# Overview/reminder of basic concepts in statistics and genetics

# SISG, Visscher-Goddard Module

A <u>discrete</u> random variable can assume only a countable number of values

Probability mass function:

$$p(x) = P(X = x)$$

Expected value:

$$\mu = E(X) = \sum x p(x)$$

As a function of random variable:

$$E[h(X)] = \sum h(x)p(x)$$

<u>Variance</u>:

$$Var(X) = E[(X - \mu)^2]$$

A <u>discrete</u> random variable can assume only a countable number of values

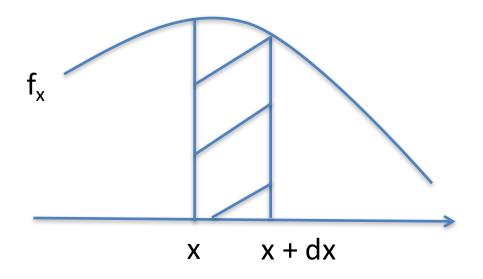
allele x:

$$p(x) = \left\{ \begin{array}{cc} p & x = 1\\ 1 - p & x = 0 \end{array} \right\}$$

$$E(X) = 0(1 - p) + 1(p) = p$$

$$Var(X) = p - p^2 = p(1 - p)$$

A <u>continuous</u> random variable can be any value within a range



probability of being in shaded area

$$= f_X(x)dx$$

the interval should contribute

$$= x f_X(x) dx$$

the expected value and variance

$$E(X) = \mu_X = \int_{-\infty}^{\infty} x f_X(x) dx \qquad Var(X) = E((X - \mu_x)^2)$$

Covariance

Let X and Y be a pair of continuous random variables, with respective means  $\mu_x$  and  $\mu_y$ . The expected value of (X -  $\mu_x$ )(Y -  $\mu_y$ ) is called the covariance between X and Y.

$$Cov(X,Y) = E\left[\left(X - \mu_x\right)\left(Y - \mu_y\right)\right]$$

If the random variables X and Y are independent, then the covariance between them is 0. However, the converse is not true.

# Summary (co)variance rules

$$Var(x) = E[x - E(x)]^{2}$$
$$Var(cx) = c^{2}Var(x)$$
$$Var(x + y) = Var(x) + Var(y) + 2Cov(x, y)$$
$$Var(x + c) = Var(x)$$

$$Cov(x, y) = E[(x - E(x)(y - E(y))]$$
$$Cov(cx, y) = cCov(x, y)$$
$$Cov(x, y + z) = Cov(x, y) + Cov(x, z)$$

# **Bayes' Theorem**

Identify people who are liable to suffer from a genetic disease later in life. 1 in 1000 people are a carrier of the disease

No test is perfect - probability that a carrier tests negative is 1%

- probability that a non-carrier tests positive is 5%

- A = the event "the patient is a carrier"
- B = the event "the test result is positive"

Hence: P(A) = 0.001; P(A') = 0.999; P(B|A) = 0.99; P(B|A') = 0.05

A patient has a positive result. **Q**: What is the probability that the patient is a carrier?

#### Answer

$$P(A \mid B) = \frac{P(B \mid A)P(A)}{P(B \mid A)P(A) + P(B \mid A')P(A')}$$
$$= \frac{0.99 * 0.001}{(0.99 * 0.001) + (0.05 * 0.999)} = 0.0194$$

# Hardy-Weinberg equilibrium

Mathematical relation between **allele** frequencies and the **genotype** frequencies is:

		AA: p <sup>2</sup>	Aa: 2pq	aa: q²
	Allele Frequency	A p		a q
Allele	Frequency	AA		Аа
А	р	p²		pq
а	q	aA		аа
		qp		q²

## **HWE and SNPs**

If SNP genotypes are coded X = 0, 1 and 2 (alleles) and the allele frequency is *p*, then:

 $E(X) = (1-p)^{2*}0 + 2p(1-p)^{*}1 + p^{2*}2 = 2p$ 

 $\operatorname{var}(X) = (1-p)^{2*}(0-2p)^{2} + 2p(1-p)^{*}(1-2p)^{2} + p^{2*}(2-2p)^{2} = 2p(1-p)$