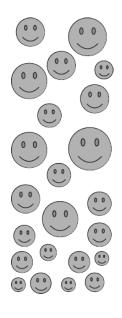
Quantitative genetics

Measuring heritable variation

- Natural selection relies on heritable variation
- Quantitative genetics focuses on variation in phenotypes
- ► Not looking at specific genes

Measuring heritable variation



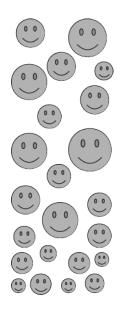
Components of phenotypic variation,

$$V_P = V_G + V_E$$
.

- $ightharpoonup V_P = \mathsf{phenotypic} \ \mathsf{variance}$
- $ightharpoonup V_G = ext{genetic variance}$
- $ightharpoonup V_E=$ environmental variance

Total phenotypic variance is the sum of genetic variance and environmental variance.

Measuring heritable variation



Components of phenotypic variation,

$$V_P = V_G + V_E$$
.

We can break down V_G even further,

$$V_G = V_A + V_D + V_I.$$

- $ightharpoonup V_{\mathcal{A}}=\mathsf{additive}\;\mathsf{genetic}\;\mathsf{variance}$
- $ightharpoonup V_G = ext{dominance variance}$
- $ightharpoonup V_E = ext{epistatic variance}$

Additive genetic variation



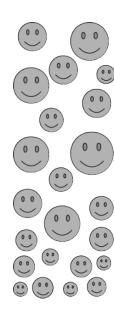
Additive genetic variation (V_A) is the variation in phenotype attributable to the summed effects of individual alleles.

- Summed effects of individual alleles are independent of other alleles (hence 'additive').
- Natural selection can only see additive variation because dominance and epistasis break down from parent to offspring.



The heritability of a phenotype determines the relative importance of heredity in determining the phenotypic value of a trait:

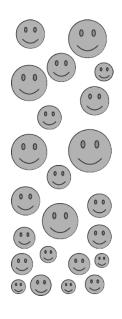
Heritability measures the proportion of the total phenotypic variation in a trait that is attributable to genetic as opposed to environmental effects.



The heritability of a phenotype determines the relative importance of heredity in determining the phenotypic value of a trait:

Heritability measures the proportion of the total phenotypic variation in a trait that is attributable to genetic as opposed to environmental effects. There are 2 types of heritability:

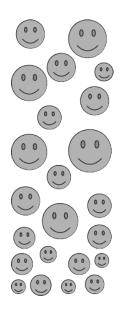
- Broad sense heritability: easier to measure, less informative
 - ▶ Narrow sense heritability: harder to measure, more informative



Broad sense heritability

$$H^2 = \frac{V_G}{V_P}$$

Total genetic variation divided by total phenotypic variation (must be between 0 and 1).



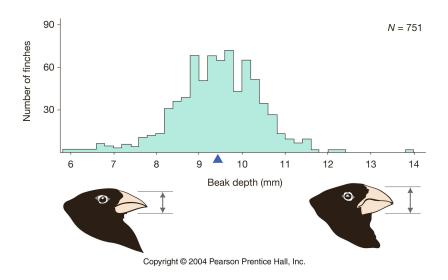
Narrow sense heritability

$$h^2 = \frac{V_A}{V_P}$$

Estimated 2 ways:

- ► Resemblance between relatives
- Response of a population to selection

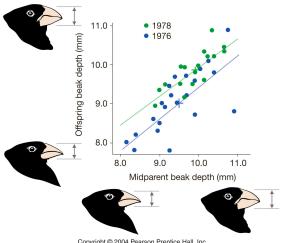
Postulate 1: Individual variation in beak depth



¹Image: Freeman, S., & Herron, J. C. (2007). Evolutionary analysis (Vol. 834). Upper Saddle River, NJ: Pearson Prentice Hall. Page 82.

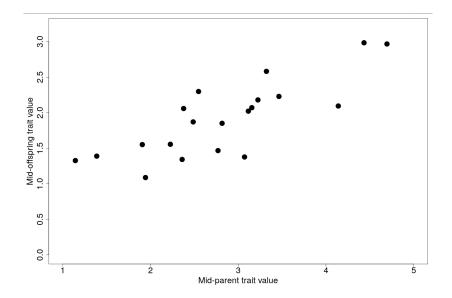
Postulate 2: Beak depth is heritable

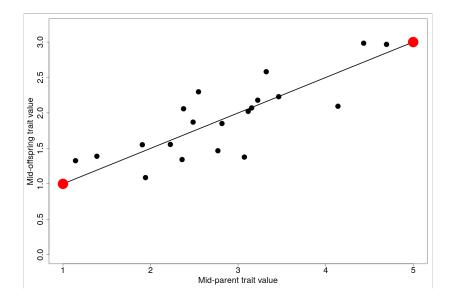
Parents with deeper beaks tend to have offspring with deeper beaks, and vice versa.



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¹Image: Freeman, S., & Herron, J. C. (2007). Evolutionary analysis (Vol. 834). Upper Saddle River, NJ: Pearson Prentice Hall. Page 83.





Narrow sense heritability can be estimated from the slope of the mid-parent to offspring regression line,

$$h^2 = \frac{rise}{run} = \frac{y_2 - y_1}{x_2 - x_1}.$$

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For the previous example,

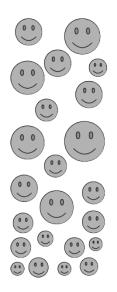
$$h^2 = \frac{rise}{run} = \frac{3-1}{5-1} = \frac{2}{4} = 0.5.$$

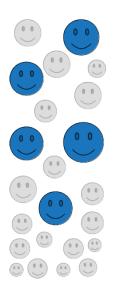
Remember $h^2 = V_A/V_P$, so we can also find, $V_A = h^2 \times V_P$.

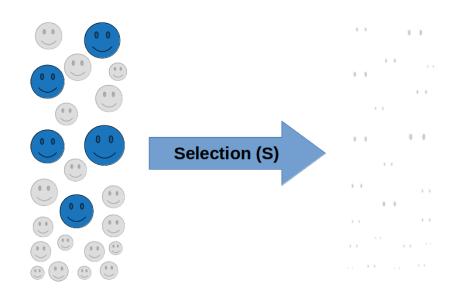
Phenotypic response (R) of a trait to selection (S) in a generation depends on h^2 ,

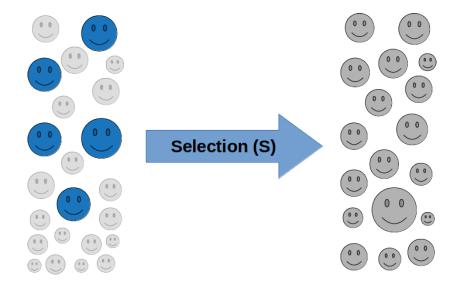
$$h^2 \times S = R$$
.

Given $h^2 = 0.5$, if we select for an increase of 2 in mean parent trait, we will get an actual change of 1 in offspring.









Before selection

- $ightharpoonup ar{P}$: Mean phenotype of population
- P*: Mean phenotype of selected parents
- ► $S = P^* \bar{P}$: Selection differential

$$h^2 = \frac{R}{S} = \frac{O^* - \bar{O}}{P^* - \bar{P}}$$

After selection

- O*: Mean phenotype for offspring of selected parents
- $ightharpoonup R = O^* \bar{O}$: Selection differential

Before selection

- $ightharpoonup \bar{P}$: Mean phenotype of population
- P*: Mean phenotype of selected parents
- ► $S = P^* \bar{P}$: Selection differential

After selection

- $ightharpoonup ar{O}$: Mean phenotype for offspring of all population
- O*: Mean phenotype for offspring of selected parents
- $ightharpoonup R = O^* \bar{O}$: Selection differential

$$h^2 = \frac{R}{S} = \frac{O^* - \bar{O}}{P^* - \bar{P}}$$

Rearrange the above to get the Breeder's equation:

$$R = h^2 S$$

Example of heritability in response to selection

Suppose you are a plant breeder attempting to increase flower width in a horticultural species.

- ▶ Measure mean corolla width of population to be 2 cm
- ▶ Select parents with a mean corolla width of 2.8 cm
- ▶ Next generation, find mean corolla width is 2.2 cm

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Note that we can assume $\bar{O} = \bar{P}$.

Predicting evolutionary change

- Can also look at multiple traits simultaneously (multivariate)
- Breeder's equation is foundational for food security
 - ► Increasing crop¹ & agricultural² yield
 - ► Increasing crop drought resistance³
- Sustainable Development Goal (SDG) 2.

¹Cobb, J. N., et al. (2019). Enhancing the rate of genetic gain in public-sector plant breeding programs: lessons from the breeder's equation. *Theoretical and applied genetics* 132:627-645.

²Hill, W. G., & Kirkpatrick, M. (2010). What animal breeding has taught us about evolution. Annual review of ecology, evolution, and systematics, 41:1-19.

³Cooper, M., & Messina, C. D. (2023). Breeding crops for drought-affected environments and improved climate resilience. *The Plant Cell*, 35:162-186.