

POPULATION GENETICS

Evolution Lectures 4

POPULATION GENETICS

The study of the rules governing the maintenance and transmission of genetic variation in natural populations.

Some Definitions

Population: A freely interbreeding group of individuals.

Gene Pool: The sum total of genetic information present in a population at any given point in time.

Phenotype: A morphological, physiological, biochemical, or behavioral characteristic of an individual organism.

Genotype: The genetic constitution of an individual organism.

Locus: A site on a chromosome, or the gene that occupies the site.

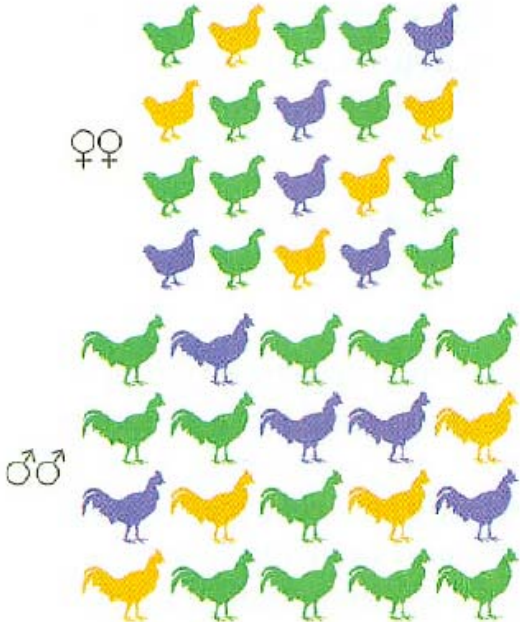
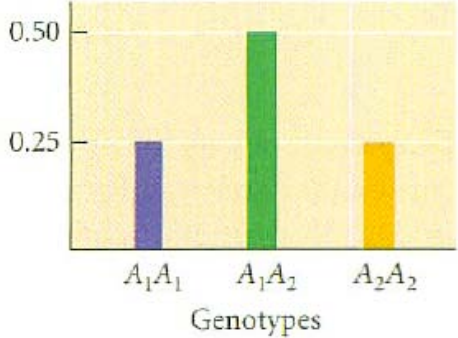
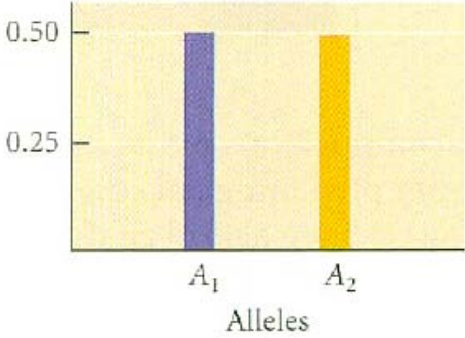
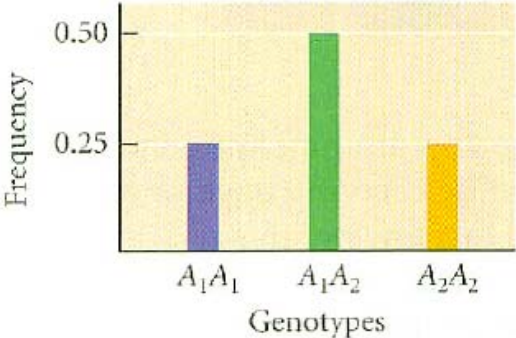
Gene: A nucleic acid sequence that encodes a product with a distinct function in the organism.

Allele: A particular form of a gene.

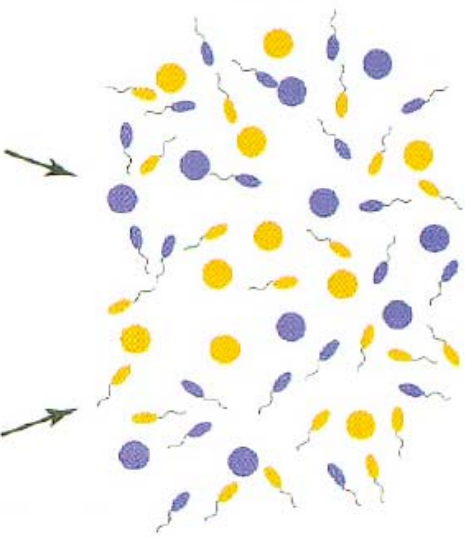
Gene (Allele) Frequency: The relative proportion of a particular allele at a single locus in a population (a number between 0 and 1).

Genotype Frequency: The relative proportion of a particular genotype in a population (a number between 0 and 1).

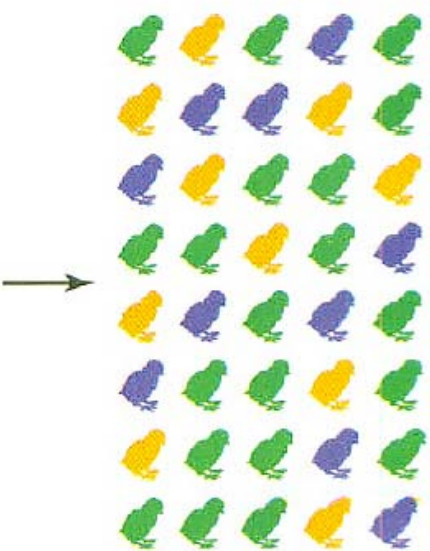
Genotype vs. Allele frequency



Genotype frequencies, parental generation



Allele frequencies among gametes



Genotype frequencies, offspring generation

Calculation of allele frequency

- Lets assume:
 - In a population 400 people have the genotype TT, 400 have Tt and 200 have tt
 - So they have 2000 allele total
 - Of these 1200 are T and 800 are t
 - If frequency of T is p, then $p=0.60$
 - If frequency of t is q, then $q=0.40$

Calculation of allele frequency

- By counting
 - $T = 800 \text{ in } TT + 400 \text{ in } Tt = 1200/2000 = 0.60$
 - $t = 400 \text{ in } Tt + 400 \text{ in } tt = 800/2000 = 0.40$
 - Or frequency of an allele=frequency of homozygote for that allele + $\frac{1}{2}$ frequency of heterozygote
- Using genotype frequency
 - $T = 0.40 TT + \frac{1}{2} (0.40 Tt) = 0.40 + 0.20 = 0.60$
 - $t = 0.20 tt + \frac{1}{2} (0.40 Tt) = 0.20 + 0.20 = 0.40$

Conservation of allele frequency

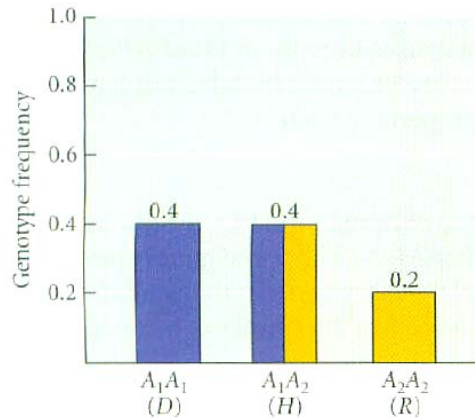
$\begin{matrix} \nearrow \\ \text{♀} \end{matrix}$ / $\begin{matrix} \searrow \\ \text{♂} \end{matrix}$	TT=0.40	Tt=0.40	tt=0.20
TT=0.40	0.160	0.160	0.08
Tt=0.40	0.160	0.160	0.08
tt=0.20	0.08	0.08	0.04

In F1; TT=0.36, Tt=0.48; tt=0.16

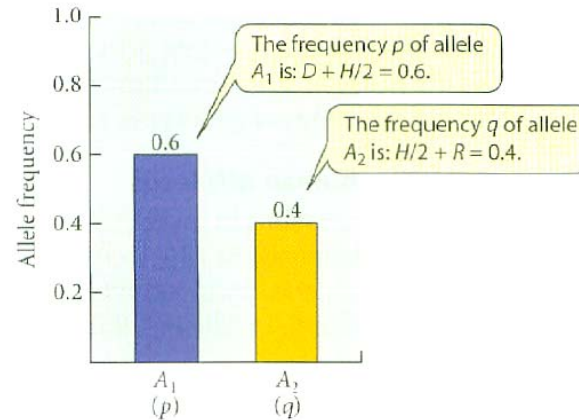
But; T=0.60 and t=0.40

Conservation of allele frequency

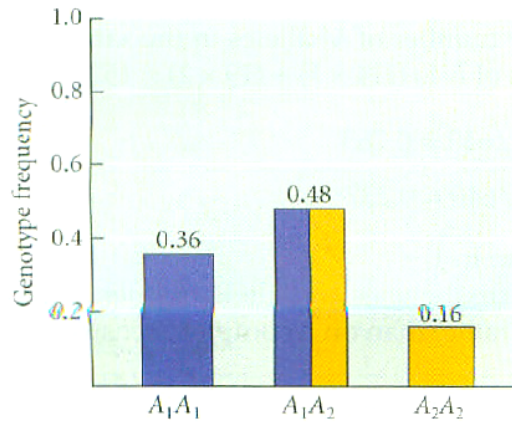
(A) Parental genotype frequencies
(not in equilibrium)



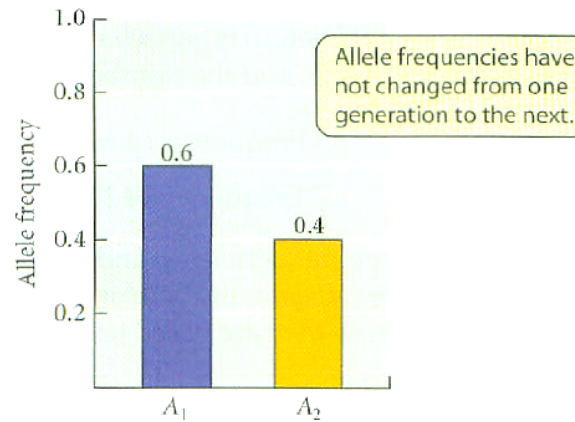
(B) Parental allele frequencies



(D) Offspring genotype frequencies



(E) Offspring allele frequencies



Calculation of allele frequency

- In F1; $TT=0.36$, $Tt=0.48$; $tt=0.16$
- And $T=0.60$ and $t=0.40$
- Also if $p=T$ and $q=t$
- Then $p^2=0.36$, $q^2=0.16$ and $2pq=0.48$

The Hardy-Weinberg Equilibrium

A single generation of random mating establishes H-W equilibrium genotype frequencies, and neither these frequencies nor the gene frequencies will change in subsequent generations.

$$p^2 + 2pq + q^2 = 1$$

Hardy-Weinberg assumptions

- Mating is random (with respect to the locus).
- The population is infinitely large.
- Genes are not added from outside the population (no gene flow or migration).
- Genes do not change from one allelic state to another (no mutation).
- All individuals have equal probabilities of survival and reproduction (no selection).

Implications of the Hardy-Weinberg equilibrium

- A random mating population with no external forces acting on it will reach the equilibrium H-W frequencies in a single generation, and these frequencies remain constant there after.
- Any perturbation of the gene frequencies leads to a new equilibrium after random mating.
- The amount of **heterozygosity** is maximized when the gene frequencies are intermediate.
 - **$2pq$ has a maximum value of 0.5 when**
 - **$p = q = 0.5$**

Genotype frequency distribution

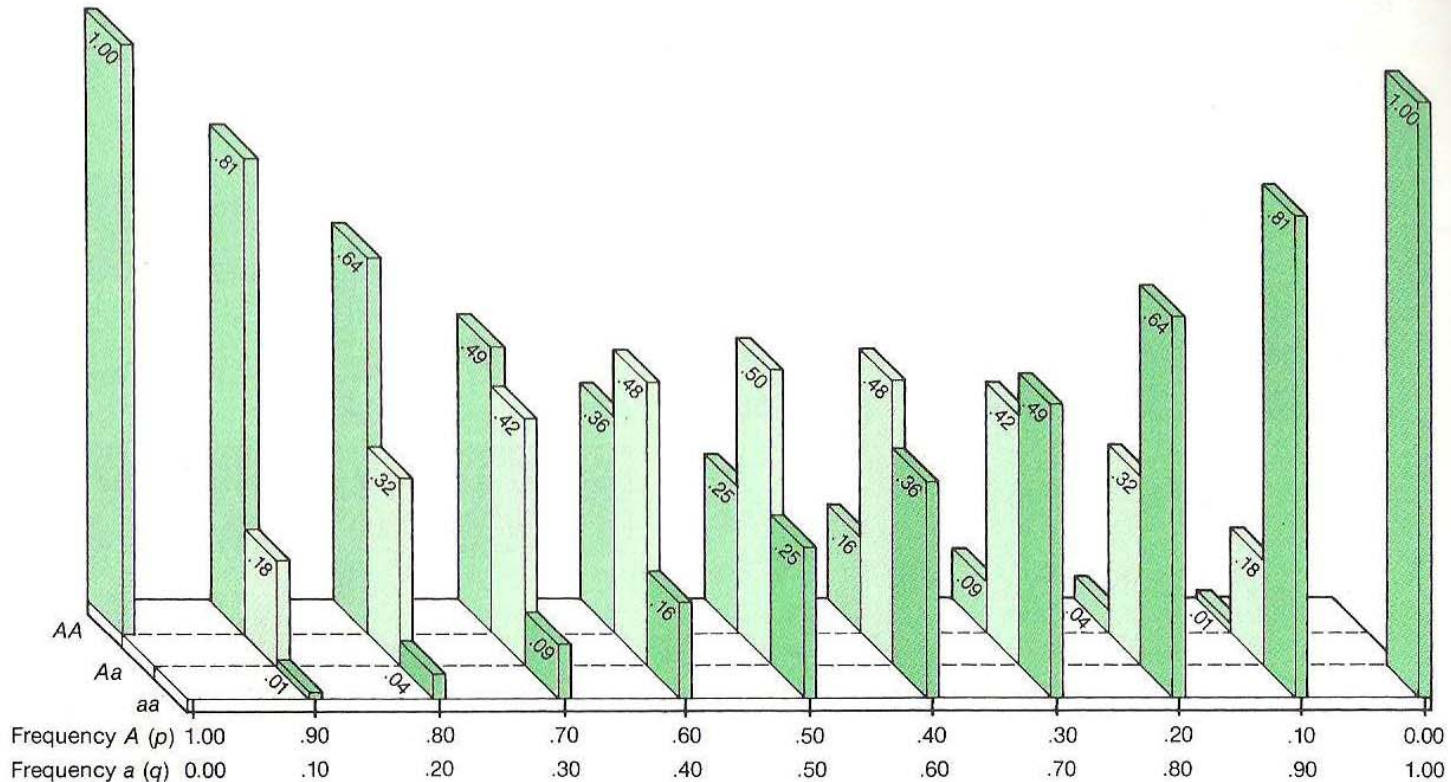
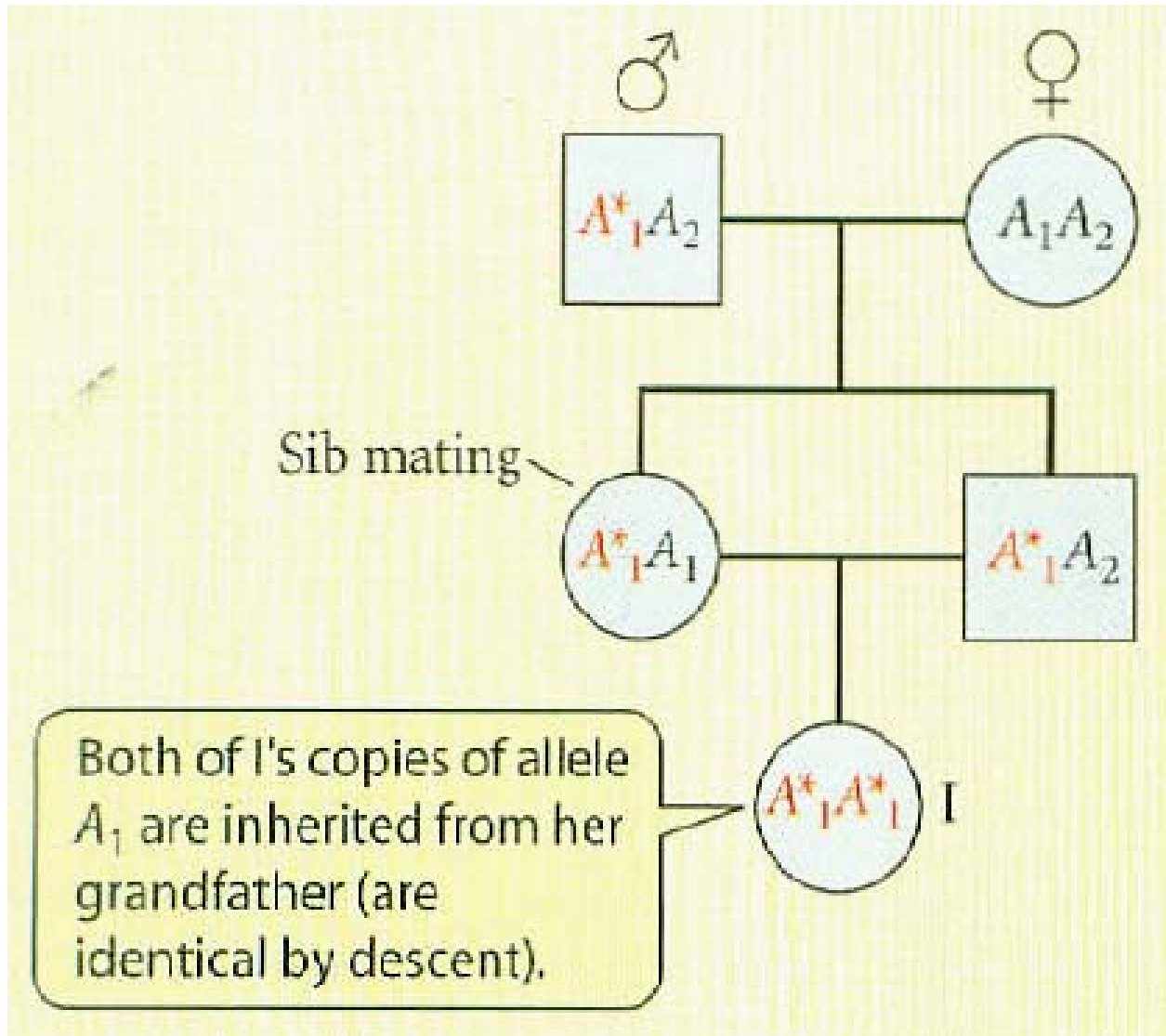


FIGURE 21-4 Genotypic frequencies at Hardy-Weinberg equilibrium for a variety of allele frequencies of A (p) and a (q). (Adapted from Wallace, B., 1970. *Genetic Load: Its Biological and Conceptual Aspects*. Prentice Hall, Englewood Cliffs, NJ.)

Factors changing equilibrium

Inbreeding

Example



Calculations

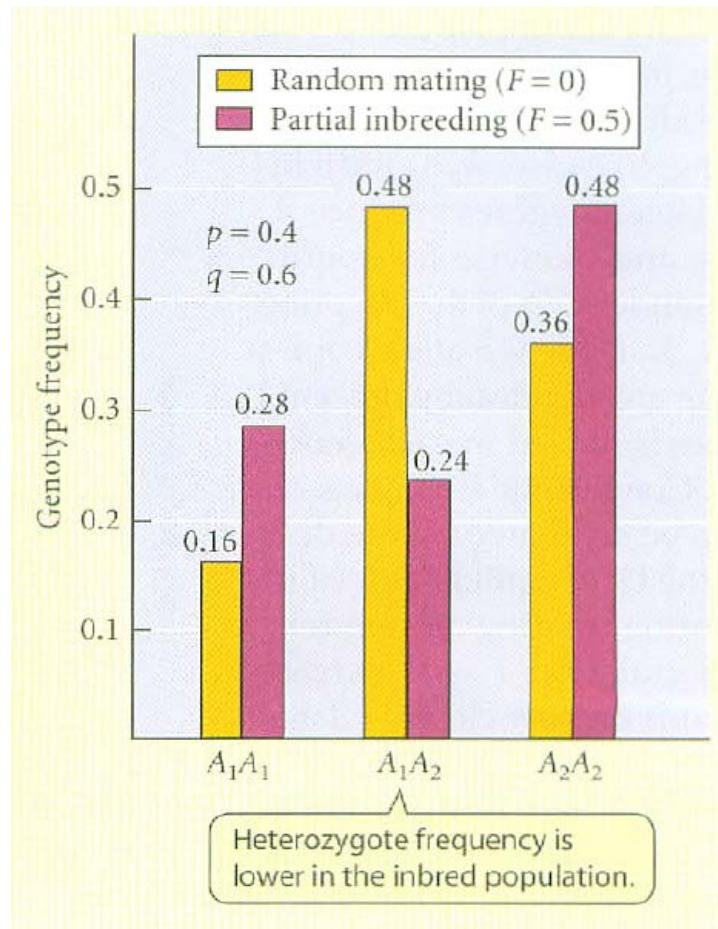
- Definition first
 - Autozygous: individuals homozygous with the identical allele by descent
 - Allozygous: either heterozygous or homozygous individuals with non-identical allele
 - Inbreeding coefficient (F) is the probability of an individual taken at random from a population will be autozygous
 - In a randomly mating population $F=0$ and in an all inbred population $F=1$

Calculations

In a population with some inbreeding $1-F$ is the allozygous frequency if F is autozygous frequency

	Allozygous	Autozygous	Genotype frequency
A_1A_1	$p^2(1-F)$	$+pF$	$=p^2+Fpq$
A_1A_2	$2pq(1-F)$		$2pq(1-F)$
A_2A_2	$q^2(1-F)$	$+qF$	$=q^2+Fpq$

In a graph



Find: How to calculate inbreeding frequency in a population?

Factors changing equilibrium

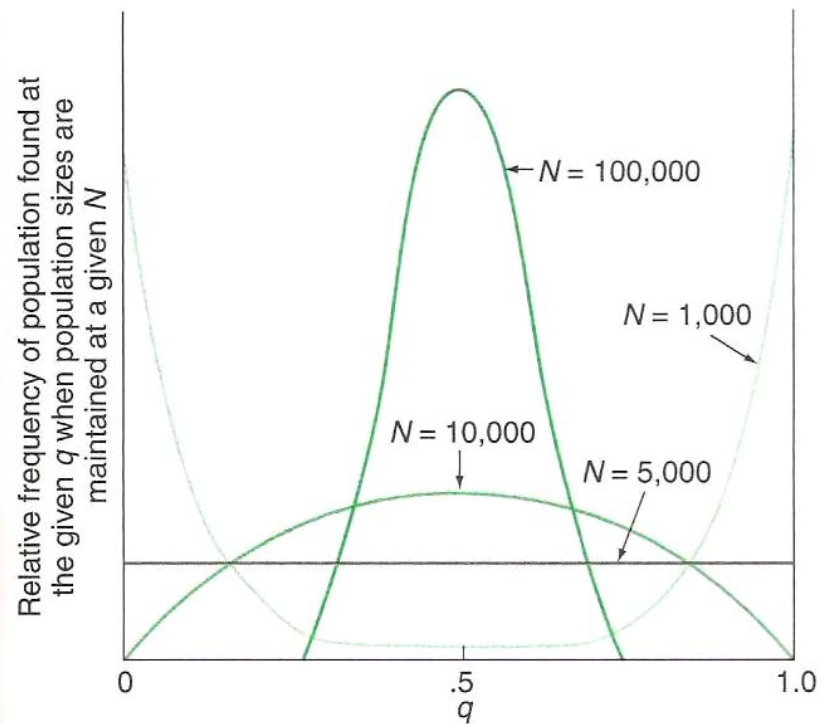
Genetic drift

GENETIC DRIFT

- Alteration of gene frequencies due to chance (**stochastic**) effects.
- Most important in *small* populations.
- Tends to reduce genetic variation as the result of extinction of alleles.
- Generally does not produce a fit between organism and environment; can, in fact, result in nonadaptive or maladaptive changes.

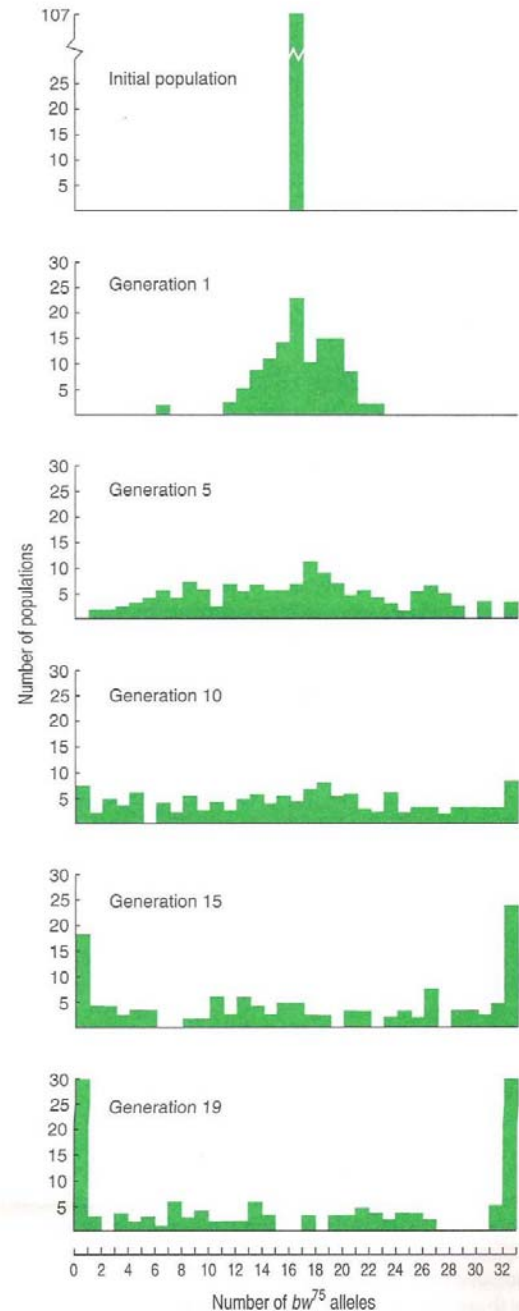
How does it work?

- Calculated by standard deviation $\sigma = \sqrt{pq/N}$ (p is the frequency of one allele, q is another; N is the number of genes sampled)
- For diploid population $\sigma = \sqrt{pq/2N}$
- If we start $p=q=0.5$ and $N=100000$
- Then $\sigma = .001$
- That is the frequency of either p or q will vary 0.5 ± 0.001
- This is genetic drift and is greater in smaller population



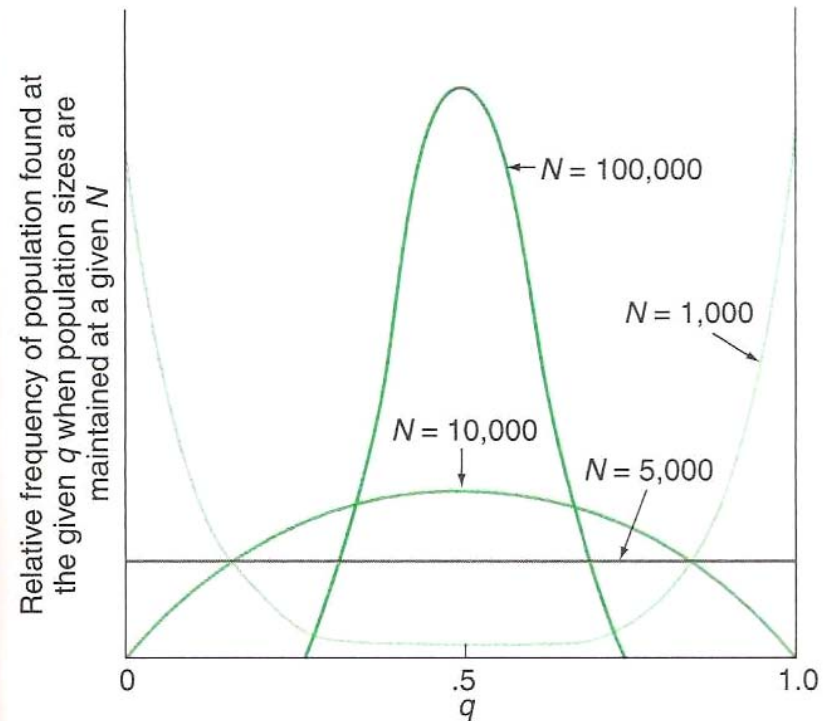
Example

- Laboratory expt by Buri (1956)
- Set up 107 separate *D. melanogaster* lines, each with two brown alleles bw and bw^{75}
- Started each generation with 16 parents (8 males and 8 females, random selection)
- Therefore each generation started with $16 \times 2 = 32$ gene copies
- Continued for 19 generations
- See the changes in the allele frequency in the first generation
- At 19th 30 populations lost all bw^{75} allele and 28 have been fixed.
- Remember: No selection pressure



How does it work?

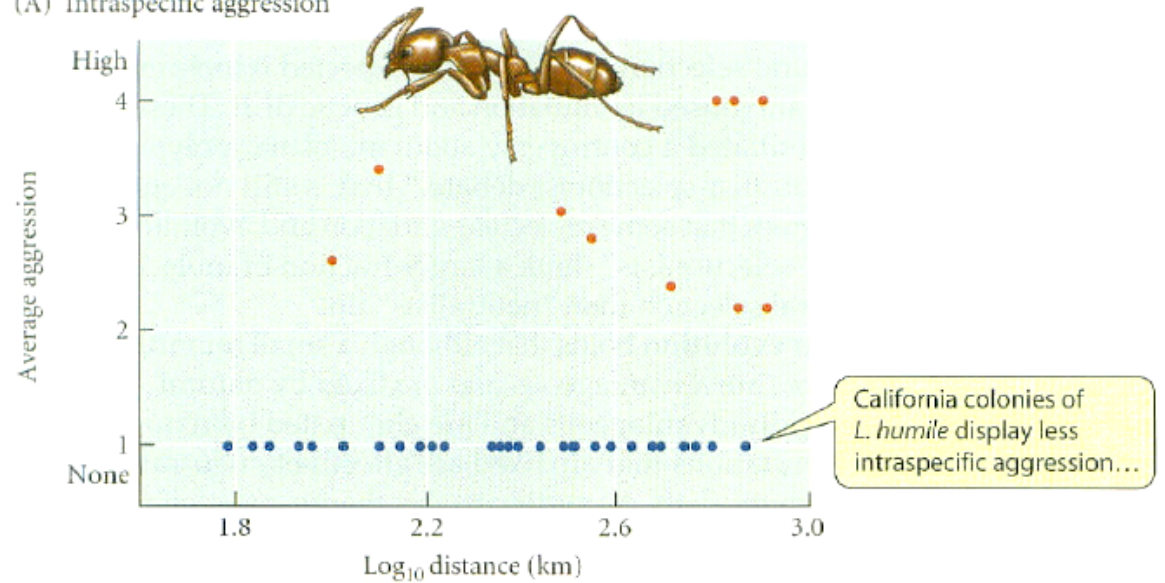
- If we start $p=q=0.5$ and $N=214$
- Then $\sigma=.034$
- That is the frequency of either p or q will vary 0.5 ± 0.03
- This is just one generation
- In the next
 - at one end 0.47 ± 0.03
 - and at the other 0.53 ± 0.04
- Do the calculations for few more generations
- This is what is called Founder's Effect



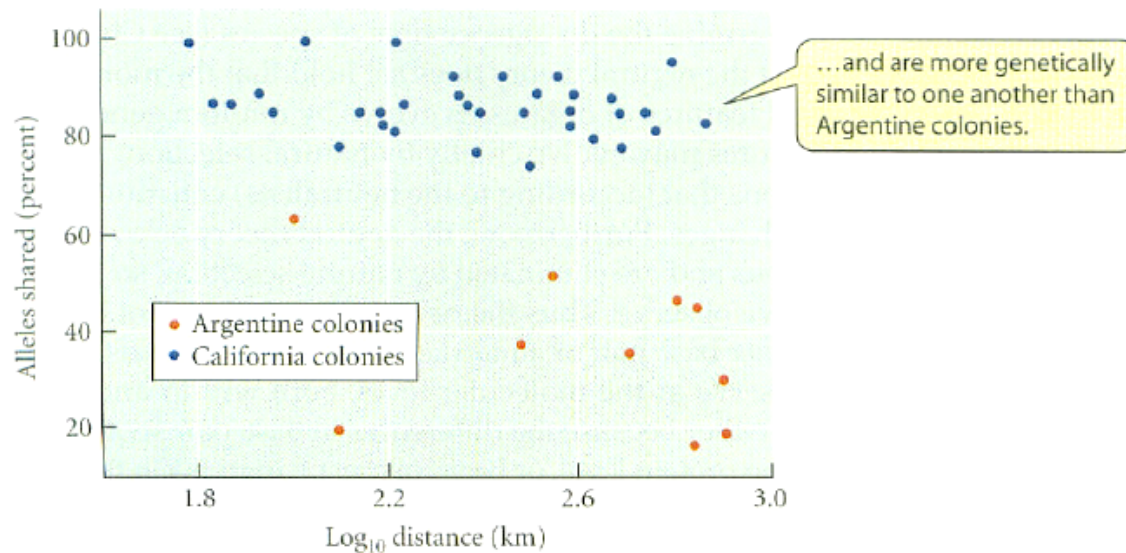
Example

- Argentine ant (*Linepithema humile*)
- Introduced in California accidentally
- Highly aggressive in Argentina, between colonies
- Each colony has different odor
- But they form super colony in California
- Small group brought had same colony odor

(A) Intraspecific aggression



(B) Genetic similarity



Coalescence

