

## ALLELE AND PHENOTYPE FREQUENCIES IN ROCK POCKET MOUSE POPULATIONS

### INTRODUCTION

The tiny rock pocket mouse weighs just 15 grams, about as much as a handful of paperclips. A typical rock pocket mouse is 172 millimeters long from nose to rump, which is shorter than an average pencil. Its impact on science, however, has been enormous. What’s so special about this little mouse?

Populations of rock pocket mice are found all over the Sonoran Desert in the southwestern United States. Two varieties occur widely in the area—a light-colored variety and a dark-colored variety. Similarly, there are two major colors of substrate, or surface materials, which make up the rocky desert floor. Most of the desert landscape consists of light-colored sand and granite. Here and there, however, separated by several kilometers of light-colored substrate, are patches of dark volcanic rocks that formed from cooling lava. These areas of dark volcanic rock range in age from one thousand to more than one million years old.

Dr. Michael Nachman of the University of Arizona and his colleagues have spent many years researching the genetics of fur color in rock pocket mice. In particular, they were interested in understanding the forces that shape genetic variation in natural populations.

Investigating the adaptive value of different coat colors in rock pocket mice is an example of how scientists are attempting to connect genotype with phenotype for fitness-related traits. In this type of research, investigators try to find the underlying gene or genes for a given adaptation. Examples of other fitness-related traits currently being investigated are resistance to the pesticide warfarin in rats, tolerance to heavy metals in plants, and antibiotic resistance in bacteria.

### MATERIALS

calculator

a computer and the Selection Coefficient Excel file found under “Survival of the Fittest—Battling Beetles” at <http://www.biointeractive.org/activities/index.html>

### PROCEDURE

1. Watch the short film, *The Making of the Fittest: Natural Selection and Adaptation*. As you watch, record the following information:

a. What specific trait was studied in this investigation?

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b. How does this trait affect the survival of the mice in different environments?

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c. What is the genetic basis of the trait?

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2. After watching the film, complete Parts 1–3 that follow.

**PART 1: REVIEWING THE PRINCIPLES OF HARDY-WEINBERG**

The genetic definition of evolution is a change to a population’s gene pool. Gene pool is defined as the total number of alleles present in a population at any given point in time. According to the Hardy-Weinberg theorem, a population is in equilibrium (and is therefore *not* evolving) when all of the following conditions are true:

1. The population is very large and well mixed.
2. There is no migration.
3. There are no mutations.
4. Mating is random.
5. There is no natural selection.

To determine whether a population’s gene pool is changing, we need to be able to calculate allelic frequencies. Suppose, for example, a gene has two alleles, *A* and *a*. Each individual has one of three genotypes: *AA*, *Aa*, or *aa*. If the population is in equilibrium, the overall number of *A* alleles and *a* alleles in the gene pool will remain constant as will the proportion of the population with each genotype. If allele frequencies or genotype frequencies change over time, then evolution is occurring.

Two equations are used to calculate the frequency of alleles in a population, where **p** represents the frequency of the dominant allele and **q** represents the frequency of the recessive allele:

**$p + q = 1.0$**

and

**$p^2 + 2pq + q^2 = 1.0$**

The first equation says that if there are only two alleles for a gene, one dominant and one recessive, then 100% of the alleles are either dominant (*p*) or recessive (*q*).

- The second equation says that 100% of individuals in the population will have one of these genotypes: *AA*, *Aa*, and *aa*. Let’s look at each of these one by one to understand the equation:
- If *p* represents the frequency of the *A* allele, then the frequency of the genotype *AA* will be (*p*)×(*p*) or *p*<sup>2</sup>.
- If *q* represents the frequency of the *a* allele, then the frequency of the genotype *aa* will be (*q*)×(*q*) or *q*<sup>2</sup>.
- For heterozygotes, we must allow for either the mother or the father to contribute the dominant and recessive alleles. You can think of it as allowing for both genotypes *Aa* and *aA*. So, the frequency of the heterozygous genotype is calculated as 2*pq*.

In rock pocket mice, several genes code for fur color. Each of those genes have several possible alleles. That’s why there is a range of fur color from very dark to light. For simplicity, we will work with two alleles at one gene. The allele for dark-colored fur (*D*) is dominant to the allele for light-colored fur (*d*). In this scenario, individual rock pocket mice can have one of three genotypes and one of two phenotypes, as summarized in the table below.

	<b>Genotype</b>	<b>Phenotype</b>
Homozygous dominant	<i>DD</i>	Dark
Heterozygous	<i>Dd</i>	Dark
Homozygous recessive	<i>dd</i>	Light

So, applying Hardy-Weinberg:

- p** = the frequency of the dominant allele (*D*)
- q** = the frequency of the recessive allele (*d*)
- p<sup>2</sup>** = the frequency of *DD*
- 2pq** = the frequency of *Dd*
- q<sup>2</sup>** = the frequency of *dd*

This can also be expressed as:

(the frequency of the *DD* genotype) + (the frequency of the *Dd* genotype) + (the frequency of the *dd* genotype) = 1

**SAMPLE PROBLEM**

In a hypothetical population consisting of 100 rock pocket mice, there are 81 individuals with light, sandy-colored fur. Their genotype is *dd*. The other 19 individuals are dark colored and have either genotype *DD* or genotype *Dd*.

Find *p* and *q* for this population and calculate the frequency of heterozygous genotypes in the population.

It is easy to calculate *q*<sup>2</sup>.

$q^2 = 81/100 = 0.81$  or 81%

Next, calculate *q*.

$q = \text{square root } (0.81) = 0.9$

Now calculate *p* using the equation  $p + q = 1$

$p + (0.9) = 1$

$p = 0.1$

Now, to calculate the frequency of heterozygous genotypes, we need to calculate  $2pq$ .

$2pq = 2(0.1)(0.9) = 2(0.09)$

$2pq = 0.18$

**QUESTIONS**

1. If there are 12 rock pocket mice with dark-colored fur and four with light-colored fur in a population, what is the value of *q*? Remember that light-colored fur is recessive.

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2. If the frequency of *p* in a population is 60% (0.6), what is the frequency of *q*?

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3. In a population of 1,000 rock pocket mice, 360 have dark-colored fur. The others have light-colored fur. If the population is at Hardy-Weinberg equilibrium, what percentage of mice in the population are homozygous dominant, dark-colored mice?

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**PART 2: APPLYING HARDY-WEINBERG TO ROCK POCKET MOUSE FIELD DATA**

Dr. Nachman and his colleagues collected rock pocket mice across 35-kilometers of the Arizona Sonoran Desert that included both dark, rocky lava outcrops and light rocky granite areas. Substrate color and coat color frequencies were recorded for each location. Each site was separated from any of the others by at least 8 kilometers. A total of 225 mice were trapped. His data is summarized below:

Collecting site	Substrate color	Number of mice	Phenotype	
			Light	Dark
1	Light	6	6	0
2	Light	85	80	5
3	Dark	7	0	7
4	Dark	5	0	5
5	Dark	45	3	42
6	Light	77	34	43

(Source of data: Hoekstra, H, K. Drumm, and M. Nachman. Ecological Genetics of Adaptive Color Polymorphism in Pocket Mice Geographic Variation in Selected And Neutral Genes.2004. *Evolution*, 58(6), pp. 1329-1344.)

**QUESTIONS**

1. Calculate the overall frequencies of light-colored mice and dark-colored mice caught on light-colored substrates.

Frequency = (number of mice of one color)/(total number of mice)

Frequency of light-colored mice \_\_\_\_\_ Frequency of dark-colored mice \_\_\_\_\_

2. Calculate the overall frequencies of light-colored mice and dark mice caught on dark-colored substrates.

Frequency = (number of mice of one color)/(total number of mice)

Frequency of light-colored mice \_\_\_\_\_ Frequency of dark-colored mice \_\_\_\_\_

3. Using the Hardy-Weinberg equation and data from the table above, determine the number of mice with the *DD* and *Dd* genotypes on the rocky, light granite substrate.

Frequency of mice with *dd* genotype on light substrate \_\_\_\_\_

Frequency of mice with *DD* genotype on light substrate \_\_\_\_\_

Frequency of mice with *Dd* genotype on light substrate \_\_\_\_\_

4. Using the Hardy-Weinberg equation and data from the table above, determine the number of mice with the *DD* and *Dd* genotypes on the rocky, dark lava substrate.

Frequency of mice with *dd* genotype on dark substrate \_\_\_\_\_

Frequency of mice with *DD* genotype on dark substrate \_\_\_\_\_

Frequency of mice with *Dd* genotype on dark substrate \_\_\_\_\_

5. Which fur color seems to have the greatest overall selective advantage? Use data collected from both dark-colored and light-colored substrates to support your answer.

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6. According to the film, what environmental change gave a selective advantage for one coat color over another?

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7. In a separate study, 76 rock pocket mice were collected from four different widely-separated areas of dark lava rock. One collecting site was in Arizona. The other three were in New Mexico. Dr. Nachman and colleagues observed no significant differences in the color of the rocks in the four locations sampled. However, the dark-colored mice from the three New Mexico locations were slightly darker than the dark-colored mice from the Arizona population. The entire *Mc1r* gene was sequenced in all 76 of the mice collected.

The mutations responsible for the dark fur color in the Arizona mice were absent from the three different populations of New Mexico mice. No *Mc1r* mutations were associated with dark fur color in the New Mexico populations. These findings suggest that adaptive dark coloration has occurred at least twice in the rock pocket mouse and that these similar phenotypic changes have a different genetic basis. How does this study support the concept that natural selection is not random?

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8. To determine if the rock pocket mouse population is evolving, explain why is it necessary to collect fur color frequency data over a period of many years?

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**PART 3: HARDY-WEINBERG EXTENDED**

The Hardy-Weinberg equations can be adapted to investigate what happens to gene frequencies in a population that is evolving. To do this, it is necessary to introduce a new term, **selection coefficient**. It is defined as the relative advantage or disadvantage of a genotype with respect to survival and reproductive success. It can also be thought of as the relative selection advantage of a specific allele. For example, if there are two alleles present in a population for a particular trait and one allele is 10% more likely to survive than the other allele, then the selection coefficient for that allele is +0.1.

*The Making of the Fittest:*  
*Natural Selection and Adaptation*

**QUESTIONS**

1. Use the Excel spreadsheet to determine how the selection coefficient ( $s$ ) influences the phenotype of future generations. Substitute increasingly large numbers for  $s$ . Record each new value and describe what happens to the frequencies of  $p$  and  $q$  over the next five generations.

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2. Explain how the selection coefficient and natural selection are related.

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3. In areas with primarily dark-colored substrate, dark-colored mice have a selective advantage over light-colored mice. Therefore, mice with one or more copies of the dominant *Mc1r D* allele have a selective advantage over mice with two copies of the *Mc1r d* allele.

In the film, Dr. Sean Carroll says that with a 1% selection advantage, it takes 1000 years for 95% of the mice to have the dominant phenotype. With a 10% selection advantage, it would take just 100 years. Use the Excel spreadsheet to verify these facts.

a. Find out how many generations following the first appearance of a dark-colored mutation it would take for 95% of the mice to express the dominant dark-colored phenotype given a 1% advantage ( $s=0.01$ ). Rock pocket mice have approximately one litter of pups a year, so the number of generations will be equal to the number of years. You will not be able to use the graph on the tab called "main page" since it only goes up to 100 generations. So, you will need to look at column D of the worksheet called "main worksheet." Scroll down until the value is greater than 0.95. Record your answer below.

It would take about \_\_\_\_\_ generations.

b. Repeat the process for a 10% advantage ( $s=0.1$ ).

It would take about \_\_\_\_\_ generations.

c. What would the selection coefficient need to be for 95% of the mice to have the dominant phenotype in just 50 years? Record your answer below.

The coefficient would need to be about \_\_\_\_\_.

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