Mark scheme - Nucleotides and Nucleic Acids

20	i	wbc do not have cell walls to break open √ wbc are, individual cells / not a tissue, so no separation needed √	1 max	
	ii	disrupts / breaks down / dissolves, phospholipid bilayer / membrane √	1	ALLOW remove bilayer / membrane
		(named) protease √		
	iii	break down, histones / proteins associated with DNA \checkmark	2	ALLOW hydrolytic
		Total	4	
21	i	break / AW, cell walls √	1 (AO 1.2)	IGNORE membranes Examiner's Comments A number of candidates achieved this mark. Many discussed increasing surface area or breaking cell membranes or breaking open
				nuclei, which didn't score any marks.

				IGNORE references to incorrect protease
				Examiner's Comments
	iii	<i>idea that</i> pineapple juice contains DNA √ <i>idea that</i> pH might be too low √	1 (AO 3.4)	Many candidates correctly made the link between pineapple juice and pineapple DNA. Responses that questioned the effectiveness of pineapple protease were not credited. Not were those that suggested pineapple juice would stain the DNA.
				DO NOT CREDIT in the context of washing IGNORE lipase
	iv	(add) detergent / washing-up liquid √	1 (AO 2.7)	Examiner's Comments
			2.7)	A few candidates omitted this question, while many candidates suggested heating or safety precautions, which were not credited.
	v	<u>precipitat</u> ion √	1 (AO 2.7)	Mark first suggestion only
		Total	0	
		YES reasons why it would , work / be successful: 2 max		IGNORE additional unlikely ideas throughout e.g. detergent breaks cell wall, salt disrupts membranes.
		<u>detergent</u> , breaks / disrupts, Y1 (cell) membrane(s) / nuclear envelope		
		OR <u>detergent</u> , releases contents of, cell / nucleus √		
22		OR <u>detergent</u> , releases contents	3 max	ALLOW protease separates DNA from, protein / chromatin
22		OR <u>detergent</u> , releases contents of, cell / nucleus √	3 max	
22		OR <u>detergent</u> , releases contents of, cell / nucleus √ Y2 <u>salt</u> , helps DNA, shed water / precipitate √ <u>protease</u> breaks down, histones / proteins around DNA / proteins attached to	3 max	chromatin ALLOW ORA for N1-N4, e.g. action, should be / ought to be / needs to be, done to e.g. N1 'plant should be crushed to break

		abrasion / grinding / blender \checkmark		Examiner's Comments
		 N2 no RNAase added to remove RNA (from DNA / chromatin) √ N3 no, alcohol / ethanol, added to, precipitate DNA √ N4 temperature not low to reduce, enzyme activity / DNA break down √ 		Knowledge of the reasons for each step in a procedure to purify DNA was poor. Candidates may have been put off by the commands to suggest and justify. Essentially candidates needed to argue yes for the correct steps listed which they could explain the point of, and no for the extra steps that they realised had been omitted.
				Misconception
				Many candidates thought that a crushing stage would be needed to break cell membranes instead of cell walls. Conversely, many candidates thought the detergent would break cell walls instead of cell membranes.
				Many candidates thought that protease would break down DNA instead of its associated proteins such as histones.
				There was misunderstanding of the roles of salt and ethanol to precipitate the DNA (separate it from the aqueous solution).
		Total	3	
23		detergent (1) works as an emulsifier / attracts phospholipid molecules and water molecules (1) it will break up the plasma / nuclear membranes (1)	2	
		Total	2	
24	i	condensation √	1	If additional incorrect answer given, then 0 marks ACCEPT esterification Examiner's Comments Most candidates identified the correct reaction
				involved and stated that the chemical released was water. Esterification also gained credit for some candidates. A minority of candidates

			wrongly answered hydrolysis, with hydrogen given off.
ï	water √	1	If additional incorrect answer given, then 0 marks ACCEPT H ₂ O (correct formula only) Examiner's Comments Most candidates identified the correct reaction involved and stated that the chemical released was water. Esterification also gained credit for some candidates. A minority of candidates wrongly answered hydrolysis, with hydrogen given off.
	 phosphodiester bonds in, backbone / described √ hydrogen / H, bonds / bonding (between chains / bases) √ purine to pyrimidine / A to T and C to G √ ref to correct number of bonds between base pairs (A-T & C-G) √ 	max 3	 IGNORE antiparallel ACCEPT covalent bond in backbone DO NOT CREDIT if other bond mentioned to connect between the two chains DO NOT CREDIT H⁺ bonds IGNORE strength of bond DO NOT CREDIT thiamine / cysteine / adenosine Note: 'Two bonds between A and T and three bonds between C and G' = 2 marks (mp 3 and mp 4) 'Two hydrogen bonds between A and T and three hydrogen bonds between C and G' = 3 marks (mp 2, mp 3 and mp 4) Examiner's Comments Generally this was a well answered question with candidates recalling correctly the base pairs and the relevant number of hydrogen bonds between able to describe the correct location of the phosphodiester bond in the sugar-phosphate
	Total	5	DNA structure, incorrectly identifying them as polypeptides and then going on to list the bonds found in protein structure.

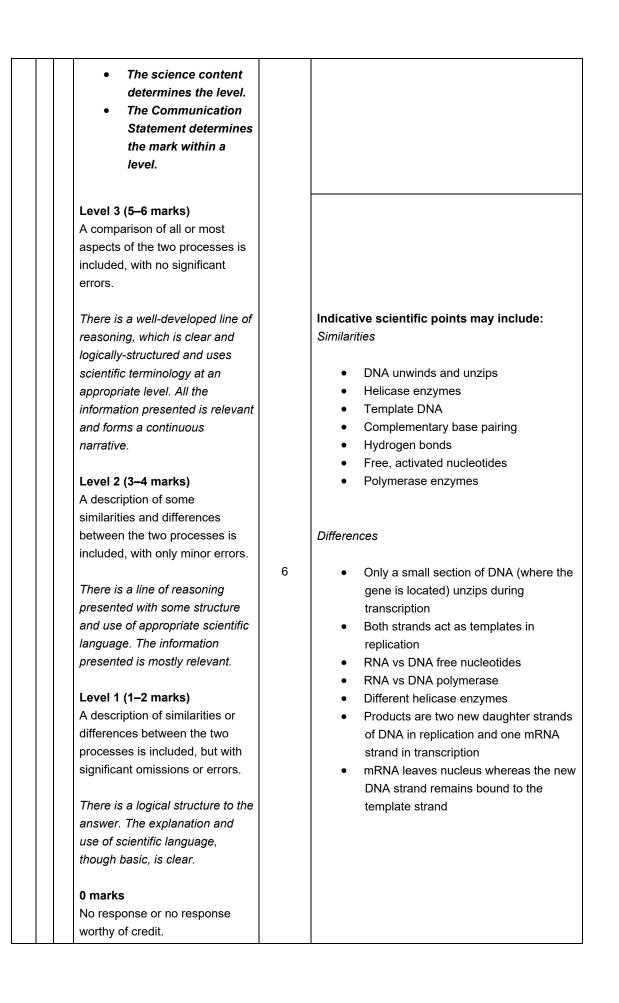
25		 1 nucleotides joined by phosphodiester bonds √ 2 hydrogen bonds between, complementary / named bases √ 3 (polynucleotides) are anti parallel / described √ 4 AVP √ 	3 max (AO1.1)	 1 ALLOW sugar phosphate backbone held with phosphodiester bonds e.g. sense / coding, strand is 5' to 3' antisense / nonsense / template, strand is 3' to 5'
		Total	3	
26		nucleotide ✓ phosphate ✓ pentose ✓ strands ✓	4	If two or more words are given for each gap do not accept contradictory responses ALLOW two Examiner's Comments The majority of candidates were able to gain at least one mark in Q19(a) and the gap fill in Q19(b) enabled the majority of candidates to show their knowledge about DNA structure gaining at least three out of the four marks available.
		Total	4	
27	i	phosphate √ deoxyribose √	3(AO 1.1 2.1)	IGNORE PO ₄ DO NOT CREDIT dioxyribose DO NOT CREDIT if any other sugar mentioned
		guanine √		IGNORE G DO NOT CREDIT if any other base mentioned
		phosphodiester √	2(AO	
	ii	phosphodiester √ condensation √	1.1 2.1)	
	ii	phosphodiester √		

	1		
			DO NOT ALLOW 'ATP has ribose' without
	deoxyribose replaced by ribose		comparison
	\checkmark		Examiner's Comments
	an OH group on carbon 2 of the		
	an OH group on carbon 2 of the, sugar / ribose		This question differentiated well. Candidates needed to recall the structure of ATP and compare that to the nucleotide shown in figure 24. Many candidates accurately identified two similarities and two differences. The most commonly stated similarities 'contain adenine' and 'contain phosphate'. Other similarities were given by more able candidates such as 'contain pentose sugars'. The most commonly stated differences were 'ATP contains three phosphates against one in the nucleotide' and 'ribose compared to deoxyribose'. Very few candidates went into further detail to describe the difference between ribose and deoxyribose. A few candidates lost credit as they did not make a clear comparison under the 'differences' section. The least able candidates appeared to have little concept that ATP is quite similar in structure to a nucleotide. Exemplar 5
			a phosphate group (s) and a nitrogeous ban DMORENCES ATP has 3 phosphate groups, the DNA nucleotide has any 1. ATP has ribose sugar, DNA has elegyribose.
	Total	4	
	<i>three from</i> adenine / A pairs with thymine / T and cytosine / C pairs with guanine / G (1) (because of) hydrogen bonding (1) <i>idea that</i> purine can only bind with pyrimidine because they are different sizes (1)		
29		3	ALLOW 2 H bonds between A and T and 3 H bonds between C and G.
	<i>idea that</i> if one base is known it can pair with only one other base (1)		
	Total	3	

30	i	circle around the two nitrogen containing rings √	1	e.g. OOO HOPOPO OO OO OO OO OO OO OO OO OO OO OO O
	ï	ADP has 2 phosphates whereas DNA nucleotide (with adenine) has 1 phosphate √ ADP has ribose whereas DNA (nucleotide with adenine) has deoxyribose √ or ADP has OH on carbon 2 of sugar whereas DNA (nucleotide with adenine) has no OH on carbon 2 of sugar √	2	Note: a clear comparison between ADP and DNA nucleotide must be made
	iii	condensation √	1	ALLOW phosphorylation
		Total	4	
		hand during the state of the state		
31		bond drawn between phosphate and carbon 3 of sugar and labelled phosphodiester bond √ two bonds drawn between bases T & A and three bonds between C & G and labelled hydrogen bonds √	2	ACCEPT just one phosphodiester bond drawn
31		and carbon 3 of sugar and labelled phosphodiester bond √ two bonds drawn between bases T & A and three bonds between C & G and	2	
31	i	and carbon 3 of sugar and labelled phosphodiester bond √ two bonds drawn between bases T & A and three bonds between C & G and labelled hydrogen bonds √		
	i	and carbon 3 of sugar and labelled phosphodiester bond √ two bonds drawn between bases T & A and three bonds between C & G and labelled hydrogen bonds √ Total	2 1	DO NOT ALLOW adenosine

		ATP used to provide energy for , (named) metabolic reactions / processes √ ATP is , not stored long term / used immediately √		ALLOW ATP produced is coupled to metabolic reactions IGNORE used for respiration unqualified ALLOW ATP is used as fast as it is produced
		Total	4	
33	i	deoxyribose √	1	Examiner's Comments Many candidates correctly identified the circled component as deoxyribose sugar. Pentose sugar or ribose were common responses that were not credited a mark.
	ii	phosphodiester √	1	Examiner's Comments The majority of candidates correctly named the bond as a phosphodiester bond. Weaker candidates gave other names that they could recall such as 'hydrogen bond' or 'covalent bond' which were not given credit.
		Total	2	
34	i	164 706 √√	2	Correct answer with no working = 2 marks If the answer is incorrect, look for a working mark: either (incorrect rounding) ALLOW 1 mark for seeing 164 705 or 164 705.88 or 164 705.9 anywhere or ALLOW 1 mark for any ref to 56 ÷ 34 (e.g. 5.6 ÷ 0.34 or 5600 ÷ 34) Examiner's Comments Most candidates recognised that the number of kB would be obtained by dividing the length of DNA by the length of a kB to arrive at the number of kB in the length of DNA. However, they were not confident converting units of cm and micrometres to standard form, and also

				failed to state the answer to the nearest whole number.
	ii	28 √√	2	Correct answer with no working = 2 marks If answer incorrect, ALLOW 1 mark for seeing 100 – 44 or 50 – 22 Examiner's Comments The majority of candidates correctly scored maximum marks for this calculation, but the most common mistake that was presented was through poor arithmetic, e.g. 100 – 44 = 66.
		Total	4	
		Please refer to the marking instructions point 10 for guidance on how to mark this question.		
35		In summary: Read through the whole answer. (Be prepared to recognise and credit unexpected approaches where they show relevance.) Using a 'best-fit' approach based on the science content of the answer, first decide which of the level descriptors, Level 1, Level 2 or Level 3, best describes the overall quality of the answer. Then, award the higher or lower mark within the level, according to the Communication Statement (shown in italics): • award the higher mark where the Communication Statement has been met. • award the lower mark where aspects of the Communication Statement have been missed.		



		Total	6	
36	i	$2.8 \times 10^3 \sqrt{\sqrt{2}}$	3 (AO2.6)	ALLOW 2.777 x 10^3 or 2.778 x 10^3 or 2.78 x 103 ALLOW 2 marks for 2777 ALLOW max 2 marks for working Each line can be awarded 1 mark: $3000\ 000\ 000\ /\ 50\ =\ 60\ 000\ 000\ (s^{-1})$ $60\ 000\ 000\ (s^{-1})\ /\ 3600\ (s)\ =\ 16\ 667\ (h^{-1})$ $16\ 667\ /\ 6\ (h)$ OR $3\ 000\ 000\ 000\ \div\ 21600\ (i.e.\ 6\ x\ 60\ x\ 60)\ =\ 138\ 889138\ 889\ \div\ 501.08\ x\ 10^6\ /\ 1080000Each line can be awarded 2 marks:300000000\ \div\ 1080000\ (ie:\ 6\ x\ 60\ x\ 60\ x\ 50)OR3\ x\ 10^9\ \div\ 1.08\ x\ 10^6$
	ii	helicase √ <u>DNA</u> polymerase √ AVP √ e.g. (DNA) ligase (DNA) gyrase	2 max (AO1.2)	ALLOW 'helixase'
	ili	 1 enzymes , are (biological) catalysts / speed up reactions √ 2 they lower the activation energy (so reactions can take place at, low / body, temperatures) √ 3 high temperatures (in living organisms), would denature, enzymes / proteins √ 	2 max (AO1.1)	ALLOW enzymes catalyse reactions ALLOW enzymes reduce time taken for reaction Examiner's Comments Where more complex calculations were required, such as in this question, the working of these calculations was often confused and difficult to follow. Candidates needed to calculate how many enzyme complexes were required to replicate the human genome in a set time. However, responses were often incomplete or

					there were bits of working arranged haphazardly around the page.
			Total	7	
37	а	i	(new DNA molecule comprises) one, original / old / parent, strand and one new strand ✓ each strand (of DNA molecule) acts as a template strand (for a new double helix) ✓	1 max	DO NOT ACCEPT 'DNA strand' instead of 'DNA molecule' Examiner's Comments Most candidates gave a perfect answer referring to the DNA molecule comprising one original or parental strand and one new strand. The most common mistake was referring to just having an old strand. As in previous years, many candidates found it difficult to express their answer in a creditworthy manner e.g. 'each strand contains one old strand and one new', 'half of the DNA is replaced with new DNA', 'half of the original strand is used to make the new one'. Most of these answers show a lack of understanding of the difference between a DNA strand and a DNA molecule. Some candidates used logic with the term 'semiconservative' and wrote about 'half the DNA staying the same', without appreciating this process compared with dispersive replication.
		ii	E1 (DNA) helicase √	4 max	Mark the first 2 enzymes mentioned NOTE only award the function mark when linked to the correct enzyme IGNORE ligase

	 F1 unzips the DNA molecule / breaks hydrogen bonds (between complementary bases) / separates the (2) strands ✓ E2 <u>DNA</u> polymerase ✓ F2 forms phosphodiester bonds / joins (adjacent) nucleotides / forms sugar-phosphate backbone ✓ 		F1 IGNORE unwinds the DNA molecule E2 DO NOT ACCEPT RNA polymerase F2 DO NOT ACCEPT forms H bonds ACCEPT checks for errors (in nucleotide sequence) Examiner's Comments
	Also creditworthy E3 gyrase √ F3 unwinds / uncoils, the DNA √		Most candidates named two enzymes, commonly helicase and DNA polymerase. A few candidates described helicase as unzipping one strand, rather than separating the strands, but most correctly described its action. The most common misconception was describing DNA polymerase as pairing up free nucleotides with the template strand, and joining or reforming hydrogen bonds. Candidates should appreciate that hydrogen bonds occur due to complementary bases pairing and then DNA polymerase joins the nucleotides 'vertically', forming the sugar-phosphate backbone with phosphodiester bonds between the adjacent nucleotides. A few candidates did not specify 'DNA' polymerase or stated RNA polymerase. Some candidates attempted to write about DNA ligase and this illustrates the importance of reading the question properly. A very few candidates suggested random enzymes such as
b	<i>tube with generation 1</i> shows (new) DNA / band, contains, light nitrogen / N ¹⁴ , and , heavy nitrogen / N ¹⁵ ✓	2 max	protease. Marks can be awarded from suitably labelled / annotated diagrams ACCEPT shows that (new) DNA is a hybrid

		 (new) DNA / band, made from only, light nitrogen / N¹⁴ √ so a, light / N¹⁴, strand of <u>DNA</u> must be a, template / parent strand, for the new molecule √ <i>tube with generation 10</i> (highest band gets thicker because) more of the <u>DNA</u> is made from only, light nitrogen / N¹⁴ √ 		Could be credited in context of generation 10 instead (but only award once) Examiner's Comments This question was answered poorly. Only a few candidates could apply the information given in the stem, or were already familiar with the experiment, to correctly describe the banding pattern.
		Total	7	
38	i	(involves) DNA polymerase (1) sugar-phosphate backbone (re)forms / condensation reaction between phosphate and sugar (1)	3	
	i	DNA winds into double helix (1)		ALLOW higher level answers e.g. role of DNA ligase in joining sugar- phosphate backbone lagging strand filled in with Okazaki fragments.
	ii	(new molecule consists of) one old strand and one new strand (1)	1	
		Total	4	
39	i	1 gene / DNA, copied / transcribed, to (m) <u>RNA</u> √	2 max	Read all and mark as prose ALLOW used as a template to create / AW, for 'copied to' ALLOW RNA, copies / takes a copy of, gene / DNA DO NOT ALLOW replicated for 'copied'
		 2 (<i>idea that</i> RNA goes to / translation is at) ribosome(s) / RER √ 3 <u>DNA</u>, is too large to / cannot / is not able to, leave <u>nucleus</u> / 		ALLOW ORA <u>'RNA</u> , is small enough to / can / is able to' or just 'RNA leaves nucleus' ALLOW nuclear membrane for 'nuclear envelope' DO NOT ALLOW leave the cell for 'leave

cross <u>nuclear</u> envelope / fit through <u>nuclear</u> pores √	nucleus' <u>Examiner's Comments</u> As a recall question this was done well,	
	particularly with respect to the sizes of mole being able or unable to leave the nucleus vi nuclear pores, and ribosome being the site translation. Some candidates confused translation with transcription.	а
	?	
	Misconception Some candidates misuse language in descr transcription.	ibing
	Correct:	
	 DNA is transcribed into mRNA. (No passive tense) mRNA is a transcript of the DNA. mRNA is a copy of the DNA. 	ote
	 Incorrect: mRNA transcribes the DNA. (Activitiense) mRNA copies the DNA. (RNA polymerase does this). DNA is converted into mRNA. 	e
	Correct final answer gets 2 marks, even i working is shown. Wrong final answer (which may include a 90 stage in the working) = ALLOW 1 mark for seeing any of these:) 252
 90 252 or 90 255	327 × 92 × 3 OR 30 084 OR 981	
or 90 258 √ √	Examiner's Comments	
	Most candidates multiplied the number of an acids in pepsin (327) by the number of time bigger that titin is compared to pepsin (92). These candidates gained 1 mark for arriving the figure 30 084. Only a minority of candidates	s gat

For answers marked by levels		understood that the question information was about the number of amino acids in a polypeptide while the question was asking for the number of bases in the equivalent DNA. Some of those who realised the distinction divided by the number of bases that code for one amino acid in error. The correct process was multiplying by 3 due to the logic that every amino acid in a polypeptide is coded for by 3 bases on DNA. Candidates who followed a different route could calculate the number of bases in DNA coding for pepsin and then multiply by 92, or could add 3 or 6 bases to their final answer for a stop and/or start codon.
of response: Read through the whole answer from start to finish, concentrating on features that make it a stronger or weaker answer using the indicative scientific content as guidance. The indicative scientific content indicates the expected parameters for candidates' answers, but be prepared to recognise and credit unexpected approaches where they show relevance.	6 max	Communication may be via bullet points, a table of comparisons, labelled diagrams or prose. Indicative scientific points may include the following: FIBROUS PROTEINS Properties: • insoluble • elongated / long / rods / filaments / ropes / strands • strong / tough • flexible IGNORE size refs / compact / coiled / bond types / hard Functions: Look for the general category or for a named protein or glycoprotein example with supporting detail. Related categories and examples are paired or grouped together: • for structure collagen in, bone / cartilage / connective • tissue / tendons / ligaments / skin / blood vessels • fibrin + role described • for protection • keratin in, skin / hair / nails
communication statement (in italics) have been met.		to give, elasticity / elastic properties

for, contraction / mechanical movement
 for, contraction / mechanical movement
• actin / myosin, in muscle
microtubules in, cilia / flagella / spindle /
cytoskeleton
GLOBULAR PROTEINS
Properties:
• soluble
 spherical / ball-shaped
 have, 3D / tertiary / 3o, shape / structure
specific / complementary (to another
molecule)
ref. conjugated / contain prosthetic group
• temperature / pH, sensitive
hydrophilic on outside
IGNORE size refs, compact, round, bond type
Functions: Look for the general functional
category name or description, or a named
protein or glycoprotein example with some
supporting detail.
enzymes / metabolic role / to catalyse
reaction(s) / to lower activation energy
 named enzyme + its specific role described
hormones / receptors / for cell signalling
named hormone / insulin + role described
opsonin / antitoxin / agglutinin + role described
fibrinogen in blood clotting
to transport substances across cell membranes
 carrier / channel / pump + role described
to transport substances in blood
haemoglobin + role described e.g. carry
• oxygen
• to, package / organise DNA
to, packago, organico Drivit

	not be clear.		Examiner's Comments
	0 marks No response or no response worthy of credit.		Candidates generally had plenty to say and almost all attempted to fulfil the command word compare by making reference to both fibrous and globular proteins. Level 3 answers fully answered the question brief by providing science content covering at least one property of a fibrous protein and one property of a globular protein and at least one function of each type of protein. The word 'property' was confused by some candidates with structure and this resulted in irrelevant material about levels of protein structure and bonding within proteins. Properties may be physical or chemical and relate to aspects like the solubility, strength, flexibility, shape and stability (for example at different temperatures) of molecules. Exempler 1 <u>fibrour proteins within the body for</u> example, keratin which makes up thing like stim, hair and nails. Globular proteins have more of a spherical shape Thuy are usually souble in water and are more metabolically active than fibrous proteins. Globular
	EITHER 1 9300 / 9700 √		<u>are bath examples of glabular pratries and</u> <u>are involved in character reactions in the body</u> Exemplar 1 is a level 3 answer that is contained within the line space, answers all aspects of the question and contains an appropriate level of science content. There is a well-developed line of reasoning, a clear and logical structure and all the material is relevant and substantiated by fact, so the communication statement for 6 marks is met. Correct answer to 2 s.f. with correct matching units = 2 marks
iv	 <u>deaths year</u>⁻¹ or <u>deaths</u> per <u>year</u> or <u>deaths</u> / <u>year</u> √ OR 3 9.3 / 9.7 √ thousand <u>deaths year</u> ⁻¹ or 4 thousand deaths per year or thousand <u>deaths</u> / <u>year</u> √ 	2	 ALLOW mark for unit even if no or wrong figure given ALLOW minus sign with number or 'fewer' with unit ALLOW from AIDS / of AIDS in unit ALLOW mp 3 so long as the word thousand appears afterwards or in the units (even if the

unit is wrong in another respect) DO NOT ALLOW '9.3 1000 deaths per year' for mp3 (but gets mp 4)
Examiner's Comments
Candidates often achieved one of the two marks available but few successfully worked through all the processes involved in arriving at an answer with appropriate units for the rate of decrease over three years. One error was for candidates to calculate not a rate (over time) but a percentage decrease. A breakdown of how to tackle this question is listed in the 'Assessment for Learning' box. This, together with sections from the three tutorial sheets listed under OCR support, could form the basis of a step-by-step worksheet on solving the problem set in this question. Additional questions could be devised using this graph to calculate rates of increase or decrease in the numbers of new diagnoses or those living with an AIDS diagnosis for different time periods.
AfL
1. Select the dash-dot line for deaths and read to the nearest half-square of the grid where values for 1995 and 1998 intercept the y axis.
2. Check the left-hand y axis label to see that these figures represent thousands.
3. Subtract one away from the other to find the difference.
4. Divide this answer by the time between the two values on the x axis, 3 years.
5. Give the answer to two significant figures.
6. Determine the units.
í
OCR support

40	i	3 bases / triplet, code for 1 (specific) amino acid √	2 max	
		Total	14	
	v	 1 decrease in new diagnoses, from 1992 / already / began before 1995 √ 2 peak / plateau, in deaths, from 1994 / already / began before 1995 √ 3 no change in / same, (rate of) increase in people living with AIDS, before / after, 1995 √ 	2 max	ALLOW from 7 since 7 aner, 1993 (instead of 1992) Examiner's Comments As specified in the question, candidates had to make use of data from the graph in their answer. Ideas from their own knowledge like improved education or increased precautions against transmission of HIV did not therefore score. Strong responses did not just look at 1995 to judge whether the introduction of a drug had an effect (pre-supposing that a change would begin from this point), but instead drew conclusions from ongoing trends that pre-dated 1995. These showed that new diagnoses were already falling, deaths had already peaked and the number living with AIDS experienced no change in its rate of increase.
		(answers must relate to data on graph)		 ALLOW when, saquinavir / drug / medicine, was introduced for '1995' in mps 1, 2 and 3 ALLOW new diagnoses decrease at same time as deaths ALLOW from / since / after, 1993 (instead of
				Tutorial sheets and quizzes are available to support the teaching of the skills listed in the specification for Maths for Biology. Three areas cover key skills needed to successfully answer this question: <u>http://www.ocr.org.uk/qualifications/by- subject/biology-related/maths-for-biology/m3- graphs/</u> (calculating a rate from a graph with time on the <i>x</i> axis) <u>http://www.ocr.org.uk/qualifications/by- subject/biology-related/maths-for-biology/m0- arithmetic-and-numerical-computation/</u> (introduction to significant figures) <u>http://www.ocr.org.uk/qualifications/by- subject/biology-related/maths-for-biology/m0- arithmetic-and-numerical-computation/ (introduction to significant figures)</u>

		sequence of, bases / triplets,		
		determines the sequence of, amino acids / primary structure		e.g. more than one codon codes for an amino acid /
		√		degenerate
		(code) non-overlapping \checkmark		code is, universal / similar in eukaryotes and prokaryotes
		AVP √		
		mechanical strength (to cells) \checkmark		
		cell, support / stability / maintains shape √		
		movement of (named), molecules / vesicles / organelles within cell		IGNORE strength unqualified
	ii	OR holding organelles in position √	3 max	ALLOW maintain internal organisation
		formation / movement, of, cilia / flagella √		
		cell movement / endocytosis / exocytosis / phagocytosis / cytokinesis / described √		
		movement of mRNA from nucleus to ribosome \checkmark		
		movement of polypeptides through the rER \checkmark	2 max	Note: this requires more detail than part ii
	iii	movement of vesicles from rER to Golgi √		
		movement of vesicles between cisternae of Golgi (cis to trans face) √		
		movement of secretory vesicles from Golgi to cell surface membrane√		
		Total	7	
41		С	1	Examiner's Comments
				Candidates found this difficult, many suggesting

			A or D. It is possible that they had misread the question and gave an option that was a correct statement about the genetic code rather than an incorrect one. Candidates should be encouraged to take care when reading questions rather than rushing into answering the question that they thought had been asked.
	Total	1	