Introduction to Multivariate Quantitative Genetics

Number of Individuals

Height

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Topics we will cover:

- Multivariate quantitative genetics
- 1. Pleiotropy & Genetic correlations
- 2. The G matrix
- 3. Genetic constraints

Selection

- 1. Empirical methods to estimate selection
- 2. Empirical results

Selection

• genetic constraints and evolutionary lines of least resistance are only relevant in the context of selection The effect of the G-matrix on the response to selection Showing the trajectory in the absence of genetics correlation

how can we estimate selection?



Value for trait 1

Breeder's equation, selection differential

Lande-Arnold Regression

- Lande and Arnold (1983) proposed a method to estimate selection on phenotypes using regression
- operationally we can obtain selection gradients by estimating the partial regression coefficients of trait values on relative fitness (multiple regression)
- easy to implement, and can be used to disentangle selection acting directly and indirectly on a trait

Lande-Arnold Regression



Lande and Arnold 1983

Partitioning selection among traits

Selection Gradient Analysis: Total selection on a trait z is the sum of its direct effect on fitness plus the indirect effects of correlated traits

$$S_{z} = cov(z, w) = \beta_{z} + r_{zy}\beta_{y}$$
$$S_{y} = cov(y, w) = r_{zy}\beta_{z} + \beta_{y}$$

Partitioning selection among traits

$$direct$$

$$S_{z} = cov(z, w) = \frac{\beta_{z}}{\beta_{z}} + r_{zy}\beta_{y}$$

$$S_{y} = cov(y, w) = r_{zy}\beta_{z} + \frac{\beta_{y}}{\beta_{y}}$$

$$direct$$

Partitioning selection among traits



Directional selection is strong



- median variance standardized directional selection gradient of 0.153 for natural selection 0.250 for sexual selection
- Median mean standardized multivariate directional selection gradient was 0.54 (mean 0.28)

Hoekstra, H.E., Hoekstra, J.M., Berrigan, D., Vignieri, S.N., Hoang, A., Hill, C.E., Beerli, P. and Kingsolver, J.G., 2001. Strength and tempo of directional selection in the wild. *PNAS*, *98*(16), pp.9157-9160. Hereford, J., Hansen, T.F. and Houle, D., 2004. Comparing strengths of directional selection: how strong is strong?. *Evolution*, *58*(10), pp.2133-2143

Problems with correlated traits

- performing multiple regression on correlated traits can be a challenge because it is hard to estimate their independent effects on fitness
- this can lead to large standard errors of estimates and inaccurate estimates of selection

 over the years solutions like dropping traits, estimating selection on PCs, etc have been proposed

Problems with correlated traits



 when traits are correlated it can be hard to estimate their independent effects on fitness

 leads to large standard errors and inaccurate estimates of selection

 dropping correlated traits from analysis is a common solution

Regularized regression

- Linear regression with a penalty added
- Coefficients are constrained to be within a certain space





Regularized regression for estimating selection

- simulation study to compare the accuracy of Lande-Arnold regression to regularized regression for estimates selection
- re-analysed published studies of selection using regularized regression

Traits	Sample Size	Variance Inflation Factor	Traits under selection	Fitness distribution
4	100	low	all	binomial
7	400	mid/high	up to half	poisson
12	1000			
17				

More accurate estimates of the total strength of selection



Sztepanacz and Houle (2024) Evolution Letters 8: 361-373

Does not improve estimates of the direction of selection

4 traits low multicollinearity (VIF ~1)

17 traits high multicollinearity (VIF ~56)





Arabidopsis phenology

Estimated selection on Arabidopsis phenology from Chong et al., (2018)

Method		OLS regression	PC regression (4 PCs)	Lasso	Ridge
	Coefficients (+/– SE)				
	Flowering time	-0.299 (0.191)	-0.181	-0.371	-0.143
	Flowering duration	0.050 (0.182)	0.186	_	0.109
	Branch number	0.080 (0.085)	0.034	_	0.083
	Rosette diameter	0.059 (0.068)	0.072	_	0.072
	Rosette leaf number	0.061 (0.097)	0.033	_	-0.006
R^2	_	0.55 [0.007, 0.939]	_	0.680 [0.081, 0.997]	0.964 [0.838, 0.997]

Sztepanacz and Houle (2024) *Evolution Letters* 8: 361-373

Stabilizing selection

- even though we often focus on directional selection, stabilizing selection is the most common form of selection operating in populations
- stabilizing selection reduces the variance in the population
- disruptive selection increases the variance (eg. selection for extremes, negative frequency dependent selection)

Quadratic regression to estimate correlational and quadratic selection

$\overline{w} \sim \beta_1 x_1 + \beta_2 x_2 + \cdots + \beta_n x_n + \frac{1}{2} \gamma_1 x_1^2 + \frac{1}{2} \gamma_2 x_2^2 + \gamma_{12} x_1 x_2 \dots$

Quadratic regression to estimate correlational and quadratic selection



disruptive selection appears to be as common as stabilizing selection

Quadratic regression to estimate correlational and quadratic selection

Table 9.1 The vector of standardized linear gradients (β) and the matrix of standardized quadratic and correlational selection gradients (γ) for male call structure components in *T. commodus*. Note that the quadratic gradients were not doubled in the original study but have been done so here. The significance of selection gradients were tested using permutation test with 9999 iterations (see text): **P* <0.05, ***P* <0.01, ****P* <0.001

		γ					
	β	CPN	CIPD	TN	ICD	DF	
CPN	0.007	0.012					
CIPD	-0.003	0.017	-0.012				
TN	0.015	0.019	0.039	-0.080			
ICD	-0.214* * *	-0.022	-0.036	0.086*	-0.160**		
DF	0.059	0.024	-0.031	0.041	-0.013	-0.094*	

Canonical rotation of correlational and quadratic selection

Table 9.2 The **M** matrix of eigenvectors derived from the canonical analysis of γ . The linear (θ_i) and quadratic (λ_i) gradients of selection acting along these eigenvectors are provided in the last two columns. We tested the significance of selection along these eigenvectors using a permutation test based on the double regression method of (Bisgaard and Ankenman 1996) but note that the results were qualitatively the same when the permutation test of Reynolds et al. (2010) was used. Randomization test: *P < 0.05, **P < 0.01, ***P < 0.001

		Μ				Selection	
	CPN	CIPD	TN	ICD	DF	θi	λ_i
m ₁	0.800	0.497	0.305	-0.057	0.130	0.028	0.035
m ₂	-0.446	0.806	-0.001	-0.091	-0.377	-0.017	-0.003
m ₃	-0.302	0.003	0.776	0.500	0.240	-0.082	-0.019
m ₄	-0.257	0.208	-0.102	-0.405	0.846	0.144***	-0.108**
m ₅	0.068	0.244	-0.543	0.758	0.258	-0.160***	-0.240***



Figure 9.3 Thin-plate spline perspective view visualization of the fitness surface on the two major axes of non-linear selection, m_4 and m_5 . The thin-plate spline was estimated using the *Tps* function in R (version 9.2.1), using the value of the smoothing parameter, λ , that minimized the GCV score. Redrawn from Brooks et al. (2005), with permission from John Wiley and Sons.

Multivariate saddles may be common



Brodie III, E.D., 1992. Correlational selection for color pattern and antipredator behavior in the garter snake Thamnophis ordinoides. Evolution, 46(5), pp.1284-1298.

Pleiotropic model of MSB

- alleles affect a focal trait and their pleiotropic effects on other traits are condensed into an effect on net fitness
- mutations may increase or decrease the value of a focal trait, but their effects on fitness are almost certainly deleterious
- individuals with more extreme values of a focal trait will tend to carry alleles that have deleterious effects with respect to net fitness



Artificial selection experiments

• Evolutionary limits are also indicative of pleiotropic effects on fitness



Apparent stabilising selection



Sztepanacz J.L., and Rundle, H.D. (2012). <u>Reduced genetic variance among high fitness individuals: inferring stabilizing</u> selection on male sexual displays in *Drosophila serrata*. **Evolution.** 66(10): 3101-3110

Apparent stabilising selection



persistence time of new mutations: 6753 generations





persistence time of new mutations:
 562 generations

Apparent stabilising selection



Correlated responses to selection were larger than responses on targeted traits



Correlated selection against extreme phenotypes



- significant quadratic (stabilizing) selection via male reproductive failure in both g_{max} and m_{max} populations
- the multivariate trait combination under significant stabilizing selection via male reproductive failure was not g_{max} or m_{max}

Key take-aways:

- genetic variation is unevenly distributed across multivariate trait combinations because of pleiotropy
- the uneven distribution of genetic variance can lead to evolutionary constraints
- we can estimate selection on multiple traits using linear or quadratic regression approaches
- Most stabilizing selection may arise through the pleiotropic effects of alleles on multiple traits

Questions?