

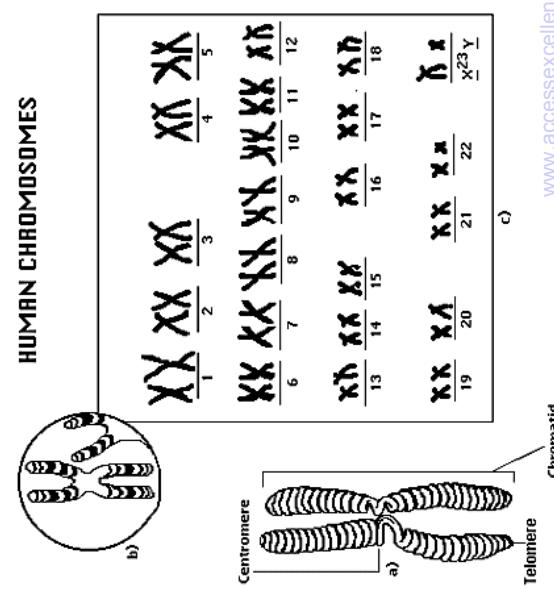
The genome

- The genome is distributed along **chromosomes**, which are made of compressed and entwined **DNA**.
- A (protein-coding) **gene** is a segment of chromosomal **DNA** that directs the synthesis of a **protein**.

The human genome

- The human genome is distributed along **23 pairs of chromosomes**
 - 22 autosomal pairs;
 - the sex chromosome pair, **XX** for females and **XY** for males.
- In each pair, one chromosome is paternally inherited, the other maternally inherited (cf. meiosis).

Chromosomes

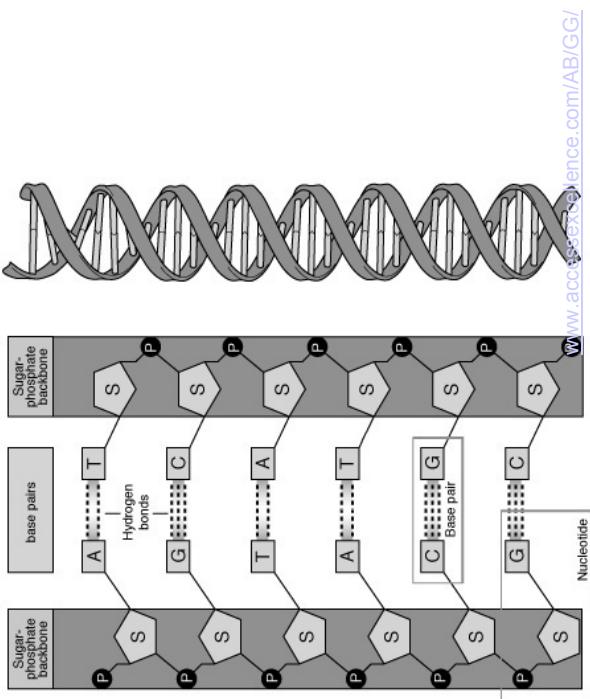


www.accessse excellence.com/AB/GG/

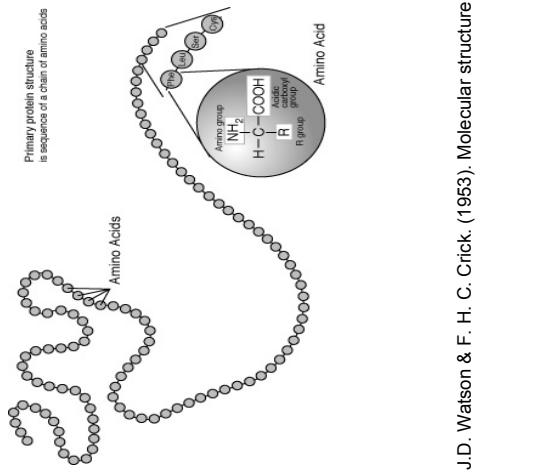
DNA structure

- A **deoxyribonucleic acid** or **DNA** molecule is a double-stranded polymer composed of four basic molecular units called nucleotides.
- Each **nucleotide** comprises
 - a phosphate group;
 - a deoxyribose sugar;
 - one of four nitrogen bases:
 - purines: **adenine (A)** and **guanine (G)**,
 - pyrimidines: **cytosine (C)** and **thymine (T)**.

DNA structure



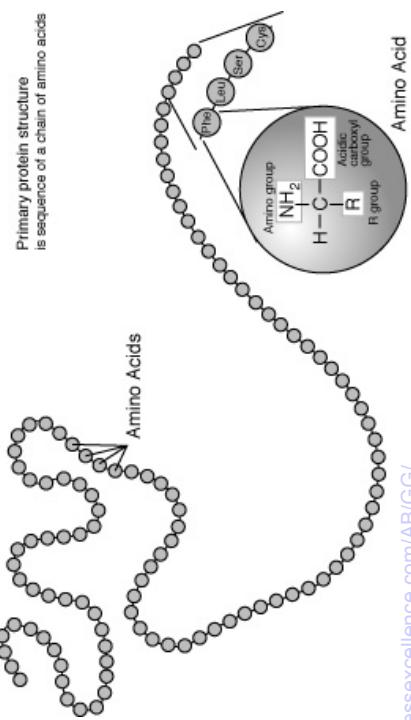
DNA replication



J.D. Watson & F. H. C. Crick. (1953). Molecular structure of Nucleic Acids. *Nature*. 171: 737-738.

"It has not escaped our notice that the specific pairing we have postulated immediately suggests a possible copying mechanism for the genetic material."

Proteins

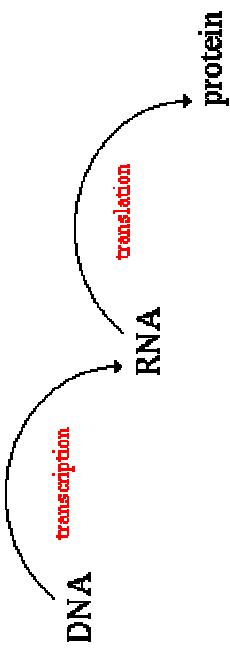


- **Proteins:** large molecules composed of one or more chains of amino acids, **polypeptides**.
- **Amino acids:** class of 20 different organic compounds containing a basic amino group ($-\text{NH}_2$) and an acidic carboxyl group ($-\text{COOH}$).
 - The order of the amino acids is determined by the **base sequence** of nucleotides in the **gene** coding for the protein.
 - E.g. hormones, enzymes, antibodies.

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Central dogma



RNA

- A **ribonucleic acid** or **RNA** molecule is a nucleic acid similar to DNA, but
 - single-stranded;
 - ribose sugar rather than deoxyribose sugar;
 - **uracil (U)** replaces thymine (T) as one of the bases.
- RNA plays an important role in protein synthesis and other chemical activities of the cell.
- Several classes of RNA molecules, including **messenger RNA (mRNA)**, transfer RNA (tRNA), ribosomal RNA (rRNA), and other small RNAs.

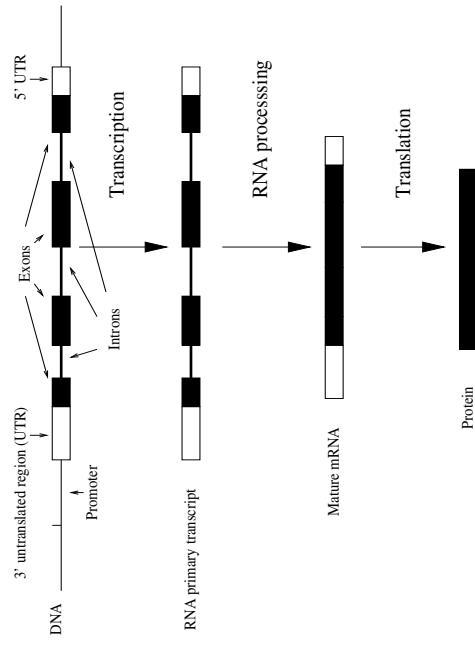
The genetic code

- DNA: sequence of **four** different nucleotides.
 - Proteins: sequence of **twenty** different amino acids.
 - The correspondence between DNA's four-letter alphabet and a protein's twenty-letter alphabet is specified by the **genetic code**, which relates nucleotide triplets or codons to **amino acids**.
- Start codon: initiation of translation (AUG, Met). Stop codons: termination of translation.
- Mapping between codons and amino acids is **many-to-one**: 64 codons but only 20 a.a.. Third base in codon is often redundant, e.g., stop codons.
-
- The diagram shows a tRNA molecule with a red "3' end" and a blue "anticodon" (UCG) at its 3' end. It is shown binding to a portion of a mRNA strand (mRNA 3') with the sequence AUAG. The 5' end of the tRNA is labeled "5'".
- The table below shows the genetic code mapping between codons (3 bases) and amino acids (1 letter).
- | 3rd base in codon | 2nd base in codon | 1st base in codon | Amino Acid |
|-------------------|-------------------|-------------------|---------------|
| U | C | A | G |
| U | C | T | Tyrosine |
| U | C | G | Cysteine |
| U | A | T | STOP |
| U | A | G | Leucine |
| C | C | A | Proline |
| C | C | T | His |
| C | C | G | Alanine |
| C | G | A | Arginine |
| C | G | T | Asparagine |
| A | A | A | Isoleucine |
| A | A | T | Threonine |
| A | A | G | Alanine |
| A | T | A | Termination |
| G | A | A | Valine |
| G | A | T | Aspartic acid |
| G | A | G | Glycine |
| G | T | A | Glutamic acid |
| G | T | G | Glycine |
- The Genetic Code**
- www.accessexcellence.com/AB/GG/

Exons and introns

- Genes comprise only about 2% of the human genome.
- The rest consists of **non-coding regions**
 - chromosomal structural integrity,
 - cell division (e.g. centromere)
 - regulatory regions: regulating when, where, and in what quantity proteins are made .
- The terms **exon** and **intron** refer to coding (translated into a protein) and non-coding DNA, respectively.

From gene to protein



Florence, 2005, Introduction

3

Differential expression

- Each cell contains a complete copy of the organism's genome.
- Cells are of many different types and states
 - E.g. blood, nerve, and skin cells, dividing cells, cancerous cells, etc.
- What makes the cells different?
 - **Differential gene expression**, i.e., **when, where, and how much each gene is expressed.**
- On average, 40% of our genes are expressed at any given time.

Alternative splicing

- There are more than 1,000,000 different human antibodies. How is this possible with only ~30,000 genes?
- **Alternative splicing** refers to the different ways of combining a gene's exons. This can produce different forms of a protein for the same gene.
- Alternative pre-mRNA splicing is an important mechanism for regulating gene expression in higher eukaryotes.
- E.g. in humans, it is estimated that approximately 30% of the genes are subject to alternative splicing.

Alleles: Different versions of the genetic code at a particular locus

- **Types of Alleles (2 of many)**
 - **Single Nucleotide Polymorphism (SNP)**
 - Allele 1: ... TCGTGCAGGTATTAGC...
 - Allele 2: ... TCGTGTCAAGGTATTAGC...
 - **Short Tandem Repeat Polymorphism (STRP)**
 - Allele 1: ... AACGTC **GAGAGACAGGGTATTAGC...**
 - Allele 2: ... AACGTC **GAGAGAGACAGGGTATTAGC...**
 - Allele 3: ... AACGTC **GAGAGAGAGACAGGGTATTAGC...**
 - ...
 - The term **allele** refer to a DNA sequence variant for a particular gene.
 - The term **genotype** refers to DNA variants at a gene or set of genes for a given individual.
 - The term **phenotype** refers to an observable trait or set of traits in an individual.
E.g. blood pressure, disease status.
- Disease-Gene vs. Disease Related Allele(s)**
- E.g., the PAH Gene codes for the enzyme **Phenylalanine Hydroxylase**

Genotype and phenotype

Microsatellite Markers

Single Nucleotide Polymorphism (SNP)

Allele 1: ... TCGTGCAGGTATTAGC...
Allele 2: ... TCGTGTCAAGGTATTAGC...

Alleles: Different versions of the genetic code at a particular locus

Short Tandem Repeat Polymorphisms (STRPs)

Repeating di-, tri-, tetra-, or penta- nucleotide units

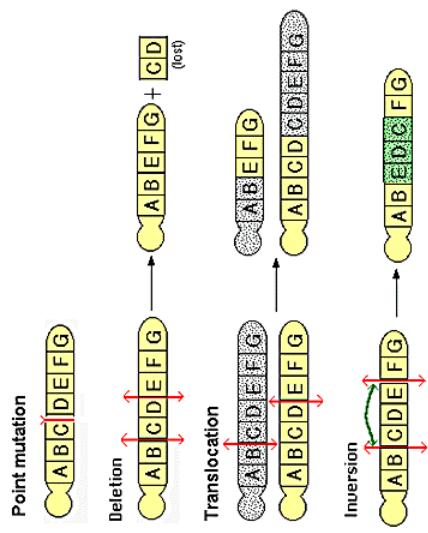
E.g., a $(CA)_n$ di-nucleotide repeat

Allele 1: ... AACGTC**CCACA...CAGAGGTATTAGC...**
Allele 2: ... AACGTC**CCACA...CACAGGGTATTAGC...**
Allele 3: ... AACGTC**CCACA...CACACAGGGTATTAGC...**
...

Genetic diseases

- **Genetic disease:** the susceptibility or resistance to the disease is influenced by variants in the DNA sequence (alleles of the disease genes) that are transmitted from parent to offspring.
 - E.g. Alzheimer's disease, asthma, cancer, color blindness, diabetes, hemophilia, sickle cell anemia, etc.

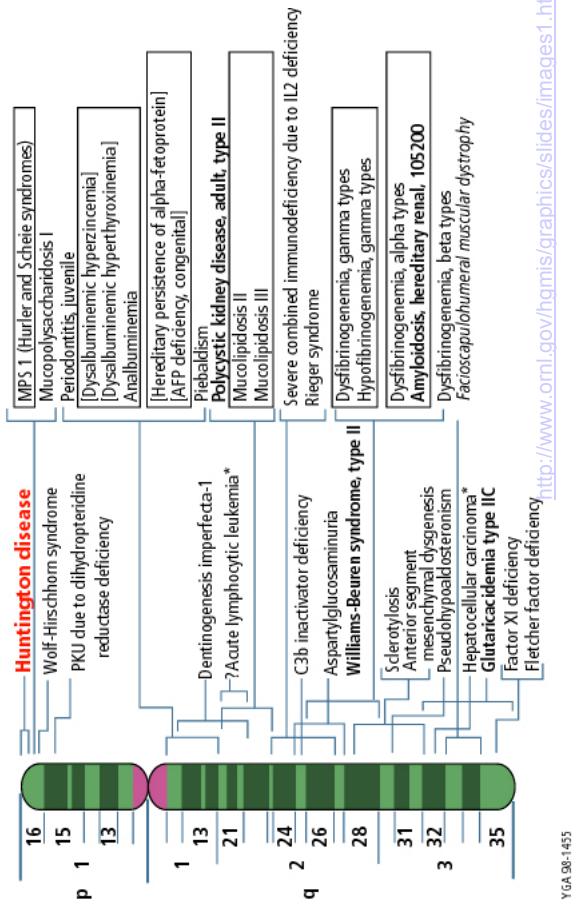
Chromosome mutations



From DNA to proteins



Chromosome 4



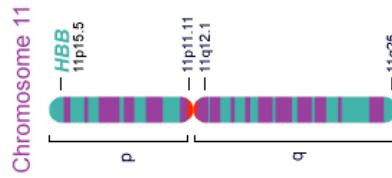
Mendelian traits

- Usually influenced by a single gene.
- Usually a simple deterministic relationship between genotype and phenotype:
0/1 penetrances.
- E.g.
 - Cystic fibrosis (autosomal recessive);
 - Hemophilia (X-linked recessive);
 - Huntington's disease (autosomal dominant);
 - Sickle cell anemia (autosomal recessive).

Sickle cell anemia

- **Phenotype:** Abnormal type of hemoglobin, HbS.
Hemoglobin is a protein found in red blood cells and involved in oxygen transport. Individuals with the disease tire easily and often develop heart failure.
- **Molecular pathology:** Point mutation in the **hemoglobin beta gene** (HBB) found on the short arm of chromosome 11, **11p15.5**.
The amino acid Glu (Glutamic acid) is replaced by Val (Valine) in the HbA beta-chain
 - mutant HbS molecule;
 - sickle-shaped red blood cells.

Sickle cell anemia



Sickle cell anemia

HBB Sequence in Normal Adult Hemoglobin (Hb A):						
Nucleotide	CTG	ACT	CCT	GAG	GAG	AAG
Amino Acid	Leu	Thr	Pro	Glu	Glu	Lys
	1	3	6	9		

HBB Sequence in Mutant Adult Hemoglobin (Hb S):						
Nucleotide	CTG	ACT	CCT	GTG	GAG	AAG
Amino Acid	Leu	Thr	Pro	Val	Glu	Lys
	1	3	6	9		

<http://www.ncbi.nlm.nih.gov/chromosomes/posters/hb/hbb.html>

Sickle cell anemia

- **Mode of inheritance:** Autosomal recessive (gene on chr. 11).
An individual is affected iff he/she has two copies of the mutant allele for HBB.

- **Prevalence**

- Overall US population: 1/72,000.
 - Higher among African Americans: 1/500 are affected and 1/12 are heterozygous carriers.

SCA and malaria

- The high incidence of SCA in certain populations is related to another disease: **malaria**.
- Heterozygotes for SCA are resistant to malaria.
- Malaria is caused by four protozoan parasites of the genus *Plasmodium*.
- *P. falciparum* is the most widespread and dangerous.
- The parasites are transmitted from person to person by female mosquito. They multiply in infected individuals and break down red blood cells.
- Worldwide incidence: each year 300-500 million cases, 1 million deaths (mainly sub-Saharan Africa).
- Symptoms: fever, headaches, etc.

Hemophilia A

- **Phenotype:** Clotting factor VIII is reduced or missing → inability to properly form blood clots.

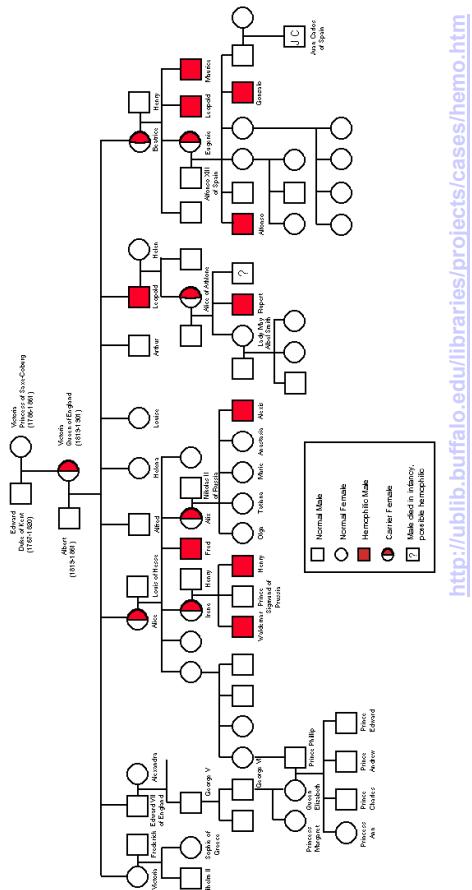
- **Mode of inheritance:** X-linked recessive.

- Males are affected iff they have one copy of the mutant allele;
- Females are affected iff they have two copies of the mutant allele.
- **Prevalence:** 1/5,000 in males, much lower in females.

Hemophilia A

The royal disease

- Multiple genes, multiple alleles of a gene.
- No clear deterministic relationship between genotype and phenotype:
 - unknown penetrances, usually not 0/1;
 - incomplete penetrances and phenocopies.
 - Unknown population genetic model, non-random mating.
 - Genetic heterogeneity.
 - Diagnostic uncertainties (e.g., late onset).
 - Gene-gene interactions.
 - Gene-environment interactions.
- E.g. Alzheimer's disease, asthma, diabetes, hypertension, etc.



Type 1 diabetes

- Type 1 or Insulin Dependent Diabetes Mellitus (IDDM).
- **Phenotype:** the body does not produce any insulin, insulin is needed to convert sugar into energy.
- **Autoimmune disease:** T-cell destruction of pancreatic beta-cells which produce insulin.
- **Prevalence:** 300,000-500,000 in the US.
- **Mode of inheritance: complex.**
 - 20 or so genes (most yet to be identified) are believed to influence susceptibility to IDDM -- the exact combination of genes differing with the individual.
 - The major histocompatibility complex (MHC) of genes (called HLA in humans) may account for about half of the total genetic risk of developing IDDM.
 - Environmental factors.

Type 1 diabetes