

# Solving genetic problems

BIOL3120 – LECTURE 6



# Where to find the problem sets?

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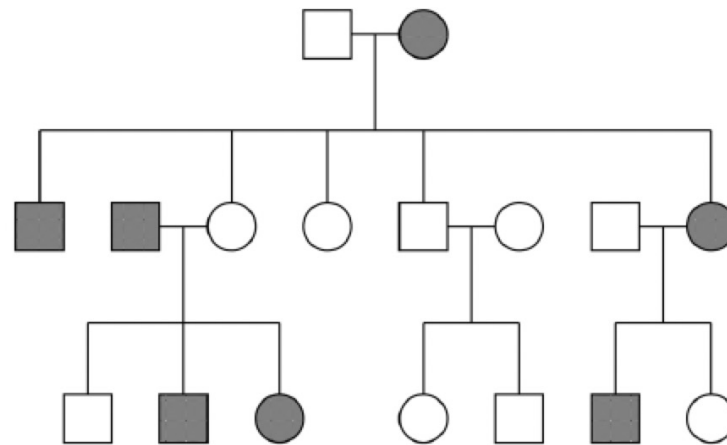
iLearn!

In the assessments tab

# AT1

## PROBLEM SET ASSESSMENT

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1. What is the *most likely* mode of inheritance for the condition shown in the above pedigree? Why?
- a) Autosomal dominant
  - b) Autosomal recessive
  - c) X-linked dominant
  - d) X-linked recessive

# Special considerations

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## Student wellbeing / special considera... ▼

If you experience events or conditions that adversely affect your academic performance, please visit the [special consideration page](#).

Visit the the [Student Support](#) webpage to see a range of services available to students at Macquarie University.

If you feel you are struggling with studies or have had disruptions in your personal life, you can speak to a trained professional at [Campus Wellbeing](#).

# Your job as geneticists is to work out which tools you can use

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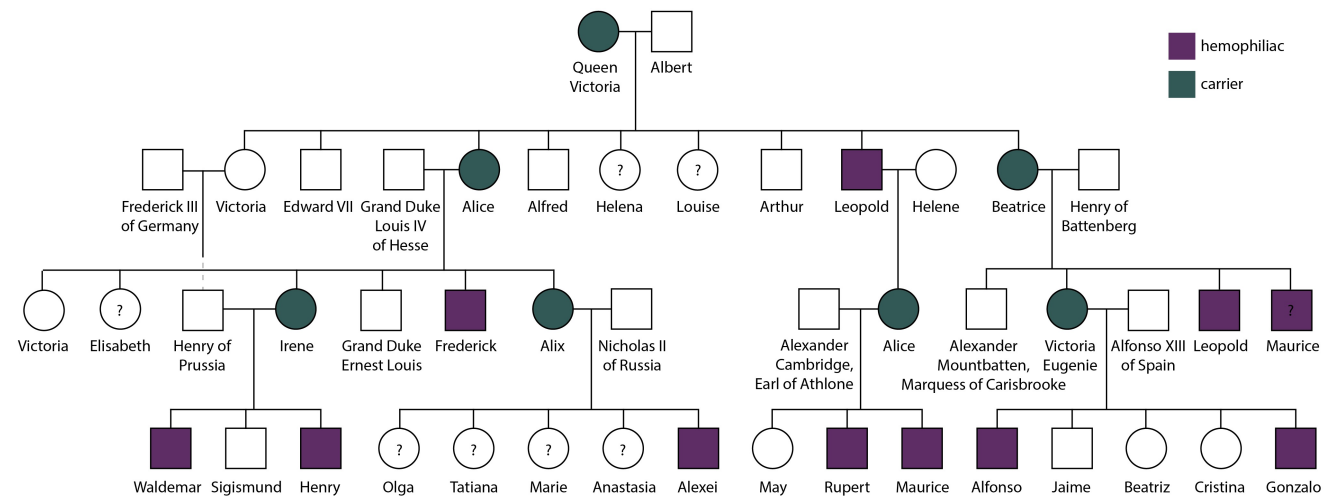
**You already know all the tools to solve genetic problems**

- Using family trees to infer inheritance patterns
- Hardy-Weinberg equilibrium
- Punnett squares to find expected phenotypes
- Simple probability



# Using family trees to infer inheritance patterns

- Useful for inferring inheritance patterns
- Useful for inferring individual's genotype



# Hardy-Weinberg equilibrium

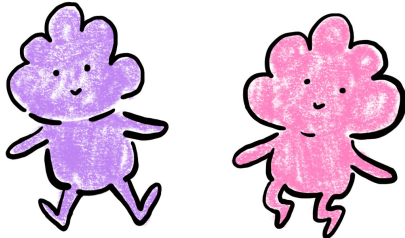
## Hardy-Weinberg equilibrium

If there are only 2 alleles  
for a trait in a Population, then:

$$P + q = 1$$

frequency of  
dominant allele

frequency of  
recessive allele



Purple is dominant to Pink

## Hardy-Weinberg equilibrium

If there are only 2 alleles  
for a trait in a Population, then:

$$P^2 + 2Pq + q^2 = 1$$

frequency of  
homozygous  
dominant  
genotype

frequency of  
heterozygous  
genotype

frequency of  
homozygous  
recessive  
genotype



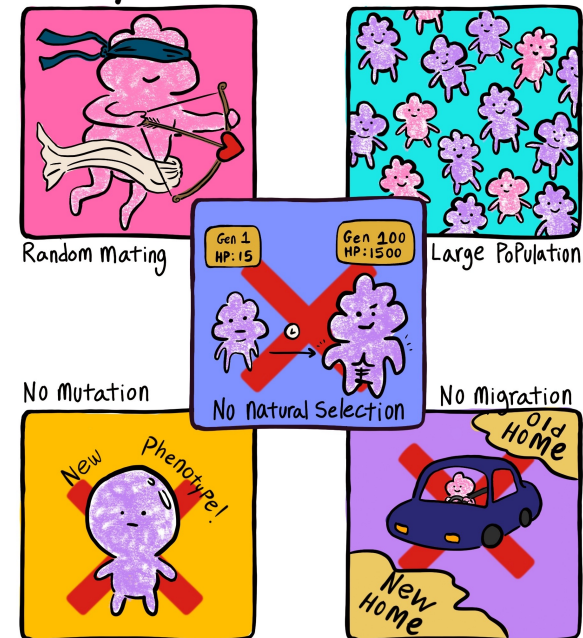
Purple is dominant to Pink



# Hardy-Weinberg equilibrium

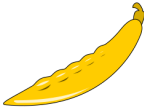





- Assumptions of Hardy-Weinberg equilibrium
  1. Large population (avoids random chance having a big impact on allele frequencies).
  2. No selection (ie no allele is necessarily more likely to get passed on than another)
  3. Mating is random (ie any individual has an equal chance of mating with any other individual).
  4. Mutation either does not occur or is in equilibrium.
  5. Immigration and emigration do not occur.

## Hardy-Weinberg Assumptions















# Punnett squares to find expected phenotypes

	g 	g
G 	Gg 	Gg 
g	gg 	gg 

## Punnett Square in Pea plant

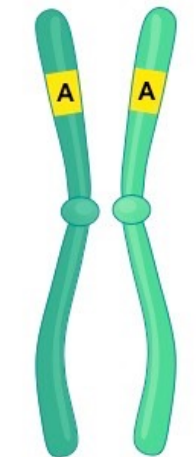
		 Round Yellow (RrYy)			
		RY	Ry	rY	ry
RY		RRYY	RRYy	RrYY	RrYy
Ry		RRYy	RRyy	RrYy	Rryy
rY		RrYY	RrYy	rrYY	rrYy
ry		RrYy	Rryy	rrYy	rryy

  
Round Yellow (RrYy)

 Round Yellow (n=9)  
 Round Green (n=3)  
 Wrinkled Yellow (n=3)  
 Wrinkled Green (n=1)

**F2 Generation**

# Punnett squares to find expected phenotypes



**Homozygous**  
*Alleles are same*



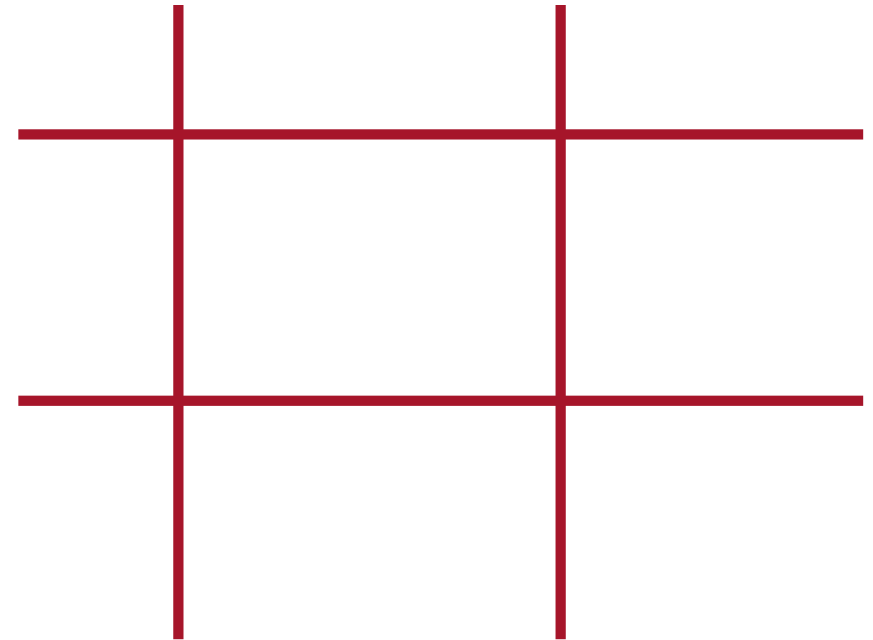
**Heterozygous**  
*Alleles are different*



**Hemizygous**  
*Only one allele  
(e.g. XY)*

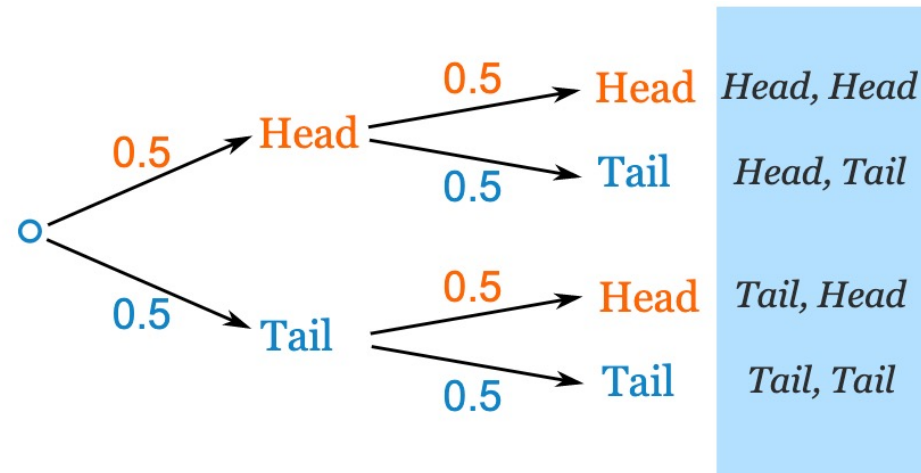
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# Simple probability

- If you need to think about a probability problem, draw a probability tree



# Solving genetic problems

## LEARNING OBJECTIVES

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At the end of this lecture you should be able to:

- List the tools available to solve genetic problems
- Think about which tools may be useful in certain circumstances
- Be prepared to work through genetic problems in week 3 and beyond