

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?

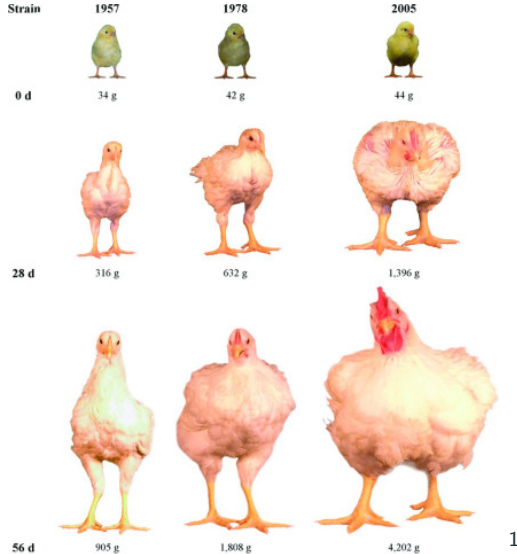
What is genetics?

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?



¹Modified from Figure 1 - Zuidhof et al. 2014

History of Genetics

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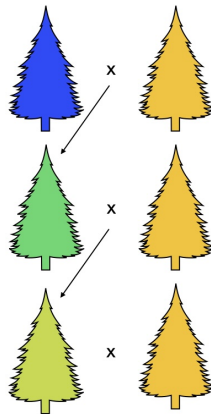
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- The inheritance of acquired characteristics was widely accepted for much of history (from Hippocrates to Aristotle to Lamarck)
- *Early microscopists thought that they had seen small humans inhabiting sperm cells!*



History of Genetics

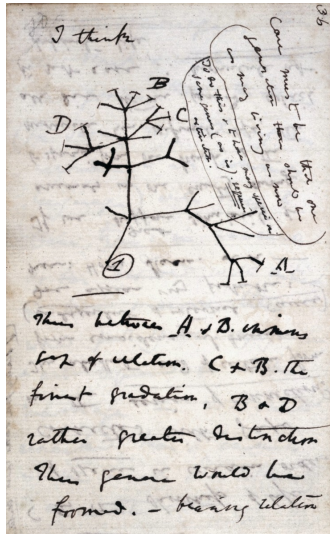
By the 19th Century, the dominant theory was **blending inheritance**

- 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!*

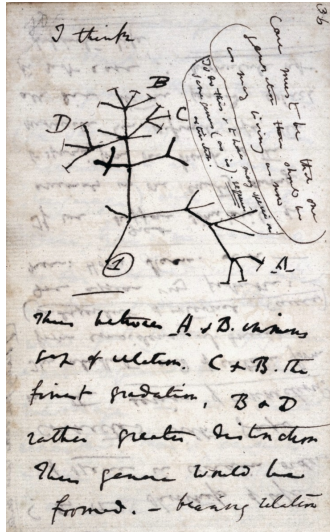


*What's the big problem with
blending inheritance?*

Darwin's Thoughts on Inheritance

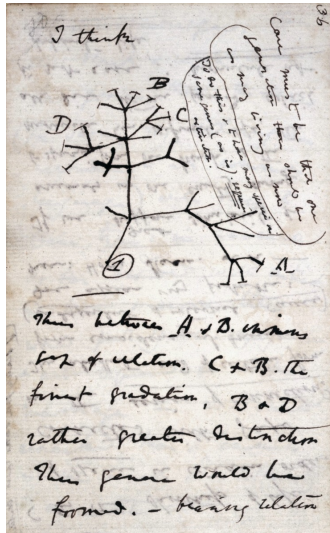


Darwin's Thoughts on Inheritance



“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” ^a

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" ^a

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" ^b

^aCh. 1, *The Origin of Species*, C. Darwin 1859

^b *Foundations of the 'Origin of Species'*, F. Darwin 1909

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- Discrete - traits that exhibit categorical differences (e.g. different leaf forms, distinct flower colour)

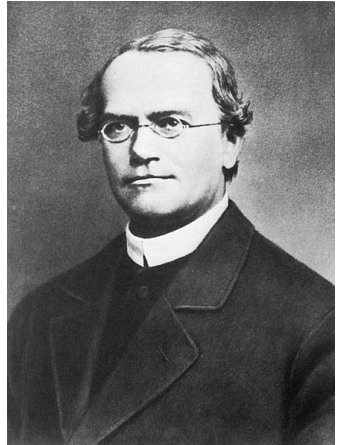
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But different modes of variation are common:

- 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

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Blending Inheritance

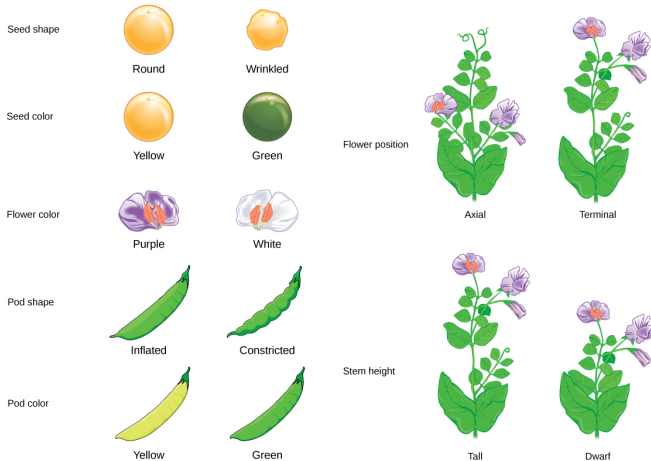
- Offspring exhibit averages of parental traits
- The "blended" traits are transmitted to offspring
- Variation is rapidly lost across generations

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

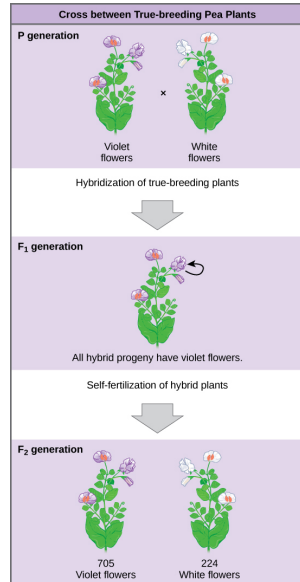
Mendel examined variation and inheritance of several discrete characteristics of pea plants



Mendel's Crosses

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

- Crossing lines produces an F₁ generation
- The patterns of variation among the F₂ generations were Mendel's focus



Mendel's Crosses



Note the 3:1 ratios of the two pea phenotypes in the F₂

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

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

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**?

Answer: *Individuals (i.e. seeds) in the F₂ generation exhibit a combination of seed colours and textures*

Mendel's Laws



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How does the image
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Answer: *The uniformity of
trait values in the F_1
generation*

Mendel's Laws



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

Mendel's Laws



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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 F₂ generation*

Mendel's Laws



I count 38 F₂ 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

		GY	
		G	Y
YG	G	GG	YG
	Y	YG	YY

*This table is an example of a Punnett square - Yellow phenotypes are shown in **bold***

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- Yellow is dominant to Green, so any offspring possessing a single Y particle will be yellow
- So $\frac{3}{4}$, (or 75% or 0.75)of the offspring are expected to be yellow
- With 38 F2 seeds, we would expect 9.5 seeds to be yellow, but **we would also expect variation around this number**

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- We use the term **homozygote** to refer to the offspring possessing the GG or the YY combinations
- We use the term **heterozygote** to refer to both YG and GY offspring as they are equivalent^a

^abut not always - see notes

Codominance

Codominance is a form of inheritance wherein both alleles in a heterozygote are equally expressed

As seen in this example from Sitka Spruce



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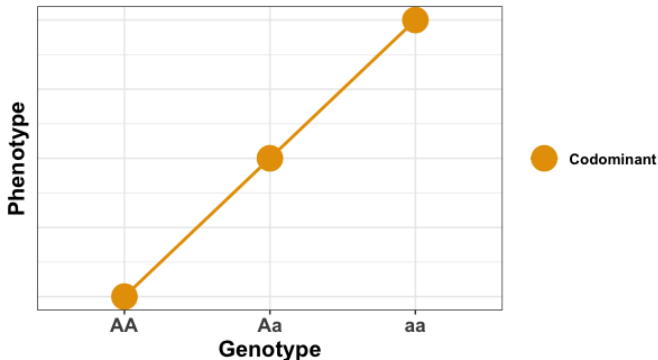
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Codominance

Codominant and full dominance are just two domains on a continuous range

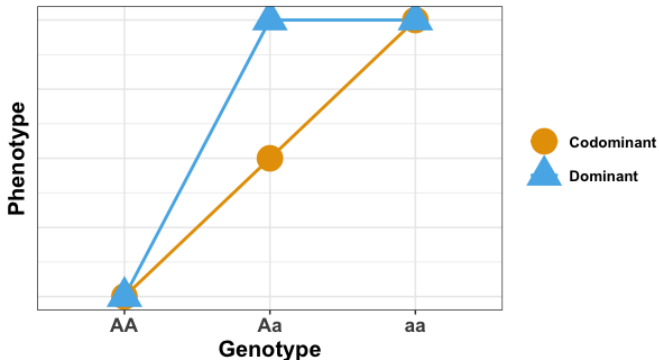
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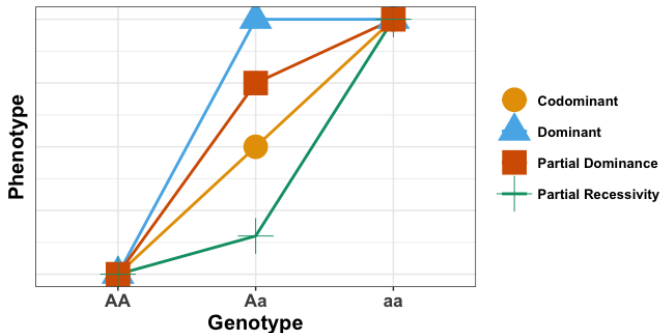
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Crossing Experiments

Leaf phenotypes in European beech, *Fagus sylvatica*



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

Test crosses are used to determine individual genotypes

In a test cross, individuals with unknown genotype (WW or Ww ?) are crossed with individuals homozygous for a recessive trait (ww)



×



If any of the offspring exhibit the recessive phenotype, the unknown parent must be... ?

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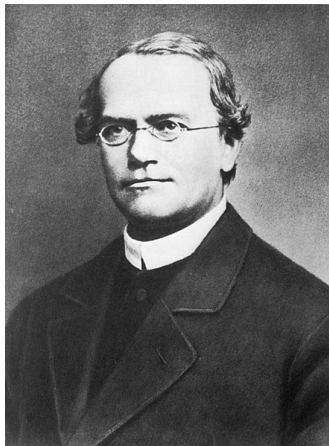
Questions?

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Let's take a short break

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

Mendel's contributions were underappreciated in his time

Mendel's findings began to be appreciated early in the 1900s, largely thanks to the work of William Bateson and Edith Rebecca Saunders

The rediscovery of Mendel's laws kicked off a scientific feud
The Biometricians v. The Mendelians

Reconciling the Mendelians and the Biometricians

Imagine a species of tree
with a codominant trait we
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In this species there is a single gene that controls height - with a pair of alleles (just like yellow v. smooth peas). The A allele does not affect height, but the a allele leads to a height of $+1$

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Pretend that we conducted a crossing experiment on these plants just like Mendel

What possible trait values could the F_2 generation would exhibit?

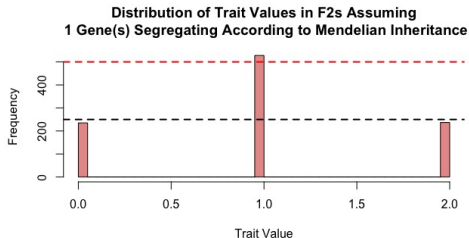
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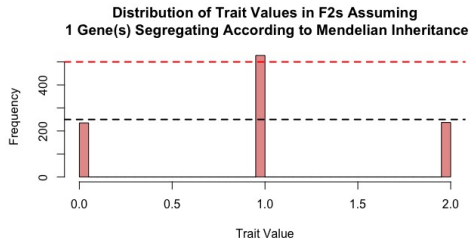


Reconciling the Mendelians and the Biometricians

Now let's say that our tree height trait is controlled by 100 genes, inherited according to the law of independent assortment

Instead of a +1 effect on height, each one has an effect of $+\frac{1}{100}$

What would the distribution of possible trait values look like now?

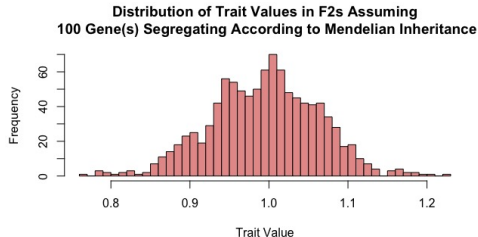
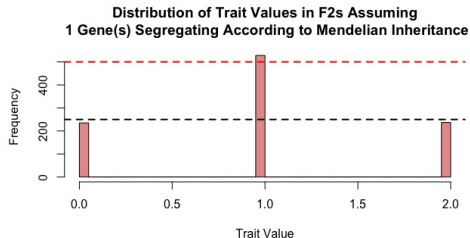


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Reconciling the Mendelians and the Biometricians

Assuming Mendel's laws and that a large number of genes controls complex traits can reconcile the Mendelian and Biometrician's arguments

This model was first formalized in 1918¹

If we take this model to the limit of an infinite number of genes, each making an infinitely small contribution to a particular trait we get **the infinitesimal model** - the basis of quantitative genetics (more on that in Module 3)

¹This idea, represents a major turning point in the history of genetics

The Units of Heredity

So a particulate theory of inheritance can explain discrete and continuous traits

But what are the particles??

The Units of Heredity

So a particulate theory of inheritance can explain discrete and continuous traits

But what are the particles??

It should be obvious that we are talking about chromosomes and DNA

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

A Question for Next Lecture...



Albinism is caused by a recessive lethal allele
Seedlings with albinism have no chlorophyll, thus no photosynthesis
How come tree populations still carry this allele?

- Basic definitions in genetics
- Basic principles and terms in classical genetics
- How Mendelian inheritance can lead to continuous trait variation
- Molecular mechanisms of Mendelian inheritance

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)
```