

DNA and Inheritance

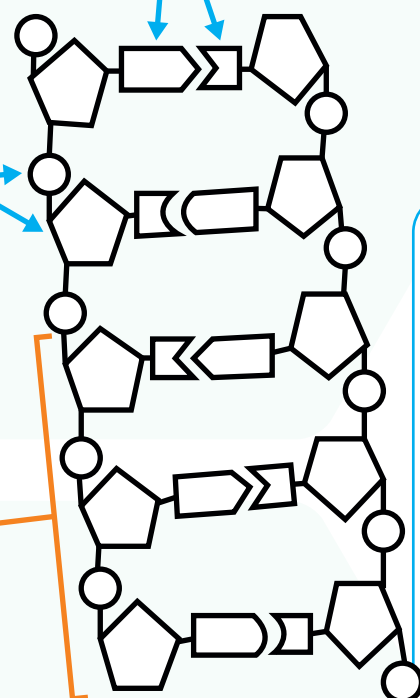
DNA structure

4 types of base connect the chains. The bases show **complementary** base pairing. **Guanine (G)** only pairs with **Cytosine (C)** and **Adenine (A)** only pairs with **Thymine (T)**.

Diagram key



DNA is constructed from **2 long chains of alternating sugar and phosphate**.



DNA is a **code**.

Reading down one of the chains the order of bases forms a triplet code. E.g. TAG

The triplet of bases will code for a particular **amino acid**.

Chains of amino acids form **proteins**. This is how DNA codes for the structure of different proteins.

3 bases form a **triplet code**: In this case TAG.

Genetic crosses - **Genes** code for characteristics. Genes can have different versions called **alleles**. These alleles are inherited through **sexual reproduction**, one from each parent so they occur in **pairs**.

Alleles can be represented by letters. This is the **Genotype**. A dominant allele is represented by a capital letter, this allele is shown in the **phenotype** (how the organism looks) whenever present.

We can use the **Punnet square** to the right to estimate the outcome of genetic crosses. In this example the **gametes** (sex cells) are shown at the top and side. The yellow boxes represent the alleles found in the sperm. As each sperm only contains 1 of a pair of alleles it contains either B or b. The green boxes represent the alleles in the egg cell, either B or b.

	B	b
B	BB	Bb
b	Bb	bb

These parents were **heterozygous**. This means that they have 2 different alleles of the same gene.

The **recessive** allele is represented by a lower-case letter. This allele must be in a **homozygous** pair (both alleles are the same i.e. bb) to be shown in the phenotype.

The punnet square shows us that:

- 3:1 ratio of dominant to recessives traits shown in the phenotype of the offspring in this cross.
- 1:2:1 ratio of homozygous dominant to heterozygous to homozygous recessive genotypes in this cross.

Genetic profiling

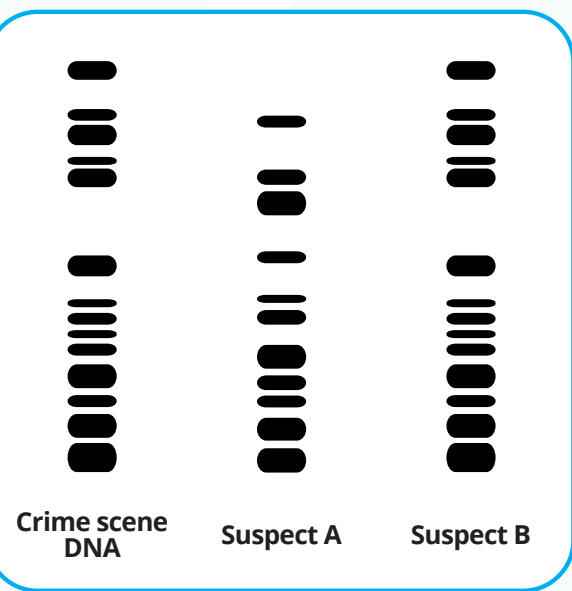
A genetic profile can be used to **compare similarities** between DNA samples.

A DNA sample is cut into **short pieces** which are then **separated into bands**.

This technique can be used in:

- criminal cases
- paternity cases
- comparing species for classification purposes.

Another benefit is **identifying genes associated with disease**. However, there are **ethical concerns** with this technology.



Genetic modification

Genetic modification allows genetic material from one organism to be transferred into the DNA of another organism.

Advantages: Genes for **disease resistance** can be transferred to crop plants to **increase yield**.

Herbicide resistant genes can also increase yield as herbicides can be used to kill competing plants (weeds).

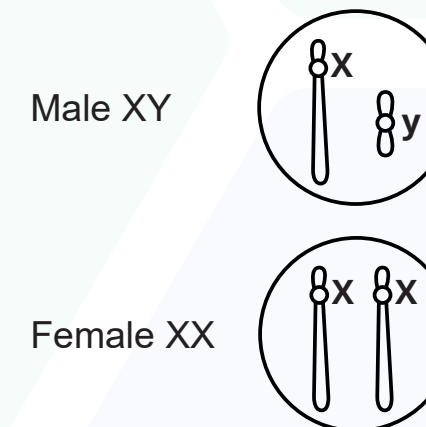
Disadvantages: Creation of super weeds if the herbicide resistance genes are taken up by weed species.

Unknown long-term effects of modifying genomes.

Unknown health effects of eating modified organisms.

Sex determination

Sex is determined in humans by chromosome **pair 23**. These chromosomes are labelled as:



The inheritance of these chromosomes can be shown using a Punnet square.

This shows that in each fertilisation there is a **50%** chance the offspring will be male or female.

	X	X
X	XX	XX
Y	XY	XY