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CEL191 LECTURE MATERIAL

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CEL191 2025

Molecular Biology & Genetics

Lecture 21

Population Genetics & Natural Selection

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Te Tari Mātai Kararehe | Department of Zoology

Lecture 21 Objectives

After you have revised this lecture you should be able to:

- ❖ Calculate genotype frequencies using Hardy-Weinberg equilibrium.
- ❖ Identify that random genetic drift occurs rapidly in small populations.
- ❖ Explain how populations evolve over time in response to selection.

POPULATIONS EVOLVE, INDIVIDUALS DON'T

- ❖ **Population:** localised group of individuals of the same species
- ❖ **Gene Pool:** total aggregate of genes (and their alleles) in the population at one time

Why might we need to estimate frequencies of genotypes in a population?

1. To predict how many individuals will inherit a genetic disease
2. To estimate the proportion of individuals who are 'carriers' of a genetic disease

HARDY-WEINBERG EQUATION

Males

Females

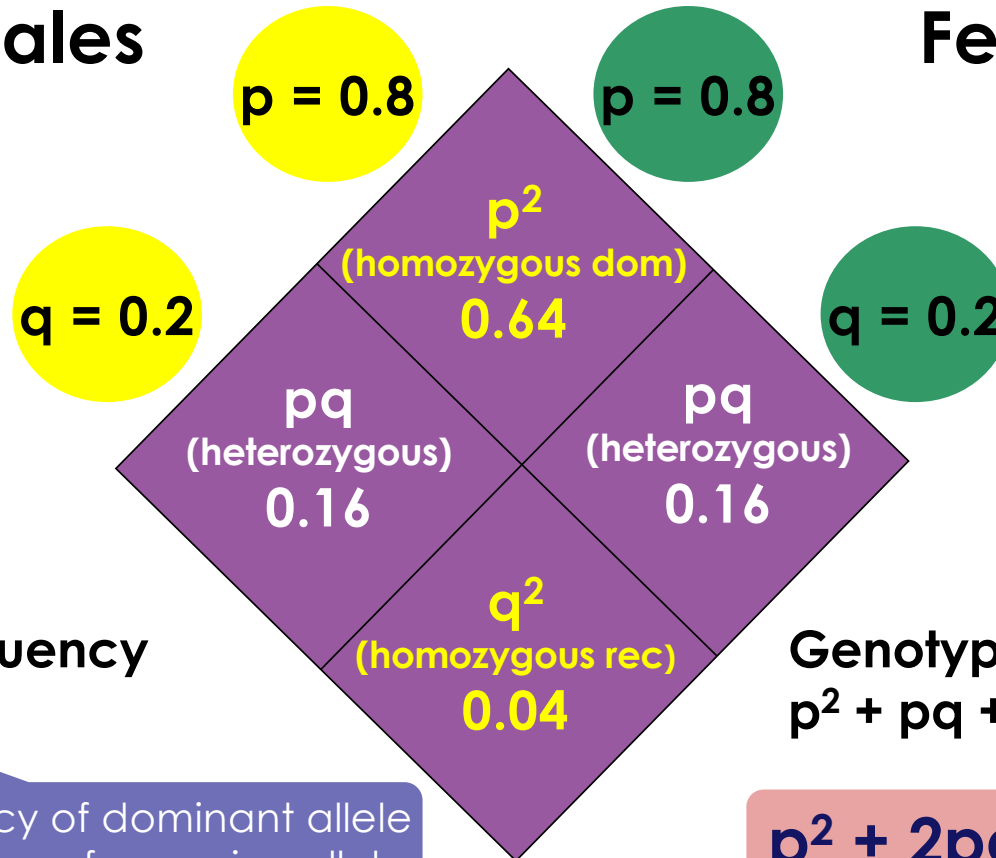
How do we
predict
genotypes in
a population?

Allele frequency
 $p + q = 1$

p = frequency of dominant allele
 q = frequency of recessive allele

Genotypic frequency
 $p^2 + pq + pq + q^2 = 1$

$$p^2 + 2pq + q^2 = 1$$



Example Question 1

1 in 2500 children are born with cystic fibrosis.
What proportion of the population are carriers for the disease?

A: Carriers = heterozygous = $2pq$therefore:

Estimate q:

$$q^2 = 1/2500$$

$$q = 1/50 = 0.02$$

(recessive allele)

Estimate p:

$$1 - q = p = 0.98$$

(dominant allele)

Estimate $2pq$:

$$2pq = 2(0.98 \times 0.02)$$

$$= 0.0392$$

= about 4% of the population are carriers

Example Question 2

If 1 in 10,000 children is born with Tay-Sachs disease, what percentage of the population are expected to be carriers for this autosomal recessive disease?

- (A) 0.10%
- (B) 0.20%
- (C) 0.99%
- (D) 1.98%

Estimate q:

$$q^2 = 1/10000$$

$$q = 1/100 = 0.01$$

Estimate p:

$$1 - q = p = 0.99$$

Estimate 2pq:

$$2pq = 2(0.99 \times 0.01) \\ = 0.0198$$

= about 2% of the population are carriers

Example Question 3

If the proportion of X chromosomes with the allele for haemophilia (a) in a population is 0.4, how many females will be affected by the disease? And how many males?

$$X^a = q = 0.4$$

$$\text{Females} = X^aX^a = q^2 = 0.4^2 = 0.16 \text{ or } 16\%$$

$$\text{Males} = X^aY = q = 0.4 = 40\%$$

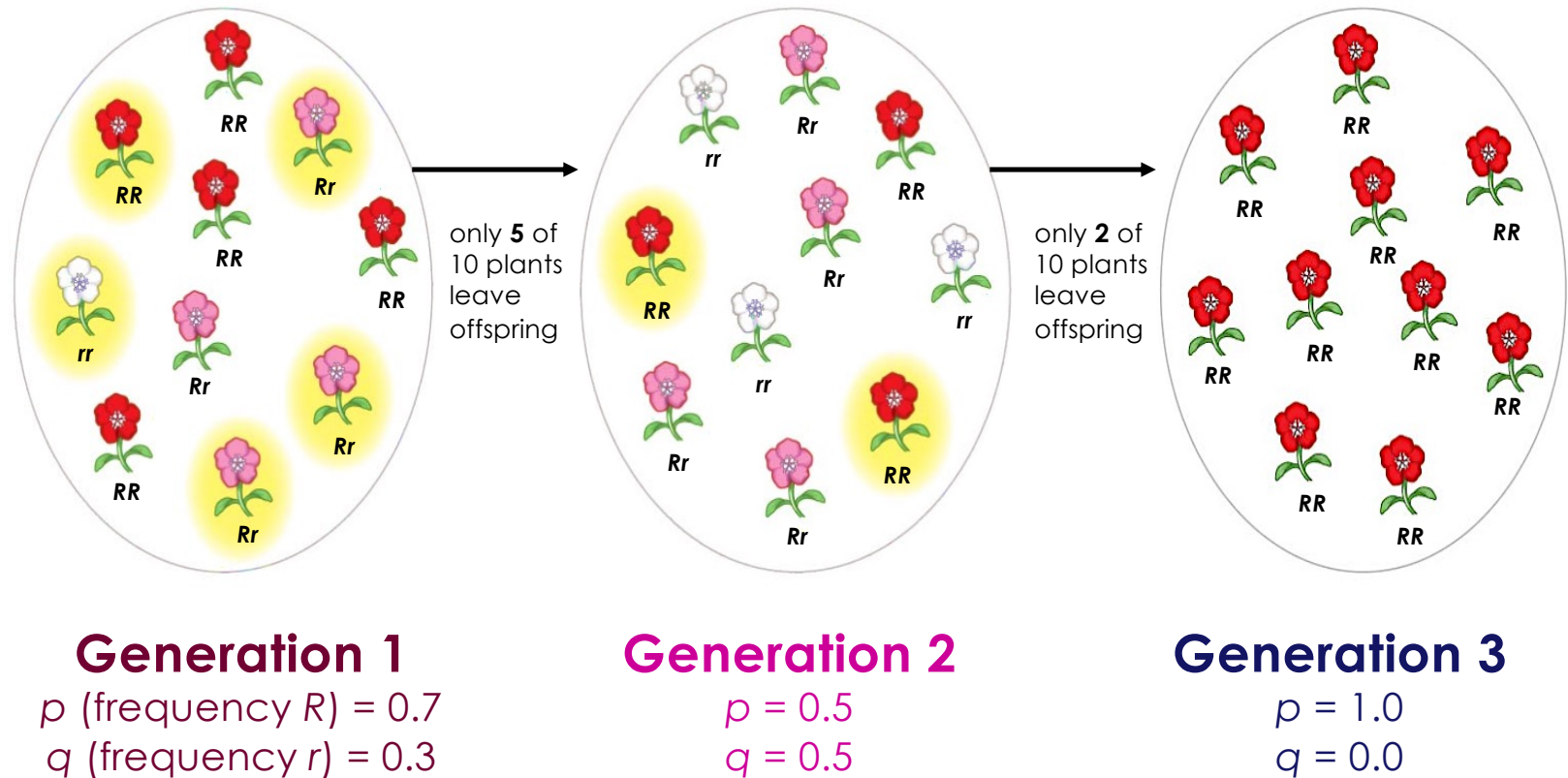
Remember, for an x-linked trait: females need to be homozygous recessive to get the disease; males need just one recessive allele

Allele frequencies can change via...

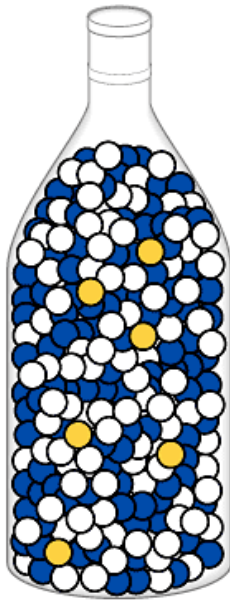
- ❖ **Non - random mating**
 - *assortative mating*
 - *inbreeding*
- ❖ **Random Genetic Drift**
- ❖ **Bottleneck Effect**
- ❖ **Founder Effect**
- ❖ **Natural Selection**
- ❖ **Gene Flow or Migration**
- ❖ **Mutation**

Random Genetic Drift

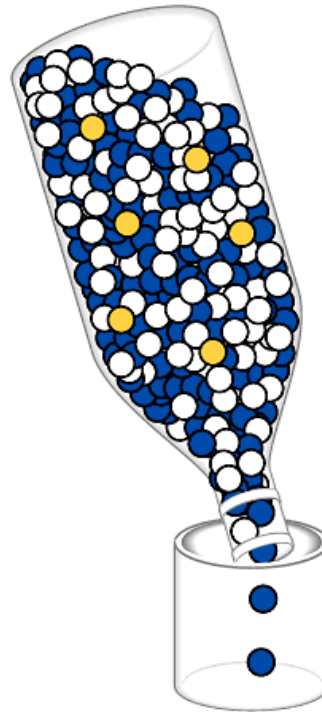
A random change in allele frequencies due to sampling error over generations



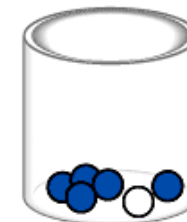
The Bottleneck Effect



Original population has equal frequencies of blue and white with a few yellow

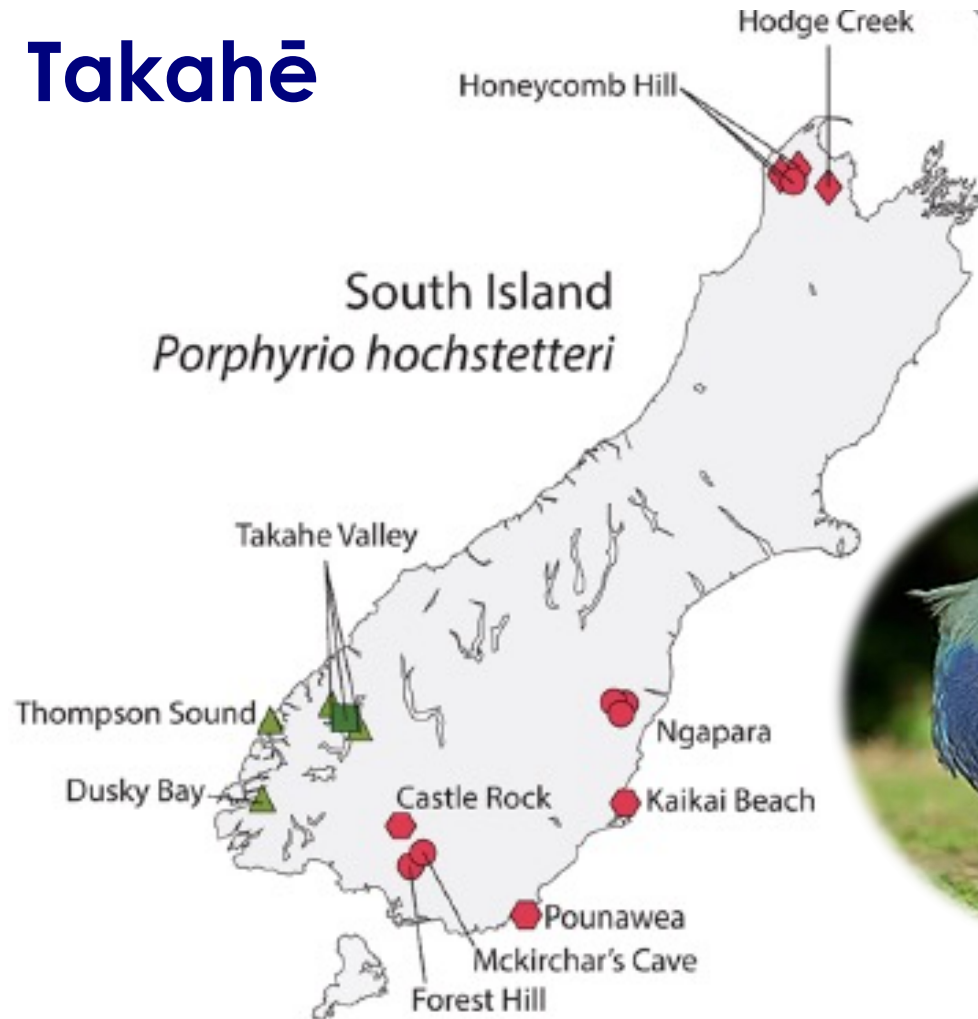


Population reduction or Bottleneck event

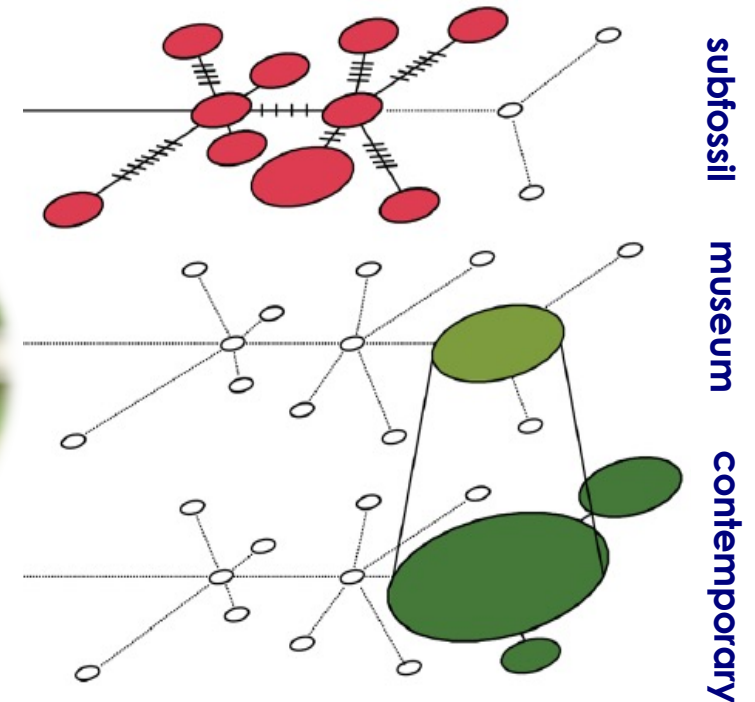


Yellow allele completely lost, white allele now rare

Takahē

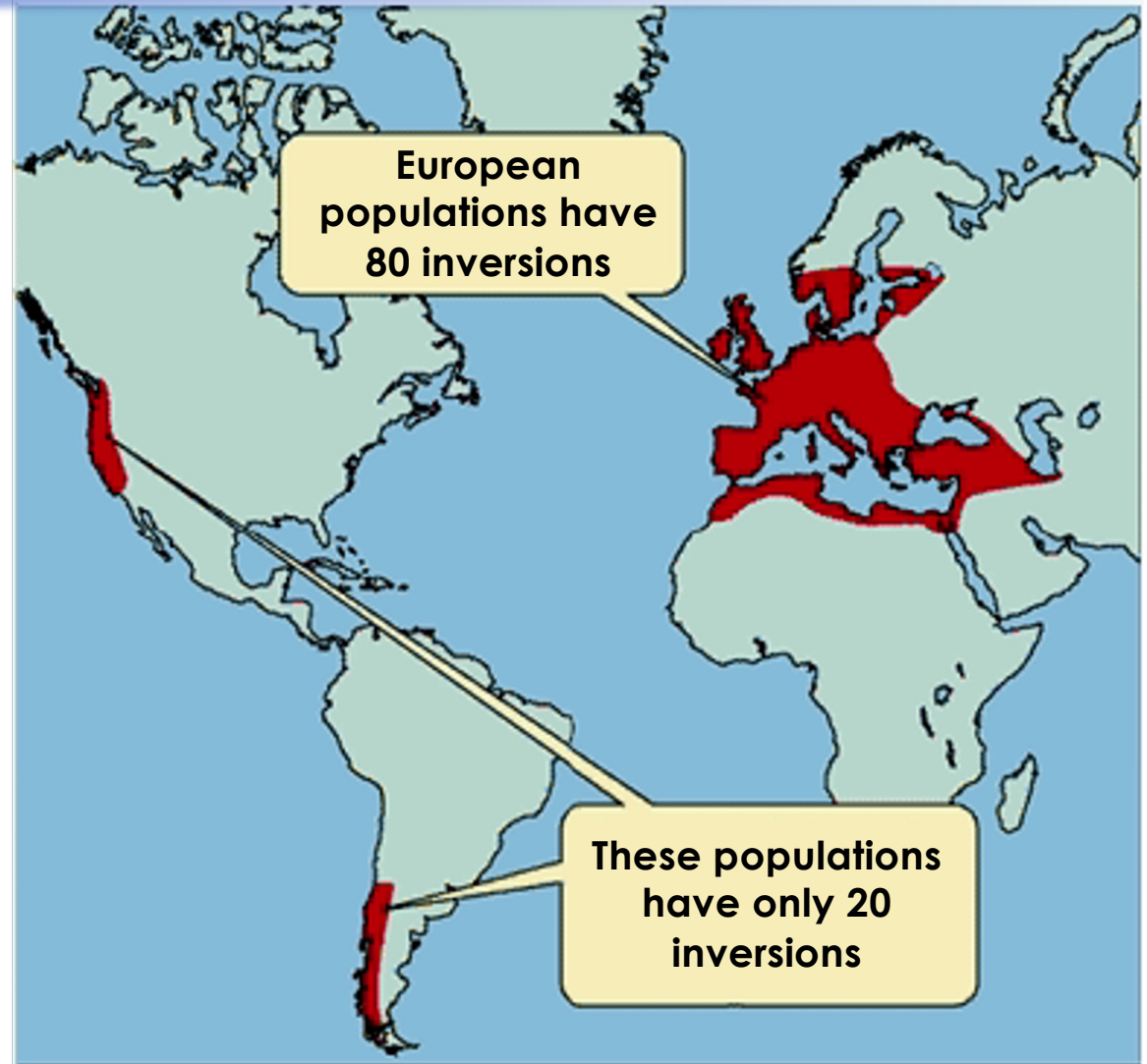


Bottlenecks in NZ's endangered birds

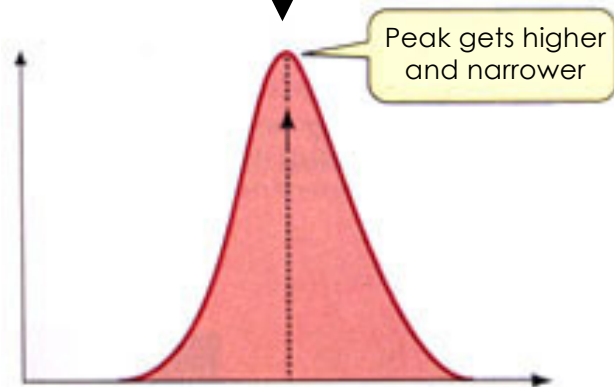
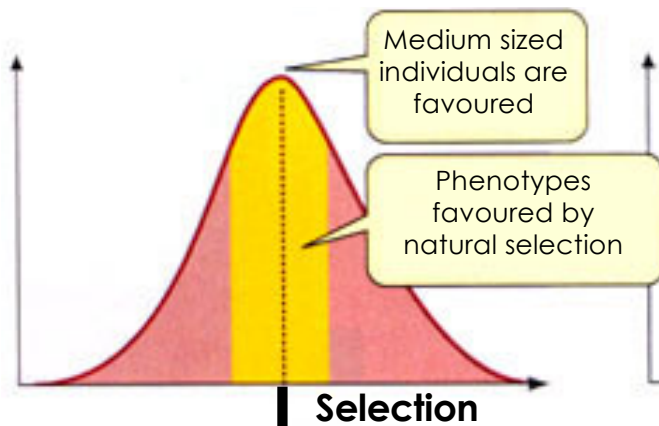


The Founder Effect

Within 20 years of arriving in the New World, populations of *D. subobscura* have increased dramatically, but have reduced genetic variation

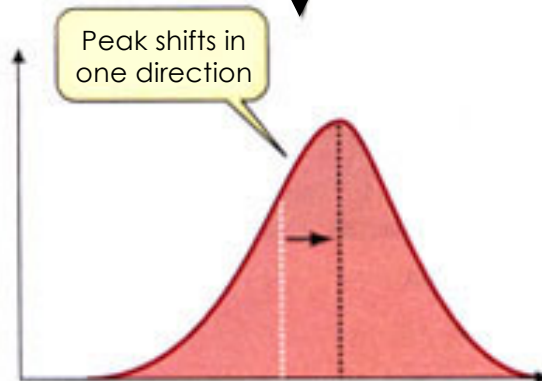
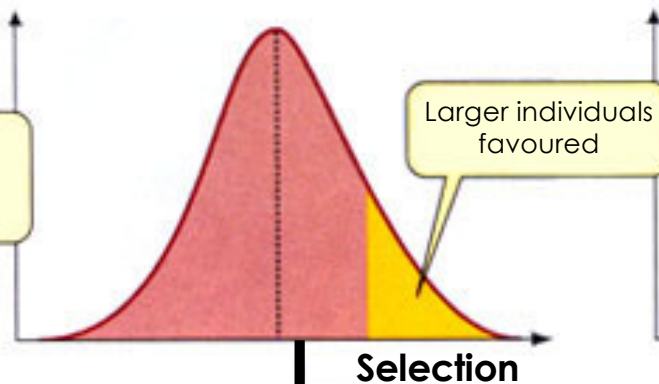


Stabilising Selection



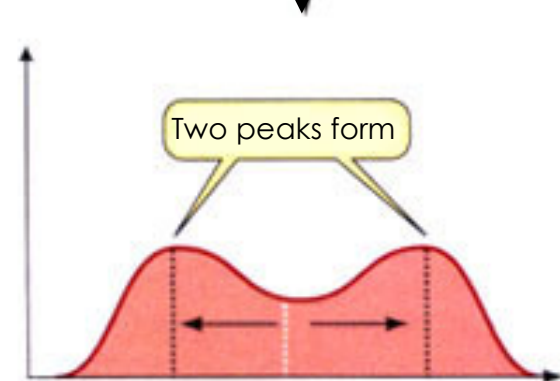
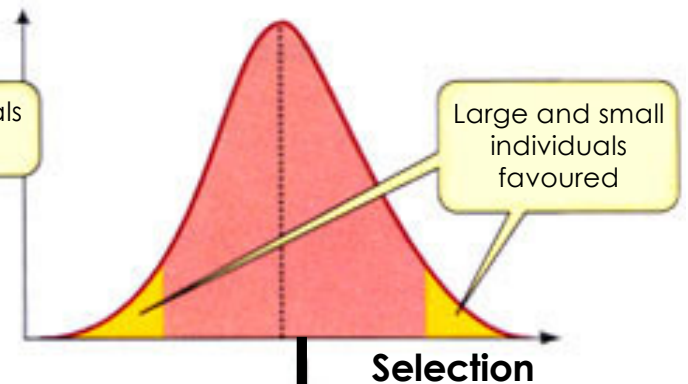
Reduces variation but does NOT change the mean

Directional Selection



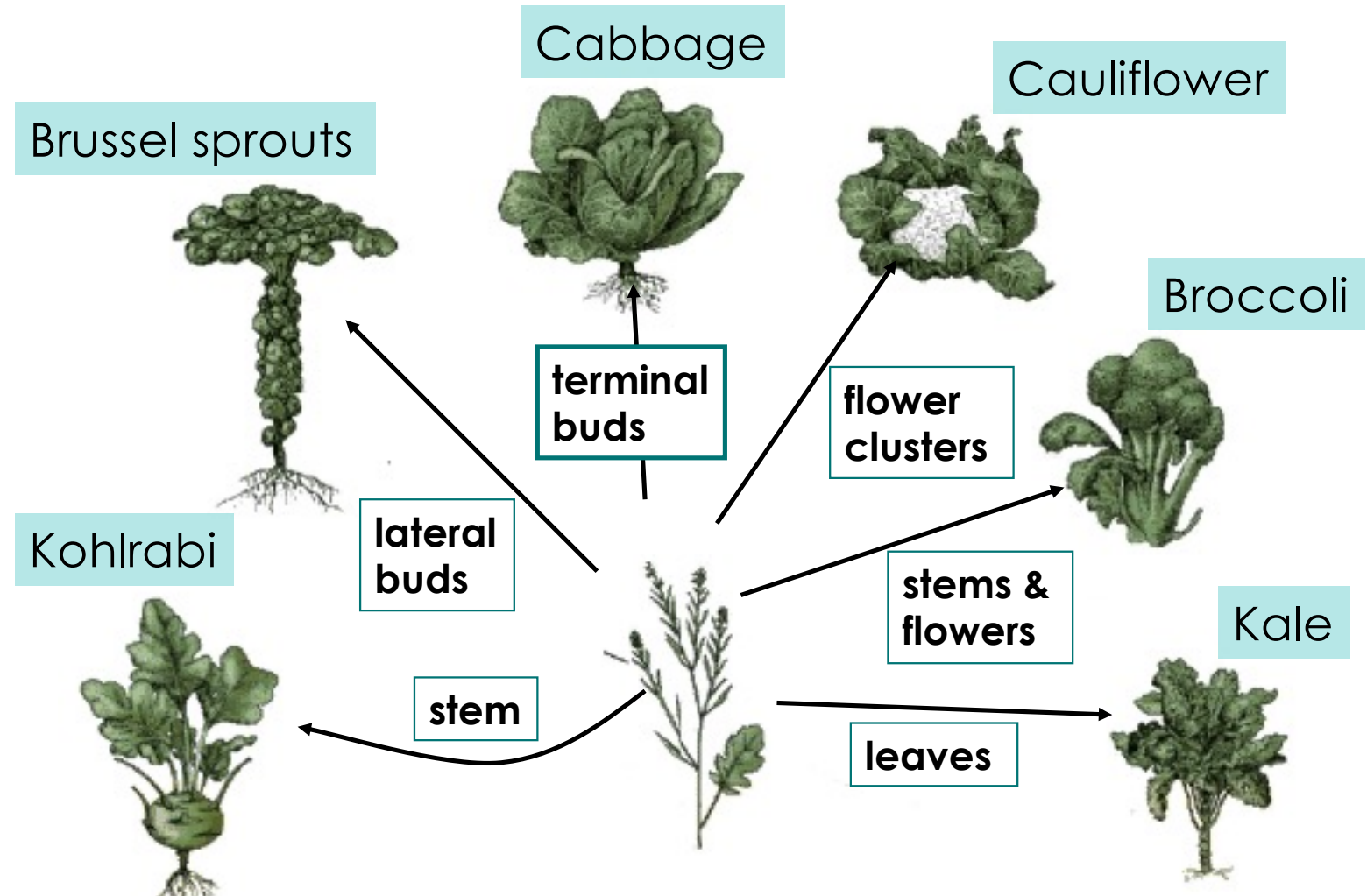
Changes the mean value towards one extreme

Disruptive Selection

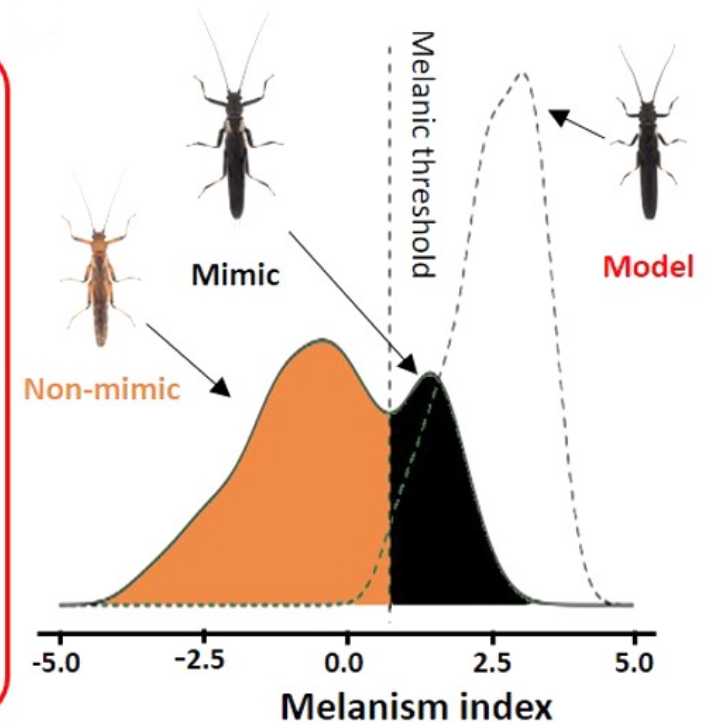
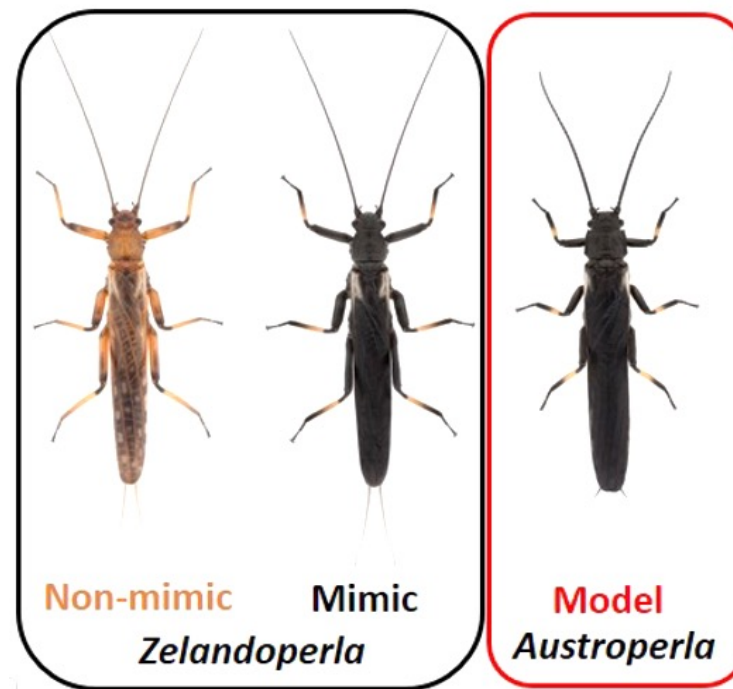


Favours the two extremes producing two peaks

Directed selection on common wild mustard

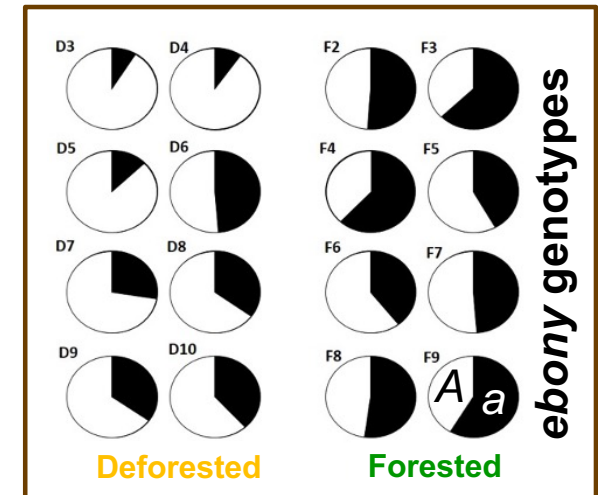
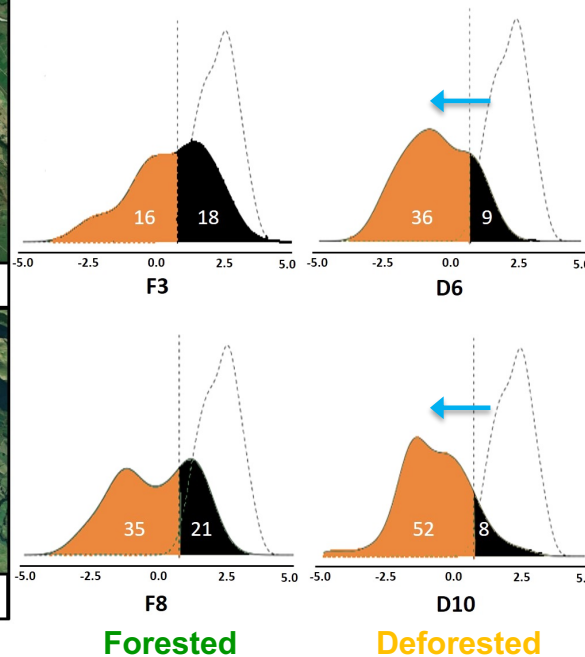
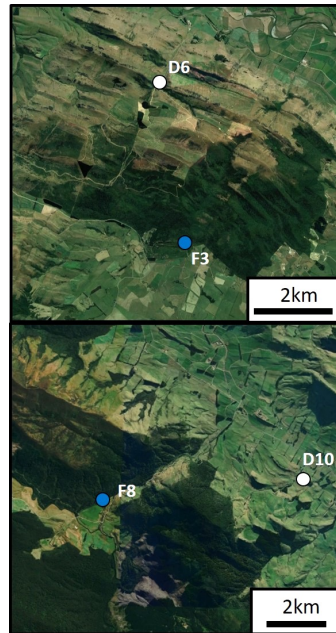
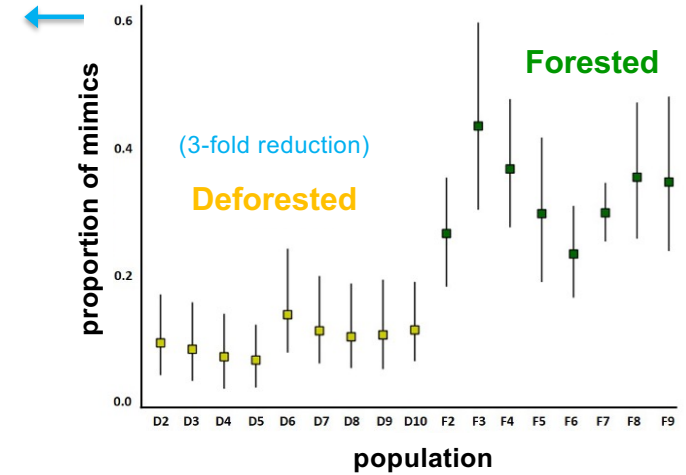


Directional selection on insect colour

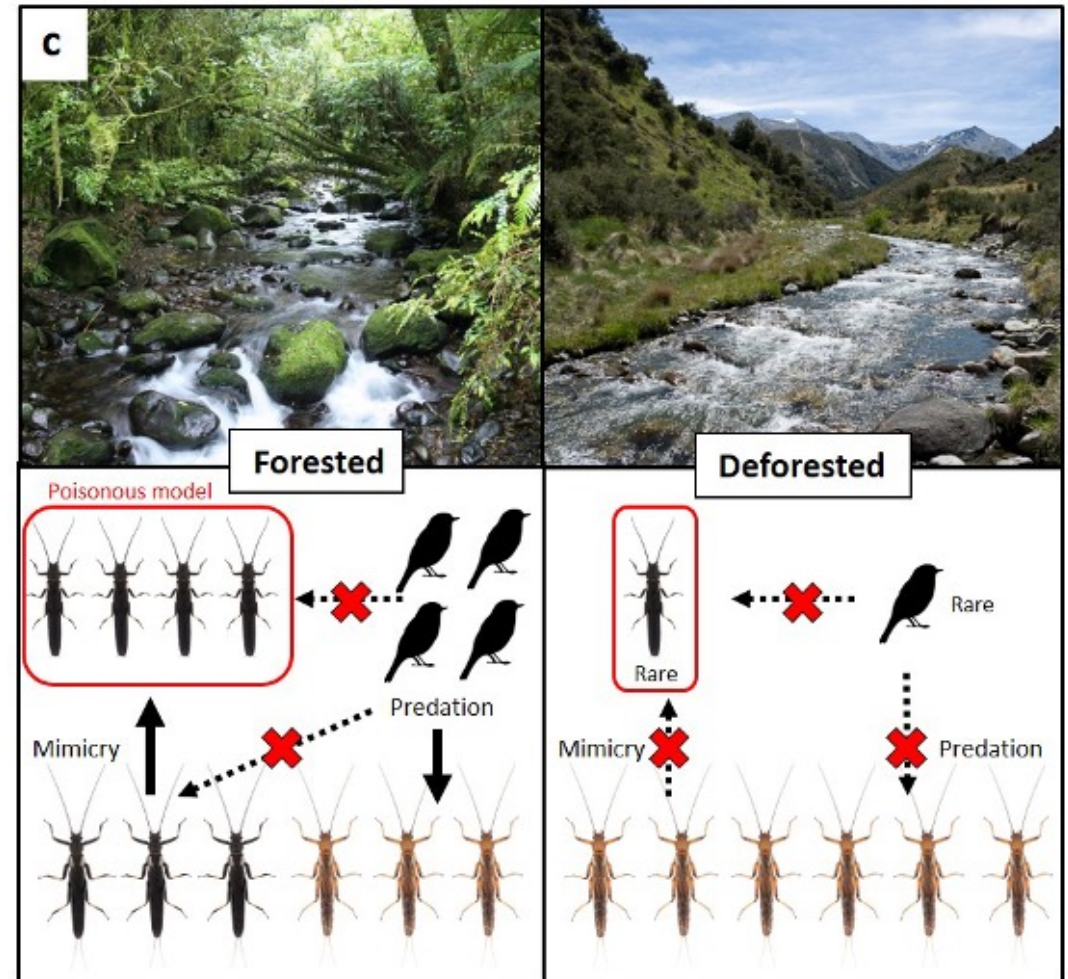


Directional selection on insect colour

Phenotypic shifts:
fewer dark individuals in
deforested habitats



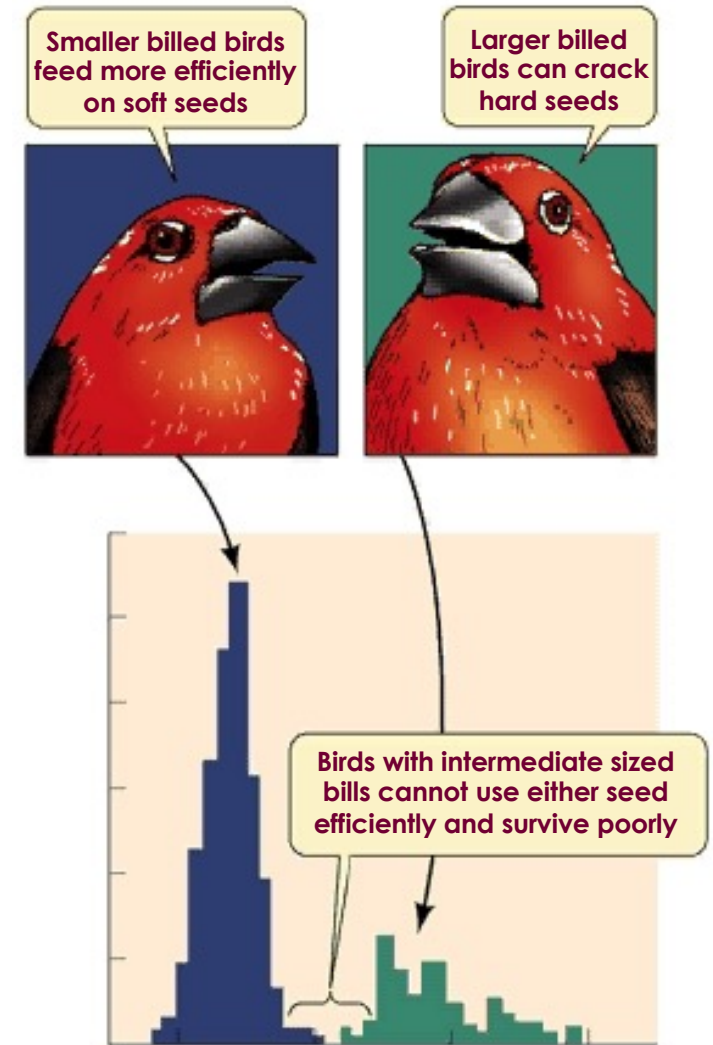
Directional selection on insect colour



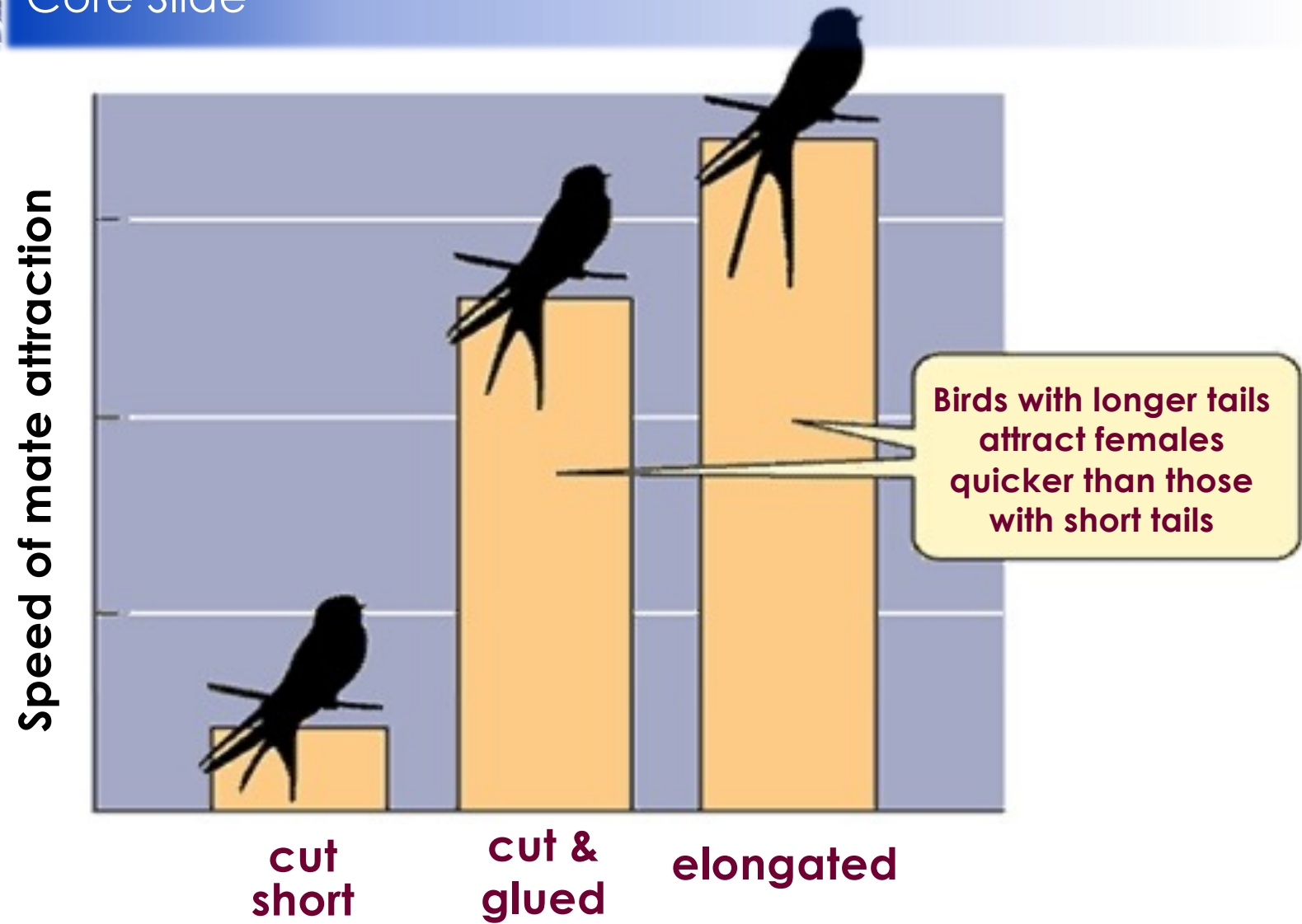
West African Black-bellied Seedcrackers display two bill morphologies

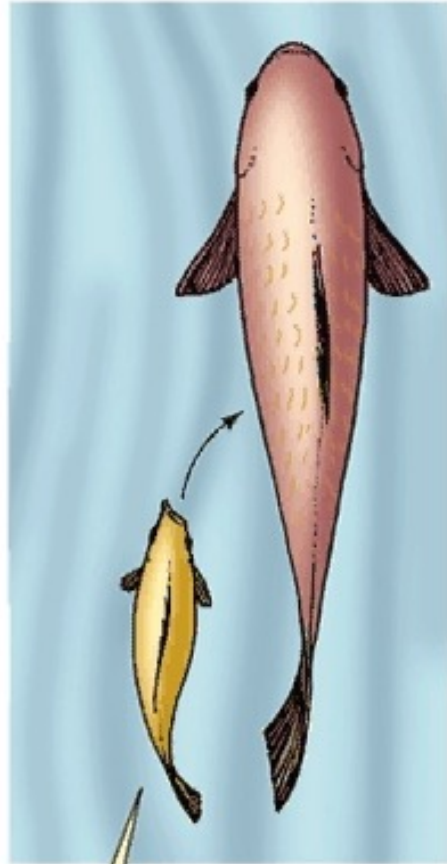
What type of selection is this?

DISRUPTIVE SELECTION

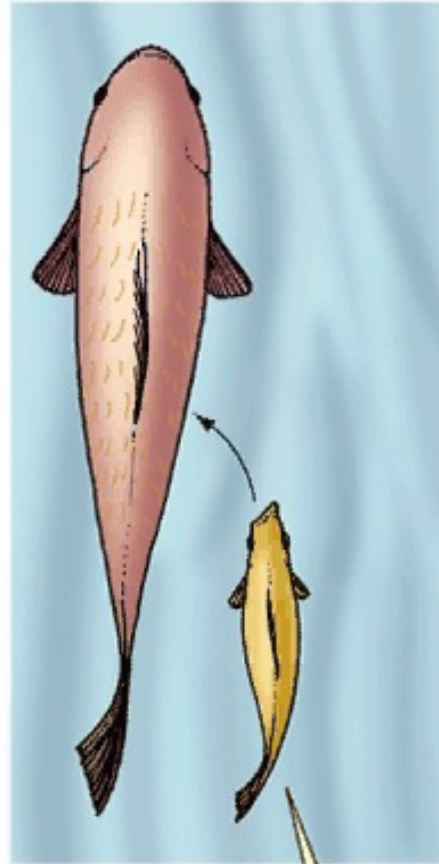


Sexual Selection





right
mouthed

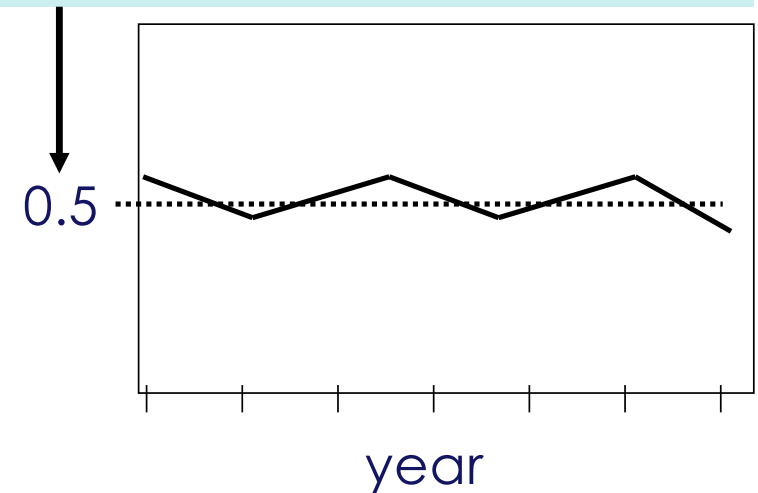


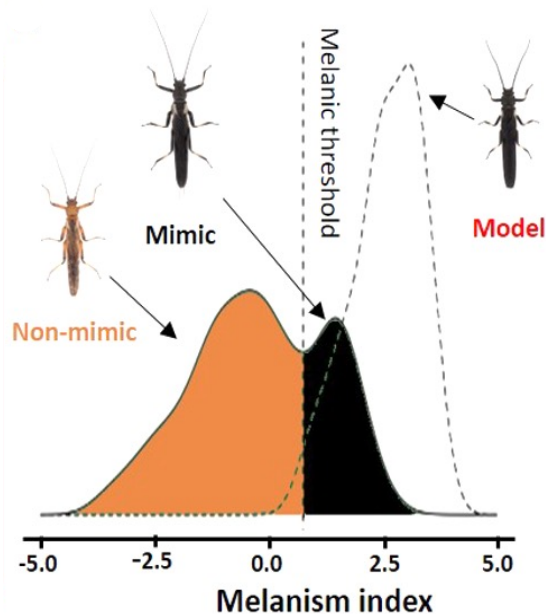
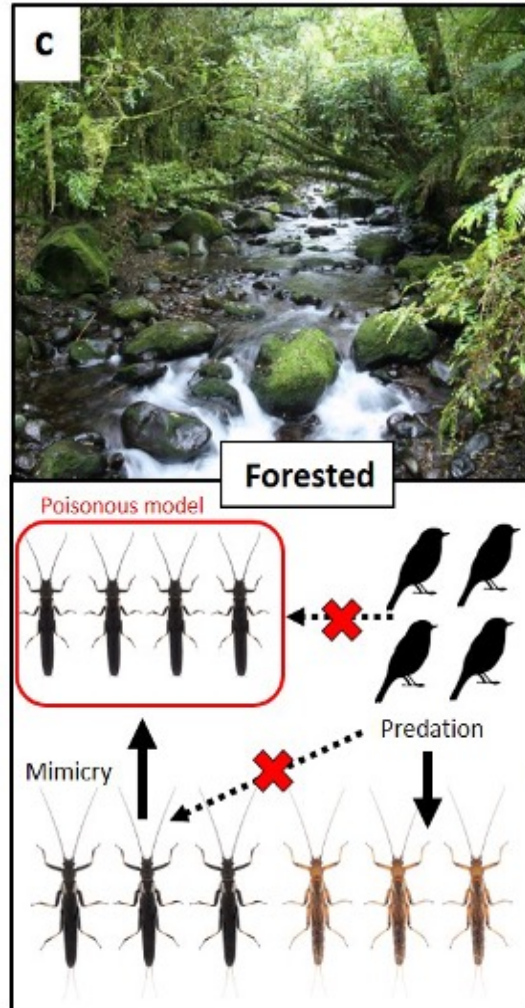
left
mouthed

Frequency dependent selection in *Perissodus microlepis*

Natural
selection
maintains an
equal
proportion of
left and right
mouthed
individuals in
the
population

Mean frequency of left mouthed
Perissodus





Frequency dependent selection?

Why are there any non-mimics in the population?

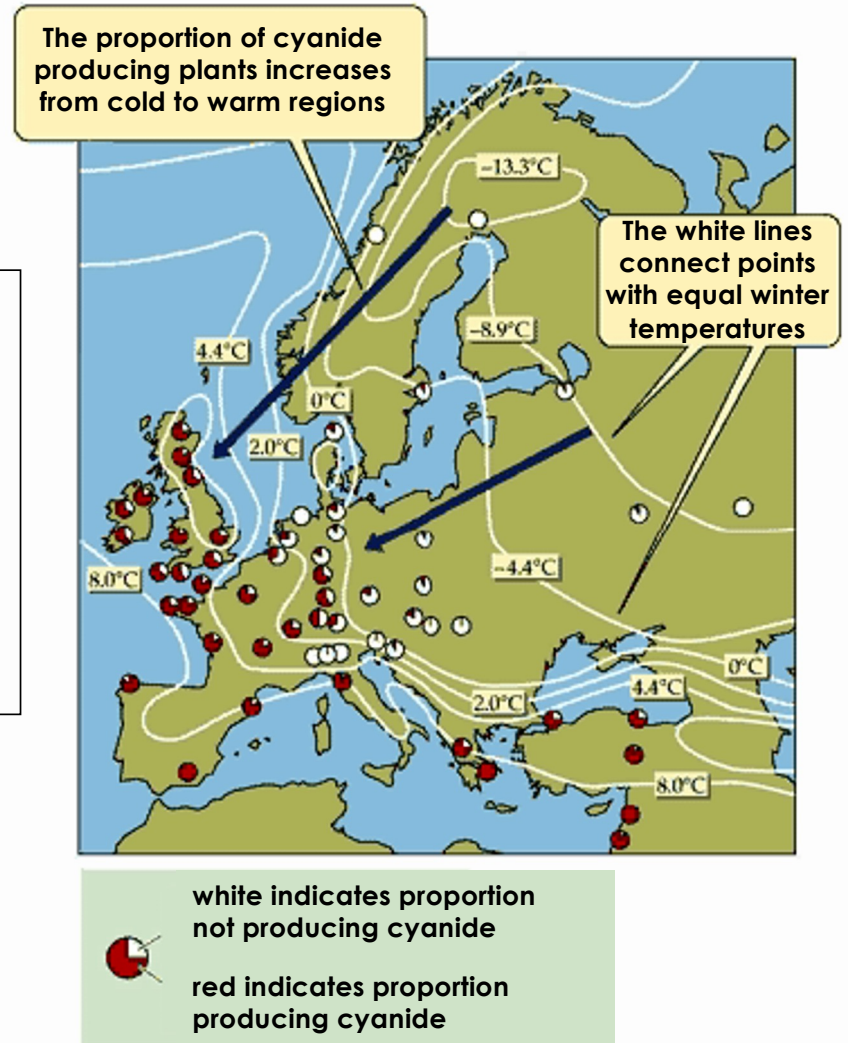
What happens if mimics start to outnumber the poisonous 'model'?

Spatial Distribution of Genetic Variation

Cyanide production in clover
Trifolium repens

The gradual geographic change in genetic/phenotypic composition is termed at:

CLINE



Mutation

Very slow to act and usually disadvantageous, its role is usually of macroevolutionary proportions

Migration

An individual from another population successfully mates (i.e. contributes gametes) to the gene pool.

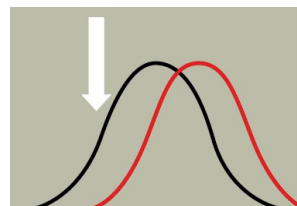
- ❖ brings new alleles
- ❖ changes proportions of existing alleles
- ❖ changes population size
- ❖ makes two populations more similar

Lecture 21 Summary

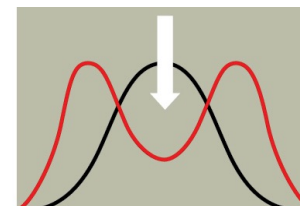
- ❖ The Hardy-Weinberg equation can be used to predict genotype frequencies in a population
- ❖ Small populations rapidly lose allelic diversity as a result of genetic drift.
- ❖ Different types of natural selection lead to contrasting shifts in the phenotypic and genotypic composition of populations
- ❖ In cases of frequency-dependent selection, the fitness of a genotype is influenced by how rare or common it is.

Objective-Based Questions

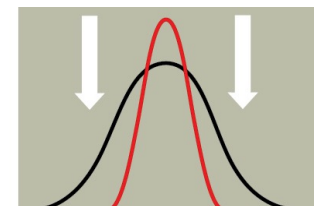
- ❖ Why would we want to estimate genotype frequencies in a population?
 - ❖ State the Hardy-Weinberg equation for calculating genotypic frequency and define the terms p , q and $2pq$.
- If $p=0.8$ in a certain population, what
- a) is the value of q ?
 - b) % of the alleles will be the dominant allele?
 - c) is the frequency of individuals that are homozygous dominant?
 - d) is the frequency of individuals that are homozygous recessive?
 - e) is the frequency of heterozygous individuals?
- ❖ If 1 in 10000 children is born with Tay-Sachs disease, what percentage of the population are expected to be carriers for this autosomal recessive disease?
 - ❖ Using an example, explain what is meant by the Founder Effect.
 - ❖ Identify the modes of selection in figures (a)-(c) on the right. Note: the large white arrows symbolize selective pressures against certain phenotypes.



(a)



(b)



(c)

If you have any questions relating to this part of the course then.....

Review the lecture resources on Blackboard, watch the lecture recordings, use the Learning Modules, read your textbook, ask during the lab, come along to a drop-in session, contact me, ask ask ask.....

As teachers we appreciate the difficulty of learning this stuff so please do not be afraid to ask for help. However, you will only get better by practising - so go through questions and understand your answers right or wrong. Get a feeling for the concepts - what are the main points? Once you understand this, the rest becomes easier.....**honest!**



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