

## LABORATORY EXERCISE 2. POPULATION EFFECTS

**Exercise Protocol****Part 2**

You may use the same trait as in Part 1 or you may choose a different one to examine the effect of offspring population size on observed phenotypic ratios. However, use the same trait for all progeny numbers.

**STEPS FOR THIS EXERCISE**

1. Return to the **FLY COLONY** program on Canvas.
2. Select your mutant phenotype in one of the drop down boxes on the Parental screen.
3. Mate the flies through the F2 generation using FOUR DIFFERENT OFFSPRING numbers:
  - a. 5000 offspring
  - b. 500
  - c. 50
  - d. 10
4. Record the results from each set of matings on the DATA SHEETS provided.
5. Answer the Exercise 2 Questions below.

Predict what the Expected ration of each cross will yield and enter in the boxes below. Then look at your total progeny numbers from the cross. Divide the number of dominant offspring by the number of recessive offspring. This is your Observed phenotypic ratio (round to the nearest decimal). Enter your observed ratio for each population below.

Expected Ratio		Observed Ratio
<input type="text"/>	5000	<input type="text"/>
<input type="text"/>	500	<input type="text"/>
<input type="text"/>	50	<input type="text"/>
<input type="text"/>	10	<input type="text"/>

Describe what you see happening to the Observed numbers versus the Expected ratios. Reflect on WHY you may be seeing something occurring.

# GENETIC CROSS

## DATA SHEET (Exercise 2) – 5000 Offspring

NAME:

TRAIT:  PHENOTYPES CROSSED:  Male  Female  X

### CROSS DIAGRAM

Parentals

	Male		Female
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

F1 Results

	Male		Female
Phenotype	<input type="text"/>		<input type="text"/>
Genotype	<input type="text"/>		<input type="text"/>

Parents for F2

	Male		Female
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

F2 Results

Predicted Segregation Ratio =  phenotype  ratio :  phenotype  ratio

Gender	Phenotype	Genotype	Exp Number	Obs Number
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

### Chi-Squared Test (enter values from F2 Generation Page, combine sexes to one phenotype)

Phenotype	Observed	Expected	O - E	( O - E ) <sup>2</sup>	( O - E ) <sup>2</sup> / E
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
TOTAL	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

Observed Chi – Squared Value =

Degrees of Freedom (df) =

Table Value (0.05) =

Overall Conclusion =

CONCLUSIONS:

# GENETIC CROSS

## DATA SHEET (Exercise 2) – 500 Offspring

NAME:

TRAIT:  PHENOTYPES CROSSED:  Male  Female  X

### CROSS DIAGRAM

**Parentals**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

**F1 Results**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>		<input type="text"/>
Genotype	<input type="text"/>		<input type="text"/>

**Parents for F2**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

**F2 Results** *Predicted Segregation Ratio* =  phenotype  ratio :  phenotype  ratio

Gender	Phenotype	Genotype	Exp Number	Obs Number
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

### Chi-Squared Test (enter values from F2 Generation Page, combine sexes to one phenotype)

Phenotype	Observed	Expected	O - E	( O - E ) <sup>2</sup>	( O - E ) <sup>2</sup> / E
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<b>TOTAL</b>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

Observed Chi – Squared Value =

Degrees of Freedom (*df*) =

Table Value (0.05) =

Overall Conclusion =

CONCLUSIONS:

# GENETIC CROSS

## DATA SHEET (Exercise 2) – 50 Offspring

NAME:

TRAIT:  PHENOTYPES CROSSED:  Male  Female  X

### CROSS DIAGRAM

Parentals

	Male		Female
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

F1 Results

	Male		Female
Phenotype	<input type="text"/>		<input type="text"/>
Genotype	<input type="text"/>		<input type="text"/>

Parents for F2

	Male		Female
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

F2 Results

Predicted Segregation Ratio =  phenotype  ratio :  phenotype  ratio

Gender	Phenotype	Genotype	Exp Number	Obs Number
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

### Chi-Squared Test (enter values from F2 Generation Page, combine sexes to one phenotype)

Phenotype	Observed	Expected	O - E	( O - E ) <sup>2</sup>	( O - E ) <sup>2</sup> / E
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
TOTAL	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

Observed Chi – Squared Value =

Degrees of Freedom (df) =

Table Value (0.05) =

Overall Conclusion =

CONCLUSIONS:

# GENETIC CROSS

## DATA SHEET (Exercise 2) – 10 Offspring

NAME:

TRAIT:  PHENOTYPES CROSSED:  Male  Female  X

### CROSS DIAGRAM

**Parentals**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

**F1 Results**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>		<input type="text"/>
Genotype	<input type="text"/>		<input type="text"/>

**Parents for F2**

	<b>Male</b>		<b>Female</b>
Phenotype	<input type="text"/>	x	<input type="text"/>
Genotype	<input type="text"/>	x	<input type="text"/>

**F2 Results** *Predicted Segregation Ratio* =  phenotype  ratio :  phenotype  ratio

Gender	Phenotype	Genotype	Exp Number	Obs Number
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

### Chi-Squared Test (enter values from F2 Generation Page, combine sexes to one phenotype)

Phenotype	Observed	Expected	O - E	( O - E ) <sup>2</sup>	( O - E ) <sup>2</sup> / E
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>
<b>TOTAL</b>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

Observed Chi – Squared Value =

Degrees of Freedom (*df*) =

Table Value (0.05) =

Overall Conclusion =

CONCLUSIONS: