

## Mark schemes

- 1** (a) Reverse transcriptase; 1
- (b) 1. Probe (base sequence) complementary (to DNA of allele A / where A is (and) binds by forming base pairs / hydrogen bonds;  
*Accept gene A*
2. So (only) this DNA labelled / has green dye / gives out (green) light;  
*Accept glows for green light* 2
- (c) (i) 1. More probe binding / more cDNA / mRNA / more allele / gene A means more light;
2. DNA (with **A**) doubles each (PCR) cycle;
3. So light (approximately) doubles / curve steepens more and more (each cycle) / curve goes up exponentially / increases even faster; 3
- (ii) (**G** because)
1. (Heterozygous) only has half the amount of probe for **A** attaching / only half the amount of DNA / allele A (to bind to);  
*Accept only one A to bind to*
2. (So,) only produced (about) half the light / glow / intensity (of **H**) (per cycle of PCR);  
*If reference to 'half' for point 1, allow 'less light' in 2.* 2
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- 2** (a) 1. To allow comparison;
2. Because different number of cells in samples / different times for incubation / numbers become easier to manipulate; 2
- (b) 203.7(%);;  
*Allow 1 mark for 21.8 / 10.7*  
*Allow 1 mark for correct answer (203.74) but not correctly to 1 dp*  
*204 = 1 mark* 2
- (c) (i) 1. (At every concentration) uptake is faster at 37°C / at higher temperature;
2. Due to faster respiration / ATP production; 2

- (ii) 1. Uptake at 37°C only small increase / levelling off / almost constant as carrier proteins full;  
*Accept 'no (significant) change'*  
*Ignore use of numbers*
2. Concentration of imatinib is not the limiting factor;

2

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**3** (a) 250 000;

1

- (b) (i) Loss of 3 bases / triplet = 2 marks;;  
*'Stop codon / code formed' = 1 mark max unless related to the last amino acid*

Loss of base(s) = 1 mark;

*eg triplet for last amino acid is changed to a stop codon / code = 2 marks*

*3 bases / triplet forms an intron = 2 marks*

*Accept: descriptions for 'intron' eg non-coding DNA*

*'Loss of codon' = 2 marks*

2

- (ii) 1. Change in tertiary structure / active site;  
*Neutral: change in 3D shape / structure*
2. (So) faulty / non-functional protein / enzyme;  
*Accept: reference to examples of loss of function eg fewer E-S complexes formed*

2

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**4** (a) Cytosine with Guanine and (Adenine) with Uracil;  
*Ignore G, C and U*

1

- (b) Two reasons, with suitable amplification;;

**Q**

Only infected cells have HIV protein on surface;

So carrier only attaches to / specific to these cells / siRNA can only enter these cells;

**OR**

siRNA (base sequence) complementary / specific to one mRNA;

*Accept idea of specificity*

Only infected cells contain mRNA of HIV / this gene / stops translation of this gene / only binds to this mRNA / destroys this mRNA;

*Accept could not inhibit other / non-HIV mRNA*

4 max

- (c) 1. Carrier binds to (protein on) HIV;

*1. Accept references to HIV membrane*

2. Prevents HIV / it binding to (receptor on human) cell;

*2. Reject references to binding to HIV protein on human cell*

2

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- (a) 1. Carriers are heterozygous / have one normal copy and one mutant copy of gene / have one recessive allele / don't have the condition;

2. Both have DNA that binds (about) half / 50% amount of probe (that non-carrier does);

3. Probe binds to dominant / healthy allele so only one copy of exon in their DNA / have one copy of gene without exon / base sequence for probe to bind to;

*3. Accept normal and gene*

*3. Accept have a deletion mutation*

3

- (b) 1. Introns not translated / not in mRNA / (exons) code for amino acids / introns do not code for amino acids;  
     1. *Accept not expressed*  
     1. *Accept polypeptide / protein for amino acids*
2. Mutations of these (exons) affect amino acid sequences (that produce) faulty protein / change tertiary structure of protein;  
     2. *Accept deletion leads to frameshift*  
     2. *In this context, accept affects protein made*
3. So important to know if parents' exons affected, rather than any other part of DNA / introns;  
     *Accept converse arguments involving - eg introns do not code for amino acids / proteins*  
     *Reject references to making amino acids, once*

3

- (c) 1. Restriction mapping / described;
2. DNA / base sequencing (of fragments) / description / name of method;

2

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- (a) 1. No effect at 25°C  
     *The question only refers to plants with GB*  
     1. *Reject same mass*
2. Keeps growing at 30°C and 35°C / up to 35°C (more than without GB);
3. Above 35°C, falls but grows more than plant without GB;  
     3. *Accept at all temperatures above 25°C more growth than without GB*

2 max

- (b) (i) Significantly different / SEs do not overlap ;  
     *Accept converse without GB*

1

(ii) (As temperature increases,)

1. Enzyme activity reduced / (some) enzymes denatured;
2. Less photosynthesis, so fewer sugars formed;
3. Less respiration / less energy / ATP for growth;
4. Less energy for named function associated with growth  
     4. *Eg mitosis, uptake of mineral ions*

4

- (c) 1. (Rubisco activase attaches to thylakoid and) this changes shape / tertiary structure (of enzyme) / blocks active site / changes active site;

*Note - question states enzyme stops working when it attaches to thylakoid, not before*

*1. Accept rubisco in this context*

2. (This) prevents substrate / RuBP entering active site / binding;

*2. Accept prevents ES complex forming*

*2. Accept no longer complementary to substrate / RuBP*

2

- (d) 1. GB prevents / reduces binding of rubiscoactivase to (thylakoid membrane);

*1. Accept enzyme instead of rubiscoactivase. Accept rubisco*

2. (Prevents it) up to 35°C;

3. (So) rubiscoactivase / enzyme remains active;

4. (So) photosynthesis / light-independent stage still happens;

*4. Accept descriptions of light-independent stage*

5. Above 35°C, some binding still occurs but less than without GB, so less reduction in growth;

4 max

- (e) 1. Looked for information / journals, on crop plants that grow at high temperatures;

*1. "other research" is minimum accepted*

*1. Accept previous experiments research with temperature resistant crops*

*Ignore simple references to looking at previous studies / other plants - need to relate to this context*

2. (Crop plants cited in this research) contain / make GB;

3. So assumed making plants produce GB makes them resistant to high temperatures;

2 max

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- (a) 1. Cell wall not formed / production inhibited;

*1. Q Accept: weakened cell wall, but do not accept 'cell wall is broken down'*

2. Lower water potential in bacterium;

*2. Accept: converse*

*2. Must be clear that the lower water potential is in the bacterium*

3. Water enters and causes lysis / expansion / pressure;

2 max

- (b) Human cells lack enzyme (**B**) / have a different enzyme / produce different fatty acids / use different substrates;

*Neutral: 'human cells do not have cell walls' as out of context*

1

- (c) 1. Change in base sequence (of DNA / gene) leading to change in amino acid sequence / primary structure (of enzyme);  
*1. Accept: different amino acids coded for*  
*1. Reject: different amino acids produced*
2. Change in hydrogen / ionic / disulphide bonds leading to change in the tertiary structure / active site (of enzyme);  
*2. Neutral: alters 3D structure / 3D shape*
3. Substrate not complementary / cannot bind (to enzyme / active site) / no enzyme-substrate complexes form;

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- (a) Restriction / endonuclease;

*Ignore specific names of restriction enzymes e.g. EcoR1*

1

- (b) (i) 1. (Acts as a) marker gene to show that the (human) gene has been taken up / expressed;

*1. Accept: gene marker*

2. (Only) implant cells / embryos that show fluorescence / contain the jellyfish gene;

2

- (ii) 1. Factor IX present in / extracted from milk;

2. Gene only expressed in mammary glands / udder / gene not expressed elsewhere;

*2. Ignore references to milk*

*The 'only' aspect is important here.*

3. Do not need to kill sheep (to obtain Factor IX);

2 max

- (c) (i) 1. Mutation / nucleus / chromosomes / DNA may be damaged / disrupts genes;  
 1. *Neutral: cell may be damaged*
2. May interfere with proteins (produced) / gene expression / translation;  
*Ignore references to hormone levels or time of implantation*

**OR**

3. Embryo / antigens foreign;  
 3. *Neutral: antigens change*
4. Embryo is rejected / attacked by immune system;  
 4. *sNeed idea that the immune system is involved if mark point 3 has not been given*  
*'Embryo foreign so rejected' = 2 marks*  
*'Embryo rejected by immune system' = 1 mark*  
*'Embryo is rejected' = 0 marks*

2 max

- (ii) 1. Saves time / money for others;
2. Same work is not repeated / methods can be compared / improved / amended / same errors are not made;

2

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- (a) 1. Adenylate cyclase activated / cAMP produced / second messenger produced;
2. Activates enzyme(s) (in cell so) glycogenolysis / gluconeogenesis occurs / glycogenesis inhibited;  
 2. *Neutral: 'glucose produced' as given in the question stem*  
*Accept: correct descriptions of these terms*

2

- (b) (i) 1. Glucose / sugar in food would affect the results;  
 1. *Accept references to starch / carbohydrate*  
*Or*
2. Food / eating would affect blood glucose (level);  
*Or*
3. (Allows time for) blood glucose (level) to return to normal;  
 3. *Neutral: allows time for insulin to act*

1 max

- (ii) Type 2 diabetes is a failure to respond to insulin / still produces insulin / is not insulin-dependent;

1

(iii) (For) – 3 max

*A maximum of three marks can be awarded for each side of the argument*

1. Avoids injections / pain of injections;
2. Long(er) lasting / permanent / (new) cells will contain / express gene;  
*Ignore references to methodology e.g. sample size not known*
3. Less need to measure blood sugar / avoids the highs and lows in blood sugar;
4. Less restriction on diet;

(Against) – 3 max

5. Rats are different to humans;
6. May have side effects on humans;  
*6. Accept: virus may be harmful / disrupt genes / cause cancer*
7. Long(er) term effects (of treatment) not known / may have caused effects after 8 months;
8. (Substitute) insulin may be rejected by the body;

4 max

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(a) (i)

1. Negative correlation;  
*Accept: description for 'negative correlation'*  
*Neutral: 'correlation'*  
*Reject: positive correlation*
2. Wide range;
3. Overlap;
4. (Graph suggests that) other factors may be involved (in age of onset);  
*2 / 3 Accept the use of figures from the graph*  
*2 / 3 Can refer to age of onset or number of CAG repeats*  
*Ignore references to methodology*

3 max

- (ii) 1. Age of onset can be high / symptoms appear later in life;  
*Accept: 'gene' for 'allele'*
2. (So) individuals have already had children / allele has been passed on;

**OR**

3. Individuals have passed on the allele / already had children;
4. Before symptoms occur;

2 max

- (b) (i) 1. Person **K**;
2. (As has) high(est) band / band that travelled a short(est) distance / (er) so  
has large(st) fragment / number of CAG repeats;  
*Must correctly link distance moved and fragment size*

2

- (ii) Run fragments of known length / CAG repeats (at the same time);  
*Accept: references to a DNA ladder / DNA markers*  
*Do not accept DNA sequencing*

1

- (iii) Homozygous / (CAG) fragments are the same length / size / mass;  
*Accept: small fragment has run off gel / travelled further*

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**[9]**