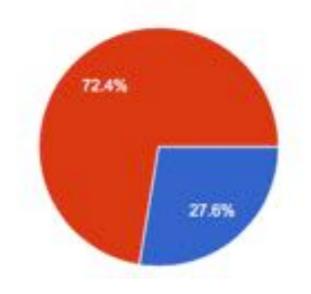
# Course website: bio393.andersenlab.org

Problem set #1 is out.

Genetics terms are online.

### Final will be on Friday March 16 1-3 PM



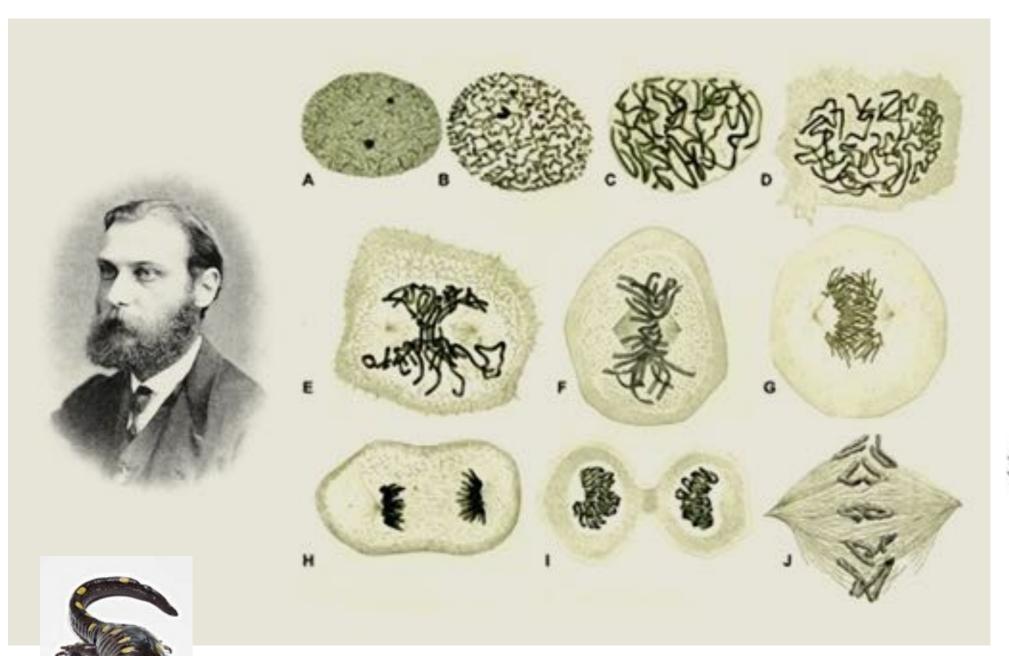
- Normal date and time: Friday March 23rd 12-2 PM
- During reading week: Friday March 16th 1-3 PM

# **Bio393: Genetic Analysis**

Chromosome theory, recombination, and mapping



# Walther Flemming stained cells





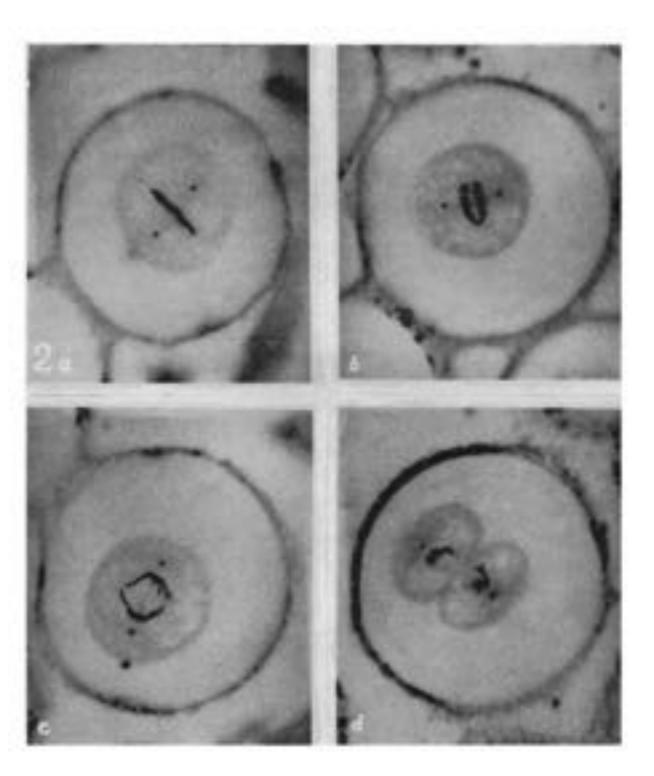
Walther Flemming, 1882

# Cells divide their chromosomes with high fidelity



Theodor Boveri

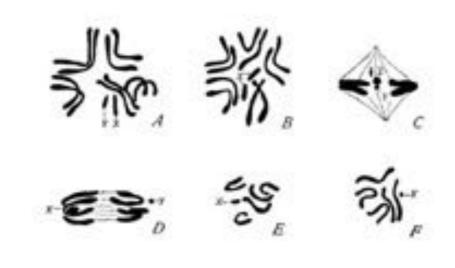




# Discovery of sex chromosomes



Nettie Stevens





Tenebrio melitor

### Gametes have half the chromosomes of the soma



Theodor Boveri



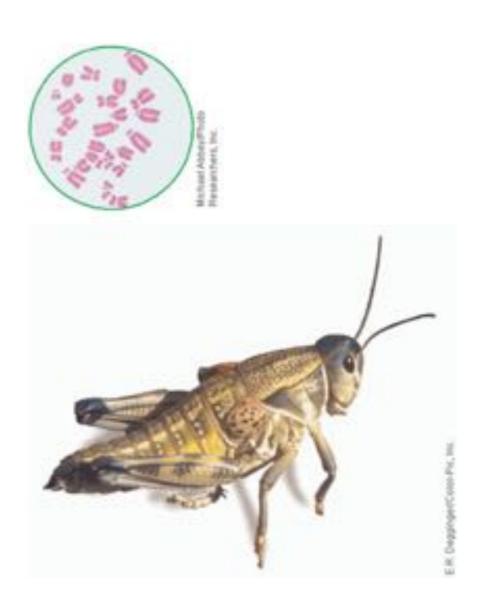
Parascaris equorum



### Discovery of a connection to Mendel's principles

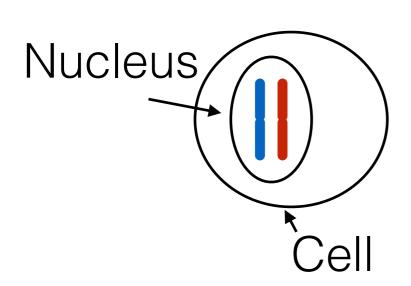


Walter Sutton



- Gametes have half chromosome complement of somatic cells
- Homolog separation to gamete was random

#### Terms for mitosis and meiosis



Ploidy (N)
Diploid (2N)
Haploid (1N)
Polyploid (>2N)
Gamete

Chromosome

Pair of homologs (2N)

Sister chromatids

#### Meiosis: A reductional division in two acts

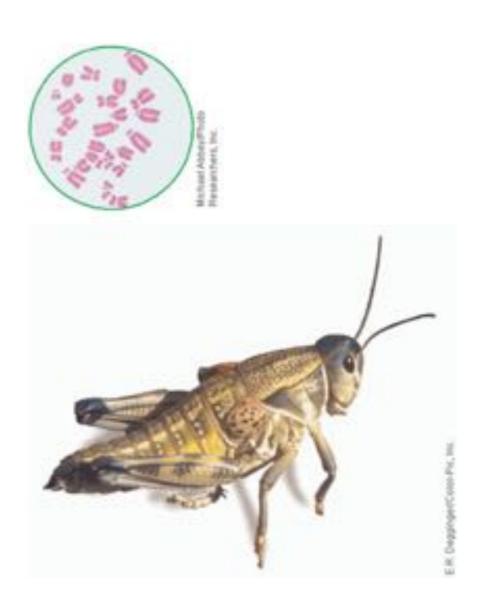
Homologs separate first Cytokinesis Telophase I Interphase Metaphase I Prophase I Metaphase II Anaphase II Prophese II Interkinesis Cytokinesis Sisters MEIOSIS II separate last

Keep track of centromere

### Discovery of a connection to Mendel's principles

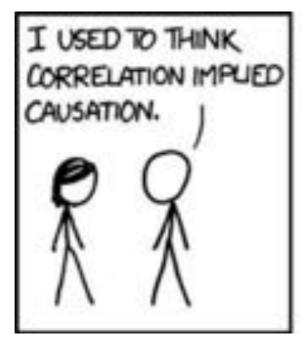


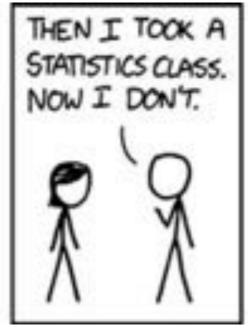
Walter Sutton

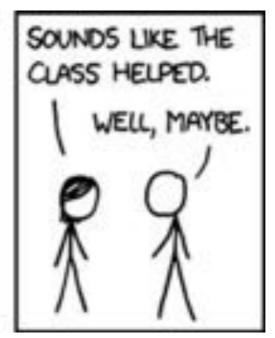


- Gametes have half chromosome complement of somatic cells
- Homolog separation to gamete was random

#### Correlation does not mean causation







xkcd.com



**Thomas Hunt Morgan** 

### Drosophila melanogaster: genetics superstar



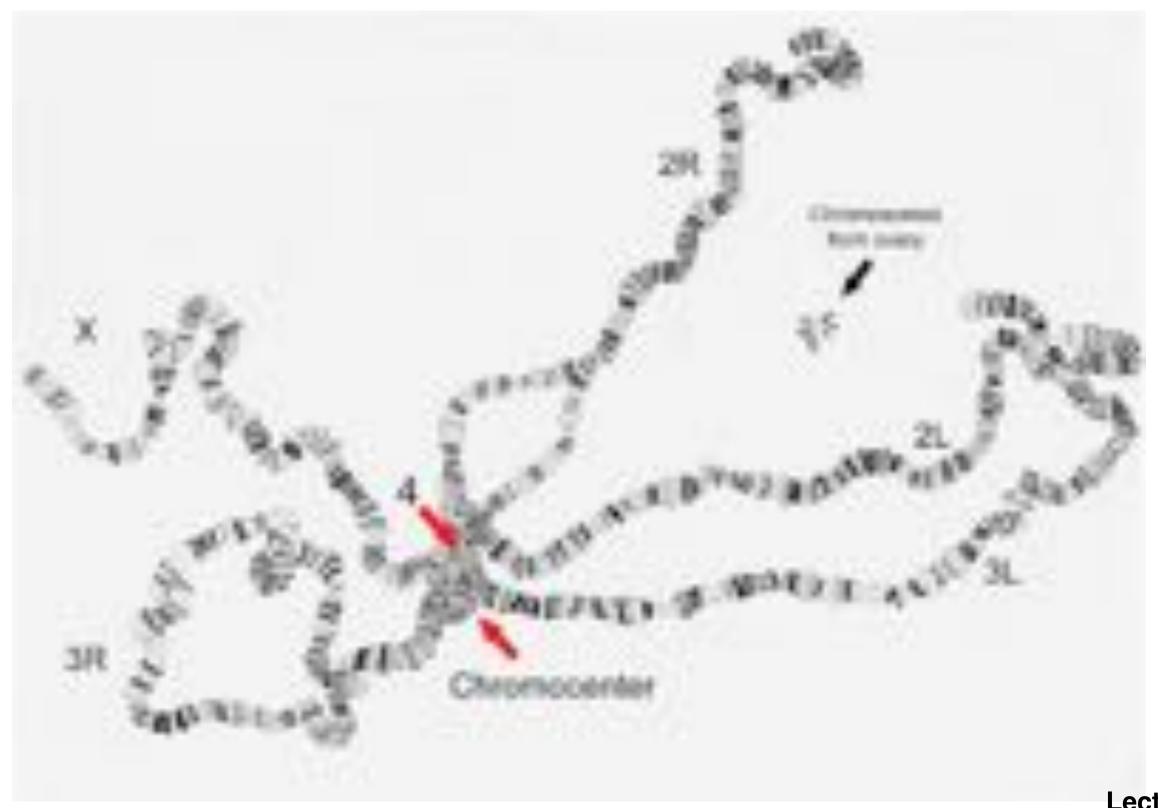


**Nettie Stevens** 

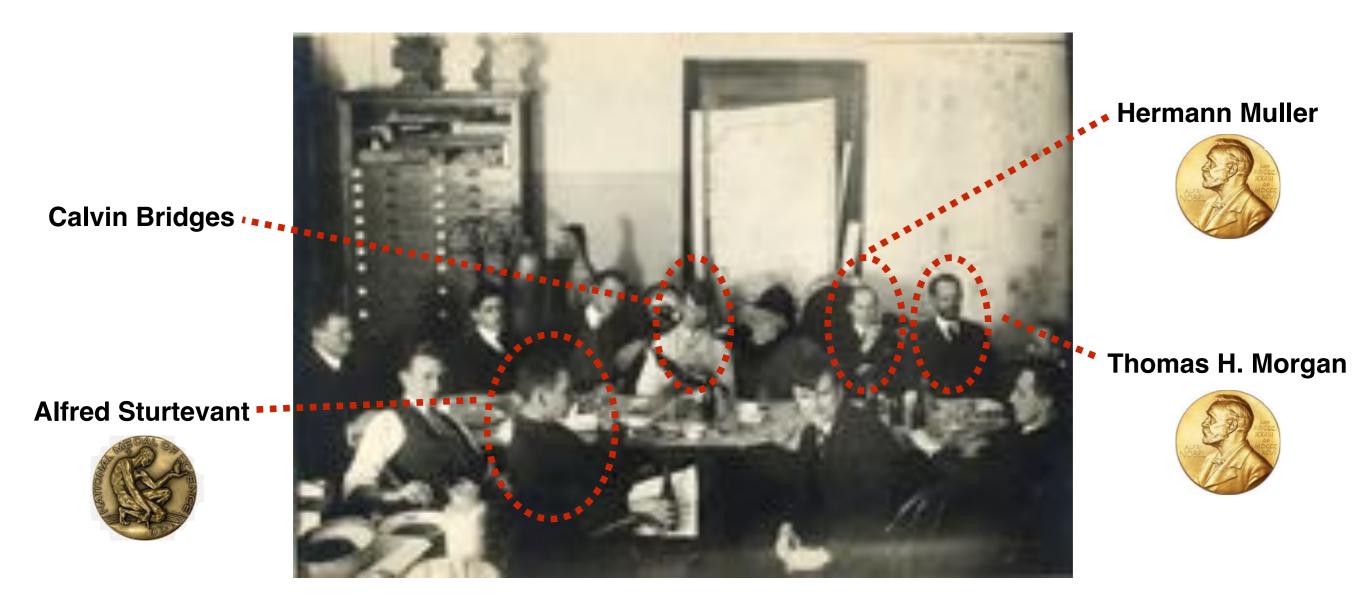


**Thomas Hunt Morgan** 

# Drosophila polytene chromosomes allow us to directly visualize genetic principles

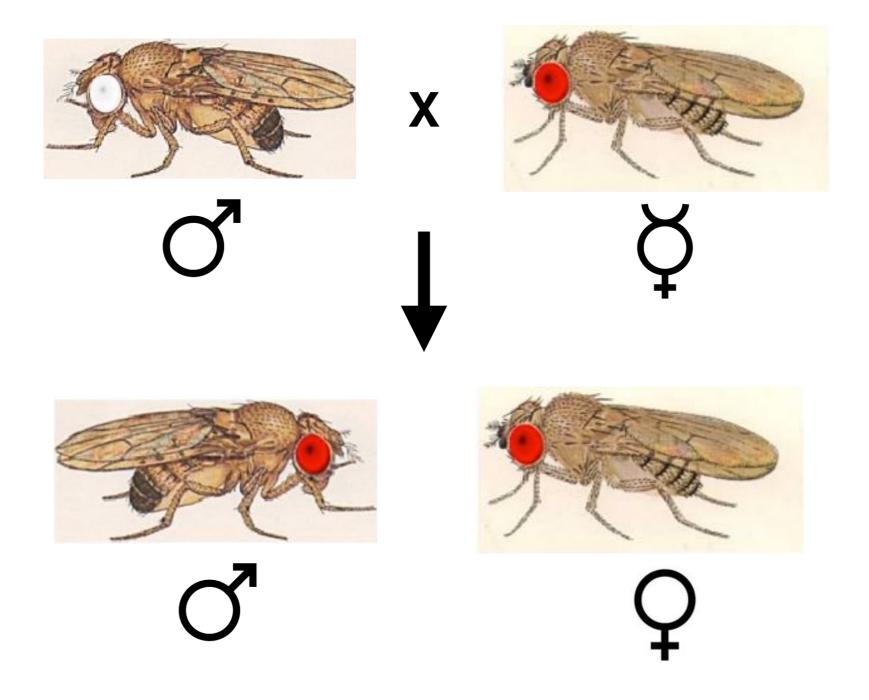


# The fly room at Columbia

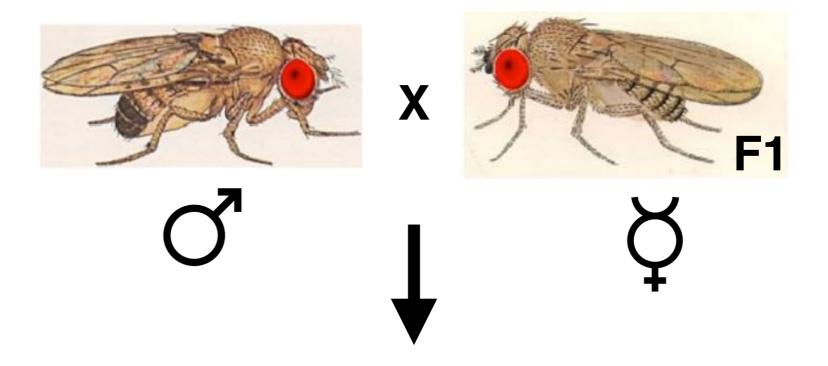


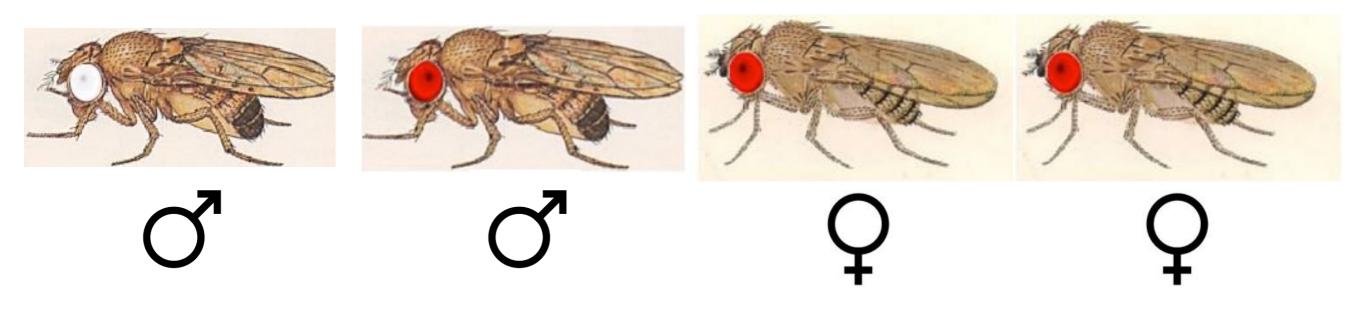
*W*<sup>+</sup>



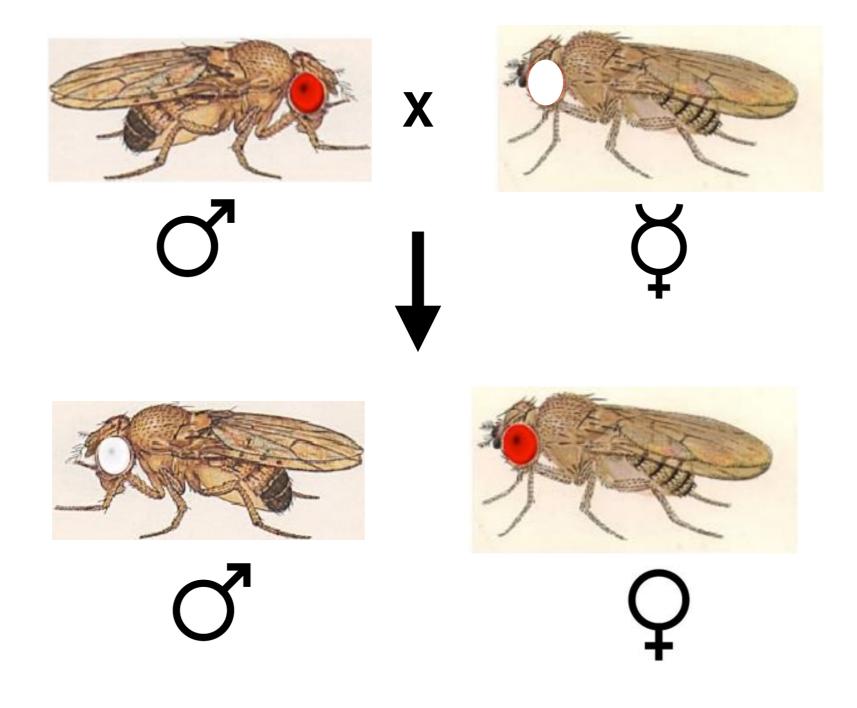


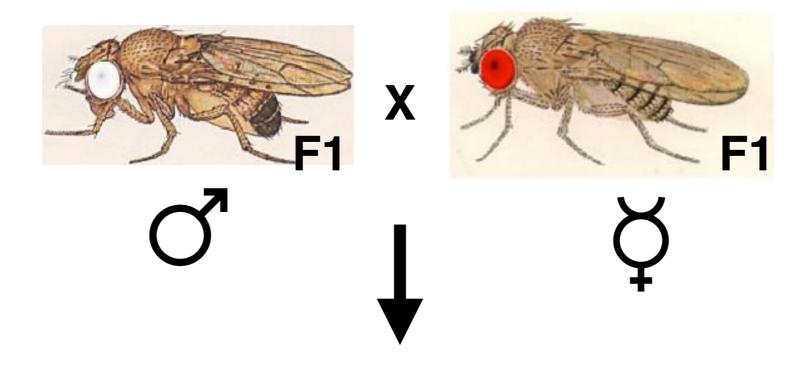
What is dominance relationship of white mutant allele?

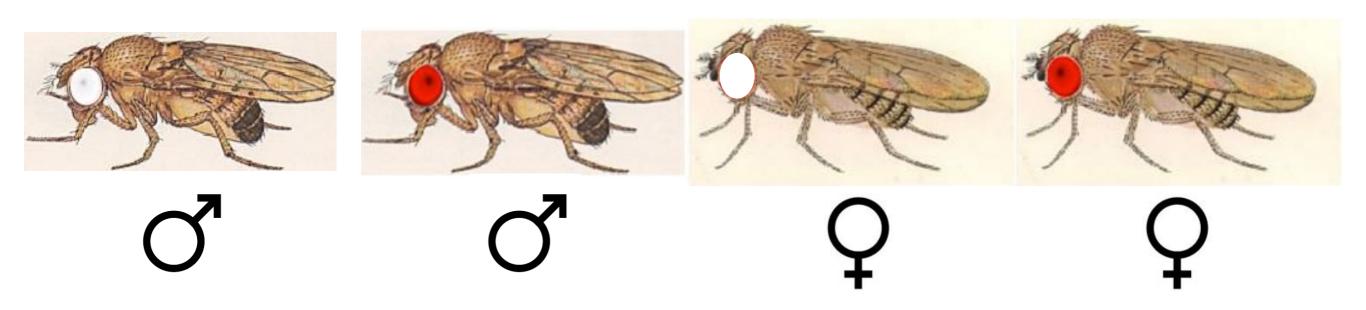




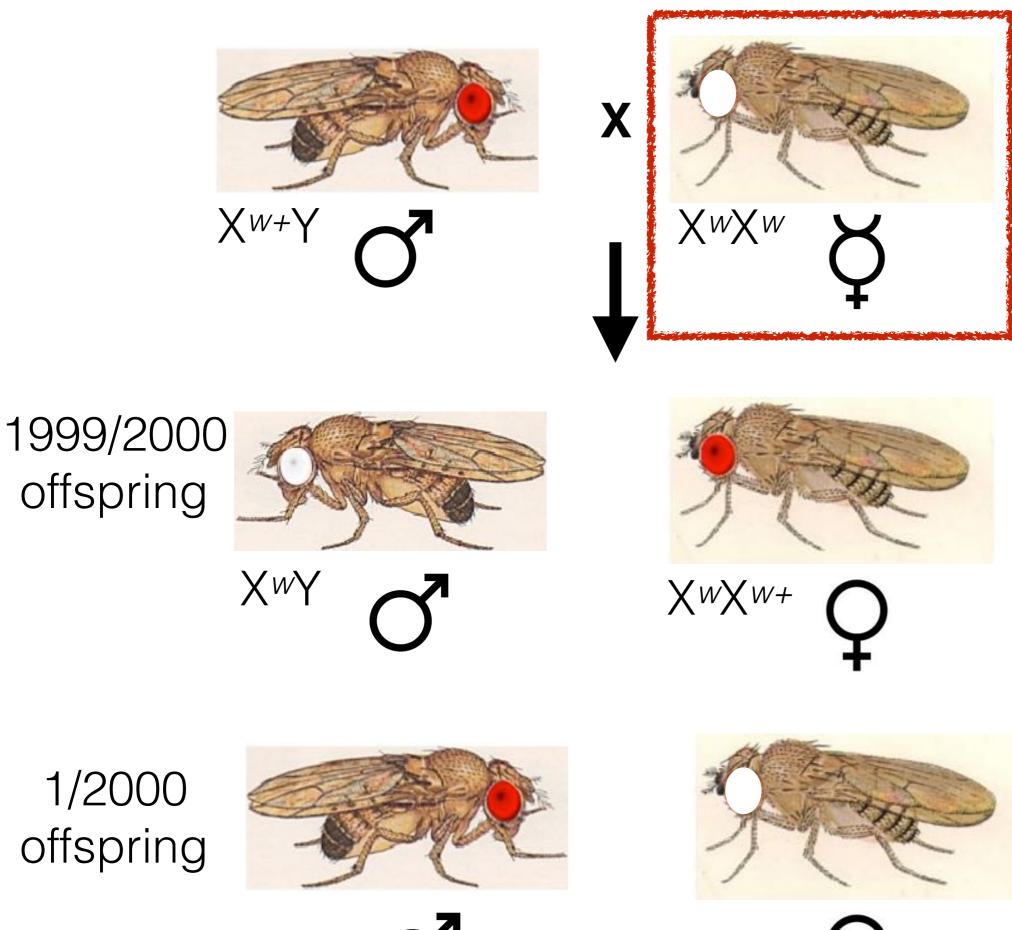
# The reciprocal cross





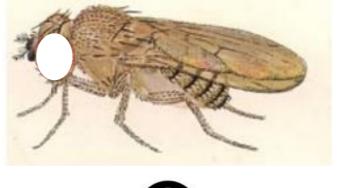


Equal ratios of each sex and eye color



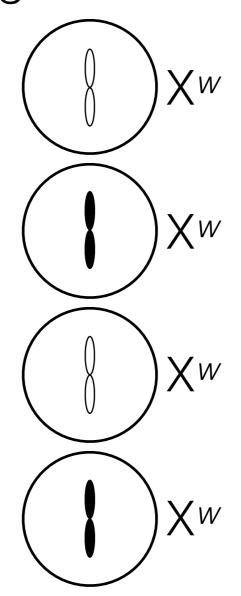
1/2000 offspring



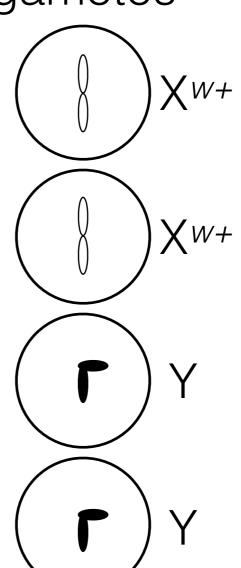




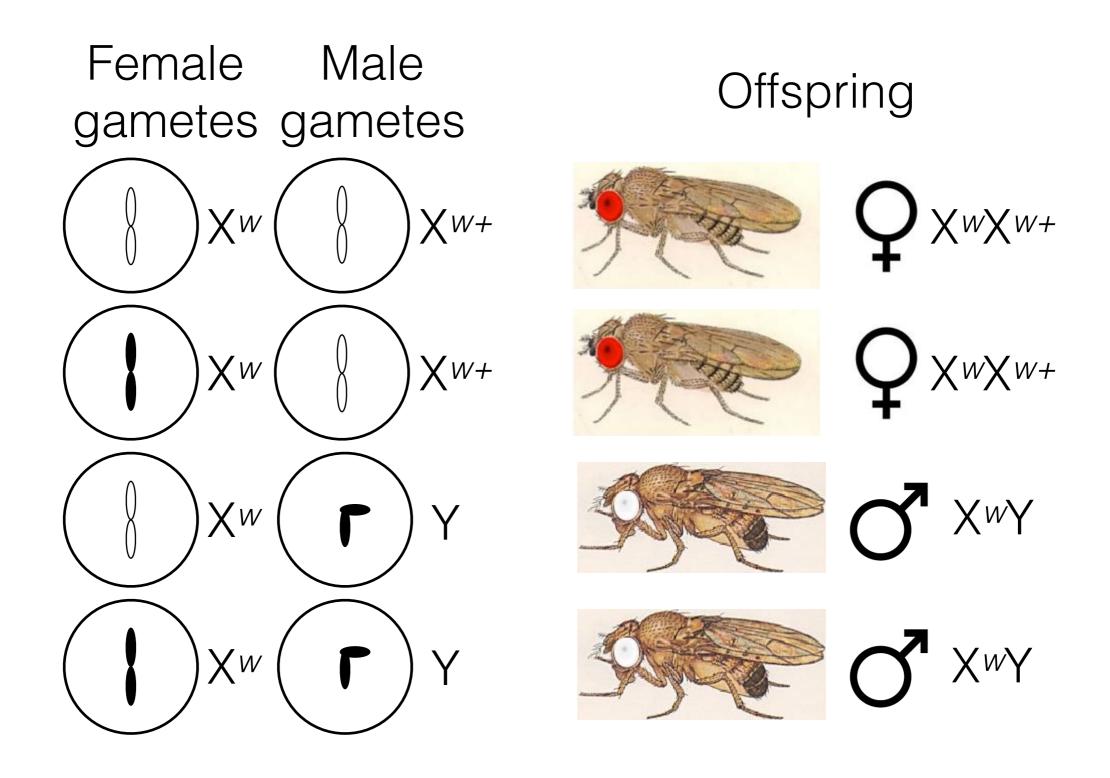
Female gametes



Male gametes

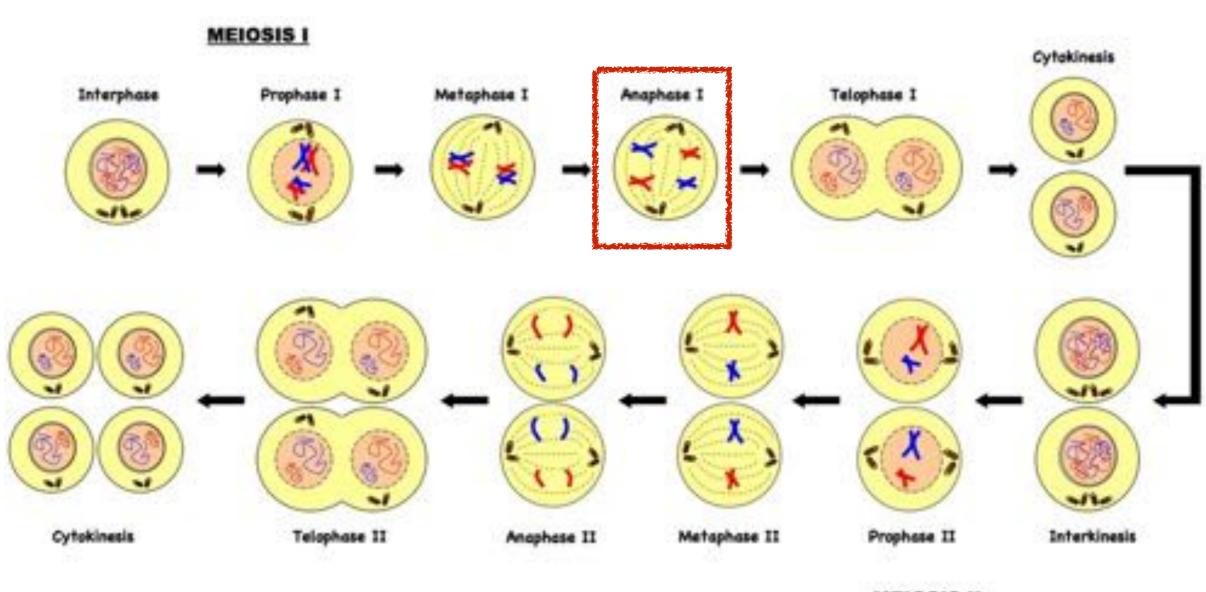


1999/2000 offspring



What is going on with the rare (1/2000) class?

# Meiotic non-disjunction I



MEIOSIS II

Meiosis I NDJ Female Male Female gametes gametes gametes XwXwXW XwXwX-Null X-Null

1999/2000 offspring

 $X^{W+}X^{W}$  $X^{W}Y$  1/2000 offspring

 $X^{w+0}$ 

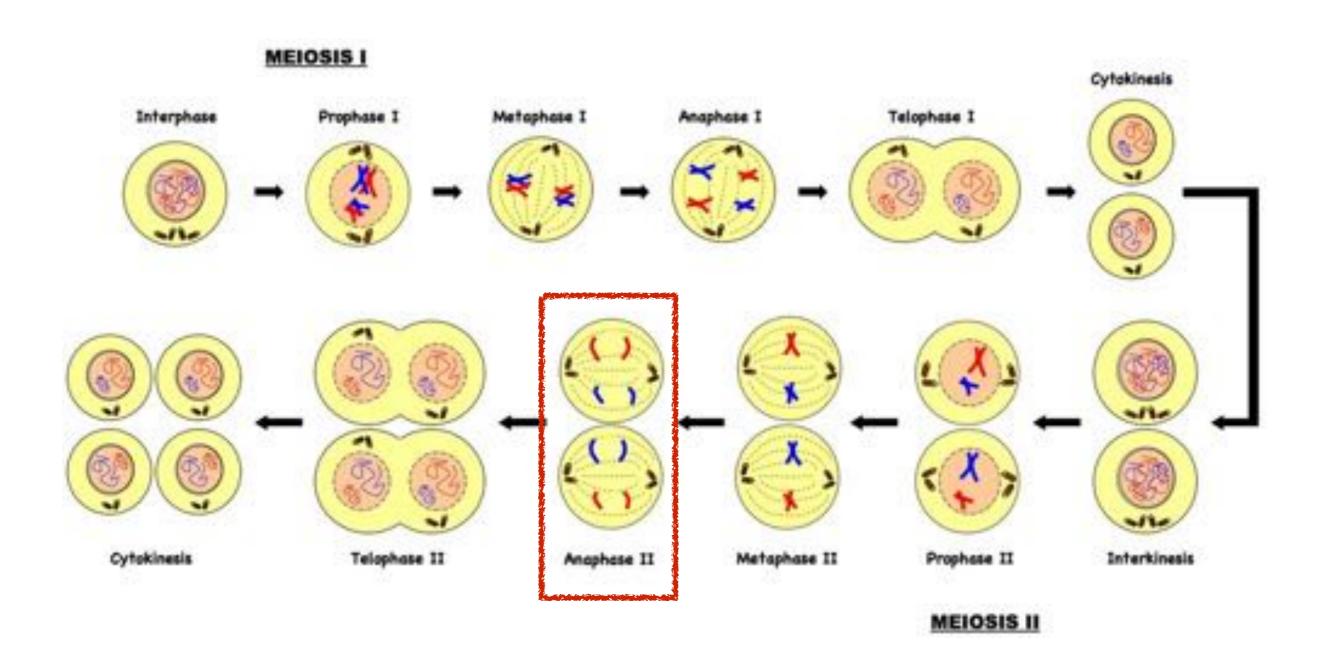
red male white female

XW+

XW+

Meiosis I NDJ Male Offspring Female gametes gametes XwXwXW+ Dead XwXwXw+  $\chi_w\chi_w$ XwXwYX-Null XW+ X-Null Dead

# Meiotic non-disjunction II



Meiosis II NDJ Female Male Female gametes gametes gametes XwXwXW+ XW XW+X-Null XW XW

1999/2000 offspring

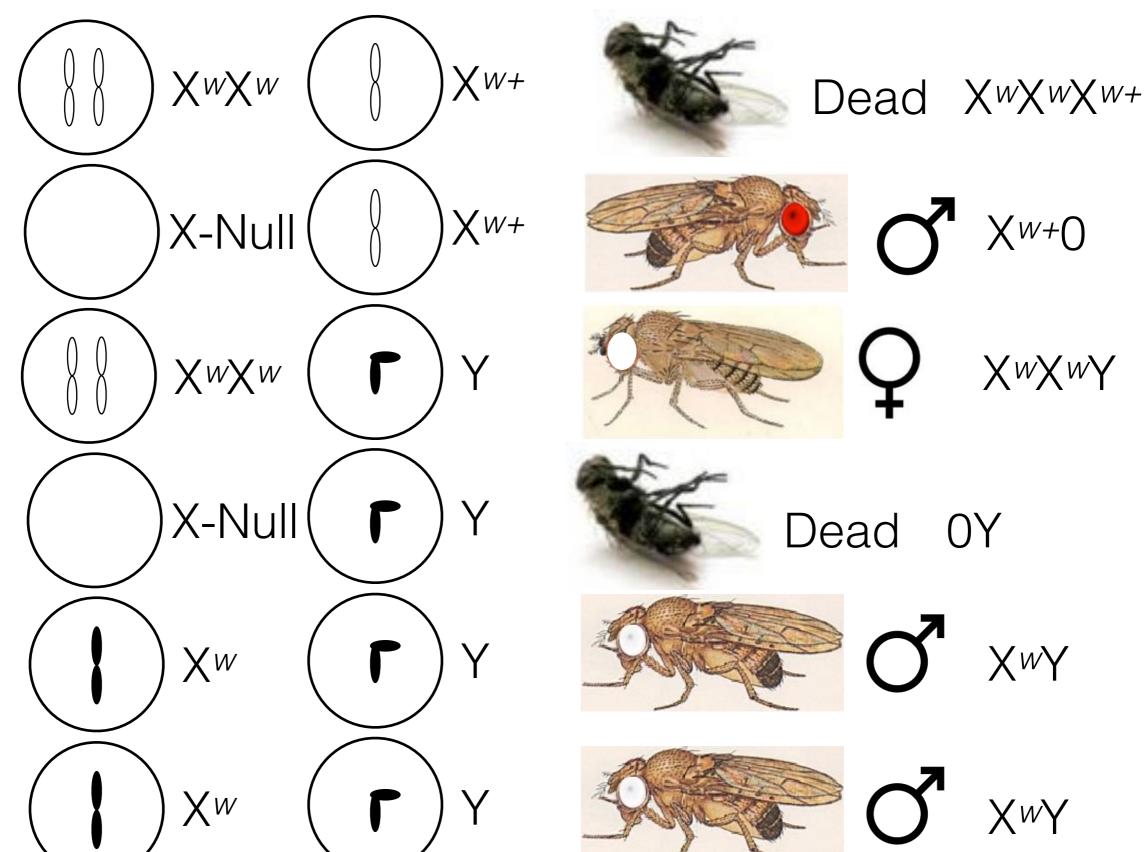
 $X^{w+}X^{w}$  $X^{w}Y$  1/2000 offspring

 $X^{w+0}$ 

red male white female

Meiosis II NDJ Male Female gametes gametes

Offspring



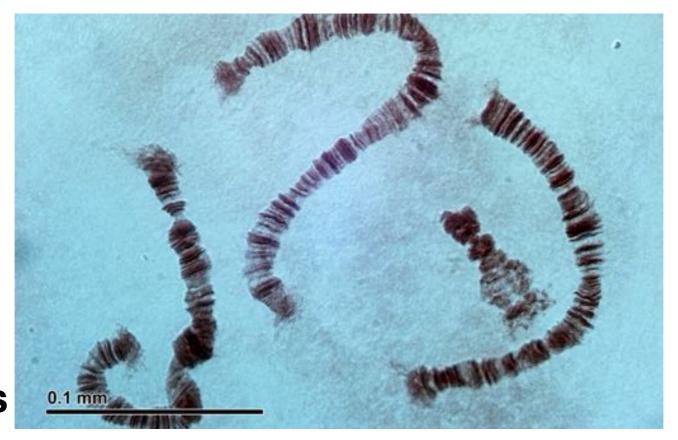
# The connections between chromosome NDJ and a trait was made by Stevens and Bridges



Nettie Stevens



**Calvin Bridges** 



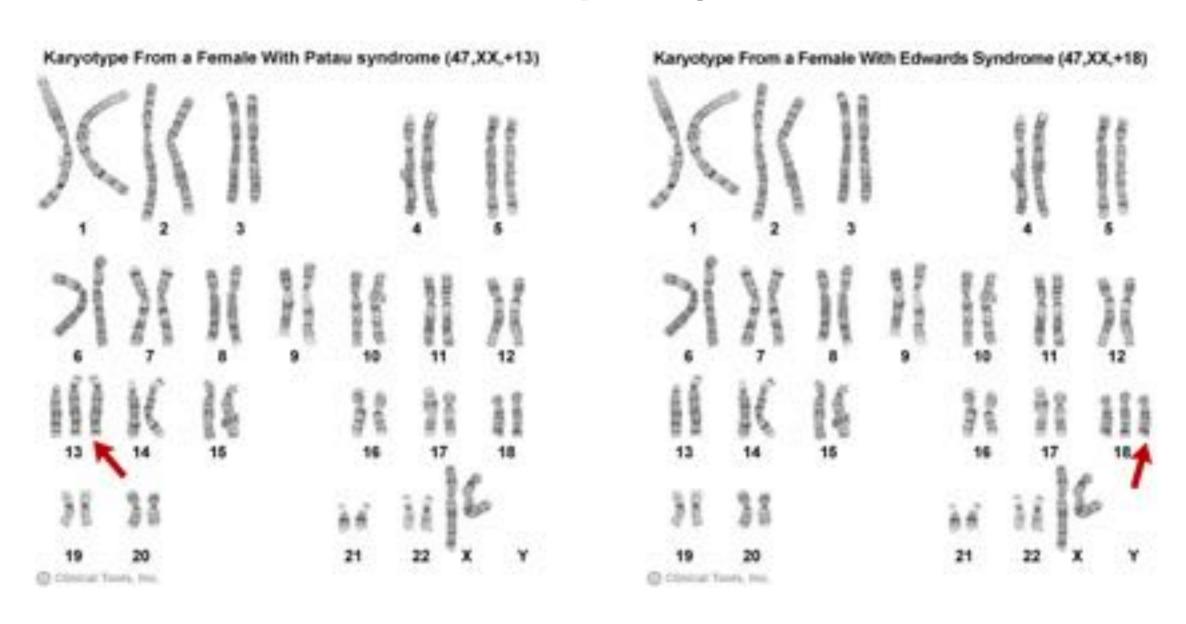
Polytene chromosomes

Why did the first cross not indicate to them that something weird was going on?

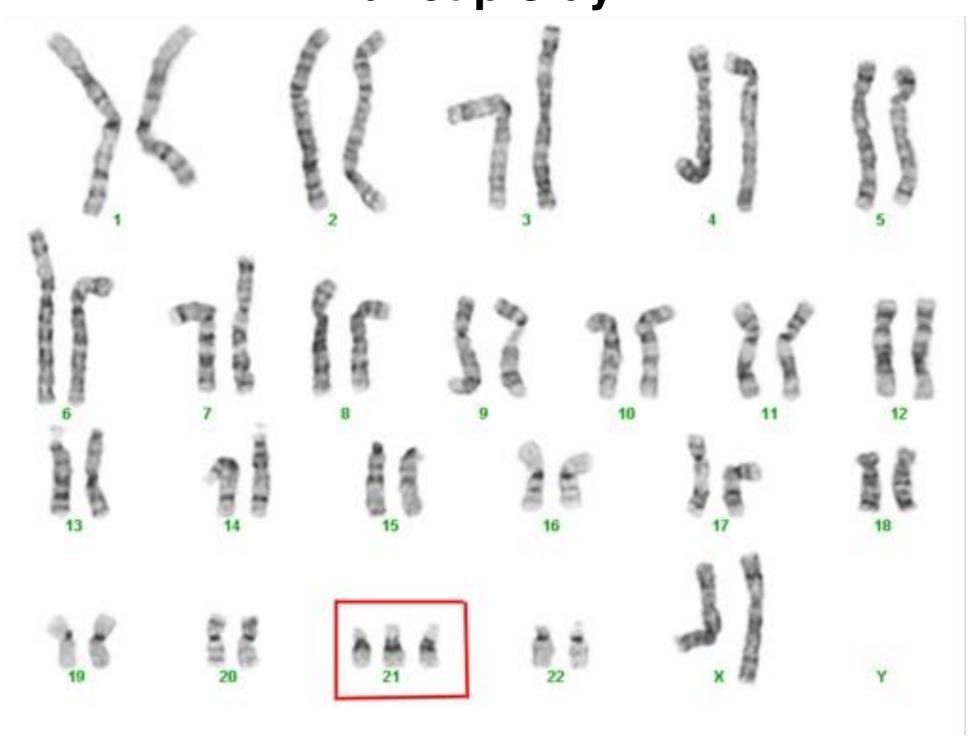
XwY Xw+Xw+

How can you tell the difference between Meiosis I NDJ and Meiosis II NDJ?

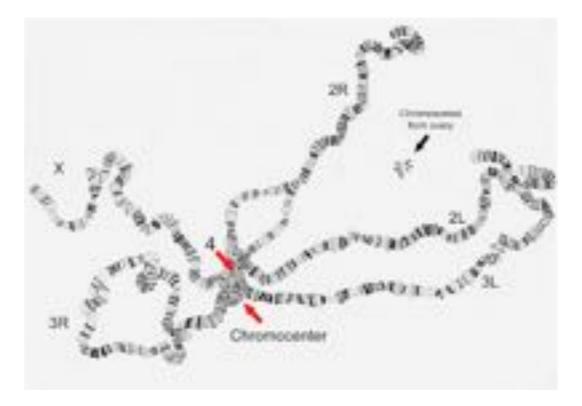
# Non-disjunction is a relatively common error not just the X chromosome aneuploidy

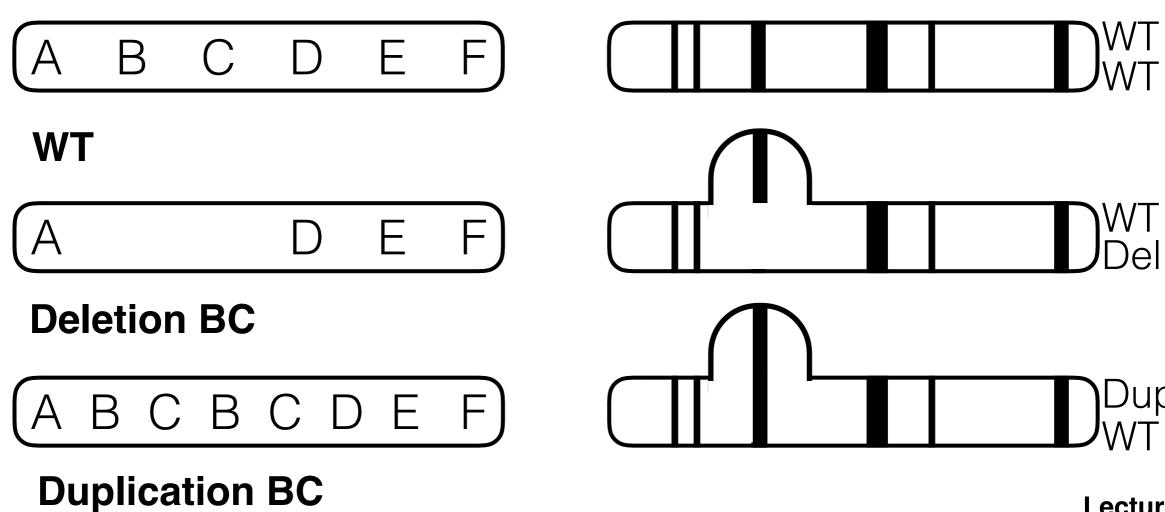


# Non-disjunction is a relatively common error not just the X chromosome aneuploidy



### **Chromosomal abnormalities**





**Lecture 2** 

### **Chromosomal abnormalities**





WT



**Inversion BCD** 

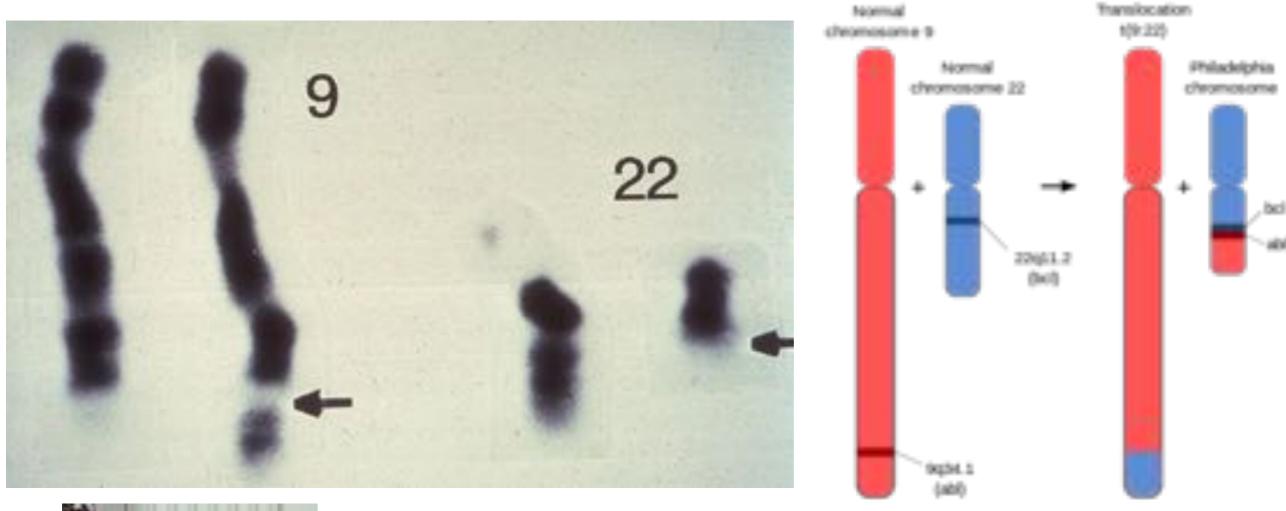


XYZDEF

Fusion of two chromosomes

**Translocation ABC-XYZ** 

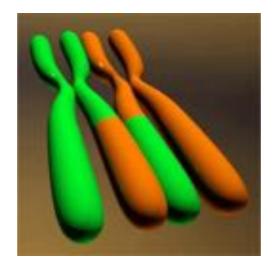
### The Philadelphia chromosome: translocation





Janet Rowley

# **Recombination and mapping**





Reginald Punnett William Bateson

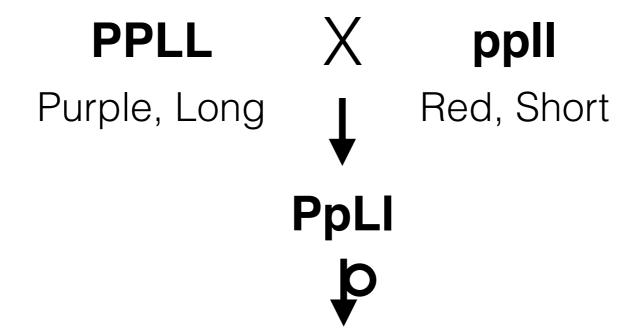


P= purple flower

p= red flower

L= long pollen

I= short pollen



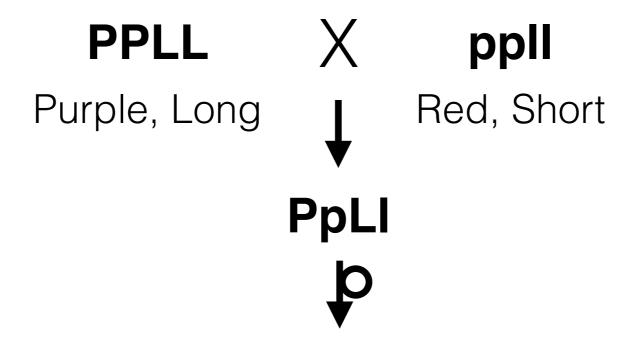
| Phenotype    | Expected number | Expected ratio |
|--------------|-----------------|----------------|
| Purple Long  | 215             | 9              |
| Purple short | 71              | 3              |
| red Long     | 71              | 3              |
| red short    | 24              | 1              |

P= purple flower

p= red flower

L= long pollen

I= short pollen



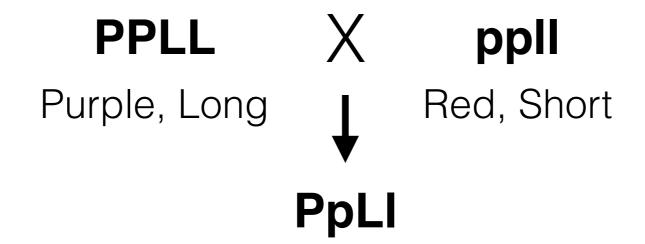
| Phenotype    | Expected number | Expected ratio | Observed number |
|--------------|-----------------|----------------|-----------------|
| Purple Long  | 215             | 9              | 284             |
| Purple short | 71              | 3              | 21              |
| red Long     | 71              | 3              | 21              |
| red short    | 24              | 1              | 55              |

P= purple flower

p= red flower

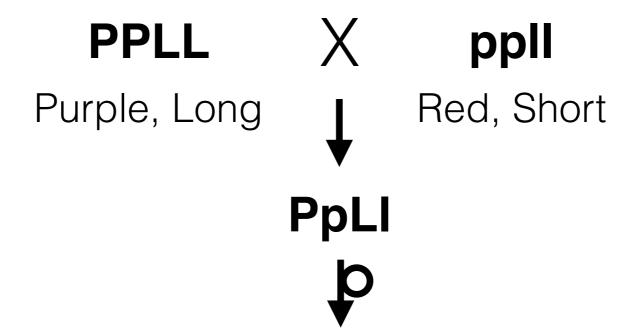
L= long pollen

I= short pollen



Parental = allelic combination found in parents (most abundant classes, always paired)

Recombinant = allelic combination NOT found in parents (least abundant classes, always paired)



| Phenotype    | Expected number | Expected ratio | Observed number |
|--------------|-----------------|----------------|-----------------|
| Purple Long  | 215             | 9              | 284             |
| Purple short | 71              | 3              | 21              |
| red Long     | 71              | 3              | 21              |
| red short    | 24              | 1              | 55              |

P= purple flower

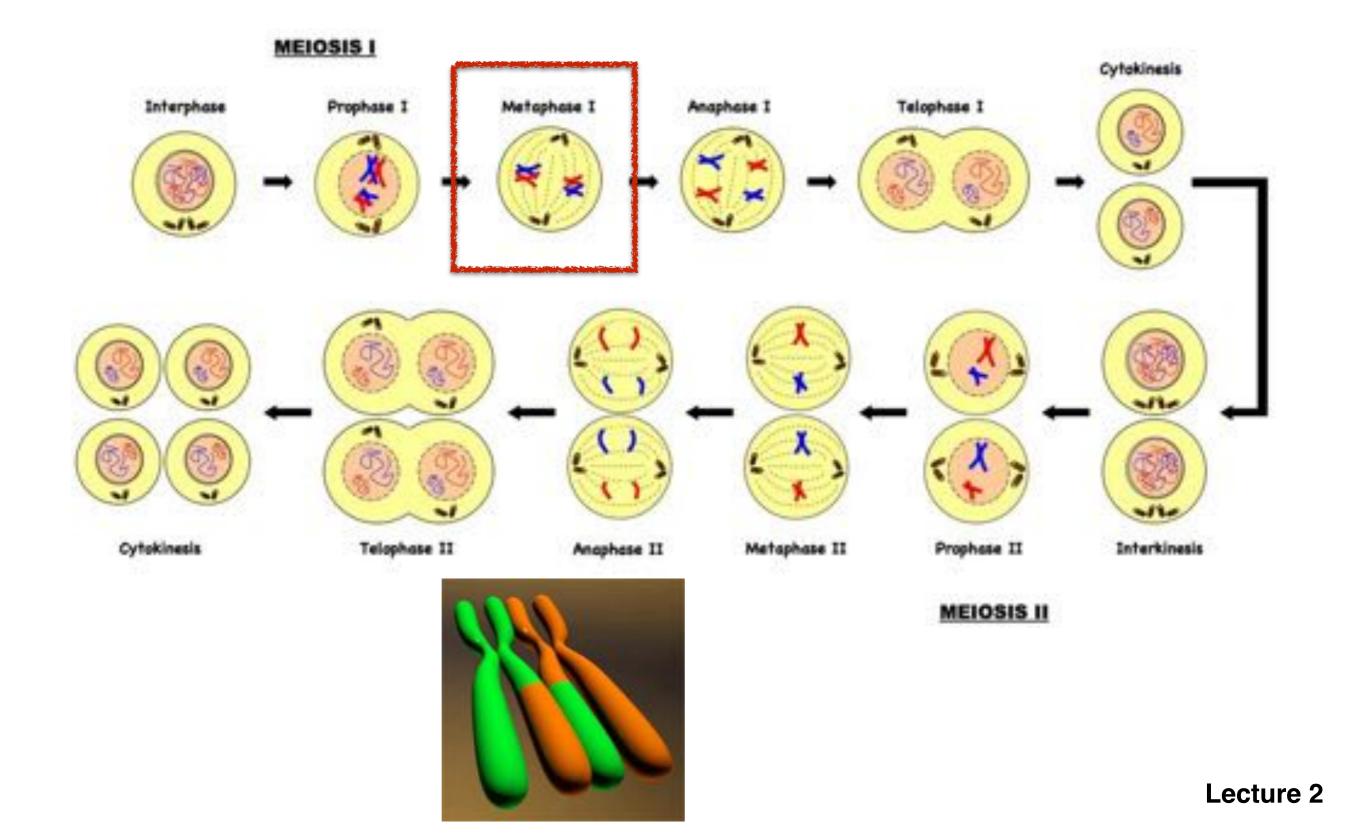
p= red flower

L= long pollen

I= short pollen

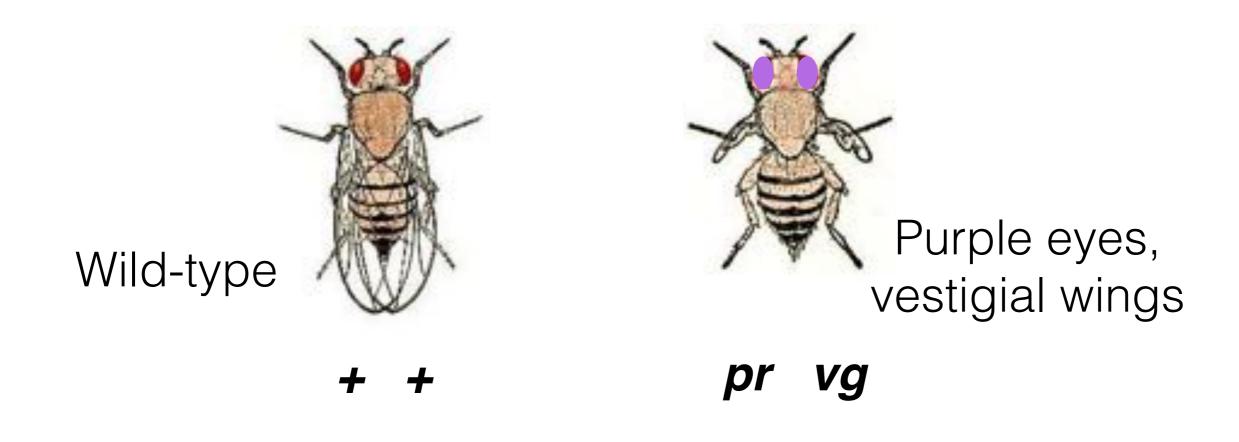
Which are recombinant and parental offspring?

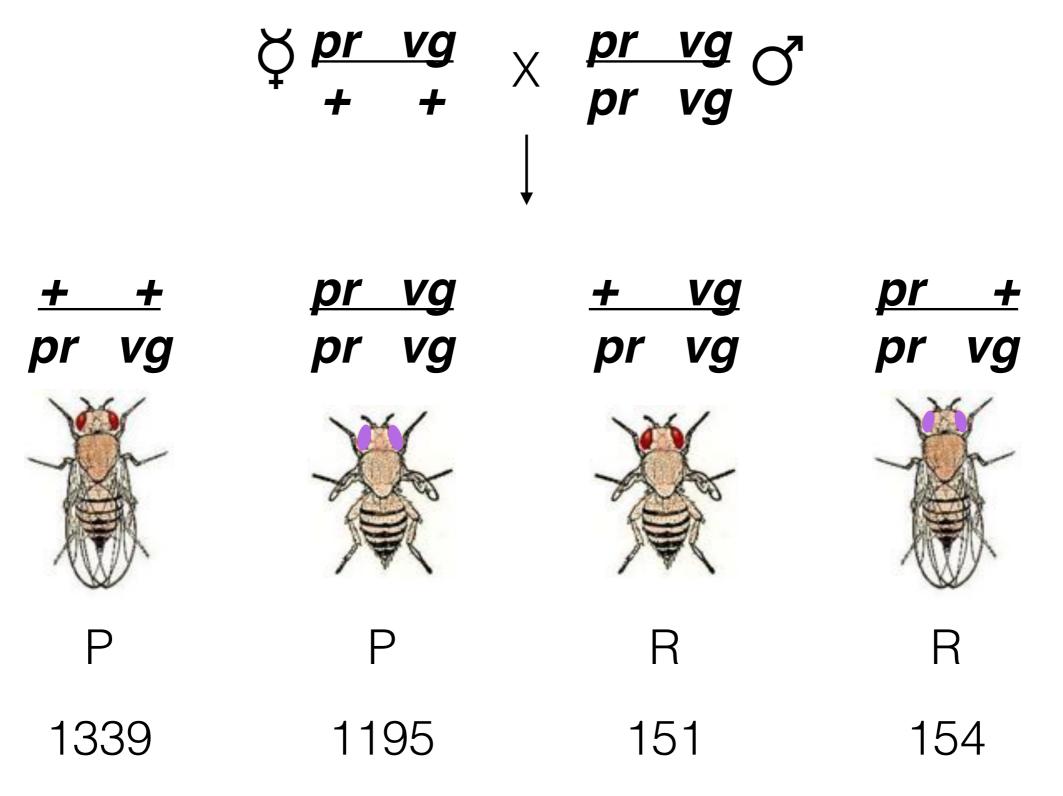
### Meiosis: A reductional division in two acts



# The fly room at Columbia







**Expectation is equal proportion of each class** 

Total = 2839



**Alfred Sturtevant** 

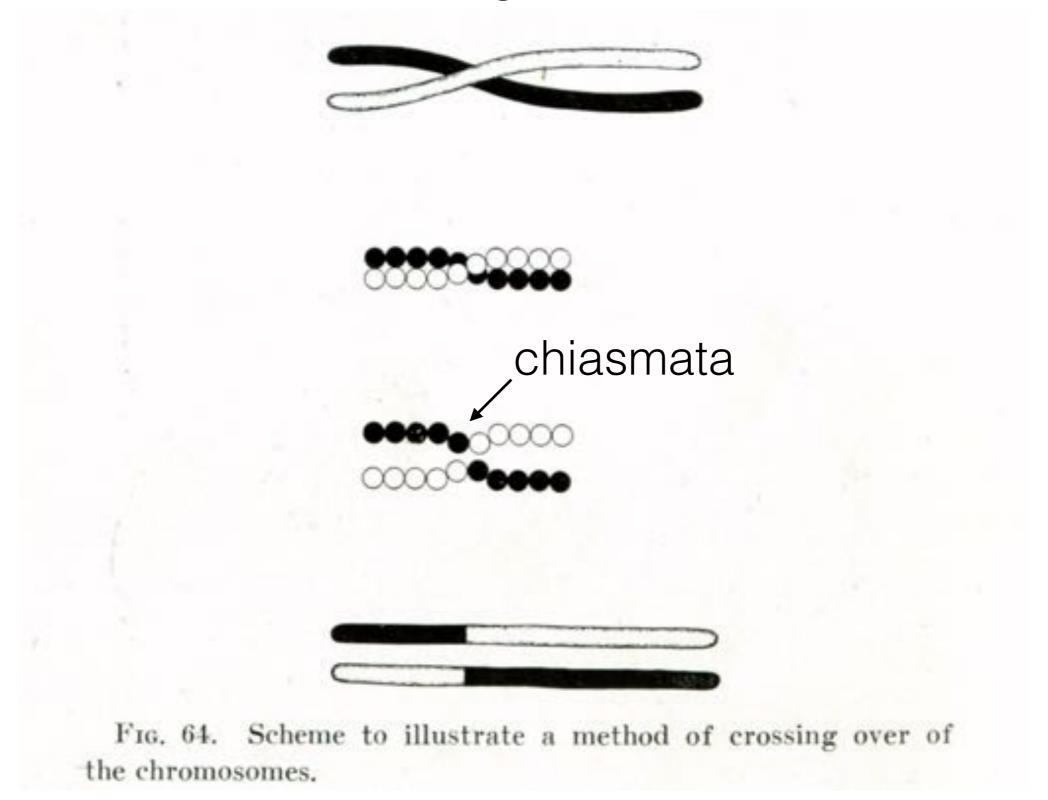
$$\frac{\text{Number of recombinants}}{\text{Total progeny}} \quad \times \quad 100 = \frac{\text{Recombination}}{\text{frequency}}$$

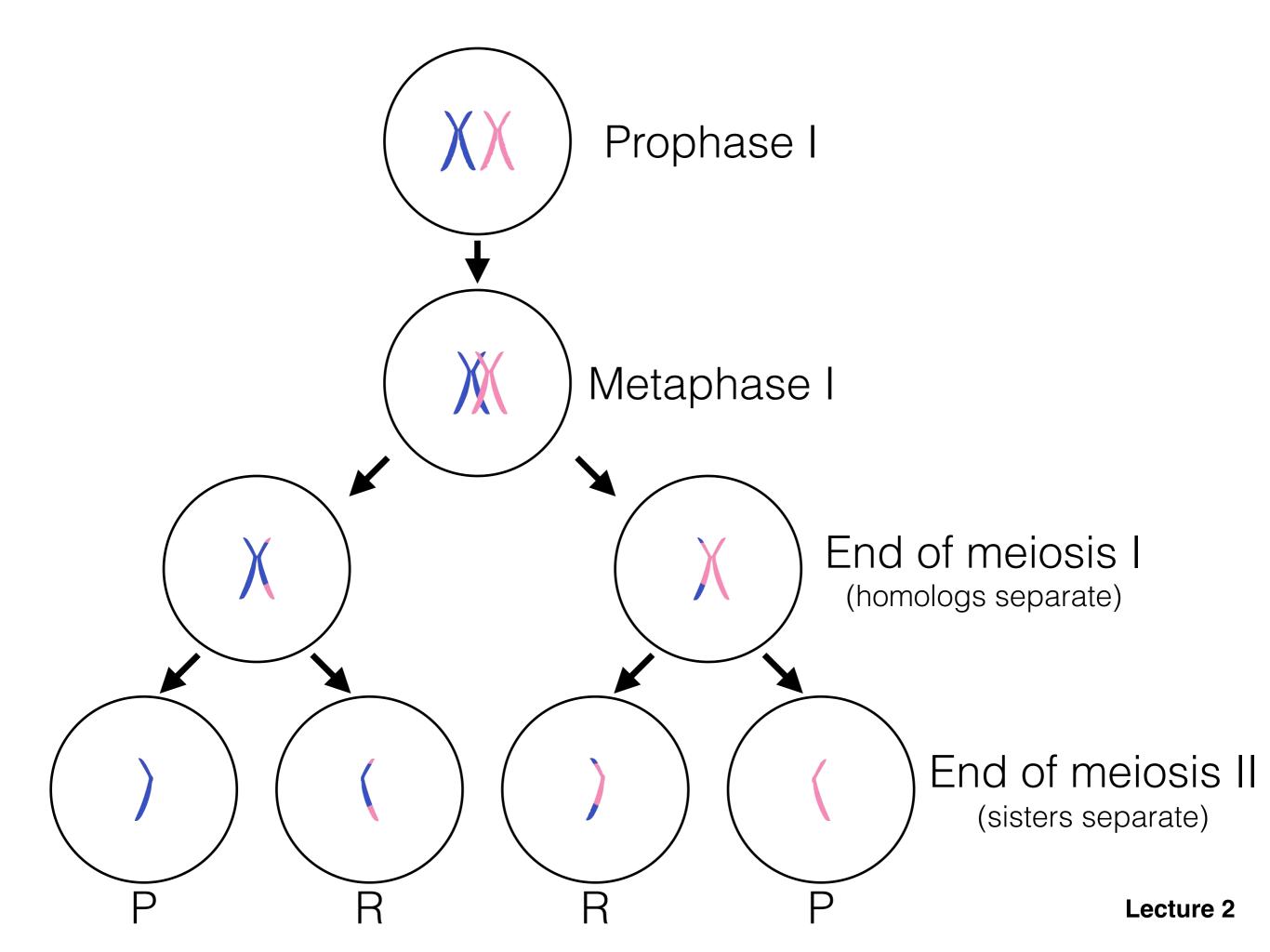
1% RF = 1 map unit = 1 centiMorgan

Total = 2839

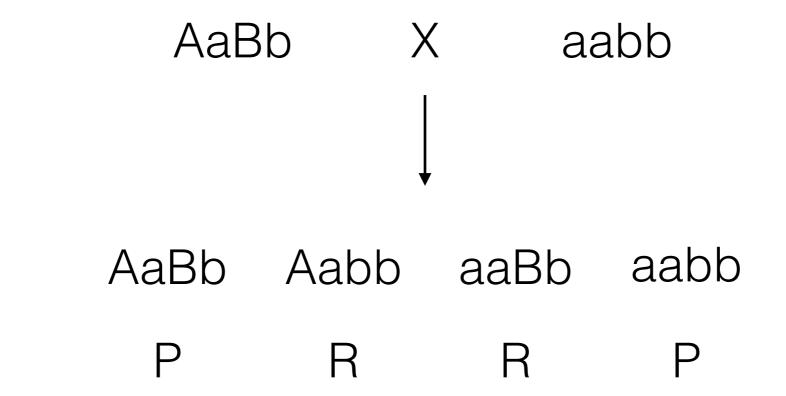
Total = 2839

# Recombination is the exchange of genetic material between homologous chromosomes



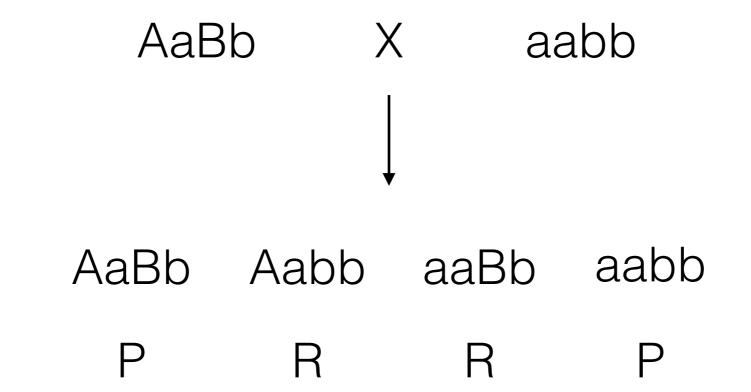


### Independent assortment defines the limit of linkage at 50 cM



All four classes occur in equal ratios

### Independent assortment defines the limit of linkage at 50 cM



All four classes occur in equal ratios

$$\frac{2^*x}{2^*x + 2^*x} \times 100 = 50\%$$

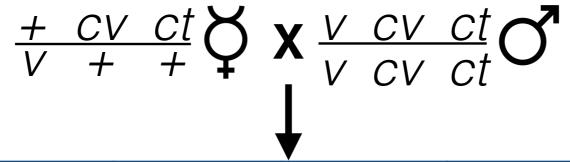
### A three-factor cross

| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
|------------------|------------------------|------------------|---------------------|
| Red              | No crossvein           | Cut wing         | 580                 |
| Vermillion       | Crossvein              | Normal wing      | 592                 |
| Red              | Crossvein              | Cut wing         | 40                  |
| Vermillion       | No crossvein           | Normal wing      | 45                  |
| Red              | Crossvein              | Normal wing      | 94                  |
| Vermillion       | No crossvein           | Cut wing         | 89                  |
| Red              | No crossvein           | Normal wing      | 5                   |
| Vermillion       | Crossvein              | Cut wing         | 3                   |

v = vermillion eyes

ct = cut wings

cv= crossveinless wings



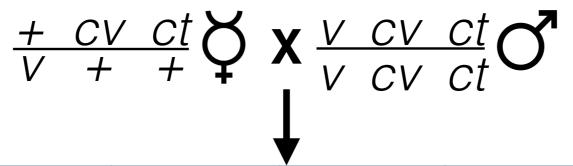
| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
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| Red              | No crossvein           | Cut wing         | 580                 |
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| Vermillion       | Crossvein              | Cut wing         | 3                   |

1. Determine parental class, label

v = vermillion eyes

ct = cut wings

cv= crossveinless wings



| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
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| Red              | No crossvein           | Normal wing      | 5                   |
| Vermillion       | Crossvein              | Cut wing         | 3                   |

| + | CV | Ct |
|---|----|----|
| V | +  | +  |
|   |    |    |

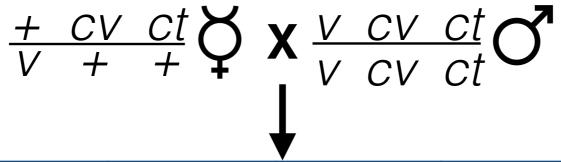
PRRRRR

- 1. Determine parental class, label
- 2. Are all classes present?

v = vermillion eyes

ct = cut wings

cv= crossveinless wings



| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
|------------------|------------------------|------------------|---------------------|
| Red              | No crossvein           | Cut wing         | 580                 |
| Vermillion       | Crossvein              | Normal wing      | 592                 |
| Red              | Crossvein              | Cut wing         | 40                  |
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| Vermillion       | Crossvein              | Cut wing         | 3                   |

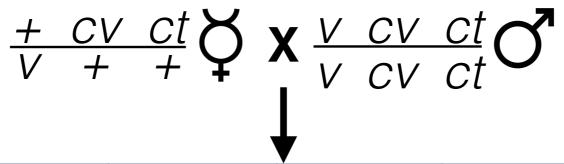
| CV<br>+ |   | FFF         |
|---------|---|-------------|
| CV<br>+ | _ | F<br>F<br>F |

- 1. Determine parental class, label
- 2. Are all classes present?
- 3. Least abundant class is double recombinant, tells gene in middle

v = vermillion eyes

ct = cut wings

cv= crossveinless wings



| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
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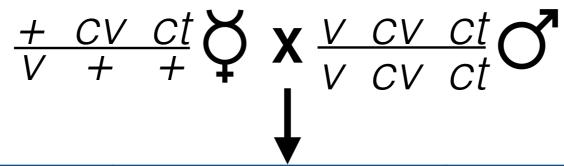
| + | CV | ct | F |
|---|----|----|---|
| V | +  | +  | F |
| + | +  | Ct | F |
| / | CV | +  | F |
| + | +  | +  | F |
| / | CV | Ct | F |
| + | CV | +  | F |
| V | +  | ct | F |
|   |    |    |   |

- 1. Determine parental class, label
- 2. Are all classes present?
- 3. Least abundant class is double recombinant, tells gene in middle
- 4. Write out the genotypes of the offspring

v = vermillion eyes

ct = cut wings

cv= crossveinless wings



| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
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| + | CV | ct |
|---|----|----|
| V | +  | +  |
| + | +  | Ct |
| V | CV | +  |
| + | +  | +  |
| V | CV | Ct |
| + | CV | +  |
| V | +  | Ct |
|   |    |    |

1448 total progeny

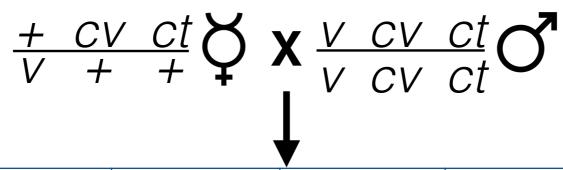
R

- 1. Determine parental class, label
- 2. Are all classes present?
- 3. Least abundant class is double recombinant, tells gene in middle
- 4. Write out the genotypes of the offspring
- 5. Calculate distance from one gene to middle gene **V to** *ct*

v = vermillion eyesct = cut wingscv= crossveinless wings+ = red eyes and normal wings

$$\frac{94+89+5+3}{1448} \times 100 = 13.2\%$$

Lecture 2



| Eye<br>Phenotype | Crossvein<br>Phenotype | Cut<br>Phenotype | Number of offspring |
|------------------|------------------------|------------------|---------------------|
| Red              | No crossvein           | Cut wing         | 580                 |
| Vermillion       | Crossvein              | Normal wing      | 592                 |
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| + | CV | ct |
|---|----|----|
| V | +  | +  |
| + | +  | Ct |
| V | CV | +  |
| + | +  | +  |
| V | CV | Ct |
| + | CV | +  |
| V | +  | Ct |
|   |    |    |

1448 total progeny

- 1. Determine parental class, label
- 2. Are all classes present?
- 3. Least abundant class is double recombinant, tells gene in middle
- 4. Write out the genotypes of the offspring
- 5. Calculate distance from one gene to middle gene
- 6. Calculate distance from the other gene to middle gene CV to Ct

v = vermillion eyes

ct = cut wings

cv= crossveinless wings

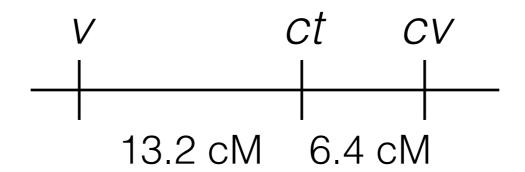
+ = red eyes and normal wings

$$\frac{40+45+5+3}{1448} \times 100 = 6.4\%$$

Lecture 2

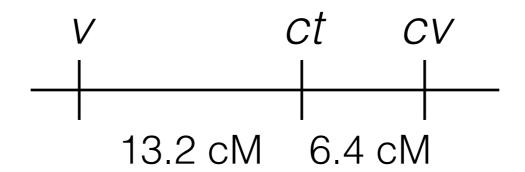
R

### Our first genetic map



- 1. Order by least abundant class
- 2. Arbitrary which genes on ends
- 3. Class *v* to *cv* undercounts because double recombinants look like parentals

### Our first genetic map



- 1. Order by least abundant class
- 2. Arbitrary which genes on ends
- 3. Class v to cv undercounts because double recombinants look like parentals

## We have a better way!