

8.4 The control of gene expression (A-Level Only) - Gene technologies 1 – Mark schemes

Q1.

- (a) Produces (c)DNA using (m)RNA;
Accept: 'converts' (m)RNA to (c)DNA.
Reject: tRNA 1
- (b) Joins nucleotides to produce (complementary strand/s of) DNA;
Accept: 'joins DNA nucleotides'. 1
- (c) 1. To remove any DNA present;
2. As this DNA would be amplified / replicated;
1. *Must be idea of removal / destruction.*
2. *Accept: idea of DNA not being used as template.* 2
- (d) 1. Ratio in range of 1.4 :1 to 1.5 :1 = **2 marks**;
2. One mark for answers which shows incorrect ratio but Shows 0.24 as a number or line on the graph
OR
Ratio in correct range, but the wrong way round
OR
Ratio in correct range but not expressed to 1
OR
Ratio shown the other way round in range
1: 0.67 to 1:0.71;
*Note: ratio not expressed to 1 in correct range may be shown in different ways, for example as:
3:2 or simply as 1.5 for one mark.* 2
- (e) Limited number of primers / nucleotides;
Accept: DNA polymerase (eventually) denatures
Accept: primers / nucleotides 'used up'. 1
- (f) 1. Base sequences differ;
2. (Different) complementary primers required;
1. *Accept: reference to either RNA or DNA base sequences but reject reference to DNA base sequence in viruses.* 2

[9]

Q2.

- (a) 1. Human DNA / human gene / HGH gene contains introns
OR
Methods 2 and 3 produce DNA / HGH without introns;
2. *E. coli* cannot remove introns / cannot splice mRNA / cannot splice pre-mRNA; 2

- (b) Faster to use gene machine than all the enzyme-catalysed reactions (involving reverse transcriptase);
Accept extra step / more steps involved in isolating mRNA 1
- (c) 1. Cut the plasmid with a restriction endonuclease;
Allow 'add base sequences to blunt ends of plasmid and HGH gene'
2. (So that) both have complementary / sticky ends;
3. (Mix together) and add ligase to join the complementary / sticky ends; 3
- (d) Can quickly identify transformed bacteria using UV light; 1
- (e) 1. Arabinose alters structure of araC protein / reduces effect of araC protein;
2. So stops / reduces inhibition of promoter gene and GFP gene is transcribed;
OR
 So stops / reduces inhibition of promoter gene and GFP is produced; 2
- [9]

Q3.

- (a) 1. (If injected into egg), gene gets into all / most of cells of silkworm;
2. So gets into cells that make silk. 2
- (b) 1. Not all eggs will successfully take up the plasmid;
2. Silkworms that have taken up gene will glow. 2
- (c) Promoter (region / gene). 1
- (d) 1. So that protein can be harvested;
2. Fibres in other cells might cause harm. 2
- [7]

Q4.

- (a) (i) Restriction endonuclease; 1
- (ii) (DNA) ligase; 1
- (b) (For those plants that contained the desired gene in the nucleus/plant DNA)
1. (DNA of desired gene) copied/replicated with host DNA/inside nucleus;
2. Passed on by mitosis/plant grows by mitosis;
3. Produces genetically identical cells/clones;
- Ignore references to protein synthesis or plasmids not taking up the gene*

1. *Accept DNA replication during mitosis*
1. and 2. *Accept converse for plants with the gene in the cytoplasm*
3. *Neutral 'identical unqualified'*
3. *Accept description, e.g., DNA is the same*

3

- (c) 1. Genetic code is universal/triplets in DNA always code for same amino acid;
2. It/insect DNA can be transcribed;
3. Can be translated (process/mechanism same in all organisms/cells);
2. *Accept (basic) transcription (process/mechanism) same in all organisms/cells;*
 2. *Accept descriptions of process*
 3. *Accept descriptions of process*

3

[8]

Q5.

- (a) (i) Does not code for amino acid/tRNA/rRNA;
- Accept 'does not code for production of protein/polypeptide'*
- Reject 'that produces/makes amino acid'*

1

- (ii) Deletion mutation;
- Accept 'deletion'*
- Ignore references to splicing*

1

- (b) (The) polymerase chain reaction;
- Accept PCR*

1

- (c) 1. Probes are single stranded / have a specific base sequence;
2. Complementary base sequence on (specific) spacer

OR

3. Complementary/specific to (particular) spacer;
4. (In white squares probe) binds (to single-stranded spacer) and glows/produces light/fluoresce;
2. *Need idea of complementary to spacer*
3. *Accept converse for dark squares*

3

- (d) 1. To see if strain is resistant to any antibiotics;
2. So can prescribe effective/right antibiotic;

OR

3. To see whether (any) vaccine works against this strain/ see which vaccine to use/ to produce specific vaccine;
4. (So) can vaccinate potential contacts/to stop spread;

OR

5. Can test other people to see if they have the same strain/ to trace where people caught TB;
6. Allowing control of spread of disease/vaccinate/treat contacts (of people with same strain) before they get TB;

Do not allow mix and match of points from different alternative pairs

2 max

[8]

Q6.

- (a) 1. Cut (DNA) at same (base) sequence / (recognition) sequence;
Accept: cut DNA at same place
2. (So) get (fragments with gene) **R** / required gene.
Accept: 'allele' for 'gene' / same gene 2
- (b) 1. Each has / they have a specific base sequence;
2. That is complementary (to allele r or R).
Accept description of 'complementary' 2
- (c) 1. Fragments L from parent rr, because all longer fragments / 195 base pair fragments;
Ignore: references to fragments that move further / less, require identification of longer / shorter or 195 / 135
Accept: (homozygous) recessive
2. Fragments N from parent RR, because all shorter fragments / 135 base pair fragments;
1 and 2 Accept: A3 for 195 and A4 for 135
2. Accept: (homozygous) dominant
3. (M from) offspring heterozygous / Rr / have both 195 and 135 base pair fragments.
Accept: have both bands / strips
Reject: primer longer / shorter 3
- (d) 1. (Cells in mitosis) chromosomes visible;
2. (So) can see which chromosome DNA probe attached to. 2
- (e) (i) 1. For comparison with resistant flies / other (two) experiments / groups;
Ignore: compare results / data / no other factors
2. To see death rate (in non-resistant) / to see effect of insecticide in non-resistant / normal flies.
Accept: 'pesticide' as 'insecticide'
Accept to see that insecticide worked / to see effect of enzyme 2
- (ii) (PM must be involved because)
1. Few resistant flies die (without inhibitor);

2. More inhibited flies die than resistant flies;
3. (PM) inhibited flies die faster (than resistant flies);
(Other factors must be involved because)
4. Some resistant flies die;
5. But (with inhibitor) still have greater resistance / die slower than non-resistant flies.

Accept: (with inhibitor) die slower than non-resistant flies

4 max

[15]

Q7.

(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

[8]

Q8.

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

3

2

[8]

Q9.

- (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
- [15]**

Q10.

- (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

[9]

Q11.

- (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

[8]

Q12.

- (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
- (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;

3 max

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 1
- [9]**

Q13.

- (a) restriction (enzyme) / endonuclease / named example; 1
- (b) unpaired bases / sticky ends / staggered;
complementary / explained; 2
- (c) *1 mark for each correct outcome*
plasmid with foreign DNA joined in ring;
ring with plasmid only; ring of foreign DNA only;
ignore linear structures 3
- [6]**

Q14.

- (a) a length of DNA;
that codes for a single protein / polypeptide; 2
- (b) by heating;
to break the H-bonds (between complementary bases); 2
- (c) (i) to allow the DNA polymerase to attach / start addition of
nucleotides / mark start and end of sequence to be
copied / prevents strands re-joining; 1
- (ii) because the sequences at the ends of the target sequence
are different / one is at the beginning and one at the end; 1
- (d) 8;
accept 7 1
- [7]**

Q15.

- (a) 1 DNA is cut;

- 2 Using restriction enzyme;
- 3 Use electrophoresis;
- 4 Separates according to length / mass;
- 5 Southern blotting / transfer to (nylon) membrane;
- 6 Make single-stranded;
- 7 Apply probe;
- 8 Radioactive / fluorescent;
- 9 Reference to tandem repeats / VNTRs / minisatellites;
- 10 Autoradiography / eq;
8 and 10 should be consistent

max 6

- (b) (i) All bands in cub which don't come from mother;
Must be in father's DNA fingerprint;
Principle that all bands in cub must come from mother and father = 1

2

- (ii) Select pairs with dissimilar DNA fingerprints;

1

- (c) (i) Cells (from panda) in faeces / gut cells / blood cells;

1

- (ii) To increase amount of DNA / only small amount present;

1

- (iii) DNA / primer has specific base-sequence;
Reference to specific / complementary base-pairing;

2

- (d) Taking samples from animals causes stress / injury to animal;

Difficult to find animals;

Pandas are dangerous / threat to human;

max 2

[15]

Q16.

- (a) Restriction (enzyme / endonuclease);

1

- (b) Move towards anode / move because charged;

Different rates of movement related to charge / size;

2

- (c) (i) Piece of DNA;
Single stranded;
Complementary to / binds to known base sequence / gene;

max 2

- (ii) DNA invisible on gel / membrane;
Allows detection;

2

[7]

Q17.

- (a) 1. DNA is cut;
2. using restriction enzyme;
3. electrophoresis;
4. separates according to length / mass / size;
5. DNA made single-stranded;
6. transfer to membrane / Southern blotting;
7. apply probe;
8. radioactive / single stranded / detected on film / fluorescent;
9. reference to tandem repeats / VNTRs / minisatellites;
10. pattern unique to every individual;

6 max

- (b) cells on toothbrush;
DNA present in cell;

2

- (c) (i) toothbrush gives small sample of DNA / need more DNA
for analysis;
PCR gives many copies;

2

- (ii) uses heat;
to separate strands;
OR
PCR replicates pieces of DNA;
because DNA has been cut;
OR
primer added in PCR;
to initiate replication

2 max

- (d) (i) PCR / amplification needed;

1

- (ii) other DNA present; need to identify 'required' DNA from rest;

2

[15]

Q18.

- (a) (i) protein / immunoglobulin;
specific to antigen;
idea of 'fit' / complementary shape;

2 max

- (ii) 1. virus contains antigen;
2. virus engulfed by phagocyte / macrophage;
3. presents antigen to B-cell;
4. memory cells / B-cell becomes activated;
5. (divides to) form clones;
6. by mitosis;
7. plasma cells produce antibodies;
8. antibodies specific to antigen;

9. correct reference to T-cells / cytokines;

6 max

- (b) 1. antibody gene located using gene probe;
2. cut using restriction enzyme;
3. at specific base pairs;
4. leaving sticky ends / unpaired bases;
5. cut maize / DNA / vector using same restriction enzyme;
6. join using DNA ligase;
7. introduce vector into maize / crop / recombinant DNA into maize;

4 max

(c) passive / person is not making own antibodies / antibodies not replaced;
memory cells not produced;

2

(d) fewer ethical difficulties / less risk of infection;

1

[15]

Q19.

(a) (i) Reverse transcriptase;

1

(ii) Idea that mRNA is present in large amounts in cell making
the protein / mRNA has been edited / does not contain
introns / mRNA codes for single protein;

1

(b) (Ligase) splices / joins two pieces of DNA / "sticky ends";

1

[3]

Q20.

(a) Endonuclease / restriction enzyme;

1

(b) DNA made of base pairs;
Each base pair is same length / occupies same distance
along backbone;

2

(c) (i) Second blank box from left labelled 6;

1

(ii) Distance moved depends on length / number of base pairs /
second longest fragment / second shortest distance identified;

1

(d) 5;

1

[6]

Q21.

(a) Mother and father both heterozygotes / Tt / carriers;
Probability of thalassaemia 1/4 and female 1/2;
Probability of both 1/8;

3

- (b) (i) Cut at same base sequence as same enzyme used;
Fragments are same length / size / have same charge; 2
- (ii) Single base occurs many times;
Sequence of 20 unlikely to occur elsewhere;
Allow one mark for establishing the principle where neither marking point clearly made. 2

[7]

Q22.

- (a) (i) Sticky ends / description;
Reference to complementary base-pairing 2
- (ii) Ligase; 1
- (b) Carrier of DNA / gene; (*context of foreign DNA*)
Into cell / other organism / host; 2
- (c) Act as marker gene;
Allows detection of cells containing plasmid / DNA; 2

[7]

Q23.

- (a) (i) Different genes / characteristics / features;
Reference to mutations;
Or
Base sequence determines protein;
Different species have different protein sequences; max 2
- (ii) Primer has different DNA sequence;
DNA specific / complementary base-pairing; 2
- (iii) Electrophoresis separates DNA;
(So they can be) identified by position on gel;
Smaller / shortest fragments travel furthest / quicker / or
reverse argument; 3
- (b) (*conventional*) Many lengths / all DNA / (*new*) one length;
Each rung is DNA of one / specific length; 2
- (c) 1 Heat DNA;
2 Breaks hydrogen bonds / separates strands;
3 Add primers;
4 Add nucleotides;
5 Cool;
6 (to allow) binding of nucleotides / primers;
7 DNA polymerase;
8 Role of (DNA) polymerase;
9 Repeat cycle many times;

Q24.

- (a) (i) Sticky ends / description;
Reference to complementary base-pairing 2
- (ii) Ligase; 1
- (b) Carrier of DNA / gene; (*context of foreign DNA*)
Into cell / other organism / host; 2
- (c) Act as marker gene;
Allows detection of cells containing plasmid / DNA; 2

[7]

Q25.

- (a) 1 (DNA altered by) mutation;
2 (mutation) changes base sequence;
3 of gene controlling cell growth / oncogene / that monitors cell division;
4 of tumour suppressor gene;
5 change protein structure / non-functional protein / protein not formed;
6 (tumour suppressor genes) produce proteins that inhibit cell division;
7 mitosis;
8 uncontrolled / rapid / abnormal (cell division);
9 malignant tumour; max 6
- (b) cancer cells die / break open;
releasing DNA; 2
- (c) normal DNA and changed DNA have different sequences;
DNA only binds to complementary sequence; 2
- (d) fewer abnormal / cancerous cells / smaller tumours;
less cell damage / less spread / fewer locations to treat; 2
- (e) mRNA base sequence has changed;
gene / DNA structure is different / has mutated;
cancer gene active / tumour suppressor gene inactive; 3

[15]

Q26.

- (a) (i) contains genes / nucleotides / sections of DNA / artificial
DNA from two species / 2 types of organisms; 1
- (ii) carries gene / DNA (into the other organism / gene carrier); 1

- (iii) expose cells to the fungus;
 non-resistant ones die, resistant ones survive;
 OR identify by adding marker gene / gene probe / (qualified)
 marker probe; description of positive result
 e.g. radioactivity / fluorescence / complementary base pairing;

2

- (b) EITHER 1 cut desired gene (from DNA) of oat plant;
 2 using restriction endonuclease / restriction enzyme;
 OR 1 use mRNA from oat which will code for resistance;
 2 and use reverse transcriptase to form desired DNA;
 OR 1 make artificial DNA with correct sequence of bases;
 2 using DNA polymerase;
 3 cut plasmid open;
 4 with (same) restriction endonuclease / restriction enzyme;
 5 ref. sticky ends / unpaired bases attached;
 6 use (DNA) ligase to join / ref. ligation;
 7 return plasmid to (bacterial) cells;
 8 use of Ca^{2+} / calcium salts / electric shock;
 (if ref. to 'insulin' allow 5 max.)

max 6

[10]

Q27.

- (a) only small amounts obtained / PCR increases the amount / mass of DNA;
 so enough DNA available for genetic fingerprinting;
- (b) (i) to separate the two strands of the DNA /
 to break the hydrogen bonds;
(Reject "unzip")
- (ii) short lengths / fragments of DNA / nucleotides /
 single stranded DNA;
- (iii) to mark beginning and / or ends of the part of DNA needed /
 for attachment of enzymes or nucleotides / initiator /
 keeps strands apart;
- (iv) would not be denatured;
 must be heated to 95 °C / must withstand high temps;
- (c) 1 DNA extracted from sample;
 2 DNA cut / hydrolysed into segments using restriction endonucleases;
 3 must leave minisatellites / required core sequences intact;
 4 DNA fragments separated using electrophoresis;
 5 detail of process e.g. mixture put into wells on gel and electric
 current passed through;
 6 immerse gel in alkaline solution / two strands of DNA separated;
 7 Southern blotting / cover with nylon / absorbent paper (to absorb DNA);
 8 DNA fixed to nylon / membrane using uv light
 9 radioactive marker / probe added (which is picked up by required
 fragments) / complementary to minisatellites;
 10 (areas with probe) identified using X-ray film / autoradiography;

2

1

1

1

2

max 6

- (d) adult 3;
this is only one which, (with number 1), can provide (all) the DNA
fragments which children have / all bars match;
(Reject 'genes')