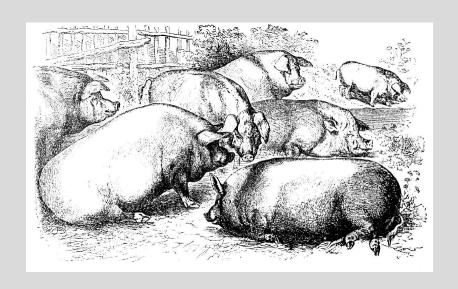
1.3 Inheritance of a Single Trait & Response to Selection



Stevan J. Arnold
Department of Integrative Biology
Oregon State University

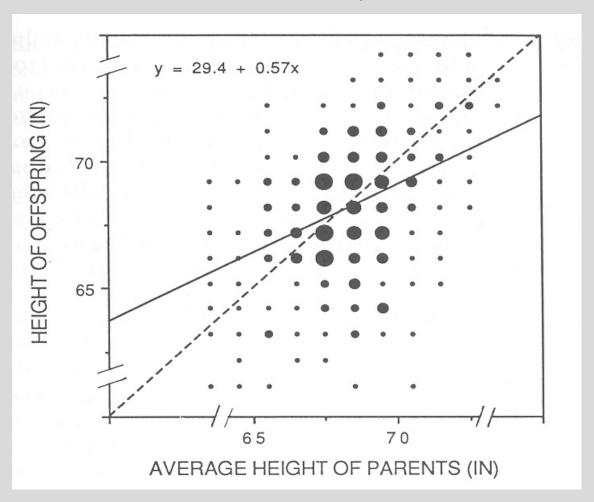
Thesis

- Most traits are affected by many genes
- We can model the inheritance of such traits with a statistical approach
- Additive genetic variance is the statistical concept that enables us to model both inheritance and response to selection

Outline

- 1. Phenotypic resemblance between parents and offspring reveals heritable variation.
- 2. To understand resemblance we need a model.
- 3. Some examples.
- 4. Why don't we run out of additive genetic variance?
- 5. Changing the trait mean with selection.

1. Phenotypic resemblance: Galton's plot



2. A Model of Phenotypic Resemblance

a. A model for phenotypic value (from Joe's lecture this morning)

$$P = \mu + {AA - 2 \choose Aa \ 0 \atop aa \ 3} + {BB \ 0.6 \atop Bb \ 0.1 \atop bb \ -0.2} + {CC \ -1 \choose Cc \ 6 \atop cc \ 6} + {DD \ 0.3 \atop Dd \ 0.3 \atop dd \ 0.7} + {EE \ -0.4 \atop Ee \ 0.3 \atop ee \ -0.3} + {environmental \atop effect}$$

b. An equivalent model, without the locus-by-locus details, for

phenotypic value z = x + e

$$z = x + e$$

phenotypic mean $\overline{z} = \overline{x} + \overline{e}$

$$\overline{z} = \overline{x} + \overline{e}$$

phenotypic variance

$$P = G + E$$

2. A Model of Phenotypic Resemblance

c. A new perspective on Galton's regression

$$h^2 = Cov(z_o, z_p)P^{-1} = Cov(x_o, x_p)P^{-1} = GP^{-1} = G/P$$

Our model identifies additive genetic variance, G, as the key statistical property responsible for Galton's regression

2. A Model of Phenotypic Resemblance

d. What about resemblance between other kinds of relatives?

We need to include domiance, d, and epistasis, i, as well as additive effects, x

$$z = x + d + i + e$$

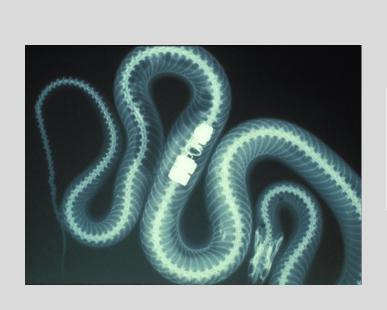
$$i = i_{A\!A} + i_{A\!D} + i_{D\!D} + i_{A\!A\!A} + i_{A\!A\!D} + i_{A\!D\!D} + i_{D\!D\!D} + \dots$$

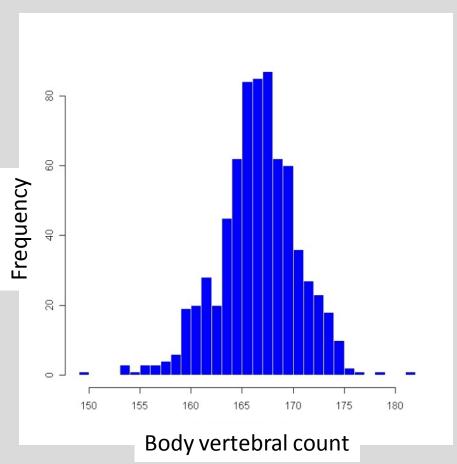
to obtain a general expression for resemblance between relatives X and Y

$$Cov(X,Y) = rG + uG_D + r^2G_{AA} + ruG_{AD} + u^2G_{DD} + r^3G_{AAA} + r^2uG_{AAD} + ru^2G_{ADD} + u^3G_{DDD} + \dots (5.6)$$

3. Some examples

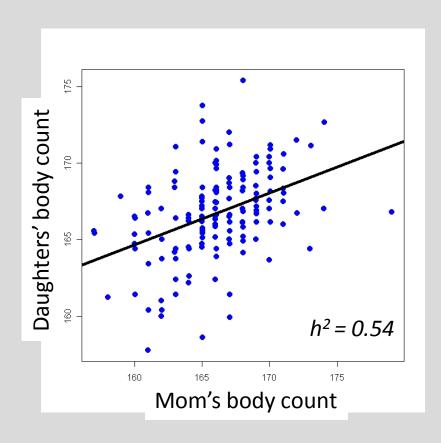
a. Mother-daughter resemblance in vertebral counts in garter snakes

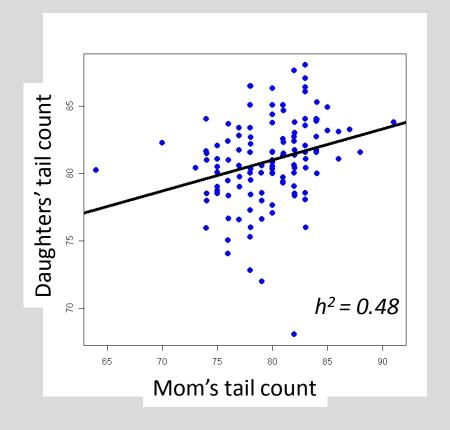




3. Some examples

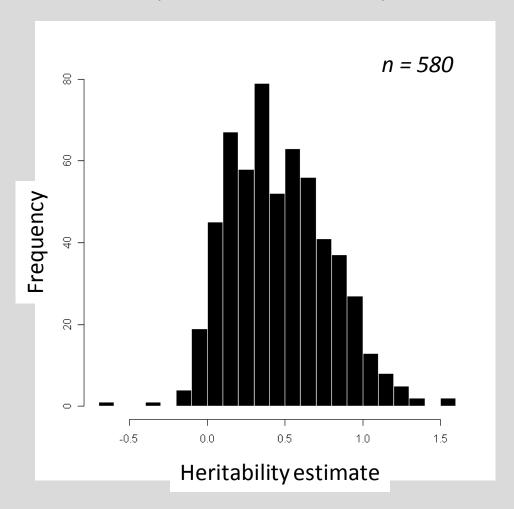
a. Mother-daughter resemblance in vertebral counts in garter snakes





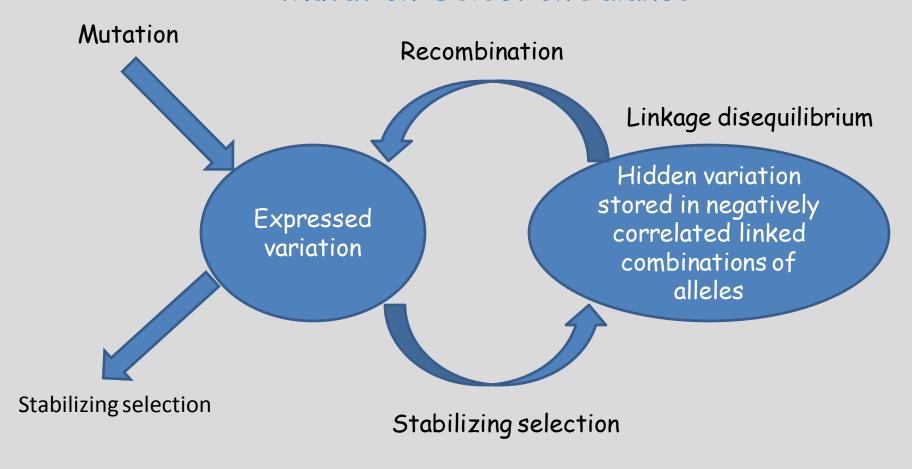
3. Some examples

b. A survey of heritability estimates



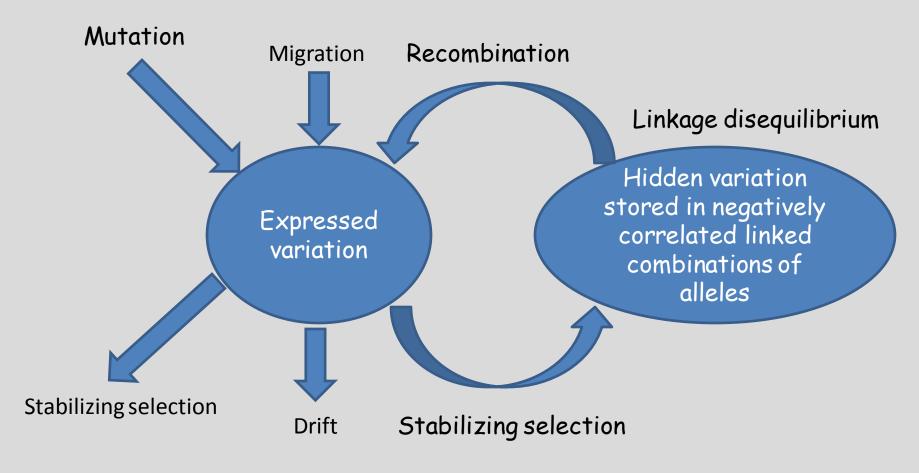
4. Why don't we run out of additive genetic variance?

Mutation-Selection Balance

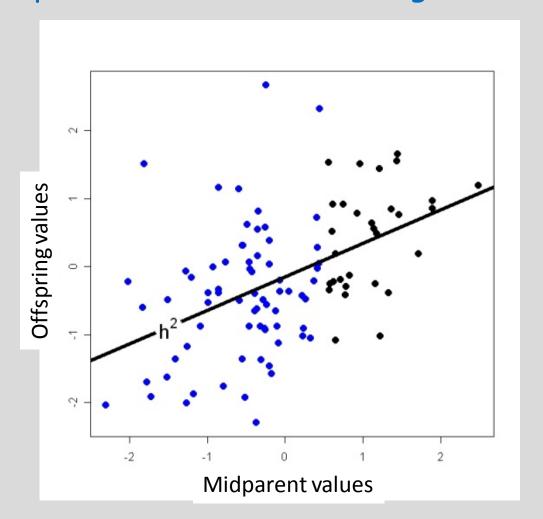


4. Why don't we run out of additive genetic variance?

Mutation-Migration-Selection-Drift Balance

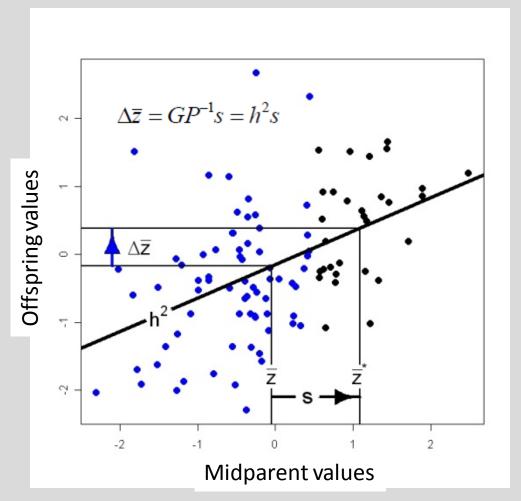


a. Response to selection as a regression problem



Heritability = h^2

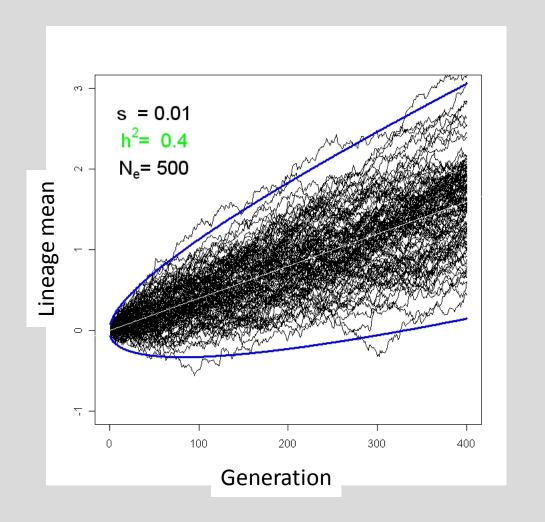
a. Response to selection as a regression problem



Selection differential

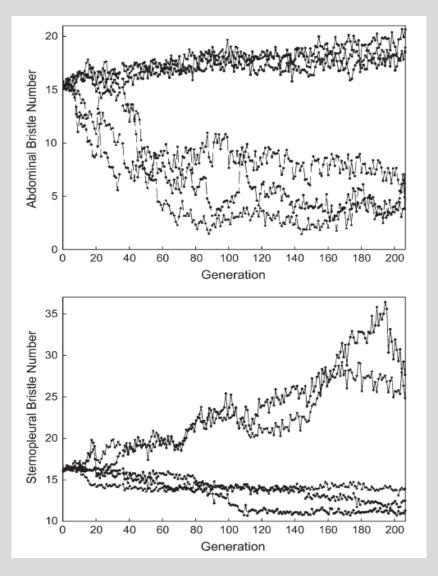
$$s = \bar{z}^* - \bar{z}$$

b. Response to selection in a finite population



Animation

c. Response to long term selection for bristle numbers in Drosophila



What have we learned?

- 1. Additive genetic variance, G, is the key to understanding resemblance between parents and offspring.
- 2. Consequently, G is also the key to modeling response to selection.
- 3. G is nibbled away by selection but restored by mutation (and migration).

References

- Galton, F. 1889. Natural Inheritance. MacMillan, New York.
- Lande, R. 1979. Quantitative genetic analysis of multivariate evolution, applied to brain: body size allometry. Evolution 33: 402-416.
- Cockerham, C. 1954. An extension of the concept of partitioning hereditary variance for analysis of covariances among relatives when epistasis is present. Genetics 39: 859-882.
- Arnold, S. J. and P. C. Phillips. 1999. Hierarchial comparison of genetic variance-covariance matrices. II. Coastal-inland divergence in the garter snake, *Thamnophis elegans*. Evolution 53:1516-1527.
- Mousseau, T. A. & D. A. Roff. 1987. Natural selection and the heritability of fitness components. Heredity 59: 181-197.
- Lande, R. 1976. The maintenance of genetic variation by mutation in a polygenic character with linked loci. Genetical Research Cambridge 26: 221-234.
- Mackay, T. F. C., R. F. Lyman & F. Lawrence. 2005. Polygenic mutation in *Drosophila melanogaster*: mapping spontaneous mutations affecting sensory bristle number.
 Genetics 170: 1723-1735