

Chap 1: POPULATION GENETICS

1.1 Segregation Principle

- Gamete randomly receive one of adult's 2 alleles (Aa) for each autosomal gene with equal probability.
- Gamete randomly receive one of the adult's two sex chromosomes (XY) with equal probability.
- Occurs **independently** for genes on different chromosomes.
- Example:

○ Aa x Aa

	A	a
A	AA	Aa
a	Aa	aa

- By **segregation principle**, E_1 =offspring get A from father, E_2 =offspring get A from mother $\rightarrow P(E_1) = P(E_2) = 0.5$
- Events are **independent**.
 - $P(AA)$ \rightarrow 25%
 - $P(AA) = P(E_1 \cap E_2) = P(E_1) \times P(E_2)$
 - multiplication law for independent events as E_1 and E_2 are independent
 - $0.5 * 0.5 = \mathbf{0.25}$
 - $P(Aa)$ \rightarrow 50%
 - $P[(E_1 \cap E'_2) \cup (E'_1 \cap E_2)]$
 - $P(E_1 \cap E'_2) + P(E'_1 \cap E_2)$ as mutually exclusive
 - $[P(E_1) \times P(E'_2)] + [P(E'_1) \times P(E_2)]$ as E_1 and E_2 are independent
 - $(0.5 * 0.5) + (0.5 * 0.5) = \mathbf{0.5}$
 - $P(aa)$ \rightarrow 25%

1.2 Law of probability

- Event \rightarrow something that may or may not occur.
- Complement Law:
 - $P(E') = 1 - P(E)$
- Conditional probability:
 - $P(E_1|E_2) = \frac{P(E_1 \cap E_2)}{P(E_2)}$
- Independent events: (multiplication law)
 - $P(E_1 \cap E_2) = P(E_1) \times P(E_2)$
- Mutually exclusive: (addition law)
 - $P(E_1 \cap E_2) = 0$

$$\circ P(E_1 \cup E_2) = P(E_1) + P(E_2)$$

1.3 Gene Disorders

- 4000 (rare) diseases, caused by single gene disorders/Mendelian disease
 - Depend on whether gene is:
 - Autosomal / sex-linked
 - Dominant / recessive
- Example:
 - Autosomal dominant
 - Huntington's disease (frequency 1 in 40,000)
 - Autosomal recessive
 - Sickle cell anemia (frequency 1 in 625)
 - X-linked recessive
 - Red-green color blindness (frequency 1 in 125)
- **Prevalence VS Incidence**
 - Prevalence (frequency) -> total number/proportion of **population** affected by disease **at a given time**.
 - Incidence -> total number/proportion of **new births** that are affected by disease.
 - If incidence is constant, incidence = prevalence

1.4 Genotype Number and Frequency

- N – total population size
- N^{AA} – number of AA individuals in population
- f^{AA} – frequency of AA in population $f^{AA} = \frac{N^{AA}}{N}$
- $N^{AA} + N^{Aa} + N^{aa} = N$ and $f^{AA} + f^{Aa} + f^{aa} = 1$

1.5 Allele Numbers and Frequency

- N^A – number of A alleles in population
- f^A – frequency of A allele in **allele pool** (allele pool (single allele) from everyone in population)

1.6 Calculating allele frequencies from genotype frequencies

- $N^A = 2N^{AA} + N^{Aa}$ $N^a = 2N^{aa} + N^{Aa}$
- $f^A = \frac{N^A}{2N}$ $f^a = \frac{N^a}{2N}$
- $f^A = \frac{N^A}{2N} = \frac{2N^{AA} + N^{Aa}}{2N} = \frac{N^{AA}}{N} + \frac{1}{2} \frac{N^{Aa}}{N}$
- $f^A = f^{AA} + \frac{1}{2} f^{Aa}$ $f^a = f^{aa} + \frac{1}{2} f^{Aa}$ (note: $f^A + f^a = 1$)

1.7 Modelling assumptions (assumption to derive a math model) in population genetics

- 1.7.1. Random mating
 - Adults choose mating partner random (mate selection independent of genotypes)
 - Same as random sampling (**with replacement**) from the allele pool
 - Thus:
 - **Single random mating**

- $P(\text{offspring is AA}) = (f^A)^2$
- $P(\text{offspring is Aa}) = 2f^A f^a = 2f^A(1 - f^A)$
- $P(\text{offspring is aa}) = (f^a)^2 = (1 - f^A)^2$

▪ **N random mating**

- $N^{AA} \sim \text{Bi}(N, (f^A)^2)$
- $N^{Aa} \sim \text{Bi}(N, 2f^A(1 - f^A))$
- $N^{aa} \sim \text{Bi}(N, (1 - f^A)^2)$

○ $X \sim \text{Bi}(n, p)$ n=no. trials, p=probability of success

▪ $P(X = k) = \binom{n}{k} p^k (1 - p)^{n-k}$

▪ $E(X) = np$

• $E(N^{AA}) = N(f^A)^2$

• $E(f^{AA}) = \frac{1}{N} E(N^{AA}) = (f^A)^2$

▪ $\text{Var}(X) = np(1 - p)$

• $\text{var}(N^{AA}) = N(f^A)^2(1 - (f^A)^2)$

• $\text{var}(f^{AA}) = \frac{1}{N^2} \text{var}(N^{AA})$

- Mean genotype frequencies are independent of N, but S.D. shrink as N increase.

• 1.7.2 Large population ($N \cong 10^4$) assumption

○ As N increases, genotype frequency become concentrated on mean value, ignore variability in f^{AA}

○ $f^{AA} = E(f^{AA})$