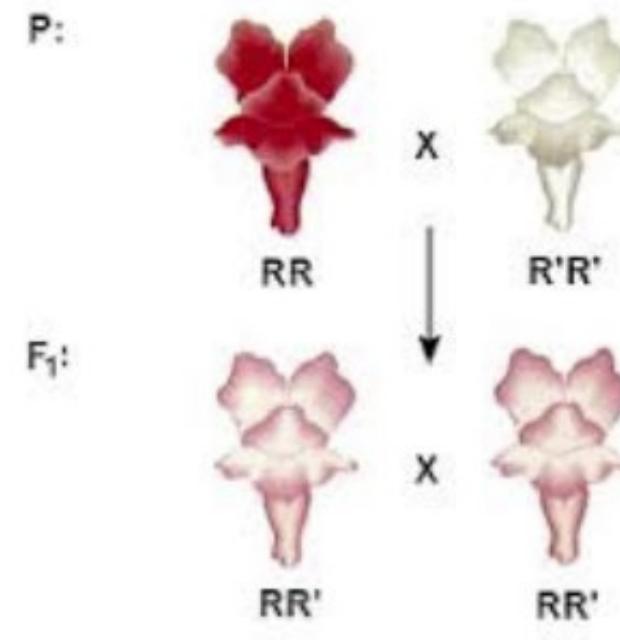
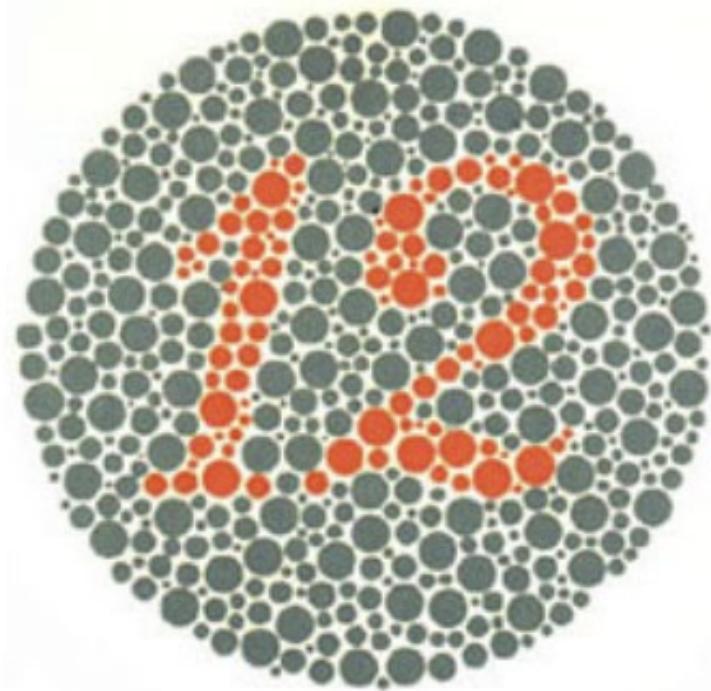


UNIT 3 | EXTENSIONS OF MENDELIAN GENETICS



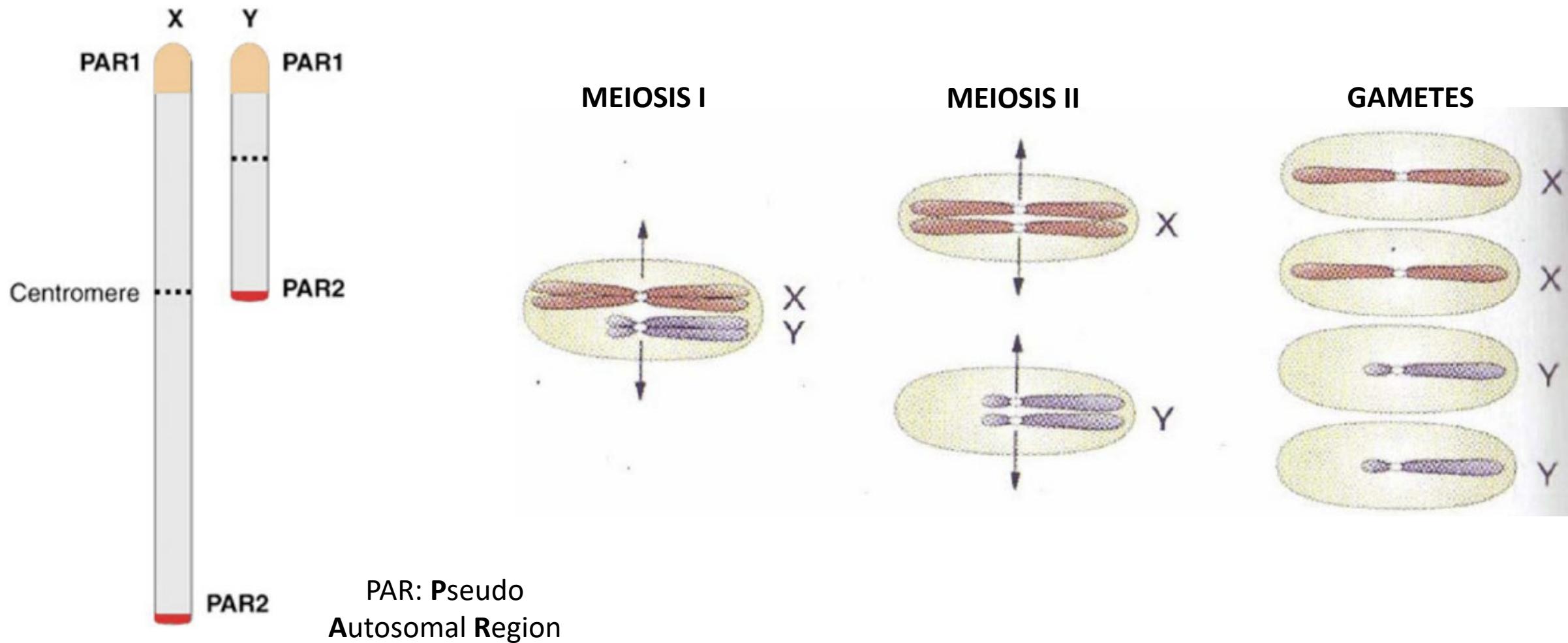
ONE GENE

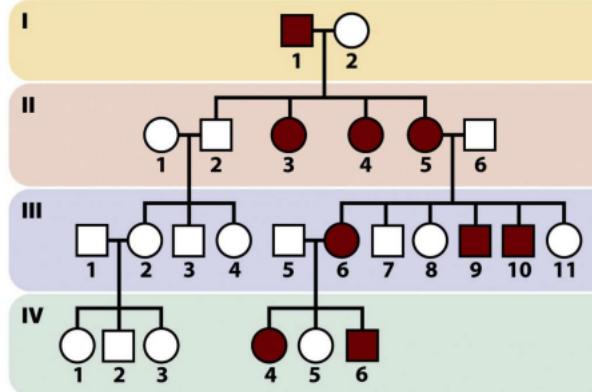
- **Sex-dependant traits:**
 - Sex-linked (chromosome X)
 - Holandric (chromosome Y)
 - Sex-limited
 - Sex-influenced
- **Incomplete dominance**
- **Codominance**
- **Allelic series**
- **Lethal genes**

SEVERAL GENES

- **Gene interaction**
- **Duplicate interaction**
- **Dilution genes**
- **Epistasis**
 - Dominant
 - Recessive
 - Duplicate dominant
 - Duplicate recessive
 - Dominant-recessive

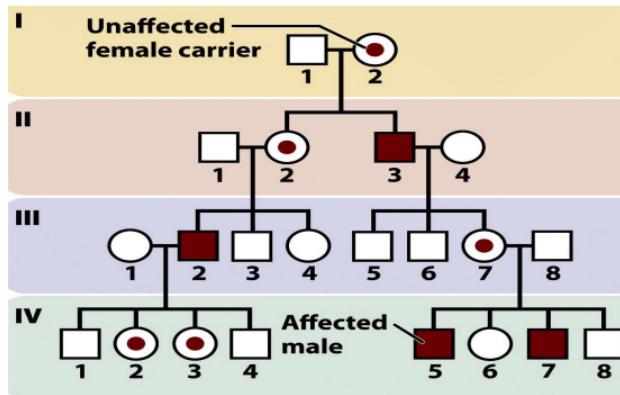
SEX IS INHERITED IN A MENDELIAN WAY





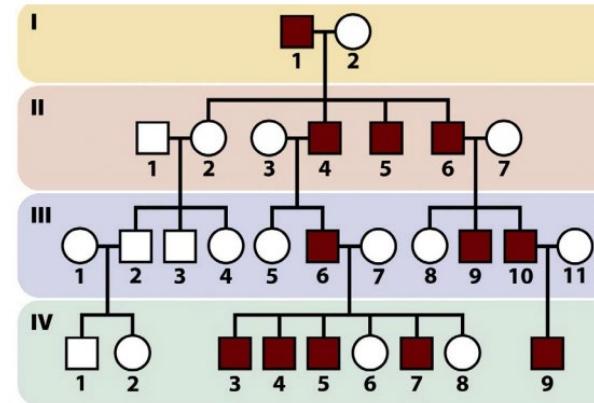
X-LINKED (DOMINANT)

- More females are affected
- Do not skip generations
- Affected males have an affected mother
- Affected females have either an affected mother or father
- Males do not pass it to their male offspring



X-LINKED (RECESSIVE)

- More males are affected
- Tend to skip generations
- Males do not pass it to their male offspring
- Affected males have carrier female offspring

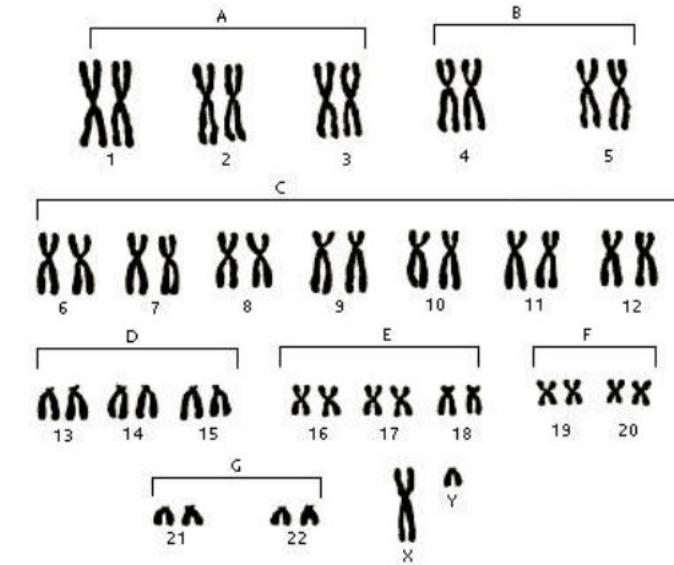
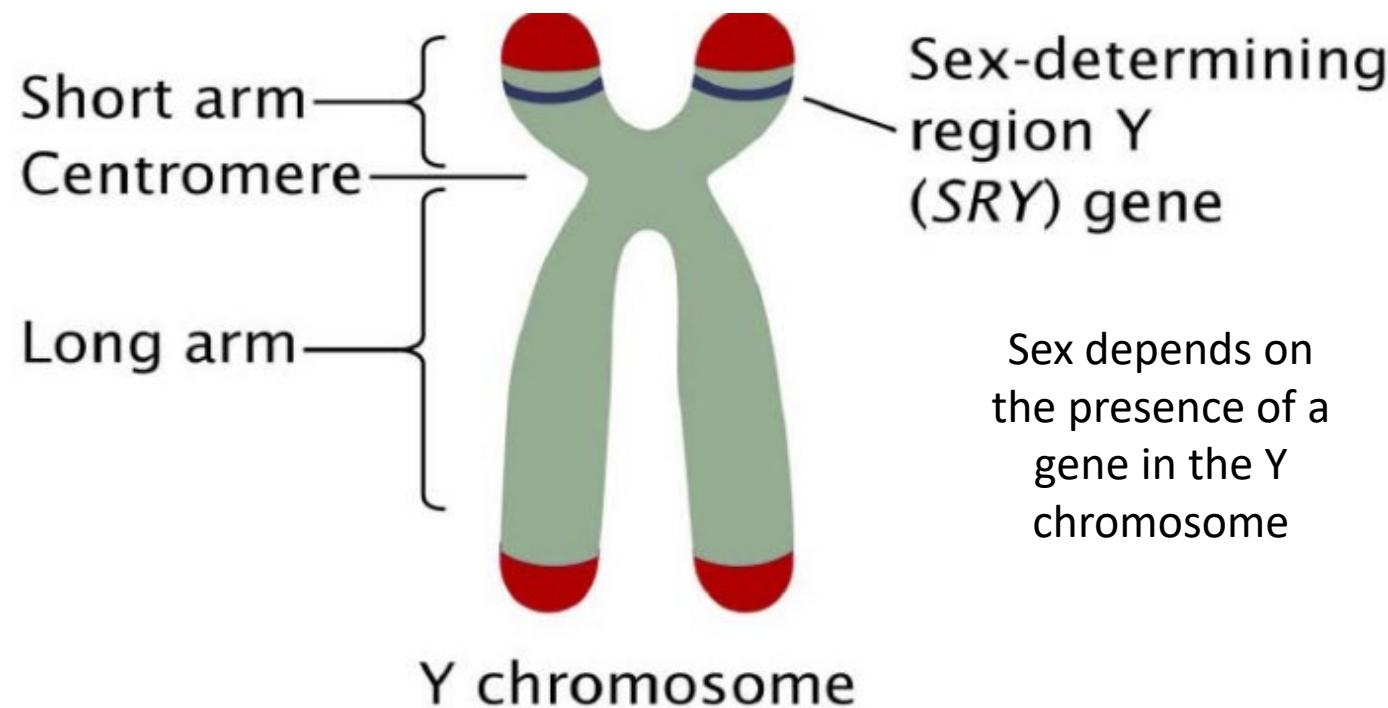


HOLANDRIC

- Only males are affected
- Males pass it to their male offspring
- Do not skip generations

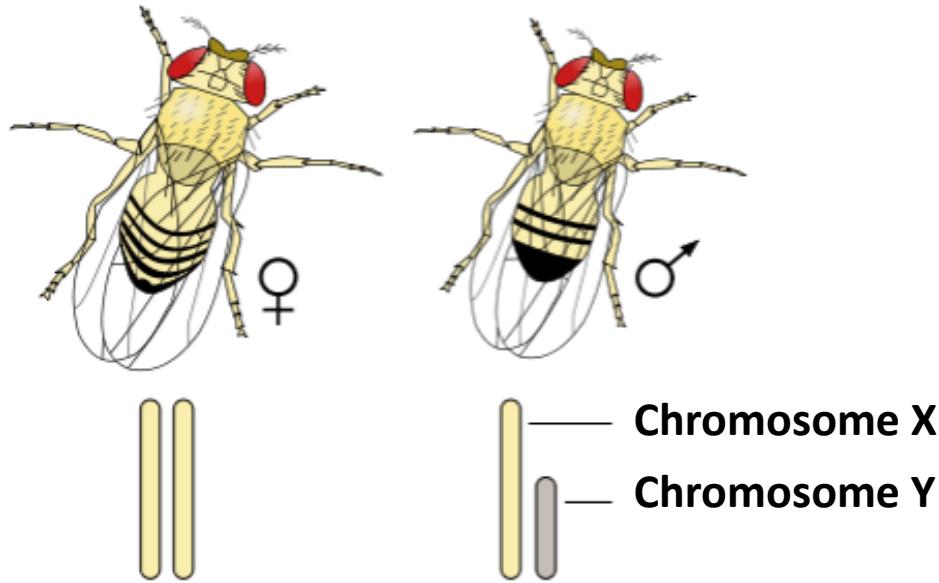
SEX-DETERMINATION SYSTEMS

(active Y)



SEX-DETERMINATION SYSTEMS

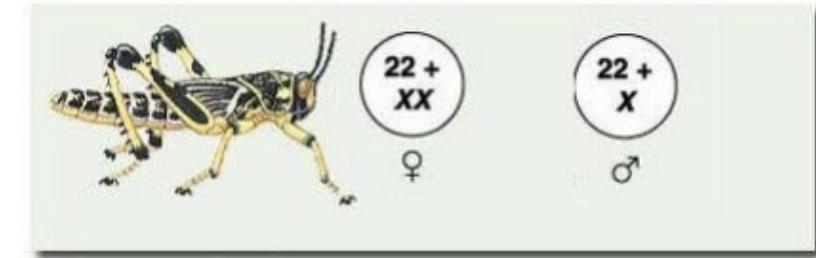
(X:A ratio)



Sex depends on the ration between number of X chromosomes and set of autosomes:

XX: 2/2 → female

XY: 1/2 (<1) → male

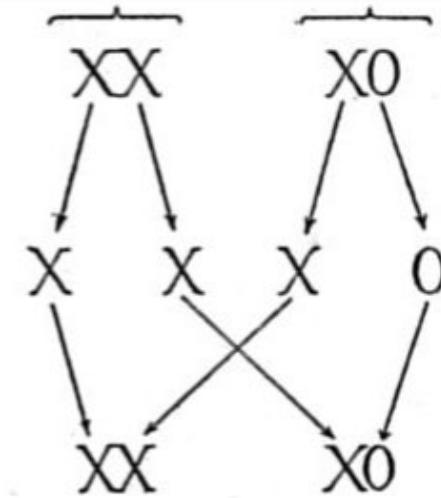


Diploid Nuclei

Gametes

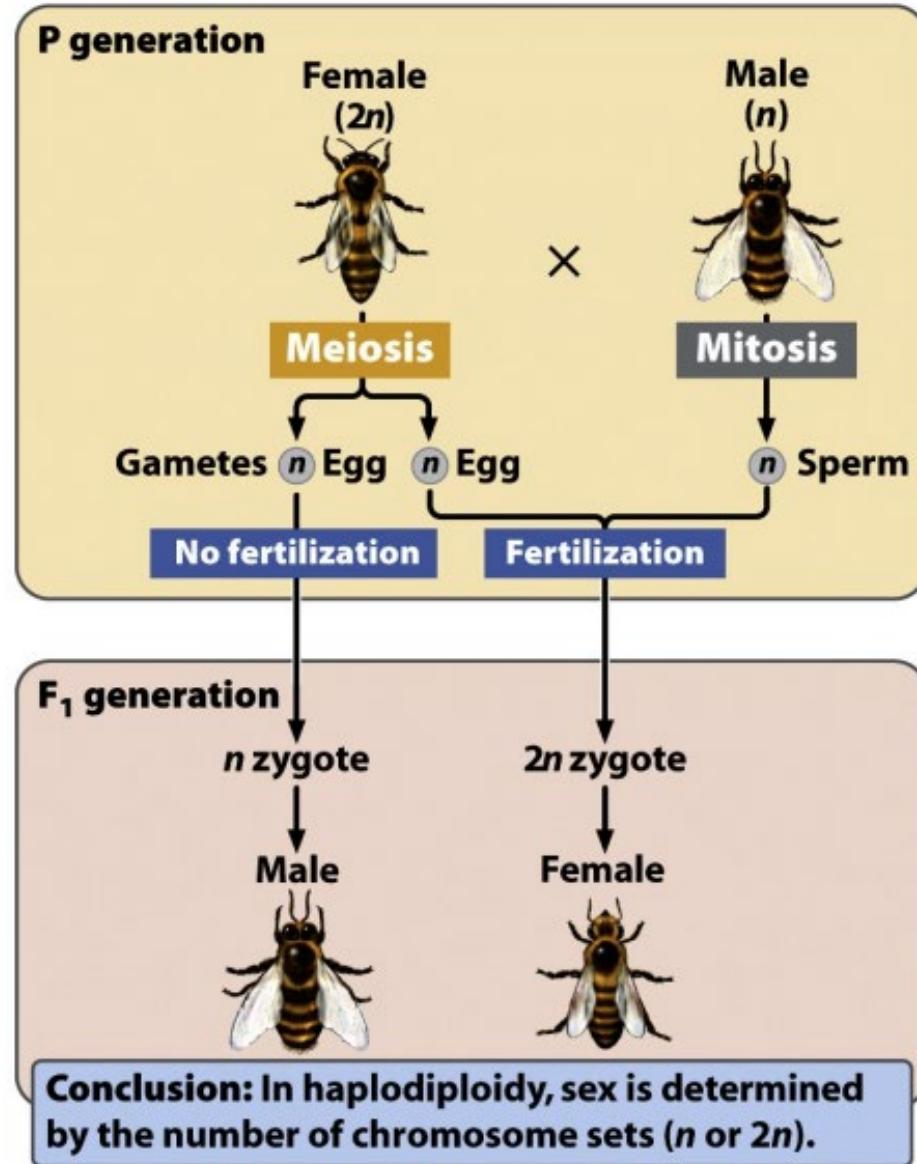
Fertilization

Zygotes

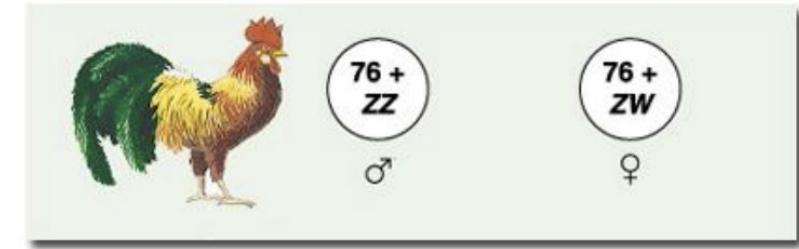
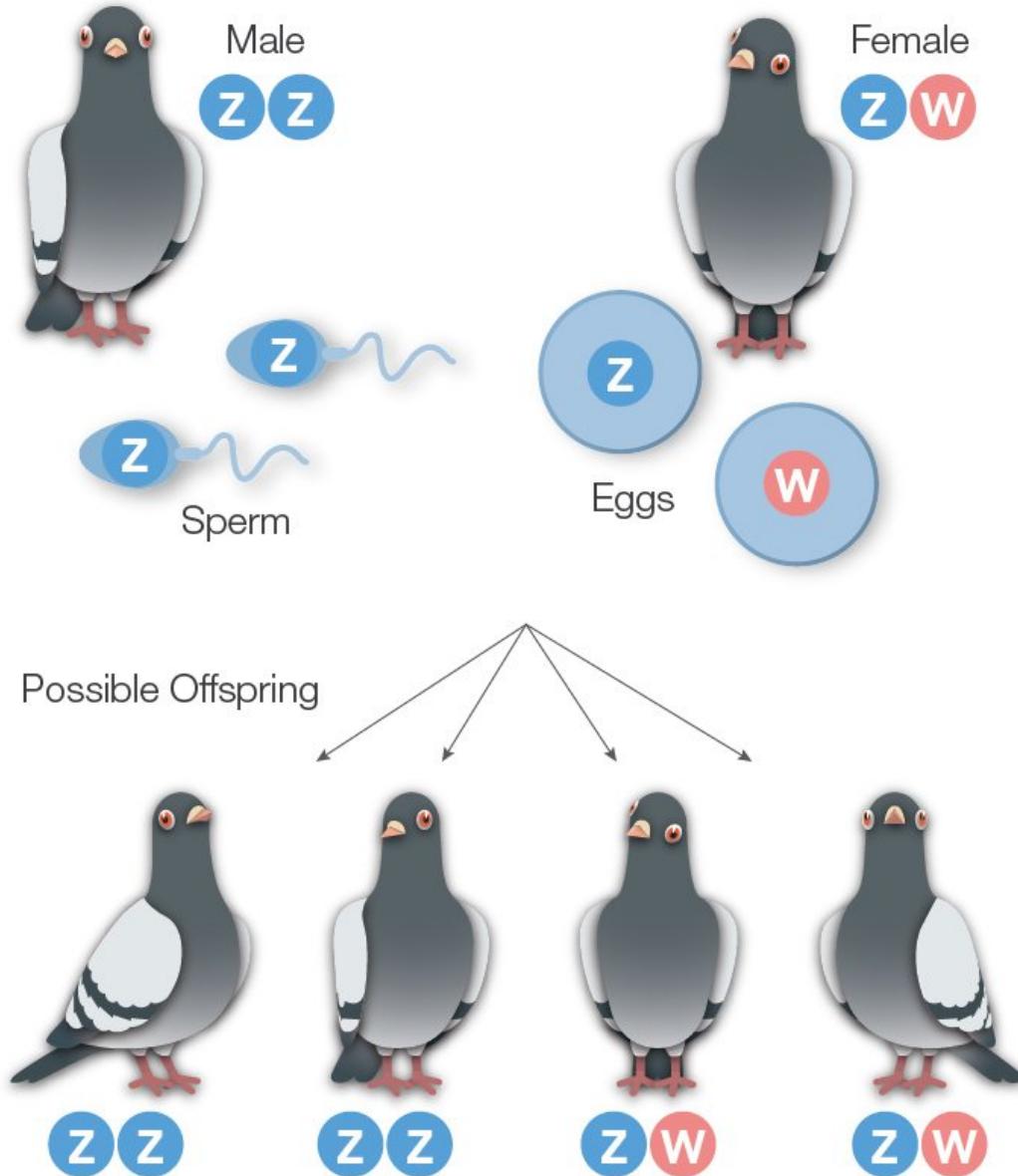


SEX-DETERMINATION SYSTEMS

(Haplodiploidy)

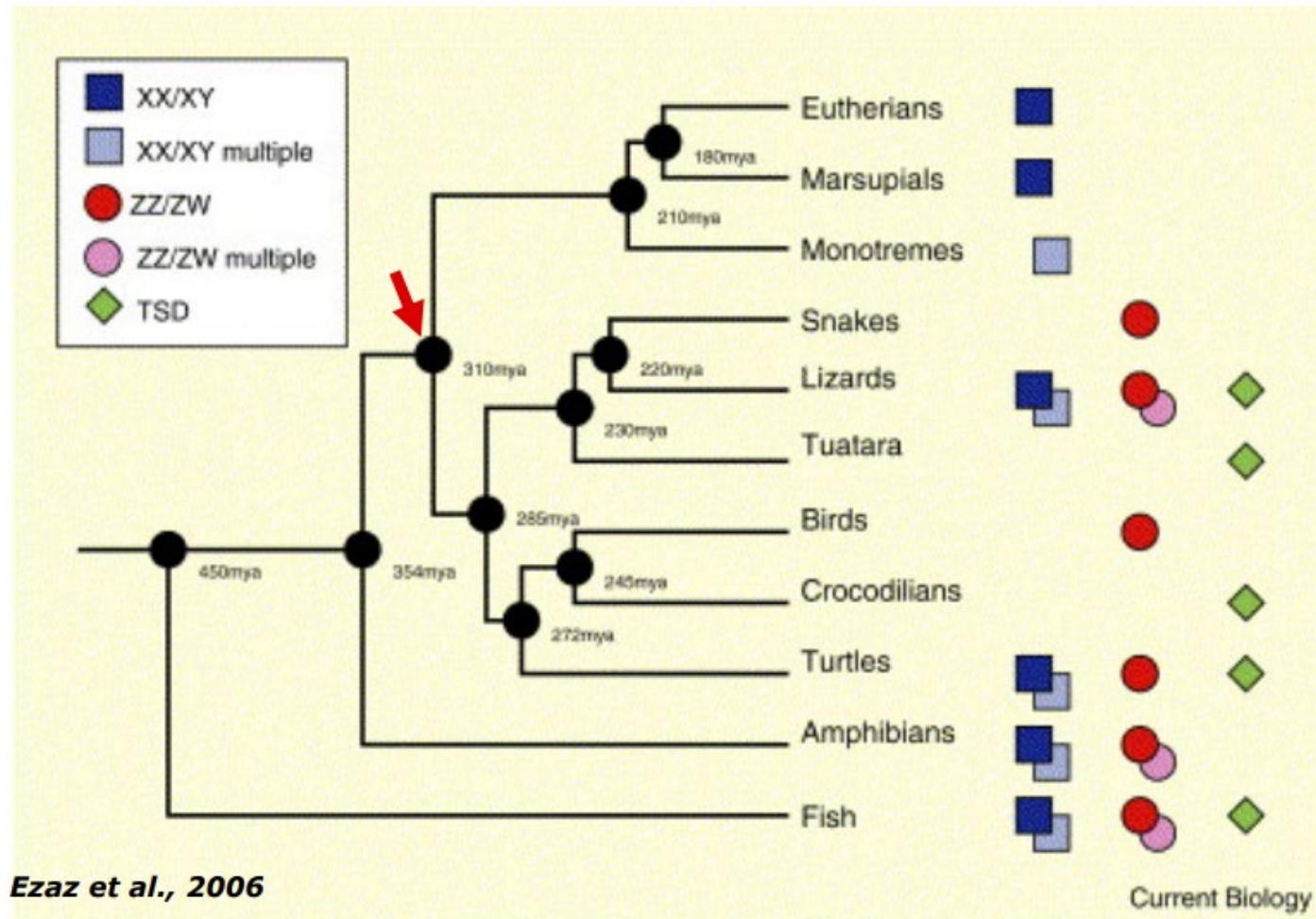


SEX-DETERMINATION SYSTEMS

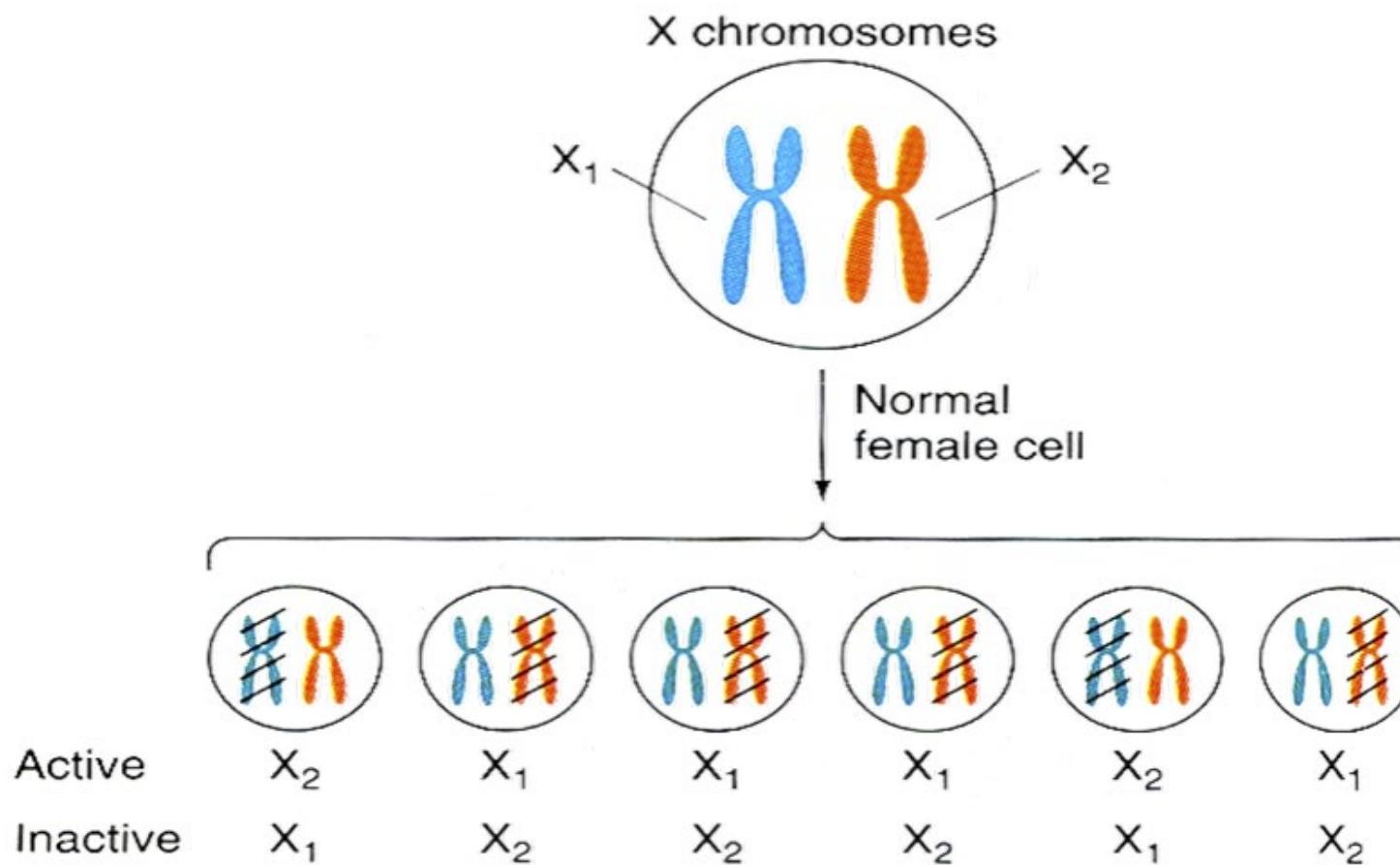


Homogametic males |
Heterogametic females

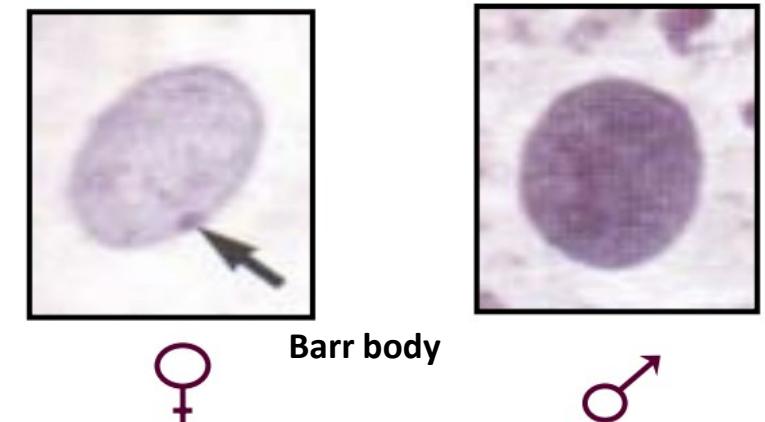
SEX-DETERMINATION SYSTEMS



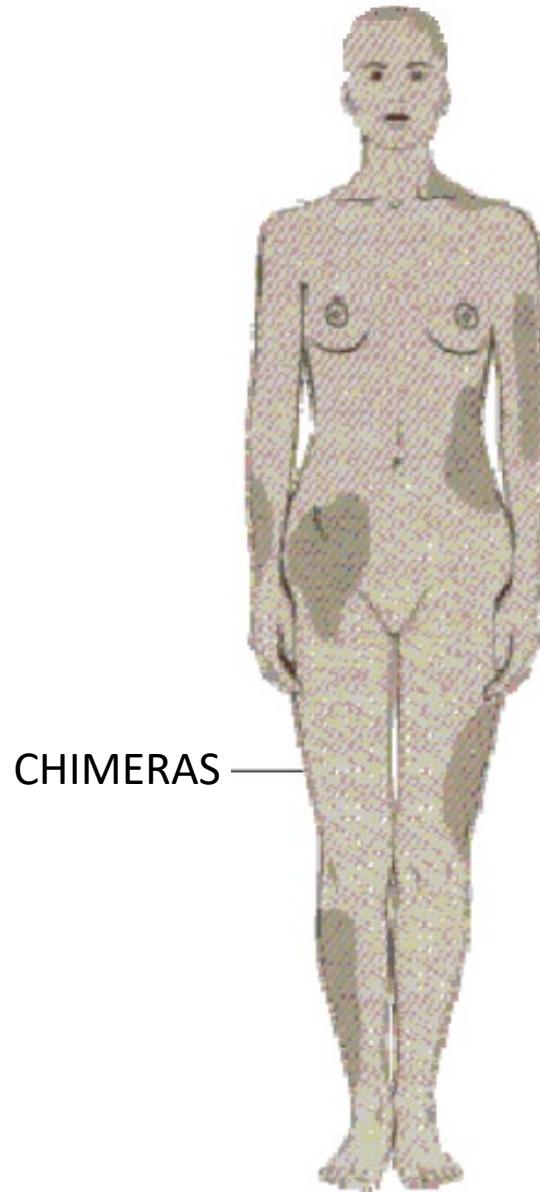
X INACTIVATION



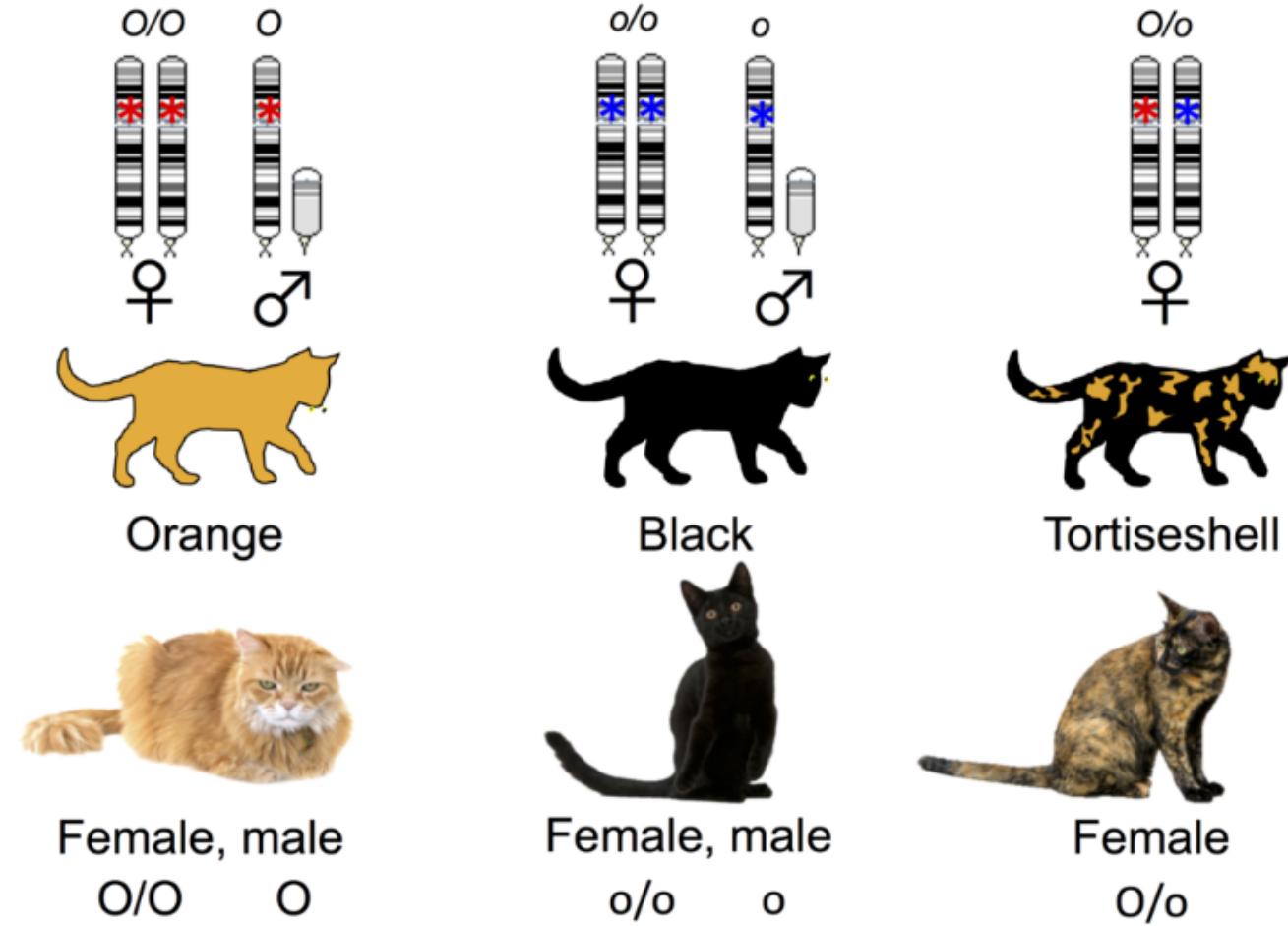
Inactivation occurs at random, so that X_1 is active in approximately half the cells



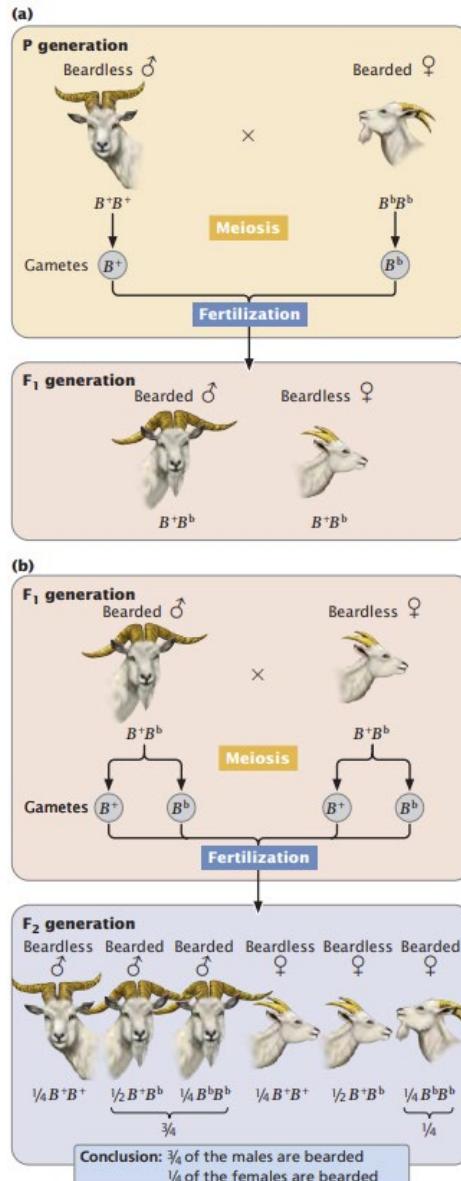
X INACTIVATION



Locus O (Orange): On Chromosome X

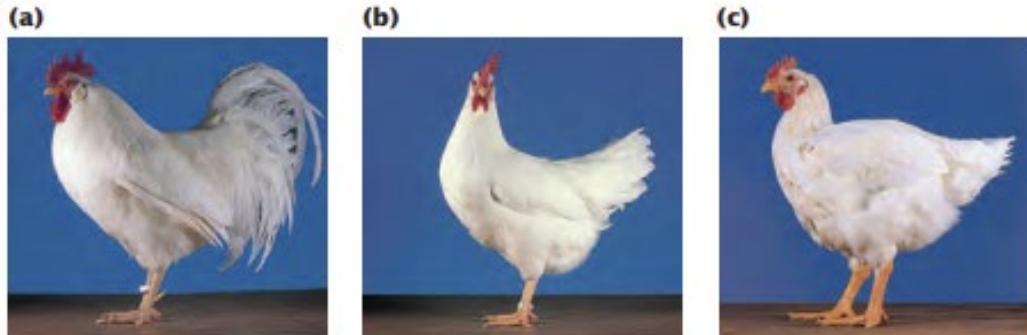


SEX-INFLUENCED AND SEX-LIMITED TRAITS



5.12 Genes that encode sex-influenced traits are inherited according to Mendel's principles but are expressed differently in males and females.

Genotype	Fenotype	
	Males	Females
$a'a'$	Bald	Bald
$a'a$	Bald	Hair
aa	Hair	Hair



5.13 A sex-limited characteristic is encoded by autosomal genes that are **expressed in only one sex**. An example is cock feathering in chickens, an autosomal recessive trait that is limited to males.
(a) Cock-feathered male. (b) Hen-feathered female. (c) Hen-feathered male.
[Larry Lefever/Grant Heilman Photography.]

INCOMPLETE DOMINANCE

$R^1 R^1$ red \times $R^2 R^2$ white

P₁



$R^1 R^2$ pink

F₁

Hybrids exhibit an intermediate phenotype

$R^1 R^2 \times R^1 R^2$

F₁ × F₁



1/4 $R^1 R^1$ red

1/2 $R^1 R^2$ pink

1/4 $R^2 R^2$ white



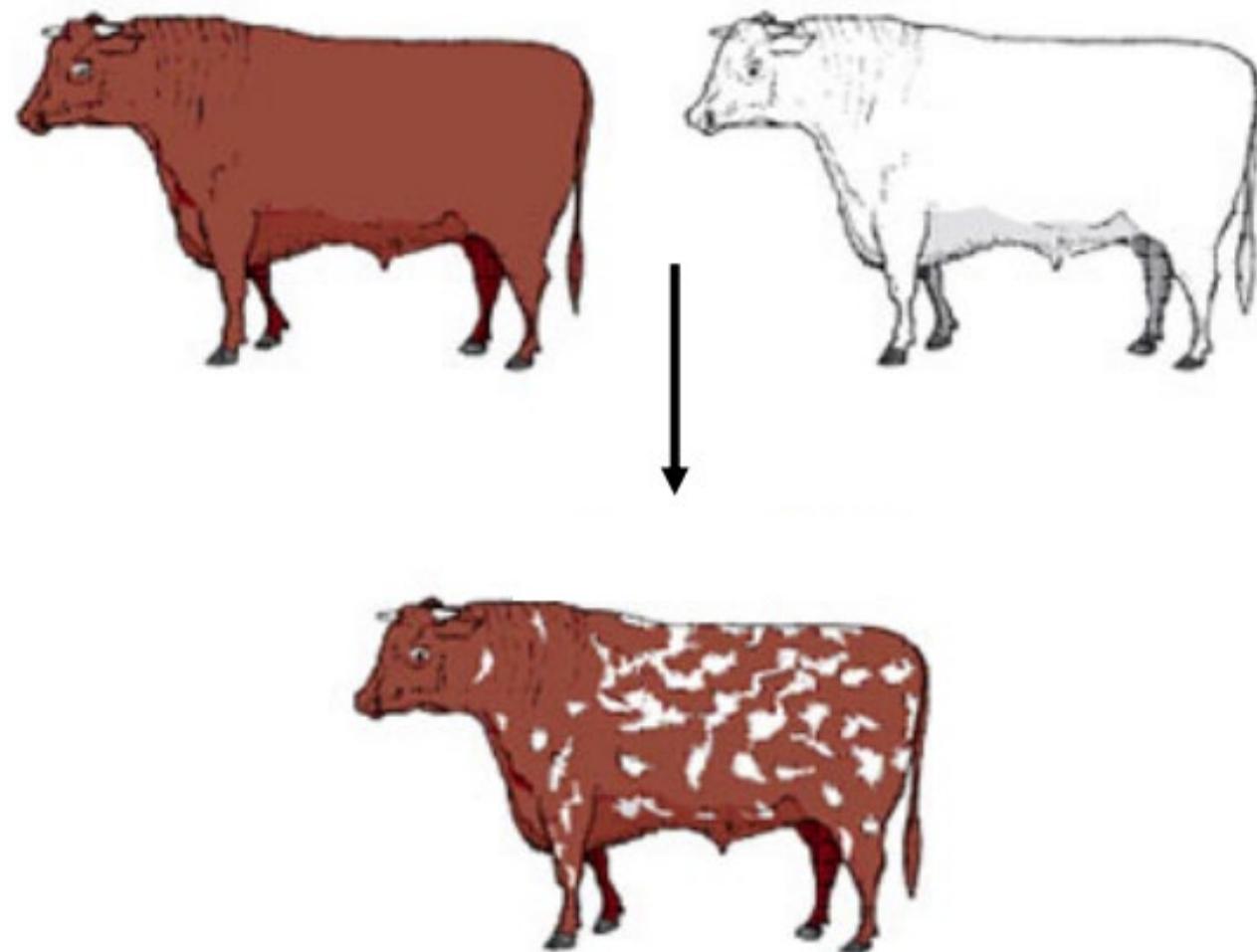
F₂



Phenotypic proportions = Genotypic proportions

CODOMINANCE

Hybrids exhibit dominant and recessive phenotypes



ALLELIC SERIES

Locus C (Albino, Colorpoint)

$C > c^b > c^b c^s > c^s > c/c$



Pigmented

Burmese

Tonkinese

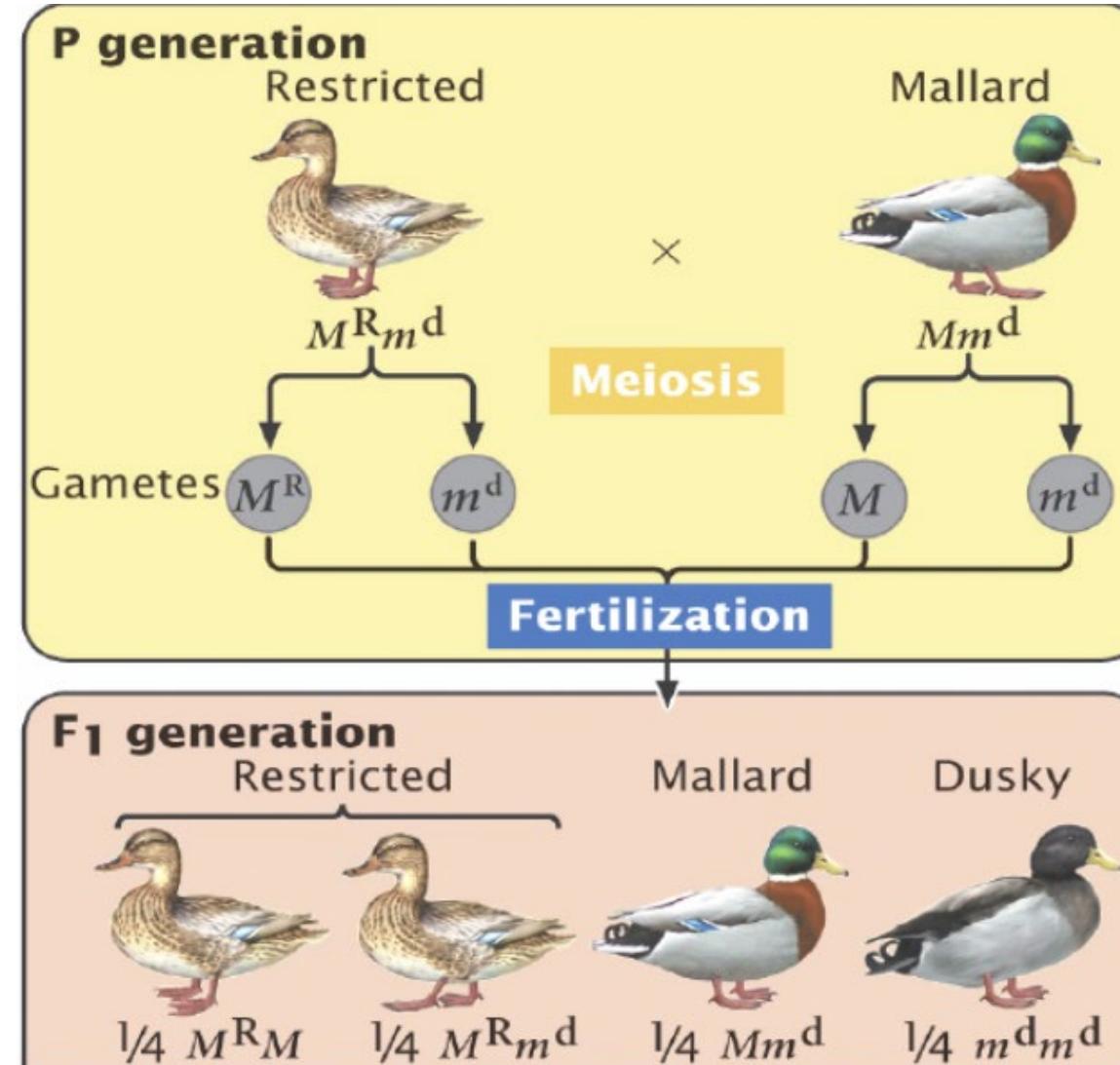
Siamese

Albino

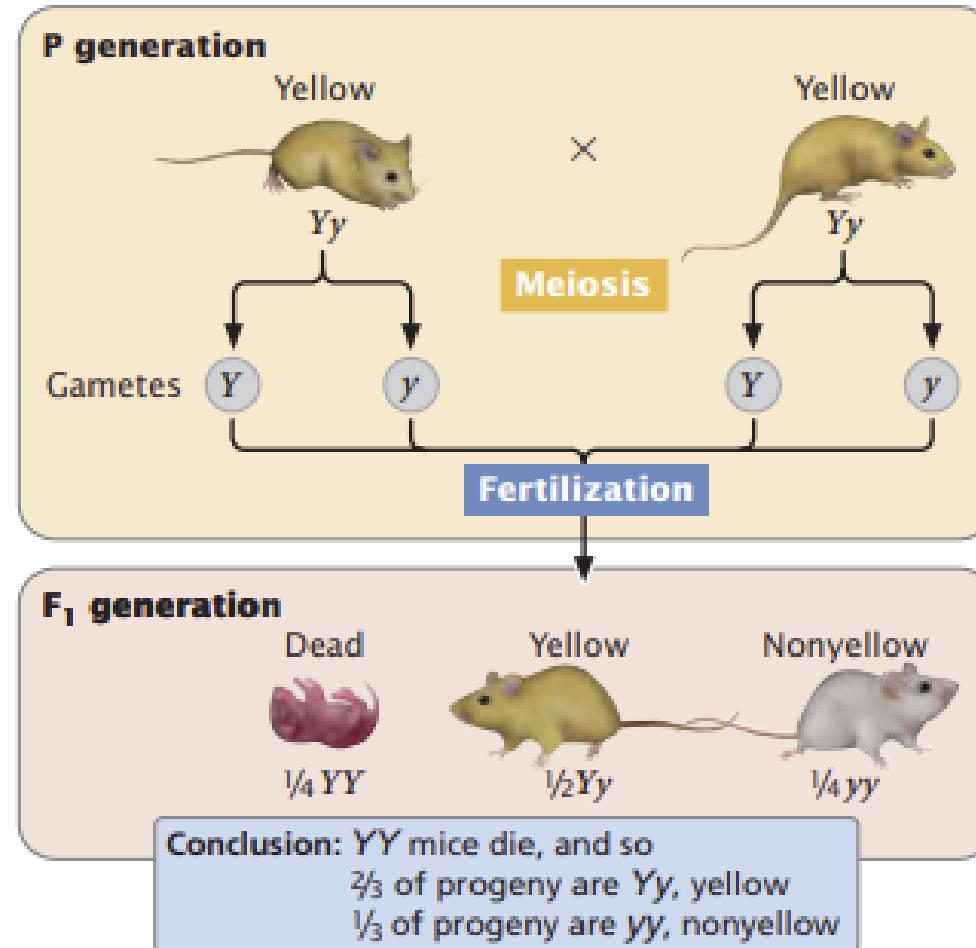
“Colorpoint”



ALLELIC SERIES



LETHAL GENES



5.1 The 2 : 1 ratio produced by a cross between two yellow mice results from a lethal allele.

GENE INTERACTION

(separate metabolic pathways)

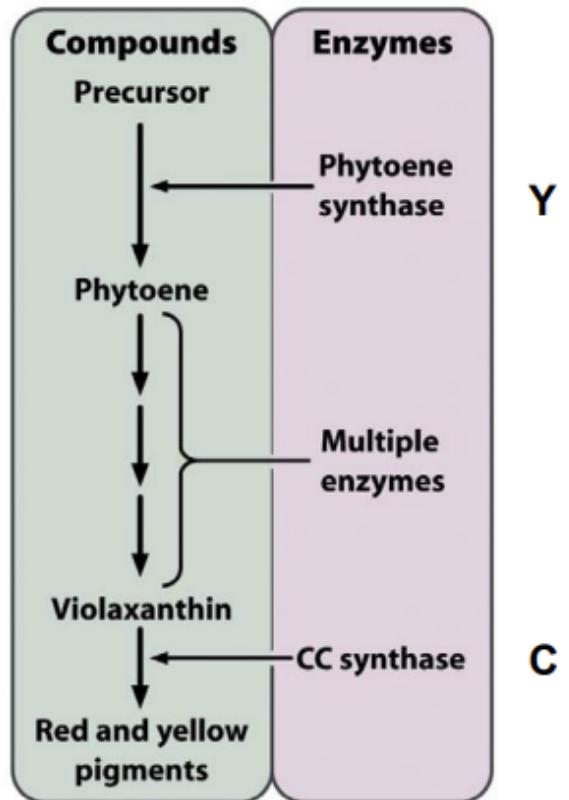
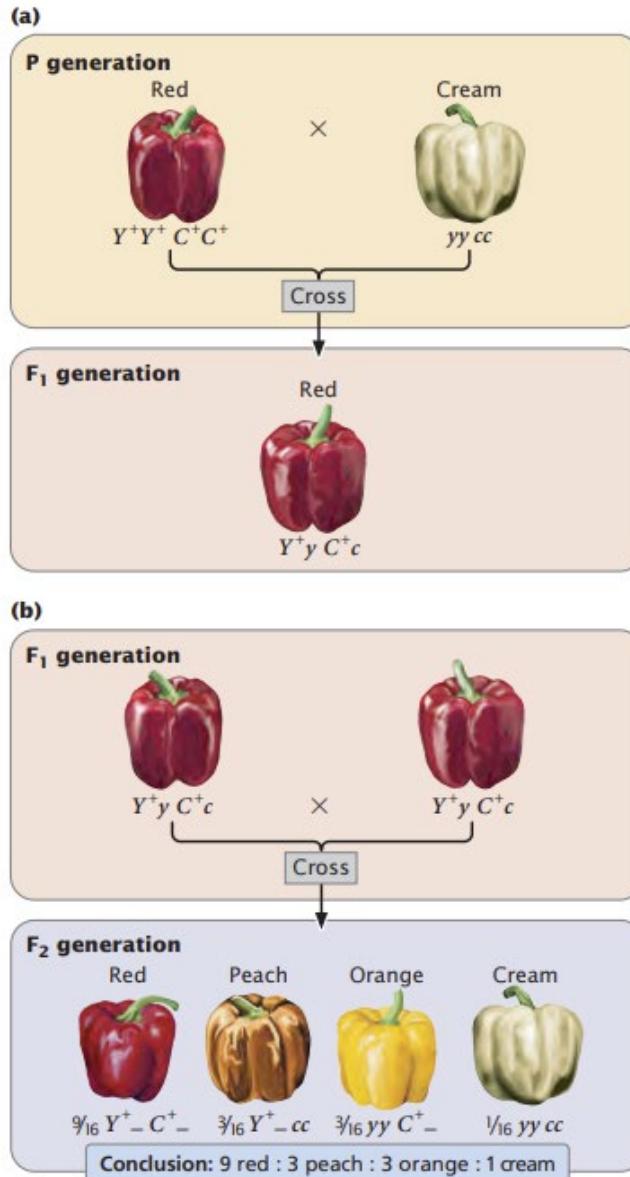


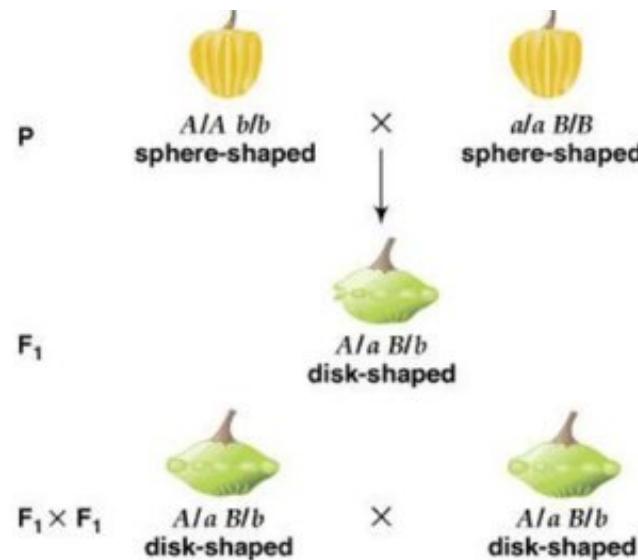
Figure 5-7
Genetics: A Conceptual Approach, Third Edition
© 2009 W.H. Freeman and Company



5.7 Gene interaction in which two loci determine a single characteristic, fruit color, in the pepper *Capsicum annuum*.

DUPLICATE INTERACTION

(separate metabolic pathways)



Here the A and B genes
Both control fruit shape.
One big A OR one big B
Gives sphere-shaped.
But the **combination** one
Big A AND one big B
interacts to give disk-
shaped.

F_2 ratio for $A/a \times A/a$	F_2 ratio for $B/b \times B/b$	Combined F_2 ratios	F_2 phenotypic proportions
$\frac{3}{4} A/-$	$\frac{3}{4} B/-$	$\frac{9}{16} A/-\ B/-$	$\frac{9}{16}$ disk-shaped
$\frac{1}{4} a/a$	$\frac{1}{4} b/b$	$\frac{3}{16} A/-\ b/b$ $\frac{3}{16} a/a\ B/-$	$\frac{3}{16}$ sphere-shaped $\frac{3}{16}$ sphere-shaped $\frac{1}{16}$ long-shaped

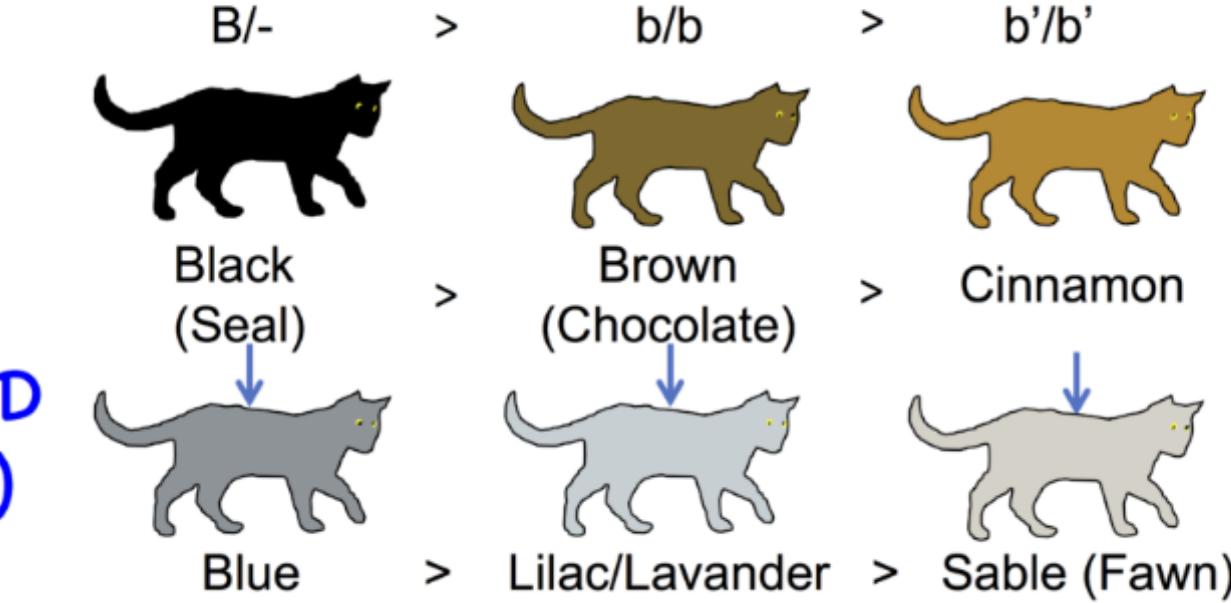
Probability calculations for each row:

- Row 1: $\frac{3}{4} \times \frac{3}{4} = \frac{9}{16}$ disk-shaped
- Row 2: $\frac{1}{4} \times \frac{3}{4} = \frac{3}{16}$ sphere-shaped
- Row 3: $\frac{1}{4} \times \frac{1}{4} = \frac{1}{16}$ long-shaped

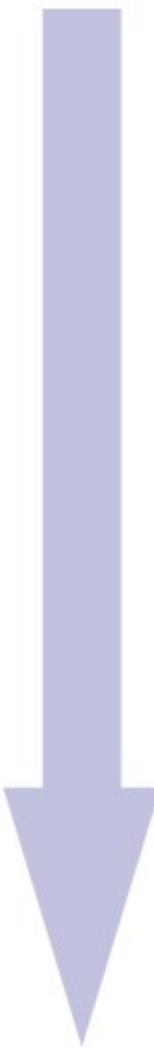
DILUTION GENES

Locus B (Brown)

**Locus D
(Dilute)**
 d/d



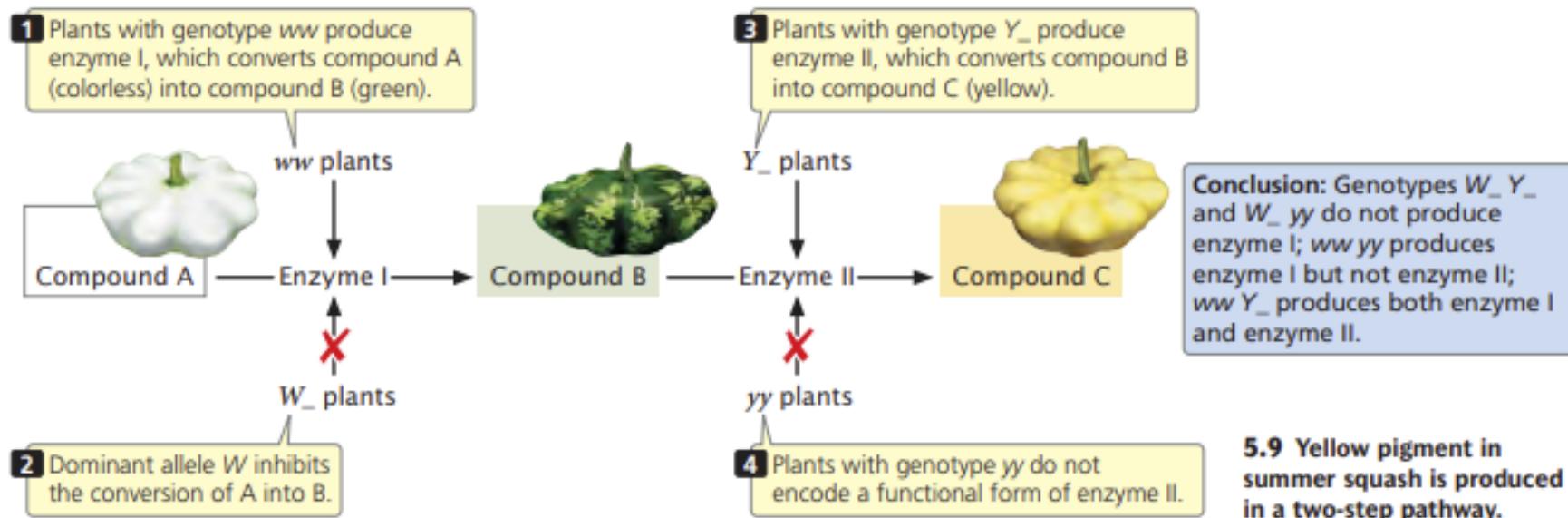
DILUTION GENES



Genotype	Phenotype
Chestnut + D/D	Chestnut
Chestnut + D/d	Palomino
Chestnut + d/d	Cremello (nearly white)
Bay + D/D	Bay
Bay + D/d	Dun or buckskin
Bay+ d/d	Perlino (nearly white)

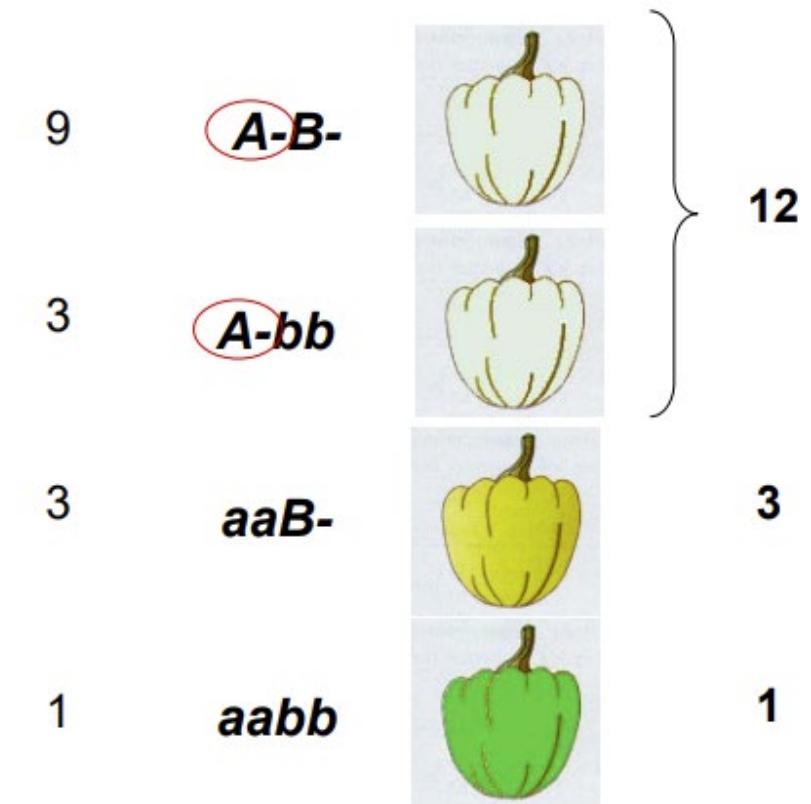
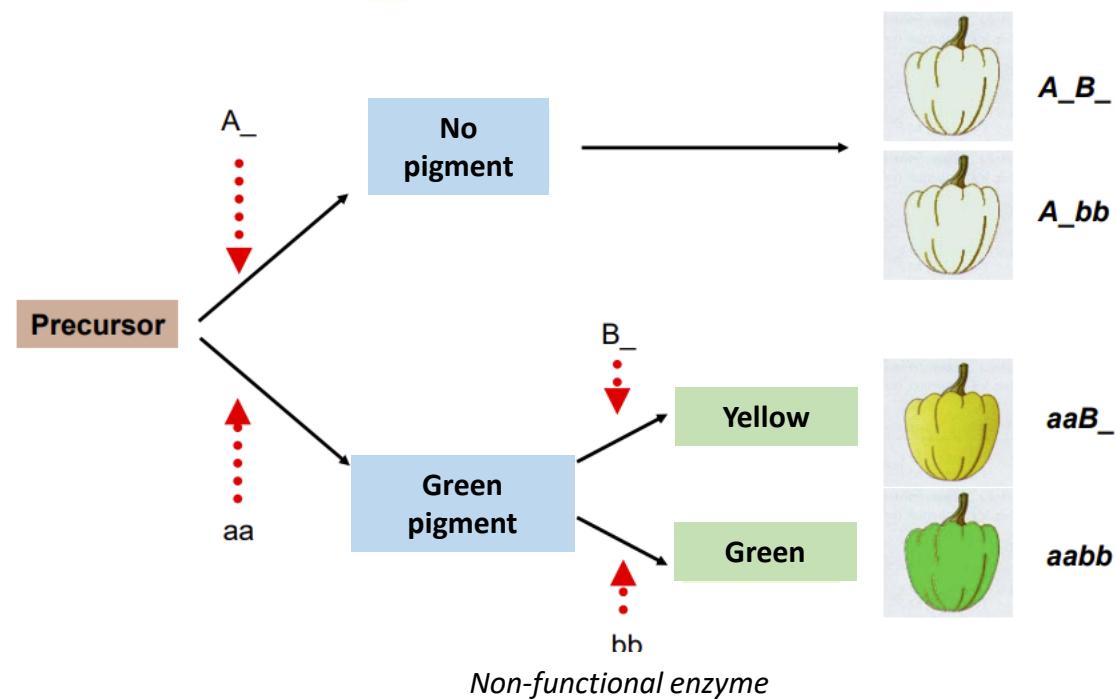
DOMINANT EPISTASIS

(same metabolic pathway)



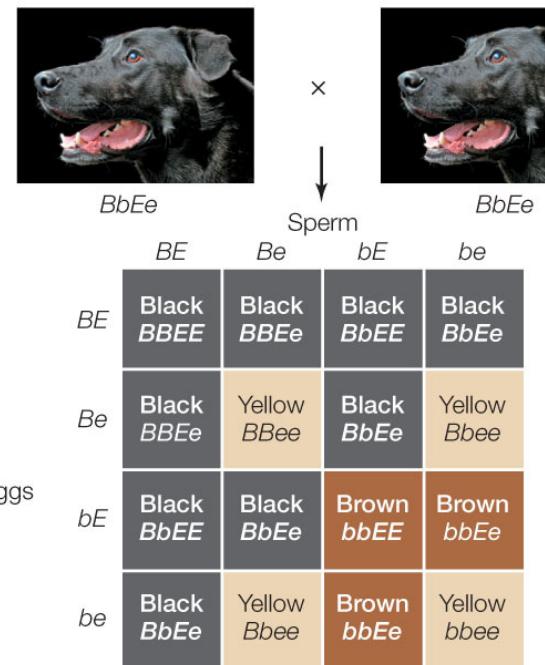
DOMINANT EPISTASIS

(same metabolic pathway)



RECESSIVE EPISTASIS

(same metabolic pathway)



8.12 Black: © Silense/istock. 8.12 Chocolate:
© Erik Lam/istock. 8.12 Yellow: © Gelpi
JM/Shutterstock.

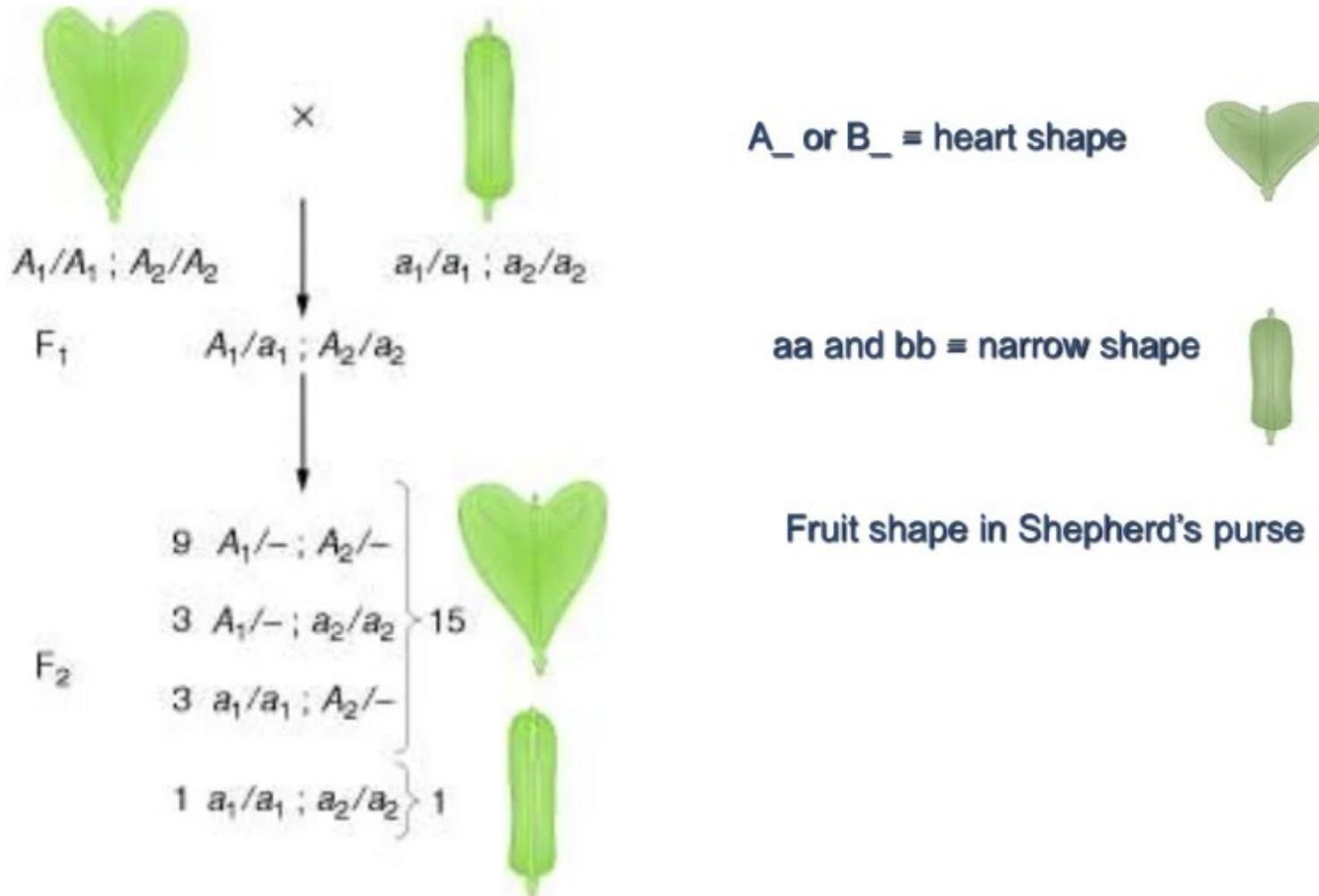
RECESSIVE EPISTASIS

(same metabolic pathway)



DUPLICATE DOMINANT EPISTASIS

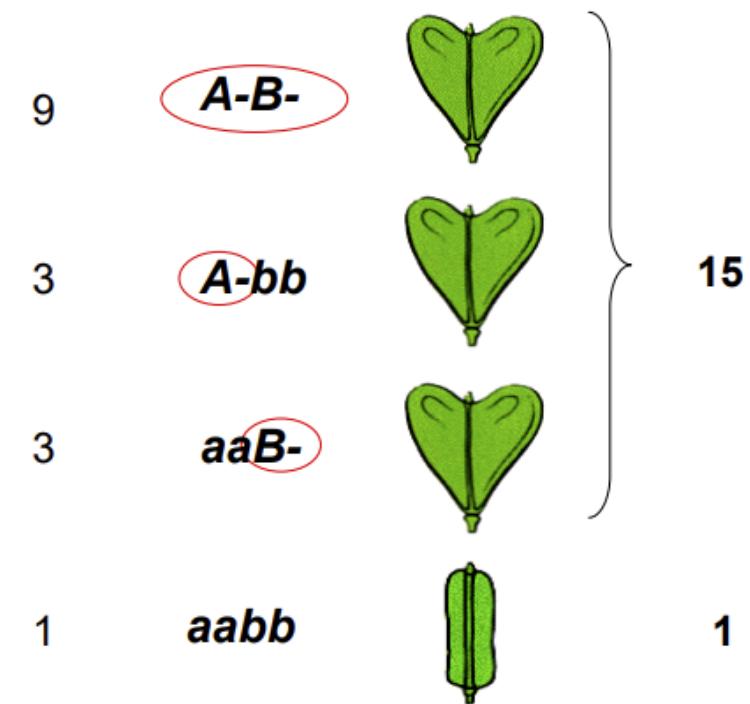
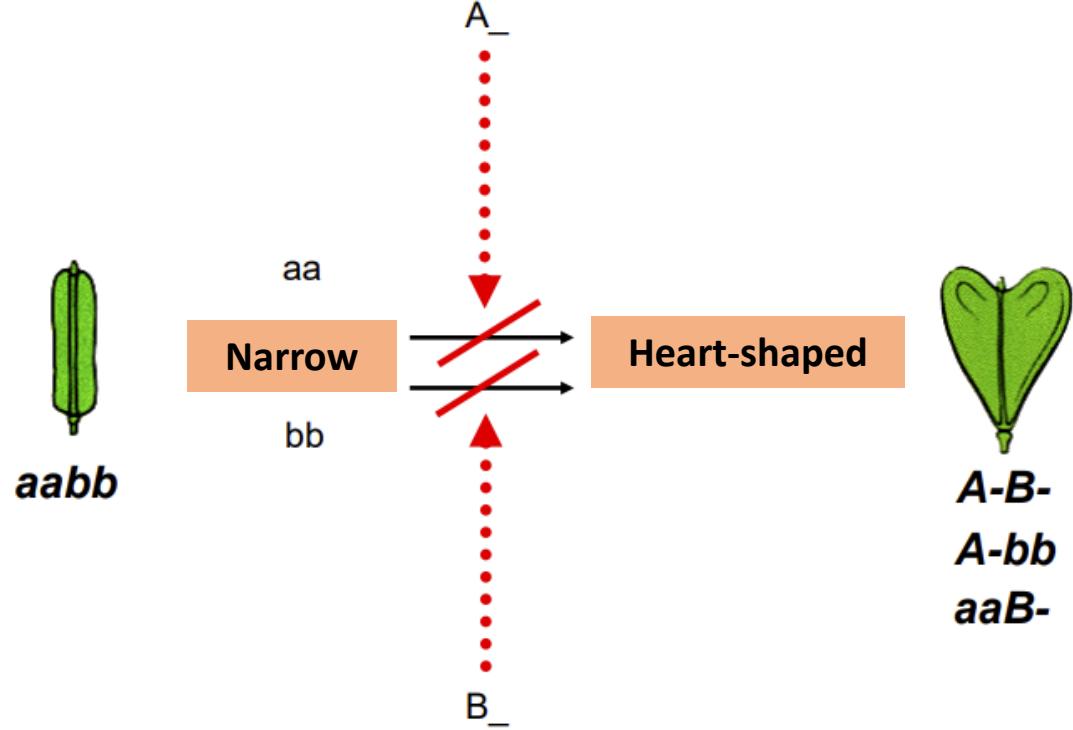
(same metabolic pathway)



DUPLICATE DOMINANT EPISTASIS

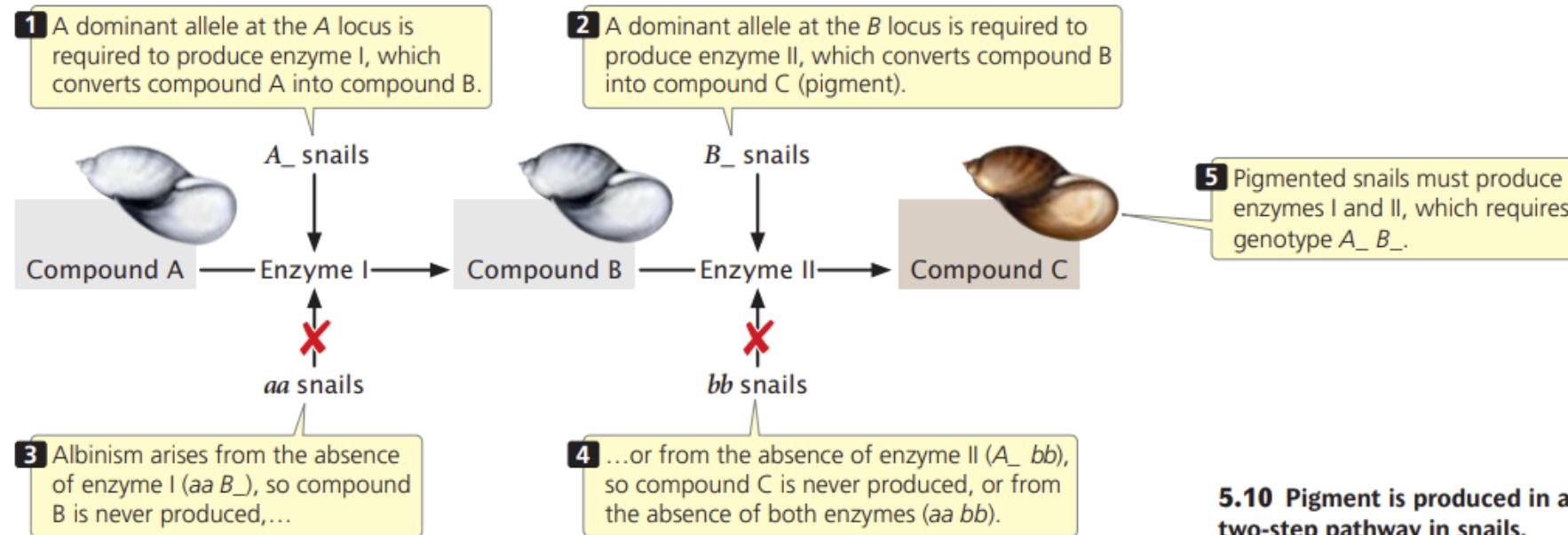
(same metabolic pathway)

In some cases:
duplicate
genes



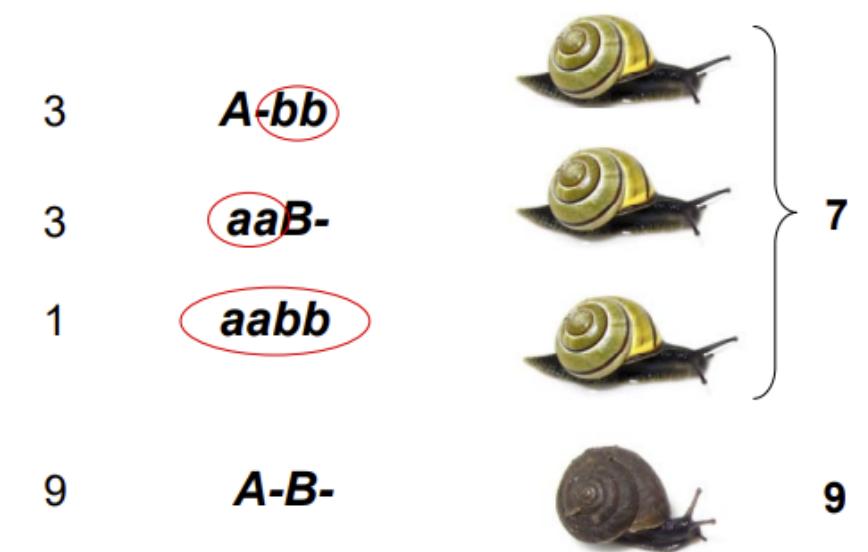
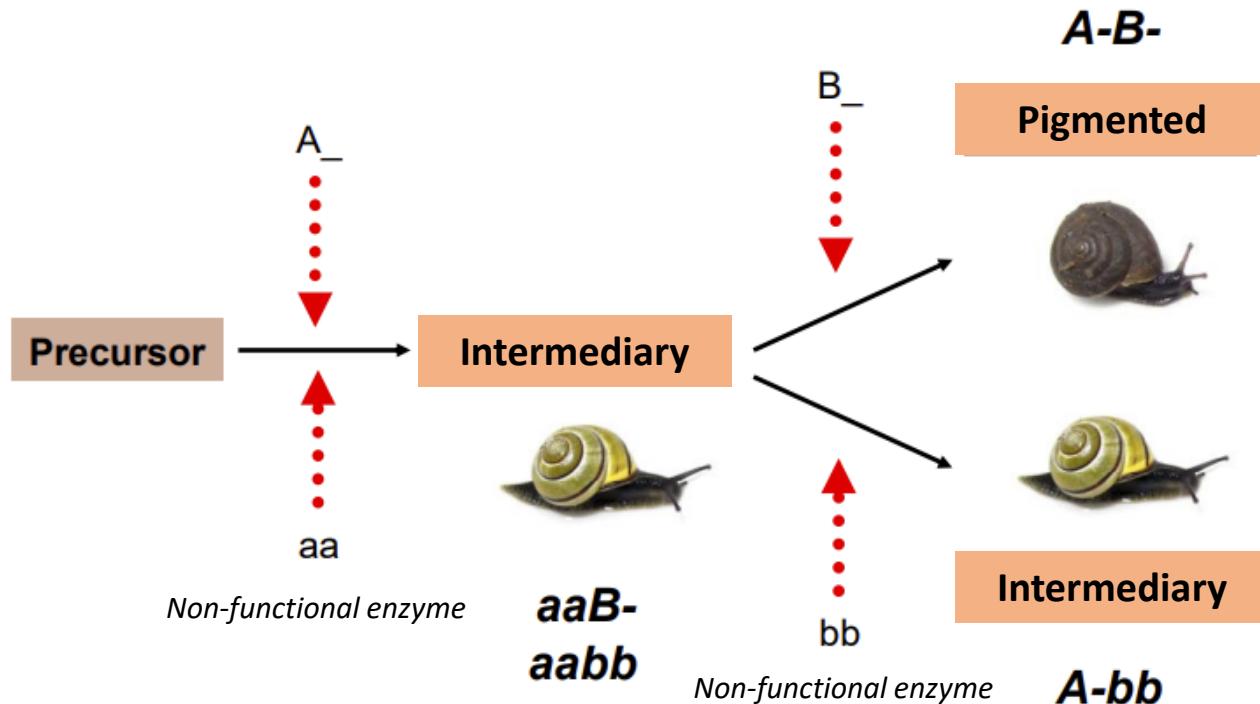
DUPLICATE RECESSIVE EPISTASIS

(same metabolic pathway)



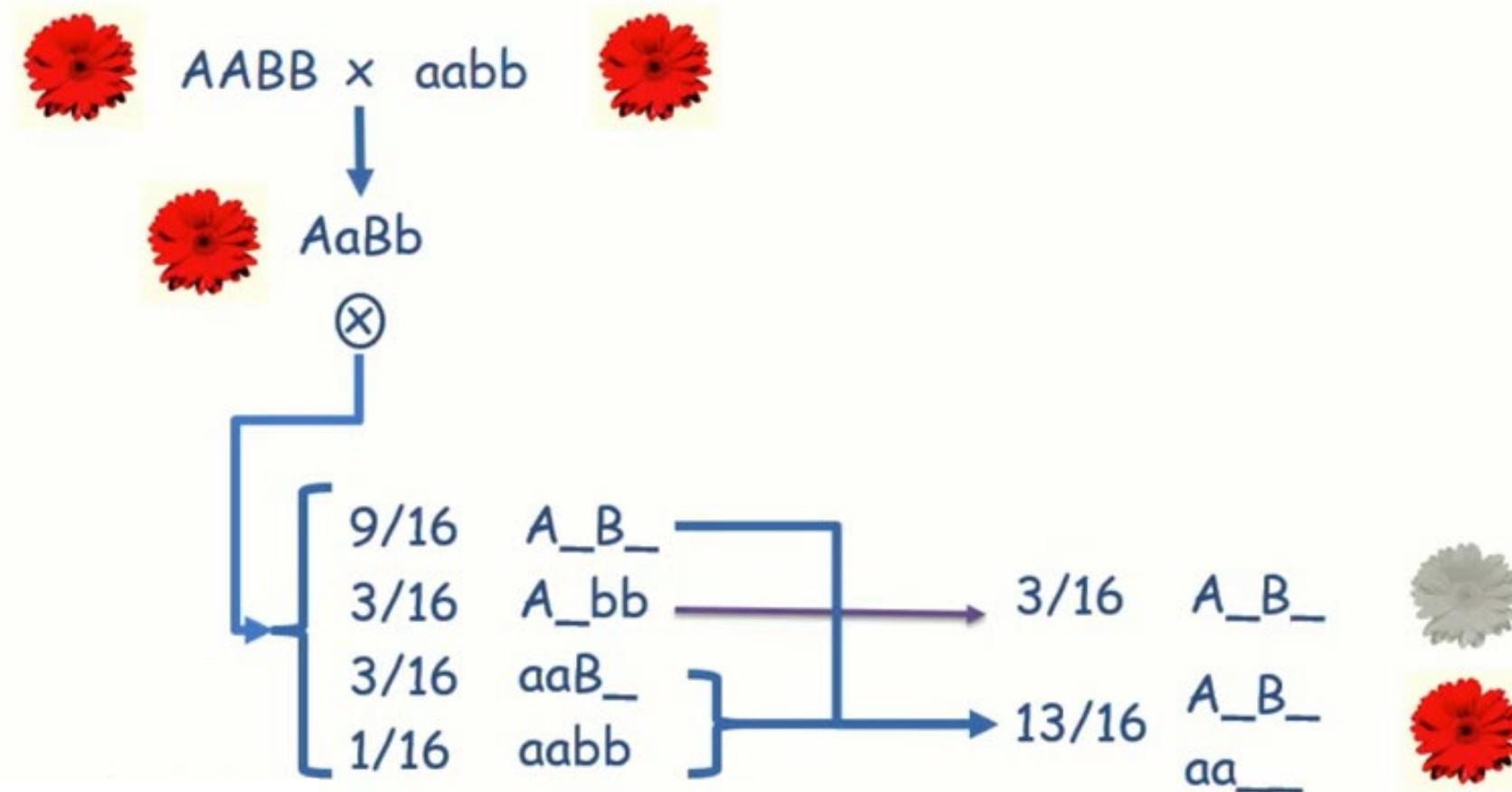
DUPLICATE RECESSIVE EPISTASIS

(same metabolic pathway)



DOMINANT-RECESSIVE EPISTASIS

(same metabolic pathway)



DOMINANT-RECESSIVE EPISTASIS

(same metabolic pathway)

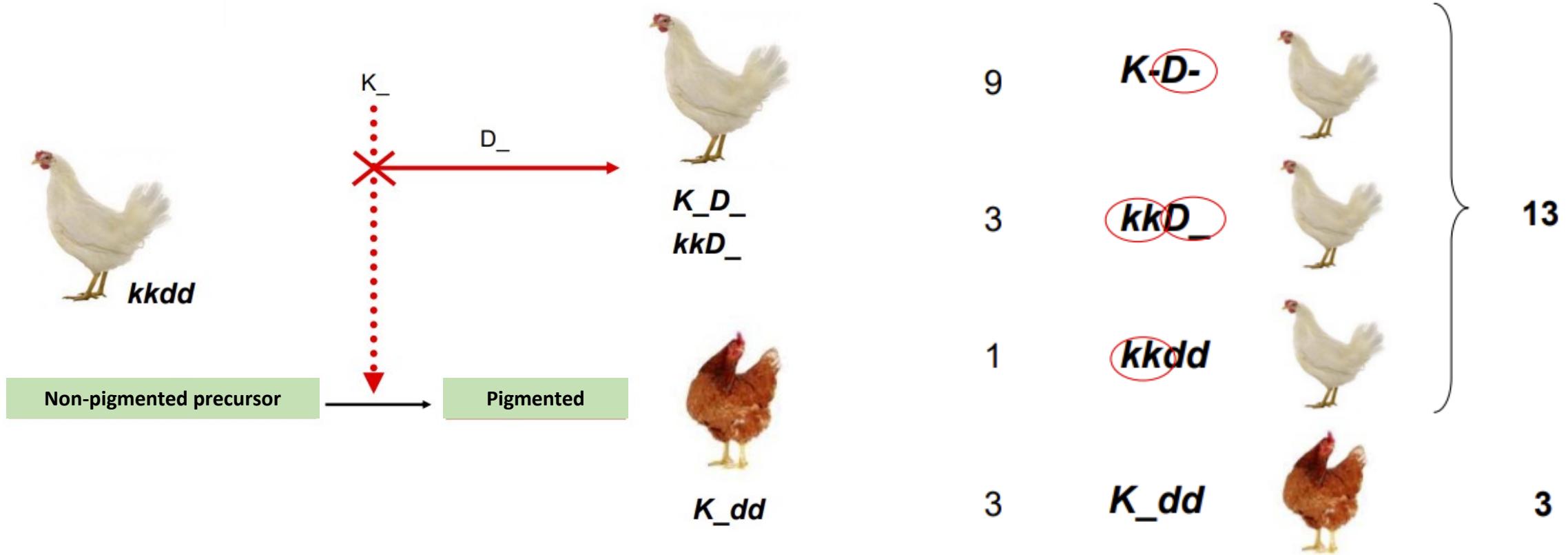


Table 5.2

Modified dihybrid phenotypic ratios due to gene interaction

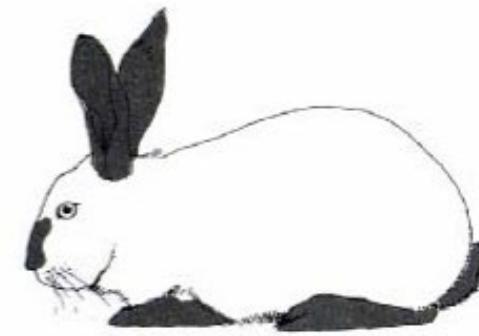
Ratio*	Genotype				Type of Interaction	Example
	$A_B_$	A_bb	$aaB_$	$aa bb$		
9 : 3 : 3 : 1	9	3	3	1	None	Seed shape and seed color in peas
9 : 3 : 4	9	3	4		Recessive epistasis	Coat color in Labrador retrievers
12 : 3 : 1	12		3	1	Dominant epistasis	Color in squash
9 : 7	9		7		Duplicate recessive epistasis	Albinism in snails
9 : 6 : 1	9	6		1	Duplicate interaction	—
15 : 1		15		1	Duplicate dominant epistasis	—
13 : 3	13		3		Dominant and recessive epistasis	—

*Each ratio is produced by a dihybrid cross ($Aa Bb \times Aa Bb$). Shaded bars represent combinations of genotypes that give the same phenotype.

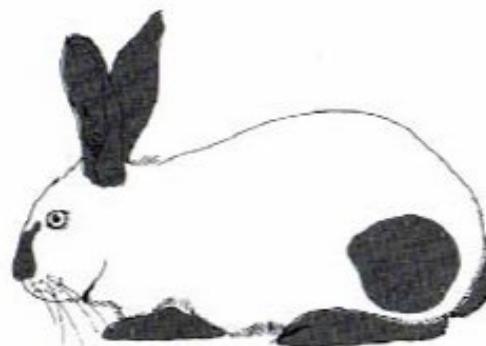
PHENOCOPY



White extremities,
reared at $>30^{\circ}\text{C}$

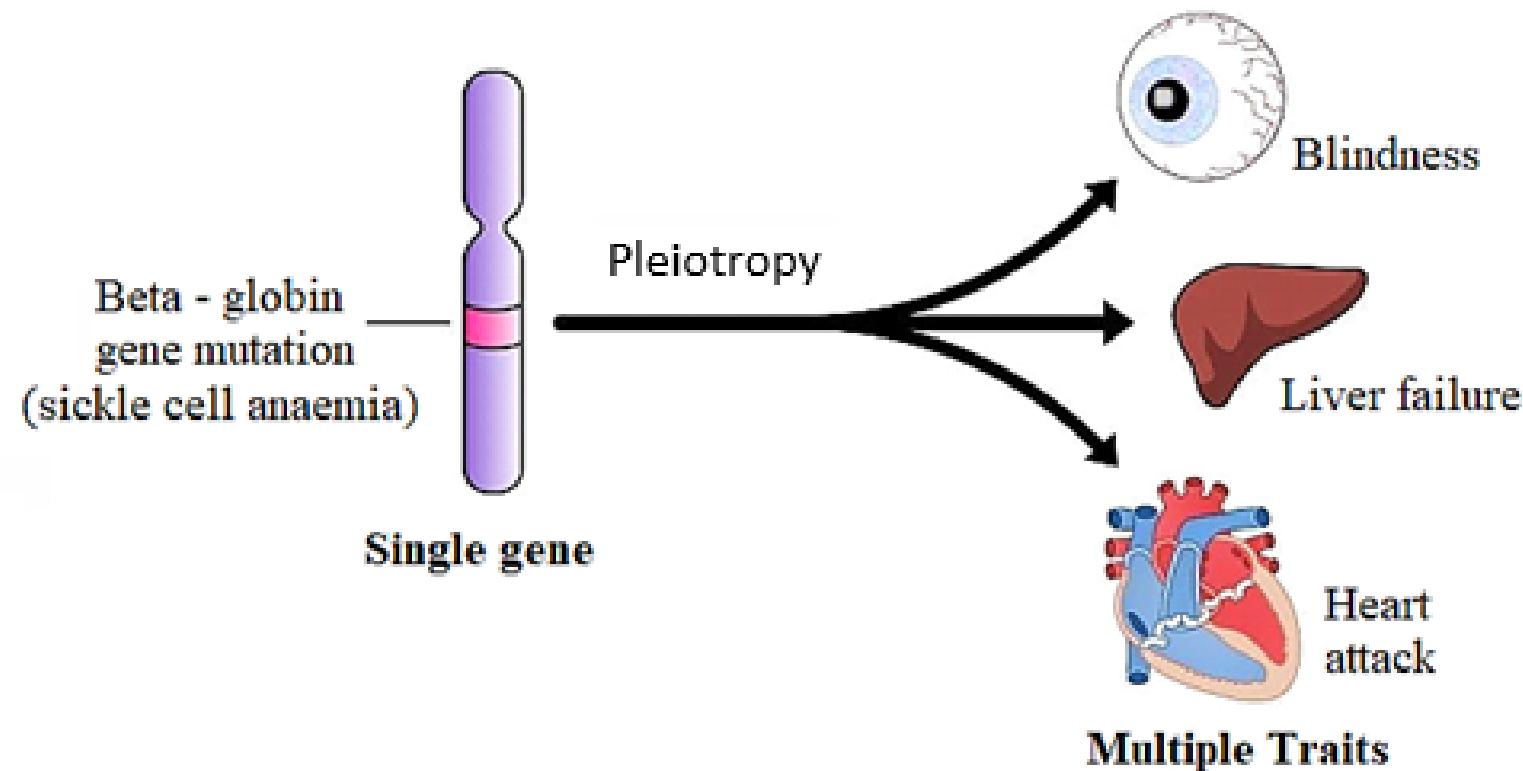


Normal
Himalayan pattern,
reared at 25°C



Himalayan pattern
with dark patch
on flank, reared at
 25°C , flank cooled
to below 25°C

PLEIOTROPY



PENETRANCE AND EXPRESSIVITY

a) Complete penetrance compared with incomplete penetrance

Complete penetrance

Identical known genotypes yield 100% expected phenotype



Incomplete penetrance

Identical known genotypes yield <100% expected phenotype



b) Constant expressivity compared with variable expressivity

Constant expressivity

Identical known genotypes with no expressivity effect yield 100% expected phenotype



Variable expressivity

Identical known genotypes with an expressivity effect yield a range of phenotypes



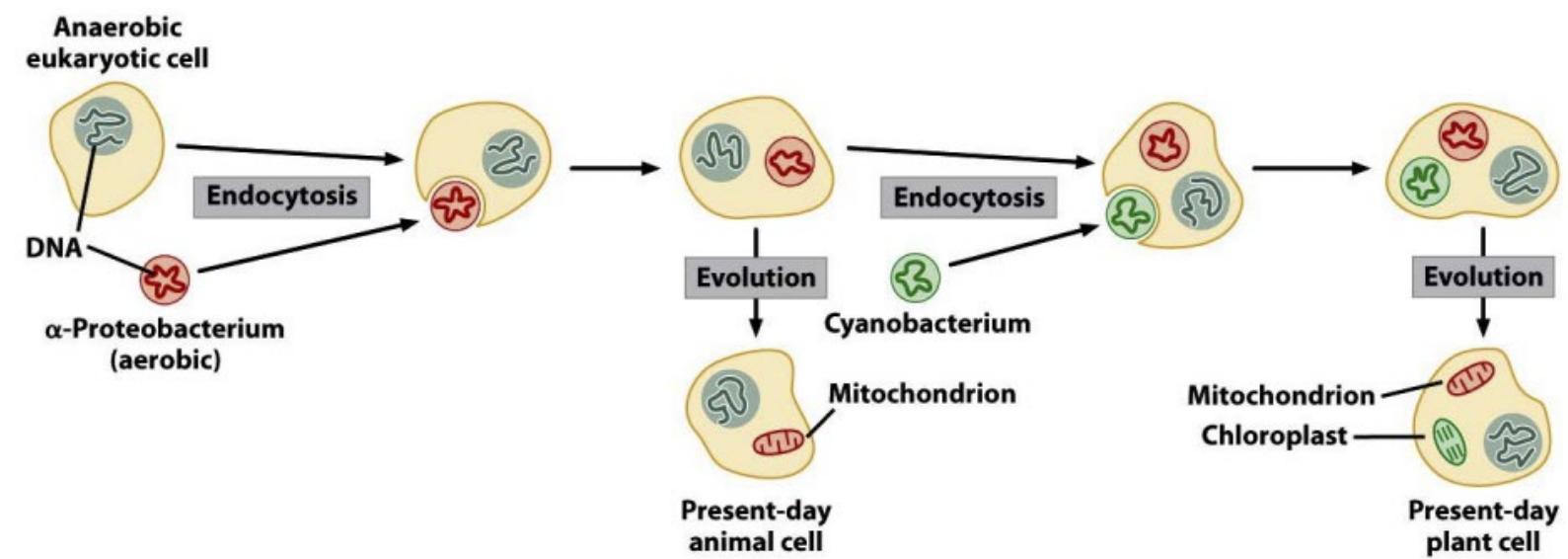
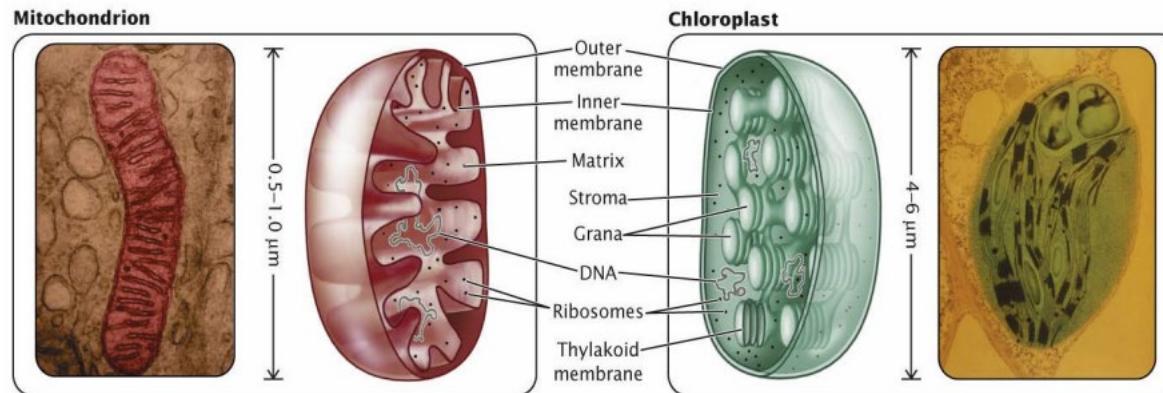
c) Incomplete penetrance with variable expressivity

Incomplete penetrance with variable expressivity

Identical known genotypes produce a broad range of phenotypes, due to varying degrees of gene activation and expression



EXTRANUCLEAR INHERITANCE



EXTRANUCLEAR INHERITANCE

