

## **A-Level Biology**

### **Gene Mutations**

**Mark Scheme** 

Time available: 56 minutes Marks available: 41 marks

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#### Mark schemes

1.

2.

(a) Box 2.

An inversion will result in a change in the number of DNA bases. Reject if more than one box with tick. Ignore crossed-out ticks

- (b) 1. (Increased) methylation (of tumour suppressor genes); Accept abnormal methylation or hypermethylation Ignore decreased acetylation of histones
  - 2. Mutation (in tumour suppressor genes);
  - Tumour suppressor genes are not transcribed/expressed OR
     Amino acid sequence/primary structure altered;
     Accept mRNA for transcription/transcribed
     Accept tertiary structure altered
     Accept different amino acid
     Ignore reference to protein not being formed
  - 4. (Results in) rapid/uncontrollable cell division; Accept cell division cannot be regulated Ignore growth
- (c) 1. Correct answer of  $1.9/1.93 \times 10^{25} = 2$  marks;; Accept  $2 \times 10^{25} = 2$  marks Ignore any numbers after 1.93
  - Incorrect answer but shows 84 = 1 mark
    OR
    28 × 3 = 1 mark
    OR
    Incorrect answer but shows 672 divided by 8 = 1 mark;
- (a) 1. Change in (sequence of) amino acid(s)/primary structure; *Reject amino acids are formed. Reject amino acids code.*
  - 2. Change in hydrogen/ionic/disulfide bonds;
  - 3. Alters <u>tertiary/3<sup>0</sup></u> structure; *Reject active site. Ignore quaternary. Ignore 3D.*

2

[6]

1

3 max

(b) 1. Produce healthy (red blood) cells **OR** 

Produce (normal) polypeptide/haemoglobin;

Produce only healthy (red blood) cells is only equivalent to mark point 1. Accept produce 'normal'/non-SCD cells.

Ignore type of stem cell e.g. pluripotent.

- No sickle/faulty/SCD (red blood) cells (produced)
   OR
   No defective polypeptide/haemoglobin;
- Stem/marrow cells (continuously) divide/replicate
   OR

Less chance of rejection (from brother/sister); Differentiate is not equivalent to divide/replicate. Ignore type of stem cell e.g. pluripotent.

(c)

Max 2 marks for marking points 1, 2 and 3

(For gene therapy)

1. No destruction of bone marrow **OR** 

No destruction of stem cells;

Accept no destruction of faulty bone marrow unless context indicates this is against gene therapy.

2. Donors are not required;

Stating 'only own cells used' is not equivalent.

3. Less/no chance of rejection (own stem cells);

(Against gene therapy)

- 4. Sickle/faulty (red blood) cells still produced
- Immune response against genetically modified cells/virus OR Long-term effect not known (as is new treatment) OR Virus could cause side effects;

Accept 'virus could cause problems' or 'risk(s) with virus'.

3 max

3

[9]

3.

(i) 4;

(a)

1

		(ii)	1. Char <i>1. R</i> e	nge in amino acid / (sequence of) amino acids / primary structure; eject = different amino acids are 'formed'			
			2. Char activ	nge in hydrogen / ionic / disulphide bonds alters tertiary structure / re site (of enzyme);			
			2. Al poin	Iters 3D structure on its own is not enough for this marking t.			
			3. Subs	strate not complementary / cannot bind (to enzyme / active site) / no vme- substrate complexes form;			
					3		
	(b)	1. Lack of skin pigment / pale / light skin / albino;					
		2.	Lack of co	ack of coordination / muscles action affected;			
				2 max			
	(c)	Four	Founder effect / colonies split off / migration / interbreeding;				
		Allow description of interbreeding e.g. reproduction between					
			indiv	viduals from different populations			
					1	r1	
4.							
	(a)	Subs	Substitution;				
		Accept inversion or translocation					
			Igno	re 'point mutation'			
					1		
	(b)						
		<i>Max 2 marks for mark points 2, 3 and 4</i>					
		1.	(VO <sub>2 max</sub> <b>a</b>	nd CS activity) increased for both groups;			
		2.	No statistical test, <b>so</b> do not know if <b>differences</b> are significant <b>OR</b>				
			No statistical test, <b>so differences</b> could be due to chance;				
			Igno	re standard deviation			
			Acce	ept correct named statistical test eg t-test			
		3.	Only 8 weeks training				
				id not loot long			
			maining di	iu not last iong,			
		4.	Might not I	be true for all types of training/exercise/females;			
					3 max		

#### (c) In Group C:

1. Less mitochondrial replication/production;

1. and 2. Accept converse for Group T

 Less transcription (of genes) for mitochondrial proteins/CS OR Less translation of (mRNA into) mitochondrial proteins;

Accept less CS/enzyme is produced

#### (d)

5.

#### Max 2 marks for mark points 3, 4 and 5

Ignore any answers relating to sample size or duration of investigation Ignore 'correlation does not mean causation' unless qualified

For (no mark)

- 1. (From Figure 2 Group T have) increased CS activity for Krebs cycle;
- (from Figure 1 Group T have) increased VO<sub>2 max</sub> so more oxygen (available)
   OR
   (from Figure 1 Group T have) increased VO<sub>2 max</sub> so more aerobic respiration

(non **Figure** 1 Group T have) increased VO<sub>2 max</sub> so more aerobic respiration OR (from Figure 1 Group T have) increased VO

(from Figure 1 Group T have) increased  $VO_{2 max}$  so delayed anaerobic respiration;

Accept 'less lactate' for delayed anaerobic respiration

#### Against (no mark)

 No correlation between (percentage change in) VO<sub>2 max</sub> and CS activity OR No correlation on Figure 3;

# It might not be thymine causing the increase OR There may be other differences in the control region (of Group T) that cause the increase;

Ignore 'could be due to lifestyle/diet changes'

 VO<sub>2 max</sub>/CS activity not the only measures of ability to exercise for longer; Accept ideas that they did not measure ability to exercise for longer

3 max

2

[9]

(a) 1. Replacement of a base by a different base (in DNA);

1

- (b) 1. (Depends on) size / mass (of protein);
  - 2. (Depends on) charge (of protein); Accept for 2 marks 'Smaller / more highly charged move further'
- (C) 1. Each protein has a different tertiary structure; 2. (Each) antibody has a specific antigen / binding / variable region / site; 3. So, (each antibody) forms different antigen-antibody complex OR (each antibody) only binds to complementary (protein); 3 (d) 1. Less NL3; 2. More NR2A and NR2B; 2 (e) 1. Higher ratio NR2B to NR2A with mutation; Accept 'more' as equivalent to 'ratio' 2. (Perhaps) better memory in mice with mutation; 2

2 max

[10]