## 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

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

The phenotype of this worm is Dpy



The phenotype of these worms is wild type

Mating symbols:

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

NAME:	

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.

dpy X	+	Parental Generation Genotype
$\overline{dpy}$	+	
		List the gametes each parent can produce
		Give the genotype of the F1 progeny

Give the phenotype of the F1 progeny

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 det	ermine linkage:
Show the self cross for the F1 worms lactors above:	cking the dpy gene from the
List the gen carry the dp	otype of the worm that does not by gene
——— What gamet	tes can this worm make?
	What are the genotypes of the progeny from this worm?
What are the	e phenotype(s) of these progeny?
Can you determine the ratio of each phe YES, give the ratio with corresponding not.	~ ~
This cross does not tell us anything about WHY NOT?	at the linkage of <i>dpy</i> to <i>unc</i> .
If the worms resulting from this cross are have to work through this cross?	re uninformative, why did you

Show the self cross for the F1 worms CONTAINING the <i>dpy</i> gene:	
List the go	enotype of the worm that carries ene
— What pare	ental gametes can this worm make?
What reco	ombinant gametes can this worm
What are the genotypes of the progeny	y from this worm?
What are the phenotype(s) of these pro	ogeny?
Can you determine the ratio of each particle. YES, give the ratio with corresponding not. HINT: Do you know how far appearance chromosome?	g phenotype. If NO, explain why

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 worm	s to determine linkage:
Show the self cross for the wor	rms lacking the dpy gene from the cross
	st the genotype of the worm that does not rry the $dpy$ gene
W	hat gametes can this worm make?
	What are the genotypes of the progeny from this worm?
W	That are the phenotype(s) of these progeny?
•	each phenotype from this self cross? If sponding phenotype. If NO, explain why
This cross does not tell us anyt WHY NOT?	thing about the linkage of <i>dpy</i> to <i>unc</i> .

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?