

# Cow Exercise 1 - Answer Key

#### Goal

In this exercise, you will use StarGenetics, a software tool that simulates mating experiments, to analyze the nature and mode of inheritance of specific genetic traits.

## Learning Objectives

After completing this exercise, you will be able to:

- 1. Identify the phenotype of a cow within StarGenetics.
- 2. Determine whether a phenotype is dominant or recessive relative to another phenotype through the analysis of results from genetic crosses.
- 3. Use Punnett Squares to predict and confirm expected genotypic and phenotypic ratios.
- 4. Explain the effect of sample size on observed phenotypic ratios.
- 5. Infer and assign genotypes of individual organisms using proper nomenclature of alleles.
- 6. Determine whether an allele is autosomal or sex-linked through the analysis of results from genetic crosses.

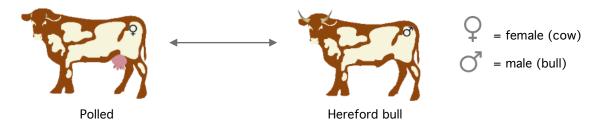
## Getting started with StarGenetics

- To get StarGenetics, please navigate to: http://star.mit.edu/genetics/.
- Click on the **Start** button to launch the application.
- Click **Trust** when a prompt appears asking if you trust the certificate.
- Click on File > New in the main menu.
- Click on the Cow Exercise 1 file.

In the **Strains** box, you will find two Hereford cows, called **Hereford cow** and **Hereford bull**, and one cow that does not look like the others, called **Polled**. The Hereford cows are homozygous (true-breeding) for the Polled allele.

#### 1 What is the sex of the Polled cow?

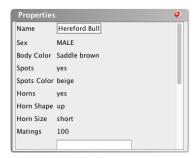
• The sex symbol is indicated within the body of each cow.



#### **Answer**

# **2** What is the phenotype of the Polled cow?

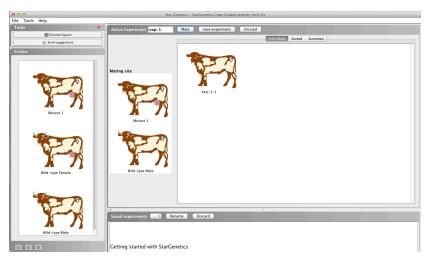
• Compare the color and features of the cows to determine which features are different. Hereford cows and bulls have brown bodies, beige spots, and horns. You can also click on a cow and look in the **Properties** box.



#### **Answer**

- **3** Is the Polled phenotype recessive or dominant to the Hereford phenotype?
  - Mate the Polled cow with a Hereford bull by dragging both cows into the Mating site.
  - Click on the Mate button.

What do you see? Let's take a look:



You can see all the cows resulting from this cross within the **Individual** tab or a summary of the experiment by clicking on the **Summary** tab. Within the **Summary** tab, all of the organisms are organized by <u>phenotype</u>.

a) How many progeny did you obtain? What phenotype(s) do you observe?

## Answer

**b)** Fill in the Punnett Squares below with the expected results in the F1 generation for each of the following situations:

If the Polled phenotype is dominant to the Hereford phenotype:

- 1. What results would you expect if the Polled cow is homozygous for the Polled allele?
- 2. What results would you expect if the Polled cow is heterozygous for the Polled allele?

If the Polled phenotype is recessive to the Hereford phenotype:

3. What results would you expect if the Polled cow is homozygous for the Polled allele?

Hint: Remember that the Hereford cows are homozygous (true-breeding) for the Polled allele.

1. Homozygous	s Dominant	2. Heteroz	zygous	3. Homozygous	recessive	
	Hereford		Hereford		Hereford	
Polled		Polled		Polled		
Expected Genotypic Ratio: Expected Phenotypic Ratio:						
<b>c)</b> Can you conclude whether the Polled phenotype is dominant or recessive relative to the Hereford phenotype after analyzing the progeny that you obtained? Explain.						
Answer						
d) Increase the number of Polled phenotype domina  • To add additional off appropriate number of the appropriate number.  • To start a new exper	int or recessive spring to a cros of matings (or c ber of additiona	to the Hereford phe is, click on the Add rosses) you would li I offspring. Repeat u	more matings ike to perform in intil you have at	button and select t the pop-up window least 12 cows.	he to add	
automatically saved	for you and can	be found in the Sav	ved Experiment	ts panel.		
Answer						
e) Based on your results	, what is the ge	notype of the Polled	d cow?			
Answer						

**4 a)** Can a sex-linked allele produce the results that you obtained in question 3? Fill in the Punnett Square below with the expected results in the case that the Polled allele is sex-linked.

The hallmark of a sex-linked allele is differential partitioning of the allele in females and males. Males have one X chromosome and, as a result, carry one copy of each sex-linked allele. On the other hand, females have two X chromosomes and, as a result, carry two copies of each sex-linked allele.

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Here	ford					
Polled						
Expected Genotypic Ra	atio:					
Expected Phenotypic F	Ratio:					
Answer						
<b>b)</b> To distinguish between the sex-linked and autosomal possibilities, cross a F1 Polled Cow with a F1 Polled Bull. Fill in the Punnett Squares below with the expected progeny from this cross if the Polled allele is sex-linked or autosomal. Indicate the expected genotypic and phenotypic ratios you would obtain in each case.						
Sex-Linked Autosomal		somal				
	F1 Polled Cow		F1 Polled Cow			
F1 Polled Male		F1 Polled Male				
Expected Genotypic Ratio: Expected Phenotypic Ratio:		<u> </u>				
c) Perform the F1 x F1 cross in part (b) in StarGenetics. How many progeny did you observe in each of the phenotypic classes? Can you conclude whether the Polled allele is autosomal or sex-linked? Explain.						
Answer						
Hereford						
Hereford males Polled females						
Polled remales Polled males						
Conclusions:						

**d)** Now generate more progeny for your cross. Does your answer change as you increase the number of progeny in your cross? Fill in your results below. Is the Polled allele autosomal or sex-linked? Explain.

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Answer
Hereford females
Hereford males
Polled females
Polled males
Conclusions:
BONUS QUESTION:
Can you think of another informative cross to distinguish between the sex-linked and autosomal possibilities? This cross must be different from the F1 Polled Cow x F1 Polled Bull cross in Question 4b. Specify your informative cross and how it enables you to distinguish between the two possibilities. Then, perform your cross in StarGenetics, and in the answer box, describe your results and explain how your results led you to your conclusion.
Answer

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