

A-Level Biology

Genetic Diversity and Natural Selection

Mark Scheme

Time available: 62 minutes Marks available: 54 marks

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Mark schemes

1.	(a)	1.	Different primary structure/amino acid sequence;		
		2.	Different tertiary structure/shape of active site;		
		3.	Enzyme-substrate complexes more likely (with enzyme from AD ^F allele); Accept converse for AD ^S Accept is more complementary	3	
	(b)	Avoids bias			
		OR			
		Results (likely to be) reliable/repeatable;			
	(c)	1.	Flies with AD ^F /allele have selective advantage (in presence of alcohol); Accept converse for AD ^S Accept description of selective advantage		
		2.	So insects (with AD ^F more likely to) reproduce;		
		3.	Pass on ADF (allele/gene);		
		4.	(So) <u>allele</u> frequency increases;	4	
	(d)	Ansv	wer = Directional selection	1	[9]

2.

(a)

Histogram

- 1. Linear scale for *y* axis;
- 2. Linear scale for *x* axis;
- 3. Correct bar widths **and** touching;
- 4. All bar heights plotted accurately;

OR

Bar chart accept for 3 marks,

- 5. Linear scale for *y* axis;
- 6. Labelled bars of equal width **and** not touching;
- 7. All bar heights plotted accurately;

OR

Graph accept for 2 marks,

- 8. Linear scale for *y* axis;
- All co-ordinates plotted accurately for frequency density; *Reject answers where data for frequency density and birth mass not used*
- (b) Correct answer for **2 marks** = 20 000;;

Accept for 1 mark, rearranged equation (eg number of babies = frequency density × range of mass)

- (c) 1. Survival increases as the birth mass increases;
 - 2. Survival decreases with smoking;
 - 3. Effect of smoking (on number) similar at all birth masses;

4

2

3

(a) Type of selection

1. Directional;

Reason:

3.

2. One extreme selected/removed/favoured/chosen **OR**

One extreme allowed to breed;

Ignore references to adaptations/natural selection Accept large fish/small fish for 'extreme'

- (b) 1. As a baseline/control;
 - 2. To show effect of no selection

 OR
 To show what happens in a normal population/naturally
 OR
 To show effect of/compare with tank A/tank C;
 Ignore reference to type of selection
 Accept not removing/not catching/not fishing for 'selection'
 Accept genetic drift for 'no selection'
 Accept no fishing/no selection/no caught fish for 'normal population'
 Accept to compare with other results
- (c) Correct answer for 2 marks

(How much greater) 1.6 to 1.7;;

Accept for 1 mark, 1.2 : 1 **and** 2 : 1

Accept for 1 mark, 4.1 : 3.4 and 4.8 : 2.4 Accept $\frac{5}{3}$ for 2 marks

2

2

2

(d) Not supported because

- 1. (Sea) fishing reduces (mean) mass of fish;
- Because large fish removed
 OR
 Because small fish escape/put back
 OR
 Because fishing (model) like Tank C;

But

- Information from (only) one species
 OR
 Sea fishing catches other/different (types of) species;
- 4. No statistical test;
- 5. Size of tank may affect fish growth;
- 6. Fish in tanks are all same age/sea fish not all the same age;

No measure of number of fish (removed)/ only measured mean mass
OR
No measure of (total) yield of fish
OR
No measure of reproductive success of fish;

8. Removal of 90% of population is unlikely to be replicated in the sea fishing;

Sea fish do not have life cycle of one year OR Sea fish do not reproduce all at the same time; 2 max for "But"

3 max

- 4.
- (a) Locus;

Accept: loci

- (b) Differences in DNA / differences in base sequence of DNA; Accept: number of different alleles / size/variation in gene pool Reject: genes
- (c) 1. Jack Russell (genetic) diversity is (significantly) greatest;
 2. Bull terrier (genetic) diversity is (significantly) smallest / is most inbred;
 - 3. Miniature terrier and Airedale terriers are similar;

1-3: do not credit just a list of values

4. Standard deviations do not overlap / do overlap with correct ref to significance;

Reference to significance must be relevant to examples given

- (d) 1. (Bull terrier) breeding has included a genetic bottleneck/ small population/more inbreeding/ greater selection (pressure);
 Accept: founder effect
 - 2. Reduced number of different alleles/size of gene pool; Reject: decrease in number of genes Ignore ref to mutations
 - OR
 - Miniature (terrier) breeding has included more outbreeding/less selection (pressure);
 - 4. Increased number of different alleles/larger gene pool/more variety of alleles;

Reject if genes used instead of alleles Reject: lower frequency of alleles Ignore ref to mutations

[7]

2

1

1

Max 3

(a) 2 of the following pairs:

Mark for explanation must be paired with correct change in structure

- 1. Larger leaves;
- 2. Photosynthesis;

OR

5.

6.

Accept converse descriptions of leaves, root and stem: longer root, taller stem, smaller leaves

- 3. Larger / bigger / thicker root;
- 4. Storage;

OR

- 5. Stem shorter / absent; Accept converse correct explanation
- 6. Less energy used in stem growth / more energy for producing sugar;

(b) Beet ready quicker / less time required / allows land to be used again / harvested earlier; Allow more crops / many harvests. Ignore references to yield / profit 1 (C) 1. (Diversity) reduced / fewer different alleles / less variation / smaller gene pool; 2. As <u>alleles</u> have been chosen / rejected; 2 [7] Difference in DNA / base sequence / difference in alleles / genes / gene pool; (a) Neutral: 'fewer alleles' unless qualified e.g. fewer different alleles. 1 (b) Environmental; Accept: Environment 1 (C) Reduced (genetic diversity); As fewer different / varied alleles / genes / reduced gene pool; 2 [4]

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4 max



7.

There are several alternatives for marking points 2 and 3

2 max