DNA, RNA and Protein Synthesis

Protein synthesis isn't easy, but imagine the warm, glowing feeling you'll get once you've worked your way through all these questions on it. Can't imagine it? Never mind, give the questions a go and you'll soon feel it.

1

	The average human cell contains around 2 m of DNA. The average human cell nucleus is only 10 μm in diameter.	
1.1	Describe how the DNA in a eukaryotic cell is arranged so that it can fit into the nucleus.	
1.2	Prokaryotic DNA is not stored in a nucleus. Give one other difference between the way that DNA is arranged in eukaryotic cells and prokaryotic.	(2 marks) yotic cells.
	It is estimated that the DNA in a human cell contains around 20 000 protein-encoding genes. These genes correspond to only 1.5% of the DNA sequence.	(1 mark)
1.3	Using the information provided above, calculate the average length of a human gene in metres.	
	Length of gene =	m (2 marks)
1.4	What name is given to the complete set of genes present in a cell?	(2 marks)
1.5	What name is given to the complete set of proteins that a cell is able to produce?	(1 mark)
		(1 mark)
	In prokaryotes, around 90% of the DNA is protein-encoding.	(
1.6	Suggest two reasons why prokaryotes have a much greater percentage of protein-encoding DNA than humans.	
	1	
	2	
1.7	Protein-encoding DNA leads to the production of mRNA. Other parts of the DNA encode functional RNA. Give two examples of functional RNA.	(2 marks)
	1	
	2	(2 marks)

2	Leigh syndrome is a metabolic disorder that affects the central nervous system. It can be caused by a mutation in the MT-ATP6 gene, which is located in the mitochondrial DNA.										
2.1		Give two differences between the structure of DNA found in the mitochondria and the structure of DNA found in the nucleus.									
	1										
	2										
	2		own the DM to many wife both to mean	(2 marks)							
2.2	What name is give	n to the location that a gene of	occupies on a particular DNA molecule?								
	A Intron										
	B Exon										
	C Allele										
	D Locus	and the state of t									
	D Locus			(1 mark)							
	Table 1 contains so	ome of the DNA codons that	code for particular amino acids.								
			Table 1								
		Amino acid	DNA codon								
		Isoleucine	ATT, ATC, ATA								
		Glutamic acid	GAA, GAG								
		Leucine	CTG, TTA, TTG								
		Methionine	ATG								
		Valine	GTT, GTC, GTA, GTG								
		Arginine	CGG, AGA								
		Alanine	GCT, GCC, GCA, GCG								
2.3	Give one piece of	evidence from Table 1 that sh	nows the genetic code is degenerate.								
	14			(1 mark)							
	Figure 1 shows one of the mutations in the MT-ATP6 gene that can cause Leigh syndrome.										
	Figure 1										
	Original gene: CAA CCA ATA GCC CTG GCC GTA										
	Mutated gene: CAA CCA ATA GCC CGG GCC GTA										
			53 154 155 156 157 158								
2.4	Describe the effect that the mutation shown in Figure 1 will have on the mRNA sequence product the MT-ATP6 gene.										
	5'05'2 5'50'2 2-50'0 5'5 1 35'0										
				(1 mark)							

2.5 Using Table 1 for reference, describe the effect that the mutation shown in Figure 1 will have on the amino acid sequence produced from the MT-ATP6 gene.	ie
2.6 MT-ATP6 codes for a subunit of ATP synthase, an enzyme involved in respiration. Explain how a change in its amino acid sequence could affect the function of ATP synthase.	mar
2.7 Describe how the mRNA produced from the MT-ATP6 gene is translated into a protein.	nark
Transcriptomics involves studying the RNA present in a cell. One technique involved in transcriptomics is described in Figure 2 .	arks
All of the mRNA is extracted from a cell. An enzyme is used to convert the mRNA into complementary DNA (cDNA). The sequence of the cDNA molecules is determined. This allows the mRNA molecules to be identified. The sequence of the cDNA molecules is determined. This allows the mRNA molecules to be identified.	le
Describe how mRNA is produced from DNA by RNA polymerase.	
(4 mar	

A team of scientists have developed a new drug. The team used the method in **Figure 2** to investigate how the levels of three different mRNA molecules changed when eukaryotic cells were treated with the drug.

Figure 3 shows two images. One represents the cDNA for one of the mRNA molecules. The other represents the original DNA strand from which the mRNA was produced in the nucleus.

	cDNA	Figure 3	Original DNA	
	e luncii ATP sendinse	a always sure		ing line
3.2	Explain why the cDNA and the origina	ll DNA shown in Figu	re 3 are different.	
	18/41			
				(2 marks)
	The results of the scientists' experimen		4.	
	7,	Figure 4		
	6-		Key:	
	- vel		Untreated =	
	Relative mRNA level		Treated =	
	절 4-			
	¥ 3 -			
	ita 2 -			
	ž ₁			
	0			
	mRNA 1	mRNA 2 mR1	NA 3	
	The scientists hypothesised that the new Method 1: By preventing RNA polymethod 2: By destroying particular materials and the scientists hypothesised that the new Method 1: By preventing RNA polymethod 2: By destroying particular materials and the scientists hypothesised that the new Method 1: By preventing RNA polymethod 2: By destroying particular materials and the scientists hypothesised that the new Method 1: By preventing RNA polymethod 3: By destroying particular materials and the scientists hypothesised that the new Method 1: By preventing RNA polymethod 3: By destroying particular materials and the scientists hypothesised that the new Method 3: By destroying particular materials and the scientists hypothesised that the new Method 3: By destroying particular materials and the scientists hypothesis hypothesi	nerase from working.	ole methods of action:	
3.3	With reference to Figure 4, explain wh	ny the drug cannot be a	acting via Method 1.	
	- A - Control -		needyes e	
				(2 marks)
2 4		4 11-i1	if the drug acts via Mathod 2	(2 marks)
3.4	Explain how the results shown in Figure	re 4 can be explained	if the drug acts via Method 2.	
				(2 marks)
				(2 marks)
			0 30 52 12	Score



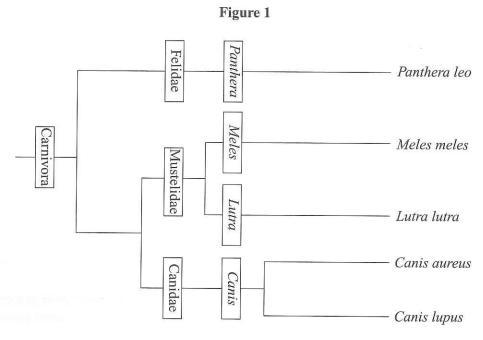
Make sure you know the basics of DNA and RNA structure, as well as the relationship between the base sequence in DNA, mRNA and tRNA. If you do, questions like 2.4 will be easy marks in the exam. It's important to remember that U replaces T in RNA too.

35

Diversity, Classification and Variation — 1

It is due to variations in the genetic code that there is such a great diversity of life on Earth. And because there's so much diversity, scientists find it easier to classify organisms into groups. There's a lot to remember for this section, but don't worry — these questions are here to help you make sure you're all set for your exams.

1 Figure 1 is a phylogenetic tree. It shows how different species from the order Carnivora are related.



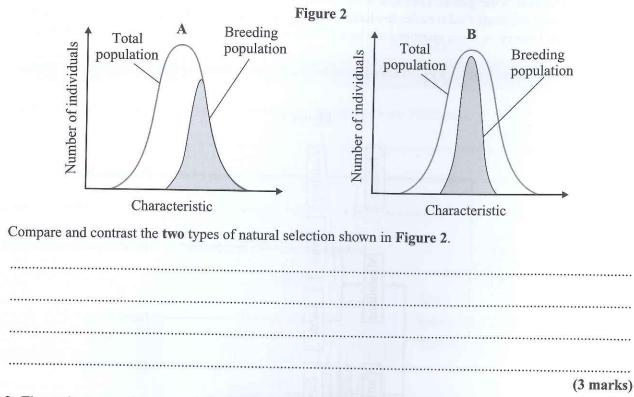
1.1	Wha	it is the defin	nition of a species?	
		······		
	*****			(1 mark)
1.2			nus is genera. rent genera are represented in Figure 1 ?	(1 11111)
	•••••			(1 mark)
1.3	Whi	ch two speci	es in Figure 1 are most closely related? Give a reason for your answer.	(1111111)
			· · · · · · · · · · · · · · · · · · ·	
				(1 mowle)
1.4	Wha	t taxon is rep	presented by the groups Felidae, Mustelidae and Canidae in Figure 1?	(1 mark)
	A	Kingdom		
	В	Phylum		
	C	Class		
	D	Family		

(1 mark)

2.1

2 Species become better adapted to their environment via the process of natural selection.

Figure 2 shows two populations (A and B) experiencing natural selection.



In **Figure 2**, one graph represents the selection on height faced by a plant population in a dense forest. The other graph represents the selection on height faced by a plant population in open grassland.

2.2 Using this information, state which population (A or B) is from the forest. Explain your answer.

2.3 Suggest why the plant population in open grassland is undergoing a different type of selection

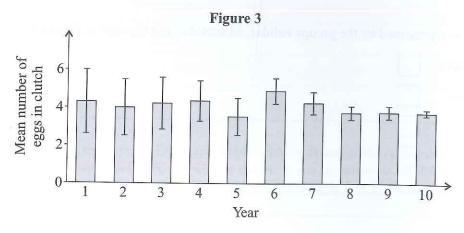
to the one in the dense forest.

(1 mark)

Clutch size is the number of eggs laid by a female bird during one breeding season.

Figure 3 shows the mean number of eggs in a clutch over several years for a bird population.

The error bars indicate standard deviation.



		iges shown in Figure 3 .	

			(3 ma
When classifying	mining -		
when classifying	organisms, scientis	sts often look at the proteins found within the organisms.	
3.1 Explain why prote	eins can be analysed	d to help classify organisms.	
C-:	12000 12100		(3 mar
They counted the	ed the amino acid se	equence of a protein in four species, to that in humans.	
	is presented in Tab		ns.
		Table 1	
	Species	Number of differences in amino acid sequence compared to human protein	
	A	25	
	В	7	
	С	1	
	D	7	
Using the information	on in Table 1 discu	ss the scientists' results.	
	Table 1, discu	ass the scientists' results.	
It was concluded that	species B and D w	vere more algority to the	(2 marks
Explain why this is n	ot a valid conclusion	on for this data.	numans.
		·····	
		-	(1 mark)
Standard d. i.i.	(16)		
to change deviation	n is a measure of the	e variation around the mean. Error bars can be used	Score
The second of th	deviation in graphs.	rut simply — the shorter the har the	
standard deviation		Put simply — the shorter the bar, the smaller the out the data is. Not all error bars represent standard ad the question carefully to figure out what they mean.	

Diversity, Classification and Variation — 2

1 Scientists investigated the diversity of plants in an ungrazed field. The data obtained is shown in **Table 1**.

Table 1

Species	Number of individual plants counted in different quadrats					
Rapeseed	24	46	32	28	32	
Common sunflower	1	0	2	1	1	pop emisjika
Common poppy	8	12	6	10	8	
Creeping thistle	13	14	7	15	13	

1.1	Complete	Table 1	to	show	the	mean	number	of	each	species	counted.
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(1 mark)

1.2 Using the mean values you added to **Table 1** and the formula provided below, calculate the index of diversity for plants in this field. Show your working.

$$d = \frac{N(N-1)}{\sum n(n-1)}$$

where N = total number of organisms of all species and n = total number of organisms of one species

d =	***************************************
	(2 marks)

The scientists then gathered data in a second field on which farm animals were allowed to graze. The mean number of creeping thistle was much lower in the second field than in the first.

	The mean name of the contract
1.3	The scientists wanted to determine if the difference in means between the two fields was significant. State which statistical test they could have used to determine this. Explain your choice.
	(2 marks)
1.4	Further investigations in both fields showed the overall biodiversity of the grazed field to be lower than that of the ungrazed field. Suggest an explanation for this.
	(2 marks)

2	Scientists wanted to investigate the impact of different farming practices on ladybird biodiversit To do so, they counted the number of different ladybird species on organic and conventional farmathem to compare the species richness of the ladybirds in the different types of farmathem.	me
2.	1 Explain why it may have been more useful for the scientists to compare indexes of diversity for their investigation.	
		(2 marks)
	The scientists' data can be seen in Figure 1. The error bars indicate standard deviation.	250
	Figure 1	
2.2	Wean number 12 - Organic Conventional Farm type	ladybird

2.3	Describe and explain two ways in which the scientists could have ensured that the results they obtwere representative of the farms sampled.	ained
	1	
	2	
		2 marks)
3	A student wanted to investigate the effectiveness of different types of antibacterial hand sanitiser against a type of bacteria found on the surface of the skin. She was provided with paper discs, three different types of hand sanitiser, a bottle of bacterial broth culture and an agar plate.	
3.1	The agar plate that the student used would have first been autoclaved. Explain why.	
	P) moulto)

3.2	Describe a method that the student could use for her investigation. Include details of the aseptic techniques she should carry out.	
	Figure 2 shows the student's results.	(5 marks
	Paper discs B C	——— Agar plate
3.3	Disc A did not contain any antibacterial hand sanitiser. Explain why it was used.	
		(2
	The area of the inhibition zone surrounding each paper disc indicates the effective antibacterial hand sanitiser.	(2 mark eness of the
2 1	Complete Table 2 by calculating the areas of the inhibition zones in Figure 2 .	NIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII
3.4	Table 2	= The area of a circle can = be calculated using the
	Disc. Area of Inhibition Zone / mm ²	formula πr^2 , where $r = \text{radius of the circle.}$

Disc	Area of Inhibition Zone / mm ²
В	The second secon
C	
D	

(1 mark)

		•••••••			***********	**********		
			************	***********	••••••	***********		
		**********		***************************************				
			••••••	***********	***********			
The diploid num	ber for human	cells is	46. Using	the form	ula prov	ided be	low, calculate the po	(4 mar ossible
number of differ number of comb		ns of ch	iromosome	s followi	ng meio	sis in h	umans.	
where $n =$ the nu	mber of homolo	ogous c	hromosom	e pairs				
	numbei	r of diff	ferent comb	oinations	of chroi	nosome	es =	
atau syndrome	is a rare chromo	neomal	disorder					(1 ma
Figure 3 shows t				Patau syn	drome.			
			Figu	ire 3				
	10	B B	20		20	2.5		
	1	2		a a -	4	5	8	
	# # 5 G	900	12 %	1 54	8 8	2.2		
	6	7	8 9	10	11	12		
	483	38	á II	23	AH	12		
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	19	20	24	46	ğ	8		
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