Chapter 8 Variance

Why variance ?
Components of variance

$$P = G + E$$

 $V_P = V_G + V_E + 2 \ Cov(G, E)$
 $= V_A + V_D + V_I + V_E$ (if $Cov(G, E) = 0$)

Table 8.1	Components	of	variance.
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Variance component	Symbol	Value whose variance is measured
Phenotypic	V_{P}	Phenotypic value
Genotypic	V_{C}	Genotypic value
Additive	V_A	Breeding value
Dominance	V_D^{α}	Dominance deviation
Interaction	V_I	Interaction deviation
Environmental	V_E	Environmental deviation

Heritability in the broad sense

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

Heritability in the narrow sense

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

Heritability measures the relative importance of heredity in determining phenotypic value.

 $V_P = V_A + V_D + V_I + V_E$

The partitioning of the variance into its components allows us to estimate the relative importance of various determinants of phenotype, in particular the role of heredity versus environment or nature and nurture.

Estimation of the degree of genetic determination, V_{G}/V_{P}

Environmental variance can't be removed because

Elimination of genotypic variance can be achieved experimentally.

Inbred lines and their F progeny are genetically uniform populations.

$$AA \times aa \downarrow Aa$$

Random bred populations are genetically mixed populations.

Example 8.1 : $h_b^2 = 49\%$ under the assumptions (a) the environmental variance is the same in all genotypes, (b) different genotypes show different sensitivities to the environment (genotype by environment interaction).

Population	Components	Observed variance
Mixed	$V_G + V_F$	0.366
Uniform	V_{F}	0.186
Difference	V_{G}	0.180
	$V_G^{\prime}/V_P =$	0.180/0.366 = 49%

Homozygote is more sensitive to environment than heterozygote.

Genetic components of variance

- G = A + D + I
- $V_G = V_A + V_D + V_I$
 - V_A is the variance of breeding value and is the chief cause of resemblance between relative and therefore the chief determinant of the observable genetic properties of the population and of the response of the population to selection.

 V_A/V_P is the heritability of the character.

Additive and dominance variance (under H-W equilibrium)

Note that the means of both breeding value and dominance deviations are zero (pg 116 and pg 119)

$$V_{A} = 2pq[a + d(q - p)]^{2} \dots [eqn \ 8.3 \ (a)]$$

$$V_{D} = (2pqd)^{2} \dots [eqn \ 8.4]$$

$$IF \quad d = 0 \quad , \quad V_{A} = 2pqa^{2} \quad , \quad and \quad V_{D} = 0$$

$$IF \quad d = a \quad , \quad V_{A} = 8pq^{3}a^{2}$$

$$IF \quad p = q \quad , \quad V_{A} = \frac{1}{2}a^{2} \quad , \quad and \quad V_{D} = \frac{1}{4}d^{2}$$

Both V_A and V_D depend on the genotypic values and gene frequencies.

Total genetic variance

G = A + D

$$V_G = V_A + V_D + 2 Cov_{AD}$$

Under H-W equilibrium, the covariance between additive and dominance effect is zero (pg 126)

$$V_G = 2pq[a+d(q-p)]^2 + [2pqd]^2$$

Example 8.2 : (pg 127)

	q = 0.1	q = 0.4	
Genotypic, V_G	1.1664	7.1424	
Additive, V_A	1.0368	6.2208	
Dominance, V_D	0.1296	0.9216	

The gene frequency and the degree of dominance influence the magnitude of the genetic components of variance (Fig 8.1)

The total phenotypic variance can then be partitioned into V_A , V_D , V_I , and V_E (Table 8.2)

Table 8.2 Partitioning of the variance of four characters in Drosophila melanogaster.Components as percentages of the total phenotypic variance.

		Character			
		(1) Bristles	(2) Thorax	(3) Ovary	(4) Eggs
Phenotypic	V_{P}	100	100	100	100
Additive genetic	VA	52	43	30	18
Non-additive genetic	$V_D + V_I$	9	6	35	44
Environmental	VE	39	51	35	38

Characters and sources of data:

 Number of bristles on 4th + 5th abdominal segments (Clayton, Morris, and Robertson, 1957; Reeve and Robertson, 1954).

(2) Length of thorax (Robertson, 1957b).

(3) Size of ovaries, i.e., number of ovarioles in both ovaries. (Robertson, 1957a).

(4) Number of eggs laid in 4 days (4th to 8th after emergence) (Robertson, 1957b).

For multiple loci, if no linkage disequilibrium

$$V_G = \sum V_{A_i} + \sum V_{D_i} + \sum V_{I_i}$$

Interaction variance

$$V_I = V_{AA} + V_{AD} + V_{DA} + V_{DD}$$

Interaction involving larger numbers of loci contribute so little variance that they can be ignored,.....

There is no doubt that interaction between loci controlling quantitative characters is a frequent occurrence.....(pg 190).

Variance due to linkage disequilibrium

 $D_{AB} = P_{AB} - P_A P_B \qquad (pg \ 16)$

Considering loci A and B,

$$G = G_A + G_B$$

$$V_G = V_A + V_B + 2 Cov(A, B)$$

Covariance can be positive or negative

 $(Cov(a,B) > 0 \quad or \quad Cov(A,B) < 0)$

There are two forms of non-random mating that generate disequilibrium

(1) selection

(2) assortative mating

Two sorts of covariance :

$$D_t = D_0 (1 - C)^t$$
 (pg 18)

Correlation and interaction between genotype and environment

Correlation

The better genotypes are given better environments.

Example (1) milk yield in cows

(2) human intelligence

 $V_P = V_G + V_E + 2 Cov(G, E)$

How to estimate Cov(G, E) ?

The covariance is best regarded as part of genetic variance because

Interaction

Genotype by environment interaction :

 $V_P = V_G + V_E + 2 Cov(G, E) + V_{GE}$

The environment variance is a property of the genotype.

But the source of variation is environmental and not genetic. It is therefore logical to regard any variance due to $G \times E$ as being part of environmental variance included in any estimate of V_E .

Environmental sensitivity Environmental value : Environmental sensitivity : Example 8.3 :

Environmental variance

All variation of non-genetic origin is environmental variance.

Environmental variance is a source of error that reduces precision in genetic studies and the aim of experimenter or breeder is therefore to reduce it as much as possible by careful management or proper design of experiments.

Multiple measurements : repeatability What is repeatability ? Temporal repetition and spatial repetition

Spatial environmental variance V_{Es}

General environmental variance V_{Eg}

Gain from multiple measurements :

Increasing the number of measurements reduces the amount of variance due to special environment that appears in phenotypic variance, and this reduction of the phenotypic variance represents the gain in accuracy.

$$V_{P(n)} = V_G + V_{Eg} + \frac{1}{n}V_{Es}$$

Example 8.6 :

The advantage of breeding programmes from the gain in accuracy is the increased proportion of additive genetic variance (two assumptions).

Prediction of future performance Example 8.7

Summary of variance partitioning

Homework: (1) The variances of leaf number in the F_1 and F_2 generations of a cross of tobacco varieties were calculated in Problem 6.1. The variances were 1.46 in the F_1 and 5.97 in the F_2 . Estimate the degree of genetic determination in the F_2 generation. What assumptions have to be made to do this? (2) Calculate the amounts of the additive genetic and dominance variance arising from genes referred to in the population specified in last homework.