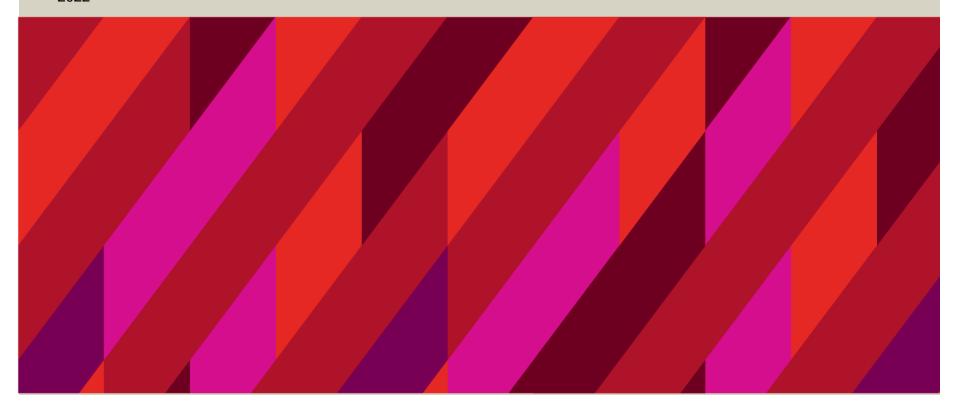


Quantitative Genetics (1)

LECTURE 5: HERITABILITY

2022



Background

Quantifying VG

-so far limited to single locus

H, A, ne, P,

- -Looking at genetics from a phenotypic level
- -Used to attribute causality to traits (nature vs. nurture)

Causal agents of phenotypes

- -Alleles segregating at many loci
- -Interactions among alleles (dominance and/or epistasis)
- -Environmental effects
- -Interactions & correlations between genetic and environmental effects

Lecture Outline

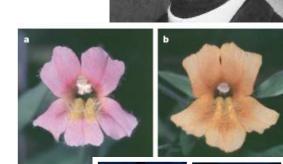
- Quantitative and Qualitative Traits
- History of Heritability
- Heritability (H²) and Genetic Causes
- Relationship with Conservation Biology
- Worked Example for H² (ANOVA)
- GxE Interaction
- Phenotypic Plasticity

'Mendelian loci'

- Blood Type (A, B, O, AB)
- Flower colour (yellow, pink)
- Thumb shape
- Coat colour (black, orange –X linked)
- Goat beard (beard, no beard)
- Phenotypes take discrete values
- Predictable using Mendel's laws

22,480 known Mendelian traits in man (as of 14 Aug 2014), OMIM: http://www.omim.org/

3,057 known Mendelian traits in animals, OMIA: http://omia.angis.org.au/home/









Quantitative genetics

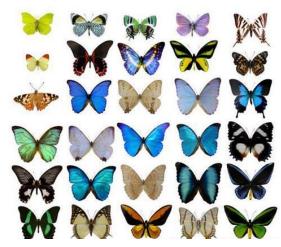
- The genetic appraisal and analysis of 'complex', 'continuous', or 'quantitative' traits, where:
- A quantitative trait = a trait whose phenotypic expression is determined by many individual genes (i.e. variation at many individual loci) (polygenic)





Examples of Quantitative Traits:

- Morphological traits:
 - Body size (height, weight)
 - Colouration (e.g. Bird plumage)
- Physiological traits
 - Enzyme activity
 - Blood pressure, cardiovascular performance
- Behavioural traits
 - Aggression, mate preferences, IQ etc.
- Life history traits
 - Developmental rates, age at maturity, etc.



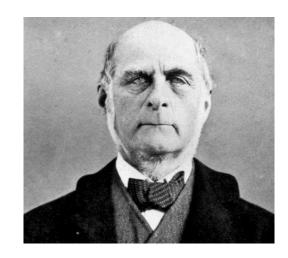






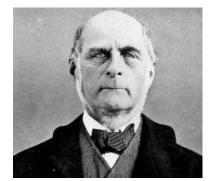
Francis Galton: 'the apostle of quantification'

- the weather (1863)
- height (1889)
- finger print patterns (1888; 1892; 1893)
- beauty (1909)
- boredom (1909)
- criminal characteristics (1885)
- the effectiveness of prayers (1872)

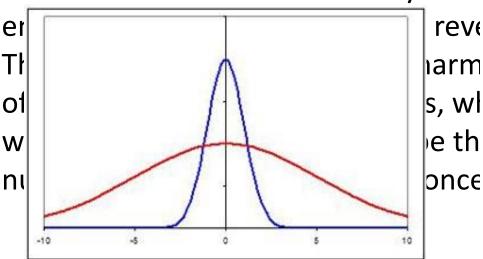


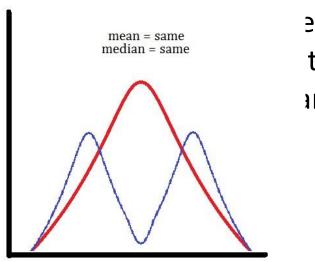
Introduced the terms 'nature' and 'nurture' (English Men of Science, 1874)

Francis Galton
Interested in *trait variation* rather than trait means



"It is difficult to understand why statisticians commonly limit their



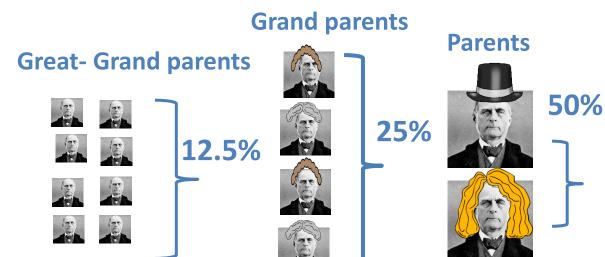


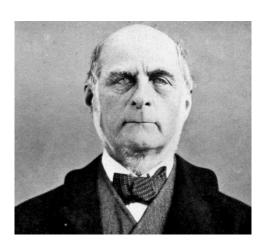
ews. tive

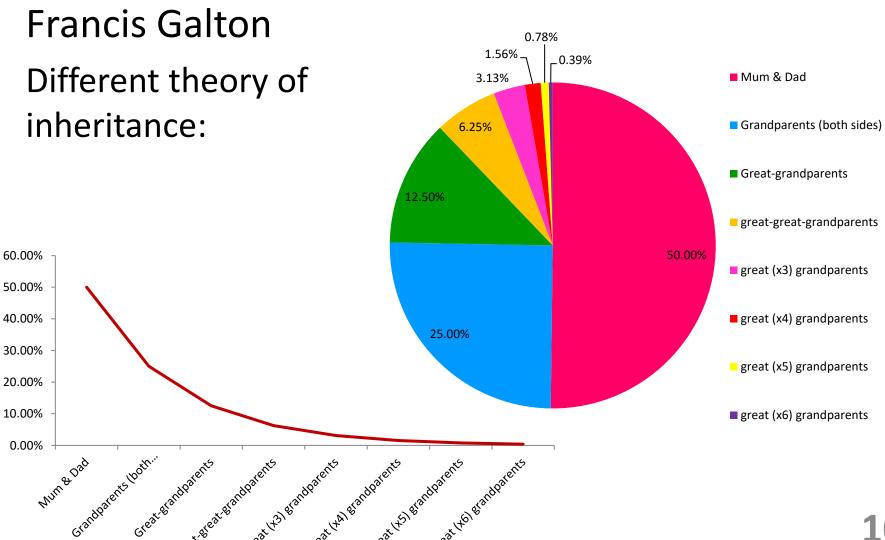
Francis Galton

Different theory of inheritance:

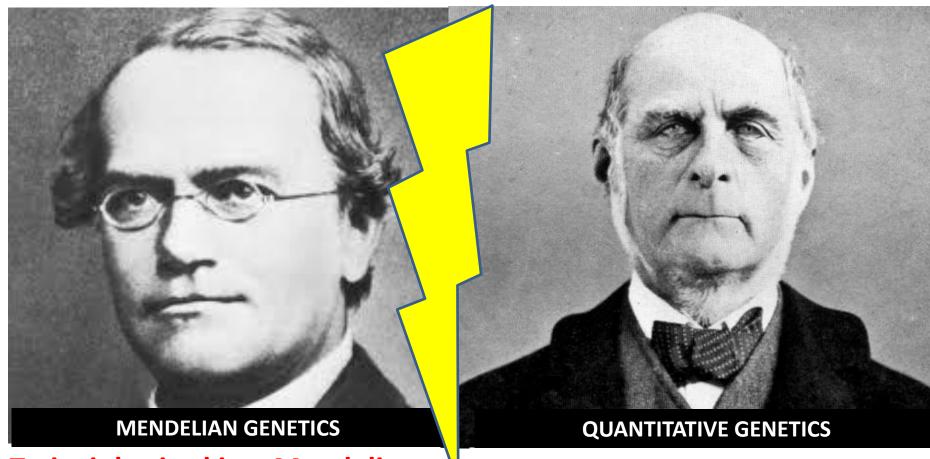
The parents of an individual together contribute on an average 50 percent of the total inherited characters, the 4 grandparents together 25 percent, the 3rd generation of ancestors together 12.5 percent, etc.







History: Galton vs. Mendel

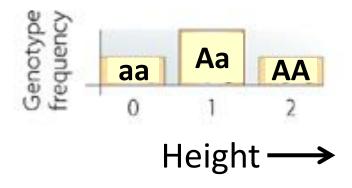


Traits inherited in a Mendelian fashion. Phenotypes discrete characteristics

Traits inherited as complex averages of ancestral phenotypes. Continuously varying phenotypes

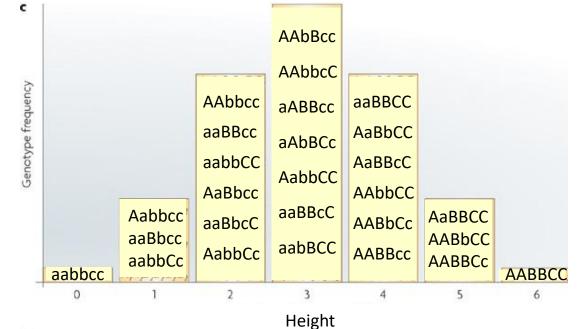
Quantitative traits and Mendelian laws

One locus(A):

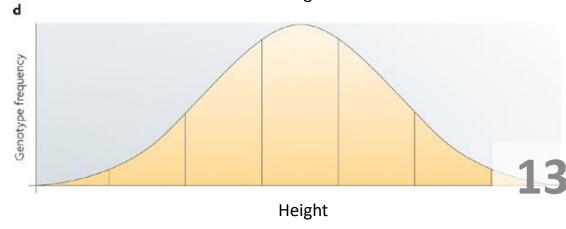


Quantitative traits and Mendelian laws

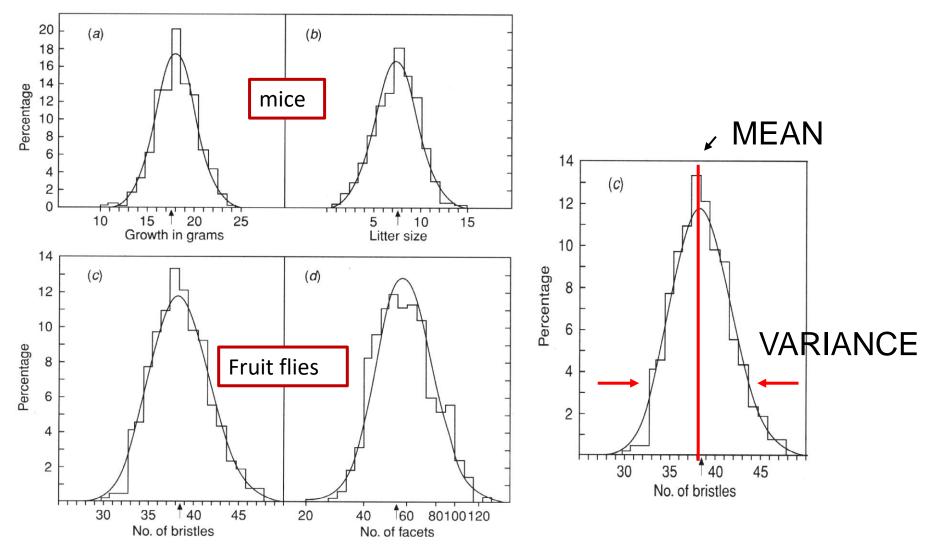
Three Loci (A, B & C):



Many loci:



Normal Distribution of Phenotypes



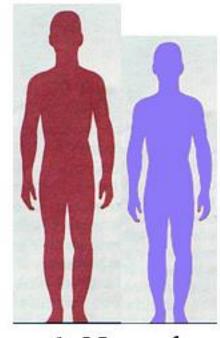
In: Introduction to Quantitative Genetics Falconer & Mackay 1996

Nature versus Nurture?



- Two influences on the expression of quantitative traits:
 - (1) Genetic (effects of many genes)
 - (2) Environmental (effects of environment)
- E.g. height:

Difference due to different genes



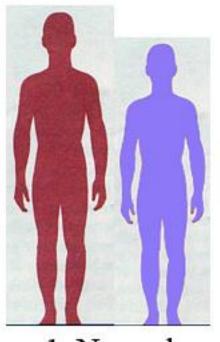
1. Normal development

Nature versus Nurture?

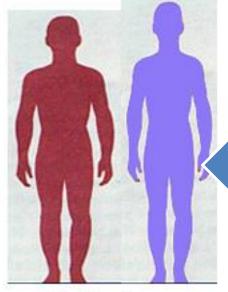
- Two influences on the expression of quantitative traits:
 - (1) Genetic (effects of many genes)
 - (2) Environmental (effects of environment)

• E.g. height:

Difference due to different genes



1. Normal development

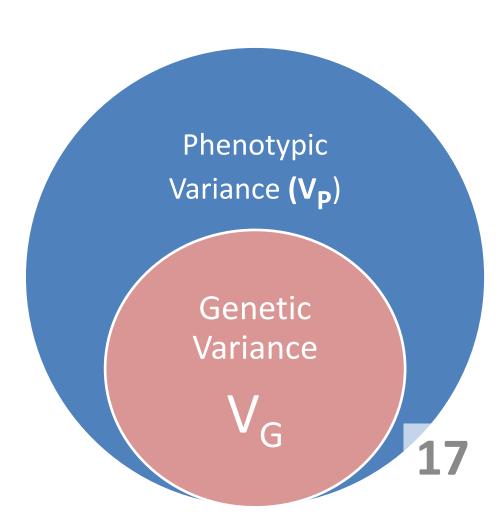


2. Red guy smoked!

Difference due to genes & env.

Partitioning trait variation:

- Most traits influenced by both genes and environment (nature & nurture)
- •If we represent all the **phenotypic variation** in the population as a solid circle, we can recognise the portion that is due to different genes
- This is called 'genetic variance'



Heritability (H²)

What Does it Measure?

Measure of how much *variation* in a phenotype is

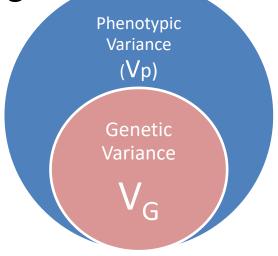
caused by variation in genotype

Assuming additivity:

VP = VG + VE

 $H^2 = VG/VP$

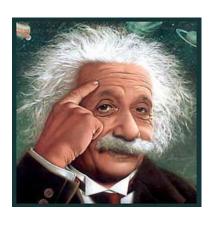




Proportion of Genetic variance to phenotypic variance

Heritability and Causation

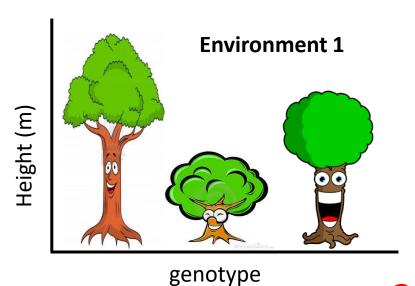
- Heritability can be used to make causal claims about the relative effects of genes (nature) and the environment (nurture)
- Which is more causally important?

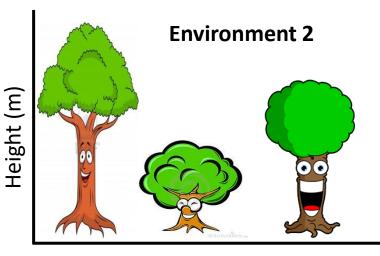






Specific types of causal claims

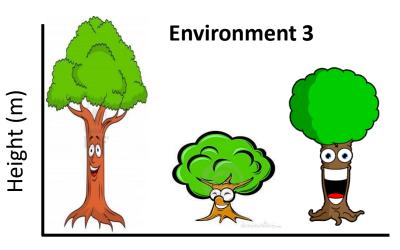




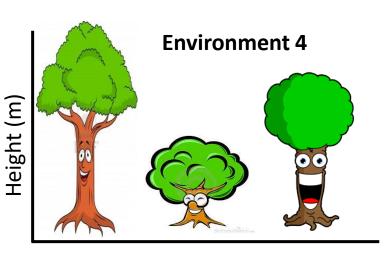
Потуре

genotype

H² HIGH

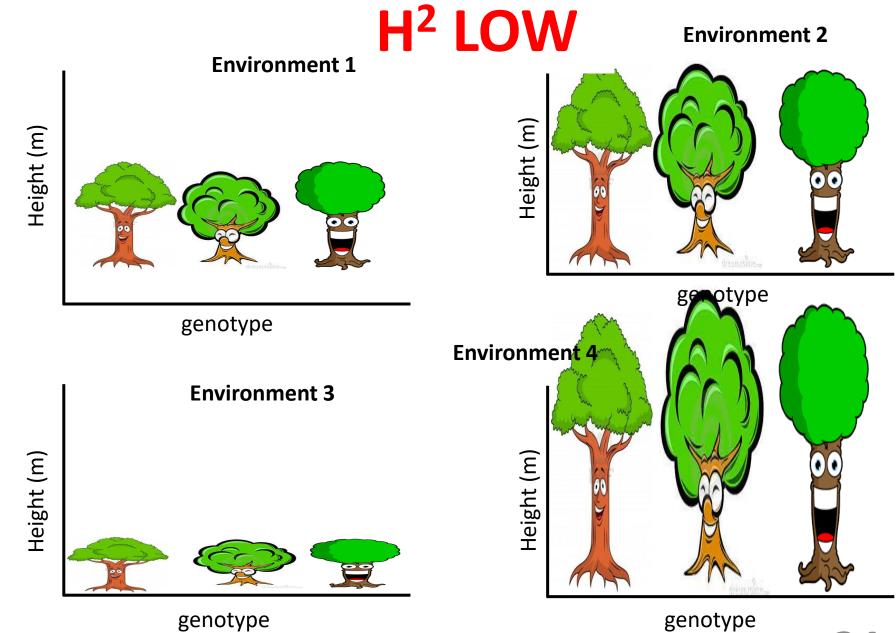


genotype

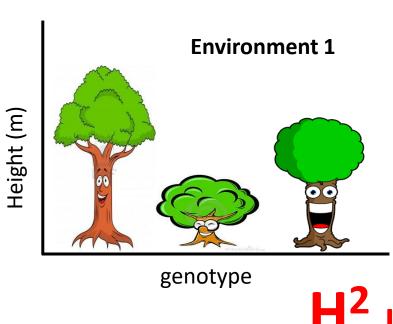


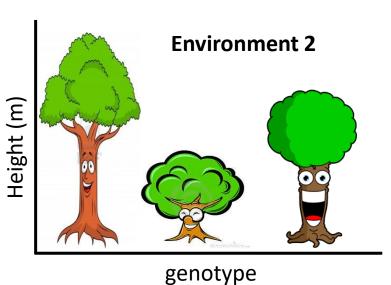
genotype

Environmental or Genetic?



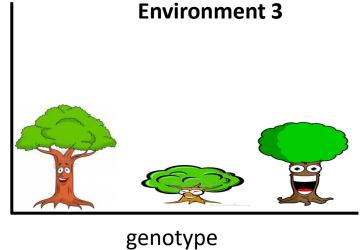
Environmental or Genetic?





H² INTERMEDIATE

Environment 4



Height (m)

Height (m)

genotype

Environmental or Genetic?

22

How does this relate to Conservation Biology?

Knowledge of quantitative fitness traits

- Clutch size
- Resistance to disease / introduced organisms
- Resistance to climate change
- Lifetime fecundity
- Age of first litter
- Days to flowering



Where do the values come from?

Heritability of postchallenge survival, an indicator of disease resistance, was estimated to be 0.35 ± 0.09 .

Silverstein, J. T., et al. (2009). Rainbow trout resistance to bacterial cold-water disease is moderately heritable and is not adversely correlated with growth. *Journal of animal science* 87(3), 860-867.

$$H^2 = VG / VP$$

$$VP = VG + VE$$

Broad-sense heritabilities ranged from 0.55 in the F2 population (Fargo, N.D.) to 0.95 in the F_{2:4} progenies at

León, A. J., Lee, M., & Andrade, F. H. (2001). Quantitative trait loci for growing degree days to flowering and photoperiod response in sunflower (Helianthus annuus L.). *Theoretical and Applied Genetics*, 102(4), 497-503.

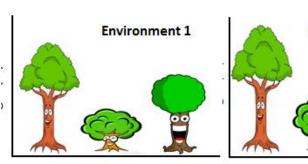
VG?

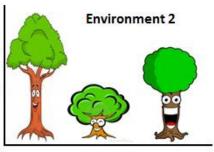
VE?

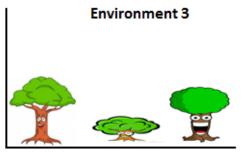
and the heritability estimates (95 % confidence intervals) for boldness and aggressiveness were <u>0.76</u> (0.49, 0.90) and <u>0.36</u> (0.10, 0.72) respectively. Furthermore, there were

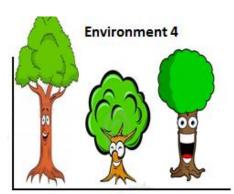
Ariyomo, T. O., Carter, M., & Watt, P. J. (2013). Heritability of Boldness and Aggressiveness in the Zebrafish. *Behavior genetics*, 43(2), 161-167.

Where do the values come from?









gen	oty	pe

genotype

genotype

genotype



	E1 (75)	E2 (75)	E3 (75)	E4 (75)
G1 (100)				
G2 (100)				
G3 (100)				

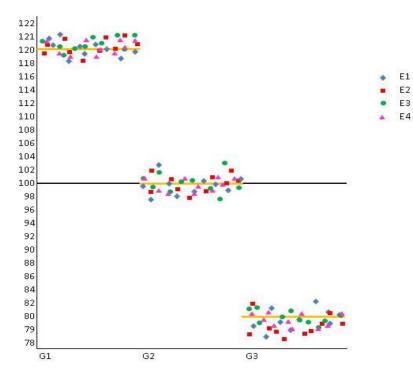
The height of the trees are measured...

	E1 (n _e =75)	$E2 (n_e = 75)$	E3 $(n_e = 75)$	E4 (n _e =75)	Genotype Means
G1 (n _g =100)	μ= 120	μ= 118	μ= 119	μ= 123	$\mu_{G1}=120$
$G2 (n_g=100)$	μ= 100	μ= 101	μ= 98	μ= 101	$\mu_{G2}=100$
$G3 (n_g=100)$	μ= 80	μ= 81	μ= 78	μ= 81	$\mu_{G3}=80$
Environment Means	$\mu_{E1} = 100$	$\mu_{E2} = 100$	$\mu_{E3} = 98.33$	$\mu_{E4} = 101.67$	X = 100

Trees with G1 genotype are significantly taller than G2, who in turn are taller than G3. Differences between environments are negligible.

	E1 (n _e =75)	$E2 (n_e = 75)$	E3 $(n_e = 75)$	E4 $(n_e = 75)$	N=300
G1 (n _g =100)	μ= 120	μ= 118	μ= 119	μ= 123	$\mu_{G1} = 120$
G2 (n _g =100)	μ= 100	μ= 101	μ= 98	μ= 101	$\mu_{G2} = 100$
G3 (n _g =100)	μ= 80	μ= 81	μ= 78	μ= 81	$\mu_{G3} = 80$
N=300	$\mu_{E1} = 100$	$\mu_{E2}=100$	$\mu_{E3} = 98.33$	$\mu_{E4} = 101.67$	X = 100

E3



$$VG = \frac{(\mu_{G1} - \ddot{X})^2 + (\mu_{G2} - \ddot{X})^2 + (\mu_{G3} - \ddot{X})^2}{n_G}$$

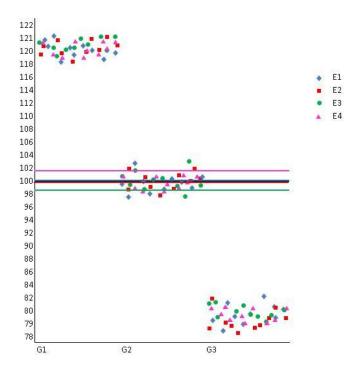
VG=
$$\frac{(\underline{120} - \underline{100})^2 + (\underline{100} - \underline{100})^2 + (\underline{80} - \underline{100})^2}{3}$$

$$VG = 266.67$$

	E1 (n _e =75)	$E2 (n_e = 75)$	E3 $(n_e = 75)$	E4 (n _e =75)	N=300
G1 (n _g =100)	μ= 120	μ= 118	μ= 119	μ= 123	$\mu_{G1}=120$
$G2 (n_g=100)$	μ= 100	μ= 101	μ= 98	μ= 101	$\mu_{G2} = 100$
G3 (n _g =100)	μ= 80	μ= 81	μ= 78	μ= 81	$\mu_{G2}=80$
N=300	$\mu_{E1} = 100$	$\mu_{E2}=100$	$\mu_{E3} = 98.33$	$\mu_{E4} = 101.67$	$\ddot{\mathbf{X}} = 100$

E1 E2

E3



$$VE = \begin{array}{c} (\mu_{E1} - \ddot{X})^2 + (\mu_{E2} - \ddot{X})^2 + (\mu_{E3} - \ddot{X})^{-2} + (\mu_{E4} - \ddot{X})^{-2} \\ \\ n_E \end{array}$$

VE=
$$\frac{(\underline{100} - \underline{100})^2 + (\underline{100} - \underline{100})^2 + (\underline{98.33} - \underline{100})^2 + (\underline{101.67} - \underline{100})^2}{4}$$

$$VE = 1.38$$

Getting the heritability estimate

The phenotypic variance is assumed to be additive

$$VP = VG + VE$$

Heritability (H²) is the proportion of the genotypic variance from the total phenotypic variance

$$H^2 = VG / VP$$
 or $VG / (VE + VG)$

Usually a high heritability estimate accurately reflects an example in which differences in genotype are the major cause of differences in phenotype

VG = 266.67

VE = 1.38

VP = 268.05

Genetic var (H²)

= VG / VP

= 266.67/268.05

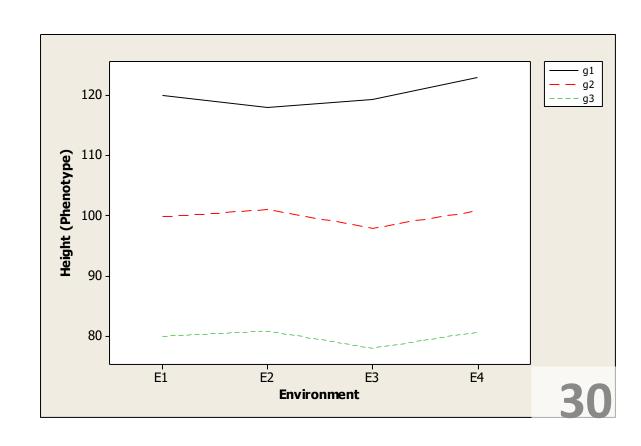
= 0.9948

Environmental var

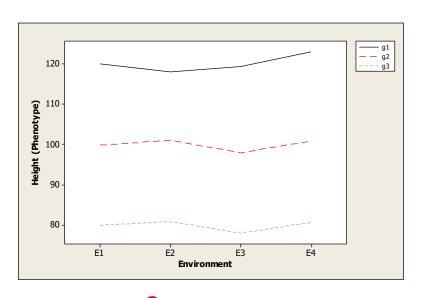
= VE / VP

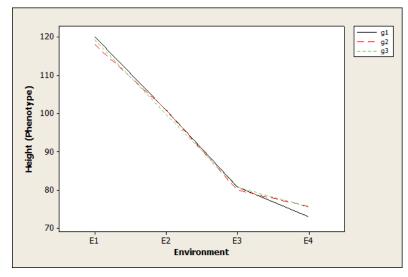
=1.38/268.05

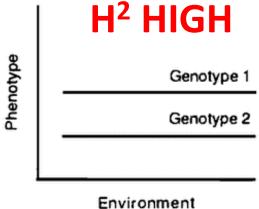
=0.0052

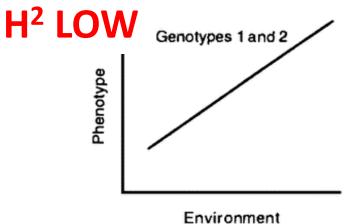


Norm of Reaction





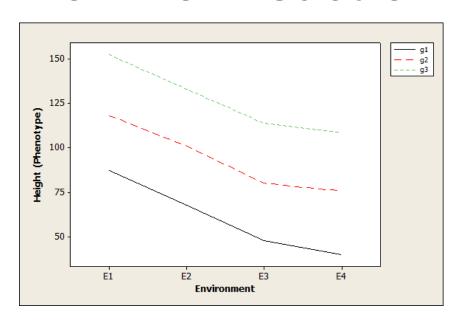




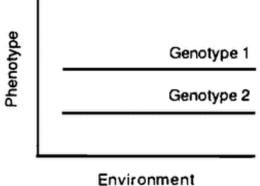
a) Biological determinism

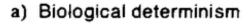
b) Social determinism

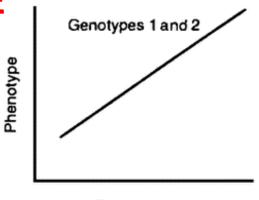
Norm of Reaction







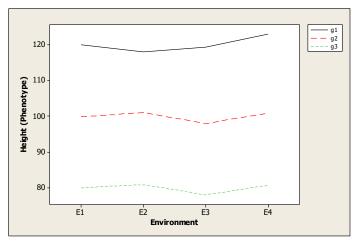




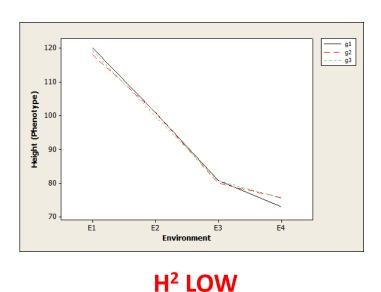
Environment

b) Social determinism

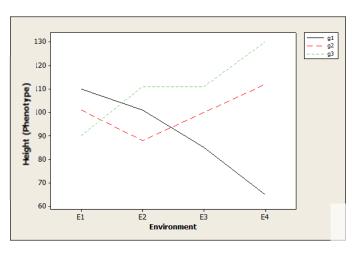
Norm of Reaction - GxE (Interaction)



H² HIGH



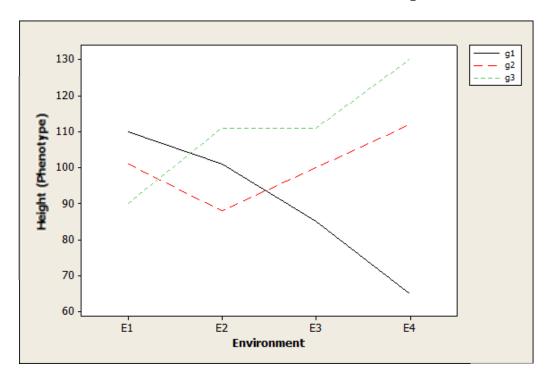
H² INTERMEDIATE



33

H² ??

$H^2 = 0.968$ (VERY HIGH!)



$$VE = 8.8089$$

$$VP = 275.4789$$

$$H^2 = 0.968$$

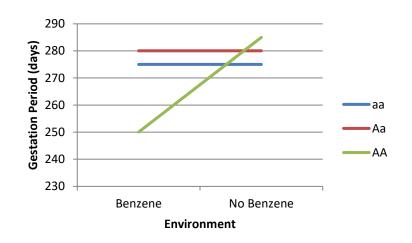
	E1	E2	E3	E4	Means
G1	110	100	85	65	90
G2	100	88	100	112	100
G3	90	110	110	130	110
Means					X =
	100	99.33333	98.33333	102.3333	100

$$VG = \frac{(\mu_{G1} - \ddot{X})^2 + (\mu_{G2} - \ddot{X})^2 + (\mu_{G3} - \ddot{X})^2}{vG} = \frac{n_G}{n_G}$$

$$VE = \frac{(\mu_{E1} - \ddot{X})^2 + (\mu_{E2} - \ddot{X})^2 + (\mu_{E3} - \ddot{X})^2 + (\mu_{E4} - \ddot{X})^2}{n_E}$$

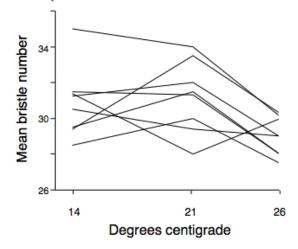
$$VE = 8.8089$$

- Trait expression varies over different environments may not be directional
- Genome produces different phenotypes depending on developmental environment
- Example 1: benzene exposure is significantly associated with shorter gestation periods in pregnant women possessing the AA allele of CYP1A1, while there is no association for those with Aa or aa (Wang et al. 2000)





- Trait expression varies over different environments may not be directional
- Genome produces different phenotypes depending on developmental environment
- Example 2: Bristle number in drosophila is determined by genotypic differences, and temperature. How temperature affects bristle number depends on the flies genetic background (Gupta and Lewontin 1981)





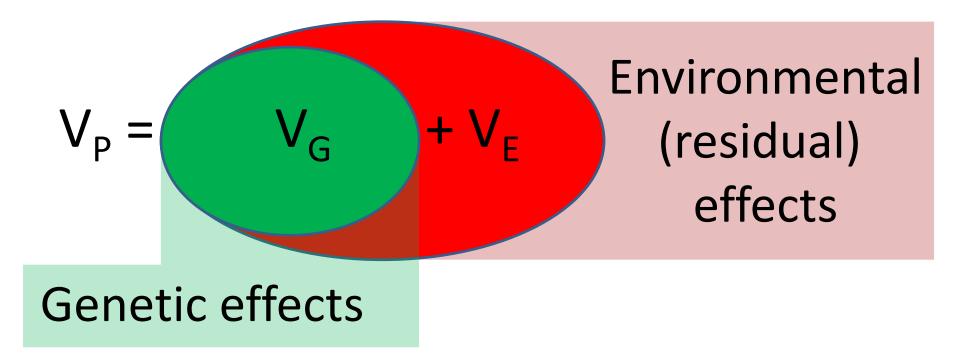
- GxE doesn't always mean an high heritability
- Could also result in low H² if not accounted for



- Demonstrates when interaction occurs and is unaccounted for – one cannot get a good understanding without looking at the NOR.
- GxE can lead to a heritability estimate that does not accurately reflect the contribution of environmental or genetic variation
- Can add GxE term into heritability model so it no longer skews results

Non-Additive

NATURE... versus ... NURTURE

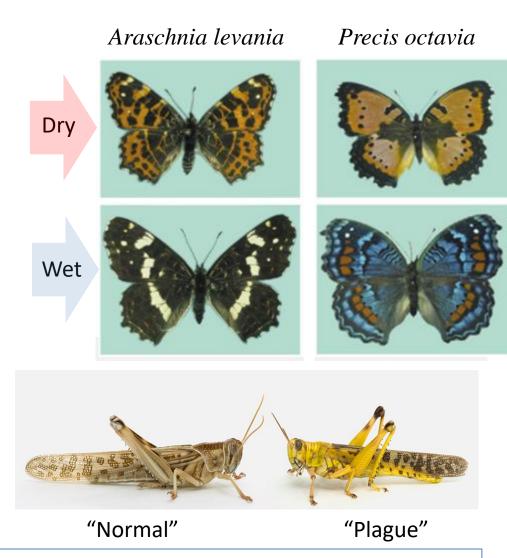


....assuming that genetic and environmental effects are additive

Biol334 Lecture 8 Slide 38

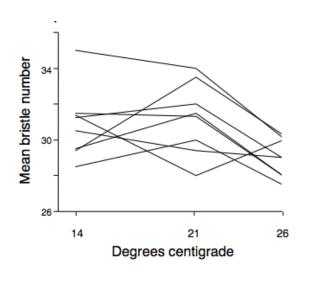
Phenotypic plasticity

The ability of a single genome to produce different phenotypes depending upon the environment experienced during development



Slide 39

GxE = A form of phenotypic plasticity
Single genotype produces multiple phenotypes,
depending on environmental interaction.





Summary

- Quantitative traits are characterised by heritability estimates (H²)
- Estimates partition the relative effects of environmental variation (VE) and genetic variation (VG) on differences in a phenotype (VP) within a population.

$$VP = VG + VE$$

 $H^2 = VG / VP$

- Can estimate the influence of genes just by studying the phenotype (no molecular techniques needed)
- Conservation biologists are interested in the genetic basis of fitness traits
- Limitations to estimates: non-additivity (GxE interaction)

$$VP = VG + VE + V_{GXE}$$

 $H^2 = VG / VP$