

# Genetics

Botany 3rd sem

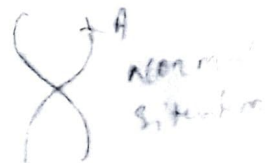
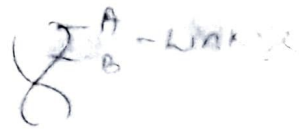
Second unit

# TYPES OF CYTOPLASMIC INHERITANCE

- Cytoplasmic inheritance are of three type
- Cytoplasmic inheritance involving essential organelle like chloroplast and mitochondria called as organellar inheritance.  
(Chloroplast and mitochondria show maternal inheritance)
  - Maternal effect depending indirectly on nuclear genes and involving no known cytoplasmic heredity unit ~~and~~ involving called as predetermination. In this maternal effect is determined before fertilisation.  
eg - shell coiling in snail (*Limnaea pergrana*)
  - Cytoplasmic inheritance involving dispensable and infective hereditary particle in cytoplasm which may or may not depend on nuclear gene called as ~~Dauer~~ modification.  
eg - sigma particle in *Drosophila*, kappa particle in *Paramecium*

## Linkage

- It is phenomenon of certain (2 or more) genes staying together during inheritance through generation with out changes or separations as they are present on same chromosome
- It is first discovered by Bateson & Punnett (1905) sweet pea (*Lathyrus odoratus*).
- They give coupling and repulsion phenomenon.
- Linkage can be thought of exception of ~~not~~ independent assortment.
- Morgan defined linkage as follows  
All the genes present on ~~same~~ a chromosome have the tendency to maintain original combination and to enter one and the same gamete





→ The tendency of two alleles to stay together is called coupling

→ The tendency of two alleles to avoid one another is called Repulsion.

\* If the two alleles, such as A and B come from the same parent they tend to enter the same gamete & transmit together (Coupling phase)

\* If the same alleles come from different parent they tend to enter into diff<sup>n</sup> gamete. (Repulsion phase)

A  
B  
A  
B  
a  
b  
AABB x aabb  
Cis

Repulsion phase  
AAbb x aaBB  
Trans

### SUTTON DUE of Linkage

→ Sutton in 1903 predicated that each chromosome contains a gene during meiosis chromosome move in to gamete as unit.

→ Hence all the gene which are situated in the same chromosome will be linked together as a result each species would have a specific number of group corresponds with the number of chromosome

\* Sc Bateson and punnet

Coupling and Repulsion hypothesis

→ B. P while working on sweet pea observe that flower colour and pollen shape tend to remain together & do not assort independently

Blue flower X Red flower  
long pollen. Round pollen

Blue long 7	Blue Round 1	Red long 1	Red Round 7
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→ Bateson and Punnett suggested that a gene comes from parents tends to same gamete & to be inherited together.

→ This phenomenon is called coupling. Similarly the gene comes from 2 diff<sup>n</sup> parent tends to enter diff<sup>n</sup> gametes and to be inherited separately & independently. This phenomenon is known as Repulsion.

## CHROMOSOMAL THEORY OF LINKAGE

Proposed by Morgan and Castle (1911)  
According to this theory.

- Gene that show in linkage are situated on same chromosome. (in parental combination, except for crossing over)
- Linkage genes are arranged in linear order.
- The distance bet<sup>n</sup> the linked genes in the chromosome determines the strength of the linkage.
- The closely located genes show greater linkage than the distant genes.
- Linkage genes depend on the original combination during inheritance.
- The linked genes show two types of arrangement: Cis and Trans.

Cis-arrangement - Dominant alleles of both the gene are present on the chromosome while their recessive alleles are present over the its homologous chromosome.

Trans-arr<sup>n</sup> - presence of dominant allele of one gene and recessive allele of the other gene on one chromosome (reverse arr<sup>n</sup>) of this genes present over its homologue.

# TYPES OF LINKAGE

There are two types of linkage

(1) Complete linkage

(2) Incomplete linkage

## COMPLETE LINKAGE

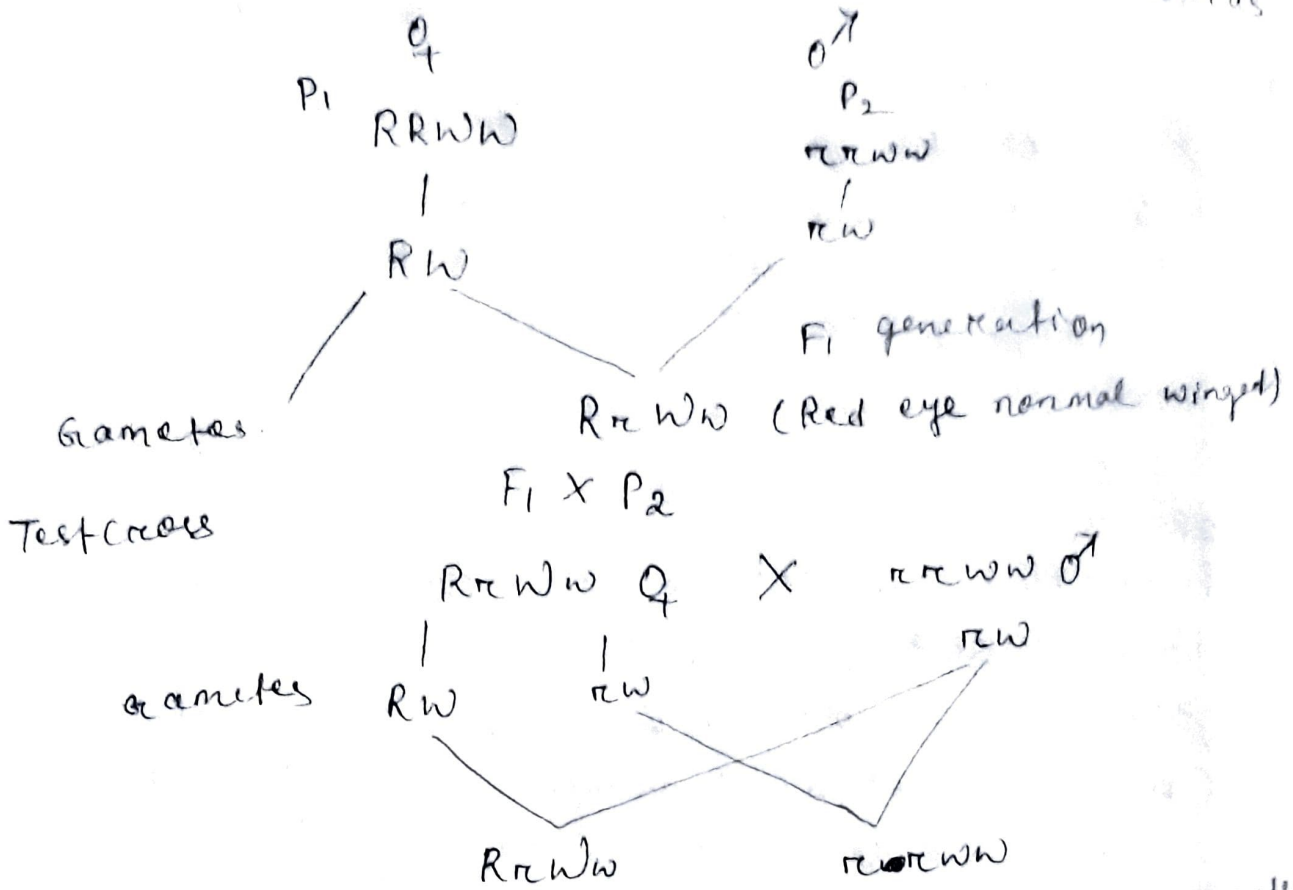
- Linkage in which gene always show parental combination. It never forms new combination.
- Crossing over is absent in it because gene are also located very closely on chromosome.
- \* When the two or more genes remain together closely for a number of generations, the linkage is said to be complete.
- \* It produce only parental combination
- \* It is very rare and found in only male *Drosophila* & female silkworm moth

ex-

- A Red eyes normal wings ( $RRWW$ ) female *Drosophila* is crossed with male *Drosophila* have purple eyes and vestigial wings ( $rrww$ )
- In  $F_1$  generation offspring are ( $RrWw$ ) Heterozygous Red eyed & normal winged.
- Now Heterozygous female fly ( $RrWw$ ) is self crossed with a ( $rrww$ ) *Drosophila* male (Having P.E and v.w)
- In  $F_2$  generation red eyes, normal wings (50%) and vestigial wings, purple eye (50%) in the ratio of 1:1
- No recombinant types are formed because linkage is complete & no crossing over occurs.



parents Red eyed, normal wings ♀ × purple eye, vestigial wings ♂



Due to complete linkage, instead of four gametes only two gametes are formed.

Test cross progeny	(R.E, N.W)	(P.E, V.W)
Ratio	1	1
Recombinants	Nil	

→ How ever, test cross bet<sup>n</sup> blue and long  
(BBkk) and double recessive (bbkk) give blue  
long (Hybrid) (43.7%)  
Red round (43.7%)  
Blue round (6.37%)  
Red long (6.37%)

→ The parental combination are 27.4% each  
due to linkage in genes on two homologous  
chromosome.

→ While in case of new combinations (12.6%) the  
genes get separated due to breaking of  
chromosome at the time of crossing over  
in prophase-1 of Meiosis

→ New combination due to incomplete linkage

## SIGNIFICANCE OF LINKAGE

- Linkage plays an important role in determining the nature of scope of hybridization and selection programmes
- It holds the parental character together and also restricts the appearance of new recombination
- It helps in maintaining the valuable traits of newly developed variety
- It allows the plant and animal breeding to combine all the desirable traits in a single variety.

## CROSSING OVER

- Crossing over is the exchange of chromosomal part between non-sister chromatids of a homologous pair resulting in recombination of gene.
  - The non-sister chromatids in which exchange of segments takes place are known as cross overs or recombinants while other chromatids not involved in exchange of segments are called non-crossover or parental type.
  - The term crossing over was first introduced by T. H. Morgan and Castell (1912) & defined it as the separation of linked gene ~~cross~~
  - The chiasma were first observed by Janssens (1909)
  - The no of chisma is variable. min<sup>m</sup> one chisma/bivalent is the rule. The highest number of chismata in long chromosomes of *Vicia faba* has been observed to be 12/bivalent
- There are two types of crossing over depending on the cell types
- meiotic crossing over / germinal cross over
  - Mitotic crossing over / somatic cross over



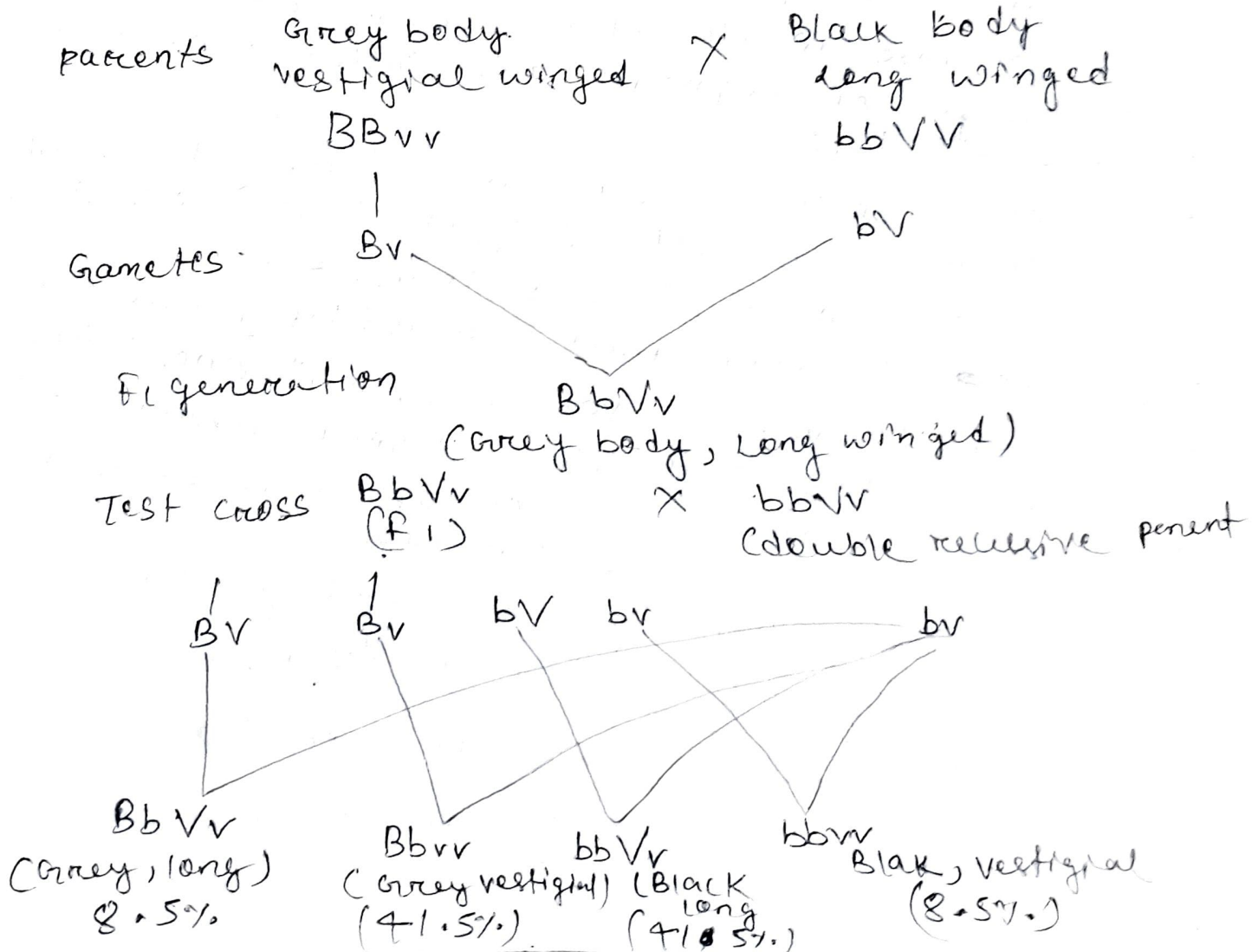
(i) Meiotic / germinal crossing over

- It occurs in all sexually reproducing organisms when they undergo meiosis during gametogenesis.
- It is primarily responsible for variation among the individuals of a species.

(ii) Mitotic / somatic crossing over

- It is rare occurrence in somatic cell during mitotic cell division.
- It has no significance from the point of view of inheritance.
- It is observed in Drosophila and Aspergillus nidulans.

\* Crossing over is absent in female silk worms and male Drosophila during sperm formation.  
ex - crossing over was reported in Drosophila by T. H. Morgan



parent in  $F_1$  generation show (BbVv) Grey body, long wing  
Drosophila

parental combination BbVv (Grey, vestigial) (41.5%)  
bbVv (Black, long) (41.5%)

recombinant combination BbVv (Grey, long) (8.5%)  
bbv (Black, vestigial) (8.5%)

→  $\frac{\text{no. of recombinant progeny in the test cross}}{\text{Total no. of progeny in the test cross}}$

OR  
$$CF = \frac{\text{no. R. Progeny in t.c.} \times 100}{\text{Total no R.P in t.c.}}$$

## MECHANISMS OF CROSSING OVER

crossing over occurs during meiosis of gametogenesis

- ① Synapsis
- ② Duplication chromosome
- ③ Exchange of chromosomal part
- ④ Termination

### Synapsis (meiosis)

→ During the zygotene stage of prophase-I the homologous chromosome move towards the each other and come to present side by side longitudinally

This pairing of homologous chromosome is Synapsis

→ The paired chromosome are called bivalent.

→ In some organism, a network of filamentous structure appears bet<sup>n</sup> two chromosome during the process during the process of pairing called Synaptonemal complex help in the synapsis

# DUPLICATION OF CHROMOSOME

- The bivalent undergoes duplication during pachyene stage.
- Each two homologous chromosomes splits longitudinally into chromatids.
- Hence 4 chromatids are produced & called tetrad <sup>stage</sup>.
- The two chromatids of a chromosome are attached to a single centromere so called sister chromatids.

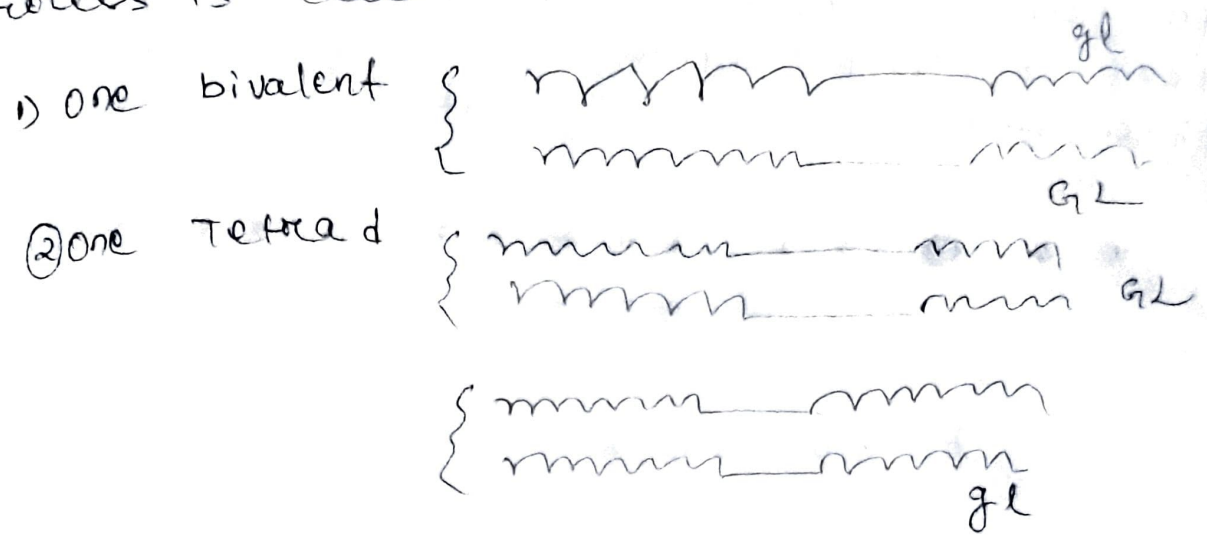
## Exchange of chromosomal part / crossing over

- The non-sister chromatids of a <sup>homologous</sup> chromosome intercross to each other at certain point called as chiasmata (~~centromere~~).
- At the chiasma, the chromatids break by endonuclease enzyme.
- The broken segments of non-sister chromatids are exchanged and fused with the other chromatids in presence of an enzyme ligase.
- The exchange of chromosomal segment between non-sister chromatids is called crossing over.



# TERMINIALIZATION

- After the exchange of the chromosomal segment the non-sister chromatids repel each other
- The separation begins from the centromere and moves towards the end of chromosome
- During separation of chromosome, the chiasmata move towards end of the chromosome. This process is called terminalisation.



# Significance of crossing over

→ crossing over provides direct proof for the linear arrangement of gene.

→ The segment of homologous chromosome are interchanged through crossing-over. Hence crossing over provides origin of new character and genetic variation.

→ crossing over leads to construction of linkage map and genetic mapping of chromosome.

→ crossing over plays an important role in the field of breeding to improved the varieties of animals and plants.

## Characteristic of sex-linked inheritance

→ Individual showing a recessive sex-linked trait is noticeable higher in heterogametic sex (male *Drosophila* & human & female birds) the homogametic sex (female *Drosophila* & humans & male birds)

→ In male Human & *Drosophila* the sex linked traits are not transmitted directly male parents to their male progeny.

→ In male human or *Drosophila* transmits its sex linked gene to all its daughters. These daughter transmitted this gene to half of their male progeny

→ Hence a sex-linked gene passes from male to female then back to male. This inheritance pattern is known as criss-cross inheritance

## Cases

→ The inheritance is cause by mainly two reason.

- The location of a gene in the X chromosome
- The absence of its allele in the Y chromosome

## Partial sex linkage

The human X and Y chromosome are morphologically distinct but they pair during meiosis in male cell the pairing occurs in the 2 telomeric region, which are called pseudoautosomal regions (PAR),

There are 2 types

① PAR 1

② PAR 2

→ The genes located in PAR1 & PAR2 are also present in the X chromosome. But these genes do not show the typical inheritance patterns for sex-linkage. Because these genes have alleles in the Y chromosome as well.

→ As a result inheritance process responsible that of autosomal genes. These phenomenon is called partial sex-linkage and the chromosome region involved in it are referred to as pseudoautosomal regions.

### Factor influenced crossing over

- 1- sex - in drosophilla crossing over is completely suppressed in male but very high in female. There is a tendency of reduction of crossing over in male mammals.
- 2- mutation - Gower first discovered that mutation reduced crossing over in all the chromosomes of drosophilla.
- 3- inversion - crossing over is suppressed due to inversion.
- 4- Temperature - plough shown that when a drosophilla is subjected to high & low temperature variation, the percentage of crossing over in certain parts of the chromosome is increased.
- 5- X-ray effect - muller demonstrated that X-ray radiation increases crossing over near centromere.
- 6- Age - Bridges demonstrated when the female drosophilla becomes older, the rate of crossing-over increases.
- 7- Nutrition - High calcium diet in young drosophilla decreases crossing over but diet deficient in calcium increase crossing over.
- 8- Heterochromatin - It decreases crossing over. (Dark/Deep colour of chromosome strands)

### Chromosome maps →

The phenomenon of linkage and crossing over has established following facts

- (i) The genes are arranged in a linear order in a chromosome.
- (ii) The linkage group in an organism are equivalent to the number of chromosome pairs.
- (iii) Frequency of crossing over or recombination bet<sup>n</sup> genes depends upon the distance between of crossing over are more bet<sup>n</sup> 2 distinctly placed genes similarly the change of crossing over are less bet<sup>n</sup> 2 closely placed genes.



→ The strength of linkage is inversely proportional to the distance betn linked gene

→ The position of gene is very definite in chromatids  
Basing the following observation. Sturtevant in (1913) developed the idea the frequency of crossing over can be caused as a title for determination of relative distance betn the genes in a linkage group.

### Chromosome maps

→ The chromosome map is the graphic representation of the relative distance betn the linked genes expressed in the percentage of recombination among the genes in a linkage group.

construction of chromosome maps (linkage maps) process.

The construction of chromosome map includes following

① Determination of linkage group:-

The exact no. of chromosome of a given species is determined. The total no. of genes are determined by hybridisation experiment in betn wild & mutant strain.

→ The no. of phenotypic trait which remain always linked is determined. The different linkage group of that species are found out.

② % Determination of map distance:-

The distance betn genes is determined by a unit called "morgan unit" or "map unit"

1 morgan unit represent 100% crossing over

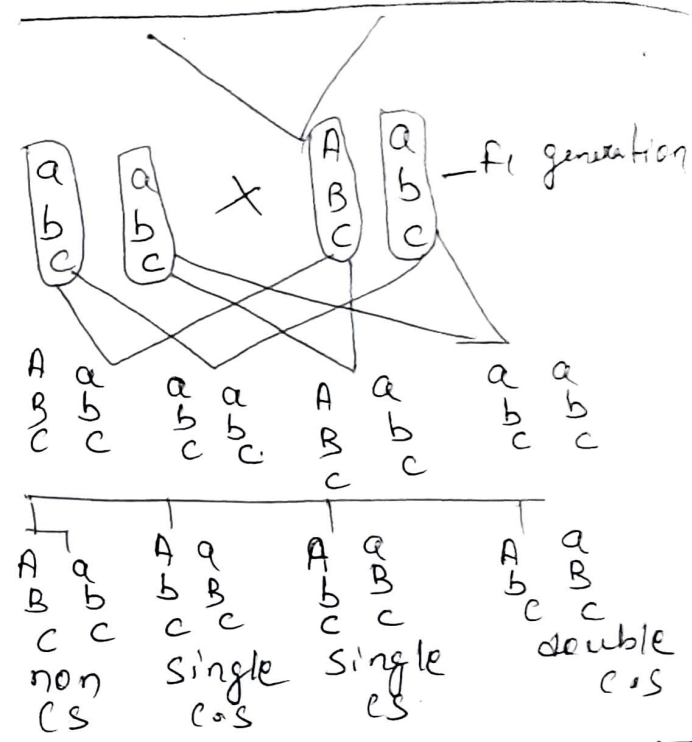
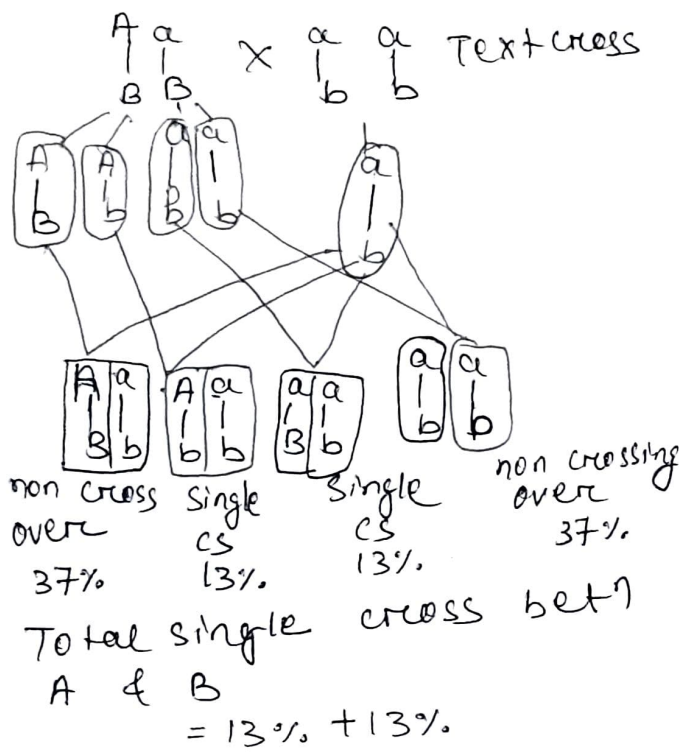
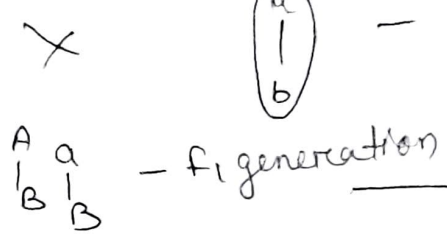
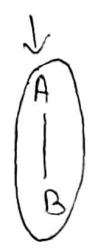
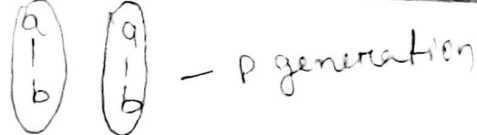
1 crossing over is expressed in 1 centimorgan or 1 map unit.

### Expt

- Suppose the percentage of crossing over betn gene A & B is 26%. then the distance betn A & B is 26 cm or map unit.

③ 2 point of test cross:-

The percentage of crossing over betn linked gene is capa by test cross



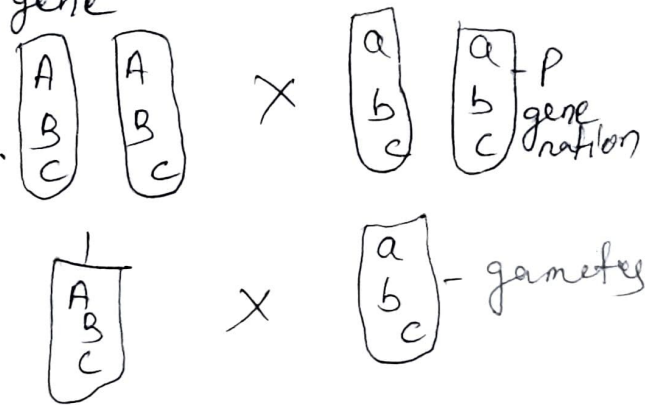
36%	36%	9%	9%	4%	4%	7%	7%

These = 26% distance bet<sup>n</sup> A & B = 36

Double crossing over don't occur bet<sup>n</sup> the genes in which the distance 5cm & less

4) Three point test cross

A Three point of test crossing over (involving 8 genes) gives the accurate information about the distance bet<sup>n</sup> the gene



Non-crossing over (C.O) = 72%  
 Total single C.O bet<sup>n</sup> A & B = 18%  
 Total single C.O bet<sup>n</sup> B & C = 8%  
 Total double C.O bet<sup>n</sup> B & C = 2%  
 Distance bet<sup>n</sup> A & B = Total single C.O + Total double C.O bet<sup>n</sup> A & B = 18% + 2% = 20%  
 Distance bet<sup>n</sup> B & C = Total single C.O + Total double C.O bet<sup>n</sup> B & C = 8% + 2% = 10%  
 10 cm or map unit



### 5 - Determination of gene order

Distance bet<sup>n</sup> A & C = single c.o + single c.o between 2  
bet<sup>n</sup> A & B + double crossing over  
 $= 20\% + 10\% - 2(2\%)$   
 $= 30\% - 4\% = 26\%$

$= 26$  or map unit

### 5 Determination of gene order

After knowing the relative distance bet<sup>n</sup> the genes of linkage group, it becomes easy to place in their proper linear order

Suppose the distance bet<sup>n</sup> the genes

$$A-B = 12 \text{ cm}$$

$$B-C = 7 \text{ cm}$$

$$A-C = 5 \text{ cm}$$

Case-I let gene A is in the middle (B-A-C)

$$\boxed{B \ 12 \ A} \quad \boxed{A \ 5 \ C} = 17 \neq 7$$

$\boxed{B \ 7 \ C}$  The distance B-C are not equable

Case-II let gene 'B' is in the middle (A-B-C)

$$\boxed{A \ 12 \ B} \quad \boxed{B \ 7 \ C}$$

$$\boxed{A \ 5 \ C}$$

The distance A-C are not equable

Case III  $\boxed{A \ 5 \ C} \quad \boxed{C \ 7 \ B}$

$$\boxed{A \ 12 \ B}$$

The distance bet<sup>n</sup> A-B are equable therefore (C) must be in middle.

Combine map segment  
 Finally the different segments of a complete genetic map.

$$\boxed{A \ 5 \ C \ 7 \ B'}$$

$$\boxed{D \ 4 \ E \ 8 \ F}$$

$$\boxed{A \ 20 \ C}$$

### Interference

The tendency of one crossing over (c.o) to interference with the c.o is called interference  
 → Because one chiasmata form reduced production adjacent chiasmata

→ The strength of interference is expressed in the form of co-efficient of coincidence

$$\text{co-efficient of coincidence} = \frac{\% \text{ actual double c.o}}{\% \text{ expected double c.o}}$$

### EXP-

Suppose expected double c.o are 2% or 0.2% & observed c.o is 1%

The co-efficient of coincidence

$$= \frac{1\%}{0.2\%} = 0.5\% \text{ or } 50\%$$

co-efficient of ~~interference~~  
 Interferences = 1.0



# Genetics

unit-11

No-1

- 1- The Linkage was first discovered (1x10-10) by \_\_\_\_\_ and \_\_\_\_\_
- 2- Linkage is exception of \_\_\_\_\_ principle.
- 3- crossing over is absent \_\_\_\_\_ type of linkage.
- 4- crossing over is exchange of \_\_\_\_\_ part
- 5- The genes are arranged in linear order in a \_\_\_\_\_
- 6- The strength of linkage is inversely proportional to the distance between \_\_\_\_\_.
- 7- The distance bet<sup>n</sup> genes is determined by a unit called '\_\_\_\_\_'
- 8- Who is the father of linkage \_\_\_\_\_.
- 9- New combination is occurs in \_\_\_\_\_ type of linkage.
- 10- The x and y chromosome are pair during \_\_\_\_\_ Division.

No-2

## Short note

(3x5-15)

- 1- crossing over
- 2- coincidence
- 3- interference
- 4- chromosome
- 5- cis and Trans-arrangement of gene.

No-3

## Long question

(8x5-40)

- 1- Briefly describe the linkage & its significance.
- 2- explanation of complete & incomplete linkage.
- 3- what is the chromosomal theory of linkage.
- 4- Describe the two factor & 3 factor crosses.
- 5- Chromosome maps
- 6- Describe the characteristic of sex-linked inheritance.