Genetics Problem Set 2 Feb 15, 2010 (concepts from chapter 3, text)

 You are interested in the mode of transmission of coat color in mice. You start by crossing a true breeding gray mouse with a true-breeding white mouse. The offspring are all gray. After letting these gray offspring interbreed you note a total of 114 gray mice and 31 white mice from several litters.

a. Propose a mechanism to explain the results. Diagram the crosses and

compare the observed results with the expected. (indicate parental, F1, and F2 generations; show possible gametes and zygotic combinations) Bucause of the F. (all gray) and F2 (3 gray I white) Approx. ratios, this looks like a genetic trait in which one gene Controls this aspect of fur color. Two allelos exist in which gragis dominant to while. I way while b. Describe the phenotypes and ratios of the F3 generations

WWXWW -> 3 gray: I whill

Gray Gray

3 ww xww -> all while

c. Suggest a molecular mechanism for gray and white fur color.

Since white is nessive the "W" sen may encode to Since white is nessive the "W" sen may encode a content walled allows the pignant walled smehow restricts the making of gray pignant.

In fruit flies (Drosophila melaogaster) a recessive mutant allele w, results in white eyes, whereas the normal, wild-type eye color is red. A second recessive mutant allele, b, results in black body color, whereas the normal, wild-type body color is gray.

a. Diagram a cross of a true-breeding gray-bodied, white-eyed fly with a true-breeding black-bodied, red-eyed fly. p BB ww x bbwl red
Show the Parental and F1 genotypes

b. Show the Parental and F1 genotypes.

c. Use a Punnett square to show F1 gametes and F2 zygotic See nut page combinations.

d. Determine the expected ratio of F2 phenotypes.

Expect 9:3 3 Black/red : black/white

a-b.
P. BB www. X bb WW

(Bw)
F. BENIE

b &	BENG BEWIN BENJACO	B b www.	be War	66 ww .
p M	B6WW	B b. Was	De Miles	
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BW	Bowin	8 B Wee	85 WW Bolder	Bb Wee 1
	\\\\@	Sans of	MA	6w 18.

6. 9. gray/white yed

2. gray/white yed

2. black/cod-yed

1. black/white yed

1. black/white yed

2

	7	B?	Wired	X	bbww Genoty	De was	100 % BBW	bolk dun	bek
€.	gave	st cros	s involving gray-bodie ar fly?	one of	the gray-bod eyed flies. V	lied, red-eye	ed F2 flies		
	actor c ccDDI		as perform	ed usir	ng a plant wi	h genotype			
a.	Wha	t shoul La BB	d the geno CCddee	type of	the other pa	rent plant be	97		
		the po		etes fo	med by Par	ents.			
C.	Show	the ge	notype of t Aa Bb (he F1 p	olant. dE=				
d.			different <u>ga</u> 32. <i>9</i>		can be produ <i>∤∂</i> S	iced in the f	F1 plant?	n= 5	
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following genotype? UuWwXxYyZz (n=5 heterozygous gene pairs). For example UwxYZ could be found in one gamete…how
many other unique gametes. $2^5 = 32$ (31 offers his ides $u_{\omega x}$)
b. What is the probability that two mice with the genotype above will produce offspring with the same genotype (UuWwXxYyZz)?
5. When performing a testcross on a maize plant giving all red kernels you note on the offspring plant the development of ears of corn with both red and white kernels (on the same ear). a. What, specifically, might you hypothesize about your original maize plant with red kernels? (show the genotypes in the cross and resultant offspring) The was he have your allows the form of the control of
 b. You count 2 rows of kernels. The results are: 15 red: 35 white. Later you count a whole ear of corn noting 240 red: 260 white.
What is your null hypothesis? Perform the X^2 analysis on both sets of data. What do you note about the value (o-e) in both data sets? (i.e. how far from the expected value were you in each case)? In each case what is your p value? Decide whether you would reject or fail to reject your null hypothesis in each case.
Why might the same ear of corn give you data with different "goodness of fit"
Sample of the law of corn so the newson we reject the probably reflected more variation related to small we reduce sample. When we increased the sample to the reduce shown in dark gray. Predict the mode of inheritance of the phenotype and note the most probable genotypes of each individual. Assume that one gene controls the phenotype (the disease) and alleles are A and a for the 2 alleles for the gene. Would any in generation V (the offspring of

4.

generation IV) be expected to have the disorder? If yes, under what circumstances? aa This is a recessive hait. most lilly individuals have affected officery Since now of the Chiden in IX are objected its likely
their Mom was AA and they are betaggets. The could pass
the tilt to generation V if the maded with another
they have 4 children together. For a and b below, sketch a nediarra for
the family showing the the family showing the parents and children. Show the expected number of affected children in each case. More than one correct answer may be possible. a. The disorder is recessive and is controlled by a gene that is NOT on one of the sex-determining chromosomes (autosomal recessive) 50% expected to b. The disorder is dominant and is controlled by a gene that is NOT be offered on one of the sex-determining chromosomes (autosomal dominant) aa. 50% hd 100% Affected

Test cross gives 2 phenotypes suggesting Plant showing the dom. Phinotype is heteroziggows

> WW X WW -> 1/2 WW red red While 1/2 www while

* null hypoth = Difference in 0- ; e values is due to

1st count

15 + 35 = 50

Expect 1:1

25: 25

 $\chi^{2} = \frac{(15 - 25)^{2}}{25} + \frac{(35 - 25)^{2}}{25}$ $\chi^{2} = 4 + 4$

X2 = 8

df = 1

p7.01 - repect null hypoth

| v-e|=10

2rd Count 240 + 240 = 500 Expect 1:1 250: 250

 $\chi^{2} = (240 - 250)^{2} + (240 - 250)^{2}$

. 4 . 4

x = 0.8

. Z L P L - 5

Fail to rejuct

Absolute devistion 15 SAME - but

Sample 2 15 much Larger (10x)