

Name:

Class:

Taxonomy QP

Author:

Date:

Time: 53

Marks: 46

Comments:

IVIT.(a	a)	1.	Kingdom, Phylum, Class, Order, Family;		
		2.	Luscinia svecica; 1 mark for each correct column Allow Genus and Species if both placed in box for species but not if both placed in genus box	2	
	(b)) Number of different alleles of each gene; Accept number of different base sequences (found) in each gene		1	
	(c)	1.	Has greater proportion of genes / percentage of genes showing diversity;		
		2.	Percentage is 35% compared with 28% / proportion is 0.35 compared with 0.28; Allow correct figures that are not rounded up, i.e., 34.9% / 0.349 and 27.8% / 0.278	2	[5]
M2.		(a)	Kingdom, phylum and class;; Lose 1 mark for each error (i.e. omission or incorrect response). Sequence not essential.	2	
	(b)	(i)	Shows evolutionary relationship;	1	
		(ii)	26;	1	
	(c)	(i)	 Base sequence will be similar / some bases in common; These bases will bind together / hydrogen bonds / complementary pairs; Do not accept same here. 		

		argument is being made.	2	
	(ii)	Relationship is closer / more complementary bases / more base pairs forming more hydrogen bonds;		
		 More heat energy needed (to separate bonds); Do not allow stronger hydrogen bonds. Not higher temperature as this is in question. 	2	[8]
М3.	(a)	(i) Order, Family, Genus. (all correct = 2 marks; 2 correct = 1 mark)		
		(all correct – 2 marks, 2 correct – 1 mark)	2	
	(ii)	3 concentric circles in Carnivora, labelled Felidae, Panthera and L;	1	
(t	o) (i)	large groups split into smaller groups (which do not overlap);	1	
	(ii)	(phylogenetic) based on evolutionary history; shows ancestry of groups / points of divergence / example, e.g. reptiles and birds separated after mammals / reptiles and birds more closely related than mammals; (hierarchical) based on shared characteristics (seen today);	3	[7]
M4. (a)	(i)	 Groups within groups; accept idea of larger groups at the top / smaller groups at the bottom No overlap (between groups); 	2	
	(ii)	(Grouped according to) evolutionary links / history / relationships /		

common ancestry;

Neutral: closely related

Accept converse providing that it is clear that the converse

- (b) (i) 1. (Only) one amino acid different / least differences / similar amino acid sequence / similar primary structure;
 - 2. (So) similar DNA sequence / base sequence;

2

- Compared with humans / not compared with each other;
 Accept: degenerate code / more than one triplet (codes) for an amino acid
 - Differences may be at different positions / different amino acids affected / does not show where the differences are (in the sequence);

1 max

(iii) 1. All organisms respire / have cytochrome c;

Accept: converse arguments for haemoglobin

- 1. Accept 'more' instead of 'all'
- 1. Accept 'animals' instead of organisms?
- 2. (Cytochrome c structure) is more conserved / less varied (between organisms);
 - 2. Neutral: cytochrome c is conserved

1 max

[7]

M5. (a) Is species specific / allows recognition of same species;

Greater similarity in calls the closer the relationship (between the species);

Accept: 'Similar species have similar calls' as first marking point.

Reference to courtship on its own is not sufficient for a mark. Must refer to relationship for second marking point.

2

(b) G. americana and G. monachus; Highest percentage (DNA hybridisation) / more bases are similar / complementary / more hydrogen bonds / more base pairings;

Second marking point can be awarded without first marking

- (c) 1. More closely related (species) have more similarities in amino acid sequence / primary structure;
 - 2. In same protein / named protein e.g. albumin;

OR

- 3. Similar species have a similar immune response to a protein / named protein;
- 4. More closely related (species) produce more 'precipitate' / antibody-antigen (complexes) / agglutination;

Accept: 'Similar species have similarities in amino acid sequence' for first marking point.

Accept: Converse for marking points 1, 4 and 5.

Marking point 5 is for measuring the extent of the immune response.

2 max

[6]

M6. (a) group of organisms with similar features; can (interbreed to) produce fertile offspring;

2

(b) directional selection; any TWO from selection against one extreme / for one extreme; against broadest beaks in B and narrowest beaks in A / for narrowest in B and broadest in A; whole distribution / range / mean / mode / median is shifted towards favoured extreme;

3 max

[5]

M7.(a) 1. No interbreeding / gene pools are separate / geographic(al) isolation;

Accept: all marks if answer written in context of producing increased diversity of plants

- 1 Do not award this mark in context of new species being formed and then not interbreeding
- 1 Accept reproductive isolation as an alternative to no interbreeding

- 2. Mutation;
 - 2 Accept: genetic variation
- 3. Different selection pressures / different foods / niches / habitats;
 - 3 Accept: different environment / biotic / abiotic conditions or <u>named</u> condition
 - 3 Neutral: different climates
- 4. Adapted organisms survive and breed / differential reproductive success;
- 5. Change / increase in allele frequency / frequencies;

5

1

(b) Similar / same environmental / abiotic / biotic factors / similar / same selection pressures / no isolation / gene flow can occur (within a species);

Accept: same environment

[6]

- **M8.** (i) Taxon **A** there is more than one level / taxon below it / genus only has species / only has one level / taxon above it;
 - (ii) Taxon **C** there is more than one level / taxon above it / phylum only has kingdom / only has one level taxon above it;

[2]