

Culture media

- ↳ Broth
- ↳ Plate / Solid agar / Gel.

Turbidity

Bachchan College
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Gene mapping in phages → transduction

~~Open system!~~
~~Complex system!~~ → Research!
~~Research!~~
No University and, Vantage
Rashed Rahman

2) Strain of T2 bacteriophage.

Plaque
trans-
apparente

→ At least 2 genders.

$\frac{1}{2} \times \frac{1}{2}$

九

Infect both

1 and 3/2

一、接

1. Gene Gene Gene

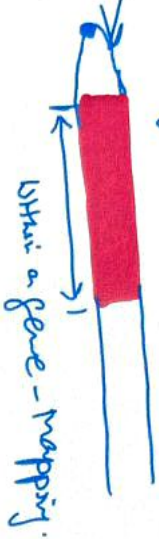
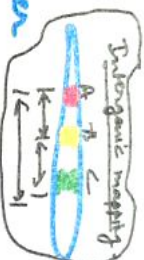
Intergenic mapping

In 1956 and 1960 - Seymour Benzer

→ Fine structure of gene → Physically on a chromosome.

Whether the gene is monocistronic or polycistronic

Cistron is size of cistron.



Benzer's mapping technique

① Wild type T₄ phage → normally produce small plaques
with round edges.

② → Mutants → K or X-mutant → rapid lysis

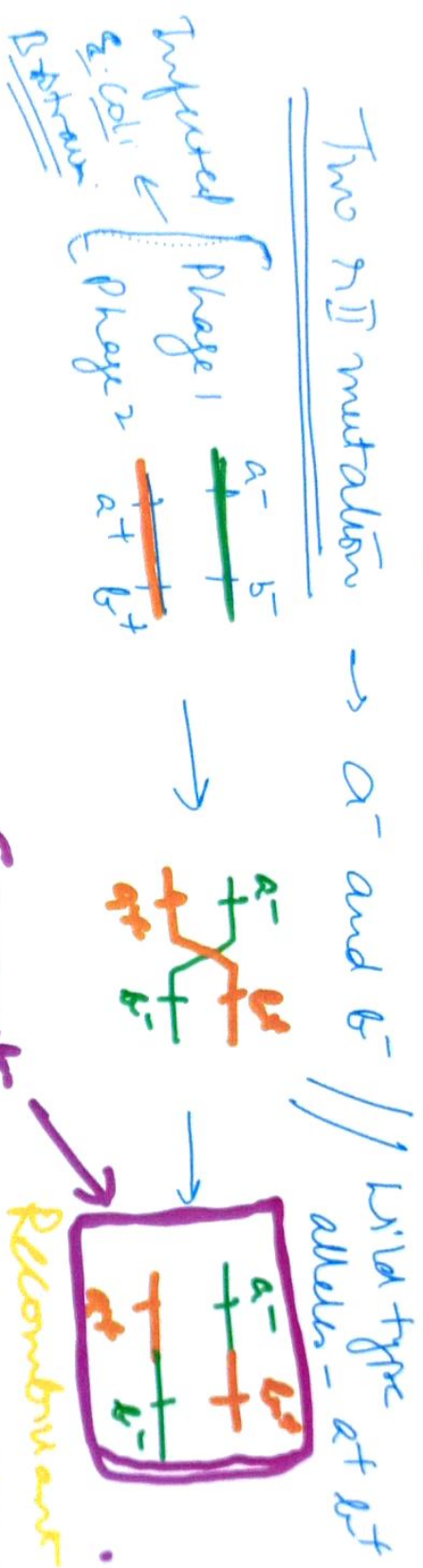
Produces larger plaques and sharply defined edges

Wild type T₄ phage produces typical plaques on E. coli strains K and B
the mutants → K and B mutants → produce K plaques on B strain
(bacteriophages) but no plaque on K strain
E. coli

Benzel collected thousands of π II mutants

He simultaneously infected bacterial cells with two diff. mutants and looked for recombinant progenies!

↳ Having mutations at various different locations!



Recomb. progeny Phages

↳ Growing $E. coli$ K strain.

Each recombinant produces an equal no. of double mutants (a^-b^-) and wild type

Chromosomes — a^+b^+

So the no. of recombinant progenies should be twice no. of wild types plaques that appear on K strain

$$R.F. = \frac{2 \times \text{No. of Plaques on E. coli K strain}}{\text{Total no. of Plaques on E. coli B strain}}$$

(Recombinant frequency)

↓ R.F.

Proportional to physical distance along the chromosome

Revealing the positions of the diff. mutations w.r.t. to

HII region of the phage chromosome.

∴ more the

Benzel's procedure for mapping λ mutants.

Friday, 23 October 2020 2:05 PM

$\lambda \rightarrow$ Rapid lysis

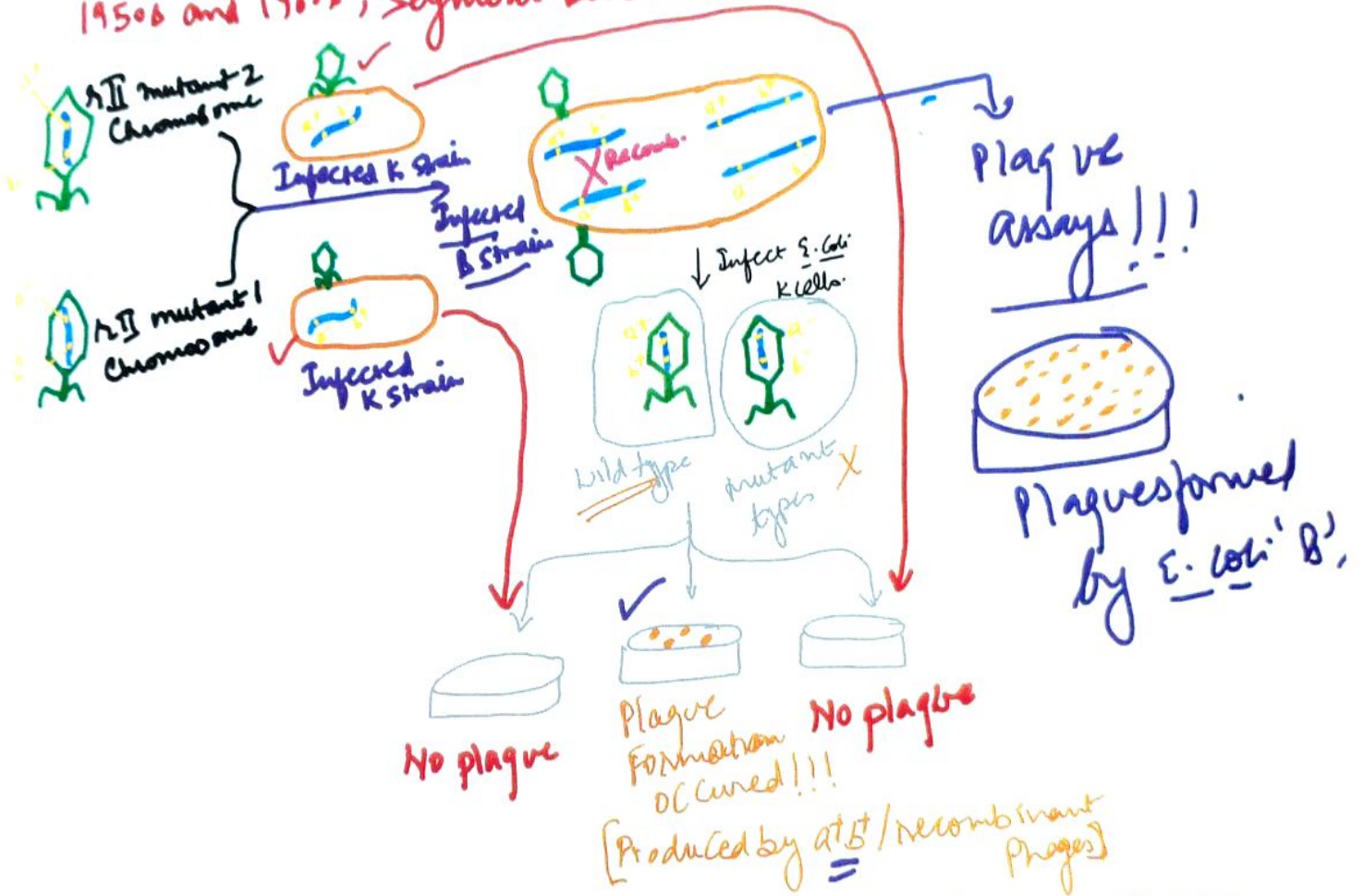
II sub group of mutant population

Mutant type bacteriophages has characteristic

$\rightarrow$ Can infect only 'B' strain of E. coli.

Whereas wild type bacteriophages could infect both 'K' as well as 'B' strains of E. coli

1950s and 1960s, Seymour Benzer



$$\text{Recombination frequency (\%)} = \frac{2 \times \text{No. of plaques on E. coli K strain}}{\text{Total no. of plaques on E. coli B strain}}$$

Complementation = $\frac{a^+ b^+}{a^- b^-}$

we conducted mutation - ? π II belong to same gene or different genes
 this exp then the relationship b/w gene and gene
 was not known, only it known that gene is
 "functional unit of heredity"

Complementation - 2 states $\leftarrow$ Cis - $\frac{a^- b^+}{a^+ b^-}$
 Trans - $\frac{a^- a^-}{b^+ b^+}$

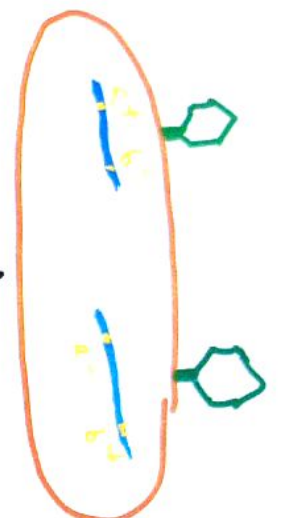
Travis Metz. $\frac{a^+}{a^-} \frac{b^+}{b^-} \rightarrow$ non-functional allele



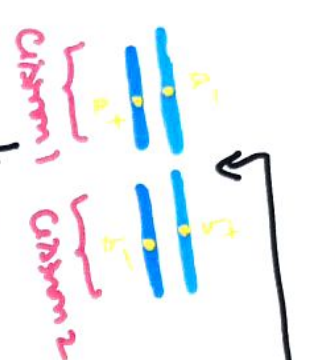
Functional protein can be formed.



! important!

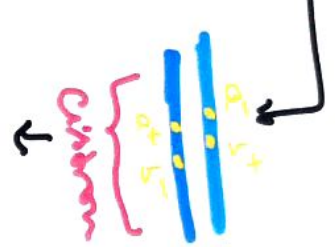


Simultaneous by EcoRI & Sma



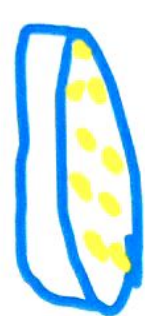
Protein B

one of the allele
Since two different loci (a & b)
is in linkage (a⁺, b⁺) therefore
a functional protein / peptide chains
are going to be formed



No functional protein formed!

Since both mutations
on either of the chromosome
is present or located at
anywhere in the gene locus
therefore a functional
peptide / protein can never
be formed.



Protein
will form



no protein