

StarGenetics Tutorial 1

This tutorial is designed to illustrate how to analyze genetic data using the StarGenetics software tool.

Learning Objectives

During this tutorial, two questions will be answered:

- How do we know if an allele is dominant or recessive?
- How do we know if an allele is autosomal or sex-linked?

Getting started with StarGenetics

- To get to StarGenetics, please navigate to: http://web.mit.edu/star/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 on the main menu.
- Click on the StarGenetics Tutorial 1 file.

When you first open this tutorial, you will see one mutant fly, called **Mutant 1**, and two wild type flies (female and male) within the **Strains** box.

1 What is the <u>sex</u> of the mutant fly?

Let's take a closer look at the mutant fly and compare it with one of the wild type flies.



The sex symbol is indicated within the body of each fly. For example, the wild type, indicated above, is female and the mutant fly is male (see sex symbol legend above).

2 What is the phenotype of the mutant fly?

Wild type flies have red eyes, brown bodies and full size antennae and wings. If you compare the color and features of the wild type and Mutant 1 fly, you will see that both have red eyes, full size antennae and wings. The one exception is body color. Mutant 1's body color is grey instead of brown (wild type body color). You can also obtain the same information by clicking on the fly and looking at the **Properties** window. The characteristics that differ from wild type are indicated within the **Properties** box as well as some other relevant information (sex, number of matings, etc).



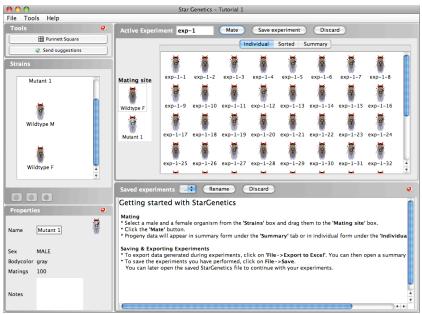


3 Next, we will address the first learning objective of this tutorial: *is the allele that confers Mutant* 1's grey body color recessive or dominant?

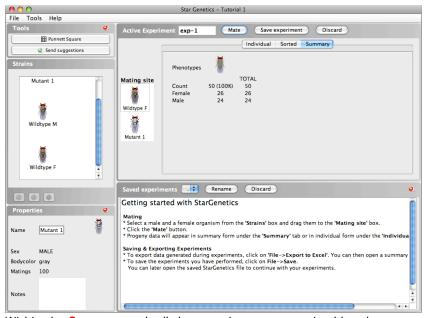
The easiest way to answer this question is to mate the mutant fly in question, Mutant 1, with a wild type fly. Let's set up this mating:

- Drag both the Mutant 1 and the female wild type fly into the Mating site.
- Click on the Mate button.

What do you see?



You can see all the flies resulting from this cross within the **Individual** tab or you can see a summary of the experiment by clicking on the **Summary** tab.

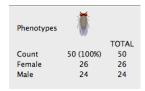


Within the **Summary** tab all the organisms are organized by <u>phenotypes</u>.



How many phenotypes do you see?

Only one phenotype is present in the F1 progeny. This phenotype resembles the wild type parent.



You can see that there are a total of 50 progeny: 21 female and 29 male flies.

What do these results tell you about the gray body color allele?

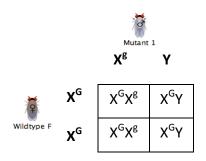
If all the flies have the phenotype of the wild type parent, then this means that the gray body color allele is recessive to wild type.

4 We will now address the second learning objective of this tutorial: *is the allele that confers Mutant* 1's grey body color autosomal or X-linked?

From the first cross, it seems like the grey body color allele might be autosomal. However, from this one cross it is not possible to make this conclusion.

Can a sex-linked grey body allele lead to the results obtained in our first cross?

Since we know that the grey body color is recessive, we can discern that if the allele was sex-linked, then the genotype would be as follows:



Expected genotypic ratio:

2 XGXg: 2 XGY

Expected phenotypic ratio:

All female and male progeny would be wild type

The hallmark of a X-linked allele is differential partitioning of the allele in female versus males, since males only carry one copy of each allele. To determine if this is the case, we will need to perform one of the two possible informative crosses.

POSSIBLE CROSS 1: One possible cross is an F1 female to an F1 male. These are the results we obtain when we perform this cross:

Phenotypes	₩	***************************************	
			TOTAL
Count	10 (20%)	40 (80%)	50
Female	0	22	22
Male	10	18	28



We observe a 1:4 ratio of gray body flies to wild type flies. All the gray body flies are male. This differential partitioning of the gray body color allele indicates that the gray body color gene is X-linked. Here is the Punnett Square for this cross (F1 female X F1 male) if we assume that the gray body color allele is X-linked.

X^G Y

 \mathbf{X}^{G}

 $\mathbf{X}^{\mathbf{g}}$

X ^G X ^G	X ^G Y
X^GX^g	X ^g Y

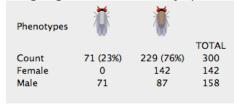
F1 male: X^GY F1 female: X^GX^g

Expected genotypic ratio:

1 X^GX^G: 1 X^GX^g: 1 X^GY: 1 X^gY <u>Expected phenotypic ratio</u>: 3 wild type: 1 gray body color

All females are wild type; 1:1 ratio of gray body male flies to wild type male flies.

The ratio of wild type to gray body flies we observe with 50 F2 flies is 4:1 and not 3:1 as predicted, respectively. An increase in the number of progeny from 50 to 300 reveals a ratio that is closer to 3:1. Using larger data sets is always preferable to increase the statistical significance of your conclusions!



As previously predicted and observed when we performed this experiment with 50 F2 flies, half of the male files have gray bodies and half of them have wild type body color. Together these results indicate that the gray body color gene is X-linked.

POSSIBLE CROSS 2: Mating two F1 flies is not the only possible cross we could have used to test whether the gray body gene is autosomal or X-linked. Mating of a F1 female to one of the original wild type male parents, will also provide the necessary information to discern between autosomal or X-linkage mode of inheritance.

X^G Y

 \mathbf{X}^{G}

Xg

X ^G X ^G	X ^G Y
X ^G X ^g	X ^g Y

Wild type male (provided in the Strain box): X^GY

F1 female: X^GX^g

Expected genotypic ratio:

1 X^GX^G: 1 X^GX^g: 1 X^GY: 1 X^gY <u>Expected phenotypic ratio</u>:

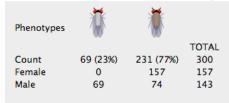
3:1 ratio of wild type to gray body color

Ver. 6



All females are wild type; 1:1 ratio of wild type to gray body color ratio for male flies.

The results we obtain match our expected results:



Therefore, either crossing a F1 female to a F1 male or crossing a F1 female to the original wild type male will lead you to the same conclusion: the gray body color gene is X-linked.

Your approach to performing the crosses for this type of genetic analysis is up to you. The important steps are 1) to evaluate your initial experimental results, 2) to construct a hypothesis that can explain your results (in this case by using Punnett Square analysis), and then 3) to test your hypothesis by performing additional experiments.

Hope you enjoyed this tutorial!