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NEW ZEALAND QUALIFICATIONS AUTHORITY
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Level 2 Biology, 2016

91157 Demonstrate understanding of genetic variation and change

9.30 a.m. Friday 18 November 2016
Credits: Four

Achievement	Achievement with Merit	Achievement with Excellence
Demonstrate understanding of genetic variation and change.	Demonstrate in-depth understanding of genetic variation and change.	Demonstrate comprehensive understanding of genetic variation and change.

Check that the National Student Number (NSN) on your admission slip is the same as the number at the top of this page.

You should attempt ALL the questions in this booklet.

If you need more space for any answer, use the page(s) provided at the back of this booklet and clearly number the question.

Check that this booklet has pages 2–11 in the correct order and that none of these pages is blank.

YOU MUST HAND THIS BOOKLET TO THE SUPERVISOR AT THE END OF THE EXAMINATION.

TOTAL

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QUESTION ONE: INHERITANCE AND MEIOSIS

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Roses display complete dominance in both their flower colour and in their susceptibility to some diseases. The allele for red petals (R) is dominant to the allele for white petals (r). In addition, the allele for healthy leaves (H) is dominant to the allele for being susceptible to leaf lesions (h). Leaf lesions are spots on the leaf that are very prone to disease and injury. The genes for petal colour and healthy leaves are located on different chromosomes.



Leaf with lesions.

<https://edis.ifas.ufl.edu/pp267>



<http://www.tophdwallpapersland.com/red-white-rose-wallpaper.htm>

A rose that was homozygous for both red petals and healthy leaves was crossed with a white rose that was susceptible to leaf lesions.

- (a) State the genotype of the F₁ generation this cross produces.

- (b) Use the Punnett square below to show the gametes of the F₁ cross, and all of the possible genotypes of the F₂ generation.

		F ₁ gametes			
F ₁ gametes					

- (c) Describe the predicted phenotype ratios produced by this cross.

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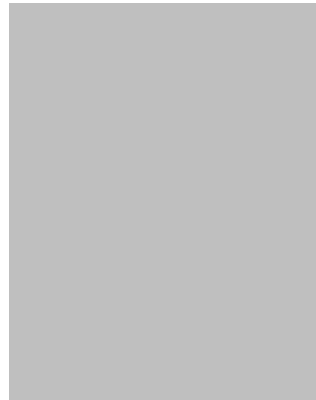
- Your answer should include:

- You may use diagrams in your answer.

QUESTION TWO: NATURAL SELECTION IN MOA

A large body mass is an advantage in cooler climates because its low surface area to volume ratio helps animals to retain heat. Many examples of this, such as polar bears, walrus and large polar sea mammals, are seen today.

Fossil evidence shows that during the last ice age, the population of heavy-footed moa, *Pachyornis elephantopus*, contained much larger individuals than the same species of moa that existed during warmer times. As the ice age ended and temperatures warmed, the fossil evidence shows that the heavy-footed moa's body mass became smaller again.



[http://collections.tepapa.govt.nz/search.aspx?term=Heavy-footed moa](http://collections.tepapa.govt.nz/search.aspx?term=Heavy-footed%20moa)

Change in moa body mass over time



Moa body mass data calculated from femur bone circumferences.

Worthy, Trevor H. and Richard N. Holdaway, 2002. *The Lost World of the Moa, Prehistoric life in New Zealand* (Indiana University Press, Bloomington), Table 5.6, p. 20.

The large body mass allele may have entered the population via a mutation.

Discuss how the allele for large body mass became established in the heavy-footed moa gene pool during the last ice age.

Your answer should include:

- a description of what a gene pool is
- a description of what a mutation is and an explanation of how it affects genetic variation in a species
- a discussion of the process of natural selection and how it affected both the body mass and the gene pool of the heavy-footed moa
- a discussion, with justified reasons, why the body mass of the heavy-footed moa returned to a smaller mass once the climate warmed again.

Many of New Zealand's native species have suffered population bottlenecks due to hunting, introduced predators, and habitat destruction. The Department of Conservation has successfully saved some of these species from extinction by moving several breeding pairs from mainland populations to predator-free islands. However, maintaining genetic diversity on island populations can be difficult for many species of flightless birds, such as the takahe, *Porphyrio hochstetteri*.

www.nzbirdsonline.org.nz/species/south-island-takahe

- a description of what genetic diversity is
- an explanation of how allele frequency in a population is affected by genetic drift and migration
- a discussion of how migration and genetic drift affect genetic diversity of flightless birds on small island populations compared to larger mainland populations.

Extra paper if required.
Write the question number(s) if applicable.

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