**Breeding Bunnies Lab** **Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Period: \_\_\_\_\_\_\_**

**Procedure:**

1. The brown beans represent the *F* allele, the **dominant** trait for **fur**.
2. The white beans represent the *f* allele, the **recessive** trait for **no fur.**
3. Place 50 brown beans and 50 white beans into the Breeding Grounds cup and shake them up.
4. Close your eyes and remove two beans from the cup. If you draw two brown beans, place them in the *FF* circle on your **Breeding Placemat**. If you draw a brown and a white bean, place them in the *Ff* circle. If you draw two white beans, place them in the *ff* circle.
5. *Ff* and *FF* bunnies are born with fur and are able to survive cold winters, while *ff* bunnies are born without fur and do not survive long enough to reproduce. Use the table below as a reference.

|  |  |  |  |
| --- | --- | --- | --- |
| **Beans Removed** | **Result** | **Genotype** | **Description** |
| Two brown beans | ***FF*** | Fur | Homozygous dominant |
| Two white beans | ***ff*** | No fur | Homozygous recessive |
| One brown / one white | ***Ff*** | Fur | Heterozygous |

Question: Why are Ff bunnies born with fur even though they have one recessive allele?

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1. Continue drawing beans, two at a time, until the Breeding Grounds cup is empty. Each pair of beans you draw represents one bunny.
2. Once you have placed all 50 pairs of beans in the correct circles, record your data in the **Breeding Results Data Table** for Generation 1.

Hint: To calculate the number of *F* and *f* alleles, simply count the number of brown (*F*) beans and the number of white (*f*) beans. For Generation 1, the total number of alleles should equal 100 because 50 bunnies were born with 2 alleles each.
3. Place the alleles of the surviving bunnies (*FF* and *Ff*) back into your Breeding Grounds cup.
4. Place the alleles of the deceased bunnies (*ff*) into the Graveyard cup.
These beans will remain there for the rest of the lab.
5. Repeat steps 4 through 10 until you reach 10 generations of bunnies.
6. Calculate the gene frequencies for the *F* and *f* alleles for each generation.
7. Divide the number of *F* alleles by the total number of alleles. Use decimal form to record your frequency.
8. Repeat for the *f* alleles.
9. For Generation 1, the allele frequencies of *F* and *f* alleles will be 0.5 (50 divided by 100). Repeat for all 10 generations.

Question: Does the total number of *F* alleles ever change? Why do you think this is?

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**Breeding Bunnies Lab- Data Chart Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Period: \_\_\_\_\_\_\_**

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Generation | Number of *FF* Bunnies | Number of *Ff* Bunnies | Number of *ff* Bunnies | Number of *F* Alleles | Number of *f* Alleles | Total Number of Alleles | Gene Frequency of *F* (decimal) | Gene Frequency of *f* (decimal) |
| 1 |  |  |  |  |  |  |  |  |
| 2 |  |  |  |  |  |  |  |  |
| 3 |  |  |  |  |  |  |  |  |
| 4 |  |  |  |  |  |  |  |  |
| 5 |  |  |  |  |  |  |  |  |
| 6 |  |  |  |  |  |  |  |  |
| 7 |  |  |  |  |  |  |  |  |
| 8 |  |  |  |  |  |  |  |  |
| 9 |  |  |  |  |  |  |  |  |
| 10 |  |  |  |  |  |  |  |  |

**Breeding Bunnies Lab- Graph Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Period: \_\_\_\_\_\_\_**

Gene Frequency



**Legend**

***F* allele**

**Frequency**

***f* allele**

**Frequency**

Generation #

**Gene Frequencies of *F* and *f* Alleles**

**Bunnies Breeding Lab- Conclusion Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_ Period: \_\_\_\_\_\_\_**

**Conclusion:** How does natural selection affect the frequency of undesirable or deadly genes in a population over time?

**Claim**:\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

**Evidence**:\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

**Reasoning**:\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_

**Reflections:**

1. What happens to the frequency of the *F* allele as the number of *F* alleles changes?

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1. Do you think all of the groups in your class got the same numbers as you in their data tables? Why or why not?

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1. In this experiment, it is assumed that new bunnies do not enter the population and that bunnies do not leave the population except by dying. It is also assumed that bunnies choose mates randomly. Is this how real-life populations of organisms work? How could this experiment be changed to simulate a real-life population of bunnies?

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1. What are some other traits in a population of bunnies in a cold, snowy environment that could be considered undesirable?

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1. In a real-life scenario involving these bunnies, what would happen to the recessive *f* alleles over time? Why or why wouldn’t the *f* alleles be eliminated completely?

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