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91159



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**Mana Tohu Mātauranga o Aotearoa** New Zealand Qualifications Authority

# Level 2 Biology 2024

## 91159 Demonstrate understanding of gene expression

Credits: Four

Achievement	Achievement with Merit	Achievement with Excellence
Demonstrate understanding of gene expression.	Demonstrate in-depth understanding of gene expression.	Demonstrate comprehensive understanding of gene expression.

Check that the National Student Number (NSN) on your admission slip is the same as the number at the top of this page.

#### You should attempt ALL the questions in this booklet.

If you need more room for any answer, use the extra space provided at the back of this booklet.

Check that this booklet has pages 2–16 in the correct order and that none of these pages is blank.

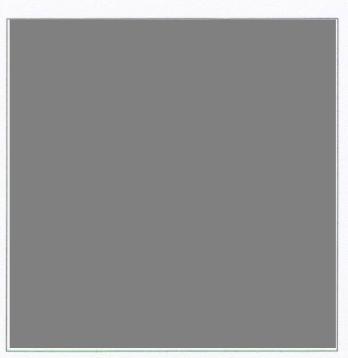
Do not write in the margins (﴿﴿﴿﴿﴿﴾). This area will be cut off when the booklet is marked.

YOU MUST HAND THIS BOOKLET TO THE SUPERVISOR AT THE END OF THE EXAMINATION.

nt TOTAL

### QUESTION ONE: Protein synthesis

	ONE: Protein synthesis te the table below to describe three ways in water.	hich mRNA and DNA are different from
ifference	DNA	mRNA
1	DNA contains deoxyribase Sugar, a phas phente and basks	mand contains ribose sugar, a phosphate and bases
2	DNA consist of Admine, thymine, quanine, and cytosine. (A,T, G, C) A pairs with T G pairs with C	mRNA in a mRNA (hain, the Thymine (T) bases are swapped out by wricle. (U) mRNA like DNA, mRNA has Adnine. quanine and Cytesine too.
3	DNA has the Strands in a double helix Colouble Stranded	MRHA is only one stranolongle which combines with one of DNA'S stranols
	. 11	
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Overview of protein synthesis.

Compared to mRNA, the structure of tRNA increases its stability, which increases the period of time that tRNA lasts in the cytoplasm. In contrast, the majority of mRNA molecules break down within a few hours after their release from the nucleus into the cytoplasm.

(b) Compare mRNA and tRNA, including their function, structure, stability, and role in protein synthesis.

In your answer, include discussion of:

- the function of both mRNA and tRNA
- how mRNA and tRNA work together to carry out protein synthesis
- the significance of tRNA being stable and mRNA being unstable.

mkNA (messanger RNA) is a type of RNA which sends signals (or messages) to parts of the body.

Like mkNA, tRNA (transfer RNA) is also a type of RNA which plays a role in the transport transfer of bases. Hessanger RN They are both used in protein synthesis.

Protein synthesis is the prosess where proteins amino acids (made & from proteins) are formed and polypeptide (amino acid choins are created, proteins code for different

7 Transcription is first phase proteinsynthe froits and functions throughout the body. Protein synthesis begins , with DNA. DNA is made up of two strands, held together in a double helix shape. The stre is called the coding strand 5' to 3' and the other template strand which runs in the opposite direction: from 3' to 5' enzyme called primase DUX pollmerase unzips these two strands alon new complementary strand (MRNA) to combine with the orgine strand. The bases MRNA compliments itself with the following the base-pairing rule States Adidine pairs with Thymine and pairs with Cytosine except mRNA strand, thymine is Uricle. mRNA is transt the cytoplasm for the protein synthesis synthesis: transcription and anstation the mall is then the ribosome where transport (tRNA) takes each "codon" of bases of three bases)

and to These sets of three bases or 'triplets'
are the building blecks to amino acids.
tRNA continues to do this in the translation
phase ultimately resulting in a
pelypeptide chain (large chain at
amno acids.)
it is important that mRNA is unstable and tRNA is stable as mRNA needs
to be able to adapt itself to complement the template strand of DNA and be
transfered around the body, whereas
fRNA has a more careful and percise
rate of forming codons and triplets

#### **QUESTION TWO: Mutations**

The table below shows the abbreviations for some of the amino acids and DNA triplets that code for them.

Amino acid	Abbreviation	Triplets on the DNA template strand
Valine	val	CAA, CAC, CAG, CAT
Proline	pro	GGA, GGC, GGG, GGT
Threonine	thr	TGA, TGC, TGG, TGT
Histidine	his	GTA, GTG
Glutamic acid	glu	CTC, CTT
Leucine	leu	AAC, AAT, GAA, GAC, GAG, GAT

One function of haemoglobin is to assist with transport of oxygen.

There are many different variants of haemoglobin. The sequence of bases in DNA for the normal gene and for the mutated version that codes for the  $\beta$ -globin polypeptide are shown in table (i) below.



Haemoglobin molecule.

- (a) Complete the following TWO tables to show the mRNA, tRNA, and amino acid sequences, using the resource information above.
  - (i) Code for **normal** β-globin polypeptide

Triplet number	1	2	3	4
DNA template strand	TGA	GGA	CTC	CTC
mRNA (codon)	ACU	CCU	GAG	aøa
tRNA (anticodon)	TGA	GGA	CTC	CTC
Amino acid	thr	pro	glu	glu

#### (ii) Code with **mutated** $\beta$ -globin polypeptide

Triplet number	1	2	3	4
DNA template strand	TGA	GGA	CAC	CTC
mRNA (codon)	ACU	CCU	aya	GAG
tRNA (anticodon)	TGA	GGA	CAC	CTC
Amino acid	thr	pro	val	glu

(b) Discuss the effect of this mutation on the amino acid sequence and the functioning of the final β-globin polypeptide.

In your answer, include discussion of:

· the causes of mutations

- · the mutation type and the severity of this type of mutation compared to other types
- the effect of this mutation on the functioning of the final haemoglobin beta polypeptide.

A mutation is a perminant change in the base Sequence of DNA and are caused by mutagens (factors that can alter DNA). Some mutagens include UV light, enzu radiation, and chemicals. These can cause point mutations re which can result it different proteins being formed and could also cause harm however, some mutations do not. The three point mutations are Insertion, deletion and substitution. Insertion where a base in the template strand of DNA gets added or 'inserted' into the cade. Deletion is the opposite and occurs when an already existing base from the template strand of DNA is removed or 'deleted' from the code, Substitution is where a base from the template strand of DNA gets swapped out with another base. For exarmple: AGC -> AAC the quanine base replaced with a adinine base. codon is a set of three bases (eg: TGA) that code for a specific protein and. (eg: threonine). These proteins have different important rules throughout the body, When mutations occur, they can disrupt the Sequence of bases resulting in different
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ino acids being formed. This can be harmful as certain proteins may not be able to form or could arrise. Other times, mutations new proteins be silent mutations meaning no obseribal effect on an individual The mutation in the example on page a substition mutation as the A is swapped for an Adinine, has resulted Valine as, instead of gluthaic, Because of this mutation, glotmic and unable to forefill its purpose in the

#### QUESTION THREE: Environment and gene expression

Anaemia is a condition that develops when a person's blood produces a lower-than-normal amount of healthy red blood cells.

Iron is an essential element in all living cells. Approximately 75% of an individual's total iron is associated with a compound called 'haem' (a part of haemoglobin found in red blood cells), which is responsible for oxygen transport. Part of the metabolic pathway for haem synthesis is shown below.



Haem synthesis.

A lack of iron and high levels of lead in the body can limit the synthesis of haem. Lead is a known poison that inhibits enzymes. If haem synthesis does not occur when needed, anaemia can develop.

Discuss how genes, enzymes and materials in the environment can change the rate of haem production. In your answer, include discussion of:

a metabolic pathway

- how the metabolic pathway for haem synthesis occurs, using the terms: substrate, enzyme, gene, and final product
- how DNA mutations, low iron, and high lead can all cause anaemia.

A metabolic pathway is a series of engine reactions greats which lead to an protein being formed. The diagram above is a metabolic pathway as it includes enzymis, and substrates leading to the final product (Haem). The pathway begins with the substrate precursor which combines with a si enzyme & from there, the intermediate substrate is created and combines was the active site with enzyme 2. Eventually, the metabolic pathway will come to the end Biology 91159, 2024 and resulting in 07840 a product (in this case Haem).

However, we can see that enzyme 2 is met with lead and iron which can after the outcome of the final product. lead 15 known poision that inhibts and is able to limit the production of and limiting factor in a metabolic pathway Another factor are enzyme mutations and can be harmful for many reasons. High levels of lead, low iron ar mutations may cause result in a build up of substrate, in this case the intermediate Substrate which can be very harmful and toxic, as too much of anything can overwhelm the body. This could also result in future energines and substrates being made and utimatley, preventing If a final product at all. This +5 Extremty can be very bad as haem Synthesis won't be able to Occur resulting in the development of anaemia, and less red blood cells.

## **Achievement**

Subject: Biology

Standard: 91159

Total score: 11

Q	Grade score	Marker commentary
One	A4	This response outlines two differences between DNA and mRNA, describes protein synthesis, details the function of mRNA, and describes the transcription process.
Two	A4	This response explains what a mutation is, describes a substitution mutation, connects a negative consequence of a mutation to the final protein, and details a deletion mutation.
Three	А3	This response describes what a metabolic pathway is, describes a substrate, and identifies how lead inhibits the functioning of enzymes.