## SI Genetics- Full Discipline Demo

### Patterns of Inheritance

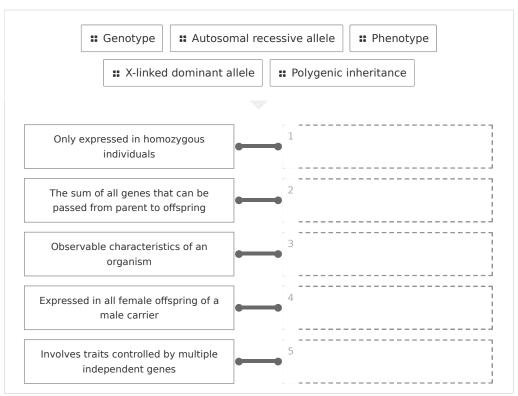
### Final Report - Answer Guide

InstitutionScience Interactive UniversitySessionSI Genetics - Full Discipline DemoCourseSI Genetics- Full Discipline Demo

**Instructor** Sales SI Demo

### Test Your Knowledge

#### Match each term with the best description.



#### Correct answers:

- 1 Autosomal recessive allele 2 Genotype 3 Phenotype
- 4 X-linked dominant allele 5 Polygenic inheritance



### Identify each statement as true or false.

# A p-value is used to det	termine which traits are dominant.
	:
	alysis that compares observed data to the
expected results to determine the asso	ociation between variables.
	:
A Punnett square for diagramming two consists of 20 cells.	independent, dominant/recessive traits
# Punnett squares diagram the poss	sible genotypes and phenotypes of offspring.
True	False
1	
orrect answers:	
1	
A chi-square (X <sup>2</sup> ) test is a statistical are expected results to determine the asset	nalysis that compares observed data to the ociation between variables.

Punnett squares diagram the possible genotypes and phenotypes of offspring.

2 A p-value is used to determine which traits are dominant.

A Punnett square for diagramming two independent, dominant/recessive traits consists of 20 cells.

### Exploration

Offspring inherit one allele from each parent in many organisms.

<ul><li>True</li></ul>			•
<ul><li>False</li></ul>			



controlled by a single gene with allele(s).	idits
one one	
• two	<b>✓</b>
• three	
four	
Male offspring expressing X-linked recessive alleles can only inherit condition from their female parent.	the
	<b>✓</b>
■ False	
Polygenic traits in humans include	
<ul><li>eye color</li></ul>	
skin color	
height	
All of the above	✓
describes a pattern of inheritance whereby heterozygous phenodisplay both alleles.	otypes
Simple dominance	
X-linked dominance	
<ul> <li>Incomplete dominance</li> </ul>	
<ul><li>Codominance</li></ul>	<b>~</b>
Only the phenotypic ratios of offspring can be determined from Punisquares.	nett
○ True	



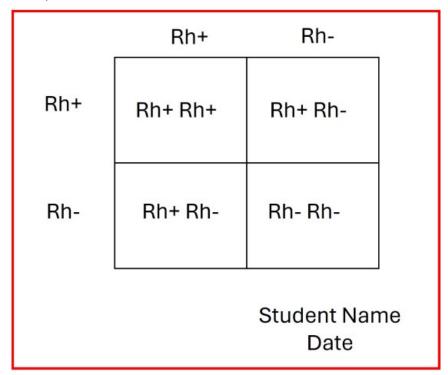
A chi-square test can be used to compare the predictions of a Punnett square to what is observed in a population.
○ True
False
Exercise 1
How do simple dominant/recessive inheritance patterns differ between autosomal and X-linked traits? How were these differences observed in the results of this exercise?
Simple autosomal dominant/recessive inheritance patterns involve traits controlled by a single gene with two alleles, one of which is completely dominant to the other, that are not influenced by the sex of the parent or offspring. Autosomal dominant alleles are expressed in the phenotype of both homozygous and heterozygous individuals. Autosomal recessive alleles are only expressed in the phenotype of homozygous individuals. X-linked dominant alleles are expressed in all female offspring of a male carrier and all male and female offspring receiving the dominant allele from a female carrier. X-linked recessive alleles are expressed in all male offspring since they only have one copy, but the alleles are only expressed in female offspring that are homozygous for the trait.  The results from the crosses in this exercise support the differences in these inheritance patterns produced by crosses between heterozygous parents. In the autosomal trait, 25% of all offspring exhibited the Rh- phenotype, whereas only male offspring inheriting the color-blindness allele from their mother exhibited the color-blind phenotype.
Were the Punnett squares created in this exercise a good predictor of offspring phenotypes resulting from the crosses performed using the digital genetics bench? Reference Photos 1 and 2 and the results recorded in Data Tables 3 and 6 in your explanation.
Note to instructors: the digital genetics bench produces unique results between students, but when sufficient offspring are generated, results should agree with Punnett square predictions. For this reason, student results and answers will vary. All answers should be supported by Data Tables 3 and 6.
Yes, both Punnett squares in Photos 1 and 2 were a good predictor of the inheritance pattern results produced by the digital genetics bench. Both the autosomal and X-linked dominant/recessive results analyses in Data Tables 3 and 6 produced p values > 0.05 meaning the differences between the expected and observed values were the result of chance and not statistically significant.



Both autosomal and X-linked recessive phenotypes frequently skip generations. Did either of your crosses support this trend? Explain your answer by referencing both the parents and offspring in each of your crosses.

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Photo 1 : Rh Factor Punnett Square (SAMPLE ANSWER BELOW)



Data Table 1: Expected Rh Factor Ratios (SAMPLE ANSWER BELOW)

(S) WHI EE / WO WER BELOW)	
Phenotypic ratio	3:1
Genotypic ratio	1:2:1
Offspring expected to be Rh+ (%)	75
Offspring expected to be Rh- (%)	25

Data Table 2: Rh Factor Results

(SAMPLE ANSWER BELOW)

	Rh+	Rh-	Total
Observed number of individuals	14	4	18
Expected number of individuals	13.5	4.5	

### Panel 1: Chi-square Calculation for Rh Factor Offspring

(SAMPLE ANSWER BELOW)

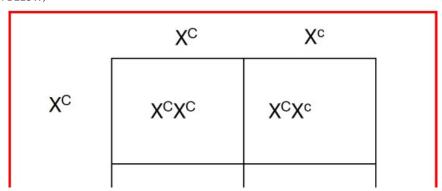
Student answers should use values from Data Table 3.

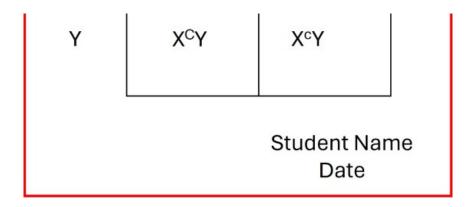
$$x^2 = \frac{(14-13.5)^2}{13.5} + \frac{(4-4.5)^2}{4.5} = 0.074$$

## Data Table 3: Rh Factor Chi-square Test (SAMPLE ANSWER BELOW)

(SAMFLE ANSWER BELOW)	1
Degrees of Freedom	
p-Value Range	.950500
Conclusion	Difference between expected and observed is not statistically significant

#### Photo 2: Color Blindness Punnett Square (SAMPLE ANSWER BELOW)





# Data Table 4: Expected Color Blindness Ratios (SAMPLE ANSWER BELOW)

(SAPITEL ANSWER DELOW)	
Phenotypic ratio (M normal: M color blind: F normal: F color blind)	1:1:2:0
Genotypic ratio (X <sup>C</sup> Y: X <sup>C</sup> Y: X <sup>C</sup> X <sup>C</sup> : X <sup>C</sup> X <sup>C</sup> )	1:1:1:1
Male offspring expected to have normal vision (%)	25
Male offspring expected to be color blind (%)	25
Female offspring expected to have normal vision (%)	50
Female offspring expected to be color blind (%)	0

# Data Table 5: Color Blindness Results

(SAMPLE ANSWER BELOW)	Male normal vision	Male color blind	Female normal vision	Female color blind	Total



Observed number of individuals	10	11	15	0	36
Expected number of individuals	9	9	18	0	

## Panel 2: Chi-square Calculation for Color Blindness Offspring (SAMPLE ANSWER BELOW)

Student answers should use values from Data Table 6.

$$x^2 = \frac{(10-9)^2}{9} + \frac{(11-9)^2}{9} + \frac{(15-18)^2}{18} = 1.06$$

### Data Table 6: Color Blindness Chi-square Test

(SAMPLE ANSWER BELOW)

Degrees of Freedom	2
p-Value Range	.950500
Conclusion	Difference between expected and observed is not statistically significant

### Exercise 2

How do your results for the eye color cross support the characteristics of polygenic inheritance patterns? Reference Graph 1 in your explanation.

Polygenic inheritance patterns involve traits controlled by multiple independent genes that produce an additive effect on the phenotype of the individual. Polygenic inheritance patterns result in a normal bell curve distribution of phenotypes within a population. The results in Graph 1 support this pattern with most of the individuals in the expected population exhibiting green, hazel, and blue green eyes while relatively few individuals have brown or light blue eyes.

Was the Punnett square created for incomplete dominance in this exercise a good predictor of offspring phenotypes resulting from the crosses performed using the digital genetics bench? Reference Photo 4 and the results recorded in Data Table 10 in your explanation.

Note to instructors: the digital genetics bench produces unique results between students, but when sufficient offspring are generated, results should agree with Punnett square predictions. For this reason, student results and answers will vary. All answers should be supported by Data Tables 12.

Yes, the Punnett square in Photo 4 was a good predictor of the hair texture results produced by the digital genetics bench. The analyses in Data Table 10 produced a p value > 0.05 meaning the differences between the expected and observed values were the result of chance and not statistically significant.



How do the ABO blood type alleles exhibit both codominant and dominant/recessive inheritance patterns? Reference your results in Data Table 11 and Photo 5 in your explanation.

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Photo 3: Eye Color Punnett Square (SAMPLE ANSWER BELOW)

J ( V )	vv)					
	$B_1B_2$	$B_1b_2$	$b_1B_2$	$b_1b_2$		
B <sub>1</sub> B <sub>2</sub>	$B_1B_1B_2B_2$	B <sub>1</sub> B <sub>1</sub> B <sub>2</sub> b <sub>2</sub>	B <sub>1</sub> b <sub>1</sub> B <sub>2</sub> B <sub>2</sub>	B <sub>1</sub> b <sub>1</sub> B <sub>2</sub> b <sub>2</sub>		
B <sub>1</sub> b <sub>2</sub>	$B_1B_1B_2b_2$	$B_1B_1b_2b_2$	$B_1b_1B_2b_2$	B <sub>1</sub> b <sub>1</sub> b <sub>2</sub> b <sub>2</sub>		
b <sub>1</sub> B <sub>2</sub>	$B_1b_1B_2B_2$	B <sub>1</sub> b <sub>1</sub> B <sub>2</sub> b <sub>2</sub>	b <sub>1</sub> b <sub>1</sub> B <sub>2</sub> B <sub>2</sub>	b <sub>1</sub> b <sub>1</sub> B <sub>2</sub> b <sub>2</sub>		
b <sub>1</sub> b <sub>2</sub>	$B_1b_1B_2b_2$	B <sub>1</sub> b <sub>1</sub> b <sub>2</sub> b <sub>2</sub>	b <sub>1</sub> b <sub>1</sub> B <sub>2</sub> b <sub>2</sub>	b <sub>1</sub> b <sub>1</sub> b <sub>2</sub> b <sub>2</sub>		
Student Name Date						

Data Table 7: Expected Eye Color Results (SAMPLE ANSWER BELOW)

Phenotypic ratio	1:4:6:4:1
Genotypic ratio	1:2:1:2:4:2:1:2:1
Offspring expected to have brown eyes (%)	6
Offspring expected to have hazel eyes (%)	25
Offspring expected to have green eyes (%)	38
Offspring expected to have blue green eyes (%)	25
Offspring expected to have light blue eyes (%)	6

Graph 1: Polygenic Eye Color Distribution (SAMPLE ANSWER BELOW)

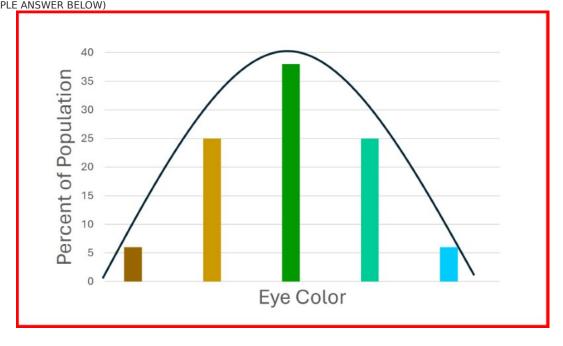


Photo 4: Hair Texture Punnett Square (SAMPLE ANSWER BELOW)

	С	S	
С	CC	CS	
S	CS	SS	
Student Name Date			

## Data Table 8: Expected Incomplete Dominance Results (SAMPLE ANSWER BELOW)

(SAMPLE ANSWER BELOW)	
Phenotypic ratio	1:2:1
Genotypic ratio	1:2:1
Offspring expected to have curly hair (%)	25
Offspring expected to have wavy hair (%)	50
Offspring expected to have straight hair (%)	25

### Data Table 9: Hair Texture Results

(SAMPLE ANSWER BELOW)

	Curly hair	Wavy hair	Straight hair	Total
Observed number of individuals	5	10	3	18
Expected number of individuals	4.5	9	4.5	

# Panel 3: Chi-square Calculation for Hair Texture Offspring $({\sf SAMPLE}\ {\sf ANSWER}\ {\sf BELOW})$

$$x^2 = \frac{(5-4.5)^2}{4.5} + \frac{(10-9)^2}{9} + \frac{(3-4.5)^2}{4.5} = 0.667$$

# Data Table 10: Hair Texture Chi-square Test (SAMPLE ANSWER BELOW)

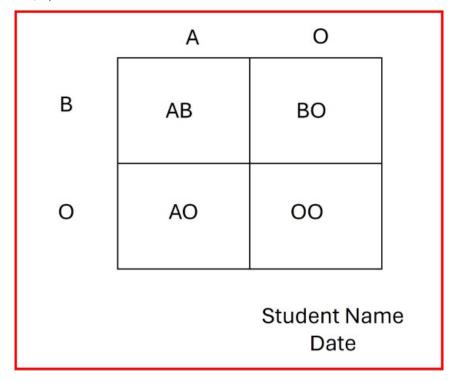
(			
Degrees of Freedom	2		
p-Value Range	0.950-0.500		
Conclusion	$\rm p > 0.05$ there is not a statistically significant difference between the expected and observed values.		



Data Table 11: Blood Type Alleles (SAMPLE ANSWER BELOW)

(SAM) EL ANSWER BELOW)			
	Genotype	Phenotype	
Codominant	AB	Type AB	
Heterozygous dominant (2)	AO, BO	Type A, Type B	
Homozygous recessive	00	Type O	

Photo 5: Blood Type Punnett Square (SAMPLE ANSWER BELOW)



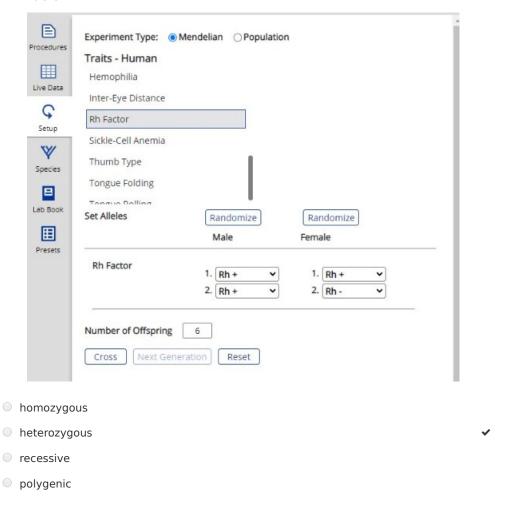
# Competency Review

Alleles are different versions of a gene that produce phenotypic variation in organisms.		
○ True	<b>✓</b>	
• False		
Examples of autosomal recessive genetic diseases include		
sickle cell anemia		
<ul><li>cystic fibrosis</li></ul>		
<ul><li>phenylketonuria</li></ul>		
All of the above	<b>~</b>	
X-linked recessive alleles are most often expressed in male offspring.  True False	*	
Polygenic inheritance patterns result in a(n) distribution of phenotypes within a population.		
<ul><li>sine curve</li></ul>		
o normal bell curve	<b>~</b>	
logarithmic curve		
<ul><li>inverse</li></ul>		

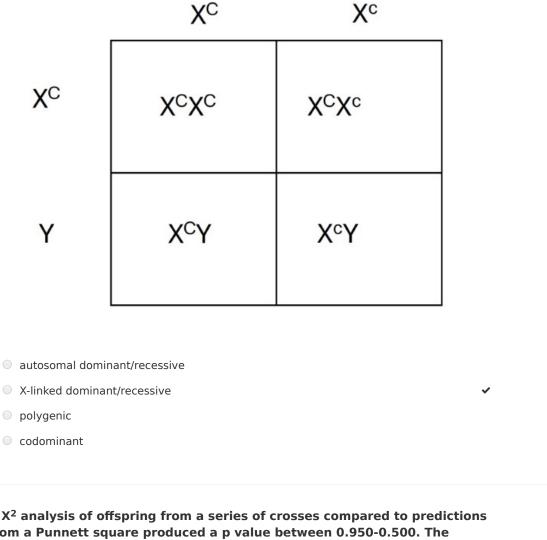


appear intermediate to the two homozygous phenotyp	es for a given anele.
Autosomal dominance	
X-link dominance	
Incomplete dominance	<b>~</b>
<ul><li>Codominance</li></ul>	
The produced by each parent are labeled on the hereical axes of a Punnett square.	orizontal and
<ul><li>alleles</li></ul>	<b>~</b>
genes	
<ul><li>offspring</li></ul>	
phenotypes	
A significant X <sup>2</sup> value implies that the difference found expected and observed data is probably not due to cha	
<ul><li>True</li></ul>	<b>~</b>

The digital genetics bench in the image below depicts a female parent that is \_\_\_\_ for Rh factor.



The Punnett square below is predicting the results of a cross involving alleles.



A X<sup>2</sup> analysis of offspring from a series of crosses compared to predictions from a Punnett square produced a p value between 0.950-0.500. The offspring sample is \_\_\_\_ the expected results.

0

- unrelated to
- significantly different from
- onot significantly different from

### **Extension Questions**

Adam is considering fathering children but has concerns because his brother suffers from the X-linked recessive genetic disease hemophilia A. Adam visits a genetics counselor seeking advice regarding the risk of having children with hemophilia A. Apply your



## knowledge of inheritance patterns to suggest how the genetics counselor should address the Adam's concerns.

(SAMPLE ANSWER BELOW)

The counselor should explain to Adam that X-linked recessive alleles such as those causing hemophilia A are expressed in the phenotypes of all male carriers because they only have one copy of the X allele. Since Adam is healthy, unlike his brother, his only X chromosome does not carry the recessive allele and he cannot transmit the genetic disease to his future children.

