

# **FRST302: Forest Genetics**

## **Lecture 1.1: Classical Genetics and its Molecular Mechanisms**

# Outline for Today

- Short history of genetics
- Mendel's laws
- Chromosomes

**What is genetics?**

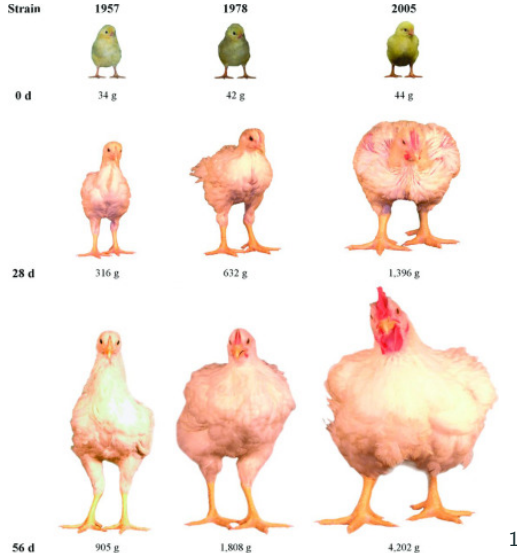
**What is genetics?**

**Genetics is the study of genes**, of variation and heredity across all branches of the tree of life

# *What are the major questions in genetics?*



# How can we apply a knowledge of genetics?



<sup>1</sup>Modified from Figure 1 - Zuidhof et al. 2014

# History of Genetics

Humans have probably pondered inheritance for all history:

- For much of history, the mechanisms of inheritance were basically unknown
- The inheritance of acquired characteristics was widely accepted for much of history (from Hippocrates to Aristotle to Lamarck)

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- The inheritance of acquired characteristics was widely accepted for much of history (from Hippocrates to Aristotle to Lamarck)
- *Early microscopists thought that they had seen small humans inhabiting sperm cells!*

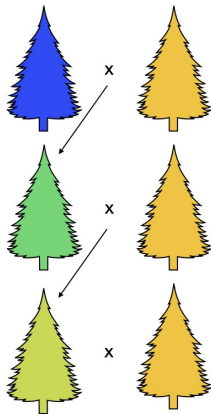




# History of Genetics

By the 19th Century, the dominant theory was **blending inheritance**

- The notion that an offspring's traits are simply the average of the parents' traits.
- This is intuitively appealing - continuously varying traits are often intermediate between their parents
- *There is one big problem with blending inheritance!*



*What's the big problem with  
blending inheritance?*

# Darwin's Thoughts on Inheritance

*“The laws governing inheritance are quite unknown; no one can say why the peculiarity in different individuals of the same species... is sometimes inherited and sometimes not so”<sup>a</sup>*

But, Darwin clearly appreciated the limitations of blending and felt the need for an alternative:

*“Each parent transmits its peculiarities, therefore if varieties allowed to cross... such varieties will be constantly demolished”<sup>b</sup>*

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<sup>a</sup>Ch. 1, *The Origin of Species*, C. Darwin 1859

<sup>b</sup>*Foundations of the 'Origin of Species'*, F. Darwin 1909

A NICE PIC OF  
DARWIN?

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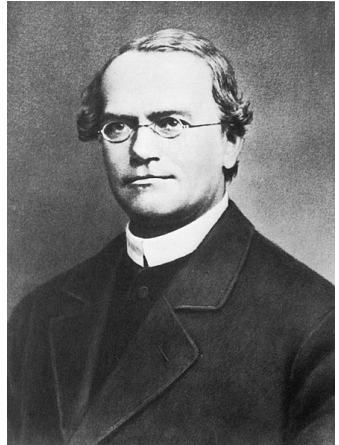
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But different modes of variation are common:

- Continuous - traits measured on a numerical scale (e.g. height, diameter, chlorophyll fluorescence)
- Discrete - traits that exhibit categorical differences (e.g. different leaf forms, distinct flower colour)
- Ordinal - discrete traits with some informative order (e.g. high, medium and low shade tolerance)

# Particulate Inheritance

Through careful experimentation analysing discrete traits in peas, Franciscan Friar Gregor Mendel found evidence supporting a model of particulate inheritance



Mmmmm...  
Peas Peas Peas Peas Peas



# Particulate Inheritance

**Particulate Inheritance:** traits are passed from parent to offspring via particles

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## Blending Inheritance

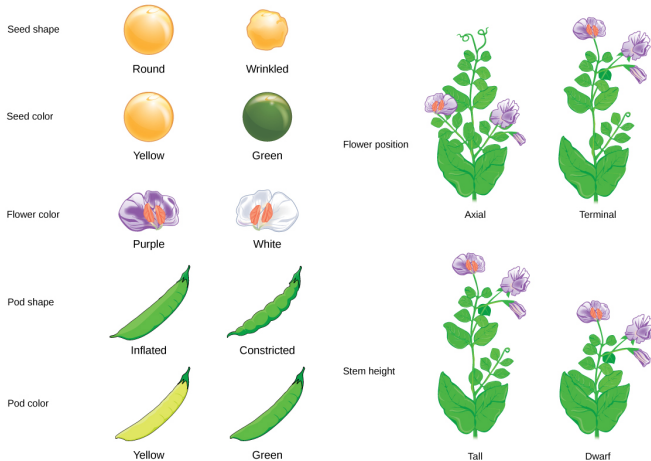
- Offspring exhibit averages of parental traits
- The "blended" traits are transmitted to offspring
- Variation is rapidly lost across generations

## Particulate Inheritance

- Offspring exhibit *combinations* of parental traits
- Parental traits can manifest in offspring (or skip generations)
- Variation is maintained over time

# Mendel's Crosses

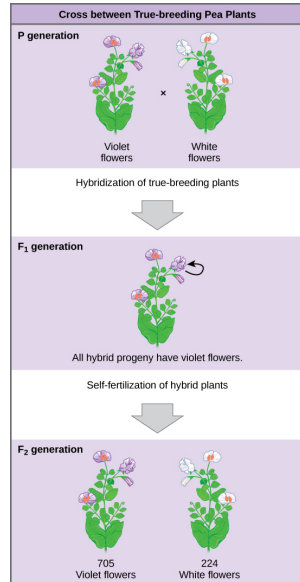
Mendel examined variation and inheritance of several discrete characteristics of pea plants



# Mendel's Crosses

Garden peas are capable of self-fertilization, so Mendel was able to generate "true" lines of peas that exhibit a particular trait/phenotype

- Crossing lines produces an F<sub>1</sub> generation
- The patterns of variation among the F<sub>2</sub> generations were Mendel's focus



# Mendel's Crosses



Note the 3:1 ratios of the two pea phenotypes in the F<sub>2</sub>

The patterns of variation that Mendel observed led him to develop three laws of inheritance

- Law of Segregation
- Law of Dominance
- Law of Independent Assortment

**The law of segregation:** each individual possesses a pair of particles for any particular trait and each parent passes one of these randomly to its offspring

**The law of dominance:** for some traits, the presence of one kind of particle masks the presence of another. Mendel referred to the **dominant** particle as masking the effects of the **recessive** particle

# Mendel's Laws

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**The law of independent assortment:** when two individuals differ in more than two pairs of traits (e.g. smooth v. wrinkly and green v. yellow), the inheritance of one pair of traits is independent of another



# Mendel's Laws



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each individual possesses a pair of particles for any particular trait and each parent passes one of these randomly to its offspring

How does the image demonstrate **the law of segregation**?

# Mendel's Laws



**The law of segregation:**  
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How does the image demonstrate **the law of segregation**?

Answer: *Individuals (i.e. seeds) in the F<sub>2</sub> generation exhibit a combination of seed colours and textures*

# Mendel's Laws



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# Mendel's Laws



**The law of dominance:**  
for some traits, the presence  
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How does the image  
demonstrate **the law of**  
**dominance**?

Answer: *The uniformity of  
trait values in the  $F_1$   
generation*

# Mendel's Laws



**The law of independent assortment:** when two individuals differ in more than two pairs of traits, the inheritance of one pair of traits is independent of another

How does the image demonstrate **the law of independent assortment**?

# Mendel's Laws



**The law of independent assortment:** when two individuals differ in more than two pairs of traits, the inheritance of one pair of traits is independent of another

How does the image demonstrate **the law of independent assortment**?

Answer: *The fact that wrinkly green peas and smooth yellow peas are seen in the F<sub>2</sub> generation*

# Mendel's Laws



I count 38 F<sub>2</sub> seeds

13 Green : 25 Yellow

9 Wrinkly : 29 Smooth

Why do we see these ratios?

# Mendelian Terminology

Remember, Mendel crossed "true" green (**G**) peas with "true" yellow (**Y**) peas.

The table below gives the results of the self-fertilization of the F1 generation

		<b>GY</b>	
		<b>G</b>	<b>Y</b>
<b>YG</b>	<b>G</b>	GG	YG
	<b>Y</b>	YG	YY



# Codominance



×



A leaf shape trait controlled by a single gene

Assuming the two individuals are homozygotes, how could you figure out if the allele for the cut leaf phenotype is dominant, recessive or codominant?

# Test Crosses

Leaf phenotypes in European beech, *Fagus sylvatica*



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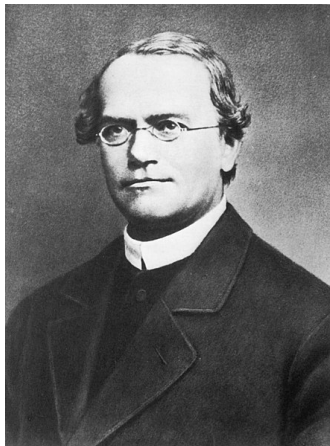


A leaf shape trait controlled by a single gene

Assuming the two individuals are homozygotes, how could you figure out if the allele for the cut leaf phenotype is dominant, recessive or codominant?

# Particulate Inheritance and Classical Genetics

- Proposed in 1865 and 1866
- 6-7 years after Darwin's Theory of Evolution
- As far as anyone knows, Darwin was totally unaware of Mendel *but see notes!*
- Represents the foundation of classical genetics
- **Classical genetics** refers to the study of genetic patterns observable from reproductive events

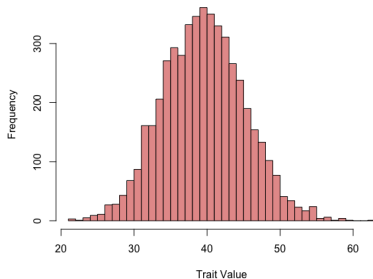


More peas please

# Reconciling the Mendelians and the Biometricians

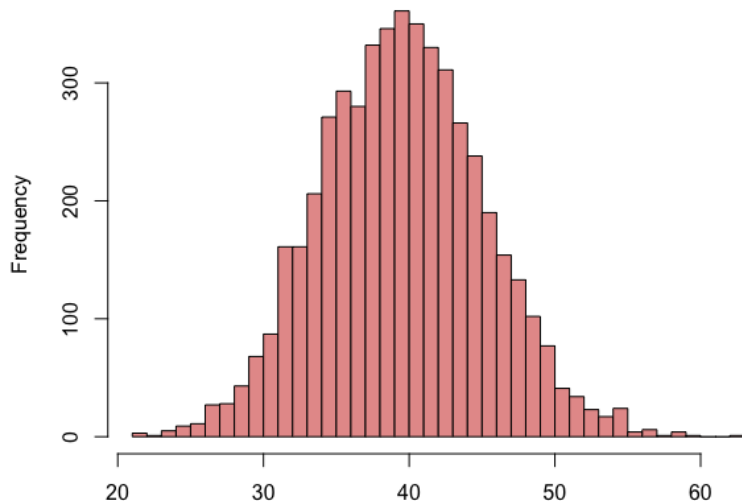
Mendel's results went largely unnoticed in his lifetime, but they were rediscovered and publicised in the early 1900s

**Distribution of Trait Values Assuming  
100 Genes Segregating According to Mendelian Inheritance**



# Reconciling the Mendelians and the Biometricians

**Distribution of Trait Values Assuming  
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# Branches of Genetics

- Behavioural genetics
- **Classical genetics**
- Developmental genetics
- **Conservation genetics**
- **Ecological genetics**
- **Evolutionary genetics**
- **Genecology**
- Genetic engineering
- Genomics
- Medical genetics
- Forensics
- Molecular genetics
- **Quantitative genetics**
- **Population genetics**
- Phylogenetics
- Statistical genetics
- Genetic epidemiology
- Archaeogenetics

**Below is the R code to make the figures on the infinitesimal model - feel free to play around with it**

```
# Demonstrate the distribution of trait values for a quantitative trait  
# Under Mendelian segregation for an arbitrary number of genes  
# Assumes random mating, constant effect sizes, constant allele frequencies  
  
nGenes = 100  
alleleFrequency = 0.2  
popSize = 5000  
effectSize = 1  
  
hist(  
  replicate(popSize ,  
            sum( 1 * rbinom(nGenes, 2, alleleFrequency) ) ),  
  col = "#e69b99",  
  xlab= " Trait - Value",  
  main= paste(" Distribution - of - Trait - Values - in - F2s - Assuming\n", nGenes ,  
              " Genes - Segregating - According - to - Mendelian - Inheritance"),  
  breaks = 40)
```