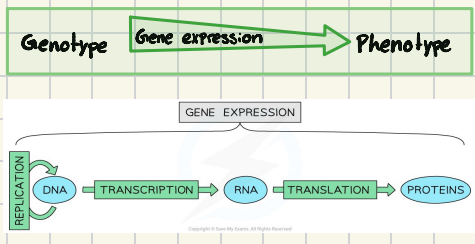
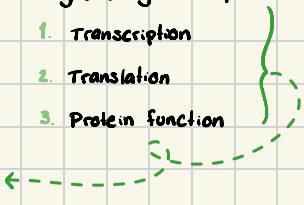


D2.2.1 — Gene expression: phenotype



Stages of gene expression

1. Transcription
2. Translation
3. Protein function



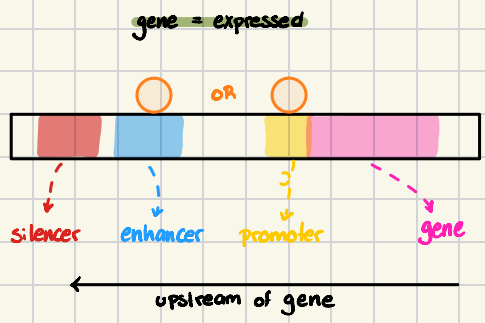
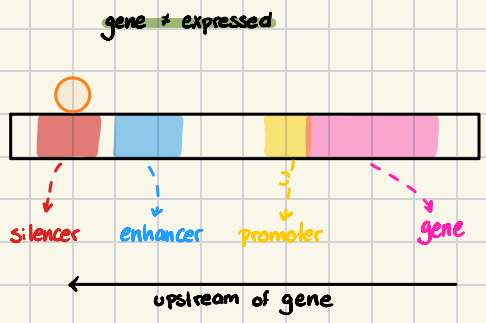
D2.2.2 — Regulation of transcription

Promotor => area/ base sequence where RNA polymerase binds to begin transcription

Transcription factor => DNA binding protein that controls transcription by binding to enhancer/silencer/promoter

Enhancer => base sequence that promotes transcription when transcription factor binds there

Silencer => base sequence that inhibits transcription when transcription factor binds there



D2.2.3 — Regulating translation

↳ Mature mRNA includes poly-A tail

- Tail shortens/degrades @ varying degrees over time

- mRNA molecules w/ shorter tails = less likely to be translated bc they = vulnerable to degradation by nuclease enzymes

o By changing Poly-A tail => cell controls how much protein is made using that mRNA

↳ Can be affected by hormones

D2.2.4 — Epigenesis

Epigenesis → stem cells differentiate into dif. tissues in multicellular organisms (process)

The development of differentiation patterns in the cells of multicellular organisms as it develops from a zygote

↳ Differentiation happens when some genes are activated & others are silenced

Epigenetic tags → chemical modifications that cause some genes to be activated & others to be silenced

↳ Does not affect genes themselves, just the pattern of expression

Epigenetics → genetic control by factors other than an individual's DNA sequence

These can be acquired through life

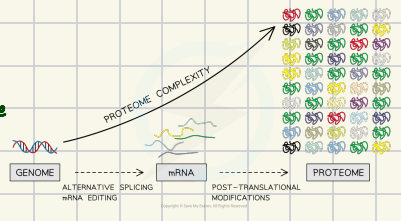
D2.2.5 — genome, transcriptome, proteome

Genome → all the genetic info. in cell

Transcriptome → all the mRNA sequences in a cell

Proteome → all the proteins produced by a cell

varies for individuals
bc cells do not transcribe all genes all the time
depends on what the cell needs @ specific times



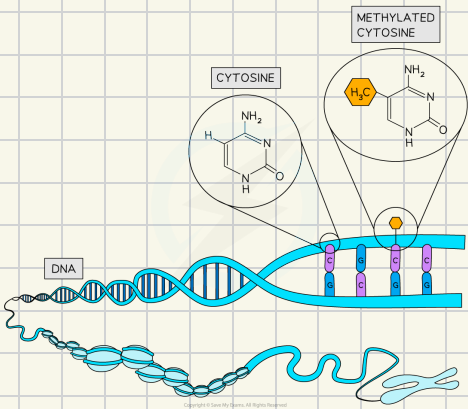
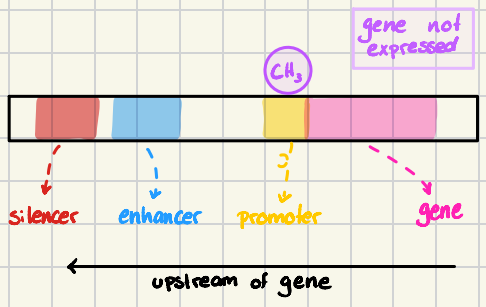
D2.2.6 — Epigenetic tags

Methylation → a type of epigenetic tag that involves addition of methyl group (to DNA or histone proteins)

↳ Methylation of promoter region of gene inhibits transcription

↳ Can change how available DNA is to transcription factors

- When CH₃ binds to histones, it "squishes" proteins together compromising RNA polymerase's ability



D2.2.7 — Epigenetic inheritance

Epigenome → all of the epigenetic tags in an organism

↳ Daughter cells of mitosis receive the genome + epigenome of parent cell

- Ex. → Neuron

1. Neuron undergoes mitosis
2. daughter cells get genome + epigenetic tags that tell cells what genes to suppress
 - i) genes suppressed = those which correspond to blood cell, cardiac cells, etc.
3. This allows daughter cells to look + function like neuron

Epigenetic inheritance → inheritance of epigenetic tags

↳ Both in mitosis — daughter cells — and meiosis — gametes — produced cells will inherit epigenome

In meiosis, however, some tags are removed
↳ "reprogramming of genome"
↳ "clean slate"

D2.2.8 — Environmental effects

↳ factors in the environment can change the pattern of genetic expression

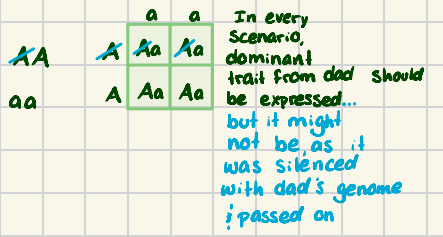
- Ex. → Air pollution decreases DNA methylation

- a) More gene transcription
- b) More immune system proteins produced [Genes that get commonly unmethylated]
 - i. Asthma
 - ii. Heart attack
 - iii. gestation

D2.2.9 — Consequences of removing epigenetic tags from ovum & sperm

Genomic imprinting → when epigenetic tags remain on DNA during gamete formation & are passed to offspring

↳ Can disrupt typical genetic inheritance if dominant allele = silenced by imprinting



Ex → Lions

↳ Female lions can carry offspring from multiple males @ same time

- Females → advantage = having lots of cubs
 - Imprinting for large litters
- Males → advantage = having biggest cub
 - Imprinting for large offspring

usually evens out

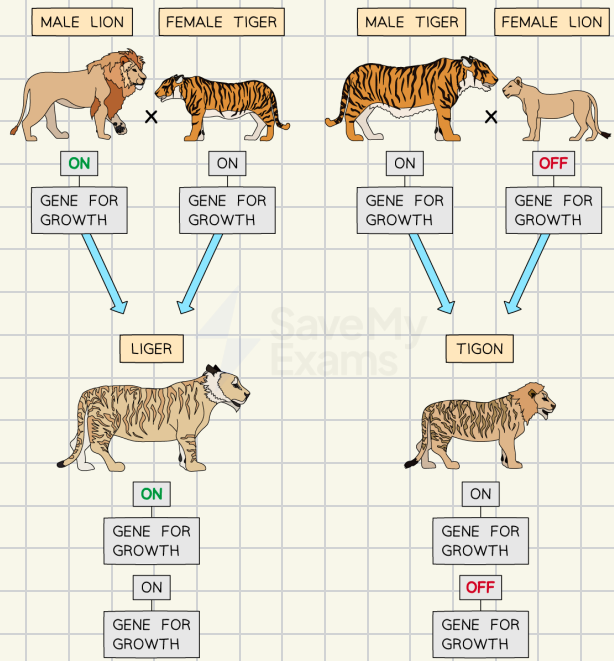
↳ Tigers & lions can interbreed

- Female tigers can only carry a litter from single male @ a time
 - No imprinting necessary in male tigers
 - Their cubs don't need to compete w/ other male's cubs

male tiger + female lion = small

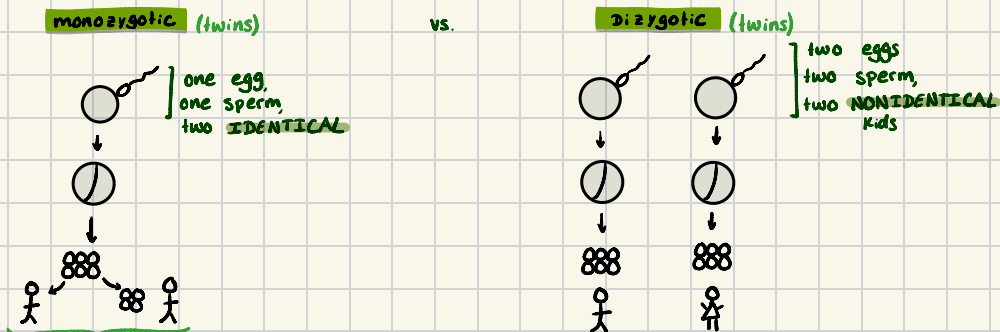
vs.

male lion + female tiger = large



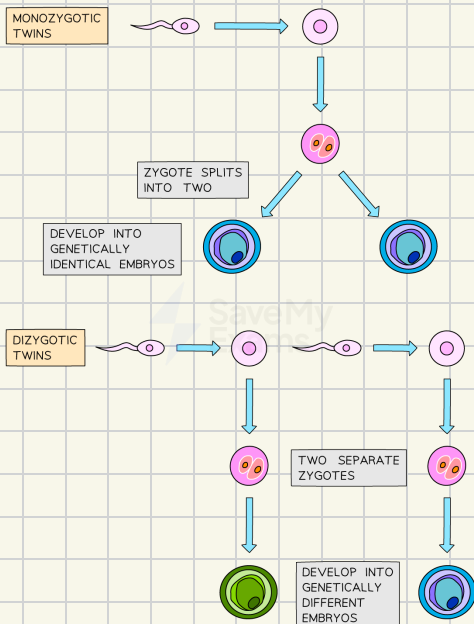
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D2.2.10 — Monozygotic twin studies



Monozygotic twins = valuable for research bc. they have same genome (DNA) but acquire dif. epigenetic tags throughout lifetime.

Nature vs. nurture
↓ genome ↓ epigenetic tags



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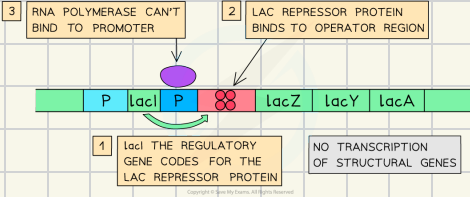
D2.2.11 — External factors

↳ Lactose = example of a **biochemical factor** that impacts gene expression

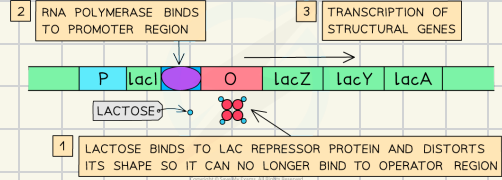
- Lactase gene is normally silenced

- when **lactose levels rise**, repressor protein = deactivated ! gene = expressed
- when **lactose levels fall**, repressor protein reactivates ! silences gene (no more lactase)

lactose absent



lactose present



↳ Oestrogen = example of **hormonal factor** that influences pattern of gene expression

- Hormone that binds to target genes ! changes their expression

- Can bind to gene coding for progesterone receptors in the endometrium

1. Oestrogen turns on this gene
2. Progesterone receptors = produced
3. Tissue becomes more responsive to progesterone

1 hormone impacting effect of another