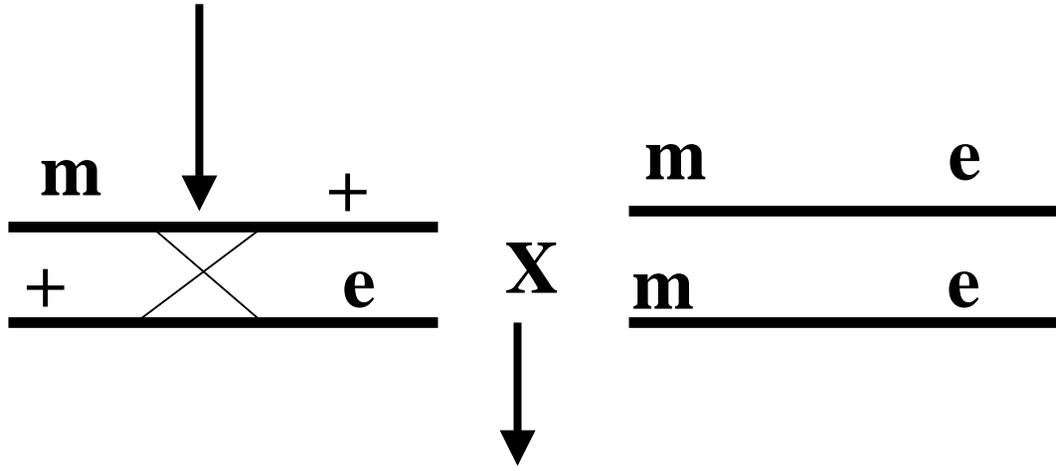
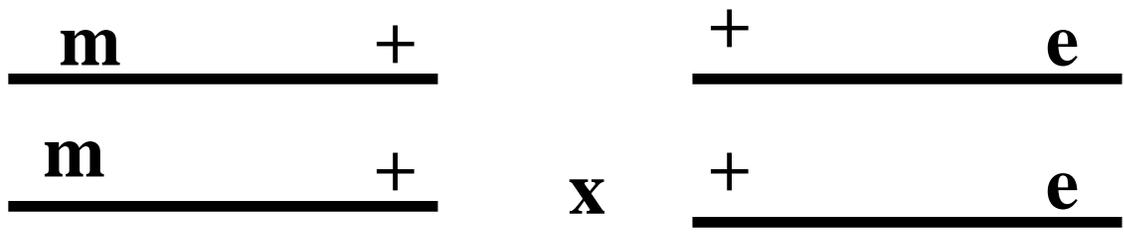
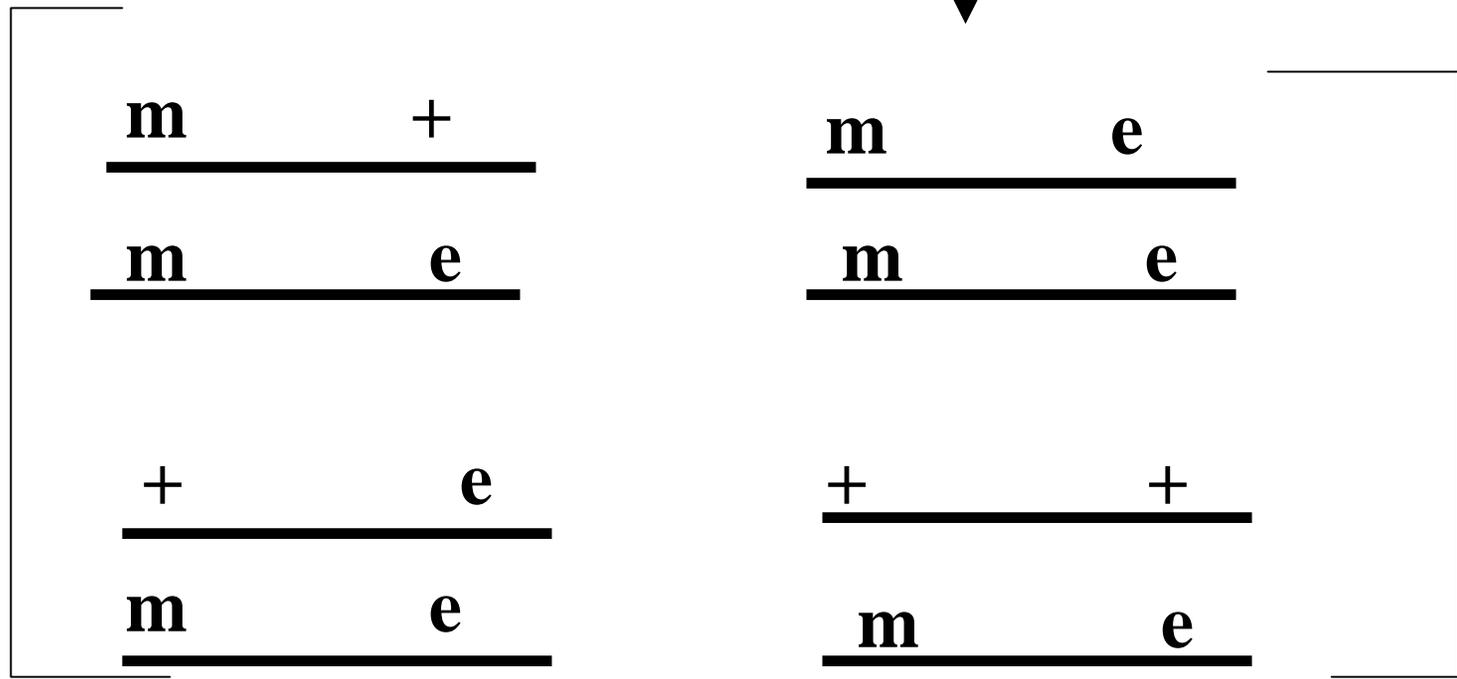


- 1. The genes for mahogany eyes and ebony body are recessive and 25 map units apart on an autosome in *Drosophila*. A mahogany-eyed female (wild type body, homozygous) was mated to an ebony-bodied (wild type eyes, homozygous) male and the resulting F1 phenotypically wild type females were mated to mahogany, ebony males.**
 - a. What percentage of the offspring would have the phenotype mahogany eyes and wild type body?**
 - b. What percentage of the offspring would have the phenotype mahogany eyes and ebony body?**
 - c. What percentage of the offspring would be phenotypically wild type for both genes?**



75%



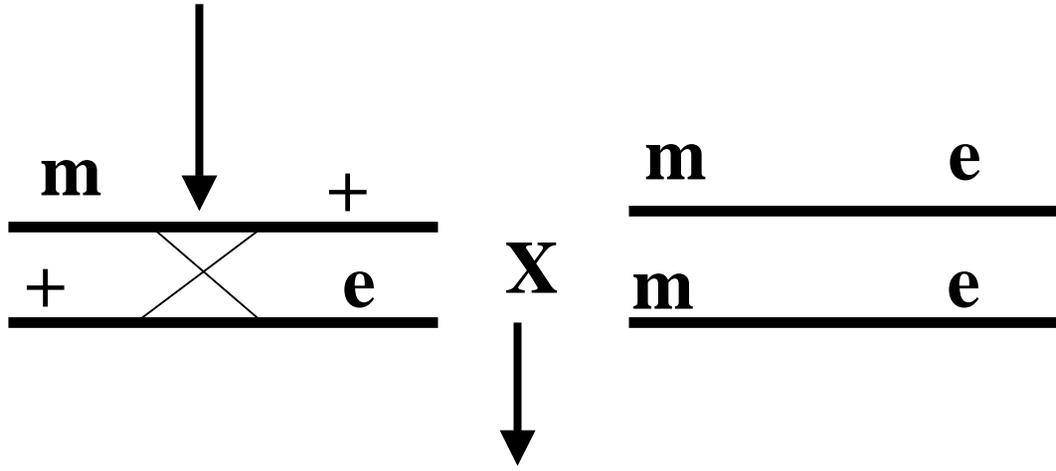
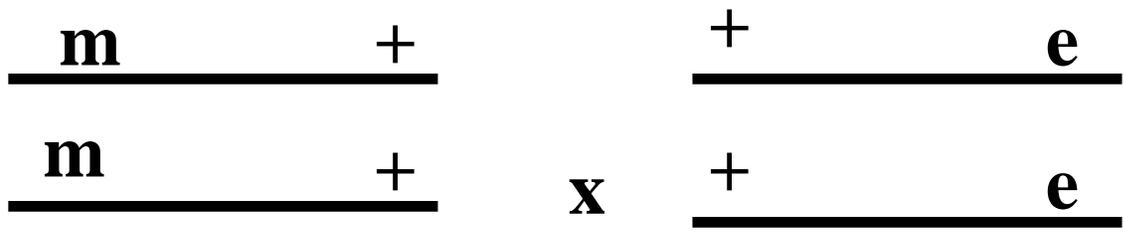
25%

a. What percentage of the offspring would have the phenotype mahogany eyes and wild type body?

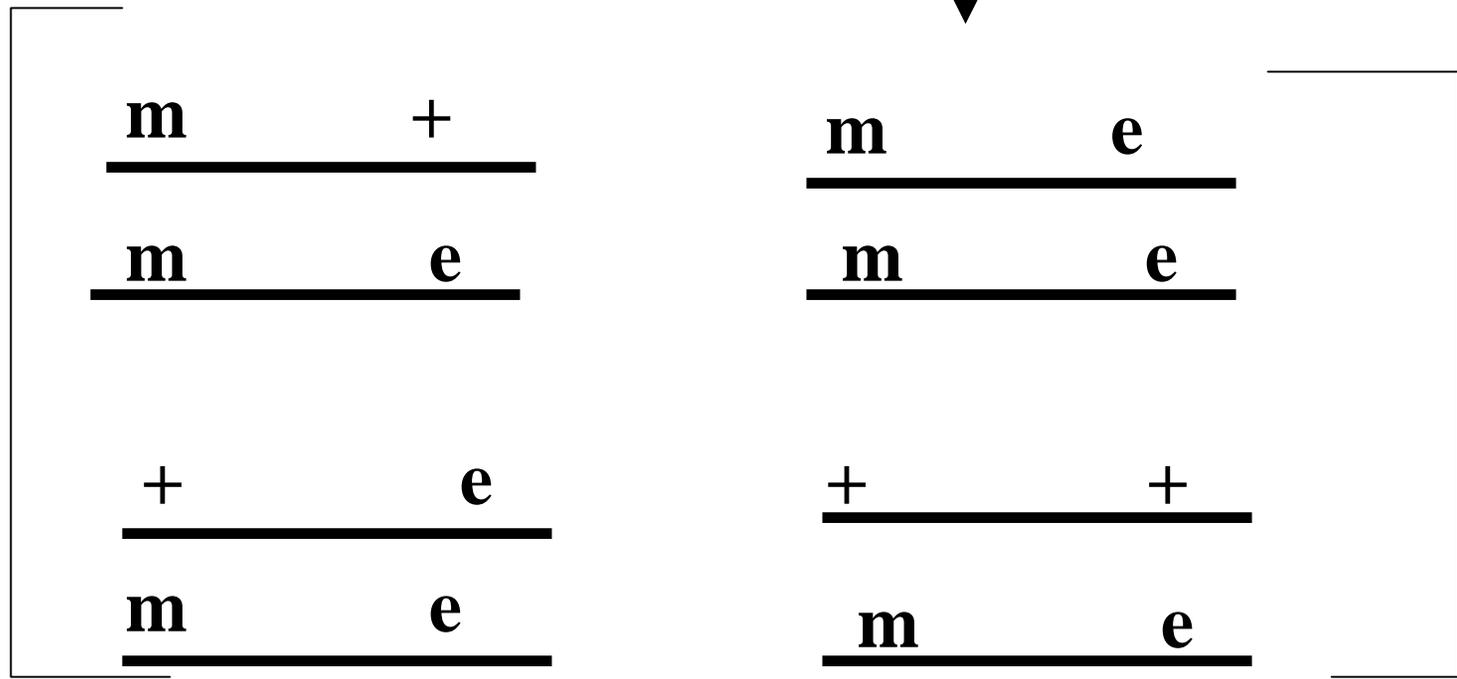
25 units means 25% recombinants formed so 75% are nonrecombinant, one-half of these or 37.5% will be mahogany and wild for body color.

b. What percentage of the offspring would have the phenotype mahogany eyes and ebony body?

Those would be the recombinants, $1/2$ of 25% = 12.5%



75%



25%

c. What percentage of the offspring would be phenotypically wild type for both genes?

These would be the recombinants, $1/2$ of $25\% = 12.5\%$

2. In corn the gene for colored (C) seeds is completely dominant to the gene for colorless (c) seeds. Similarly, for the character of the endosperm (the part of the seed that contains the food stored for the embryo), a single gene pair controls whether the endosperm is full or shrunken. Full (S) is dominant to shrunken (s). A true-breeding colored, full seeded plant was crossed with a colorless, shrunken-seeded one. The F1 colored, full plants were test-crossed to the doubly recessive type, that is, colorless and shrunken. The result was as follows:

| | |
|----------------------------|-------------|
| colored, full | 4032 |
| colored, shrunken | 149 |
| colorless, full | 152 |
| colorless, shrunken | 4035 |

- a. From these results, are the gene for color and the gene for endosperm shape linked?**
- b. What is the genotype of the F1 plants?**
- c. How many map units separate the two loci for these genes?**

a. From these results, are the gene for color and the gene for endosperm shape linked?

P₁ CCSS x ccss
↓
F₁ CcSs x ccss

If not linked would expect:

1/4 CcSs

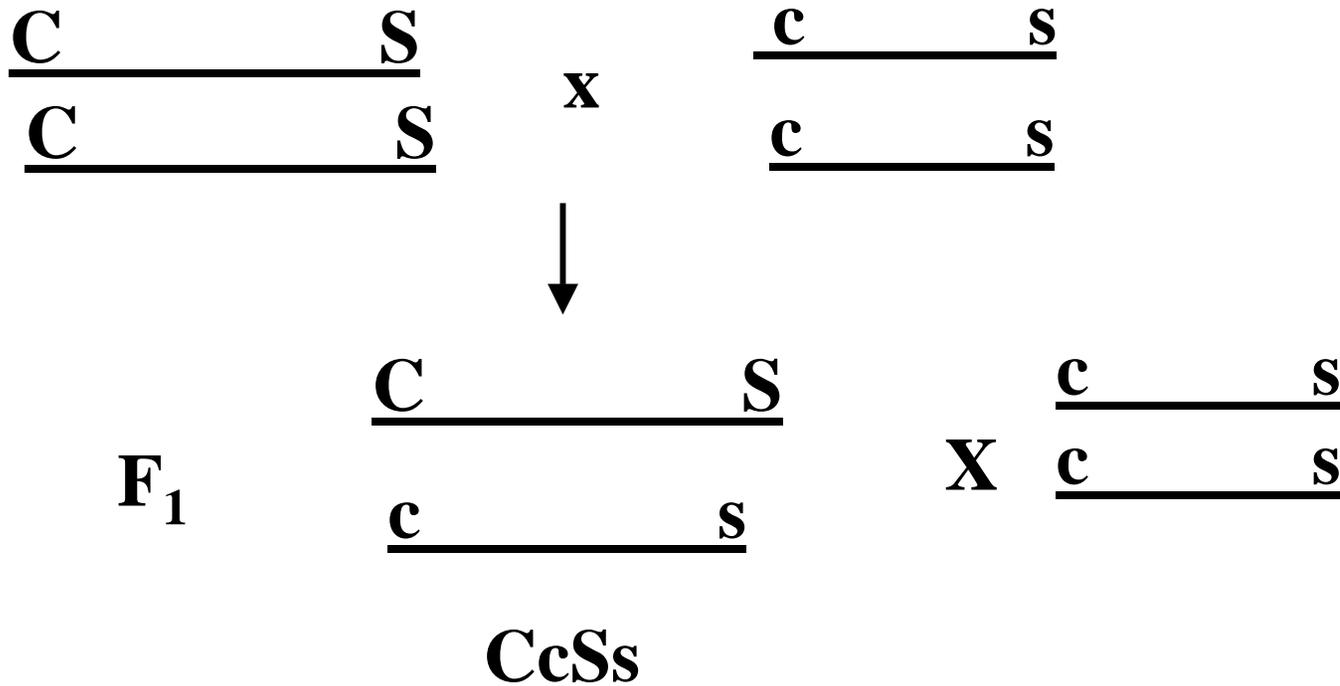
1/4 ccSs

1/4 Ccss

1/4 ccss

Must be linked

b. What is the genotype of the F1 plants?



c. How many map units separate the two loci for these genes?

| | |
|----------------------------|--------------|
| colored, full | 4032 |
| colored, shrunken | 149 |
| colorless, full | 152 |
| colorless, shrunken | 4035 |
| | ----- |
| | 8,368 |

$$\begin{aligned} \% \text{ Recombinants} &= \frac{301}{8,368} = 3.6\% = 3.6 \text{ map units} \end{aligned}$$

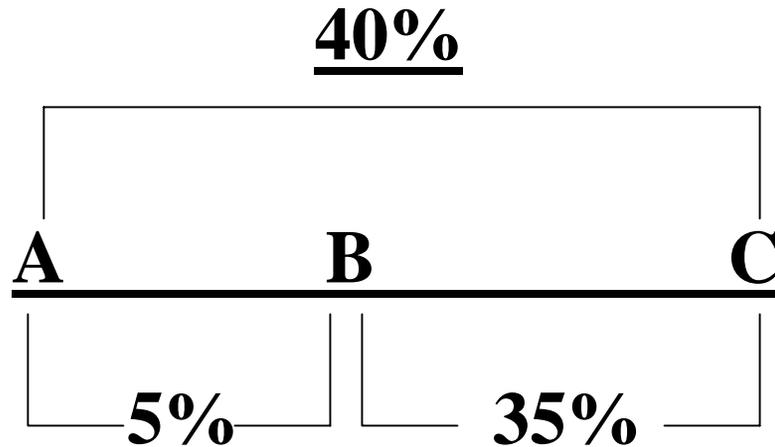
3. Below are listed the frequencies of crossing over between 3 different genes (A, B, C).

A and C = 40%

A and B = 5%

C and B = 35%

- a. Which two genes have the greatest distance between them?**
- b. Which two genes are the closest together?**
- c. How many map units is A from B?**



- Which two genes have the greatest distance between them? A and C**
- Which two genes are the closest together? A & B**
- How many map units is A from B? 5**

4. Genes A and B are linked and are 20 map units apart. An AABB individual was mated with an aabb individual.

a. List the genotypes of the gametes produced by AABB. AB

b. List the genotypes of the gametes produced by aabb. ab

c. With respect to these two genes, how many types of recombinant gametes will both of these individuals produce? 0

C.

$$\frac{\mathbf{A} \quad \mathbf{B}}{\mathbf{A} \quad \mathbf{B}} \quad \mathbf{X} \quad \frac{\mathbf{a} \quad \mathbf{b}}{\mathbf{a} \quad \mathbf{b}}$$

5. Three of the many recessive mutations in *Drosophila* that affect body color, wing shape or bristle morphology are black (b) body versus grey in the wild type, dumpy (dp) versus long wing in the wild type, and hooked (hk) bristles at the tip versus not hooked in the wild type. These genes are linked. From a cross of a dumpy female (wild type for other two traits) with a black and hooked male (wild for other trait), all the F1 were wild type for all three traits. The testcross of an F1 female with a dumpy, black, hooked male gave the following results (if a trait is not listed it is considered to be wild type in that group):

| | |
|---------------------------|------------|
| wild type | 168 |
| black | 19 |
| black, hooked | 301 |
| dumpy, hooked | 21 |
| hooked | 8 |
| hooked,dumpy,black | 172 |
| dumpy, black | 6 |
| dumpy | 305 |

Total = 1000

- a. Using genetic symbols above and below a line, what is the genotype of the F1 female?**
- b. Which groups are due to double crossing over?**
- c. What is the correct sequence of these genes?**

dp + +

dp + +

X

+ b hk

+ b hk



F₁

dp + +

+ b hk

X

dp b hk

dp b hk

b. Which groups are due to double crossing over?

| | |
|---------------------------|----------------------------|
| wild type | 168 (single cross.) |
| black | 19 (single cross) |
| black, hooked | 301 (nonrecomb.) |
| dumpy, hooked | 21 (single cross) |
| hooked | 8 (double cross) |
| hooked,dumpy,black | 172(single cross) |
| dumpy, black | 6 (double cross) |
| dumpy | 305 (nonrecomb.) |

Total = 1000

c. What is the correct sequence of these genes?

Three possibilities :

Double cross-overs

| | | |
|-----------|----------|-----------|
| dp | + | + |
| + | b | hk |

| | | | |
|-----------|----------|-----------|---|
| dp | b | + | * |
| + | + | hk | * |

| | | |
|-----------|-----------|----------|
| dp | + | + |
| + | hk | b |

| | | |
|-----------|-----------|----------|
| dp | hk | + |
| + | + | b |

| | | |
|----------|-----------|-----------|
| + | dp | + |
| b | + | hk |

| | | |
|----------|-----------|-----------|
| + | + | + |
| b | dp | hk |

6. In *Drosophila*, the three gene pairs for red eyes (cn^+) vs. cinnabar (cn), normal bristle number (rd^+) vs. reduced (rd), and long wings (vg^+) vs. vestigial (vg) are known to have their loci on chromosome 2. Suppose that a female fly that is heterozygous for all three genes with all wild type alleles on one homologue and all mutant alleles on the other is crossed with a male that is recessive for all three genes (three-point test-cross) and yields the following offspring:

| | |
|-------------------------------------|------------|
| cinnabar, reduced, vestigial | 406 |
| cinnabar, reduced, long | 46 |
| cinnabar, normal, vestigial | 28 |
| cinnabar, normal, long | 3 |
| red, normal, long | 438 |
| red, normal, vestigial | 45 |
| red, reduced, long | 33 |
| red, reduced, vestigial | 1 |

- a. Which progeny classes are the double recombinant types?**

- b. What is the correct sequence of these genes?**

- c. What is the distance between eye color (cn^+ , cn) and wing type (vg^+ , vg)?**

| | | |
|-----------|-----------|-----------|
| cn | rd | vg |
| <hr/> | | |
| + | + | + |
| <hr/> | | |

x

| | | |
|-----------|-----------|-----------|
| cn | rd | vg |
| <hr/> | | |
| cn | rd | vg |
| <hr/> | | |

a. Which progeny classes are the double recombinant types?

| | |
|-------------------------------------|------------|
| cinnabar, reduced, vestigial | 406 |
| cinnabar, reduced, long | 46 |
| cinnabar, normal, vestigial | 28 |
| cinnabar, normal, long | 3* |
| red, normal, long | 438 |
| red, normal, vestigial | 45 |
| red, reduced, long | 33 |
| red, reduced, vestigial | 1* |

b. What is the correct sequence of these genes?

cn rd vg
+ + +

cn + vg
+ rd +

rd cn vg
+ + +

rd + vg * bristle number
+ cn + * eye color
wing type

cn vg rd
+ + +

cn + rd
+ vg +

c. What is the distance between eye color (cn+, cn) and wing type (vg+,vg)?

| | | % |
|-------------------------------------|-----------------|-------------|
| cinnabar, reduced, vestigial | 406 | 40.6 |
| cinnabar, reduced, long | 46 | 4.6 |
| cinnabar, normal, vestigial | 28 | 2.8 |
| cinnabar, normal, long | 3 | 0.3 |
| red, normal, long | 438 | 43.8 |
| red, normal, vestigial | 45 | 4.5 |
| red, reduced, long | 33 | 3.3 |
| red, reduced, vestigial | <u>1</u> | 0.1 |
| | 1000 | |

| | | | |
|-----------|-----------|----------|-----------|
| rd | cn | ? | vg |
| | | | |
| + | + | | + |
| | | | |

Single cross-overs - rd cn + = 4.6%

+ + vg = +4.5%

double recomb. + .4%

9.5 units

7. In chickens the dominant allele O causes egg shell color to be bluish and the recessive genotype oo produces white eggs.

Also the allele P produces a variant in comb type referred to as pea comb. P is incompletely dominant to the allele for single comb, p, although in small wattled breeds PP and Pp are practically indistinguishable.

Both loci are located on the second chromosome.

Single



Walnut



Rose



Pea



**Assume the mating : PPoo (pea comb, white eggs)
x (single comb, blue eggs) ppOO producing the
dihybrid PpOo**

**The dihybrid is then mated to chickens with
the ppoo genotype.**

**Which two of the four possible dihybrid gametes
represent recombinant gametes?**

PO, Po , pO, po

$$\begin{array}{c} \underline{\underline{P}} \quad \underline{\underline{o}} \\ \underline{\underline{P}} \quad \underline{\underline{o}} \end{array} \quad \times \quad \begin{array}{c} \underline{\underline{p}} \quad \underline{\underline{O}} \\ \underline{\underline{p}} \quad \underline{\underline{O}} \end{array}$$

↓

$$\begin{array}{c} \underline{\underline{P}} \quad \underline{\underline{o}} \\ \underline{\underline{p}} \quad \underline{\underline{O}} \end{array} \quad \times \quad \begin{array}{c} \underline{\underline{p}} \quad \underline{\underline{o}} \\ \underline{\underline{p}} \quad \underline{\underline{o}} \end{array}$$

PO, po

Assume the following phenotypic distribution is observed from mating the dihybrids (produced from crossing the ppOO by PPoo lines) and the completely recessive ppoo:

Phenotype

| <u>Comb</u> | <u>Egg color</u> | <u># Observed</u> |
|--------------------|-------------------------|--------------------------|
| Pea | Blue | 4 |
| Pea | White | 96 |
| Single | Blue | 96 |
| Single | White | 4 |

$$\frac{p}{p} = \frac{O}{O}$$

X
↓

$$\frac{P}{P} = \frac{o}{o}$$

$$\frac{p}{P} = \frac{O}{o}$$

X
↓

$$\frac{p}{p} = \frac{o}{o}$$

NR

$$\frac{p}{p} = \frac{O}{O}$$

$$\frac{P}{P} = \frac{o}{o}$$

R

$$\frac{p}{P} = \frac{o}{O}$$

$$\frac{P}{P} = \frac{O}{O}$$

What is the recombination frequency (hence, map distance) between these two loci?

$$\frac{8}{200} = 4 \%$$

8. In rabbits the dominant allele B produces a black coat color and the recessive genotype bb produces a brown coat. Body fat color in rabbits is white if a dominant allele Y is present and yellow if the genotype is yy.

Assume the mating between rabbits with the following genotypes : Black with white fat(BBYY) and Brown with yellow fat (bbyy) which produces the dihybrid BbYy

Calculate the recombination frequency if the dihybrids are mated back to brown rabbits having yellow fat, bbyy, and the following ratio of phenotypes is observed:

| Phenotype | | # Observed |
|-------------------|------------------|-------------------|
| Coat color | Fat color | |
| Black | White | 125 |
| Black | Yellow | 75 |
| Brown | White | 85 |
| Brown | Yellow | 115 |

Phenotype

Coat color

Black

Black

Brown

Brown

Fat color

White

Yellow

White

Yellow

Observed

125

75

85

115

400

$$\frac{\mathbf{160}}{\mathbf{400}} = \mathbf{40\%}$$