

SELECTION & THE SINGLE DIALLELIC DIPLOID LOCUS. 1

• General equations

– Diploid selection to asexual/haploid selection except that parental genotypes not passed on, although their alleles are.

– Will repeatedly use the following facts from our H-W discussion :

(1) Offspring allele frequencies = those of their randomly mating parents

(2) Offspring genotype frequencies are in H-W proportions, even if their parents’ were not.

– **Relative fitnesses** of diploid genotypes: w_{AA} , w_{Aa} , w_{aa}

– frequency of A allele = p ; frequency of allele a = $q = 1 - p$.

• assume zygotes are in H-W proportions.

– Life Cycle: $\text{zygotes}_{\text{gen. } t} \xrightarrow{\text{survival}} \text{adults}_{*} \xrightarrow{\text{fecundity}} \text{gametes} \xrightarrow{\text{random union}} \text{zygotes}_{\text{gen. } t+1(')}$

– Assume all have same fecundity so relative fitness = viability

– Genotype frequencies after selection (but before reproduction)

• denoted by *

$$P_{AA}^* = \frac{\text{number of AA parents surviving}}{\text{total number of surviving parents}} = \frac{Np^2w_{AA}}{Np^2w_{AA} + N2pqw_{Aa} + Nq^2w_{aa}} = p^2 \frac{w_{AA}}{\bar{w}}$$

where $\bar{w} = p^2w_{AA} + 2pqw_{Aa} + q^2w_{aa}$ is the population **mean fitness**.

$$\text{• Similarly, } P_{Aa}^* = 2pq \frac{w_{Aa}}{\bar{w}}, \quad P_{aa}^* = q^2 \frac{w_{aa}}{\bar{w}}.$$

– Allele frequencies after selection and reproduction

• If surviving parents mate randomly, then

$$p' = p^* = P_{AA}^* + \frac{1}{2}P_{Aa}^* = p^2 \frac{w_{AA}}{\bar{w}} + pq \frac{w_{Aa}}{\bar{w}} = p \frac{(pw_{AA} + qw_{Aa})}{\bar{w}}$$

• Quantity in parentheses is **mean fitness of individuals who carry A**:

$$\bar{w}_A \equiv pw_{AA} + qw_{Aa}$$

• Can then rewrite above as, $p' = p \frac{\bar{w}_A}{\bar{w}}$.

• Similarly, $q' = q \frac{\bar{w}_a}{\bar{w}}$ where $\bar{w}_a \equiv p w_{Aa} + q w_{aa}$ is the **mean fitness of individuals who carry *a***.

– Handy Fact: $\bar{w} = p \bar{w}_A + q \bar{w}_a$ (compare with *asexual* mean fitness)

– Rate of allele frequency change ($\Delta p = p' - p$): $\Delta p = pq \frac{\bar{w}_A - \bar{w}_a}{\bar{w}}$

– Summarize:

- Equations for evolution by selection in diploids identical (in form) to asexuals/haploids.
- Allele frequency change (evolution) depends on **genetic variation** and **fitness differences**

• **Spread of an advantageous allele**

– Designate *A* to be the advantageous allele.

– Adopt the following notation:

Genotype	<i>AA</i>	<i>Aa</i>	<i>aa</i>
Fitness	$w_{AA} = 1$	$w_{Aa} = 1 - hs$	$w_{aa} = 1 - s$

– *h* measures allelic **dominance** of *A* over *a*:

- $h = 1 \Leftrightarrow 1 : 1 - s : 1 - s$ (*A* is recessive advantageous)
- $h = 0 \Leftrightarrow 1 : 1 : 1 - s$ (*A* is dominant advantageous)
- $h = 1/2 \Leftrightarrow 1 : 1 - s/2 : 1 - s$ (semi-dominant, additive, no dominance)
- $h \neq 0, 1, 1/2$ (partially dominant)

– Equations in this parameterization for allele frequency evolution:

$$\Delta p = pq \frac{s[q + h(p - q)]}{1 - sq(q + 2ph)}$$

– Application (from Y.J. Chung. Genetics. 1967)

- Selection against allele that produces stubble bristles in *Drosophila*

Genotype	+/+	+sb	sb/sb
Phenotype (Fitness)	normal	stubble	lethal in larval stages ($s = 1$)

– Two Special Cases:

(1) Dominant advantageous allele ($h = 0, \quad w_{AA} : w_{Aa} : w_{aa} \quad)$

$$\Delta p = p(1-p) \frac{s(1-p)}{1-s(1-p)^2}$$

- Dynamics: spread of allele is rapid at first; later, slows as A spreads.
- Why these dynamics?
 - Biological Intuition
 - Mathematical Explanation

(2) Recessive advantageous allele ($h = 1, \quad w_{AA} : w_{Aa} : w_{aa} \quad)$

$$\Delta p = p(1-p) \frac{sp}{1-s(1-p^2)}$$

- Dynamics: spread of allele A is slow at first; rapid later as A spreads.
- Why these dynamics?
 - Biological Intuition
 - Mathematical Explanation

– Rates of evolution: Dominant vs. Recessive Alleles

- Basic Principles:
 - rare alleles occur primarily in heterozygotes

- intuition for the magnitude of change when an allele is rare: compare heterozygous fitness to that of the common homozygote.
- Main Features:
 - Extensive time needed for spread of advantageous recessive (extensive time needed for elimination of deleterious recessive)
 - Rapid evolution from 0.1 to 0.9 in all cases
 - Multiplicative selection: produces even faster evolution (homework)
- For weak selection ($s \ll 1$), time required for a specified allele frequency change is inversely proportional to s , i.e., $t \propto 1/s$ when $s \ll 1$.