

## Mark schemes

**Q1.**

- (a) 1. Mutations/genotype/alleles;  
2. Environment/habitat

**OR**

(Natural) selection;

*Accept named different habitats, e.g. 'walls' and 'trees/trunks'.**Accept selection pressure e.g. predation.*

3. Epigenetics;  
4. Crossing over;  
*Accept recombination.*  
5. Independent segregation/assortment (of homologous chromosomes);  
*Accept 'Random assortment'.*  
6. Random fusion of gametes

**OR**

Random fertilisation;

**2 max**

- (b) 1. Provides camouflage;  
*Accept description of camouflage.*  
2. (So) not seen by predators/prey

**OR**

Less predation

**OR**

Obtain/catch (more) prey;

*Accept 'stops predation'.**Accept descriptions of reduced predation e.g. 'fewer are eaten'.***2**

(c) **Mark point 1 required for max marks**

1. (Geckos in) same habitat/environment/area

**OR**

No geographical isolation/separation (between geckos);

2. (Possibly) allopatric speciation as different (areas of same) habitat(s)/environment/area

**OR**

(Possibly) geographical isolation/separation as different (areas of same) habitat(s);

*Accept 'walls' and 'trees/trunks' as different habitats.*

3. (Could lead to) separate gene pools

**OR**

Reproductive isolation;

*Accept 'can't interbreed' in correct context, i.e., not when describing a species.*

*Accept 'no gene flow'.*

4. Mutation(s);

*Reject mutation(s) if context incorrect e.g., 'mutate to adapt', 'mutation caused by selection'.*

5. Selection for (both) extremes/colours

**OR**

Disruptive selection (occurs) as two extremes/ colours;

*Accept description of selection e.g. 'favoured'.*

*Accept selection against 'middle'*

6. (Analysis shows that) diurnal geckos are a distinct (genetic) group;

*Accept 'nocturnal' for 'diurnal'.*

7. (Genomes/DNA indicates geckos are) same species;

**5 max**

- (d) 1. Compare DNA base/nucleotide sequence

**OR**

Compare banding/position of DNA fragments;

*Idea of 'comparison' must be conveyed.*

*Accept alleles/VNTRs for 'DNA fragments'.*

*Accept genes for 'DNA fragments' in 1 but reject genes in mark point 2.*

*Ignore 'gene machine'.*

2. A distinct (group) will have different alleles/DNA/banding (from other group/s)

**OR**

If **not** distinct (group) will have similar alleles/DNA/banding (to other group/s)

*Accept genes for 'DNA fragments' in 1 but reject genes in mark point 2.*

*Accept in context of either nocturnal or diurnal group being the distinct group.*

*Accept not 'closely related' for 'distinct (group)' and 'closely related' for 'not distinct (group)'.*

*Reject 'species' for 'group'.*

3. DNA sequencing is automated/computerised

**OR**

Genetic/DNA fingerprinting is automated/ computerised

**OR**

PCR amplifies DNA/genes

**OR**

Genetic fingerprinting/electrophoresis separates fragments/genes/alleles

**OR**

Use of DNA probes/hybridisation to identify genes/alleles;

*Ignore 'gene machine'.*

- (e) 1. Marking not toxic **so** does not affect survival

**OR**

Marking not visible to predators

**OR**

Marking does not wash/rub off **so** recaptured (geckos) identified;

*Ignore births, deaths, reproduction, immigration, emigration.*

*Accept 'does not cause harm/death' for 'does not affect survival'.*

*Idea of marking affecting visibility required in relation to predators.*

2. Time/delay after release **so** (geckos) spread (in the population)

**OR**

Time/delay before recapture **so** (geckos) spread (in the population);

*Accept 'after marking' for 'after release'*

3. (Population =) (number in) first sample  $\times$  (number in) second sample divided by (number) marked in second sample / number recaptured;  
*Accept the correct equation/formula.*

**Q2.****(a) Mark as pairs: 1 and 2 OR 3 and 4**

1. Deletion/translocation;
2. Could mean triplet(s)/codon(s) missing

**OR**

Could mean amino acid(s) missing (from the polypeptide/SURF1);  
*Reject could mean an amino acid is not produced*

3. Substitution/inversion/addition/duplication/ deletion/translocation;
4. Could result in a (premature) stop triplet/codon;

**2 max****(b) Correct answer of 3 (people) = 3 marks;;;**

3.462564706/3.48/3.45 (or any correct rounding down to 1dp) =  
**2 marks** (answer not to the nearest whole number)

23 = **2 marks** (number of Faroe Islanders with nuclear mutations)

4 = **2 marks** (not factored in that only 80% of mutations are in nuclear DNA)

29 = **1 mark** (number of Faroe Islanders with LS)

**3****(c) Mark as pairs: 1 and 2 OR 3 and 4**

1. Genetic drift;
2. Frequency is higher by chance

**OR**

High frequency is not due to natural selection;

3. (Only) inbreeding/interbreeding (within a population)

**OR**

No (inter)breeding with other populations

**OR**

(Inherited from) common ancestor;

*Accept descriptions of inbreeding OR no interbreeding*

*Accept reproductively isolated*

*Accept genetic bottleneck OR founder effect*

4. Low genetic diversity

**OR**

Small gene pool

**OR**

Little gene flow

**OR**

Higher chance of inheriting allele

**OR**

Frequency of allele higher (in offspring);

**2 max**

- (d) **2 max for mark point 1 to 4 OR 5 to 7**

**Yes (no mark)**

1. Some people could be heterozygous/carriers;

2. Could prevent (human) suffering/death

**OR**

Could allow for (informed) decisions about having children;

3. (But only) in families/people with a history of LS

**OR**

(only) in families/people in the Faroe Islands (where high frequency/1: 1700);

4. Cost of screening might be cheaper than cost of treating LS;

**No (no mark)**

5. It is rare (globally)

**OR**

(Only) 1 in 40 000 (globally);

6. Caused by (too) many genes/one of 75 genes

**OR**

Would need (too) many probes/75 probes;

7. (Too) expensive to produce tests/probes (for more than 75 different genes)

**OR**

(Too) expensive to screen **all**;

**3 max**

**[10]**

**Q3.**

- (a) 1. (Colour vision involves) cones;  
*1, 2 and 3 Reject 'red cones/photoreceptors' and 'green cones/photoreceptors' only **once**.*  
*1 and 2 Each cone has a different pigment or absorbs particular wavelengths = **two** marks.*  
*1 and 3 Greater absorption by 'red sensitive' than 'green sensitive' cones = **two** marks.*

2. (Each type of) photoreceptor has a different pigment

**OR**

(Each type of) photoreceptor absorbs particular/different (range of) wavelength(s)

**OR**

(Each type of) photoreceptor stimulated by particular/different (range of) wavelength(s);

3. Greater absorption by 'red sensitive' than 'green sensitive' (cells/photoreceptors/cones)

**OR**

Provides percentage values which indicate difference in light absorption (at 600nm)

**OR**

More impulses to brain from 'red sensitive' than 'green sensitive' (photoreceptors/cones)

**OR**

More impulses along optic nerve from 'red sensitive' than 'green sensitive' (photoreceptors/cones);

*Allow approximately correct percentage values.*

*Accept suitable alternatives for 'sensitive' e.g. detecting/absorbing.*

*Accept action potentials for impulses.*

*Do **not** credit 'signals', 'messages' for third or fourth options for mark point 3.*

3

- (b) Box 3 correct - Several photoreceptors connecting to one neurone and spatial summation

1

- (c) 1. Geographical isolation/separation due to elevation/altitude;  
2. Allopatric speciation due to isolation/separation;  
3. Different selection pressures

**OR**

Different environment(s);

*Accept selection (due to) flower colour.*

*Accept different 'abiotic conditions/factors' for 'different environment(s)' or different named factor e.g. temperature, humidity but 'different altitudes' on its own is not enough.*

4. (However) some overlap in distribution

**OR**

(Both) plants found in same area/habitat/altitude;

5. So (possibly) sympatric speciation;  
*Only awarded if mark point 4 is credited.*
6. (Variation due to) mutation(s);  
*Reject mutation(s) if context incorrect e.g. 'mutate to adapt'.*
7. Reproductive isolation/separation due to different pollinators/distributions/altitudes

**OR**

Separate gene pools due to different pollinators/distributions/altitudes;

*Accept 'no gene flow' for separate gene pools.*

8. Change in allele frequency (in each population)

**OR**

Different allele frequency (in each population);

*Accept 'increase' or 'decrease' for 'change'.*

9. Different species) can no longer (interbreed to) produce fertile offspring;

5 max

[9]



**Q4.**

- (a) 1. *E. rufus* in north (west)  
**OR**  
*E. rufus* in the west  
**OR**  
*E. rufus* above river;
2. *E. rufifrons* in south  
**OR**  
*E. rufifrons* in west **and** east  
**OR**  
*E. rufifrons* below river;  
*1 and 2. Accept equivalent valid statements e.g., for 1, no E. rufus in south.*  
*1 and 2. If neither mark is awarded, accept, for one mark, 'they are separated by the river' OR 'there is no overlap in their distribution/ranges'.*  
*1 and 2. Accept converse.*  
*1 and 2. Do not penalise 'prefer'.*
3. (Actual) distribution similar to expected (distribution)  
**OR**  
 (Actual) distribution similar to environmental needs  
**OR**  
 (Actual) distribution (slightly) less than expected distribution;  
*Accept for one or both species.*
- (b) 1. Geographical isolation;  
**OR**  
 Allopatric speciation;  
*Ignore descriptions of geographical isolation.*  
*Reject sympatric.*  
*Ignore reference to two species at start.*
2. Reproductive separation/isolation  
**OR**  
 No gene flow  
**OR**  
 Gene pools separate;  
*Reproductive isolation must be at beginning of process.*  
*Accept no interbreeding but must be a separate idea from mp 6 which relates to definition of a species.*  
*Reject no inbreeding.*
3. Different selection pressures;  
**OR**  
 Different environmental/abiotic conditions/factors;

4. (Variation due to) mutation(s) (in different populations);
5. (Different/advantageous) allele/s passed on/selected  
**OR**  
Change in frequency of allele/s;
6. (Eventually different species) cannot (inter)breed to produce fertile offspring;

5 max

- (c) (Marking) does not affect survival/predation/recapture;  
*Accept. Mark does not rub/wash off/is non-toxic.*  
*Ignore 'does not harm' on its own unless it relates to survival/predation/recapture.*

1

- (d) 3;  
*Ignore any wording provided e.g. lemurs.*

1

**[10]**