

2. Mutagenesis

Screen:

Visual: flower morphology, pigment color,

Biochemical pathway mutant

auxotrophy: $his3^-$ in yeast; arg^- in *Neurospora*; $trp1^-$ in plant

Reporter gene expression (luc bioluminescent protein from firefly)

Selection:

trp pathway: 5-methylantraniolate \rightarrow 5-methyltrp (toxic)

ADH (alcohol dehydrogenase): Allyl alcohol \rightarrow acrolein aldehyde (toxic)

Lethal mutations: (such as house keeping genes)

- maintain as heterozygote

- weak hypomorph

- conditional such as temperature-sensitive mutants in T4 phase (p220-221)

Lecture 4: Mutant Characterization

Dominant and recessive mutations

Complementation tests

Biochemical pathways

Epistasis analyses

Read: 206-208; p214-215; 220-224

Fig. 7.14; 7.15; 7.20; 7.25; 7.26; 7.27

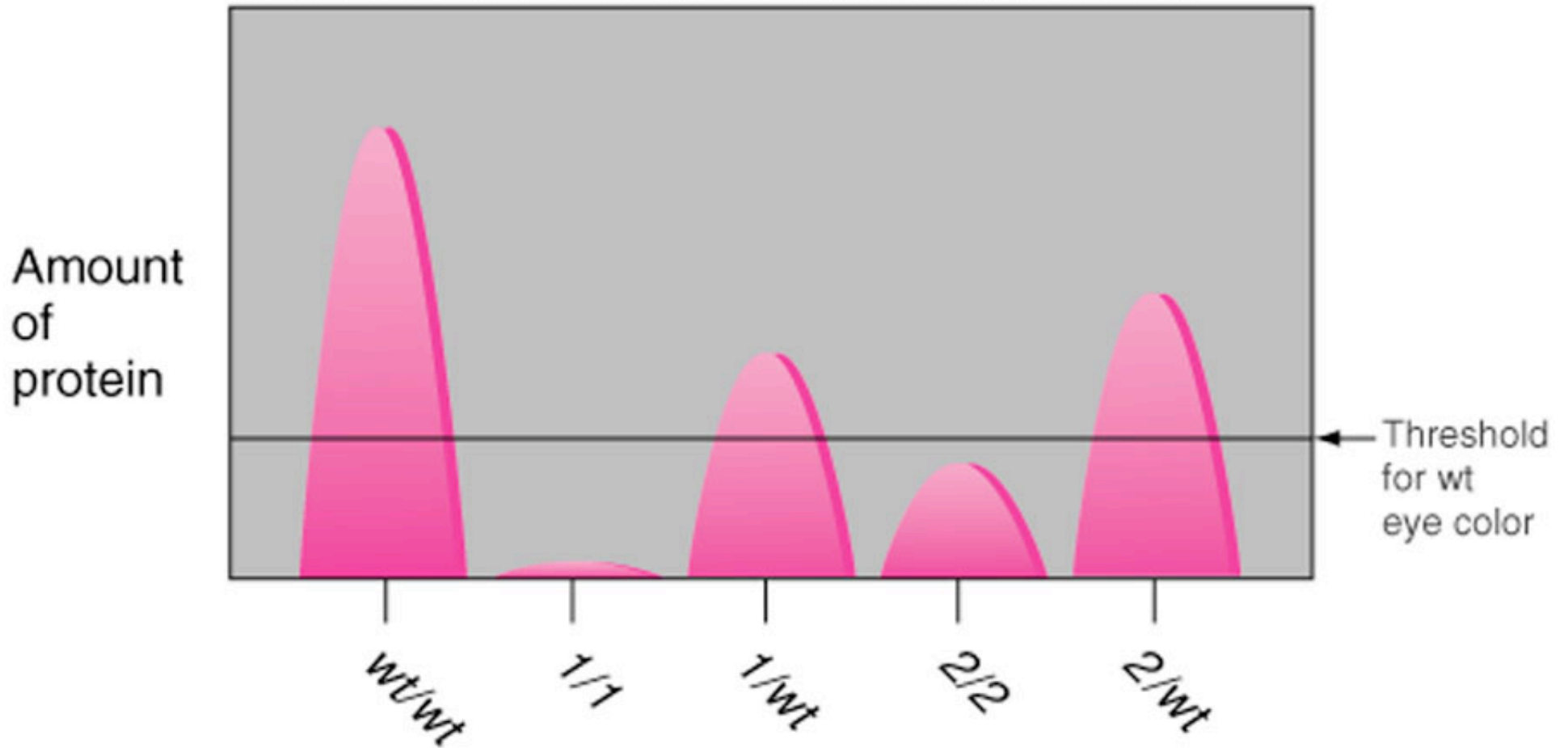
Homework#1 will be posted today!

Terminology about different mutations

- a) null mutation- complete absence of activity
- b) loss-of-function (hypomorph) - reduced activity
- c) gain-of-function (hypermorph)- increased activity
(neomorph) - new function of gene
- d) suppressors- compensate for other mutations
- e) enhancer- enhances phenotype of a mutation

Fig. 7.25

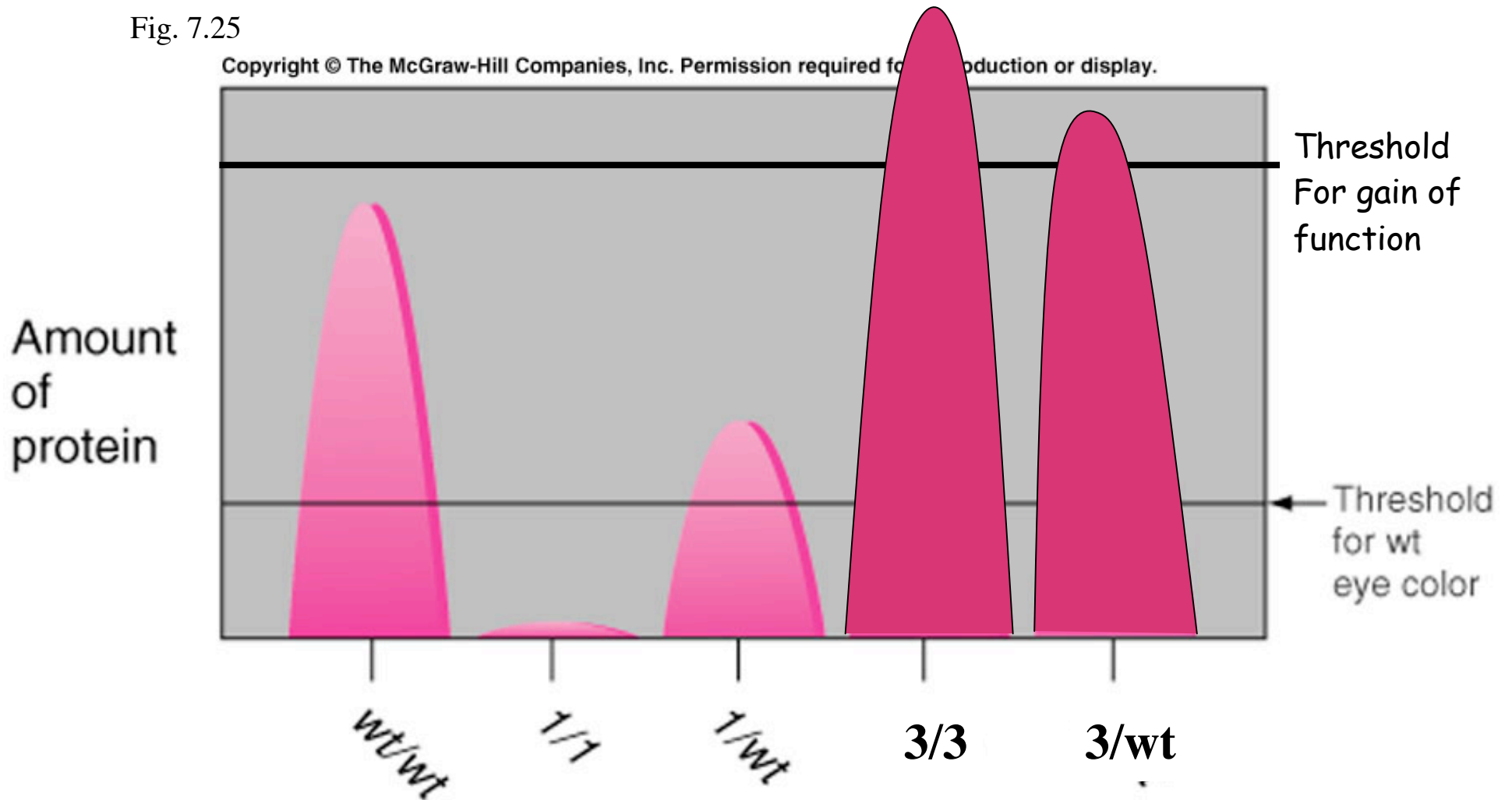
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1: null mutation; 2: hypomorphic mutation
Both 1 and 2 are recessive

Fig. 7.25

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3: hypermorphic mutation
3 is dominant

The underlying nature of recessive or dominant mutations

Recessive

hypomorph: reduced level or a protein with a weak function

Null: complete loss of function

Dominant

hypermorph: increased level or more effective activity

neomorphic: new function

dominant-negative: poisonous effect

haploid-insufficient

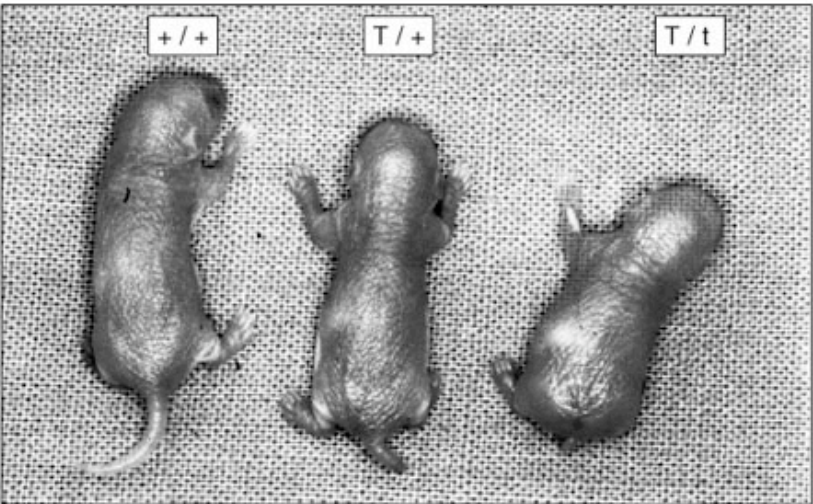
Ectopic expression

Semi-dominant

Fig. 7.27

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(a) Haploinsufficiency



(c) *Kinky*: A dominant negative mutation



(b) Dominant negative mutations

Functional enzyme	Nonfunctional enzyme			
 $d^+d^+d^+d^+$	 $d^+d^+d^+D$	 $d^+d^+Dd^+$	 $d^+Dd^+d^+$	 $Dd^+d^+d^+$
	 d^+d^+DD	 d^+DDd^+	 d^+Dd^+D	 Dd^+Dd^+
	 Dd^+d^+D	 DDd^+d^+	 d^+DDD	 Dd^+DD
	 DDd^+D	 $DDDDd^+$	 $DDDD$	

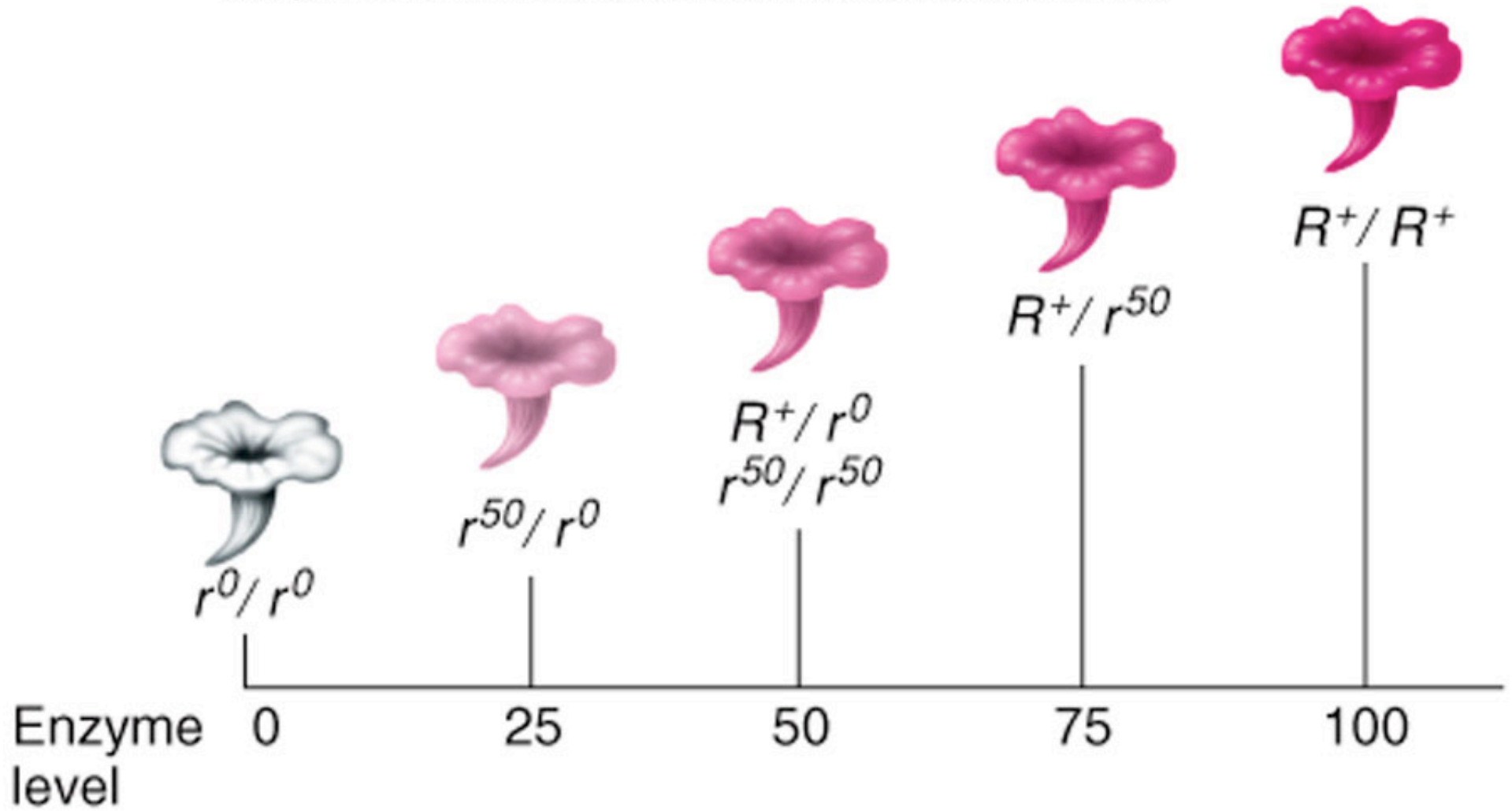
D = dominant mutant subunit
 d^+ = wild-type subunit

(d) A result of ectopic expression



Fig. 7.26

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r^0 : null; r^{50} : hypomorph; R^+ : wild type

Determine recessive or dominant nature of the mutation



C_1C_1 (Colorless mutant) \otimes $+/+$ (Red: WT)
 \downarrow
 $C_1/+$ (red: recessive)
 (colorless: dominant)
 \downarrow \otimes (self)

		<u>Recessive</u>	<u>dominant</u>
1/4	c_1/c_1	colorless	colorless
2/4	$c_1/+$	red	colorless
1/4	$+ /+$	red	red

Complementation tests among *Drosophila* eye mutants

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Fig. 7.14

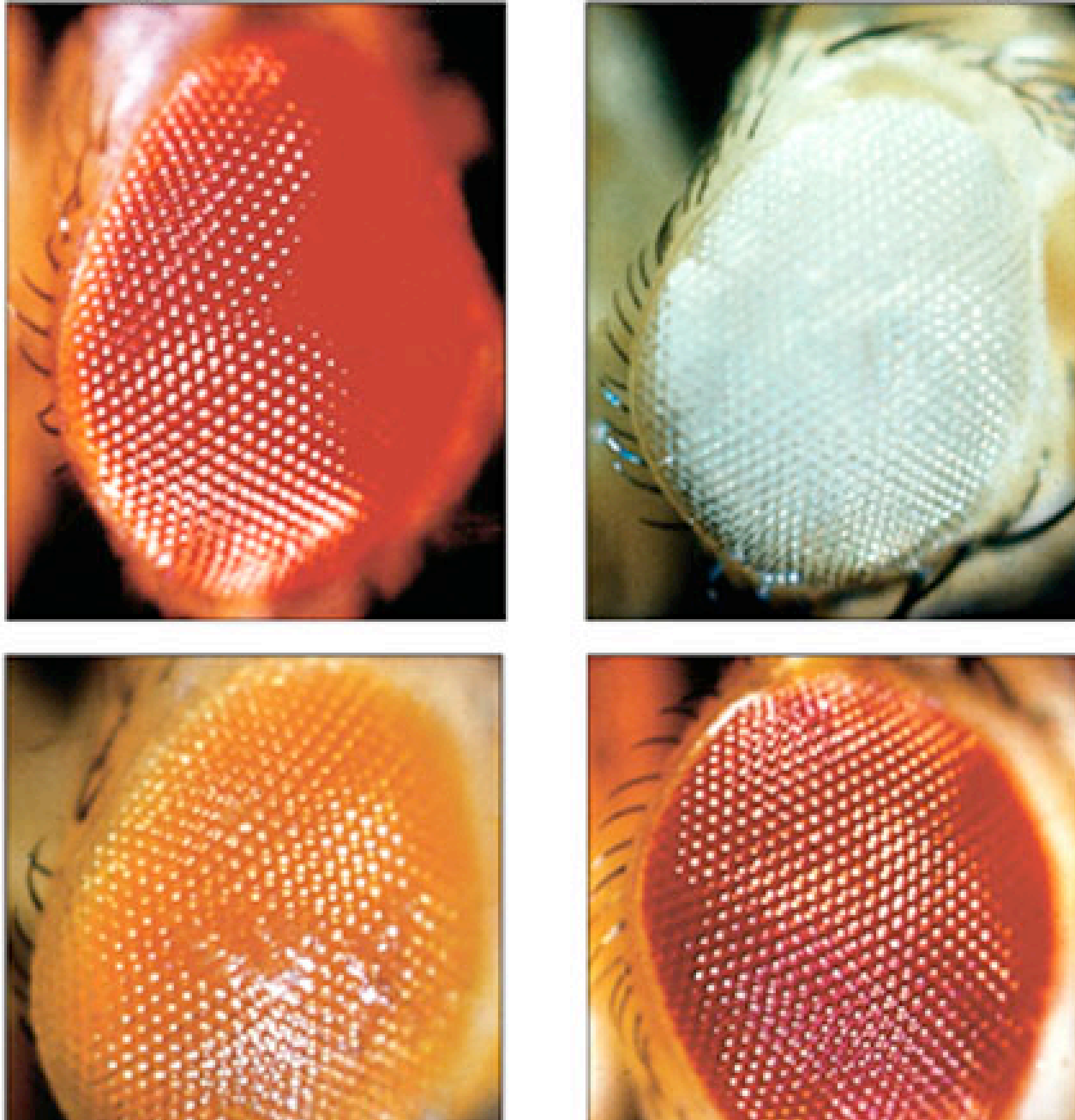
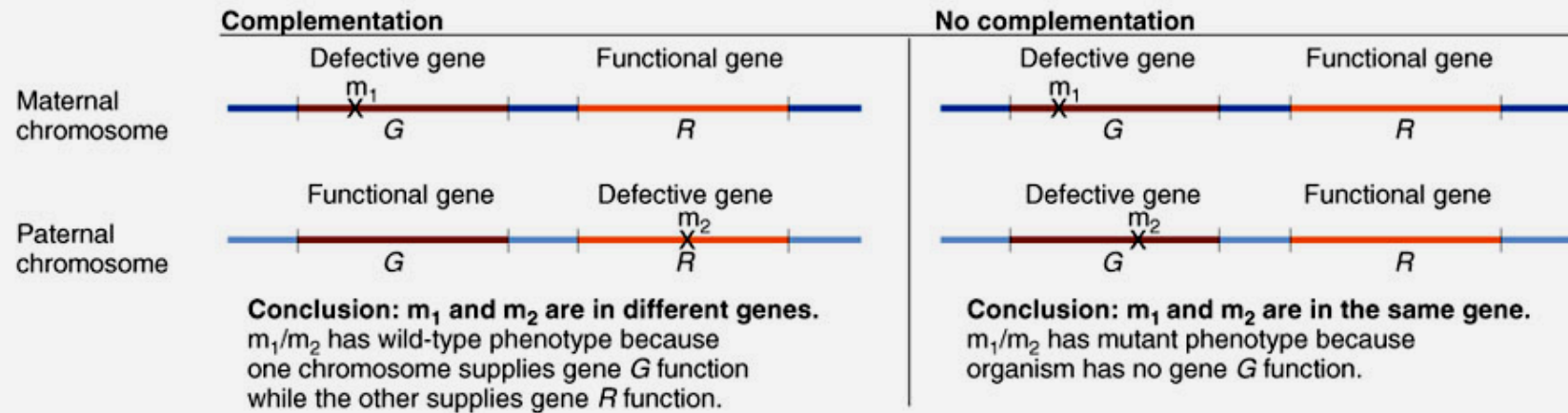


Fig. 7.15

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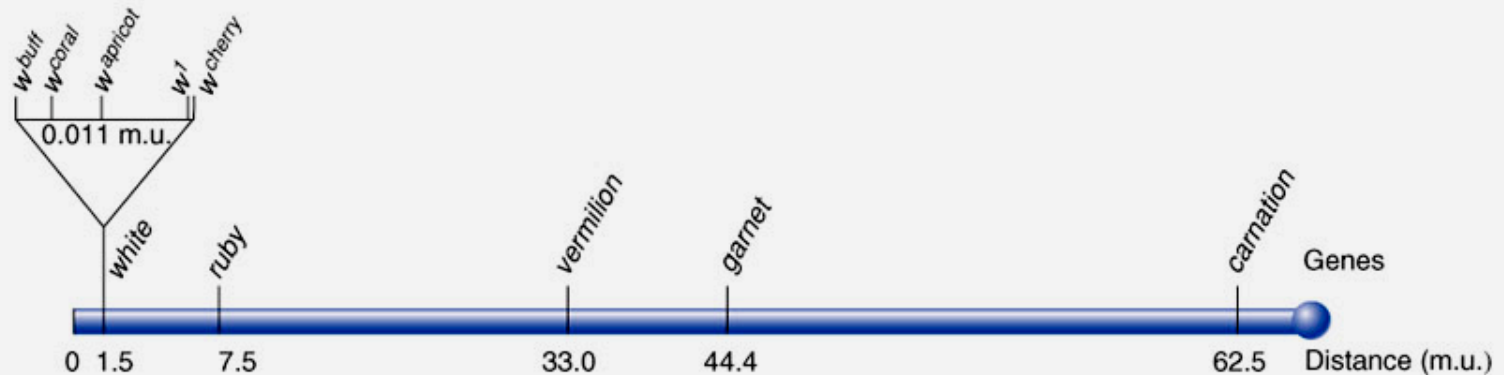
(a) Complementation testing



(b) A complementation table: X-linked eye color mutations in *Drosophila*

Mutation	white	garnet	ruby	vermillion	cherry	coral	apricot	buff	carnation
white	—	+	+	+	—	—	—	—	+
garnet		—	+	+	+	+	+	+	+
ruby			—	+	+	+	+	+	+
vermillion				—	+	+	+	+	+
cherry					—	—	—	—	+
coral						—	—	—	+
apricot							—	—	+
buff								—	+
carnation									—

(c) Genetic map: X-linked eye color mutations in *Drosophila*



Determine allelism by complementation tests

Pairwise crosses between homozygotes and examine F1 for phenotype
only applicable for recessive mutations

male	c_1c_1	c_2c_2	c_3c_3	c_4c_4	c_5c_5	c_6c_6	<u>female</u>
	Colorless	red	red	colorless	red	red	C_1C_1
		colorless	colorless	red	red	red	C_2C_2
			colorless	red	red	red	C_3C_3
				colorless	red	red	C_4C_4
					colorless	colorless	C_5C_5
						colorless	C_6C_6



Maize kernel mutants:

C_1 - C_6 : **colorless, recessive**
wt: red

Three complementation groups:

1. C_1, C_4
2. C_2, C_3
3. C_5, C_6

"Complementation group" equals "Gene"

If two mutations failed to complement,
they are alleles of the same gene
they are allelic to each other
they belong to the same complementation group

If two mutations complements each other,
they are alleles of different genes
they are not allelic to each other
they belong to different complementation groups