

Module 5: Heredity

5.1: Reproduction

How does reproduction ensure the continuity of species?

Reproduction ensures the continuity of species by ensuring the production of fertile offspring with genetic diversity for adaptability, through inheritance of genetic material.

Organism → Survives → Growth → Reproduction → Inheritance of genes → Continuity

[5.1.1] Mechanisms of Reproduction

ASEXUAL AND SEXUAL REPRODUCTION

Reproduction: The process of creating offspring, either sexually or asexually.

- **Sexual reproduction:** Reproduction which involves the combination of gametes, so that the offspring are genetically different to the parents.
- **Asexual reproduction:** Reproduction which does not involve the combination of gametes, so the offspring are identical to the parent.

| | Advantages | Disadvantages |
|-----------------------------|---|---|
| Sexual reproduction | <ul style="list-style-type: none">✓ Variations in the population lead to continuity of species✓ There are more variations, so species are better able to adapt to changing selection pressures✓ It is less likely that a selection pressure will remove all of the species | <ul style="list-style-type: none">✗ Requires more time and energy to carry out✗ Requires a mating partner, can not be done individually✗ Fewer offspring produced |
| Asexual reproduction | <ul style="list-style-type: none">✓ Allows a species to rapidly populate an environment✓ No requirement of mates✓ Quick and not energy intensive, so it can be carried out when the parent organism is under stress✓ There is no requirement to care for the offspring | <ul style="list-style-type: none">✗ Lack of diversity between species of a population. Variations, and therefore adaptations, are limited✗ Reduces ability to adapt to selection pressures✗ May lead to extinction of the population, as one selection pressure can wipe out all species, as there is no variation between them |

INTERNAL AND EXTERNAL FERTILISATION

Fertilisation: The fusion of gametes in sexual reproduction.

→ **Internal fertilisation:** Fertilisation where the combination of gametes occurs within the female body.

→ **External fertilisation:** Fertilisation where the combination of gametes occurs in the external environment.

| | | |
|------------------------|---|---|
| INTERNAL FERTILISATION | <ul style="list-style-type: none"> ✓ Increased likelihood of fertilisation as the gametes are in close proximity ✓ Less gametes produced, which reduces energy expenditure ✓ Offspring have a greater chance of survival as they are protected from the environment | <ul style="list-style-type: none"> ✗ Fewer offspring are produced over more time ✗ Mating rituals are time consuming and postpone copulation ✗ Higher risk of sexually transmitted diseases/ infections ✗ Fertilisation, gestation and parental care demand energy from the parent ✗ Breeding has to be paused for gestation (pregnancy) period |
| EXTERNAL FERTILISATION | <ul style="list-style-type: none"> ✓ A greater number of offspring can be produced ✓ The behavioural process is simpler ✓ Prevention of dehydration as it occurs in an aquatic environment ✓ Reproduction can continue without pause for gestation ✓ Offspring are dispersed, meaning less competition ✓ Many gametes released, increasing chance of fertilisation ✓ No energy spent on parental care | <ul style="list-style-type: none"> ✗ A large number of gametes must be produced. This requires a lot of energy ✗ Can only occur in aquatic environments ✗ The gametes are more susceptible to the risks of the open environment ✗ External environmental conditions are unconstant, which may impact fertilisation ✗ Absence of parental care can increase risk of predation and disease to offspring. |

Comparison of internal and external fertilisation

| Characteristic | Internal | Both | External |
|-----------------------------------|--|--|---|
| Gametes Released | Large number of sperm, Small number of eggs | | Large number of both gametes |
| Fertilisation | Occurs in female reproductive system | Sperm fertilises egg | Occurs in external aquatic environment |
| Synchronisation of gamete release | Sperm is continuously produced, Eggs are produced over a cycle | | Male and female gametes released simultaneously |
| Chance of fertilisation | High | Chance increases with proximity of gametes | Low |
| Result of fertilisation | Zygote (internal development) | The development of a zygote requires a watery medium | Zygote (external development) |
| No. of offspring | Small (but survival rate is high) | Both determined by: <ul style="list-style-type: none"> - No. of gametes produced - Survival rate | High (but survival rate is low) |
| Breeding frequency | Low, seasonal | Breeding frequency relies on environmental conditions | High |
| Parental care | Yes <ul style="list-style-type: none"> - Care of zygote during development - Care of offspring after birth | | None |

DEVELOPMENT

Development: The growth of the zygote.

When fertilised **internally**, development can be:

- **Oviparous:** External within an egg.
- **Viviparous:** Internal within the mother.
- **Ovoviviparous:** Internal within an egg within the mother.

| | | |
|--|---|---|
| Unisexual reproductive system Separate male and female individuals with separate reproductive systems. | ✓ Less energy required to maintain 1 set of reproductive organs | ✗ Mating rituals required, which take time |
| Bisexual/ hermaphrodite reproductive system Organisms have both male and female reproductive organs. | ✓ Organism doesn't need a partner of the opposite sex to reproduce ✓ Beneficial for organisms that are sedentary | ✗ A larger amount of energy is required to maintain 2 sets of reproductive organs |

SEXUAL REPRODUCTION IN ANIMALS

Gametes are produced by the parents by meiosis
↓
Each gamete is **haploid** ($\frac{1}{2}$ the total number of chromosomes)
↓
The sperm (male gamete) fertilises the ovum/ egg (female gamete)
↓
The fusion of gametes results in a **diploid zygote** (has a complete set of chromosomes). The zygote has a unique combination of genetic material from both parents.

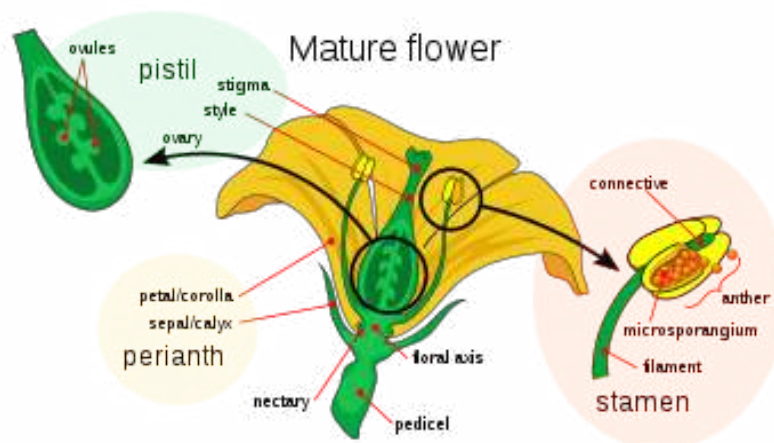
Animal fertilisation can be internal or external.

IN PLANTS

Plants are bisexual. In angiosperms, the:

MALE organs → Stamen (filament + anther)

FEMALE organs → Pistil (stigma + style + ovary)



Pollination

Pollen (the male gamete) is produced in the stamen, and held up at the anther

↓
The pollen is removed from the anther by the wind or by insects. It is transferred to the pistil of the same flower (self-pollination) or another flower (cross-pollination)
↳ [From the same plant or another one].

Fertilisation

The pollen falls on the stigma, and a pollen tube grows through the style to the ovary

↓
The pollen migrates through pollen tubes, to the ovules

↓
The pollen fertilises the ovules

Development

The fertilised ovule develops into a seed, containing the embryo



The seed protects the embryo



The fruit around it provides it with nutrients



Germination

When the fruit is removed, the seeds within it grow separate from the parent plant(s)



The embryo lies dormant



If on suitable soil, germination begins



The embryo puts out its:

- Radical (first root) for absorption of water and nutrients
- Plumule (young stem) which develops leaves for photosynthesis



The seed germinates into a seedling

Methods of pollination

| METHOD | ADAPTATIONS FOR METHOD |
|--|--|
| <ul style="list-style-type: none">• Wind | <ul style="list-style-type: none">- Protruding stigma to catch more pollen- Light-weight, small pollen- Large pollen production- Flower is not colourful to conserve energy, as it doesn't need to attract pollinators |
| <ul style="list-style-type: none">• Animals<ul style="list-style-type: none">○ Insects○ Birds | <ul style="list-style-type: none">- Colourful petals to attract pollinators- UV markings which insects are attracted to- Scent for attraction- Flower produces nectar to attract pollinators- Stigma and anthers within flower, energy not wasted in growing tall- Pollen is sticky, to attach to the pollinator and be dropped elsewhere |

Methods of seed dispersal

| METHOD | ADAPTATIONS FOR METHOD |
|---|---|
| <ul style="list-style-type: none"> • Wind | <ul style="list-style-type: none"> - Seeds are very light-weight |
| <ul style="list-style-type: none"> • Animals <ul style="list-style-type: none"> ○ Insects ○ Birds | <ul style="list-style-type: none"> - The seeds are <i>fruit bearing</i>, meaning that it is eaten by animals - Once the fruit is eaten, the seed is either <ul style="list-style-type: none"> - Left to germinate - Passes through the digestive system, excreted and germinates |
| <ul style="list-style-type: none"> • Self-dispersal | <ul style="list-style-type: none"> - The plant has an explosive mechanism, which propels the seed away from the mother plant - Propelling the seed to a different area reduces the competition for resources in one area, giving the offspring a greater chance of survival |

IN PROTISTS

Haploid (n) protists

Two haploid protist cells combine, and genetic material is combined in a new, fused nucleus. This produces a diploid zygote



The zygote undergoes meiosis to produce 4 haploid daughter cells

Diploid ($2n$) protists

Adult cells undergo meiosis, producing 4 haploid gametes



Gametes fuse during fertilisation to form a diploid zygote



The diploid zygote will grow to maturity

IN FUNGI

Mitosis splits a diploid cell into haploid two daughter cells, which each contain a single set of chromosomes



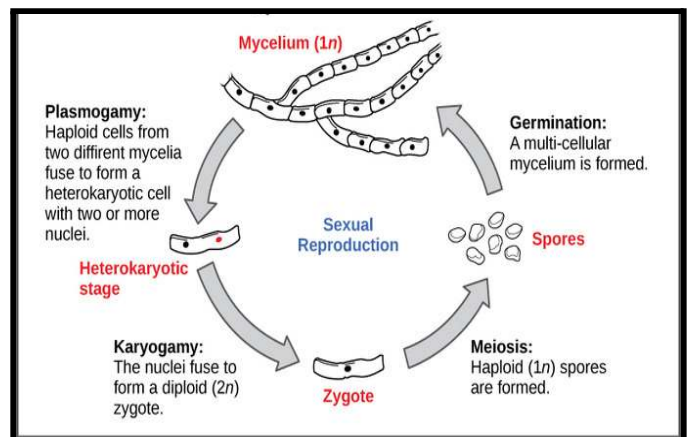
PLASMOLOGY: Two genetically different cells fuse together, combining their contents. This brings together two compatible haploid nuclei



KARYOGAMY: The two haploid nuclei fuse into one diploid nucleus. This forms a zygote cell



MEIOSIS: The zygote undergoes meiosis which produces haploid spores. These spores are distributed into the environment



ASEXUAL REPRODUCTION IN PLANTS

Vegetative propagation

Method of asexual reproduction in plants, done by **vegetative organs**, which are modified parts of the plant.

Perennating organs: Underground organs which store food.

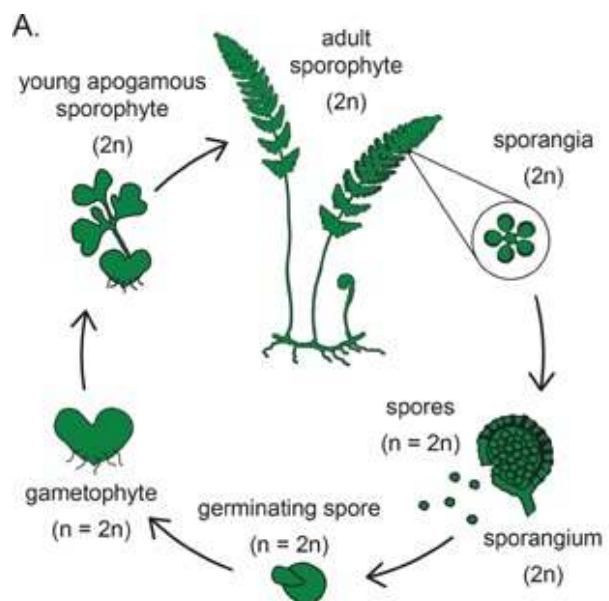
| Modified part | Vegetative organ | Perrenating? |
|---------------|---|--------------|
| Stem | - Runners Underground stems that connect plants. | NO |
| | - Rhizomes Horizontal stems that store nutrients. | YES |
| | - Tubers Develop from swollen regions of a stem, or from buds/ eyes. | NO |
| | - Bulbs Scales of bulb store food. New plants can grow from the bulb. | YES |
| Roots | - Suckers Vertical growths that can resprout from underground. | NO |

Apomixis (process)

Generative tissue gives rise to plantlets asexually.

Generative tissue includes:

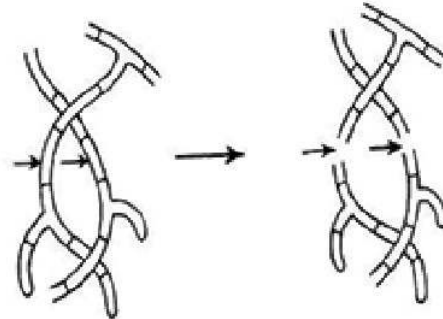
- Unfertilised ovules
- Specialised leaf tissue.



IN FUNGI

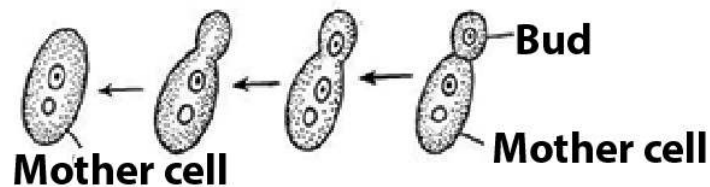
Fragmentation

Pieces of the fungal colony can break off and become separate colonies.



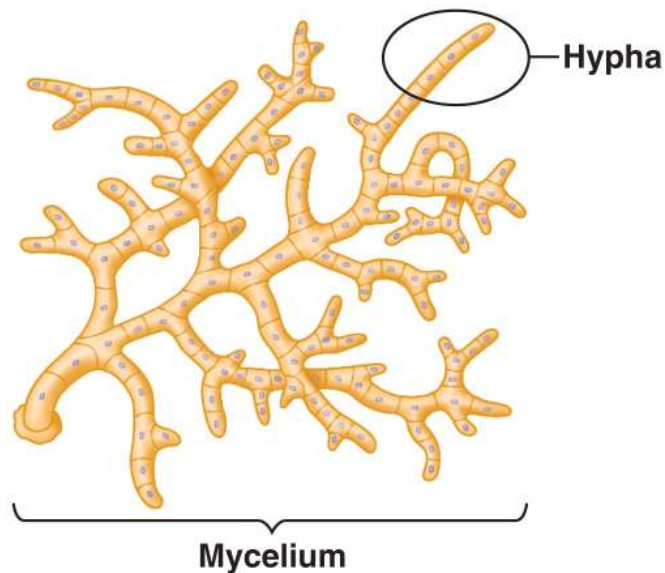
Budding

The nucleus of a fungal cell divides, and splits off from the rest of the cell by cytokinesis.



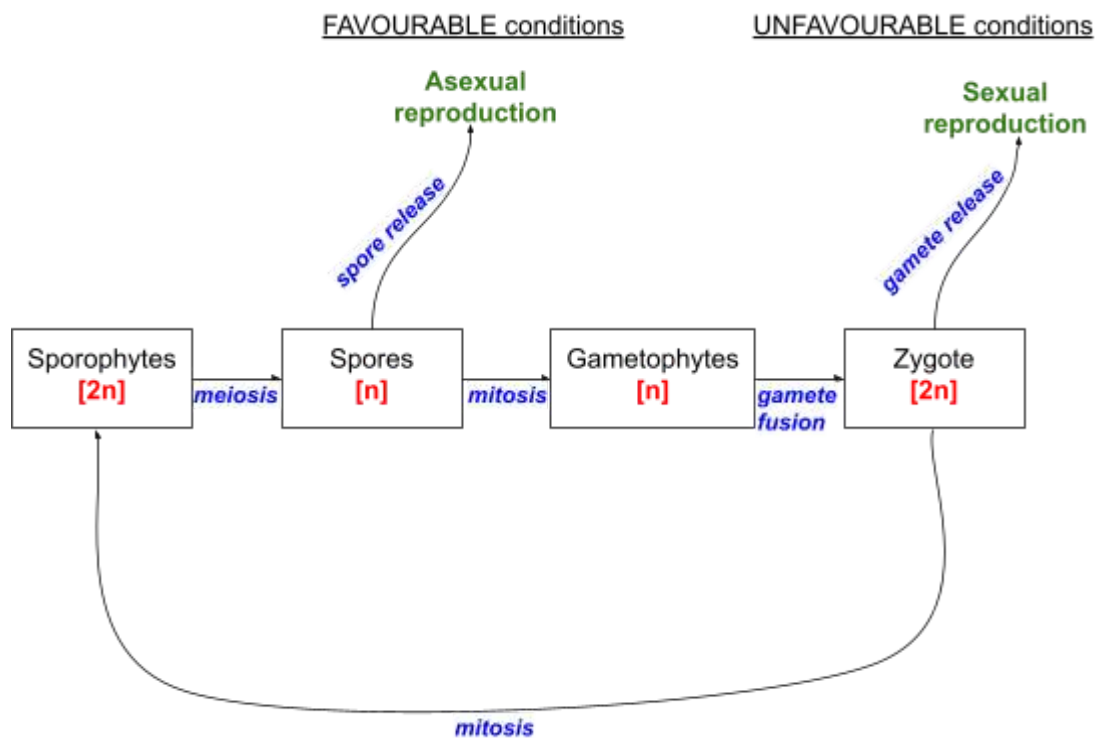
Complete budding: Separation → new cell

Incomplete budding: Bud stays attached to the mother → Keeps budding
→ hyphae → mycelium



Spores

Alternation of generation.



Fungi can reproduce both sexually and asexually within one lifetime.

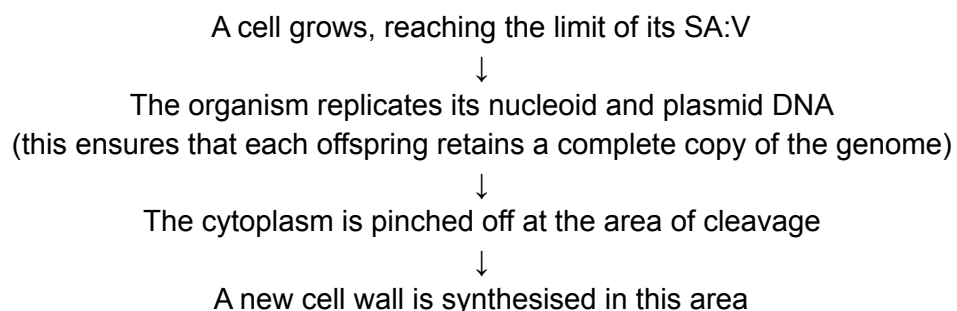
1. Haploid spores are generated from diploid sporophytes through *meiosis*
2. If the conditions are favorable, these spores will be released into the environment
→ *Asexual reproduction*
3. If the conditions are NOT favourable, the spores will form into haploid gametophytes
 - a. Both male and female gametes are produced
4. Fusion of these gametes (fertilisation) produces *zygotes*, which are released into the environment
→ *Sexual reproduction*

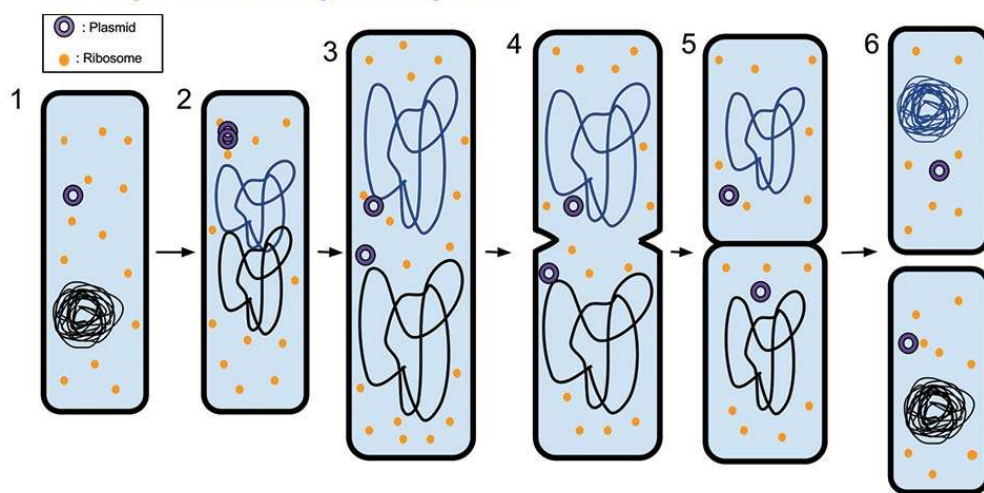
These zygotes can duplicate via mitosis, producing more sporophytes

IN BACTERIA

Binary fission

A diploid ($2n$) mother cell splits into two diploid daughter cells which are genetically identical.





IN PROTISTS

Binary fission

In the same way as bacteria, but without a cell wall.

Budding

Only a small group of amoeba called **sarcodines** can bud.

A small outgrowth develops from the body of the parent sarcodine



The offspring develop from this outgrowth and *completely separates* from the mother
(no partial separation)

[5.1.2] Fertilisation, Implantation, Pregnancy and Birth in Mammals

CONTINUITY OF SPECIES

Reproductive mechanisms ensure the *success* of producing offspring and nurturing them to reproductive maturity. This is so they can in turn **reproduce and ensure the continuity of species through generations.**

| <u>Stage of development</u> | <u>Continuity of species</u> |
|-----------------------------|---|
| • Internal fertilisation | Ensures gamete fusion |
| • Implantation | Protects embryo |
| • Pregnancy | Provides embryo with nutrients |
| • Birth | Occurs after the foetus is fully developed to ensure survival |
| • Lactation | Sustains born offspring until it reaches reproductive maturity. |

SEX HORMONES

Stages of sexual reproduction are regulated by hormones, ensuring reproductive success leading to the continuity of species.

They are responsible for:

- The development of reproductive organs
- The development of secondary sexual characteristics

In males

- Deepened voice
- Thicker body hair, and more of it
- Increased size & thickness of muscles and bones

In females

- Larger breasts
- Widening hips
- Growth of pubic hair

Types of sex hormones:

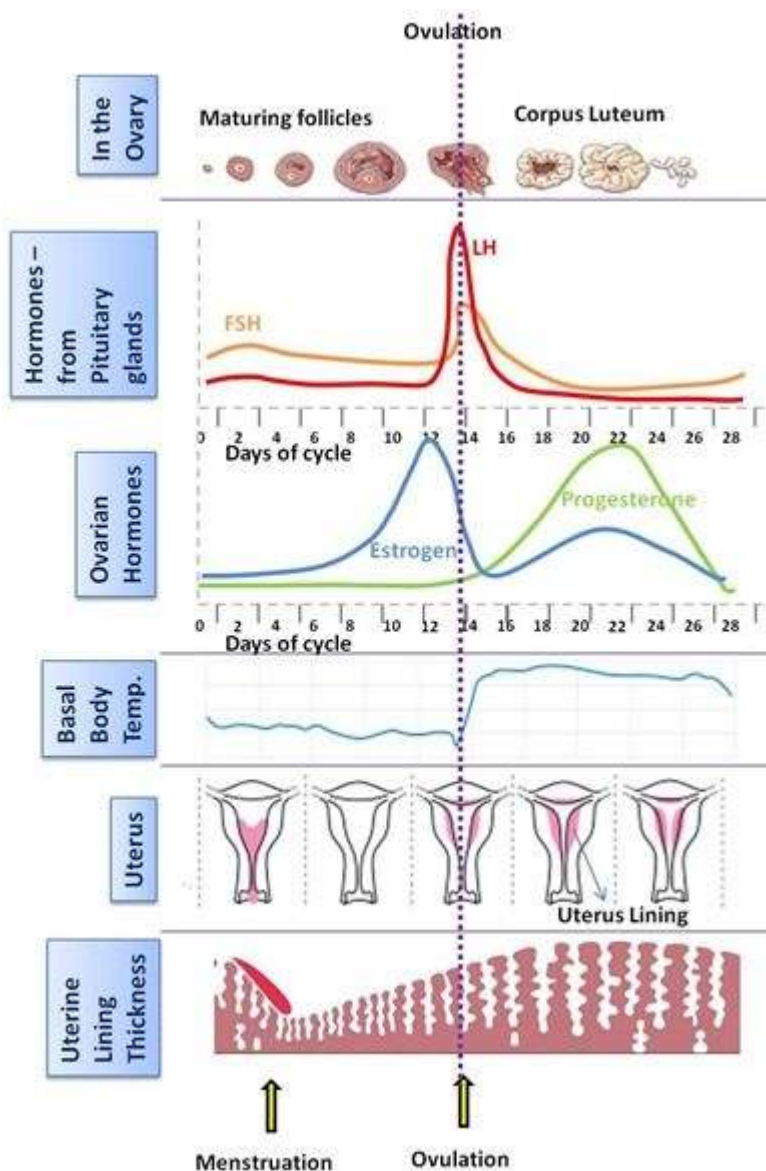
| | |
|--------------|---|
| ANDROGENS | <i>Main male hormone, but also present in females.</i> The main androgen in males is called <u>testosterone</u> , produced by the testes. Androgen is responsible for the development of sperm, and secondary sexual characteristics in males. |
| OESTROGEN | <i>Main female hormone, but also present in males.</i> Involved with the development of the female reproductive system, and secondary sexual characteristics in females. |
| PROGESTERONE | <i>Only in females.</i> Ensures that the thickness of the endometrium is maintained during pregnancy. |

MENSTRUATION AND OVULATION

Hormones involved

| Hormone | Function | Location produced |
|---------------------------------|---|-------------------------|
| 1. Follicle stimulating hormone | Maturation of follicles | Pituitary gland (brain) |
| 2. Luteinising hormone | Final maturation of follicles Development of corpus luteum | |
| 3. Oestrogen | Development of sex organs | Ovaries |
| 4. Progesterone | Controls pregnancy | |

The process



FSH rises, causing follicle cells to develop

↓

FSH boosts oestrogen production (positive feedback loop)

↓

Graafian follicle released during ovulation

↓

Oestrogen stimulates LH secretion

↓

LH inhibits FSH and oestrogen production

↓

LH causes dominant follicle to rupture and release egg

↓

Follicle develops into corpus luteum

↓

Corpus luteum produces progesterone and oestrogen

↓

LH and FSH production inhibited by progesterone. Prevents 2nd egg being released

↓

Endometrium thickens

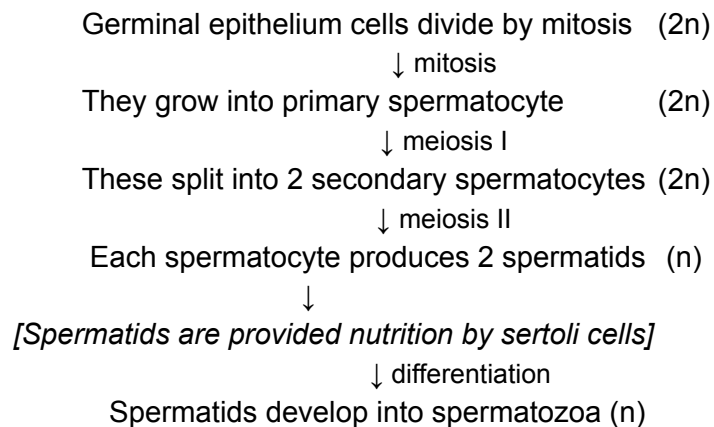
LH and FSH spike when follicle ruptures.
 Progesterone spikes when the corpus luteum is largest.
 Oestrogen spikes at both these times.

| | |
|---|---|
| <p><u>If fertilised</u></p> <p>Zygote implanted into endometrium</p> <p>↓</p> <p>Zygote produces Human Chorionic Gonadotropin (hCG)</p> <p>↓</p> <p>hCG maintains corpus luteum to produce progesterone for 12 wks</p> <p>↓</p> <p>The placenta develops fully and begins secreting progesterone</p> <p>↓</p> <p>Corpus luteum degrades, the placenta maintains the pregnancy</p> | <p><u>If NOT fertilised</u></p> <p>Corpus luteum degenerates</p> <p>↓</p> <p>Oestrogen and progesterone production stopped</p> <p>↓</p> <p>Endometrium is shed (menstrual flow)</p> <p>↓</p> <p>LH and FSH produced again</p> <p>↓</p> <p>Cycle repeats</p> |
|---|---|

GAMETOGENESIS

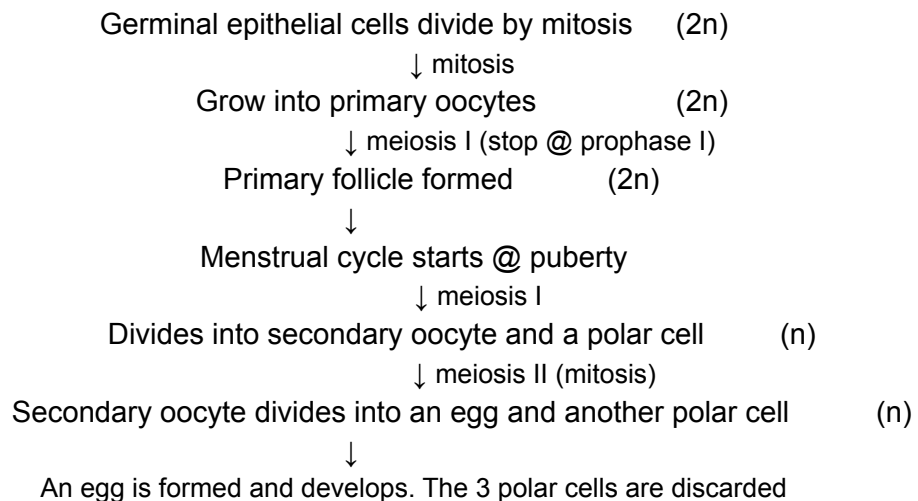
The formation of gametes from germinal epithelial cells.

Spermatogenesis



A total of 4 spermatozoa are produced in this process.

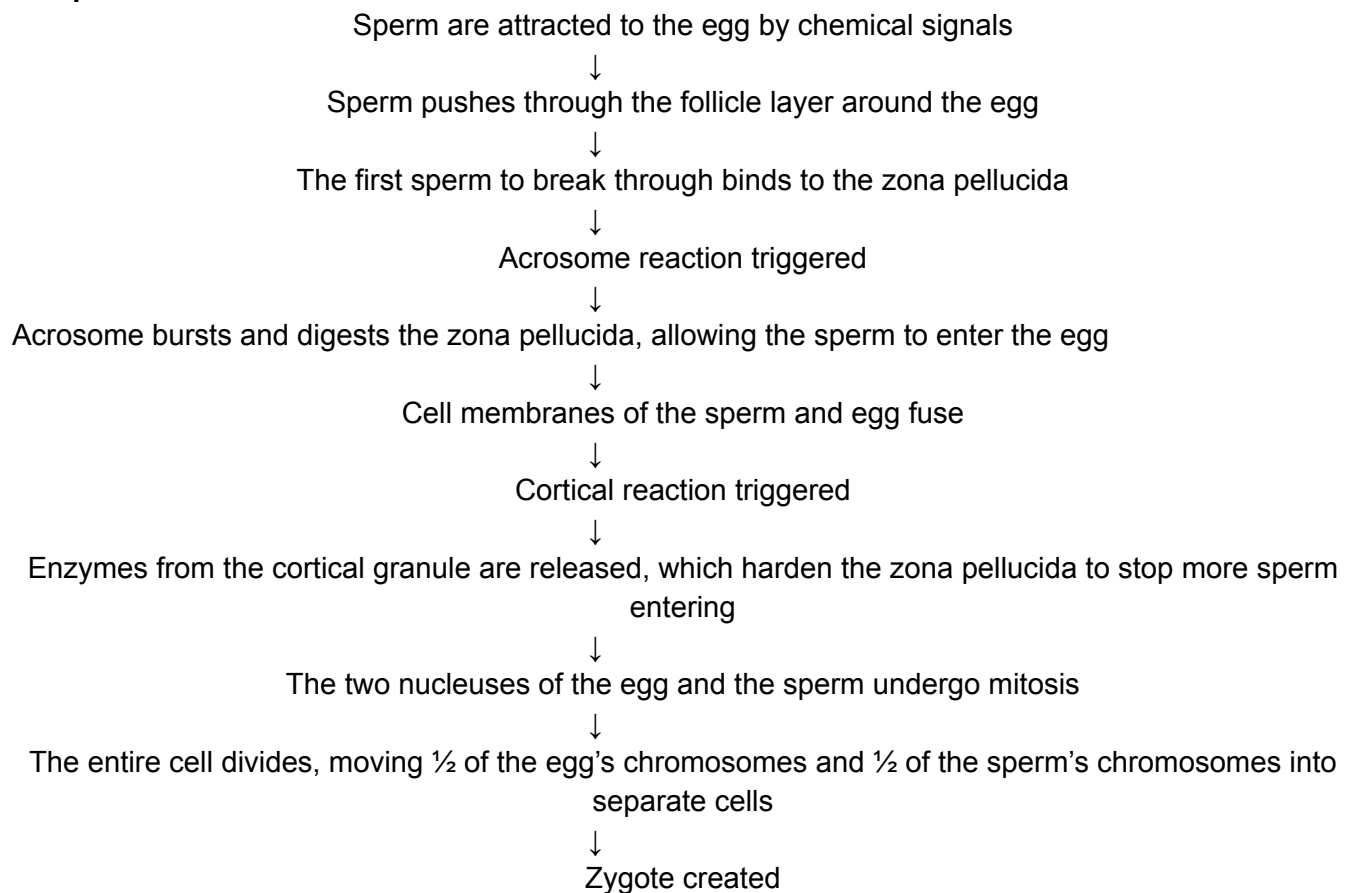
Oogenesis



FERTILISATION

Gamete fusion and the production of a zygote

The process



Breeding times

| SEASONAL BREEDERS | CONTINUOUS BREEDERS |
|--|--|
| <ul style="list-style-type: none">The female animal is ready to mate at specific periods throughout the year <p>Advantage:</p> <ul style="list-style-type: none">The offspring are produced during favourable environmental conditions | <ul style="list-style-type: none">The female animal can mate at any time during the year due to the continuous reproductive cycle <p>Advantage:</p> <ul style="list-style-type: none">Offspring are continuously produced, ensuring population growth and continuity |

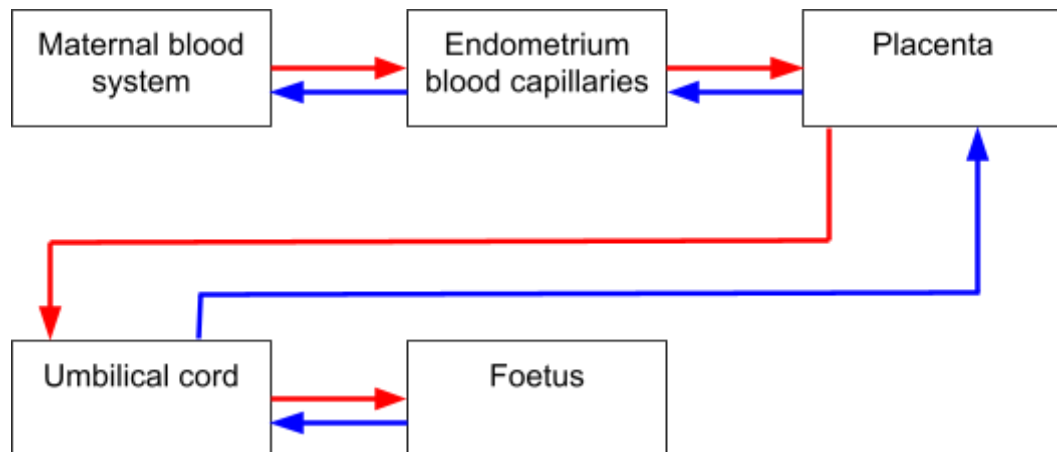
IMPLANTATION

The embryo is embedded into the uterus.

The placenta

The foetus develops an umbilical cord and placenta.

The placenta is a disk-shaped tissue which gets embedded into the uterus wall by placental villi. It connects the blood systems of the mother and the foetus without them fusing.



| Maternal → Foetus | Foetus → Maternal |
|---|--|
| <ul style="list-style-type: none">- Oxygen- Glucose- Water- Minerals- Antibodies- Hormones | <ul style="list-style-type: none">- Carbon dioxide- Urea- Water- Hormones |

PREGNANCY

Human development

| Zygote | Morula | Blastocyst | Embryo | Foetus |
|----------------|------------------------|--|--------------------------|--|
| Fertilised egg | 16 unspecialised cells | Mass of cells <ul style="list-style-type: none">- Inner forms embro- Outer forms placenta | 8wks after fertilisation | Embryo develops bone tissue. +8wks to birth |

Hormonal control of pregnancy

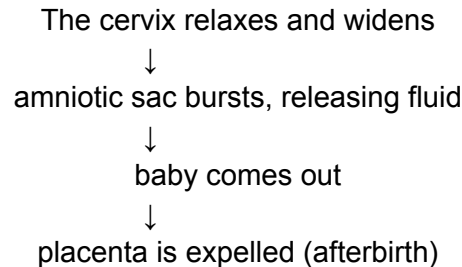
Embryos produce human chorionic gonadotropin (hCG). hCG prevents the corpus luteum from degrading, thus stimulating progesterone and maintaining a thickened endometrium.

After 12 weeks, the placenta is fully developed. It takes over the production of progesterone and oestrogen, so the corpus luteum degrades.

BIRTH

When the offspring moves into the external environment from the mother's reproductive system.

Process of birth



Hormonal control of birth

Progesterone prevents contractions in order to maintain a stable environment for the growing foetus.

In the last 1/3 of pregnancy, the placenta produces estriol.

Estriol inhibits the production of progesterone, while promoting oxytocin production. Furthermore, the rise in oestrogen increases the number of oxytocin receptors in the endometrium. Oxytocin causes the uterine muscles to contract.

Uterine contractions begin slowly, and increase due to positive feedback, becoming stronger until the baby is pushed out.

LACTATION

The production of milk to sustain the offspring after birth.

Hormonal control of lactation

The hormone prolactin stimulates the mammary glands to produce milk.

- High levels of oestrogen before birth stimulate the production of prolactin
(positive feedback)
- High levels of prolactin inhibit milk production
(negative feedback)

Milk production will only begin with the decline of prolactin. This happens with the decline of oestrogen (ie. After the birth).

[5.1.3] Manipulation of Plant and Animal Reproduction in Agriculture

Scientific understanding of the mechanisms of reproduction allowed for the development of reproductive technologies.

Reproductive technologies are any technologies which increase the rate of reproduction, ensure survival and thus continuity of species.

Biotechnology is the manipulation of living organisms for human benefit.

WHOLE-LEVEL MANIPULATION

Selective breeding (plants & animals)

Method:

Humans breed together parents with desirable traits so that at least one of these traits appears in the offspring.

Breeding can be:

- Intraspecific (between the same species)
- Interspecific (between different species)

| | |
|---|---|
| Advantages: <ul style="list-style-type: none">• If there are desirable traits in the male <i>and</i> the female, the offspring will most likely also have at least one of these traits | Disadvantages: <ul style="list-style-type: none">• Cannot guarantee the desired trait will be expressed• Reduces the genetic diversity of a population• Offspring from interspecific breeding are sterile as their gametes are not viable, thus the trait can not be passed on further |
|---|---|

CELL-LEVEL MANIPULATION

Artificial insemination (animals)

Method:

1. Semen is collected from a male
2. A sexually receptive female is detected
3. Insemination performed by using an insemination gun to shoot semen into the female's cervix

| | |
|---|--|
| Advantages: <ul style="list-style-type: none">• Able to synchronise births• Ability to choose the mates increases likelihood of passing down favourable traits• Ensurance of successful pregnancy increases yields | Disadvantages: <ul style="list-style-type: none">• Cannot guarantee the desired trait will be expressed• Reduced genetic variation |
|---|--|

Artificial pollination (plants)

Method:

1. Pollen (male gamete) removed from the stamen of one plant
2. This pollen is applied to the stigma of another plant
3. Pollen fertilises the ovum

| | |
|--|--|
| Advantages: <ul style="list-style-type: none">• Cross-breeding of favourable traits• Decreases likelihood of self-pollination• Ensures successful pollination of all plants, increasing the yield | Disadvantages: <ul style="list-style-type: none">• Cannot guarantee the desired trait will be expressed• Reduced genetic variation |
|--|--|

Cloning (plants & animals)

Method:

1. Somatic body cell of organism A is removed, and the nucleus is extracted
2. This nucleus is put into the denucleated egg of a donor
3. The egg is implanted into a surrogate mother
4. The offspring is a clone of A

| | |
|---|--|
| Advantages: <ul style="list-style-type: none">• Cloned individuals have identical requirements• The desired trait is guaranteed to be expressed | Disadvantages: <ul style="list-style-type: none">• Clones have no adaptability• Expensive• Ethical concerns• Clones have questionable health and life expectancy |
|---|--|

GENE-LEVEL MANIPULATION

Transgenesis (plants & animals)

Method:

1. A gene for a specific trait is removed from a species
2. The transgene is held in recombinant plasmid
3. The gene is injected into the zygote of another species
4. The zygote + transgene is implanted into a surrogate mother
5. The offspring express the transgene in their phenotype

| | |
|---|---|
| Advantages: <ul style="list-style-type: none">• Guaranteed expression of desired trait• Increase yield• Reduces use of harmful chemicals• Increases genetic diversity<ul style="list-style-type: none">◦ (as long as the transgenic organisms are not cloned) | Disadvantages: <ul style="list-style-type: none">• Transgenic organisms have low adaptability• Mixing of GM organisms with wild populations• Trade issues with non-GMO countries• Unknown side effects on humans• Transgenic organisms are usually cloned, decreases genetic variation |
|---|---|

IMPACT OF REPRODUCTIVE TECHNOLOGIES

Initially, due to manipulation of reproduction, there are increased variations in a species.

However, since the offspring with favourable traits are then cloned, variations decrease again.

gene level species level ecosystem level

↓ variations → ↓ genetic diversity → ↓ species diversity → ↓ long-term biodiversity

This limits the adaptability of a species to changing selection pressures, therefore limiting the continuity of species.

5.2: Cell Replication

How important is it for genetic material to be replicated exactly?

Genetic material refers to DNA.

Because DNA replication produces genetically identical offspring, all members of a species have the same genome (collection of genes) but different variations of that gene (alleles).

This means a species is defined by its genome.

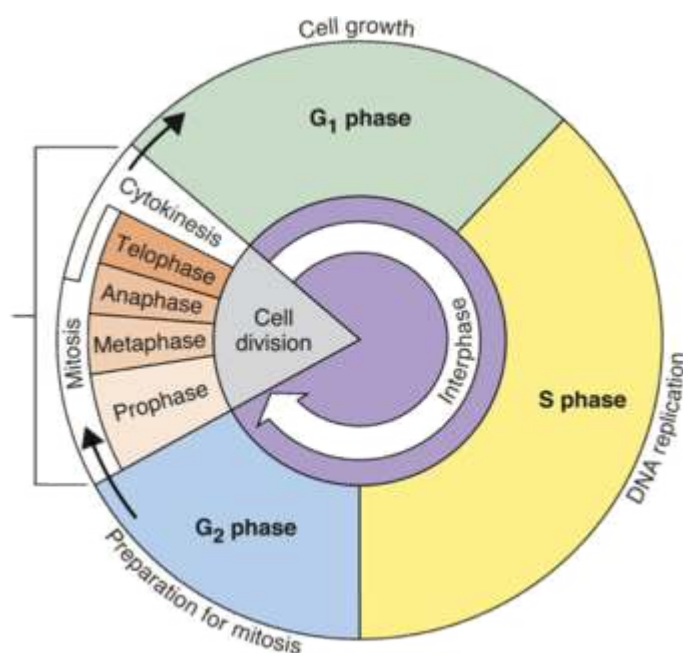
→ No two species have the same genome

Therefore, exact DNA replication is important as it ensures that the correct genes are passed on from parent to offspring, so that the species can continue in the next generation.

DNA replication → Inheritance of identical genes → Inheritance of exact genome → Continuity of species

[5.2.1] The Processes Involved in Cell Replication

THE CELL CYCLE



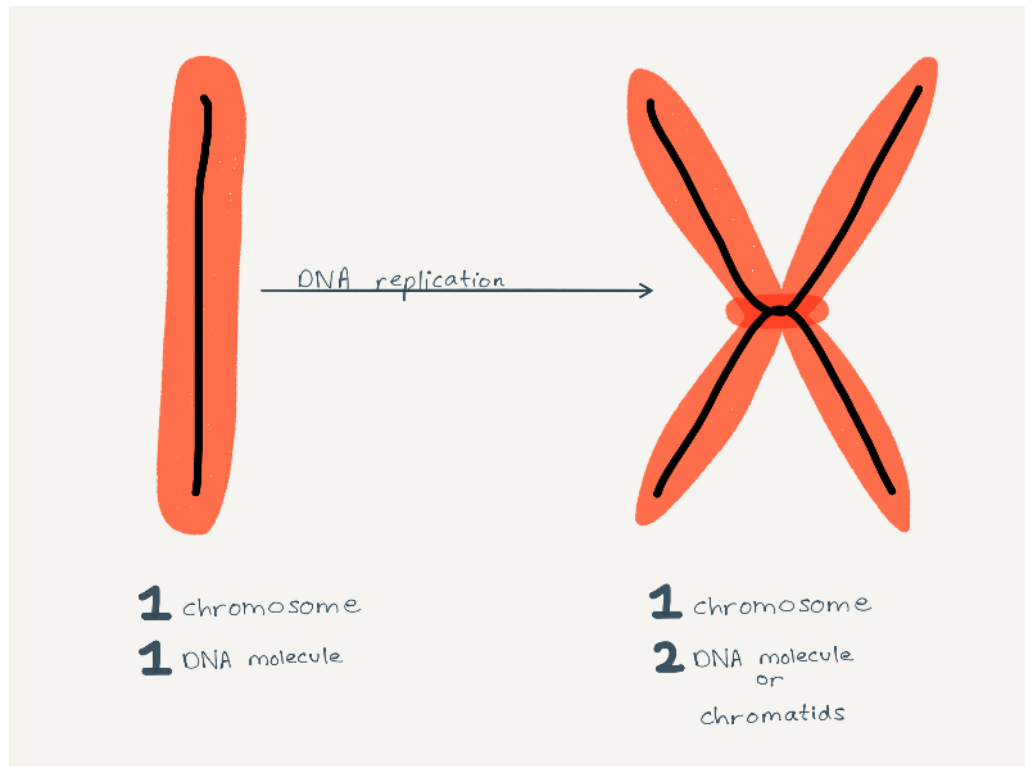
| | |
|------------------------------------|--|
| G ₁ : growth phase | Cell grows ↓ Volume increases faster than SA ↓ SA:V decreases ↓ |
| S : synthesis phase | DNA synthesis & replication ↓ Mutations ↓ |
| G ₂ : Growth phase 2 | Detection of mutations ↓ Correction of errors ↓ |
| Nuclear division | Prophase, metaphase, anaphase, telophase ↓ |
| Cytoplasmic division | Cytokinesis |

BEHAVIOUR OF CHROMOSOMES

Before replication: Many genes → 1 DNA → 1 chromosome

After replication: Many genes → 2 DNA → 1 chromosome

DNA replicates, chromosomes don't



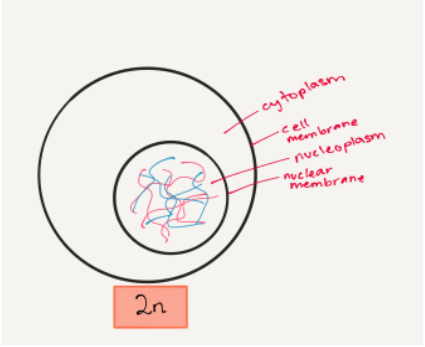
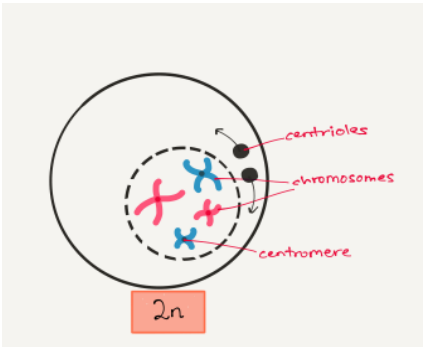
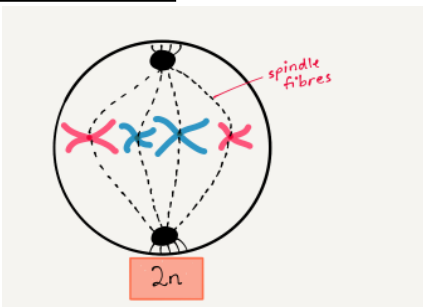
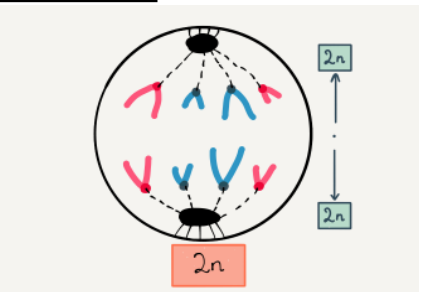
When 2 DNA are attached at the centromere, they are known as 2 sister chromatids.

MITOSIS

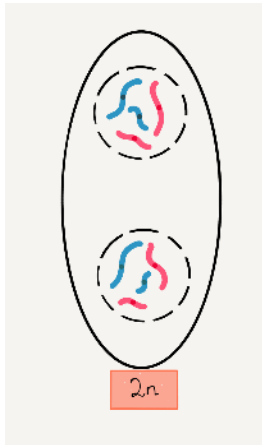
Location: Somatic (body) cells.

Result: Diploid ($2n$) parent cell creates two diploid daughter cells.

Function: Growth, repair, maintenance.

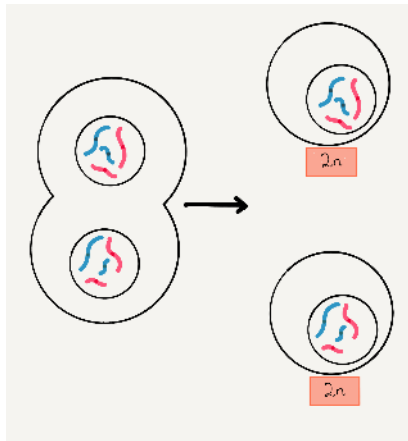
| | |
|---|--|
| <p><u>INTERPHASE</u></p>  | <ul style="list-style-type: none"> Chromosomes can not be seen individually, they are called chromatin material. <ul style="list-style-type: none"> They are thin, long and coiled together. DNA replication takes place <p>46 chromosomes DNA replication 46 chromosomes → 46 DNA 92 DNA/ chromatids</p> |
| <p><u>PROPHASE</u></p>  | <ul style="list-style-type: none"> Nuclear membrane dissolves. Centriole divides and the halves move towards opposite poles of the cell. Individual chromosomes become visible. <ul style="list-style-type: none"> Each is made of 2 chromatids due to DNA replication in interphase. |
| <p><u>METAPHASE</u></p>  | <ul style="list-style-type: none"> Chromosomes align across the equator of the cell in a single file. Spindle fibers form which connect the centrioles to the centromeres of the chromosomes. |
| <p><u>ANAPHASE</u></p>  | <ul style="list-style-type: none"> Spindle fibers contract. Chromosomes with one DNA are pulled to opposite poles of the cell. <ul style="list-style-type: none"> They are no longer chromatids since the DNA molecules are not connected. |

TELOPHASE



- Nuclear division completes.
- Nuclear membrane forms around the chromosomes at each pole.
- 2 diploid, identical nuclei are formed.

CYTOKINESIS



- Cytoplasm cleavages
- Cytoplasmic division
- 2 diploid, identical daughter cells are derived.

MEIOSIS

Location: Germline (sex) cells.

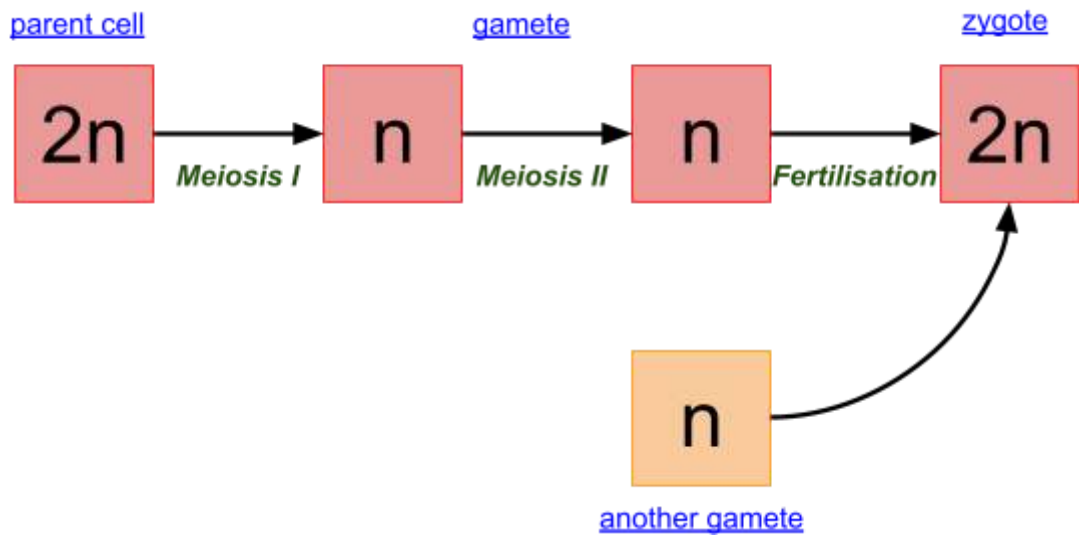
Result: Diploid parent cell creates 4 haploid (n) daughter cells.

Meiosis I: *Reduction division*. Creates haploid cells from diploid cells.

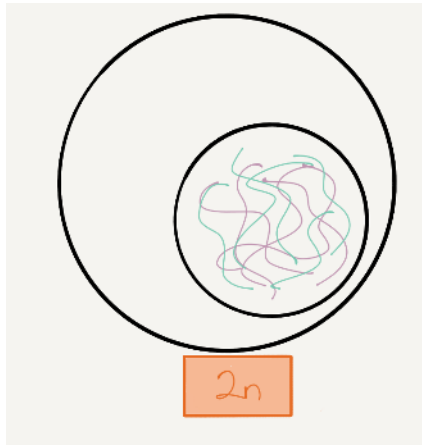
Meiosis II: Maintains haploid number in daughter cells.

Function: Gamete production for sexual reproduction.

↓
Ensures inheritance of the genome
↓
Source of variation
↓
Continuity of species



INTERPHASE



- Chromosomes can not be seen individually, they are called chromatin material.
 - They are thin, long and coiled together.

- DNA replication takes place

46 chromosomes

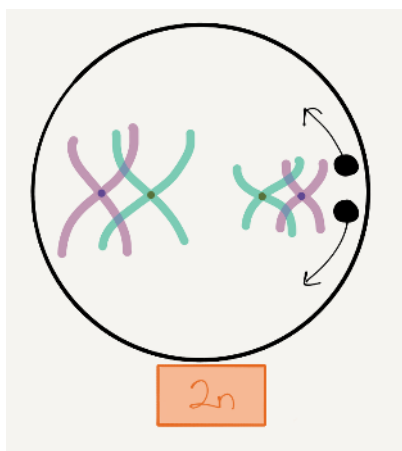
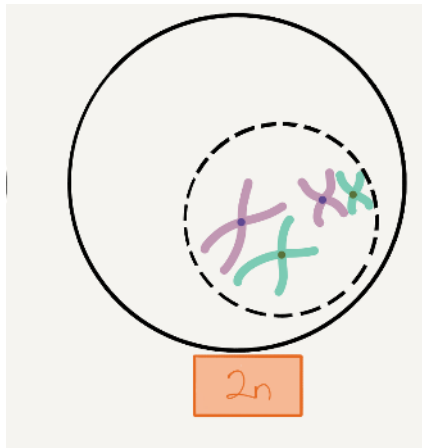
46 DNA

DNA replication
→

46 chromosomes

92 DNA/ chromatids

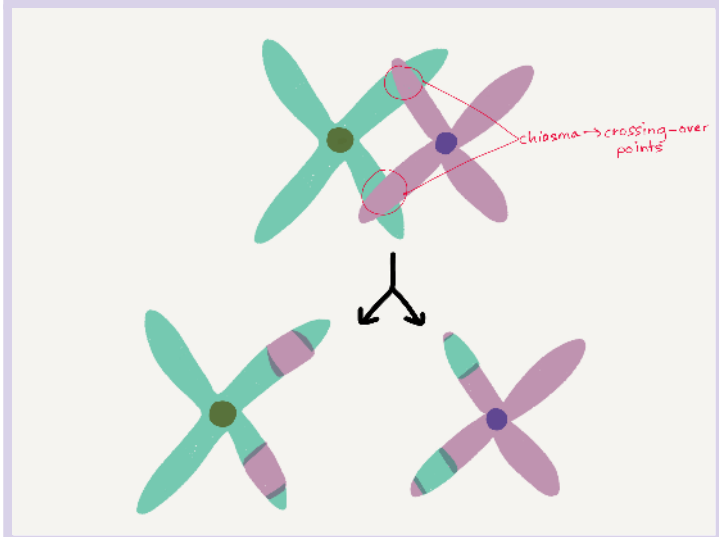
PROPHASE I



- Chromosomes with 2 chromatids appear.
- Chromosomes are arranged in homologous pairs.
- Nuclear membrane dissolves.
- Centrioles move towards the poles.

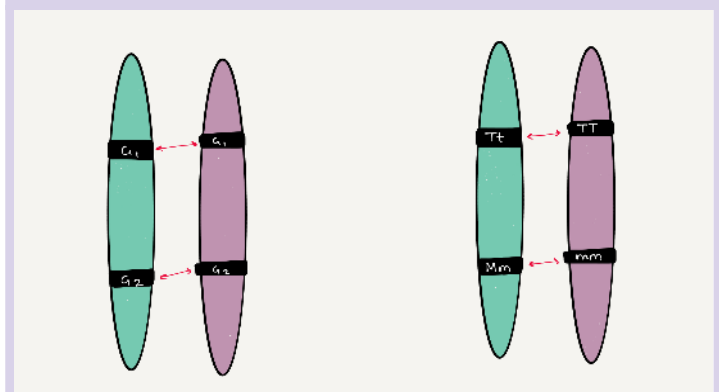
Crossing over

- Non-sister chromatids cross over and exchange maternal and paternal genes across the chiasma.
 - Creates **unique combinations of allele pairs on the same chromosome**
 - This recombination of maternal and paternal genes causes variation

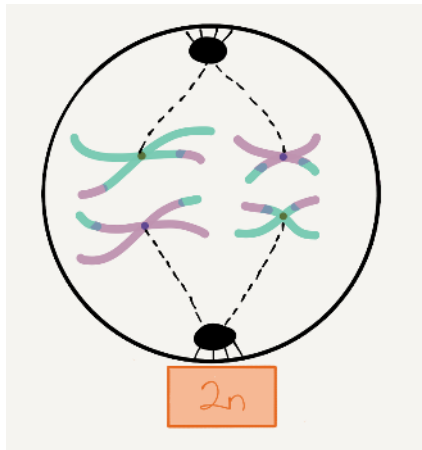


Alleles are variations of the same gene. Alleles for a common trait are found on the same locus of a chromosome.

- If a segment does not cross over in the exact area, mutations occur.



METAPHASE I



Independent assortment

- Homologous pairs of chromosomes line up in double file along the cell's equator.
- The arrangement is random, and independent of other chromosomes.

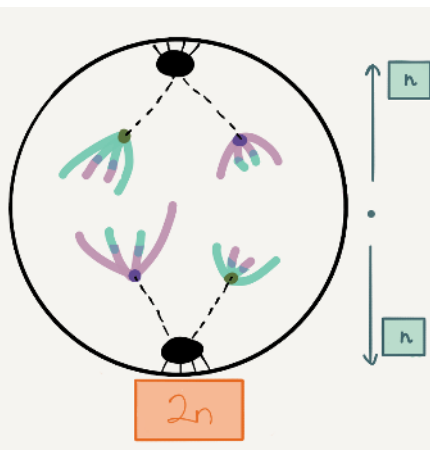
Random segregation

- Chromosomes of the homologous pairs move to the poles independently of each other.

Create a **unique combination of alleles** in the gametes.

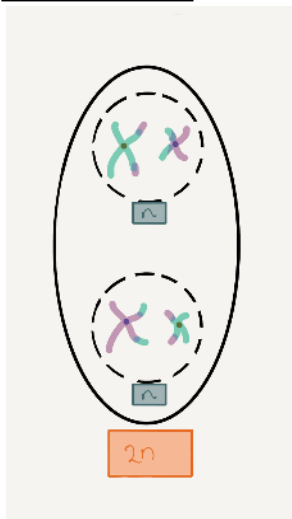
Crossing over, independent assortment and random segregation are a source of genetic variation.

ANAPHASE I



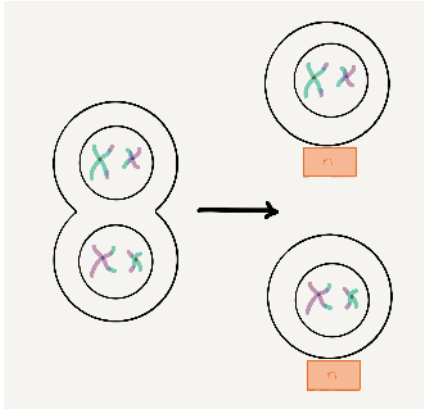
- Spindle fibers contract and pull chromosomes to the opposite poles.

TELOPHASE I



- Nuclear division completes.
- 2 haploid, non-identical nuclei are formed.

CYTOKINESIS I

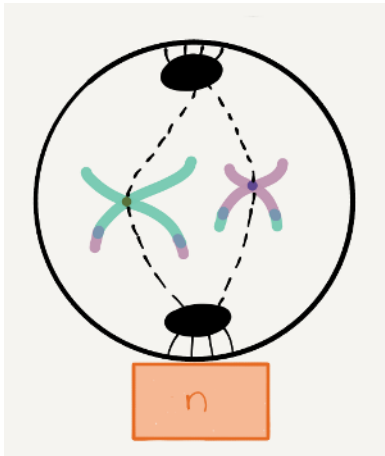


- Cytoplasmic division completes.
- 2 haploid, non-identical cells are formed.
- Meiosis I completes.

PROPHASE II

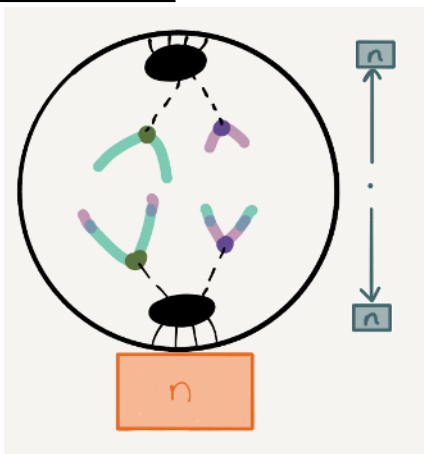
N/A

METAPHASE II



- Chromosomes line in a single file along the equator of the cell.

ANAPHASE I



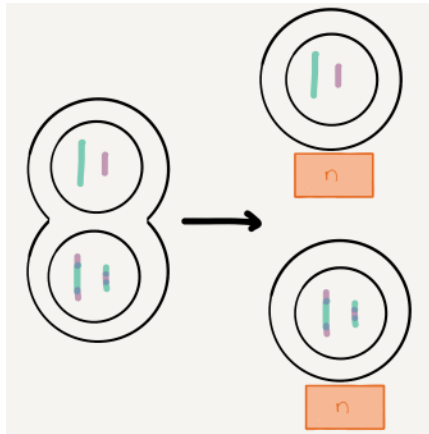
- Spindle fibers contract, pulling chromosomes to opposite poles.

TELOPHASE II



- Nuclear division completes.
- 2 haploid non-identical nuclei are formed per daughter cell.

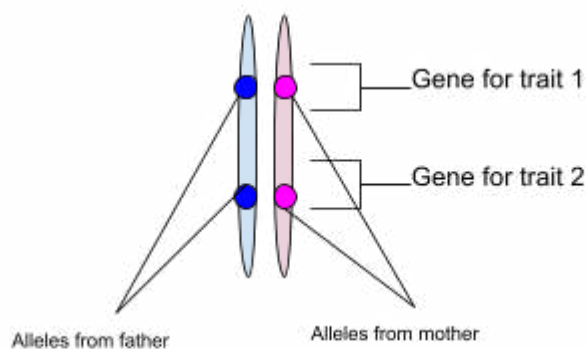
CYTOKINESIS II



- Cytoplasmic division completes.
 - 2 haploid, non-identical gametes are formed per daughter cell.
- In total, 4 haploid, non-identical gametes are formed.

When fused, the offspring will have 2 chromosomes in each pair, one containing **alleles from the father** and the other containing **alleles from the mother**.

Offspring's genotype



Furthermore, gametes are randomly fertilised, allowing for **different, random alleles to create a genotype** for a trait in the offspring. This increases genetic variation within the offspring.

Mitosis vs. Meiosis

| | <u>MITOSIS</u> | <u>MEIOSIS</u> |
|---|--|---|
| <i>Type of cell produced</i> | Somatic | Gamete |
| <i>No. of cells produced</i> | 2 | 4 |
| <i>Location</i> | Somatic cells | Germline cells |
| <i>Function</i> | Growth and repair | Gametes for sexual reproduction |
| <i>No. of chromosomes per daughter cell</i> | Diploid $2n \rightarrow 2n$ | Haploid $2n \rightarrow n \rightarrow n$ |
| <i>Degree of variation between parent cell and daughter cells</i> | Identical (Excluding variations due to mutations) | Large variation (due to random segregation and crossing over of genes) |
| <i>DNA replication</i> | Yes, in interphase. | Yes, in interphase. |

WATSON AND CRICK DNA MODEL

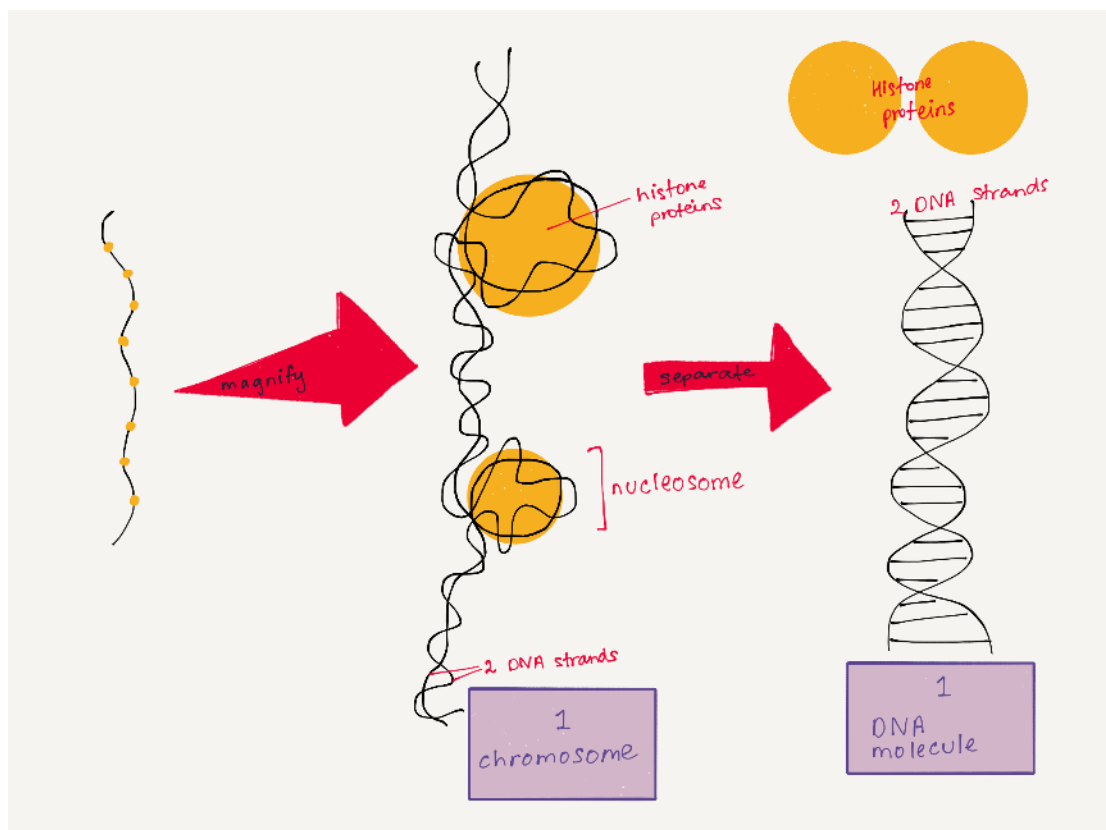
History of the DNA model

The W+C model of DNA was built in 1953.

- It was a double helix structure
- Scientists involved: James Watson, Francis Crick, Rosalind Franklin, Morris Wilkins

| Watson and Crick | vs. | Franklin and Wilkins |
|---|-----|--|
| <ul style="list-style-type: none"> • Watson was a biologist • Crick was a physicist • Perfect communication • Analytical skills | | <ul style="list-style-type: none"> • Franklin was an x-ray crystallographer • Wilkins produced DNA crystals • Franklin found angles of diffraction using crystalised DNA to create a model • Limited communication |

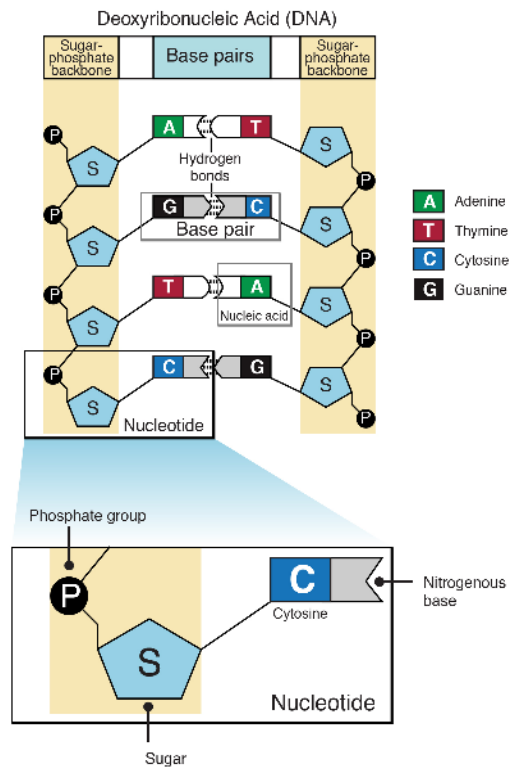
Nucleotide composition, pairing and bonding



DNA and chromosomes have a 1:1 ratio since a chromosome is a molecule of DNA wrapped around histone proteins.

Many genes → 1 DNA → 1 chromosome

DNA is a 3D molecule made of a sequence of nucleotides. Unwrapped, it looks like this:



Nitrogenous base + phosphate + sugar
= Nucleotide

The nitrogenous bases are:

1. Adenine (A) → Purines
2. Guanine (G) → Purines
3. Thymine (T) → Pyrimidines
4. Cytosine (C) → Pyrimidines

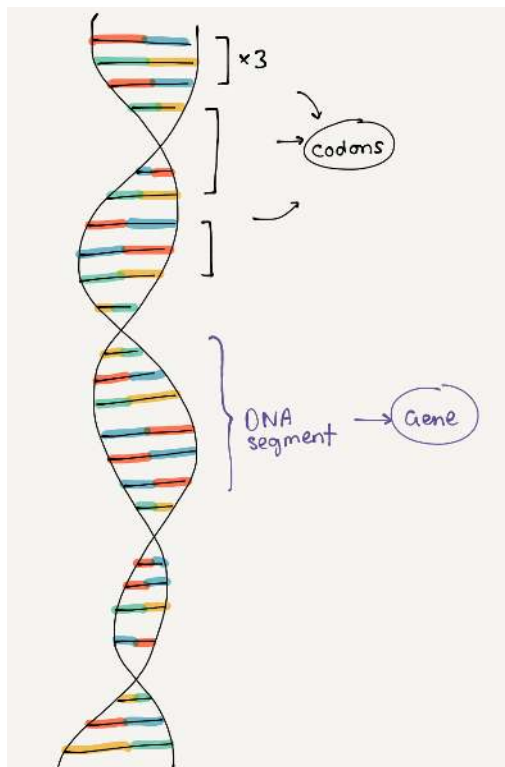
They undergo the complementary base pairings:

A = T [2 hydrogen bonds]

G ≡ C [3 hydrogen bonds]

Gene expression

The process by which the genotype shows the phenotype is called **gene expression**. It happens through protein synthesis.



GENE (DNA segment)

Sequence of nucleotides/ nitrogenous bases

Sequence of codons

↓ (each codes for an amino acid)

Sequence of amino acids

↓

Polypeptide chain

↓ (one or more chains link)

Specific protein

↓

Specific trait/ characteristic

This is how each gene determines a trait.

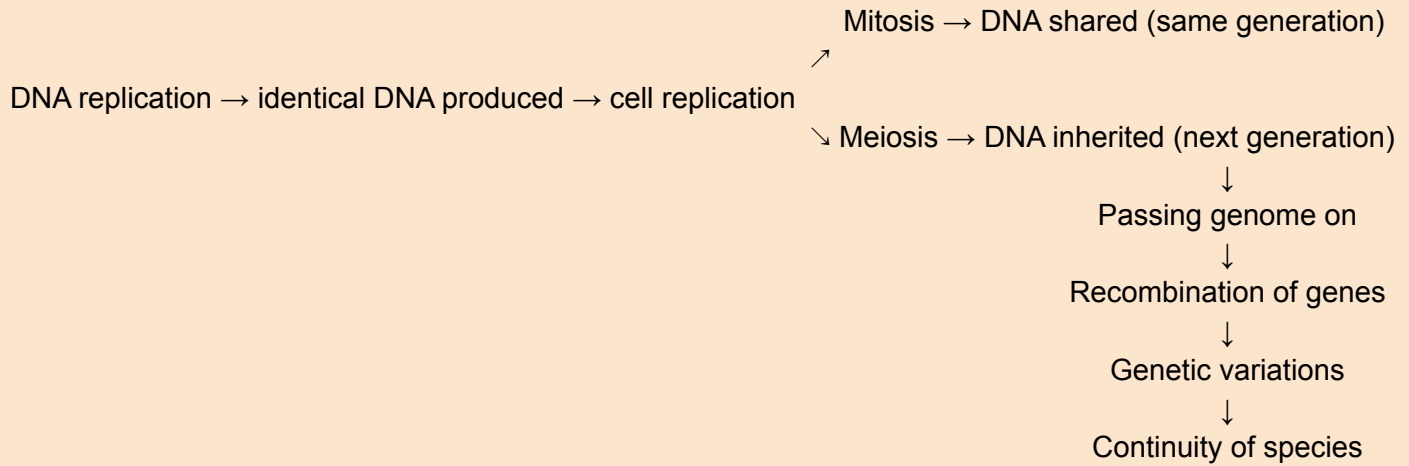
Many genes ⇒ many proteins ⇒ many traits

DNA REPLICATION IN EUKARYOTES

DNA replication occurs in the synthesis stage of interphase.

DNA REPLICATION

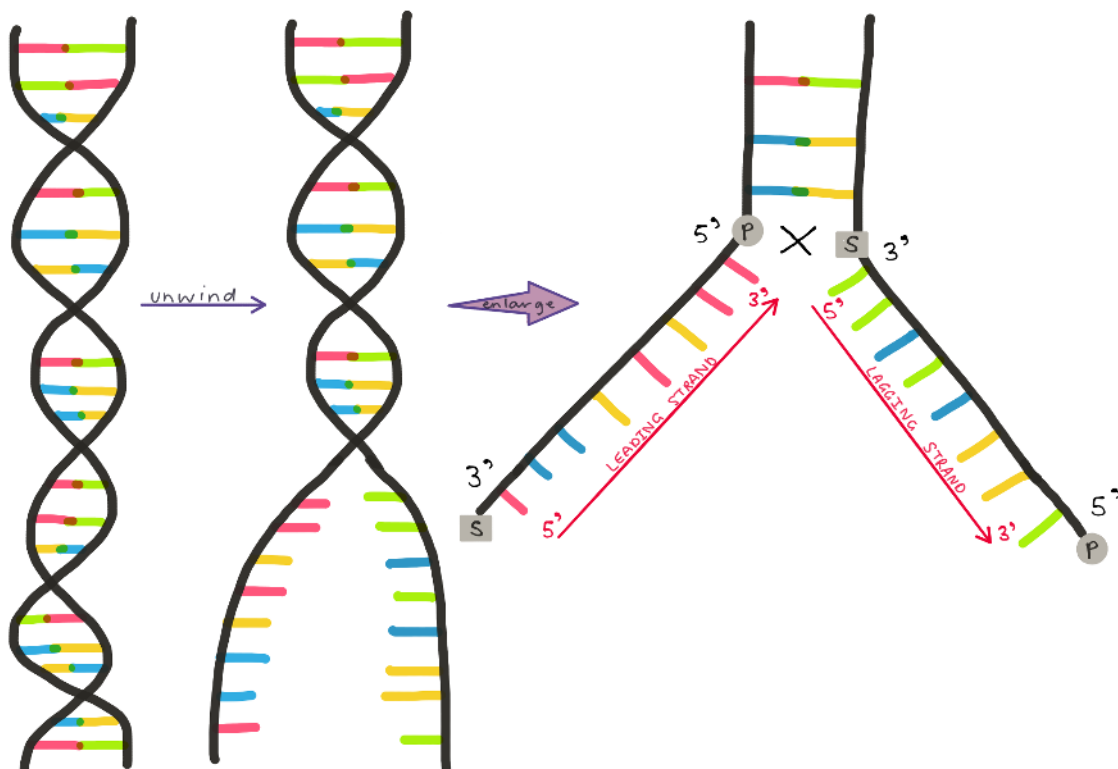
Importance



The original strand

1 molecule of DNA has 2 complementary strands which go from 3'→5' or 5'→3'.

- 3' ⇒ ends in sugar
- 5' ⇒ ends in phosphate



The strands are antiparallel.

- The *leading strand* forms continuously from 5'→3' towards the replication fork
- The *lagging strand* forms in fragments from 5'→3' away from the replication fork

The process of DNA replication

INITIATION: **Ligase** relieves the strain on a twisted DNA helix



Helicase uncoils and separates complementary strands



ELONGATION



Leading strand

DNA polymerase (III) adds nucleotides segments (primers) in the 5' → 3' *towards* the replication fork.

DNA polymerase also proofreads the strand to correct any base pair errors.



Ligase seals the gaps between DNA

Lagging strand

RNA polymerase adds a RNA primer to the strand, allowing DNA polymerase to bind.



DNA polymerase (III) adds Nucleotides next to the RNA primer in okazaki fragments.



DNA polymerase (I) replaces The RNA primer with DNA and corrects any pairing errors.



Ligase seals the gaps between DNA



The RNA primer moves to the next area, and the cycle repeats until the entire lagging strand is formed.



Strands recoil into a double-helix shape.
2 identical daughter strands produced.

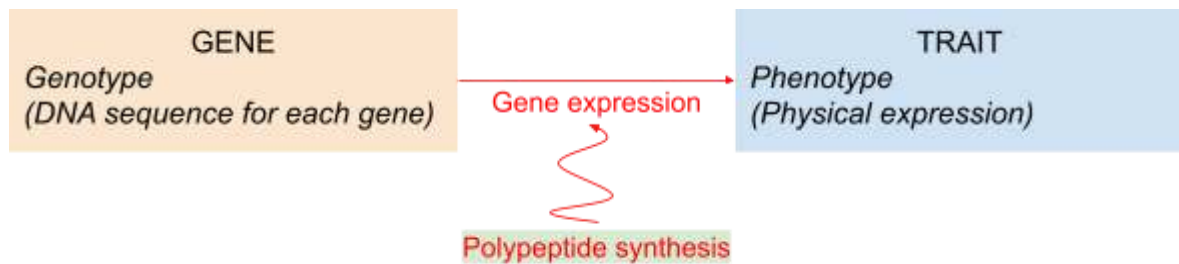


TERMINATION

The 2 new DNA strands each contain 1 old template strand and 1 new strand. This is called semi-conservative replication.

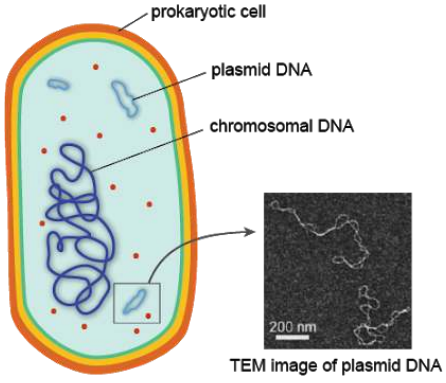
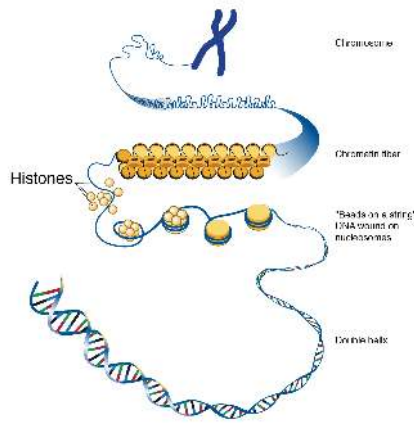
5.3: DNA and Polypeptide Synthesis

Why is polypeptide synthesis important?



Polypeptide synthesis is important because it allows for gene expression to occur so that organisms can live.

[5.3.1] DNA in Prokaryotes vs. Eukaryotes

| | <u>Prokaryotes</u> | <u>Eukaryotes</u> |
|--------------|---|--|
| DNA shape | <p>Chromosomal DNA in tight supercoils around scaffold protein.</p> <ul style="list-style-type: none"> - For essential features <p>Loop-like plasmid DNA.</p> <ul style="list-style-type: none"> - For non-essential features  <p>Conservative replication</p> | <p>Double helix coiled around histone proteins.</p> <ul style="list-style-type: none"> - DNA wrapped around histones to make nucleosomes  <p>Semi-conservative replication.</p> |
| DNA location | <p>Cytoplasm.</p> <ul style="list-style-type: none"> - Naked | <p>Nucleus</p> <ul style="list-style-type: none"> - Membrane bound |

NON-NUCLEAR DNA

Some organelles (eg. Mitochondria, Chloroplasts) have their own DNA.

Mitochondrial DNA (mtDNA) is identical through female lineage, and can be used to trace a genetic line.

- Evolves more rapidly, as replication is not checked by repair enzymes
- Abundant in every cell

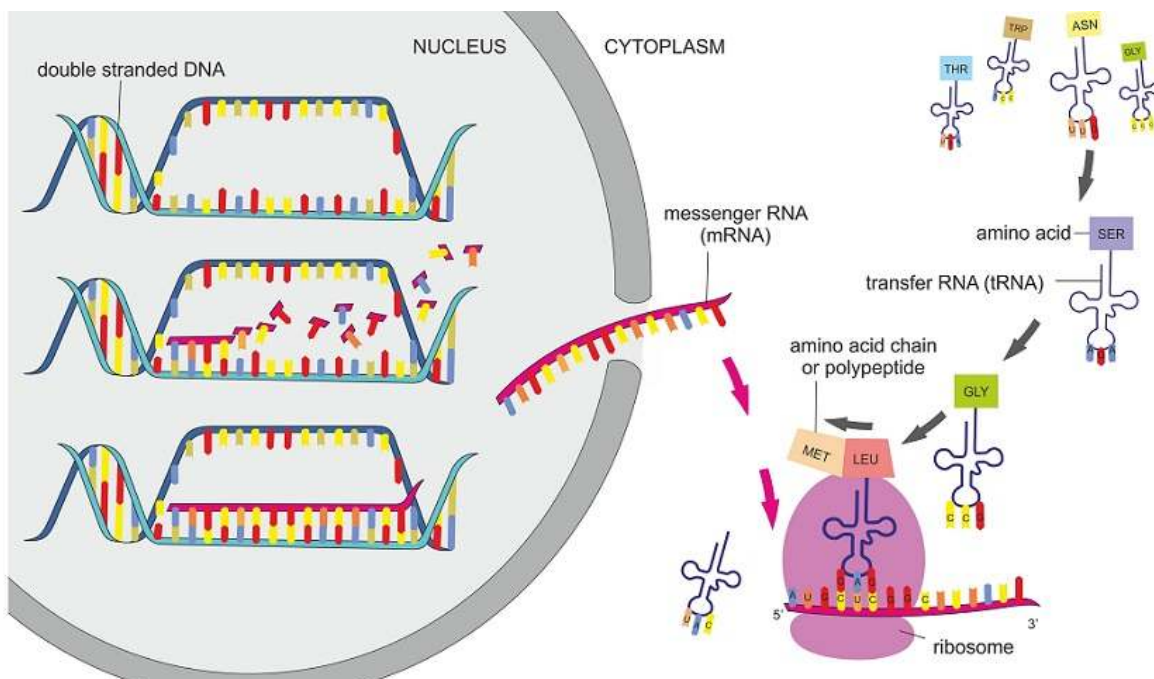
DNA vs. RNA

| DNA <i>Deoxyribonucleic acid</i> | RNA <i>Ribonucleic acid</i> |
|--|--|
| <ul style="list-style-type: none"> - Double helix - Deoxyribose sugar - 4 bases <ul style="list-style-type: none"> - Adenine - <u>Thymine</u> - Guanine - Cytosine - Stores genetic material - Undergoes replication | <ul style="list-style-type: none"> - Single helix - Ribose sugar - 4 bases <ul style="list-style-type: none"> - Adenine - <u>Uracil</u> - Guanine - Cytosine - Converts the information stored in DNA into proteins - Doesn't replicate |
| | <p>mRNA</p> <ul style="list-style-type: none"> • Takes the coding sequence for specific proteins from DNA and transports to the ribosomes <p>tRNA</p> <ul style="list-style-type: none"> • Carries amino acids to the ribosome. Specific types of amino acids correspond to the anticodon <p>rRNA</p> <ul style="list-style-type: none"> • Form the ribosome. They move along the mRNA, and bind the tRNA |

[5.3.2] POLYPEPTIDE SYNTHESIS

Function: To produce a polypeptide and then a protein

Significance: Allows gene expression to occur



Transcription

RNA polymerase enzyme unzips a segment of the DNA helix (a gene)

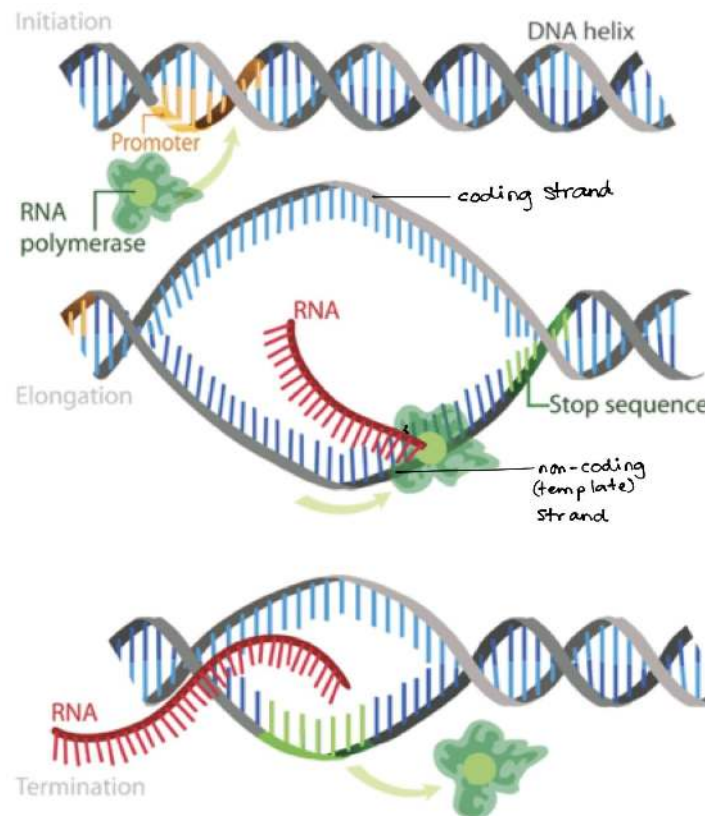
There are 2 strands: Non-coding and coding.

To replicate the coding strand, mRNA forms against the non-coding strand.

Pre-mRNA is created when free-floating nucleotides attach in complementary pairs.

- Instead of thymine, RNA uses uracil

DNA rezip and never leaves the nucleus.



Post-transcriptional modification

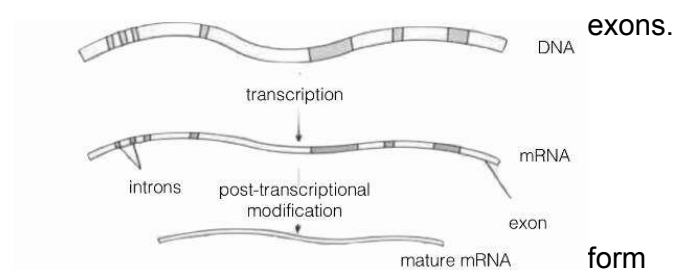
The transcribed mRNA contains introns and

Exons \Rightarrow codes for amino acids

Introns \Rightarrow non-coding regions

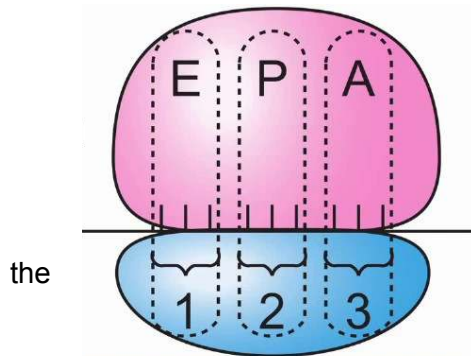
Endonuclease enzymes cuts out the introns.

The exons are connected by **ligase enzymes** to mature mRNA.



Translation

A ribosome is composed of a large and small subunit, with 3 binding sites: E, P, A.



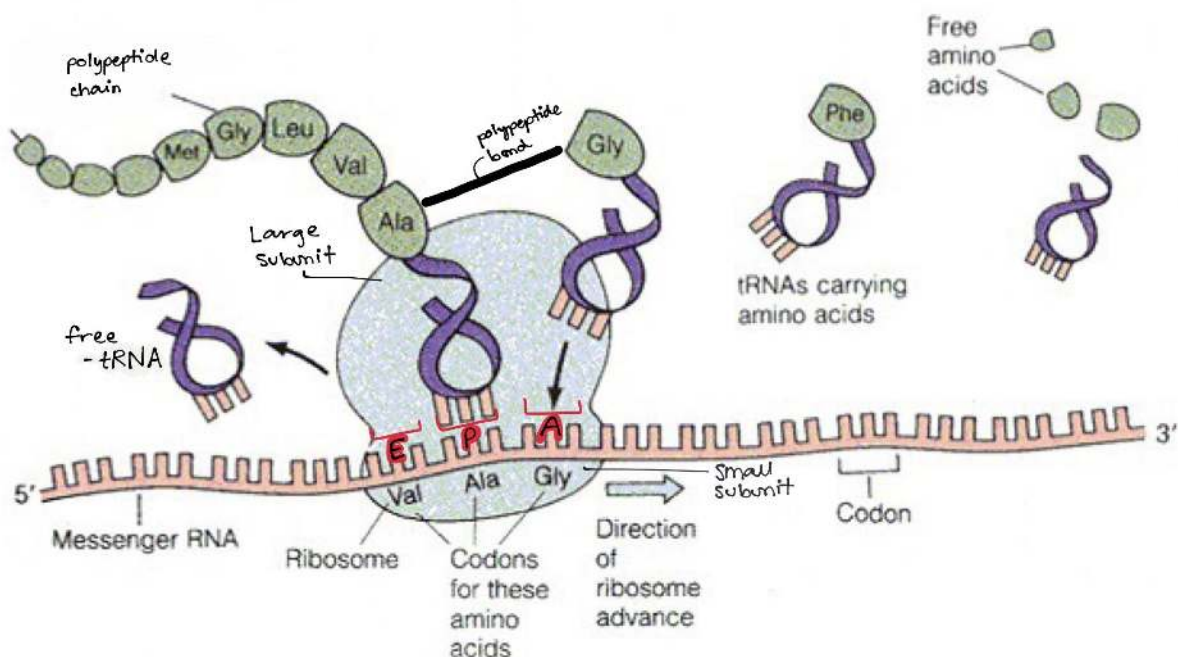
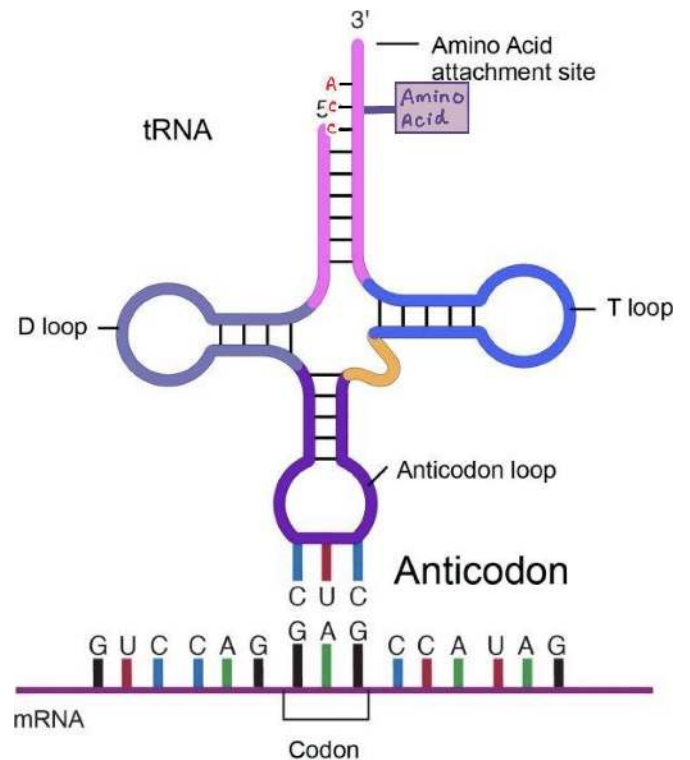
Each codon (3 base pairs) of the mRNA has a complementary tRNA which carries an amino acid.

Each codon codes for a specific amino acid.

The order and number of amino acids determines the function of protein.

| | | Second letter | | | | | |
|--------------|---|--------------------------------------|----------------------------------|--|--|------------------|--------------|
| | | U | C | A | G | | |
| First letter | U | UUU } Phe UUC UUA } Leu UUG | UCU } Ser UCC UCA } UCG | UAU } Tyr UAC UAA Stop UAG Stop | UGU } Cys UGC UGA Stop UGG Trp | U C A G | Third letter |
| | C | CUU } CUC } CUA } CUG } | CCU } CCC } CCA } CCG } | CAU } His CAC } CAA } Gln CAG } | CGU } CGC } CGA } CGG } | U C A G | |
| | A | AUU } AUC } Ile AUA AUG Met | ACU } ACC } ACA } ACG } | AAU } Asn AAC } AAA } Lys AAG } | AGU } Ser AGC } AGA } Arg AGG } | U C A G | |
| | G | GUU } GUC } GUA } GUG } | GCU } GCC } GCA } GCG } | GAU } Asp GAC } GAA } Glu GAG } | GGU } GGC } GGA } GGG } | U C A G | |

1.



1. Initiation

mRNA attaches to the small subunit. The starting codon has the sequence AUG



A complementary initiator tRNA [1] (carrying methionine) attaches to the A site and moves to the P site



The tRNA [2] of the next codon lands on the A site



A polypeptide bond is created between the two amino acids



tRNA [1] drops its amino acid and moves to the E site as a free tRNA

2. Elongation

The initiator tRNA is recycled in the cytoplasm



The tRNA [2] now carries 2 amino acids and moves to the P site



The ribosome continues to read the mRNA



tRNA [3] lands on the A site, and a polypeptide bond is created between the amino acids of tRNA [2] and [3]



tRNA [2] drops its amino acid and moves to the E site, giving tRNA [3] 3 amino acids



This cycle continues as the polypeptide chain elongates

3. Termination

The ribosome reaches a stop codon

- UAG
- UGA
- UAA

The stop codon doesn't code for a protein, but it releases the polypeptide chain from the P site by releasing chemicals.

MUTATIONS

If a mutation occurs during DNA replication, when transcribed and translated it will lead to an incorrect amino acid in the polypeptide chain, creating an abnormal protein.

This will impact the metabolism and thus the health of the organism.

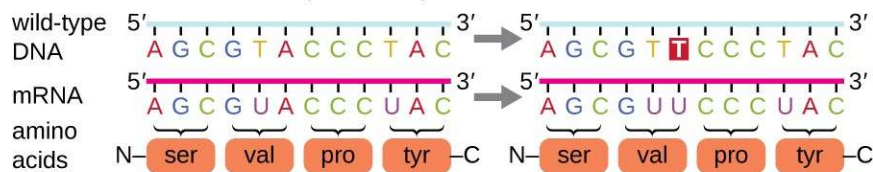
Base substitutions

1 base of the coding strand is replaced with an incorrect, non-complimentary one.

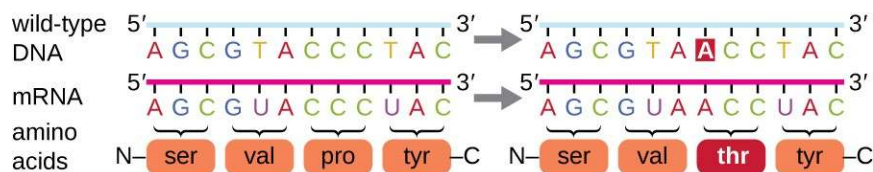
This may:

- Cause no change → **silent mutation**
- Change 1 amino acid → **missense mutation**
- Shorten the chain → **nonsense mutation**

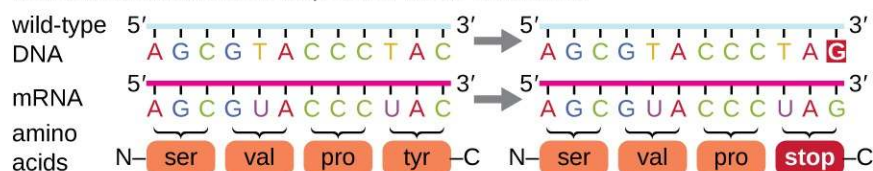
silent: has no effect on the protein sequence



missense: results in an amino acid substitution



nonsense: substitutes a stop codon for an amino acid



Base insertions and deletions

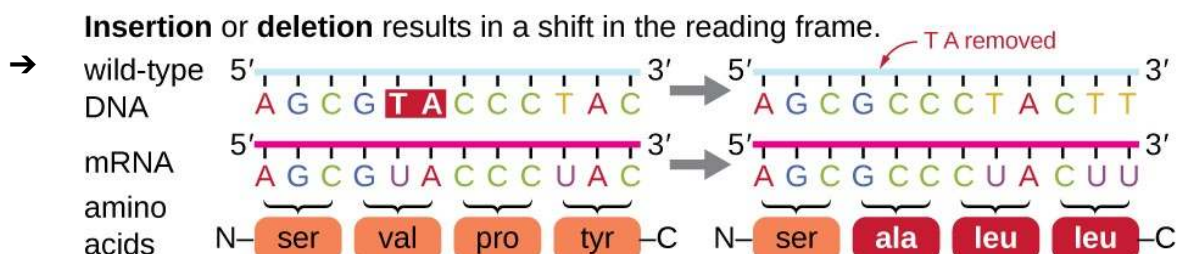
Bases are removed or added.

↓
This changes the reading frame, as bases are read in 3 (codons)

↓
Frame-shift mutation

↓
Significantly changes the amino acid sequence

↓
Non-functional protein



GENES AND THE ENVIRONMENT ON PHENOTYPE

Although the genotype is expressed as the phenotype, traits may be masked or changed by the environment.

$$\text{Genes} + \text{Environment} = \text{Phenotype}$$

The alteration may be temporary (usually) or permanent.

Genes will only express traits if the environment is suitable.

Examples:

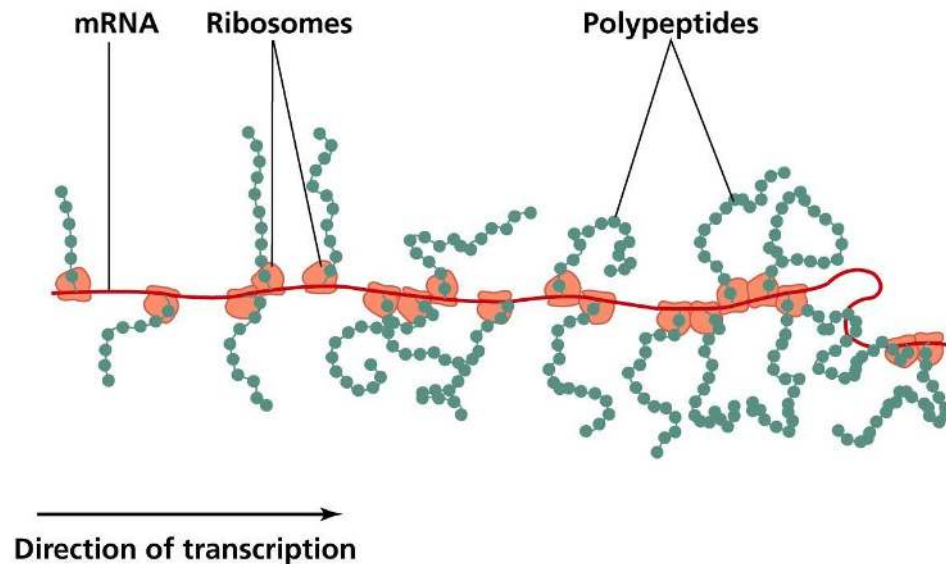
- *Having tall genes will not make an individual tall if there is not enough nutrients in the environment.*
- *The petals of hydrangea plants change colour based on soil pH*
acidic \Rightarrow blue
alkaline \Rightarrow pink
- *Siamese cats will grow dark colours on the tips of their bodies when the temperature decreases, and lose them when it increases.*

[5.3.3] Investigate the Structure and Function of Proteins in

Living Things

Polyribosomes

Multiple ribosomes can form parts of the polypeptide chain, which come together to form the complete chain.

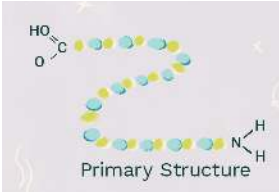
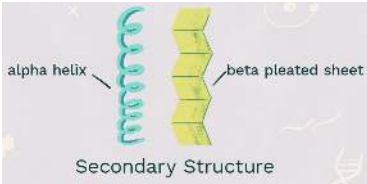




Protein conformation

One or more polypeptide chains can be structurally modified in the golgi apparatus to form specific proteins.

Amino acids (monomers) → polypeptides → proteins (polymers)

Types of proteins

| <u>PROTEIN SHAPE</u> | <u>FEATURES</u> |
|---|---|
| <p>Primary protein</p>  <p>The diagram shows a linear sequence of amino acids represented by colored spheres. The carboxyl end (HO-C=O) is on the left and the amino end (N-H) is on the right. The text 'Primary Structure' is at the bottom.</p> | <ul style="list-style-type: none">- A single, linear chain of amino acids |
| <p>Secondary protein</p>  <p>The diagram shows two types of local folding: an 'alpha helix' (a coiled blue ribbon) and a 'beta pleated sheet' (a zig-zagging yellow ribbon). The text 'Secondary Structure' is at the bottom.</p> | <ul style="list-style-type: none">- A linear sheet of amino acids- Pleated into alpha helices or beta sheets |

| | |
|---|---|
| <p>Tertiary protein</p>  <p>Tertiary Structure</p> | <ul style="list-style-type: none"> - A linear sheet of amino acids - Pleated into alpha helices or beta pleated - Folded into a 3D structure |
| <p>Quaternary protein</p>  <p>Quaternary Structure</p> | <ul style="list-style-type: none"> - 2 or more proteins (of any structure) come together to form a single protein |

Once shaped, proteins are held together by **hydrogen bonds**.

Each structure leads to the other, meaning that each protein will show characteristics of the shape before it.

Types of proteins and functions

| <u>TYPE OF PROTEIN</u> | <u>FUNCTION OF PROTEIN</u> |
|---|---|
| Neurotransmitters [primary /secondary] | Transport. Chemically transfer electrical impulses across the synapsis. |
| Lysozymes [secondary] | Storage. The fluid within the lysosome organelle. |
| Enzymes [tertiary] | Biocatalysts. Speed up biochemical reactions. |
| Haemoglobin [quaternary] | Transport. Transports oxygen |
| Hormones [quaternary] | Messengers. Chemical messengers which coordinate cell activity |
| Antibodies [quaternary] | Immunity. Fights pathogens for immunity |
| Collagen [quaternary] | Growth & repair. Muscle fiber which strengthens muscles |

5.4: Genetic Variation

How can the genetic similarities and differences within and between a species be compared?

Genetic similarities and differences can be compared at a genetic level within species and at a species level between species.

Within a population (intraspecific) :

- Similarities show common ancestry
- Differences show genetic diversity due to genetic variation

Between populations (interspecific) :

- Similarities show evolutionary relationships
- Differences show species diversity due to species variation

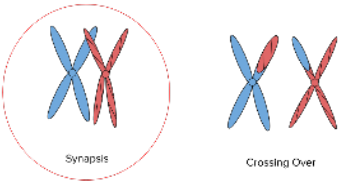
[5.4.1] Sources of Variation

Genome: The complete set of genes in a species

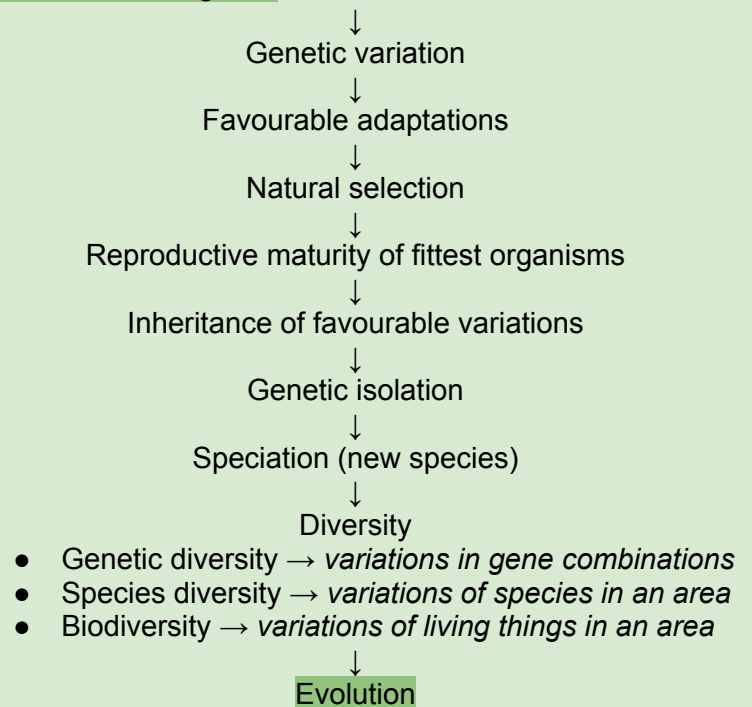
Gene pool: The complete set of alleles (gene variations) in a species.

Genetic diversity: total of all the genetic characteristics in the genetic makeup of a species.

Sources of genetic variation:

| <u>Source</u> | <u>Effect</u> |
|--|---|
| <p>Crossing over (<i>prophase</i>)</p>  | <p>Homologous chromosomes exchange genes, creating new combinations of alleles in the chromosome.</p> |
| <p>Independent assortment (<i>metaphase I</i>) & Random segregation (<i>anaphase I</i>)</p> | <p>Homologous chromosomes arrange themselves and separate independently, leading to random allele combinations in the 4 non-identical gametes.</p> |
| <p>Random fertilisation</p> | <p>Non-identical male and female gametes fuse to create an offspring with a unique combination of alleles to form unique genotypes.</p> |
| <p>Mutations</p> | <p>Produce new alleles to increase the possible variations.</p> |

Meiosis + Fertilisation + Mutations = **recombination of genes**



Therefore, genetic variation leads to evolution. (Neo-Darwinism).

[5.4.2] Mechanisms of Inheritance

Chromosomes

- **Autosome**: contains genes responsible for body traits (skin, hair, eye colour)
1-22nd homologous pairs in humans.
- **Sex-chromosomes**: contains genes for gender and secondary sexual characteristics
23rd pair in humans
 - *XX (females)*
 - *XY (males)*

Every body cell contains a complete set of **homologous chromosome pairs**.

In humans, there are 23 pairs in total.

MUTATIONS

Mutations occur in both the somatic and germline cells.

Mutations in somatic cells → expression in individual → no inheritance

Mutations in germline cells → not expressed in individual → gametes → inheritance → expressed in offspring

Gametes carry genetic variation, meaning that a trait will only be passed down if it occurs in the gametes. **Inheritance only occurs in germline cells.**

AUTOSOMAL INHERITANCE

Inheritance on germline cells where the genes inherited are on the 1st - 22nd pairs of chromosomes, and code for body characteristics.

In a specimen, there is a **gene** for each trait, and 2 variations of this gene known as **alleles**.

- The alleles can be dominant or recessive.
- Dominant alleles are always shown over recessive ones.

Eg. G_T is the gene for height.

T and t are the alleles (tall and short)

T is dominant, t is recessive.

P1: Parental generation

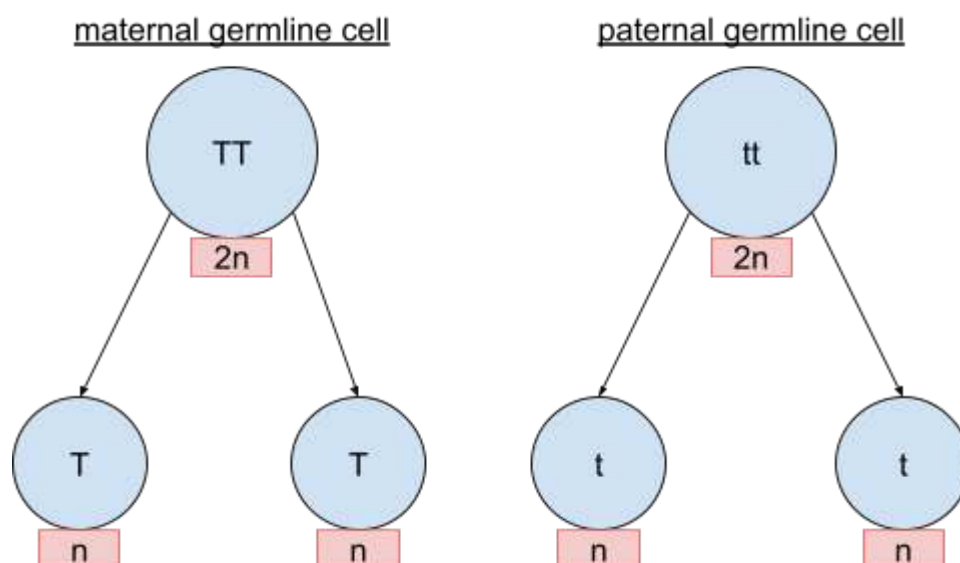
The alleles of both parents

Pure tall x Pure short

$TT \times tt$

G1: Gametes

Since only $\frac{1}{2}$ of each parent's genetic information is passed onto the offspring, they only have allele from each parent.



F1: Filial generation 1

| | T | T |
|-----|------|------|
| t | Tt | Tt |
| t | Tt | Tt |

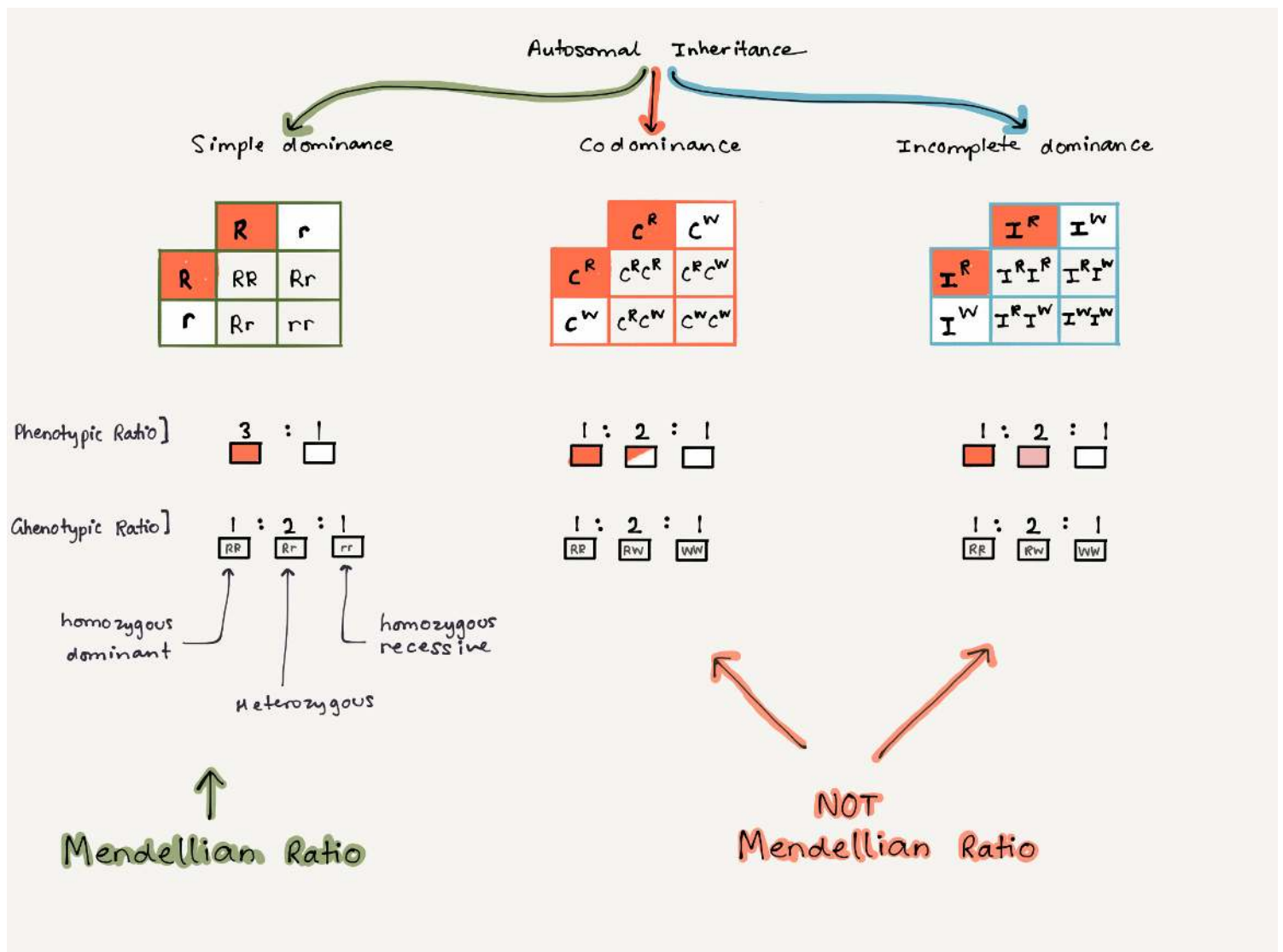
Offspring genotype: 100% heterozygous.

Offspring phenotype: 100% tall.

Simple dominance: One allele is completely dominant over the other.

Codominance: Both alleles are dominant, so they both show without blending.

Incomplete dominance: Both alleles are dominant, so they are blended in the phenotype.



Multiple alleles:

While there are only 2 alleles per genotype in an individual, some traits can be coded for more than 2 alleles.

(Aka. **Polygenic traits**)

If there are many variants, it is known as continuous distribution.

If there are few variants, it is a discrete distribution.

Example: Blood groups

There are 3 alleles that code for blood types: A, B and O

- O is recessive
- A and B are both dominant over O, but equally dominant over each other

Therefore, there are 4 blood groups for the genotypes created:

$I^A \rightarrow A$

$I^B \rightarrow B$

$i \rightarrow O$

| <u>Genotype</u> | <u>Alleles</u> | <u>Phenotype</u> |
|-----------------|-----------------|------------------|
| $I^A I^A$ | I^A | Blood type A |
| $I^A i$ | I^A and i | |
| $I^B I^B$ | I^B | Blood type B |
| $I^B i$ | I^B and i | |
| ii | i | Blood type O |
| $I^A I^B$ | I^A and I^B | Blood type AB |

SEX LINKAGE

Inheritance on germline cells where the genes inherited are on the 23rd pair of chromosomes. These genes code for gender and secondary sexual characteristics.

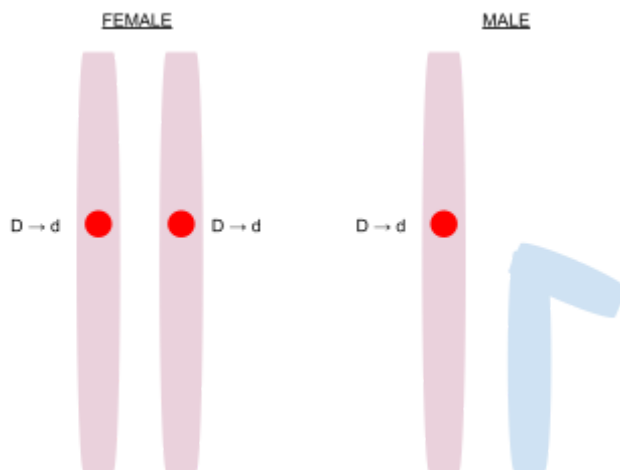
Due to a mutation, an autosomal gene can be transferred to the X chromosome of the 23rd (sex) pair.



It becomes an **X-linked gene**.



It undergoes **sex-linked inheritance**



$$X^D > X^d > Y$$

Female gametes → XX

Male gametes → XY

- Never carried on the Y chromosomes
- Only carried on the X chromosomes

X-chromosomes can be dominant or recessive, whereas Y chromosomes are always recessive.

→ Therefore, a female can be a carrier

→ Males can never be a carrier.

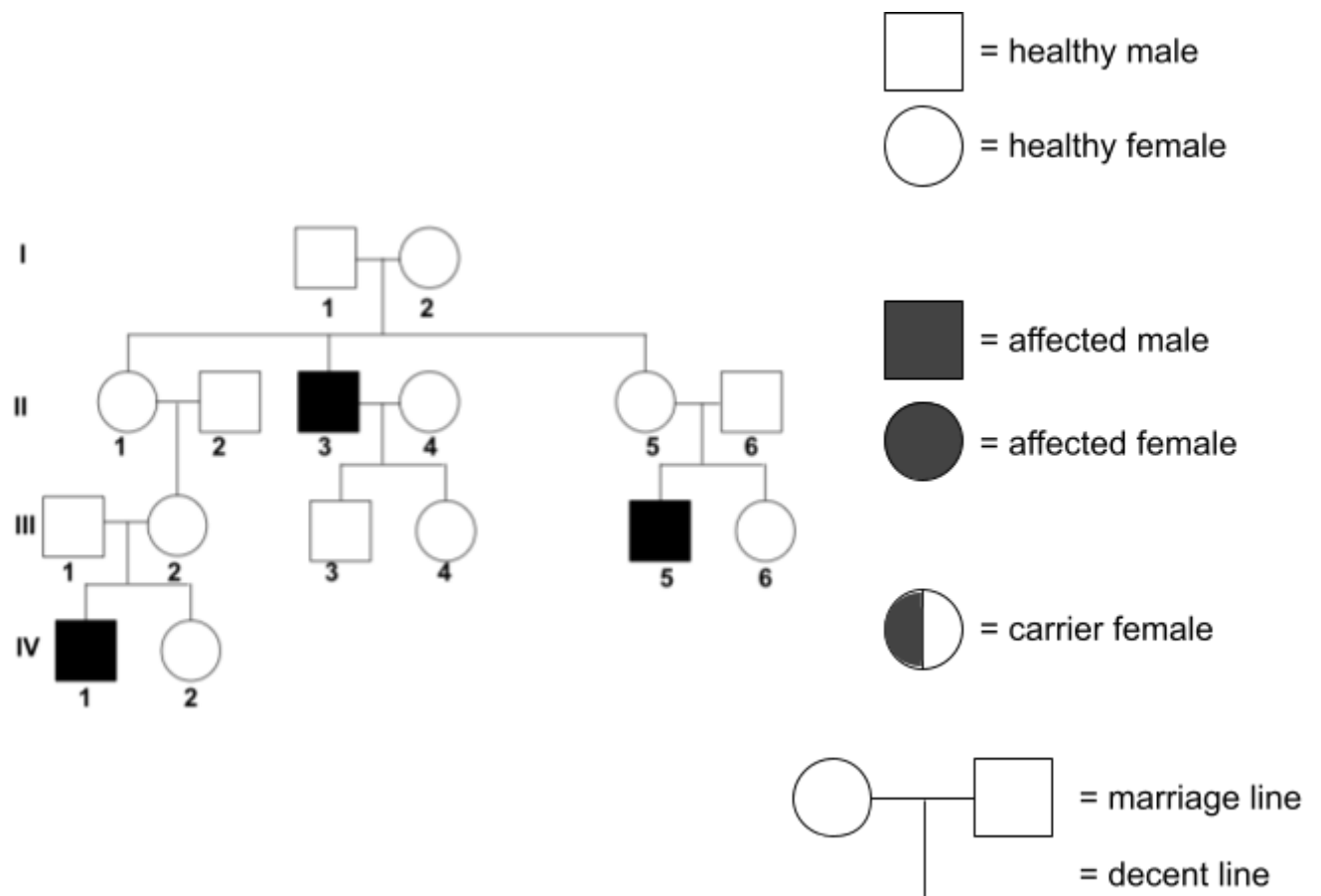
If the allele for a sex-linked disease is **recessive (e)**:

| | | |
|---|-----------------------------|-----------------------------------|
| X^E = dominant f. allele (most dominant) | X^e = recessive f. allele | Y = m. allele (most recessive) |
|---|-----------------------------|-----------------------------------|

| Genotype | Alleles | Gender | Phenotype |
|-----------|-------------|--------|-----------|
| $X^E X^E$ | X^E | Female | Normal |
| $X^e X^e$ | X^e | Female | Diseased |
| $X^E X^e$ | X^E X^e | Female | Carrier |
| $X^E Y$ | X^E Y | Male | Normal |
| $X^e Y$ | X^e Y | Male | Diseased |

PEDIGREES

Graphical representations of inheritance patterns of a trait in related individuals over generations.



Patterns of inheritance

| | | |
|---------------------|--|---|
| Autosomal dominant | → Males and females affected | |
| Autosomal recessive | → Males and females affected | |
| X-linked dominant | Male with trait: → Passes onto all daughters → Passes onto no sons | Female with trait: → Passes to both daughters and sons |
| X-linked recessive | Male with trait: → All daughters are carriers | Female with trait: → All sons are affected |

[5.4.3] Population Genetics

The study of frequency of traits/ characteristics in a population.

- ↳ Genotype
- ↳ Phenotype
- ↳ Alleles

Changes in frequency → Change to population → Genetic variation → Evolution

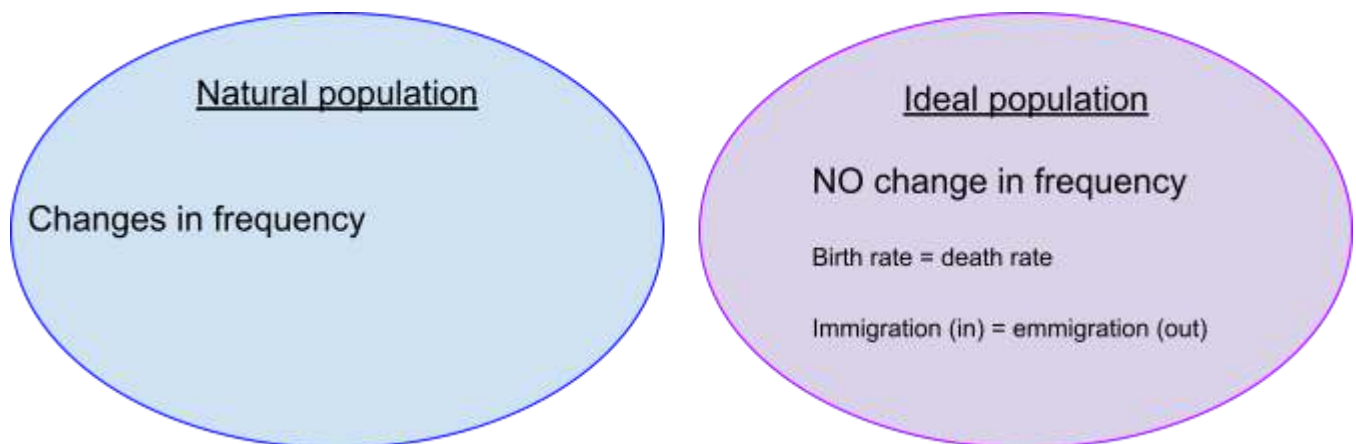
Studying population genetics allows us to:

- Predict the survival of a species through genetic variation
- Conservation management
- Predict the inheritance of diseases and disorders in a population
- Trace evolution

Mathematical model for frequency

$$\text{Frequency of specific allele/ genotype/ phenotype} = \frac{\text{No of that allele/ genotype/ phenotype.}}{\text{Total no. of alleles/ genotypes/ phenotypes}}$$

Modelling populations



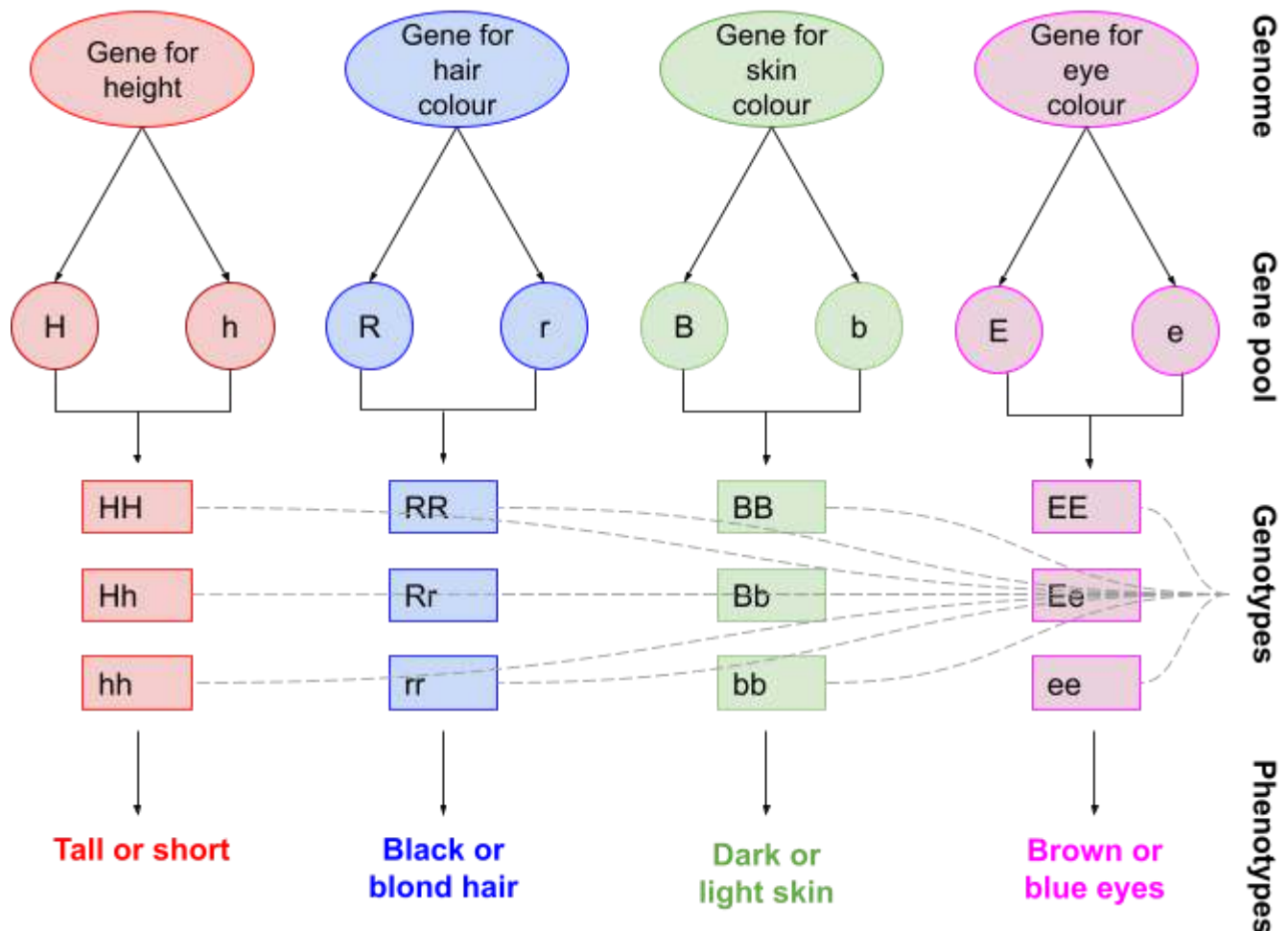
Changes in a natural population can be identified when compared to an ideal population.

→ Trends, patterns and limitations in the frequency data can be identified

Population genetics: Shows how the gene pool changes over time.

Genetic diversity: The total of all genetic characteristics in a species.

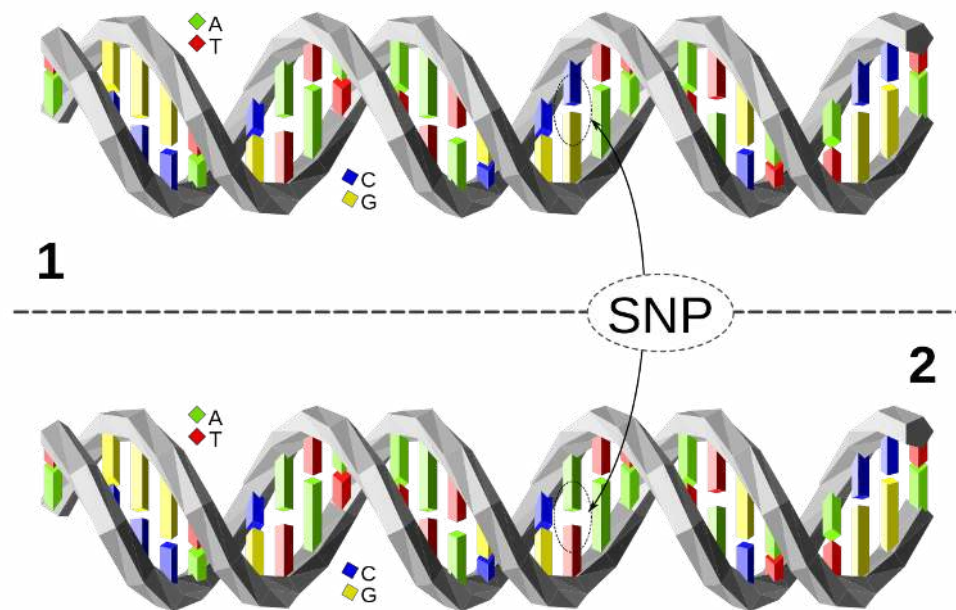
Genetic variability: The tendency for genetic traits in a population to vary between individuals.



| | |
|------------------|--|
| Genome | A collection of <u>all</u> genes responsible for <u>all</u> traits. The <u>same</u> for all members of a population. |
| Gene pool | Each gene has different variations or alleles. Collection of all the alleles in a population. |
| Genotype | Every individual inherits 2 alleles per trait. Each pair of alleles is a genotype. The genotype is <u>different</u> between members of a population. |
| Phenotype | Each genotype codes for 1 trait. The phenotypic expression of each genotype (or pair of alleles). |

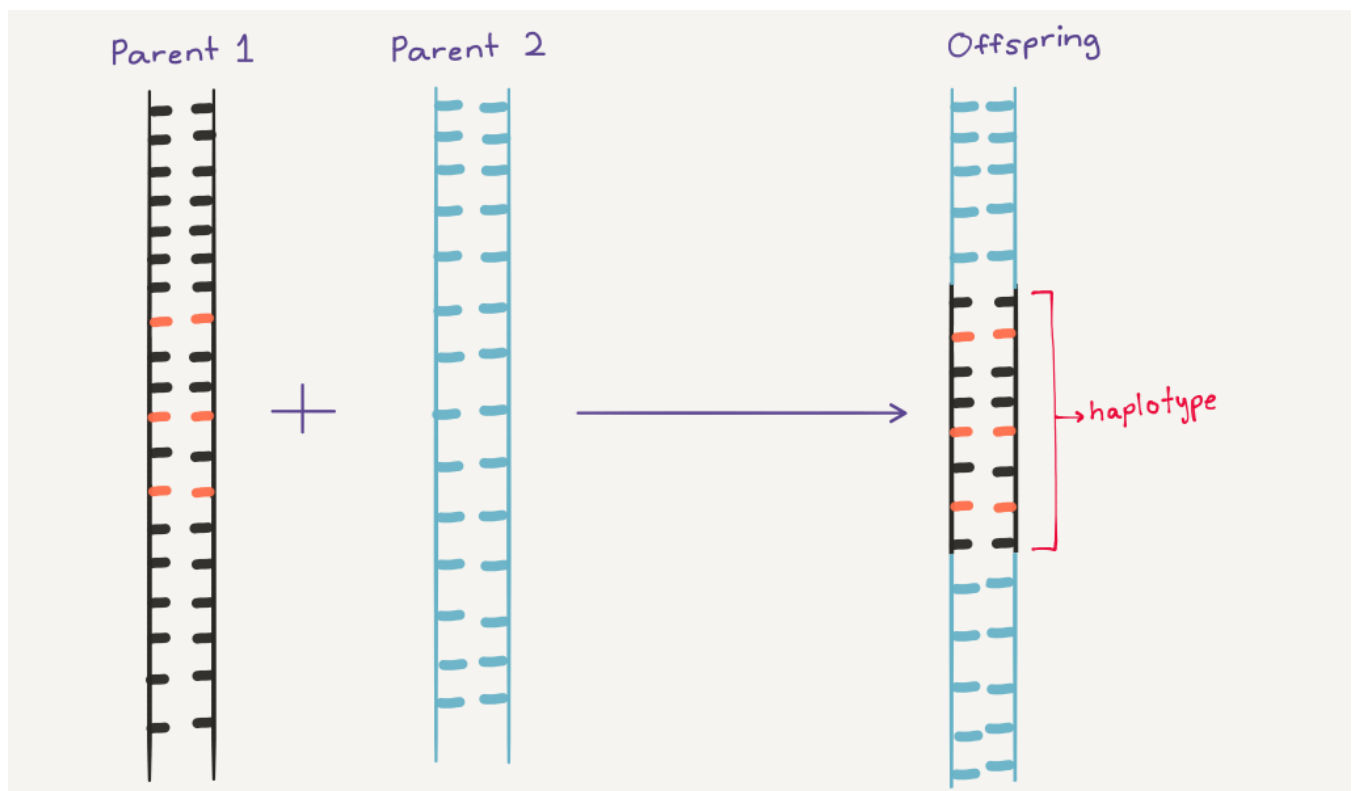
SINGLE NUCLEOTIDE POLYMORPHISM (SNPs)

When a nucleotide is replaced with an incorrect one during DNA replication, the alteration is called a **single nucleotide polymorphism** if it occurs in more than 1% of a population.



Most SNPs occur in non-coding regions of DNA.

If SNPs occur adjacent to each other on the same DNA molecule, they can be inherited together from a single parent. This group of SNPs is called a **haplotype**.



Uses of SNPs

Single nucleotide polymorphisms and **short tandem repeats** are genetic markers.

They can be used as:

- Indicators of disease susceptibility
 - ◆ Although not necessarily the cause of a disease
- Determine genetic relatedness
- Establish evolutionary relatedness

5.5: Inheritance Patterns in a Population

Can population genetic patterns be predicted with any accuracy?

Yes, using DNA analysis and genetic technology.

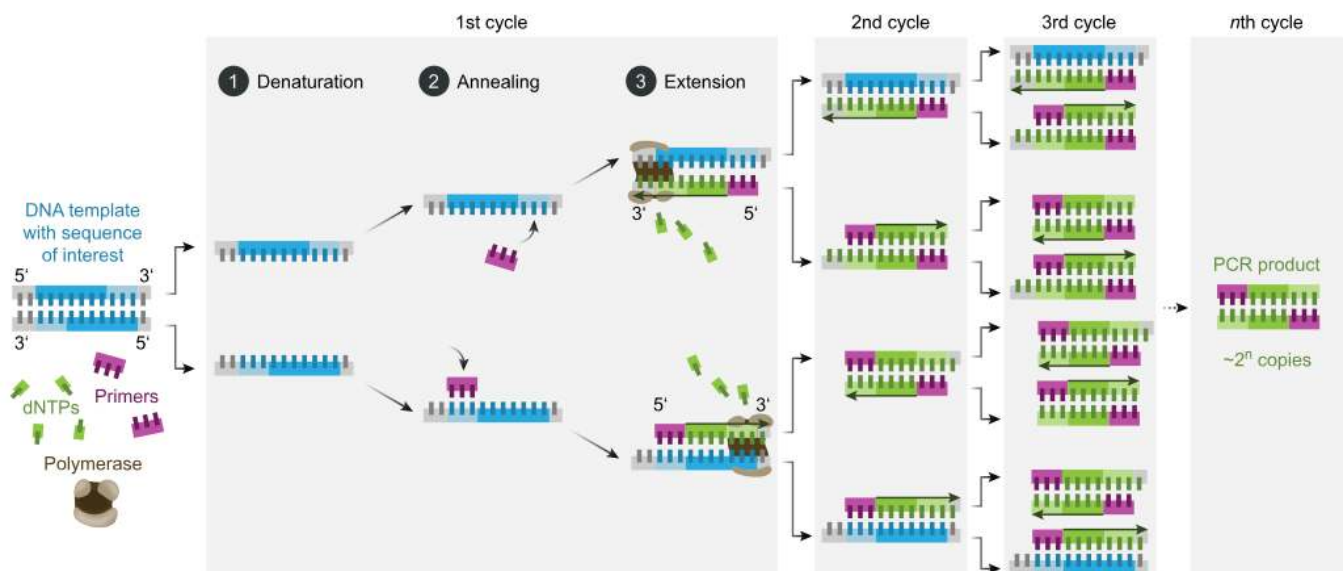
[5.5.1] Technologies Used to Determine Inheritance Patterns

The Human Genome Project

Using DNA sequencing to find the order of nucleotides in every gene of the human genome.

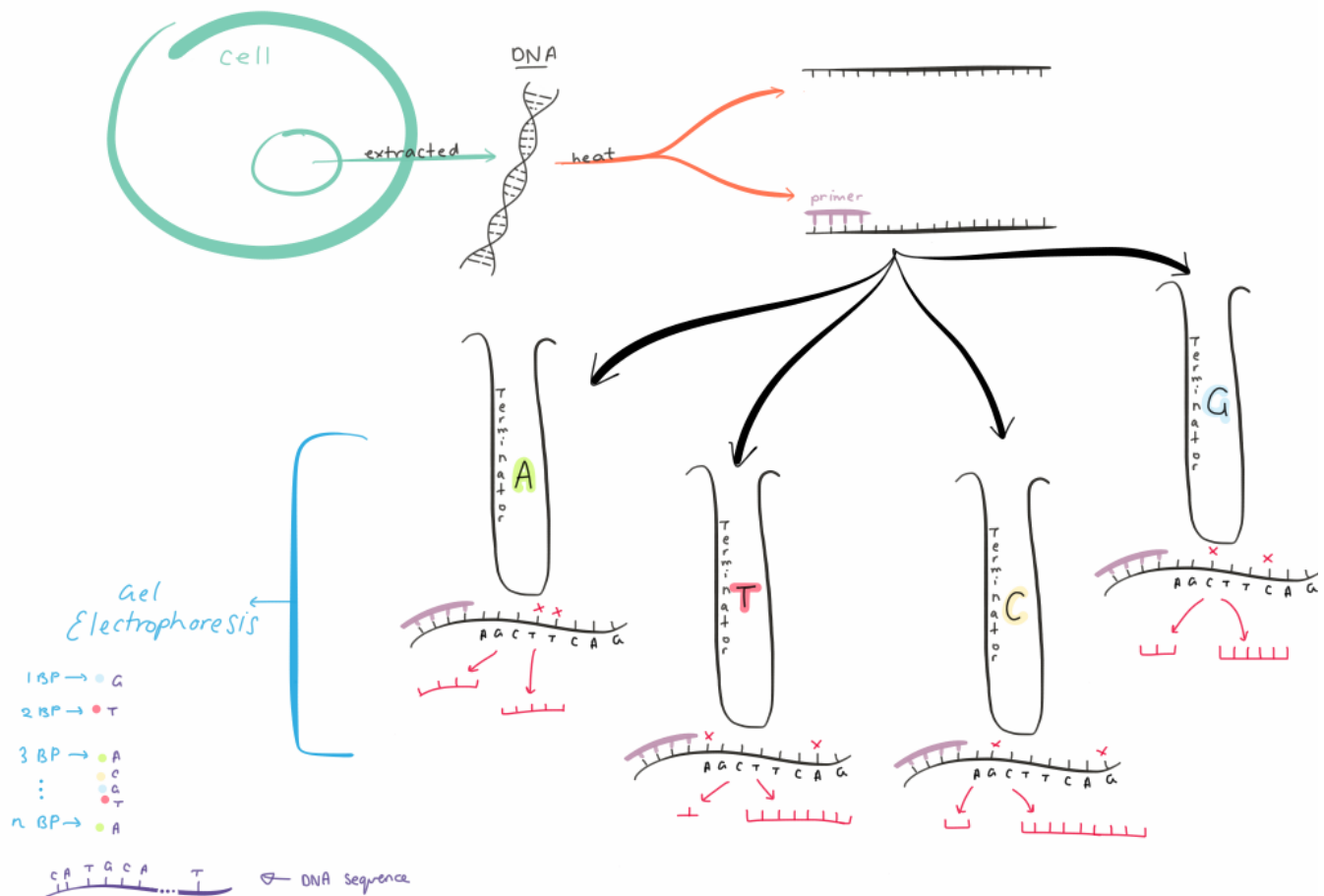
Polymerase chain reaction (PCR)

| | |
|------------|---|
| Isolating | DNA removed from cell. |
| Denaturing | DNA strands are separated into templates using heat. |
| Annealing | DNA primers bind to template strand |
| Extension | DNA polymerase attaches free-floating nucleotides onto template strands |



DNA sequencing (Sanger method)

Obtains a complete sequence of nucleotides for a specific gene or the whole genome.

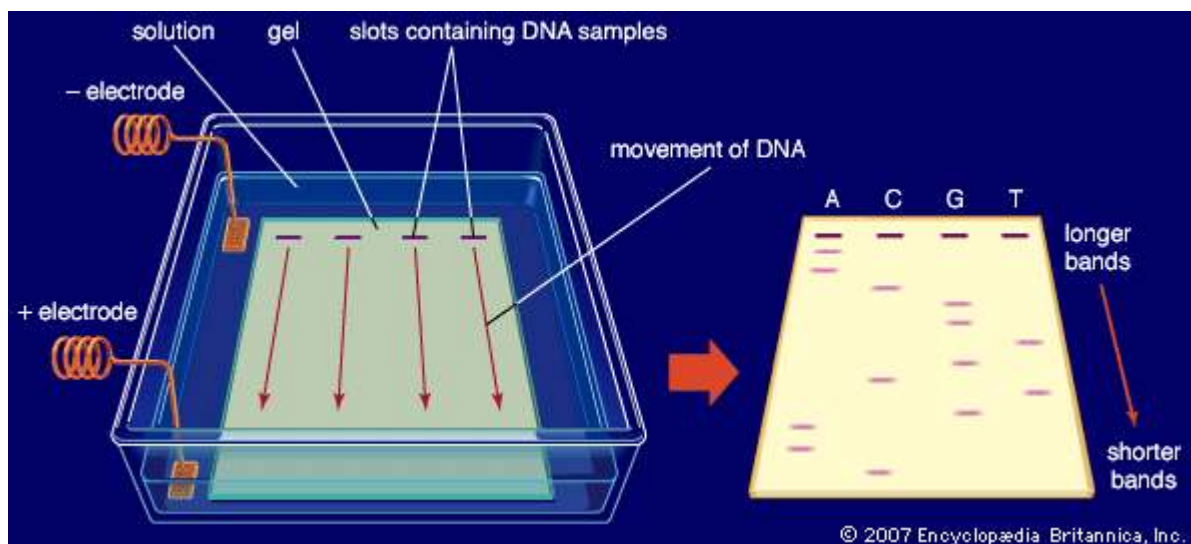
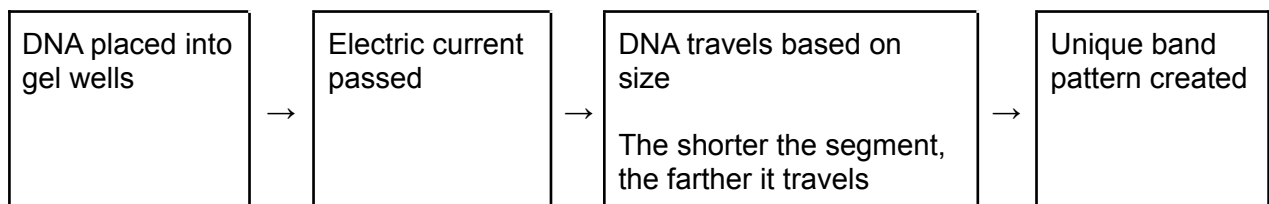


1. DNA isolated from cell
2. Strands are separated using heat, so that terminator nucleotides can attach
3. PCR used to amplify DNA of different lengths
 - a. Primer initiates beginning of replication
 - b. DNA polymerase joins free-floating nucleotides
 - c. The replication is stopped by tagged terminator nucleotides
4. The tagged nucleotides will eventually cover each base pair, creating replicated strands of differing lengths (inclusive of final base)
5. Gel electrophoresis is used to find the lengths (in base pairs) of the replicated strands
6. Laser beam used to identify the marker on each length of DNA
7. This shows the terminator nucleotide for each base pair
8. Thus, the order of the base pairs can be found by taking the complement of the terminator nucleotide

DNA profiling

Obtains band patterns using variable regions of DNA (STRs), which can be used to distinguish individuals.

1. DNA sample is extracted
2. DNA is amplified using PCR
3. Restriction endonuclease enzyme cuts out specific short tandem repeats (**STRs**)
STRs are segments of DNA that are highly variable between individuals.
4. Segments undergo gel electrophoresis



5. Radiolabelling probes are added
6. The band pattern is identified
7. Band patterns are unique to individuals (unique DNA profile)

USES OF THE TECHNOLOGIES

DNA sequencing

- Causes of genetic diseases
- Evolutionary relationships
- Cloning

DNA profiling

Disease testing

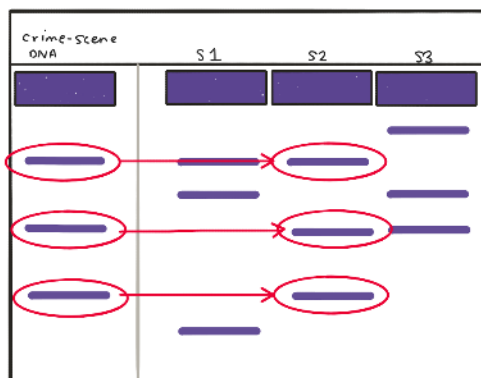
A healthy person, a carrier and a diseased person will have different lengths of DNA sequences. This can be indicated based on how far the DNA moves in electrophoresis.



This individual is most likely a carrier, since the bands match with the known carrier sample.

Forensics

The DNA in crime suspects is compared with DNA from a crime scene. If it is a 100% match, the suspect is guilty.

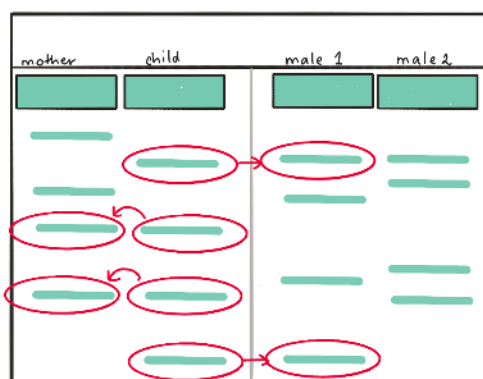


The bands of suspect 2 completely match with that from the crime scene, therefore they are probably guilty.

Paternity testing

50% of a child's DNA comes from the mother and the father.

If one parent is known, the other parent is the one which completes the child's DNA sequence.



The bands from male 1 cover the bands not covered by the known mother for the child. Therefore, he is most likely the father.

[5.5.2] Uses of Population Genetics

CONSERVATION MANAGEMENT

Trends and patterns in population data allow us to:

- Develop strategies for management of endangered species
- Observe extinct species as a comparative model for similar endangered animals

The Tasmanian Devil (endangered)

Issues:

- *Virus-induced cheek cancer*
- *Competition from introduced species*
- *Habitat loss*

Management strategies:

- *Vaccines to treat virus*
- *Raised in fenced habitats with low competition*
- *Storage of complete genome to recreate species if they go extinct*

The Woolly Mammoth (extinct)

Observation of DNA from fossils of 2 groups of woolly mammoths:

1. *Mainland group*
2. *Island group*

When compared to the mainland group, the island group had mutations impacting

- *Their sense of smell*
- *Their urine composition*

Which made it difficult for them to mark territory and mate.

This was because there was less gene flow in the island population → less genetic variation.

Thus, the island group lasted much shorter (4300 yrs) than the mainland group (45,000 yrs).

Koalas (endangered)

Tissue samples collected from koalas to identify genetic variation.

Haplotypes studies to identify disease susceptibility.

INHERITANCE OF DISEASE/ DISORDER

Newborns are screened



DNA sequenced



SNPs or haplotype identified



Large-scale database created for whole population



Genomic studies



Develops: - Preventative strategies
- Treatments

HUMAN EVOLUTION

| MULTIREGIONAL HYPOTHESIS | REPLACEMENT (OUT-OF-AFRICA) HYPOTHESIS |
|---|---|
| <p>Process</p> <p>Homoerectus migrated out of Africa ↓ Groups interbred but became reproductively isolated ↓ Different races emerged ↓ Travel prevented complete speciation</p> | <p>Process</p> <p>1st group of ancient humans left Africa ↓ 2nd group evolved into ancestors of the modern human IN Africa ↓ Group 2 outcompeted group 1 ↓ Group 2 spread into different areas and evolved into the modern human ↓ Reproductive isolation led to the races</p> |
| <p>The diagram illustrates two models of human evolution. A central vertical axis represents time in 'Years ago', with markers at 2,000,000 (Homo habilis), 1,000,000 (Homo erectus), and 50,000 (Modern humans). Both models start with 'African origin for Homo erectus' and show a 'Spread of Homo erectus throughout the World'. The Multiregional hypothesis (left) depicts continuous evolution and interbreeding between populations in Africa, Europe, Asia, and Australia. The Out of Africa hypothesis (right) shows a single migration from Africa around 50,000 years ago, with other groups being replaced or outcompeted.</p> | |
| <p>Evidence</p> <ul style="list-style-type: none"> Fossils of homosapiens show similarities to homoerectus Suggests evolutionary relationship | <p>Evidence</p> <ul style="list-style-type: none"> Carbon dating of fossils suggests ancient and modern humans existed at the same time Common genes in mt-DNA suggest that they interbred |

Using mitochondrial DNA sequencing, we can:

- Establish evolutionary history of a species
- Establish evolutionary relatedness between species

Evolutionary relatedness of two species can be determined by:

- DNA hybridisation
- % similarity of DNA sequences