

Key Area 1.1: Structure and organisation of DNA

Q5)

Q6) A Q7) C Q8) C

Q9)

A	(i)	<ol style="list-style-type: none">1. Prokaryotes have circular <u>chromosomes</u> and plasmids.2. Yeast has plasmids.3. Circular <u>chromosomes</u> in mitochondria/chloroplasts.4. Linear <u>chromosomes</u> in nucleus of eukaryotes.5. Prokaryotes have circular DNA AND eukaryotes have linear DNA (Only if point 1 or 4 not awarded).6. Linear/eukaryotic/nuclear chromosome/DNA (tightly) coiled.7. Linear/eukaryotic/nuclear chromosome/DNA packaged with/wrapped around (associated) proteins/histones. <p style="text-align: right;">(any 4)</p>
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Key Area 1.2: DNA Replication and PCR

Q1)

(a)		0.24	1	
(b)		32	1	
(c)		Inclusive scale and axes labels copied exactly from table headings 1 Points plotted and joined with a ruler 1	2	
(d)		Only donor 2 is suitable OR donor 2 is most suitable	1	
(e)	(i)	TACTGTTTAGC	1	
	(ii)	Separates strands/splits up DNA strands/breaks H bonds between strands/denatures DNA/unzips DNA 1 Any temperature from 50 - 65 1	2	NOT - splits DNA alone

Q2) A Q3) C Q4) D

Q5)

(a)	(i)	12	1	
	(ii)	1.1 to 2	1	
(b)		Taq- Takes less time to amplify (sequence)/complete a cycle. OR Replicates/cycle/process faster. OR Cheaper as less heat is required/temperature is lower. OR Taq takes 3 minutes and Pfu takes 4 minutes. OR Pfu- Proof reading/corrects errors.	1	Incorrect values negate. NOT- ensures no mistakes are made.
(c)		So it does not denature.	1	NOT- so it can work at high temperatures.

Q6)

(a)	(i)	Any TWO from: Size/mass/of muscle (tissue)/sample. Type of muscle tissue/age of fish. Temperature/pH/time. Volume/concentration/type of solution.	2	NOT - Volume of muscle (tissue). NOT - Mass/species of fish. NOT - Same solution. NOT - "Concentration"/"Volume" alone. Additional incorrect variables (e.g. Light intensity/ oxygen concentration/ CO ₂ concentration) negates 1 mark.
	(ii)	Hydrogen/ionic/disulphide/Van der Waals/hydrophobic/covalent.	1	NOT - Peptide (negates correct answer).
(b)	(i)	Correct scales (0-110 kDa and 0-50 mm) and label. (1) Correctly plotted. (1)	2	If axes transposed scales and labels mark not awarded.
	(ii)	32 kDa or whatever plotted graph shows.	1	
	(iii)	550	1	
(b)	(iv)	1, 2 and 4 each have three bands/proteins in common/at the same distance AND 3 only has one band in common with the other three species. OR They/1, 2 and 4 have more bands/proteins in common/the same/similar or converse.	1	NOT - Proteins 1, 2 and 4 all have band X.

A	(ii)	<p>a. Amplification/produces multiple copies of (target sequence of) DNA.</p> <p>b. (Heated to) 90°C - 98°C to separate strands/denature DNA/break hydrogen bonds.</p> <p>c. (Cooled to) 50°C - 65°C for primers to bind/anneal.</p> <p>d. Primers are complementary to/bind to target sequences/DNA.</p> <p>e. (Heated to) 70°C - 80°C so DNA polymerase replicates DNA/extends new DNA strand/adds nucleotides to new strand/3' end/primer.</p> <p>f. Heat tolerant DNA/Taq polymerase is used.</p> <p>g. Repeated cycles (of heating and cooling).</p> <p>Used in forensic/paternity etc.</p> <p>(any 5)</p>	<p>NOT - Copy/multiplies/replication of/ magnifies DNA</p>

Q8) A

Q9) D

Q10)

(a)		Histone	1	Not - Protein alone. Not - Associated proteins.
(b)		<p>So DNA polymerase can add nucleotides to the 3' end of the new strand</p> <p>OR</p> <p>To give DNA polymerase a start point for replication</p>	1	Not - To start/initiate replication alone.
(c)	(i)	<p>DNA polymerase adds nucleotides to the 3' end of the primer/new strand</p> <p>OR</p> <p>DNA polymerase replicates in a 5' to 3' direction.</p>	1	
	(ii)	(DNA) ligase	1	
(d)		<p>Circular chromosomes (1)</p> <p>Plasmids (1)</p> <p>If neither answer above is given then award a maximum 1 mark only for:</p> <p>Circular (DNA)</p>	2	

Key Area 1.3: Gene Expression

Q1)

(i)	1. RNA polymerase unzips/ unwinds DNA or separates DNA into two strands	1	(ii)	a. tRNA has an anticodon and an amino acid attachment site	1
	2. Hydrogen bonds between strands/base pairs break	1		b. tRNA binds/joins to/carries/collects specific/correct amino acid	1
	3. RNA polymerase aligns/brings in/joins/attaches RNA nucleotides with their complementary nucleotides/bases on DNA (template)	1		c. tRNA carries (specific) amino acid to ribosomes	1
	OR			d. anticodons are complementary/pair with codons on mRNA	1
	A to U and T to A and C to G in diagram			e. there are start and stop codons	1
	4. a primary transcript is produced	1		f. peptide bonds form between amino acids	1
	5. exons are coding and introns are non-coding (regions of the primary transcript)	1		OR	
	6. introns/non-coding regions are removed OR exons/coding regions are retained	1		a polypeptide forms	(max 4)
	7. exons are spliced/joined together to form mature mRNA)/transcript	1			

Q2)

(a)	Amino acid	1	
(b)	Protein OR Enzymes	1	
(c)	Cut/cleave AND combine polypeptide chains OR add phosphate/carbohydrate	1	NOT – <i>post translational modification</i> NOT – <i>cleave/cut</i> alone
(d)	Name: <u>Alternative</u> (RNA) splicing (1) Description: Different (combinations of/variety of) <u>exons</u> are included/ spliced together (in the mature transcript/ RNA) (1)	2	NOT – a description suggesting the order of exons is changed NOT – depends what sections are treated as exons and introns

Q3) A

Q4)

(a)		Introns	1	NOT- non-coding regions.
(b)		1,3,4/2,3,4	1	Must be in correct order and inversions not acceptable.
(c)	(i)	Shorter protein/fewer amino acids.(1) <u>Stop</u> codon is produced earlier (in the sequence) (1)	2	NOT- non-functional protein. NOT- protein is short. NOT- stop codon is produced alone.
	(ii)	Every amino acid after the mutation is changed/affected.	1	NOT- frame shift mutation alone.

Q5)

B	(i)	<ol style="list-style-type: none"> 1. Single strand of nucleotides. 2. (nucleotide) contains ribose sugar, phosphate and base. 3. Adenine, cytosine, guanine and uracil. 4. mRNA takes copy of DNA code from nucleus to ribosome. 5. 3 bases on mRNA codes for an amino acid/is a codon. 6. tRNA picks up specific/one type of amino acid. 7. tRNA carries amino acid to a ribosome. 8. tRNA has anticodon (complementary to codon) AND an amino acid attachment site. 9. rRNA (combined) with protein forms a ribosome. 10. If points 4 -9 not awarded, award point for stating mRNA, tRNA and rRNA. <p>(any 6)</p>	9	<p>NOT - Letter of bases alone.</p> <p>NOT - "DNA has thymine and RNA has uracil" alone.</p>
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(ii)	<p>a. Introns removed from primary transcript.</p> <p>b. Exons joined/spliced together to produce mature transcript.</p> <p>c. Exons coding/expressed AND introns non coding/not expressed.</p> <p>d. Alternative (RNA)splicing produces different mature transcripts.</p> <p>e. (Different mature transcripts produced) depending on which (combinations of) exons are retained/spliced together/ removed.</p> <p style="text-align: right;">(any 3)</p>	
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Q6) C

Q7)

(a)	3:5:30	1	
(b)	Protein	1	Not - Amino acids
(c)	<p>Change in the sequence may change the structure/function/shape of the ribosome</p> <p>OR</p> <p>Ribosome is not made/formed/usable (1)</p> <p>Translation can no longer occur</p> <p>OR</p> <p>mRNA cannot join to ribosome</p> <p>OR</p> <p>tRNA cannot join to mRNA (1)</p>	2	
(d)	<p>DNA double strand, (r)RNA single strand</p> <p>OR</p> <p>DNA has thymine, (r)RNA has uracil</p> <p>OR</p> <p>DNA has deoxyribose sugar, (r)RNA has ribose</p>	1	Not - T and U.

Q8)

(a)	(i)	Prokaryotes have a smaller genome OR Eukaryotes have a larger genome	1	If figures quoted they must be correct to award the mark.
	(ii)	800	1	
	(iii)	4100	1	
(b)		$2.52/2.5 \times 10^7$ OR 25 200 000	1	
(c)		Regulates transcription OR Transcribed to tRNA/rRNA	1	Not - Transcribed but not translated alone
(d)		There are different combinations of exons in the mature transcript OR Different exons are removed from the primary transcript	1	Not - any reference to a change in sequence of exons: eg different order of exons Not - Depends on which exons are treated as introns

Key Area 1.4: Cellular Differentiation

Q1)

(a)	<p>Increase in stroke volume/volume of blood pumped out of heart per heartbeat (in patients given the treatment) 1</p> <p>No effect on heart rate (of patients given the treatment) 1</p>	2	
(b)	1190	1	
(c)	<p>(i) Embryonic stem cells differentiate/develop into all/many types of cell AND adult/tissue stem cells differentiate/develop into less/narrower range of/limited cell types OR Adult stem cells are more differentiated/specialised than embryonic stem cells</p>	1	
	<p>(ii) They express/switch on the genes characteristic of that type of cell OR Certain genes/some genes are expressed/switched on (and other genes are switched off)</p>	1	NOT - genes are switched on and off
(d)	<p>Provide information on gene regulation/cell growth/cell differentiation/cell division/cell ageing/disease development OR Use as model cells to study how diseases develop OR For drug testing</p>	1	

Q2)

(a)	<p>It differentiates into/specialises into/becomes...</p> <p>many/lots of/all/wide range of cell types/tissue types</p> <p>OR</p> <p>It is pluripotent/totipotent</p>	1	NOT – multipotent
(b)	<p>Different proteins will be produced/synthesised/made (resulting in different cell types)</p> <p>OR</p> <p>Only proteins characteristic of that cell type are produced/synthesised/made</p>	1	NB : Protein coded for ≠ synthesised
(c)	<p>Repair of damaged/diseased...organs/cells/tissues</p> <p>OR</p> <p>Production of tissues for grafting/transplant</p> <p>OR</p> <p>Correct examples eg bone marrow transplants/(make) skin grafts/to treat a named disease/treat burns</p>	1	<p>NOT – cure/treat diseases alone</p> <p>NOT – research diseases</p>
(d)	<p>Embryo/it/baby/foetus/a potential life... is... destroyed/killed/not allowed to develop</p> <p>OR</p> <p>Embryos which would have been destroyed are being put to good use</p> <p>OR</p> <p>Use of stem cells for drug testing rather than animals</p> <p>OR</p> <p>Diseases could be cured</p>	1	NOT – religious reasons alone

(a)	(i)	<p>From.... start/0 – 5 weeks/over first 5 weeks it increased from <u>0</u> – 9.2 (1)</p> <p>From 5 (– 7) weeks it remained constant/levelled off (1)</p> <p>Correct values for 2 statements but no units (weeks) = 1 mark</p>	2
	(ii)	200	1
	(iii)	B	1
(b)		<p>B It/number of shoots is highest/ greatest (at 7 weeks) (1) and this is (still) increasing (1)</p> <p>OR</p> <p>C It/number of shoots... is increasing more/most rapidly (1) and B is slowing down/levelling off (1)</p>	2
(c)		<p>Greatest (average) <u>root</u> length/ Longer <u>roots</u> (1)</p> <p>More water absorbed for photolysis/photosynthesis</p> <p>OR</p> <p>More nutrients absorbed for named process eg protein synthesis/ATP production etc. (1)</p>	2

(a)	(i)	Can only differentiate/specialise/change into a few types of cell/myoblasts/muscle cells/limited variety of cells/cells of the tissue that it came from (or converse). (1)	1	NOT- can only differentiate into a limited number of cells without reference to type. NOT- multipotent alone.
	(ii)	Growth/repair/renewal of <u>muscle</u> (tissue). OR Increase number of muscle cells for growth/repair (of muscles). OR Become muscle cells for growth/repair (of muscles).	1	NOT- repair of muscle cells. NOT- growth of muscle cells. NOT- increase number of muscle cells alone.
	(iii)	Does not involve destruction/killing of a (potential) life/embryo.	1	
(b)		Testing/development of drugs/medicines. OR Study how diseases develop (or description of development of a named disease). OR Study cell growth/cell division/ cell differentiation/gene regulation.	1	NOT- descriptions of therapeutic uses. NOT- researching diseases. NOT- study cell processes.

Q5) A

Q6) B

7) D

Key Area 1.5 and 1.6: Structure of the Genome and Mutations

Q1)

(i)	1. (single gene) mutations are random changes in DNA sequences/genes/alleles/the genome	1	(ii)	a. chromosome mutation can involve changes to chromosome number/structure	1
	2. single gene mutation name AND description substitution - base/base pair/ nucleotide is replaced/substituted by another insertion - base/base pair/ nucleotide is added/inserted deletion - base/base pair/ nucleotide is removed/deleted	1		b. chromosome mutation name AND description; Translocation: genes/sections of chromosome from one chromosome become attached to another chromosome Deletion: genes/sections of chromosome deleted from chromosome Inversion: genes/sections of chromosome/rotate through 180°/flipped Duplication: genes/sections of chromosome/pieces of chromosome are duplicated/repeated	1
	3. another single gene mutation name AND description	1		c. another chromosome mutation name AND description	1
	4. If 2 or 3 not awarded - all 3 mutation names	1		d. If b or c not awarded - all 4 names but no descriptions	1
	5. Insertion/deletion results in a frameshift mutation/expansion of a nucleic acid sequence	1			
	6. (single nucleotide) substitutions include missense, nonsense and splice site mutations	1			
	7. splice site mutations can alter the mature mRNA OR result in exon removal OR result in introns remaining present	1			
		(max 4)			

Q2) B

Q3)

(a)	(i)	Deletion/insertion	1	
	(ii)	<p>Effect on lactase gene: All the codons/base sequences nucleotide sequences/ triplets/ bases/nucleotides after the mutation will change/will move along.</p> <p>OR</p> <p>All the following codons/base sequences nucleotide sequences/triplets will change. (1)</p> <p>Effect on structure of lactase: All the amino acids after the mutation may change (1)</p>	2	NOT - Amino acids produced/made.
(b)	(i)	<p>Gene is permanently switched on.</p> <p>OR</p> <p>More transcription occurred.</p> <p>OR</p> <p>Repressor molecule not produced so operator permanently switches on gene/gene not switched off.</p>	1	

Q4) C

Q5)

(a)	(i)	Hydrogen	1	
	(ii)	Introns/non-coding sequences are removed from the primary transcript OR Gene/primary transcript has introns and exons, mature transcript has (only) exons. OR RNA splicing	1	
(b)	(i)	Section of a chromosome/gene(s) is added to its homologous partner OR A gene moves from a chromosome to its homologous partner	1	
	(ii)	Beneficial mutations can occur in one of the copies of the gene/DNA (1) The other gene can still be expressed to produce its protein (1)	2	Must be clear mutation is occurring in one copy of the gene

Q6)

(a)		To show the effect of the mutations OR To show the effect of the drugs/ each drug OR To prove the drugs affect/increase chloride transport OR To compare with and without drugs	1	
(b)		Treatment - P (1) Mutation - B (1)	2	
(c)		An average was calculated OR Results are averaged	1	
(d)		Personalised medicine OR Pharmacogenetics	1	

Key Area 1.7: Evolution

Q1) B

Q2)

(a)	(i)	Sympatric	1	
	(ii)	Prevents/interrupts/stops/blocks... gene flow/gene exchange/breeding/ mating... between populations OR Prevents interbreeding	1	NOT – stops populations from mating/ breeding alone
	(iii)	(DNA) sequence data/genome analysis would be similar OR They/the two populations... can interbreed /breed together... to produce fertile offspring (or converse statement)	1	NOT – they can breed to produce fertile offspring NOT – answers referring to 2 species instead of 2 populations

Q3) B

Q4)

(a)		From 2009/from the start it increases from 10% to 66% in 2015. OR Over the first 6 years it increases from 10% to 66%. (1) Then stays constant (until 2016). (1)	2	Cannot access any marks if 66 and 2015 not mentioned. Must state % unit at least once to gain full marks. All correct values but no % unit = 1. If additional points are correctly described do not negate.
(b)		Resistant plants survive. OR Non-resistant die. (1) Pass resistance- <ul style="list-style-type: none"> genes/alleles/sequences to next generation/to offspring/vertically. OR Reproduce/breed <ul style="list-style-type: none"> and pass on resistance genes/alleles/sequences. (1) 	2	NOT- pass on characteristic/ traits/resistance.
(c)		Bacteria can exchange- <ul style="list-style-type: none"> genetic material/plasmids horizontally/in same generation and charlock cannot/charlock transfers vertically. OR Horizontal gene transfer is faster (or converse).	1	NOT- horizontal gene transfer is fast.

Q5) D

Q6) A Q7) B Q8) A

Q9) B

Q10)

(a)		Geographical (1) Prevents gene flow between populations/groups OR Prevents populations/groups interbreeding (1)	2	Not - Prevents gene flow/exchange between species Not - Prevents gene flow alone
(b)	(i)	5	1	
	(ii)	Behavioural isolation/barrier OR Ecological isolation/barrier OR Sympatric speciation	1	

Key Area 1.8: Genomic Sequencing

Q1) C Q2) B

Q3)

(a)	(i)	550	1
	(ii)	260	1
	(iii)	<p>Last common ancestor (of rats and humans) was more recent (than rats and frogs).</p> <p>OR</p> <p>Last/most recent common ancestor of rats and humans was 90 million years ago while rats and frogs was 420 million years ago.</p> <p>OR</p> <p>Rats diverged more recently from humans than from frogs.</p>	1
(b)	(i)	28	1
	(ii)	Any value from 27 to 28	1
(c)		21	1

Q4) D

Q5)

DNA sequence data	1
OR	
Fossils/fossil records	
OR	
Mutation rate	

Q6) A

Q7) A