

# Basics of Quantitative Genetics

Peter von Rohr

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## Background

- ▶ Central Dogma of Molecular Biology
  - Genotypes are the basis for phenotypic expression
- ▶ Start with simple model
  - one locus that affects quantitative trait

# Population

Population ( $N = 10$ )

$$\frac{G_1}{\overline{\overline{G_1}}}$$

$$\frac{G_1}{\overline{\overline{G_1}}}$$

$$\frac{G_1}{\overline{\overline{\overline{G_1}}}}$$

$$\frac{G_1}{\overline{\overline{G_2}}}$$

$$\frac{G_2}{\overline{\overline{G_1}}}$$

$$\frac{G_2}{\overline{\overline{G_2}}}$$

$$\frac{G_2}{\overline{\overline{G_2}}}$$

$$\frac{G_1}{\overline{\overline{G_1}}}$$

$$\frac{G_2}{\overline{\overline{G_1}}}$$

$$\frac{G_2}{\overline{\overline{G_2}}}$$

## Terminology

- ▶ **alleles**: variants occurring at a given genetic Locus
- ▶ **bi-allelic**: only two alleles, e.g.,  $G_1$  and  $G_2$  at a given locus  $G$  in population
- ▶ **genotype**: combination of two alleles at locus  $G$  in an individual
- ▶ **homozygous**: genotypes  $G_1G_1$  and  $G_2G_2$  where both alleles identical
- ▶ **heterozygous**: genotype  $G_1G_2$  different alleles

## Frequencies in Example Population

- ▶ **genotype frequencies**

$$f(G_1G_1) = \frac{4}{10} = 0.4$$

$$f(G_1G_2) = \frac{3}{10} = 0.3$$

$$f(G_2G_2) = \frac{3}{10} = 0.3$$

- ▶ **allele frequencies**

$$f(G_1) = f(G_1G_1) + \frac{1}{2} * f(G_1G_2) = 0.55$$

$$f(G_2) = f(G_2G_2) + \frac{1}{2} * f(G_1G_2) = 0.45$$

# Hardy-Weinberg Equilibrium

- ▶ **allele frequencies**

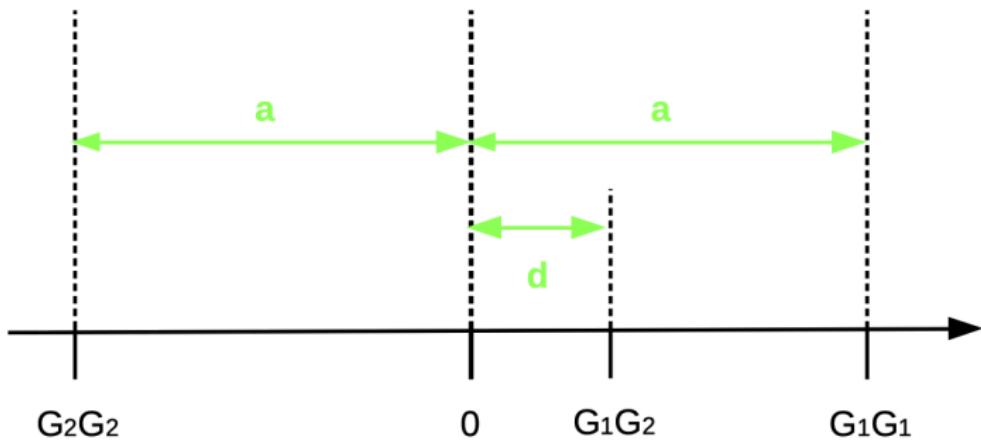
$$f(G_1) = p, f(G_2) = q = 1 - p$$

- ▶ **genotype frequencies**

Alleles	$G_1$	$G_2$
$G_1$	$f(G_1G_1) = p^2$	$f(G_1G_2) = p * q$
$G_2$	$f(G_1G_2) = p * q$	$f(G_2G_2) = q^2$

$$f(G_1G_1) = p^2, f(G_1G_2) = 2pq, f(G_2G_2) = q^2$$

## Genotypic Values



## Population Mean

- ▶ Expected value of genotypic value  $V$  as discrete random variable

$$\begin{aligned}\mu &= V_{11} * f(G_1G_1) + V_{12} * f(G_1G_2) + V_{22} * f(G_2G_2) \\&= a * p^2 + d * 2pq + (-a) * q^2 \\&= (p - q)a + 2pqd\end{aligned}$$

## Breeding Values Definition

The breeding value of an animal  $i$  is defined as two times the difference between the mean value of offsprings of animal  $i$  and the population mean.

## Derivation of Breeding value for $G_1G_1$

	Mates of $S$	
	$f(G_1) = p$	$f(G_2) = q$
Parent $S$		
$f(G_1) = 1$	$f(G_1G_1) = p$	$f(G_1G_2) = q$

## Computation of Breeding value for $G_1G_1$

$$\mu_{11} = p * a + q * d$$

The breeding value  $BV_{11}$  corresponds to

$$\begin{aligned} BV_{11} &= 2 * (\mu_{11} - \mu) \\ &= 2(pa + qd - [(p - q)a + 2pqd]) \\ &= 2(pa + qd - (p - q)a - 2pqd) \\ &= 2(qd + qa - 2pqd) \\ &= 2(qa + qd(1 - 2p)) \\ &= 2q(a + d(1 - 2p)) \\ &= 2q(a + (q - p)d) \end{aligned}$$

## Computation of Breeding value for $G_2G_2$

$$\mu_{22} = pd - qa$$

The breeding value  $BV_{22}$  corresponds to

$$\begin{aligned} BV_{22} &= 2 * (\mu_{22} - \mu) \\ &= 2 (pd - qa - [(p - q)a + 2pqd]) \\ &= 2 (pd - qa - (p - q)a - 2pqd) \\ &= 2 (pd - pa - 2pqd) \\ &= 2 (-pa + p(1 - 2q)d) \\ &= -2p(a + (q - p)d) \end{aligned}$$

## Computation of Breeding value for $G_1G_2$

$$\mu_{12} = 0.5pa + 0.5d - 0.5qa = 0.5 [(p - q)a + d]$$

The breeding value  $BV_{12}$  corresponds to

**BV12**

$$\begin{aligned}\text{BV12} &= 2 * (\mu_{12} - \mu) \\&= 2 (0.5(p - q)a + 0.5d - [(p - q)a + 2pqd]) \\&= 2 (0.5pa - 0.5qa + 0.5d - pa + qa - 2pqd) \\&= 2 (0.5(q - p)a + (0.5 - 2pq)d) \\&= (q - p)a + (1 - 4pq)d \\&= (q - p)a + (p^2 + 2pq + q^2 - 4pq)d \\&= (q - p)a + (p^2 - 2pq + q^2)d \\&= (q - p)a + (q - p)^2d \\&= (q - p) [a + (q - p)d]\end{aligned}$$