Series 2: Cross Diagrams – Linkage Analysis

There are two alleles for each trait in a diploid organism

In C. elegans gene symbols are ALWAYS italicized.

To represent two different genes on the same chromosome:

When both genes are wild-type: + is the wild type or non-mutant form of a gene:



The phenotype of this worm is wild type

When both genes are mutant:



The phenotype of this worm is double mutant Dpy and Unc phenotype.

When one gene is wild type and the other mutant:



The phenotype of this worm is Unc



The phenotype of this worm is Dpy





The phenotype of these worms is wild type

To represent two different genes on different chromosomes:

There is noticeable space between the two chromosomes



The phenotype of this worm is wild type

When both genes are mutant:

The phenotype of this worm is double mutant Dpy and Unc.

When one gene is wild type and the other mutant:



The phenotype of this worm is Unc

$$\frac{dpy +}{dpy +}$$

The phenotype of this worm is Dpy





The phenotype of these worms is wild type

Mating symbols:

- X symbolizes mating between two different individuals
- X symbolizes a self cross when the hermaphrodite worms fertilize their own eggs

To discover the chromosome on which our mutated gene of interest is located, we will use a set of reference marker genes to perform linkage testing. By completing the following cross diagrams, you will work out your expectations for ratios of progeny when two genes are on the same chromosome (linked) vs. on different chromosomes (unlinked). Your expectations will be different when:

- 1) The *dpy* gene of interest **IS** on the same chromosome as one of the *unc* markers. We should see linkage in 1 of our 5 strains.
- 2) The *dpy* gene of interest **IS NOT** on the same chromosome as one of the *unc* markers. We should see this result in 4 of our 5 strains.

Work through the expectations for each cross scenario. You should obtain different expected ratios of progeny dependent on the location of the genes. Show all work and answer all questions for full credit.

The *dpy* gene of interest IS on the same chromosome as one of the *unc* markers.

Cross 1: Dumpy hermaphrodites x wild type males to make heterozygous males. This cross will also allow you to determine if your *dpy* gene is sex-linked.

Give the phenotype	of the F1	progeny
- ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '		1 - O - J

Why must we create heterozygous males?

What phenotype would you see in these F1 progeny if *dpy* was sex linked?

Cross 2: Heterozygous males x *unc* marker strain (linked)

We will now cross the F1 progeny to the *unc* marker strains to determine what our expected progeny will look like if our *dpy* gene and *unc* marker gene are on the **SAME** autosome.

We now must self the F1 progeny to determine lin	kage:
Show the self cross for the F1 worms lacking the dicross above:	py gene from the
List the genotype of the carry the dpy gene	e worm that does not
— What gametes can this	worm make?
	ne genotypes of the om this worm?
What are the phenotyp	e(s) of these progeny?
Can you determine the ratio of each phenotype from YES, give the ratio with corresponding phenotype, not.	
This cross does not tell us anything about the linka WHY NOT?	ge of <i>dpy</i> to <i>unc</i> .
If the worms resulting from this cross are uninform have to work through this cross?	native, why did you

Show the self cross for the	ne F1 worms CONTAINING the dpy gene:
	List the genotype of the worm that carries the <i>dpy</i> gene
	— What parental gametes can this worm make?
	What recombinant gametes can this worm make?
• • • •	of the progeny from this worm? Differentiate among I gametes (most) & rarer genotypes that include one or ses.
	s) of these progeny? Differentiate those that come s, 1 recombinant gamete and 2 recombinant gametes
YES, give the ratio with	atio of each phenotype from this self cross? If corresponding phenotype. If NO, explain why w how far apart these two genes are on the

Why are double mutant *dpy unc/dpy unc* progeny highly unlikely to occur from this cross?

Cross 2: Heterozygous males x *unc* marker strain (un-linked)

We will now cross the F1 progeny to the *unc* marker strains to determine ratios in the progeny if our *dpy* gene and *unc* marker gene are on the **DIFFERENT** chromosomes.

We now must self these worms to d	letermine linkage:	
Show the self cross for the worms l on the previous page:	acking the dpy gene from the cross	
	e genotype of the worm that does not ne dpy gene	
——— What g	gametes can this worm make?	
	What are the genotypes of the progeny from this worm?	
What a	re the phenotype(s) of these progeny?	
Can you determine the ratio of each phenotype from this self cross? If YES, give the ratio with corresponding phenotype. If NO, explain why not.		
This cross does not tell us anything WHY NOT?	about the linkage of dpy to unc.	

If the worms resulting from this cross are uninformative, why did you have to work through this cross?			
Show the self cross for the	worms CONTAINING the dpy gene:		
	List the genotype of the worm that carries the <i>dpy</i> gene		
	What gametes can this worm make?		
What are the genotypes of	the progeny from this worm?		
What are the phenotype(s)	of these progeny?		

Can you determine the ratio of each phenotype from this self cross? If YES, give the ratio with corresponding phenotypes. If NO, explain why not.
What kind of assortment does this represent?