

# Deriving $\hat{q} = \sqrt{(\mu/s)}$

Step-by-Step Mathematical Derivation

## Model Assumptions

Before we begin, recall our key assumptions:

- Infinite population size (no genetic drift)
- Random mating
- Recessive deleterious allele (a)
- Mutation rate  $A \rightarrow a = \mu$  per generation
- Selection coefficient against aa = s
- Fitness:  $AA = 1$ ,  $Aa = 1$ ,  $aa = 1 - s$
- No reverse mutation ( $a \rightarrow A$  negligible)
- No migration

## 1 Define the Equilibrium Condition

At mutation-selection balance, the input of new a alleles by mutation equals the removal of a alleles by selection:

$$\text{Input by mutation} = \text{Removal by selection}$$

**Concept:** Think of this as a bathtub with water flowing in (mutation) and draining out (selection). At equilibrium, the water level stays constant because inflow equals outflow.

## 2 Calculate Input by Mutation

Each generation, a fraction  $\mu$  of A alleles mutate to become a alleles.

Number of A alleles  $\approx 2Np$  (since  $p \approx 1$  for rare deleterious alleles)

Mutations per generation =  $2Np \times \mu$

Since we're working with frequencies, we can use the proportion:

$$\Delta q_{\text{mut}} = \mu$$

**Why  $\mu$  and not  $\mu p$ ?** Because when the deleterious allele is rare, almost all alleles are A ( $p \approx 1$ ), so the mutation input is approximately  $\mu$ .

### 3 Calculate Removal by Selection

For a recessive deleterious allele, selection only acts against the aa homozygotes.

Frequency of aa genotypes =  $q^2$

Each aa individual has fitness  $(1 - s)$

Selection removes a fraction  $s$  of aa individuals

Each removed aa individual carries 2 a alleles

But we need the change in allele frequency, not individual count

Using the formula for selection against recessives:

$$\Delta q_{\text{sel}} = -sq^2$$

**Note:** The negative sign indicates that selection decreases the frequency of the deleterious allele.

### 4 Set Up the Equilibrium Equation

At equilibrium, the net change in  $q$  is zero:

$$\begin{aligned}\Delta q_{\text{mut}} + \Delta q_{\text{sel}} &= 0 \\ \mu - sq^2 &= 0\end{aligned}$$

**Critical Insight:** This simple equation captures the essence of mutation-selection balance. Mutation adds a alleles at rate  $\mu$ , while selection removes them at rate  $sq^2$ .

### 5 Solve for Equilibrium Frequency

Now we solve the equilibrium equation for  $q$ :

$$\mu - sq^2 = 0$$

$$sq^2 = \mu$$

$$q^2 = \mu/s$$

$$\hat{q} = \sqrt{(\mu/s)}$$

**Beautiful Result:** The equilibrium frequency depends on the square root of the ratio between mutation rate and selection coefficient. This makes intuitive sense - it's a balance between two opposing forces.

## 6 Interpret the Result

Let's examine what  $\hat{q} = \sqrt{(\mu/s)}$  tells us:

- If  $\mu$  increases  $\rightarrow \hat{q}$  increases (more mutation input)
- If  $s$  increases  $\rightarrow \hat{q}$  decreases (stronger selection)
- The square root means changes are dampened
- If  $\mu = s$ , then  $\hat{q} = 1$  (but this rarely happens)

**Biological Significance:** This explains why recessive genetic disorders persist in populations. Even with strong selection ( $s \approx 1$ ), they maintain equilibrium frequencies determined by their mutation rates.

## 7 Numerical Example

Let's apply the formula to a real scenario:

For cystic fibrosis:

$\mu \approx 1 \times 10^{-6}$  (estimated mutation rate)

$s = 1$  (lethal recessive)

$\hat{q} = \sqrt{(0.000001 / 1)} = \sqrt{0.000001} = 0.001$

$$\text{Carrier frequency} = 2pq \approx 2 \times 0.001 = 0.002 \text{ (1 in 500)}$$

**Note:** The actual carrier frequency for cystic fibrosis in European populations is about 1 in 25, suggesting either higher mutation rates or historical heterozygote advantage.

## 8 Comparison with Other Inheritance Patterns

It's instructive to compare with other cases:

**Dominant deleterious:**  $\hat{q} = \mu/s$

**Additive:**  $\hat{q} = \mu/(hs)$  where  $h$  is dominance coefficient

**Recessive:**  $\hat{q} = \sqrt{(\mu/s)}$  ← We derived this!

**Key Difference:** Recessive deleterious alleles reach much higher equilibrium frequencies because selection only acts against homozygotes.

## 9 Limitations and Refinements

Our derivation made several simplifying assumptions:

- Infinite population size
- Constant mutation rate
- No reverse mutation
- No migration
- Random mating
- Selection only on viability

**Real-world complexities:** In actual populations, genetic drift, fluctuating environments, and other factors modify these predictions. However, the basic  $\hat{q} = \sqrt{(\mu/s)}$  provides an excellent starting point.

## 10 Practice Derivation

Now try deriving the equilibrium for a dominant deleterious allele:

**Given:** Selection acts against both Aa and aa genotypes

**Find:** Show that  $\hat{q} = \mu/s$

**Hint:** For dominants,  $\Delta q_{\text{sel}} = -sq$

Your derivation here...

## Summary of Key Points

- Mutation-selection balance occurs when input equals removal
- For recessives:  $\Delta q_{\text{mut}} = \mu$ ,  $\Delta q_{\text{sel}} = -sq^2$
- Equilibrium:  $\mu - sq^2 = 0 \rightarrow \hat{q} = \sqrt{(\mu/s)}$
- The square root reflects that selection only acts against homozygotes
- This explains persistence of recessive genetic disorders