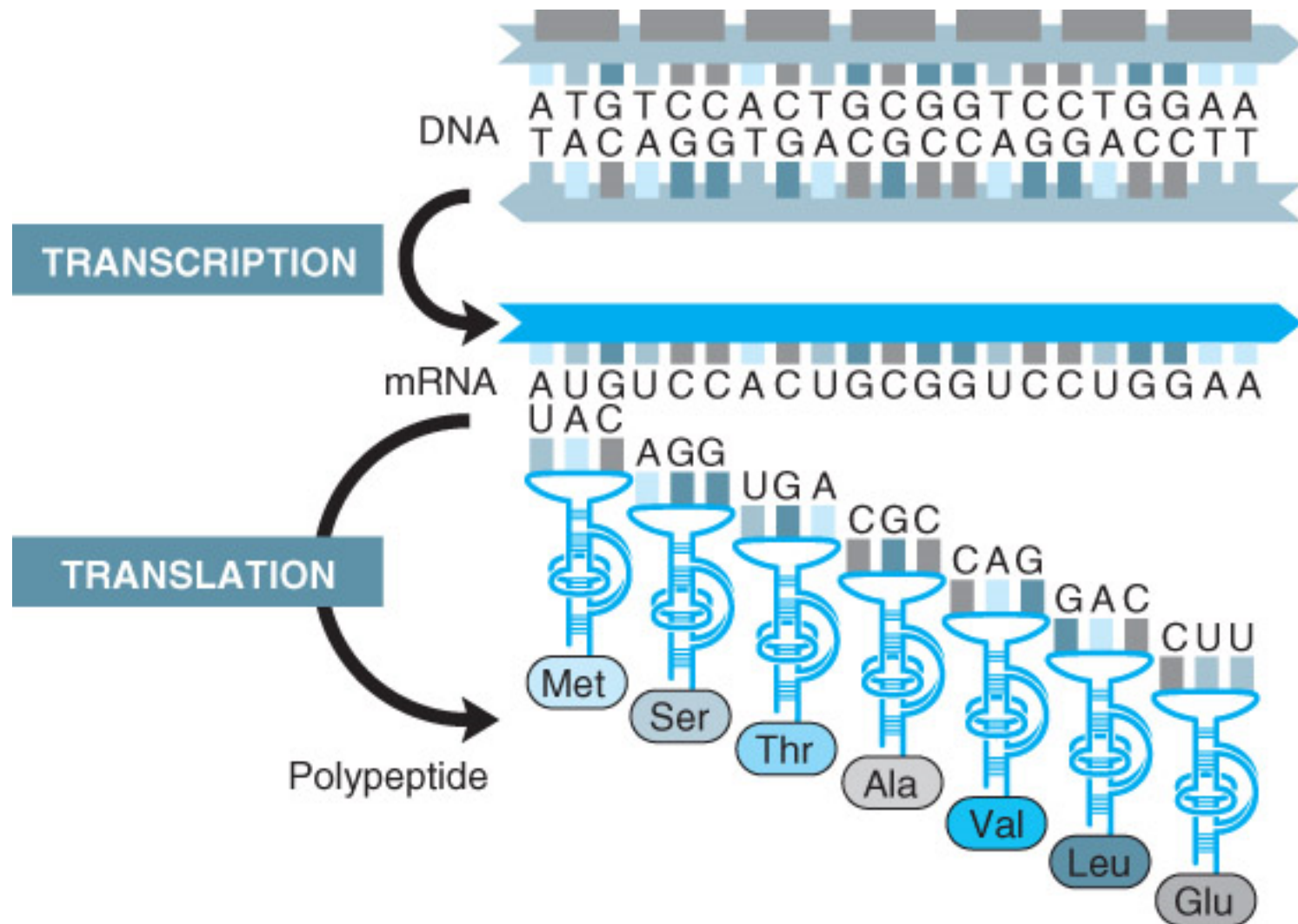


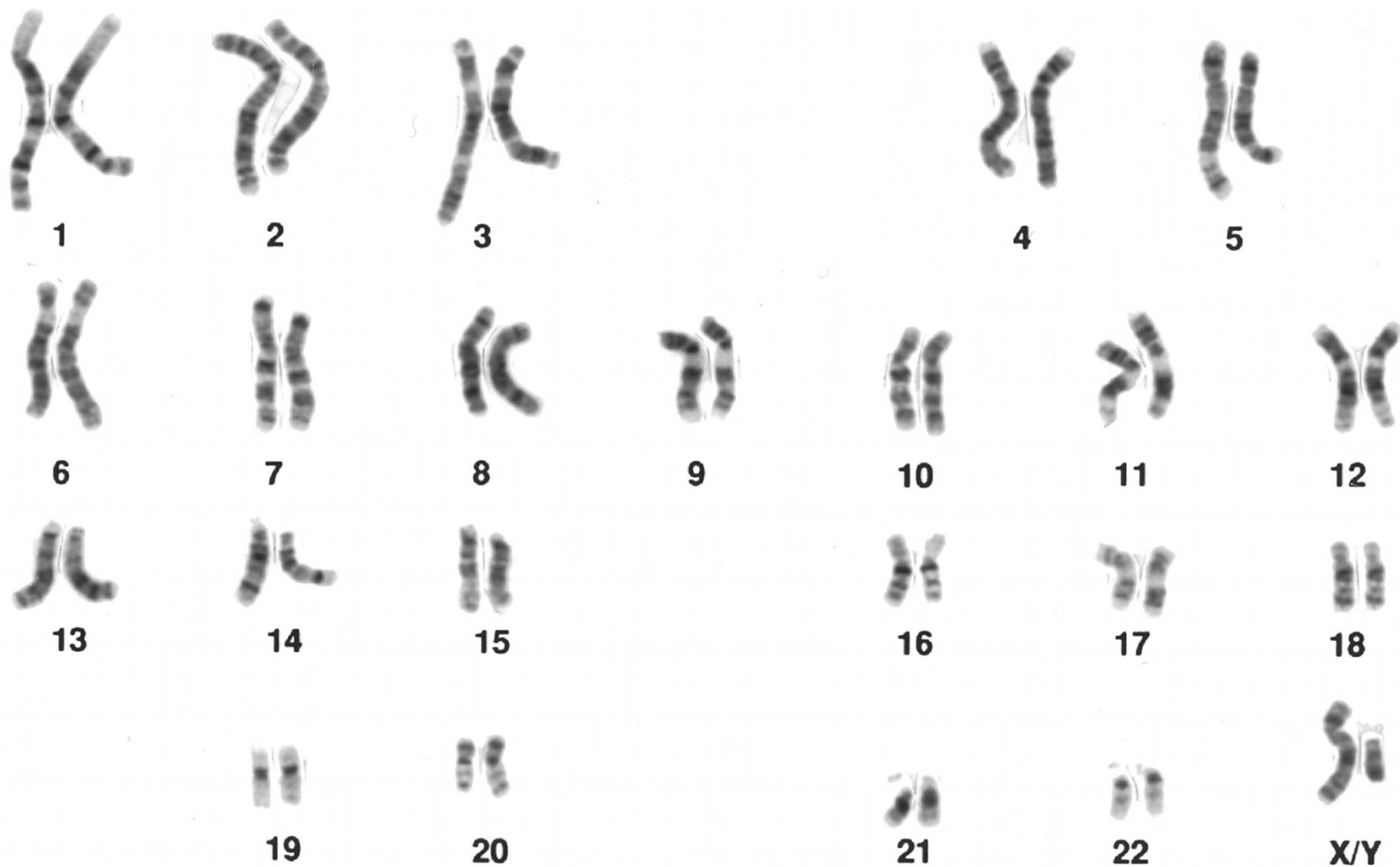
Diagram showing how a single gene encoded in DNA is transcribed and translated to produce a protein

```
>gi | 28302128 | ref | NM_000518.4 | Homo sapiens hemoglobin, beta (HBB), mRNA
ACATTTGCTTCTGACACAACCTGTGTTCACTAGCAACCTCAAACAGACACCATGGGTGCATCTGACTCCTGA
GGAGAAGTCTGCCGTTACTGCCCTGTGGGGCAAGGTGAACGTGGATGAAGTTGGTGGTGAGGCCCTGGGC
AGGCTGCTGGTGGTCTACCCTTGGACCCAGAGGTTCTTTGAGTCCTTTGGGGATCTGTCCACTCCTGATG
CTGTTATGGGCAACCCTAAGGTGAAGGCTCATGGCAAGAAAGGTCTCGGTGCCTTTAGTGATGGCCTGGC
TCACCTGGACAACCTCAAGGGCACCTTTGCCACACTGAGTGAGCTGCACTGTGACAAGCTGCACGTGGAT
CCTGAGAACTTCAGGCTCCTGGGCAACGTGCTGGTCTGTGTGCTGGCCCATCACTTTGGCAAAGAATTCA
CCCCACCAGTGCAGGCTGCCTATCAGAAAGTGGTGGCTGGTGTGGCTAATGCCCTGGCCCACAAGTATCA
CTAAGCTCGCTTTCTTGCTGTCCAATTTCTATTAAAGGTTCTTTGTTCCCTAAGTCCAACACTACTAACT
GGGGGATATTATGAAGGGCCTTGAGCATCTGGATTCTGCCTAATAAAAAACATTTATTTTCATTGC
```

The human HBB gene in FASTA format

Species 1: TAAAGACCATAGGAAATAAAGATAA
Species 2: TAACGACCAT—GGAAACAAAGATAA

Determining the similarity of two or more genes: by aligning their nucleotide sequences as well as possible; the differences due to mutation are shown in boxes



The 23 sets of chromosomes in the human genome

In Summary

