

# Chapter 3

## **Extensions to Mendel's Laws**

## **Sections to study**

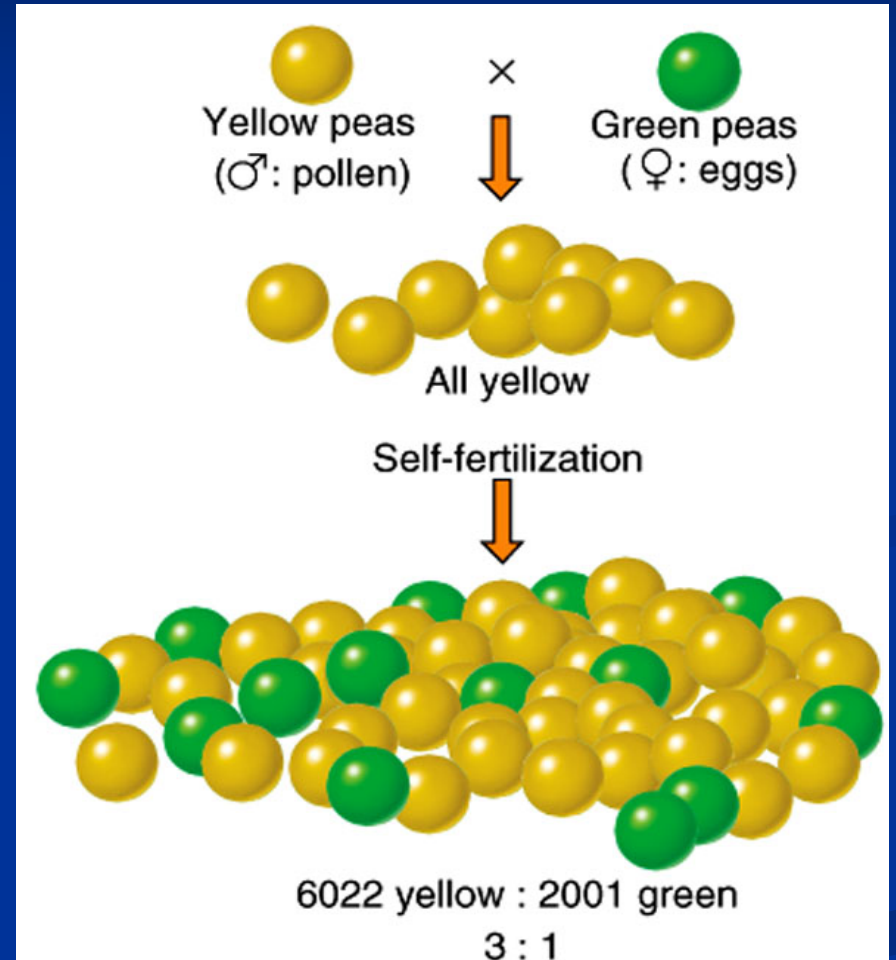
### **3.1 Extensions to Mendel for single-gene inheritance**

- **Pairs of alleles show deviations from complete dominance and recessiveness.**
- **Different forms of the gene are not limited to two alleles.**
- **One gene may determine more than one trait.**

### **3.2 Extensions to Mendel for multifactorial inheritance**

# Mendelian pattern of inheritance

- Single-gene trait
- Two alleles
- Clear-cut dominance and recessiveness
- Genotypic ratio 1: 2: 1
- Phenotypic ratio 3: 1



# Exceptions to Mendelian pattern of inheritance

## ■ **Single-gene trait**

- Pairs of alleles show deviations from complete dominance and recessiveness.
- Different forms of the gene are not limited to two alleles.
- One gene may determine more than one trait.

## ■ **Multifactorial trait**

- Traits determined by two or more genes, or by the interaction of genes with the environment.

## **3.1 Extensions to Mendel for single-gene inheritance**

# 1. Dominance is not always complete

Crosses between true-breeding strains can produce hybrids with phenotypes different from both parents.

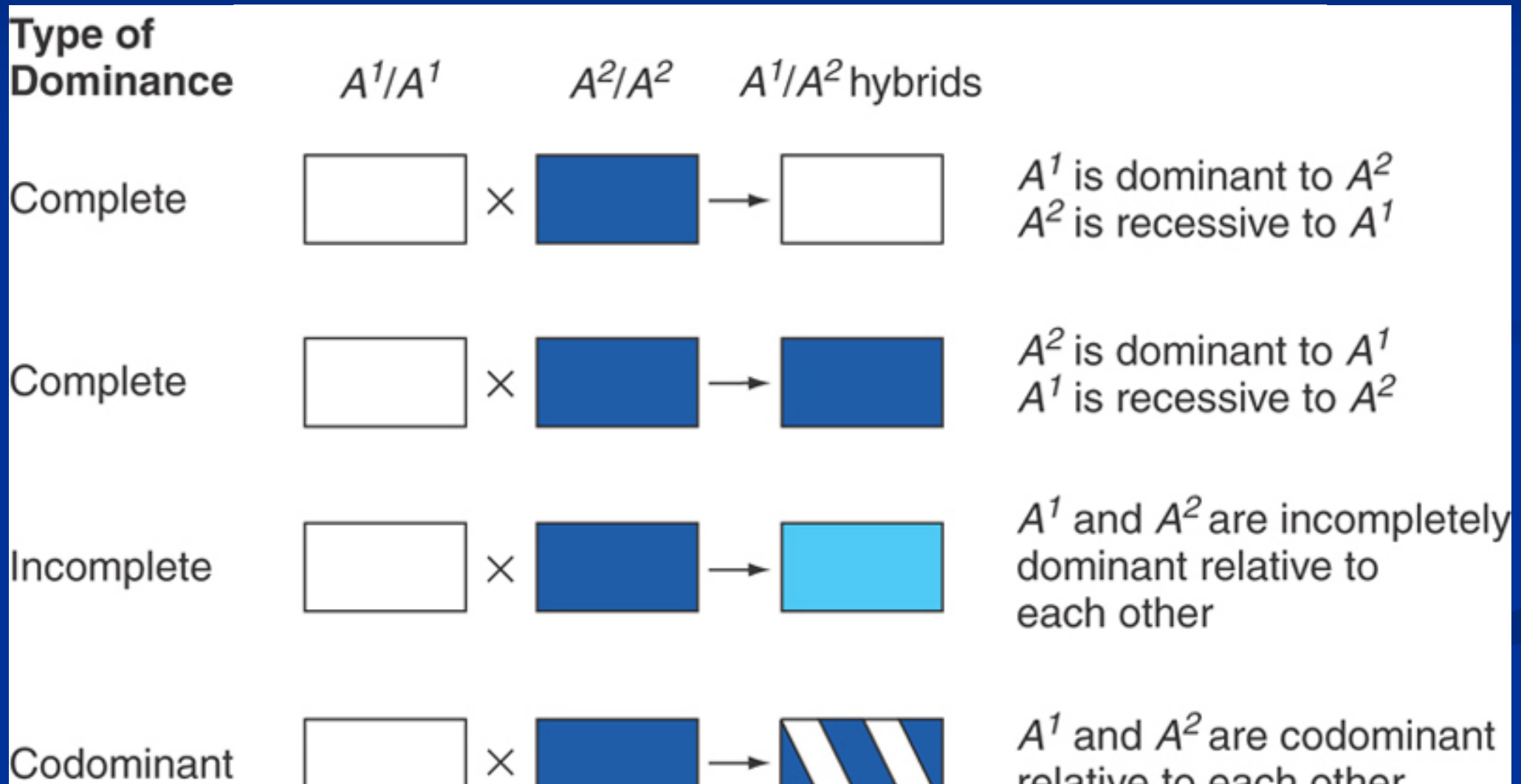


Fig. 3.2

## ■ **Codominance**

- F1 hybrids express the traits of both parents.
- Phenotypic ratios are the same as genotypic ratios.

## ■ **Incomplete dominance**

- F1 hybrids express an intermediate phenotype that differs from both parents. Neither allele is dominant or recessive to the other.
- Phenotypic ratios are the same as genotypic ratios.

# Codominant lentil coat patterns

## ■ Codominance

- F1 hybrids express the traits of both parents.
- Phenotypic ratios are the same as genotypic ratios.

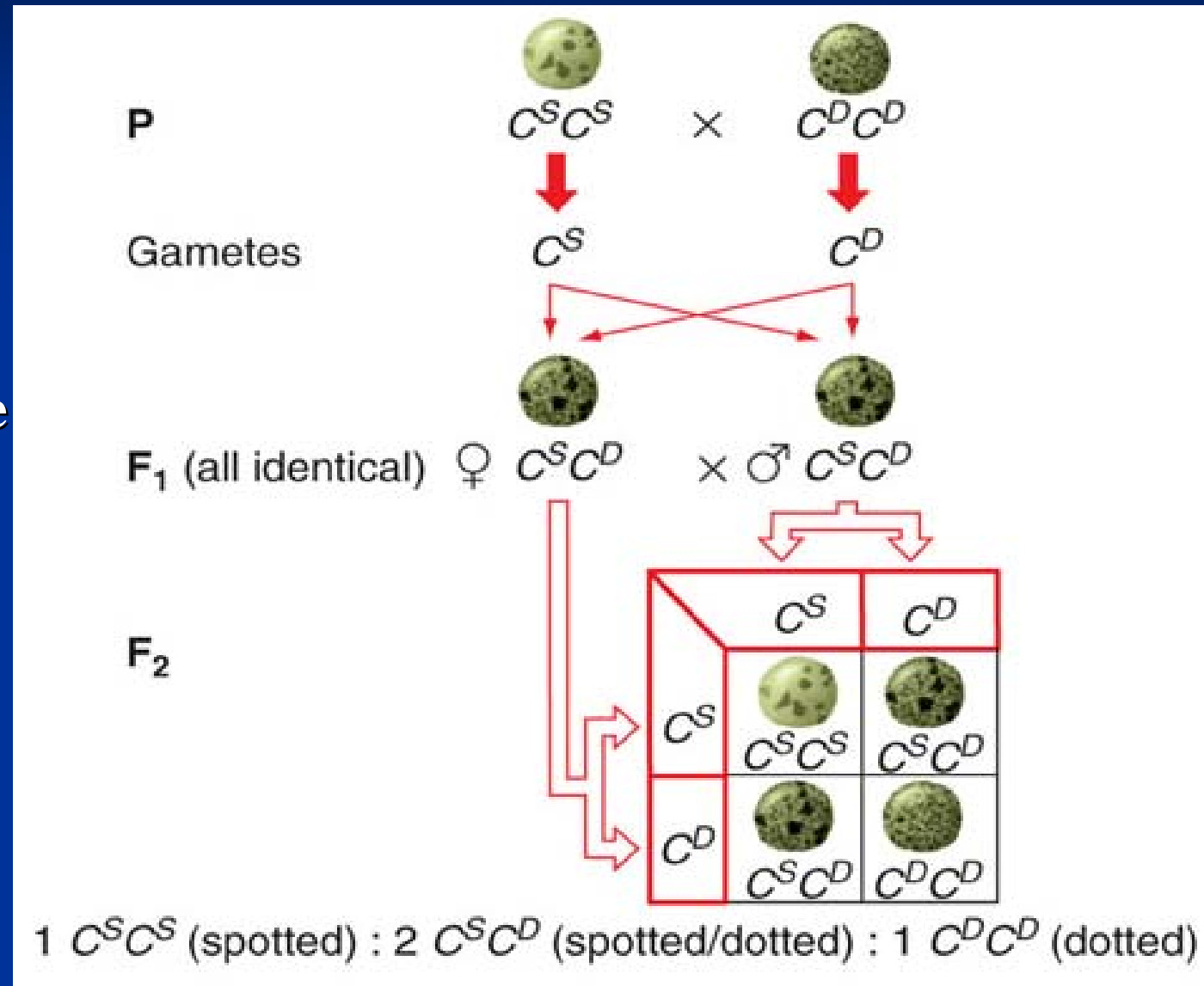


Fig. 3.4a



# Codominant blood type alleles

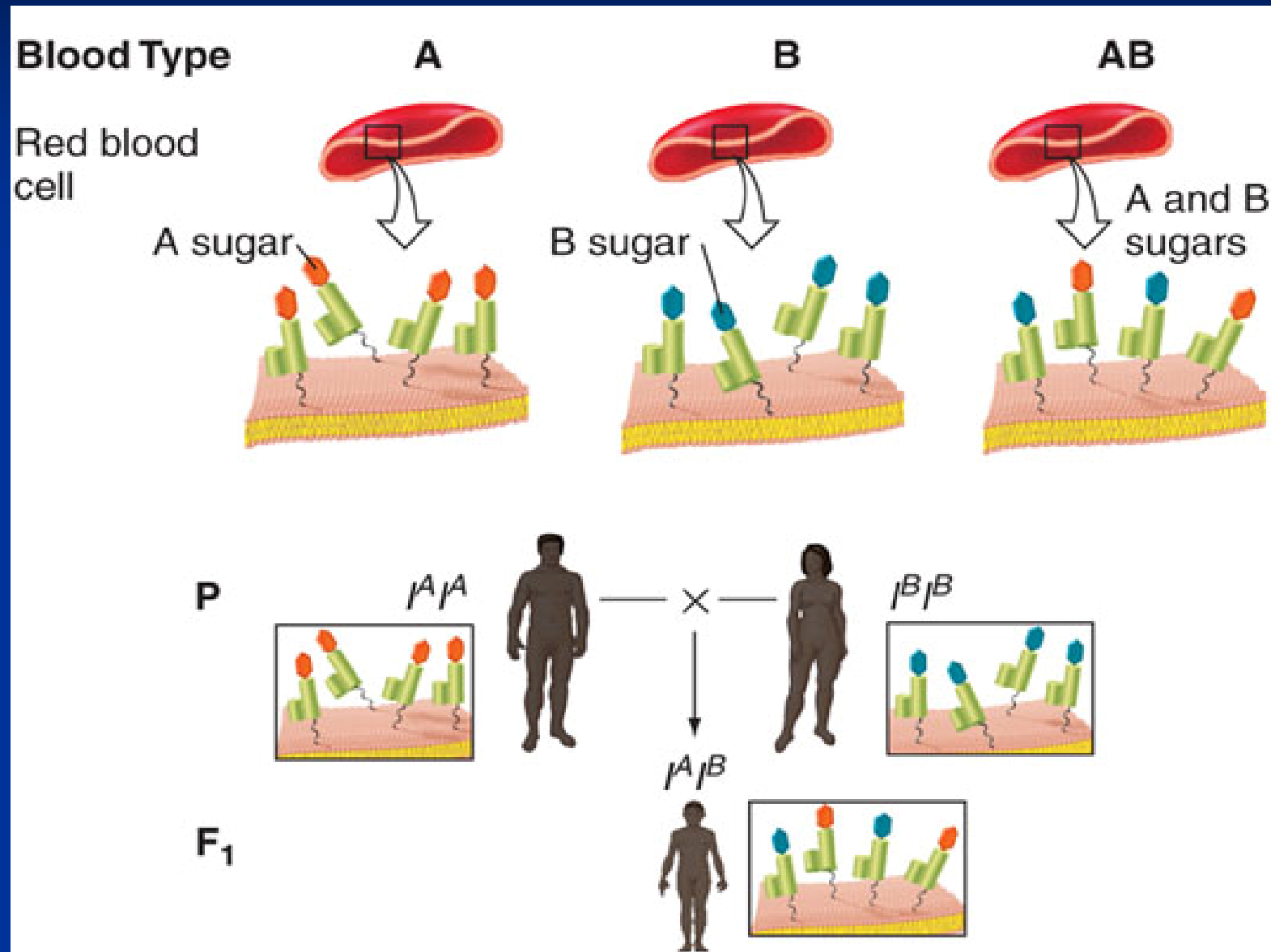
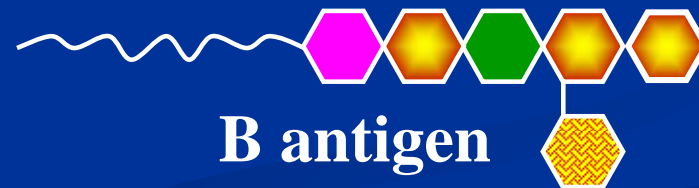








Fig. 3.4b



**GalNAc transferase**  
(*I<sup>A</sup>*)

**Gal transferase**  
(*I<sup>B</sup>*)



-  GalNAc(N-acetylgalactosamine )
-  N-Acetylglucosamine
-  Fucose
-  Glucose
-  Galactose
-  Lipid

# Incomplete dominance in snapdragons

## ■ Incomplete dominance

- F1 hybrids express an intermediate phenotype that differs from both parents. Neither allele is dominant or recessive to the other.
- Phenotypic ratios are the same as genotypic ratios.

(a) *Antirrhinum majus* (snapdragons)

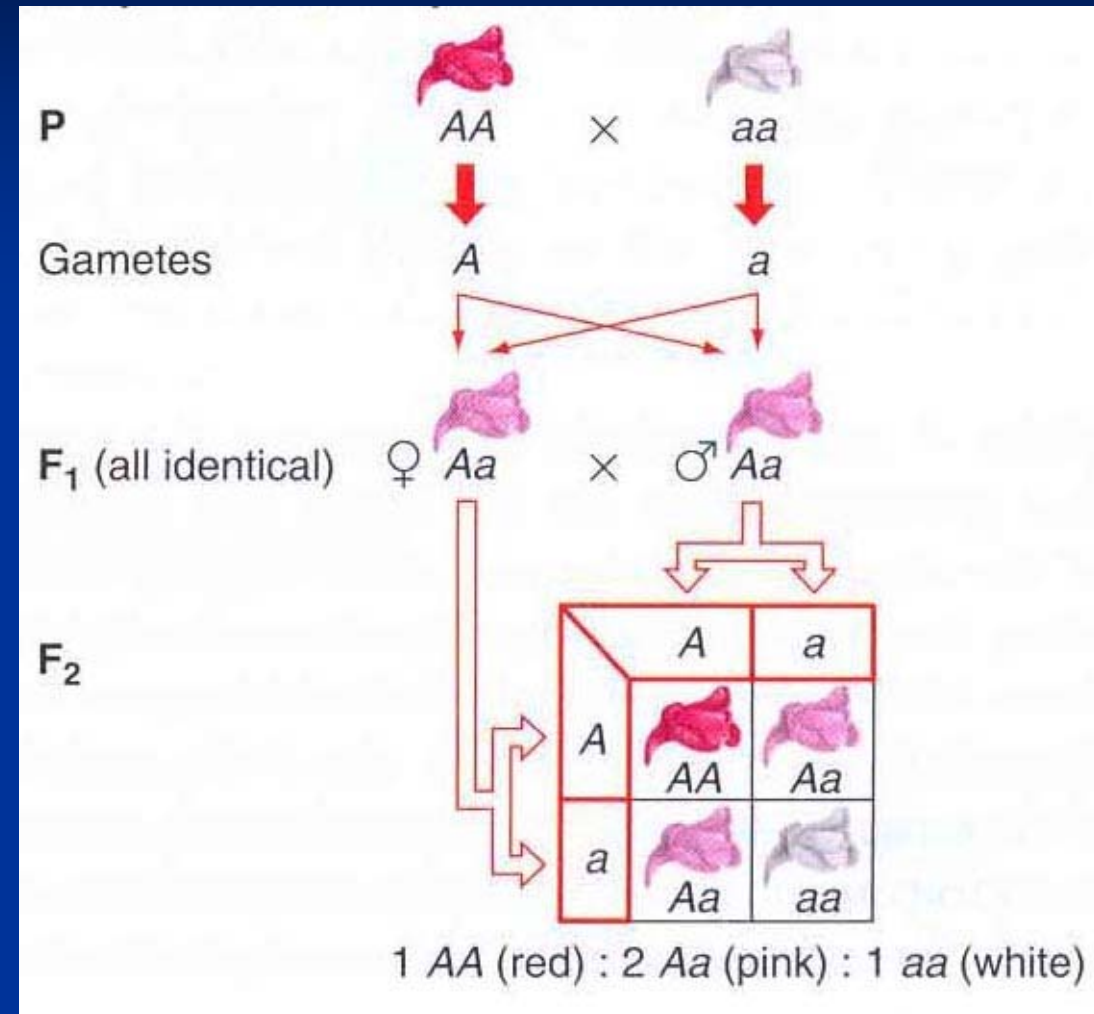


Fig. 3.3

**Incomplete dominance arises when phenotype varies in proportion to the amount of functional protein.**

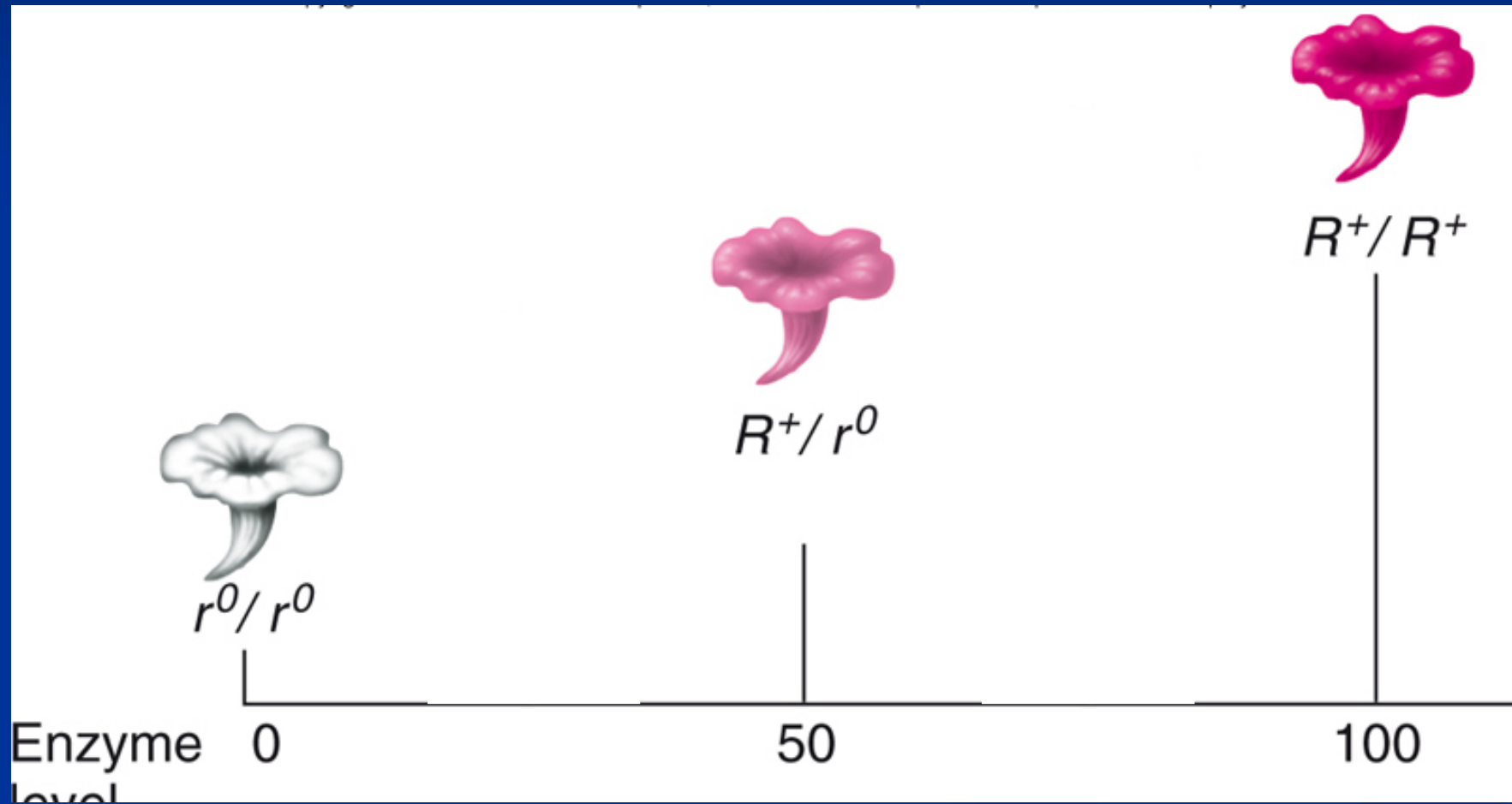


Fig. 8.30

## **Do variations on dominance relations negate Mendel's law of segregation?**

- **Mendel's law of segregation still applies.**
- **Dominance relations do not affect the segregation of alleles.**
- **Dominance relations only affect phenotype.**
  - **Gene products control expression of phenotypes differently.**

## 2. A gene may have more than two alleles

- Genes may have multiple alleles in a population.

**Human ABO blood types:** *I* (isoagglutinogen 同族凝集原)















































Phenotype	Gene	Genotype	Red blood cell surface antigen	Antibodies in serum
A	$I^A$	$I^A I^A$	A	anti-B
		$I^A i$		
B	$I^B$	$I^B I^B$	B	anti-A
		$I^B i$		
AB	$I^A, I^B$	$I^A I^B$	A and B	neither
O	$i$	$ii$	neither	anti-A and anti-B






- **Each individual carries only two of the alternative alleles.**
  - **ABO blood type**
    - **3 alleles:  $I^A$ ,  $I^B$ , and  $i$**
    - **6 possible ABO genotypes**
- **Dominance or recessiveness is always relative to a second allele.**
  - **ABO blood type**
    - **$I^A$  is completely dominant to  $i$  but codominant to  $I^B$**
    - **6 genotypes generate 4 phenotypes**



# How to establish dominance relations between multiple alleles?

- Perform reciprocal crosses between pure breeding lines of all phenotypes
- Establish dominance series

Parental Generation Parental seed coat pattern in cross Parent 1 × Parent 2	F <sub>1</sub> Generation F <sub>1</sub> phenotype	F <sub>2</sub> Generation Total F <sub>2</sub> frequencies and phenotypes	Apparent phenotypic ratio
 × 	→ 	→  798  296	3 : 1
 × 	→ 	→  123  46	3 : 1
 × 	→ 	→  283  107	3 : 1
 × 	→ 	→  1,706  522	3 : 1
 × 	→ 	→  272  72	3 : 1
 × 	→ 	→  499  147	3 : 1
 × 	→ 	→  1,597  549	3 : 1
 × 	→ 	→  182  70	3 : 1
 × 	→ 	→  168  339  157	1 : 2 : 1

**Dominance series:** marbled-1 > marbled-2 > spotted = dotted > clear

Fig. 3.6

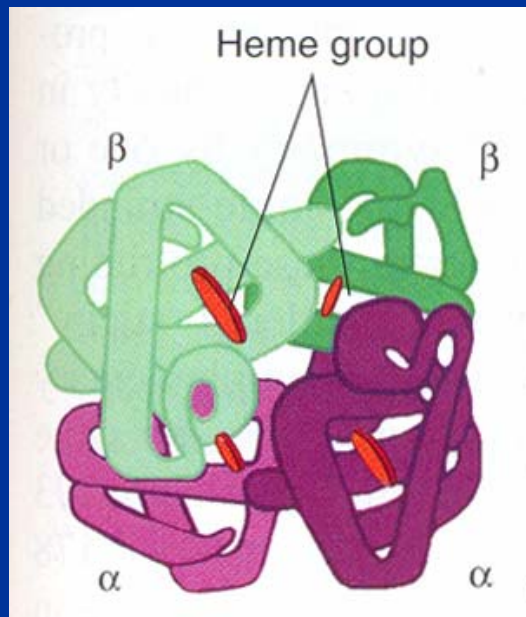


# Mutations are the source of new alleles

- Multiple alleles arise spontaneously in nature due to chance alterations in genetic material – mutations.

Wild-type  $\xrightarrow{\text{mutation}}$  Mutant form

## Example: human $\beta$ -hemoglobin



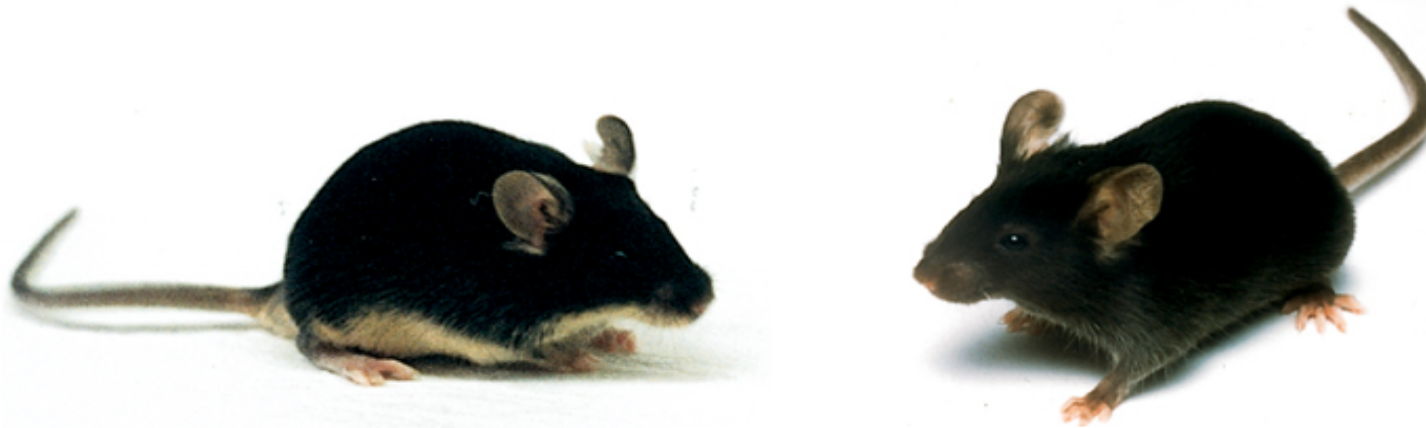
$\beta$ Chain of Hemoglobin	Sixth code	function
HbA	GAG	normal
HbS	GTG	sickled
HbC	AAG	abnormal

Fig. 9.1a, b

- Mutation rate varies from 1 in 10,000 to 1 in 1,000,000 per gamete per generation.
- **Allele frequency** is the percentage of the total number of gene copies represented by one allele.
  - **Wild-type allele** – allele whose frequency is more than 1%.
  - **Mutant allele** – allele whose frequency is less than 1%.
  - **Monomorphic** – gene with only one wild-type allele.
  - **Polymorphic** – gene with more than one wild-type allele.

# The mouse *agouti* gene: one wild-type allele, many mutant alleles (> 14)

(a) *Mus musculus* (house mouse) coat colors

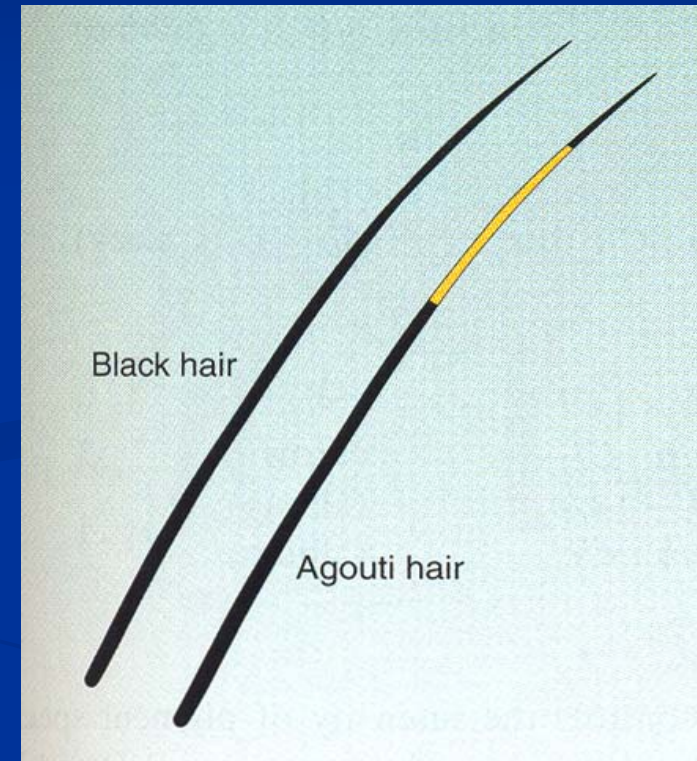


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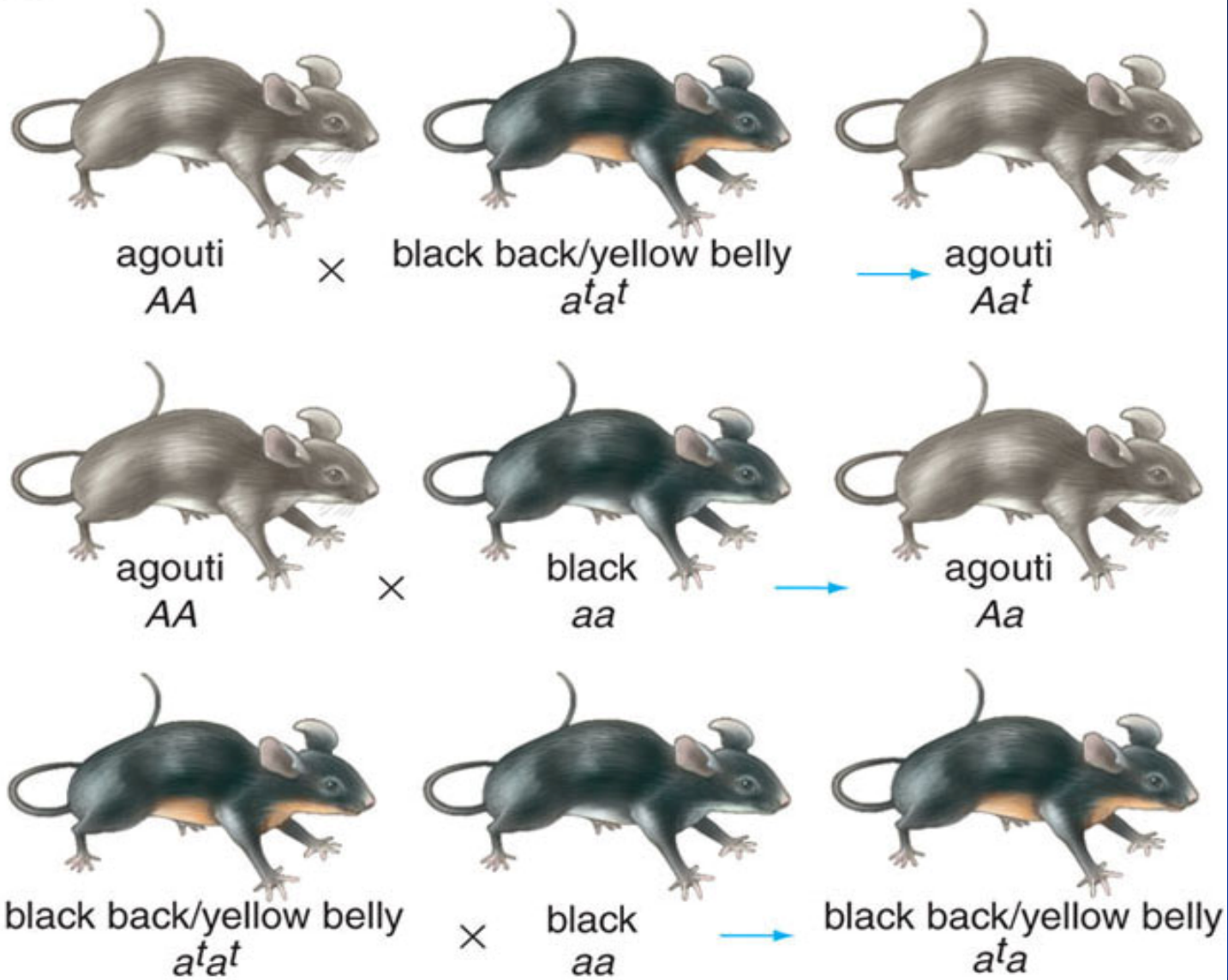
(b) Alleles of the *agouti* gene

Genotype    Phenotype

$A-$	agouti
$a^t a^t$	black/yellow
$aa$	black
$a^t a$	black/yellow



(c) Evidence for a dominance series

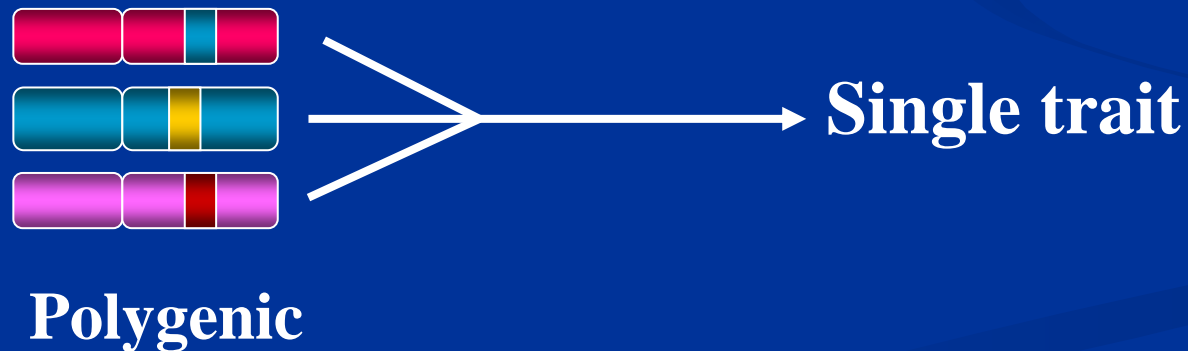
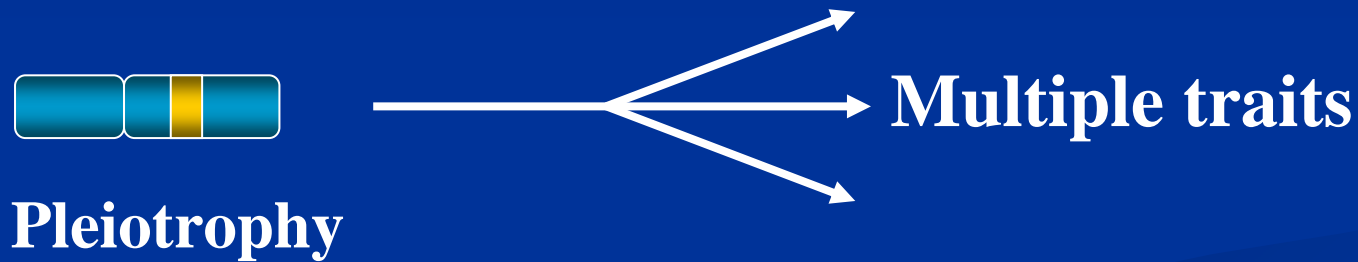


Dominance series:  $A > a^t > a$



### 3. One gene may contribute to several visible characteristics

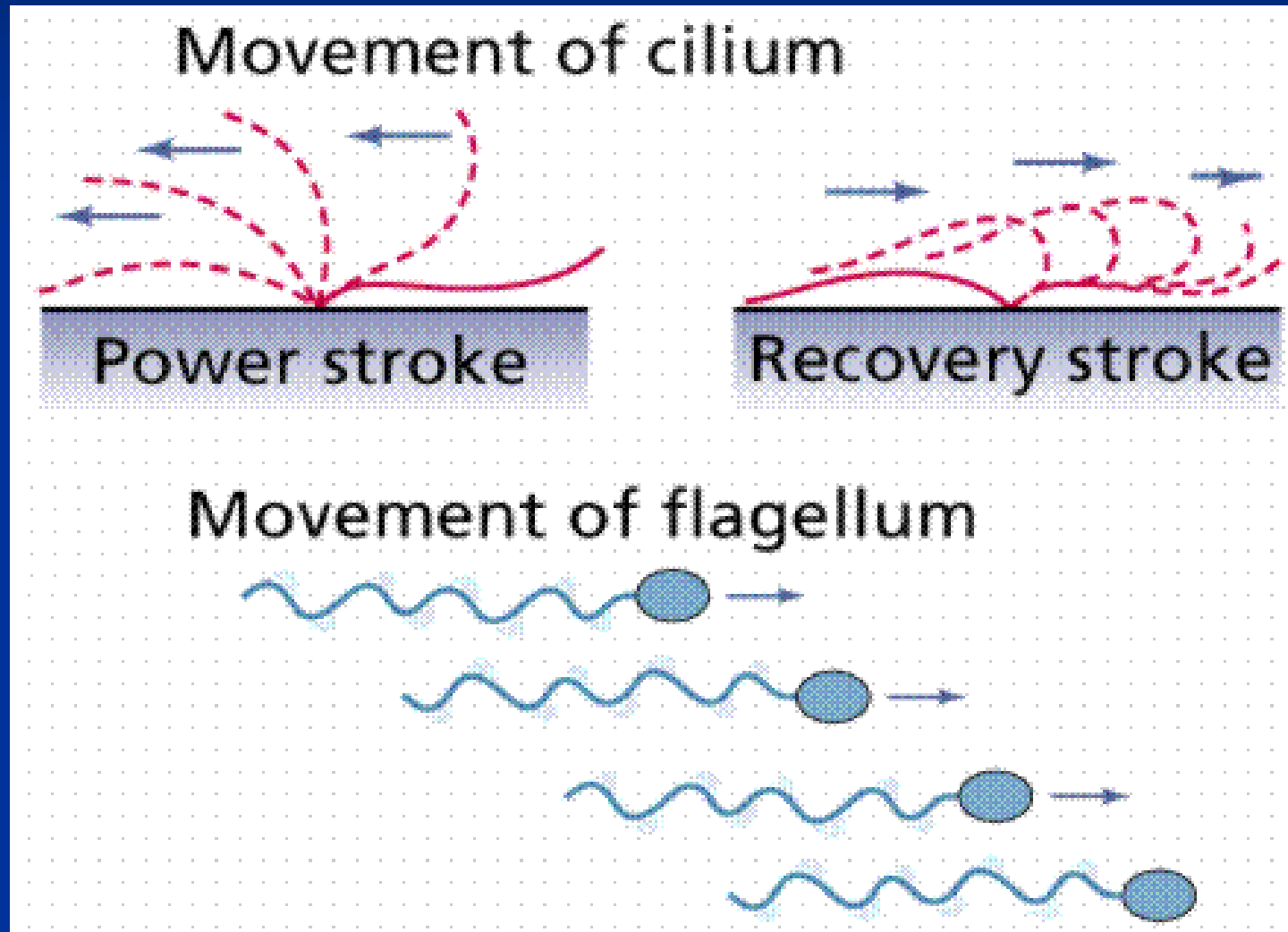
- **Pleiotropy** – A single gene determines a number of distinct and seemingly unrelated characteristics.



- **Pleiotropy** – A single gene determines a number of distinct and seemingly unrelated characteristics.
  - One recessive mutation in Maori people causes respiratory problems and infertility in men.



- Defects in cilia of respiratory system and flagella in sperm are the cause for the Maori disease.



- **Some alleles affect viability.**
  - **Alleles that affect viability often produce deviations from a 1:2:1 genotypic and 3:1 phenotypic ratio.**

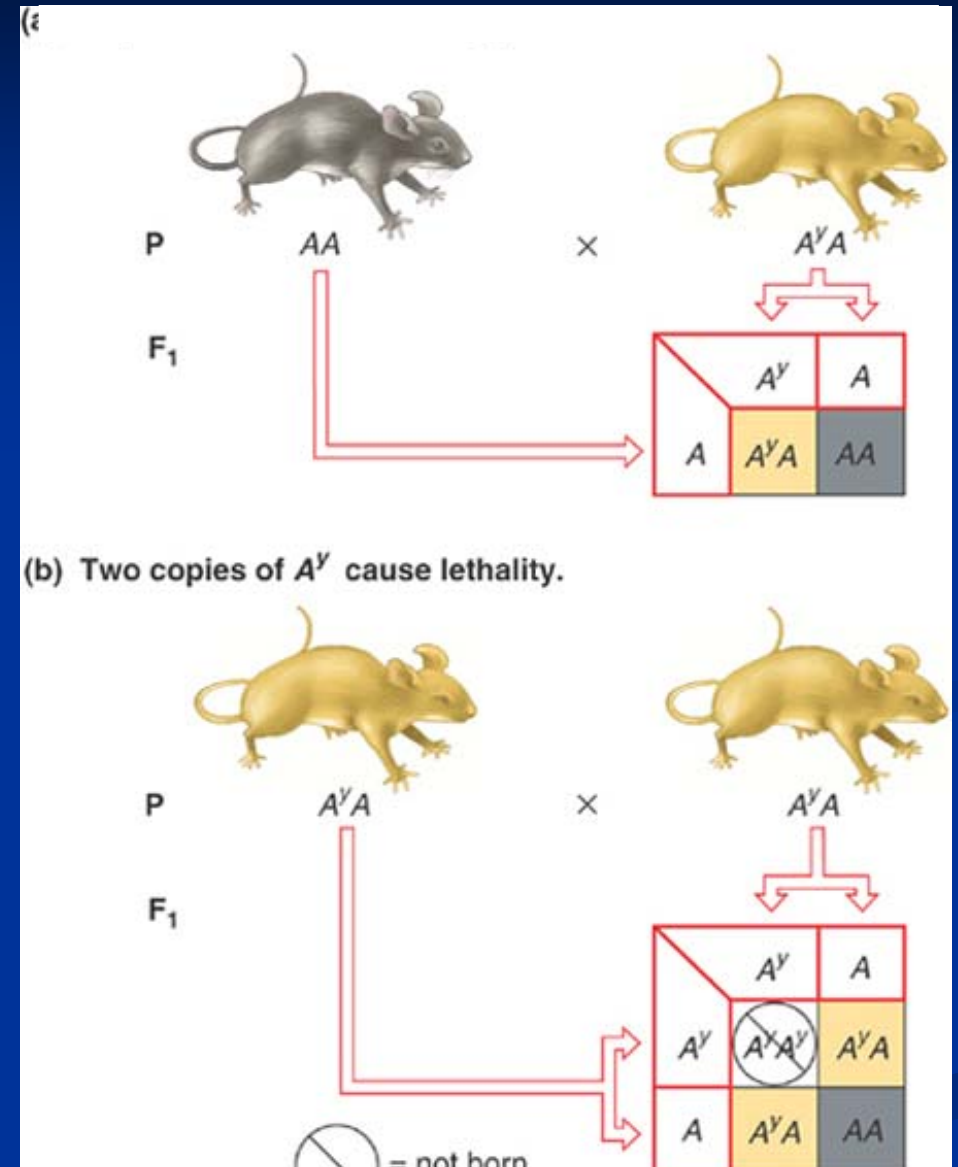


Fig. 3.9



# A dominant coat color allele also produces recessive lethality in mice

- $A^Y$  is dominant to  $A$  in determining coat color.
  - Yellow must be  $A^YA$ .
- $A^Y$  is recessive to  $A$  in producing viability.
  - Yellow mice do not breed true.
  - $A^YA^Y$  die *in utero* and do not show up as progeny.

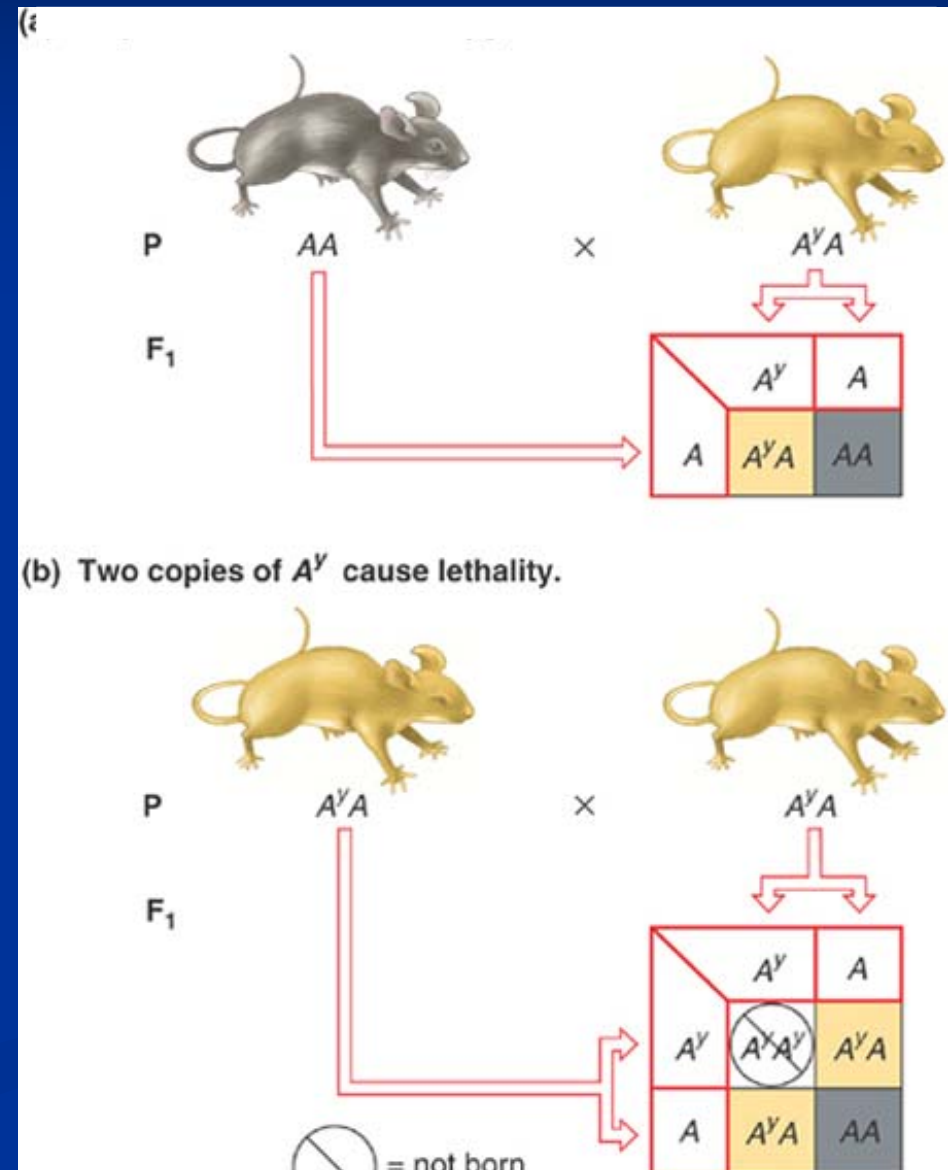


Fig. 3.9

**TABLE 3.1****For Traits Determined by One Gene: Extensions to Mendel's Analysis Explain Alterations of the 3:1 Monohybrid Ratio**

What Mendel Described	Extension	Extension's Effect on Heterozygous Phenotype	Extension's Effect on Ratios Resulting from an $F_1 \times F_1$ Cross
Complete dominance	Incomplete dominance Codominance	Unlike either homozygote	Phenotypes coincide with genotypes in a ratio of 1:2:1
Two alleles	Multiple alleles	Multiplicity of phenotypes	A series of 3:1 ratios
All alleles are equally viable	Recessive lethal alleles	No effect	2:1 instead of 3:1
One gene determines one trait	Pleiotropy: one gene influences several traits	Several traits affected in different ways, depending on dominance relations	Different ratios, depending on dominance relations for each affected trait

# Sickle-cell anemia

Mutant  $\beta$  chain (E6V) of hemoglobin forms aggregates that cause red blood cells to sickle.

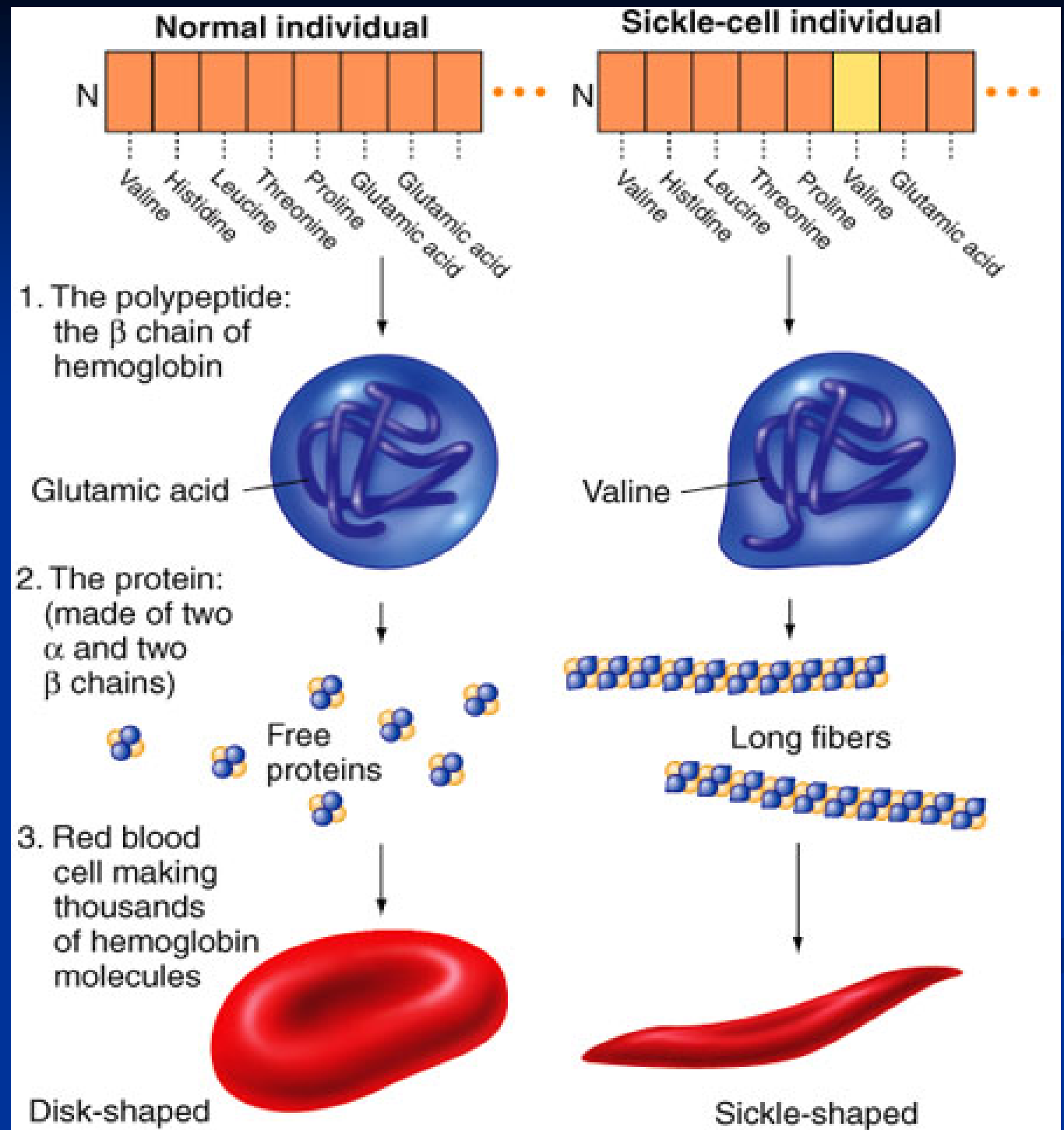
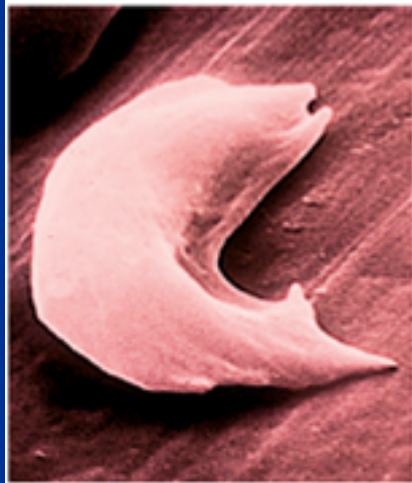
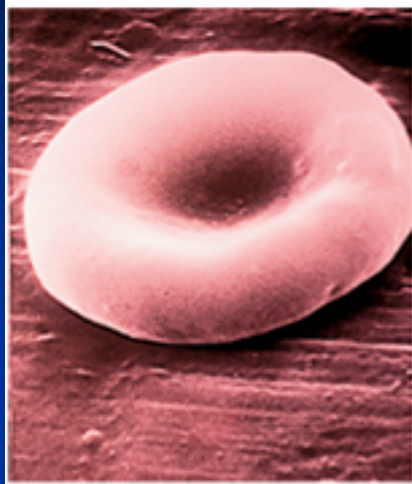


Fig. 7.25a

- **Multiple alleles**
  - Wild-type is  $Hb\beta^A$
  - More than 400 mutant alleles identified so far
  - $Hb\beta^S$  allele specifies abnormal peptide causing sickling among red blood cells
- **Pleiotropy**
  - $Hb\beta^S$  affects more than one trait
    - Muscle cramps, shortness of breath, and fatigue
    - Anemia
    - Resistance to malaria
- **Recessive lethality**
  - Homozygous  $Hb\beta^S Hb\beta^S$  individuals die at young age.
- **Different dominance relations**
  - Heterozygous  $Hb\beta^A Hb\beta^S$  individuals become sick in low-oxygen environment.



# Pleiotropy of sickle-cell anemia



(a)

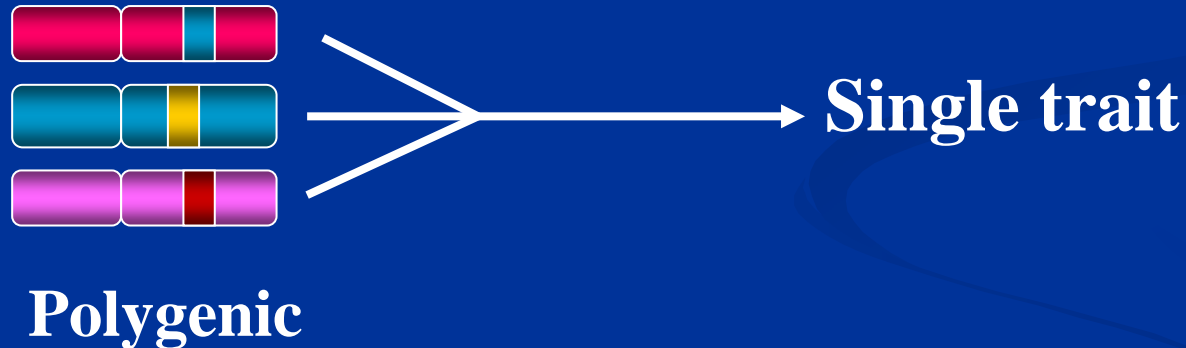
Phenotypes at Different Levels of Analysis	Normal <i>Hbβ<sup>A</sup> Hbβ<sup>A</sup></i>	Carrier <i>Hbβ<sup>A</sup> Hbβ<sup>S</sup></i>	Diseased <i>Hbβ<sup>S</sup> Hbβ<sup>S</sup></i>	Dominance Relations at Each Level of Analysis
β-globin polypeptide production				<i>Hbβ<sup>A</sup></i> and <i>Hbβ<sup>S</sup></i> are codominant
Red blood cell shape at sea level	Normal 	Normal 	Sickled cells present 	<i>Hbβ<sup>A</sup></i> is dominant <i>Hbβ<sup>S</sup></i> is recessive
Red blood cell concentration at sea level	Normal 	Normal 	Lower 	
Red blood cell shape at high altitudes	Normal 	Sickled cells present 	Severe sickling 	<i>Hbβ<sup>A</sup></i> and <i>Hbβ<sup>S</sup></i> show incomplete dominance
Red blood cell concentration at high altitudes	Normal 	Lower 	Very low, anemia 	
Susceptibility to malaria	Normal susceptibility 	Resistant 	Resistant 	<i>Hbβ<sup>S</sup></i> is dominant <i>Hbβ<sup>A</sup></i> is recessive

(b)

Fig. 3.10

## 3.2 Extensions to Mendel for multifactorial inheritance

- Novel phenotypes can emerge from the combined action of the alleles of two genes.



**P**  $AABB \times aabb$

**F1** ♀  $AaBb \times$  ♂  $AaBb$

**F2**

	$AB$	$Ab$	$aB$	$ab$
$AB$	$AABB$	$AABb$	$AaBB$	$AaBb$
$Ab$	$AABb$	$AAbb$	$AaBb$	$Aabb$
$aB$	$AbBB$	$AaBb$	$aaBB$	$aaBb$
$ab$	$AaBb$	$Aabb$	$aaBb$	$aabb$

9  $A\_B\_$

3  $A\_bb$

3  $aaB\_$

1  $aa\ bb$

# 1. Two genes interact to produce new phenotypes

- Genes A and B interact to produce new colors in lentils.

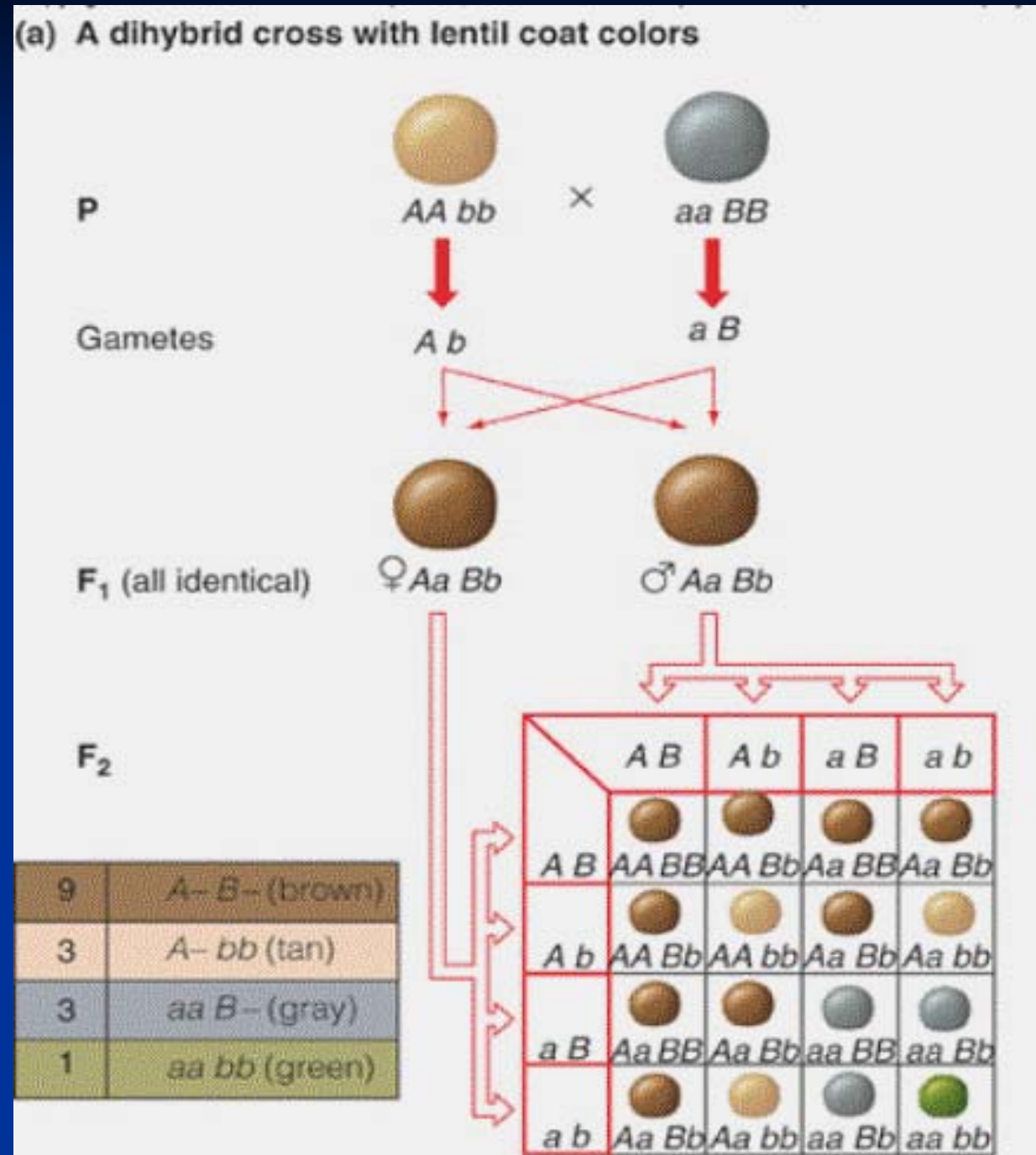


Fig. 3.11a



## 2. Complementary gene action generates flower color in sweet peas



9:7 (purple: white) ratio demonstrates that one dominant allele of two genes must be present to produce purple flowers.

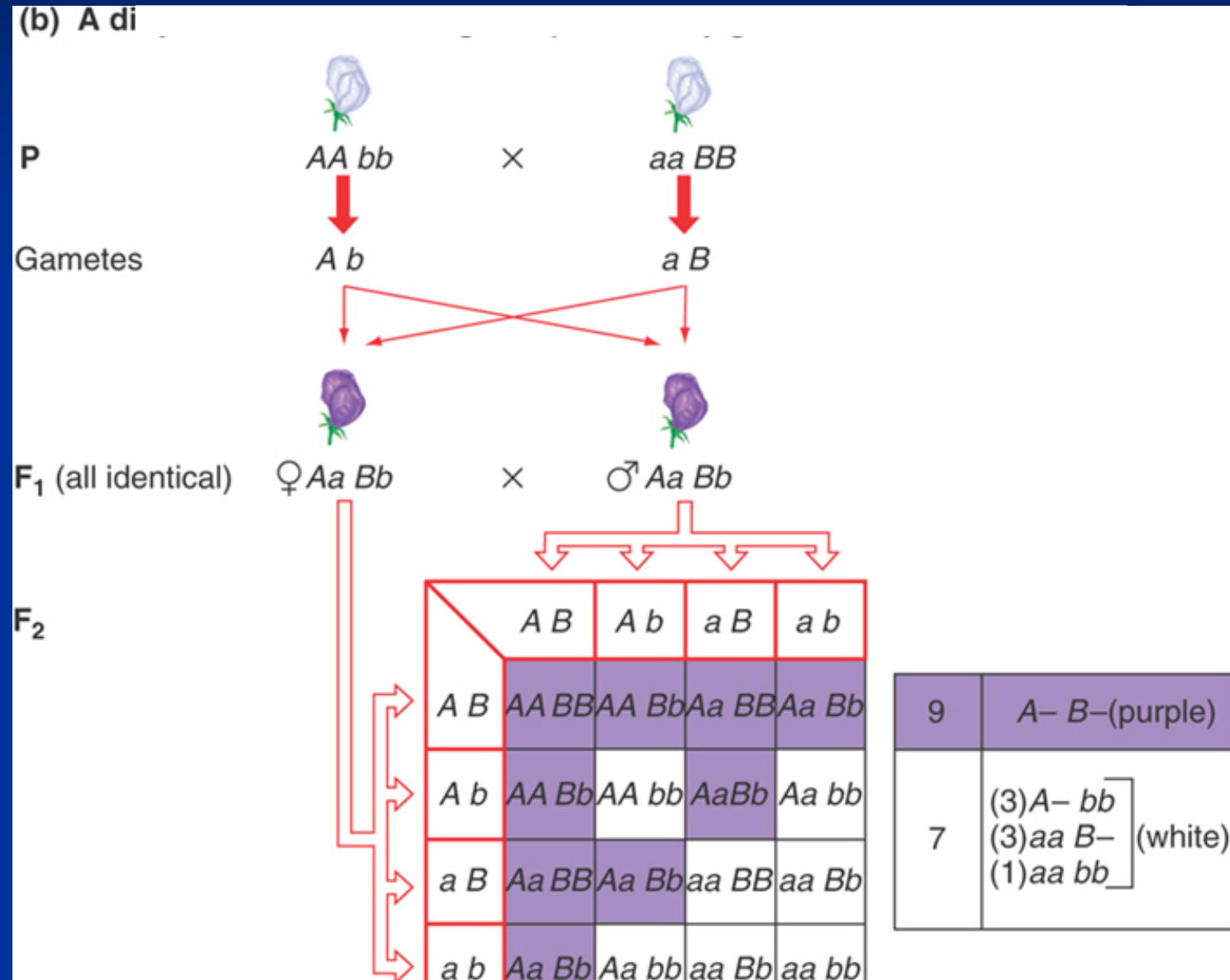


Fig. 3.12

# Two genes work in tandem to produce purple pigment

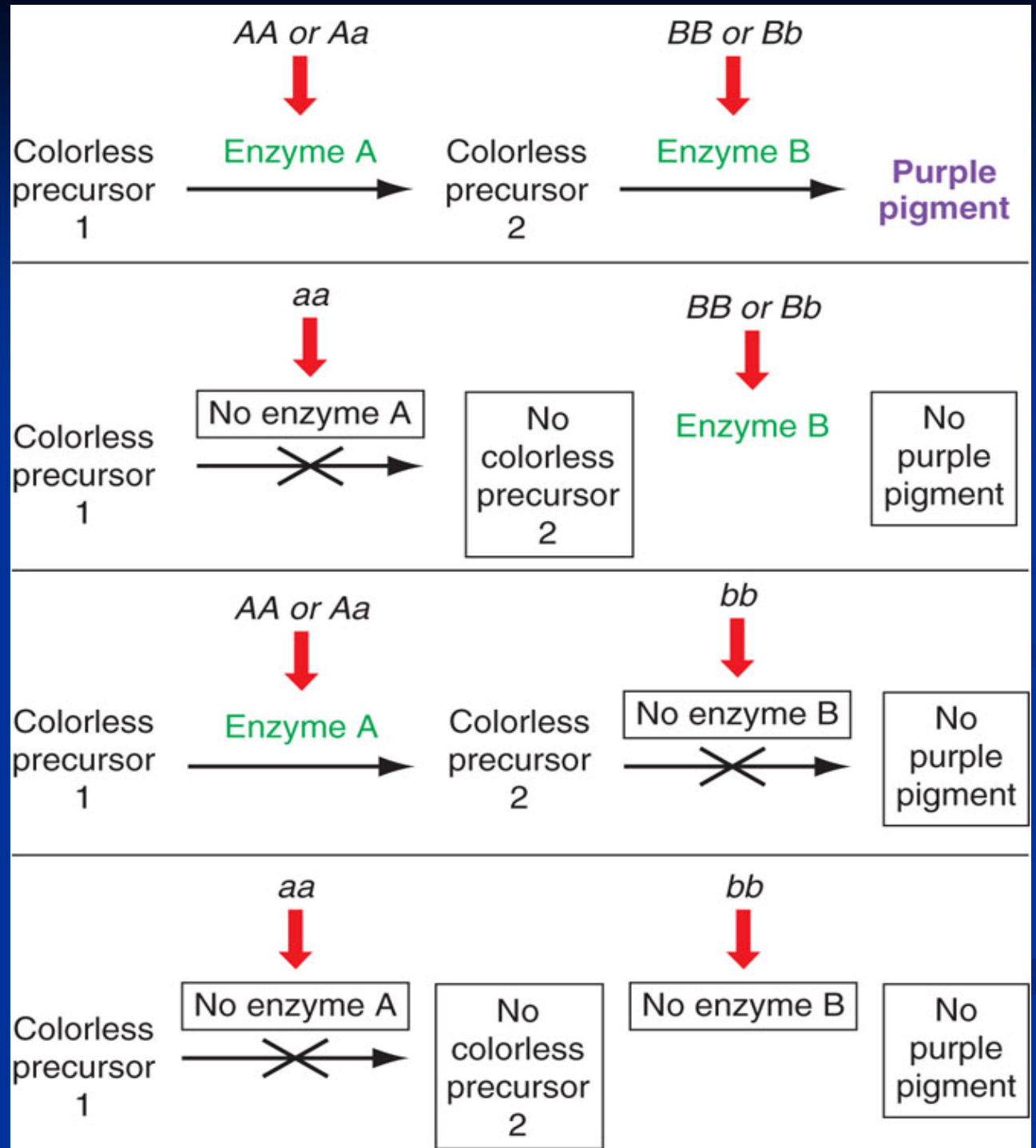
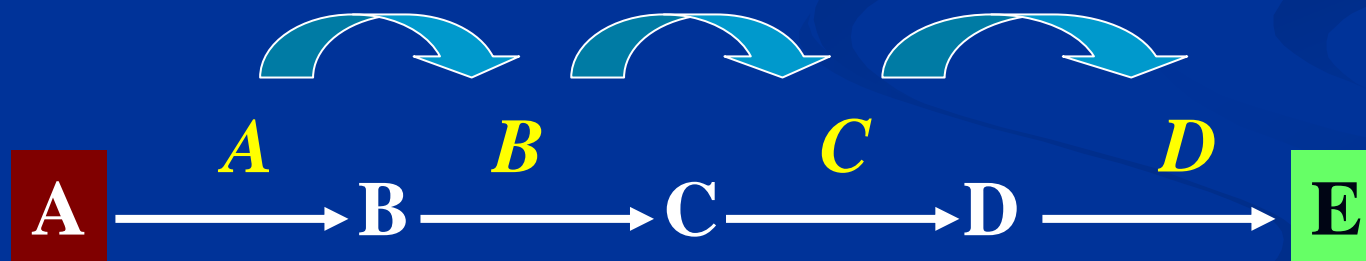


Fig. 3.13

### 3. One gene's alleles mask the effects of another gene's alleles

- **Epistasis:** A gene interaction in which the effects of an allele at one gene hide the effects of alleles at another gene.
  - **Recessive epistasis** – The recessive allele of one gene hides the effects of another gene.
  - **Dominant epistasis** – The dominant allele of one gene hides the effects of another gene.



**A:** epistatic gene

**B:** hypostatic gene

# 3-1 Recessive epistasis

## Coat color in Labrador retrievers

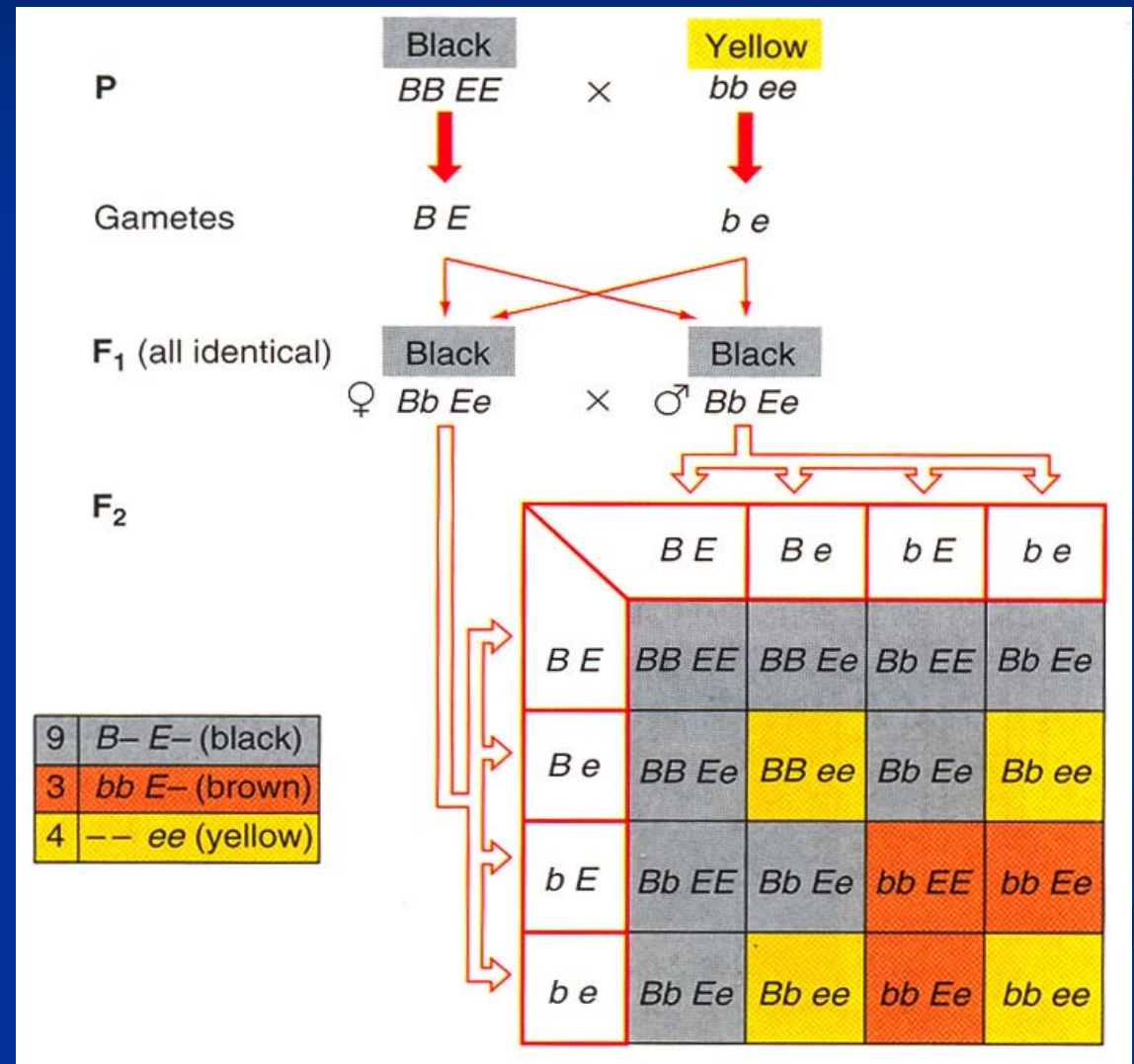
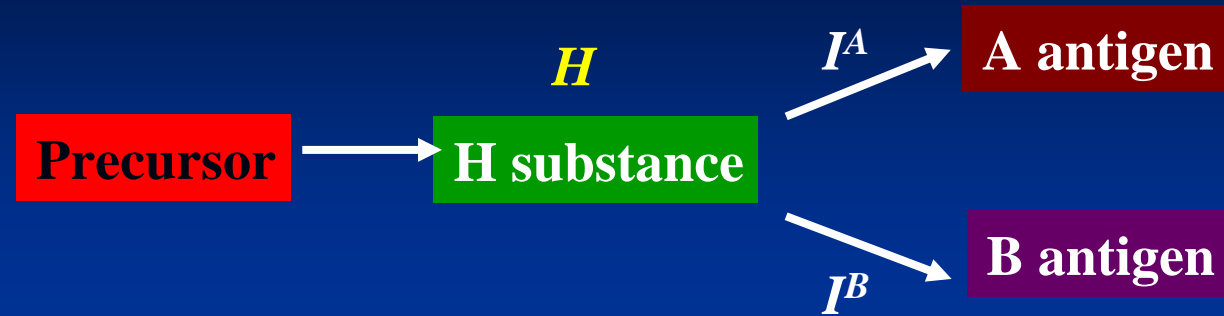


Fig. 3.14a



# Recessive epistasis in human blood types

**Bombay phenotype** – mutant recessive allele at second gene *hh* masks phenotype of ABO alleles.



**Woman**  
 (O type)  
*I<sup>A</sup>I<sup>B</sup> hh*

**Man**  
 (O type)  
*ii Hh*

**Children**  
 (A, B or O type)  
*I<sup>A</sup>i Hh*  
*I<sup>B</sup>i Hh*  
*I<sup>A</sup>i hh, I<sup>B</sup>i hh*

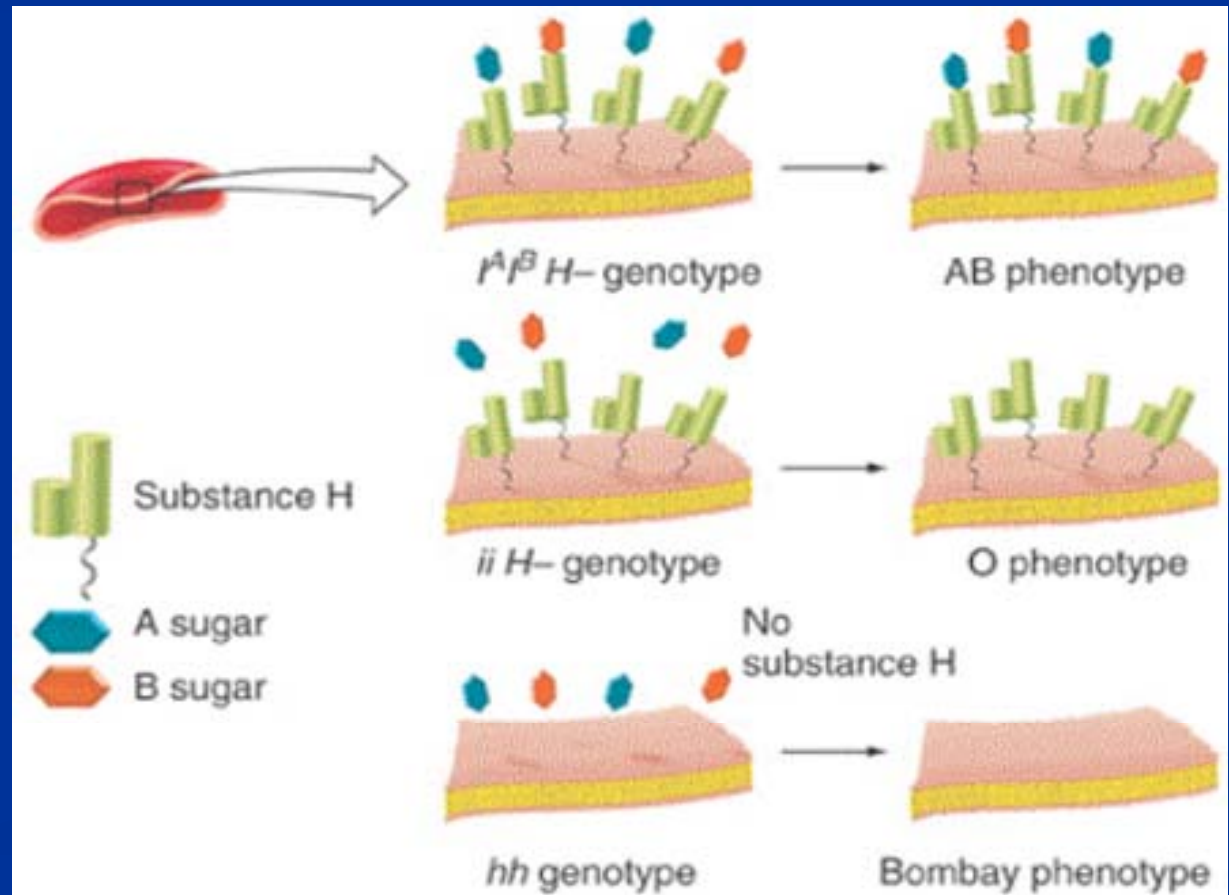
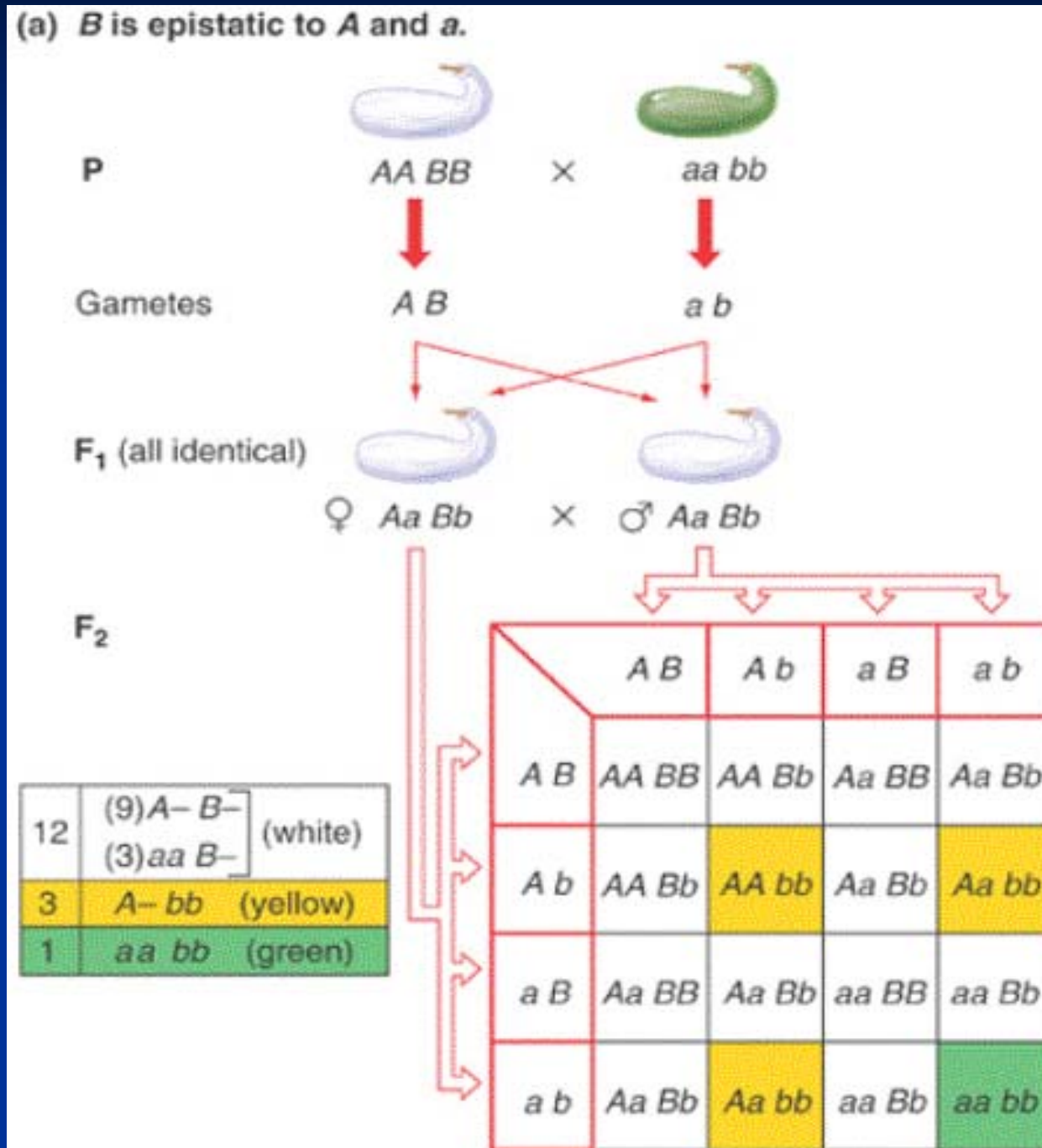


Fig. 3.14b

# 3-2 Dominant epistasis (I)

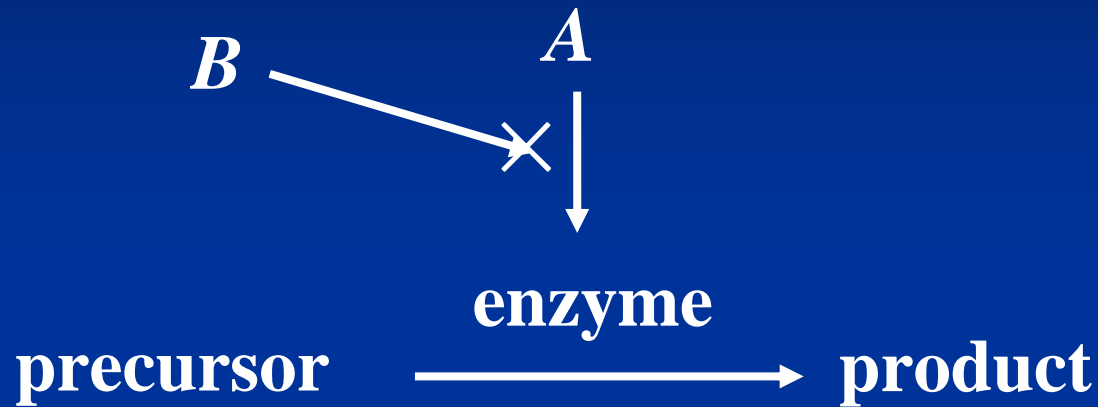


Summer squash

Fig. 3.15a

## 3-3 Dominant epistasis (II)

One gene inhibit the other gene's function

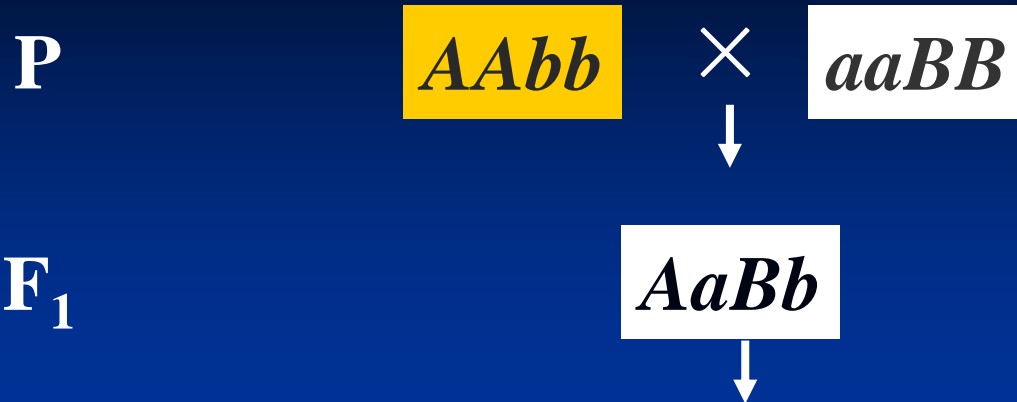


*B* gene suppress *A* gene's function.

Example: Primrose color

**A: yellow**

**B: inhibit A, white (colorless)**



Predicted F <sub>2</sub> ratio	Genotype	Phenotype	Final ratio
3/16	<i>A_bb</i>	yellow	3/16
9/16	<i>A_B_</i>	white	13/16
3/16	<i>aaB_</i>	white	
1/16	<i>aa bb</i>	white	



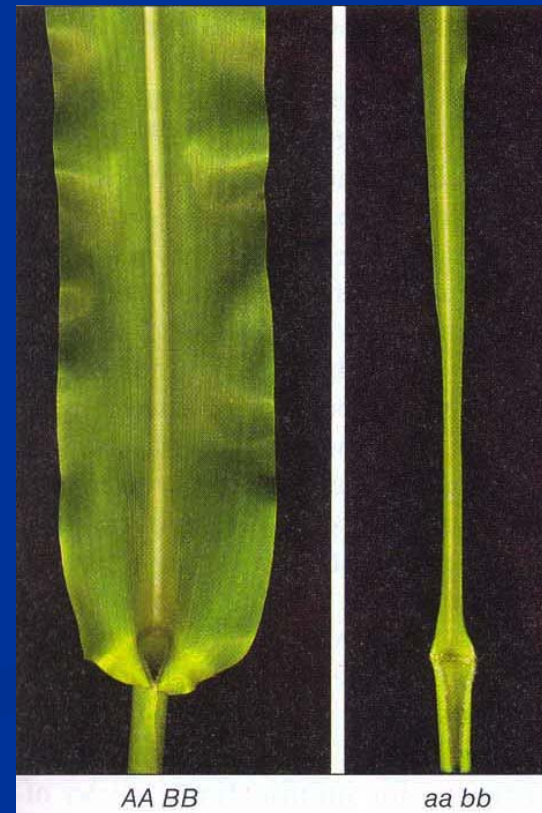
## 4. Two genes have the same phenotype

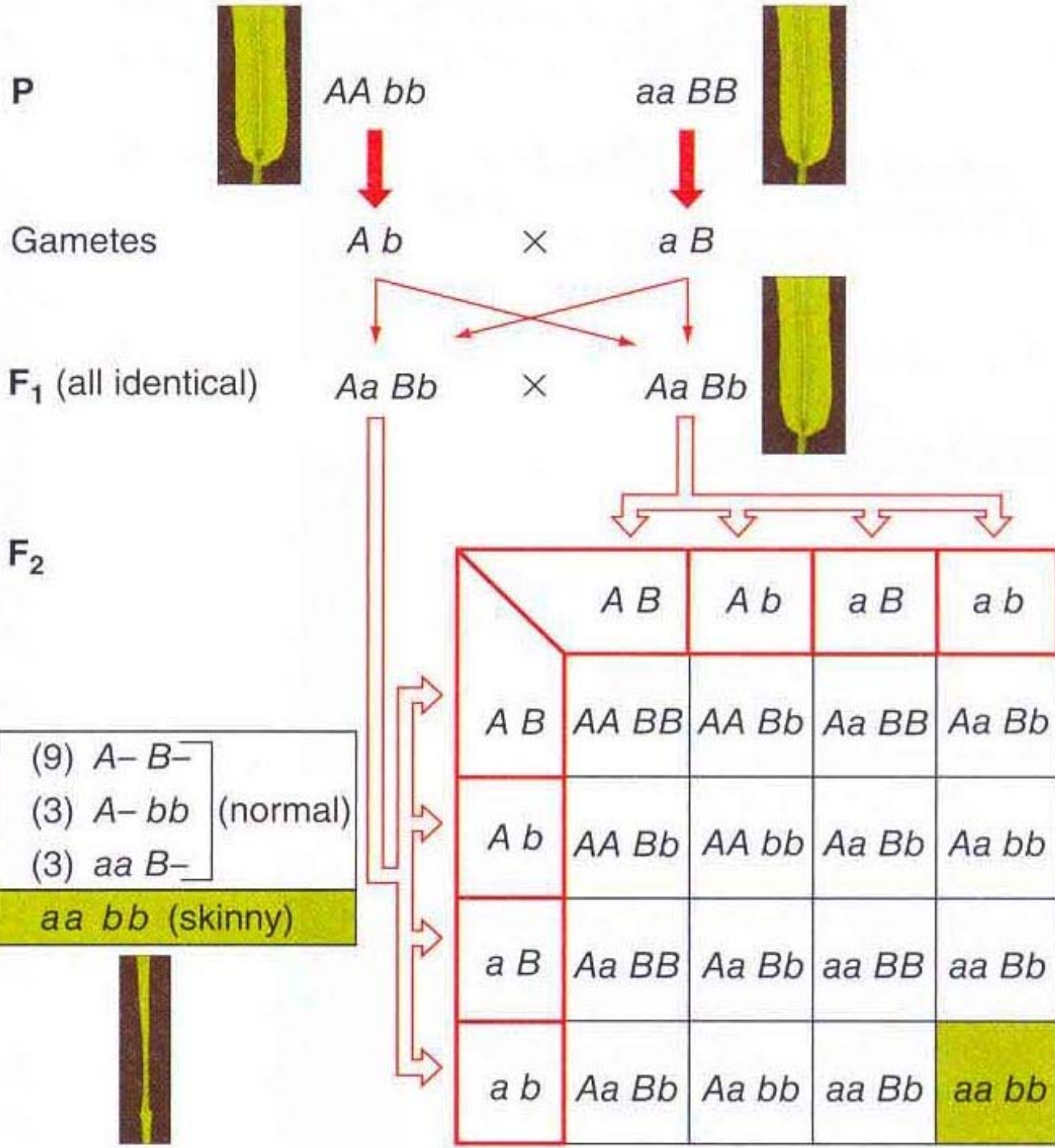
Precursor  $\xrightarrow{A \text{ or } B}$  Product

*A* and *B* gene have the same function, only one is enough.

Example:

The shape of maize leaves.





**P**      *AABB* × *aabb*

**F1**    ♀ *AaBb* × ♂ *AaBb*

**F2**

	<i>AB</i>	<i>Ab</i>	<i>aB</i>	<i>ab</i>
<i>AB</i>	<i>AABB</i>	<i>AABb</i>	<i>AaBB</i>	<i>AaBb</i>
<i>Ab</i>	<i>AABb</i>	<i>AAbb</i>	<i>AaBb</i>	<i>Aabb</i>
<i>aB</i>	<i>AbBB</i>	<i>AaBb</i>	<i>aaBB</i>	<i>aaBb</i>
<i>ab</i>	<i>AaBb</i>	<i>Aabb</i>	<i>aaBb</i>	<i>aabb</i>

9 *A\_ B\_*

3 *A\_ bb*

3 *aa B\_*

1 *aa bb*

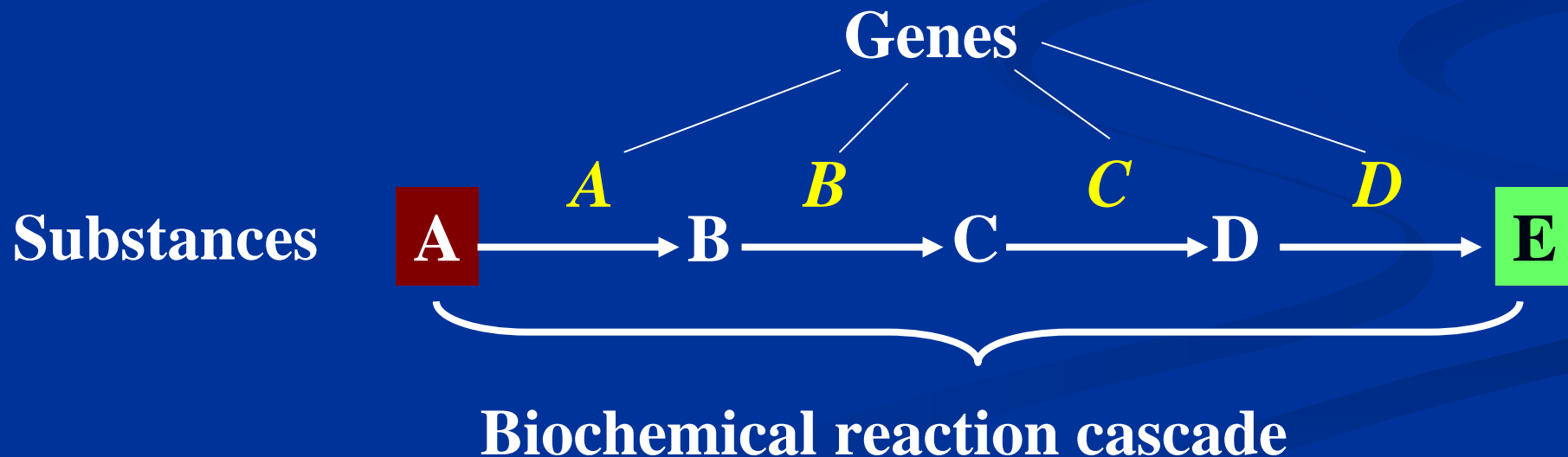


**TABLE 3.2**
**Summary of Gene Interactions**

Gene Interaction	Example	F <sub>2</sub> Genotypic Ratios from an F <sub>1</sub> Dihybrid Cross				F <sub>2</sub> Phenotypic Ratio
		A- B-	A- bb	aa B-	aa bb	
<b>None:</b> Four distinct F <sub>2</sub> phenotypes	Lentil: seed coat color (see Fig. 3.10a)	9	3	3	1	9:3:3:1
<b>Complementary:</b> One dominant allele of each of two genes is necessary to produce phenotype	Sweet pea: flower color (see Fig. 3.12b)	9	3	3	1	9:7
<b>Recessive epistasis:</b> Homozygous recessive of one gene masks both alleles of another gene	Labrador retriever: coat color (see Fig. 3.14b)	9	3	3	1	9:3:4
<b>Dominant epistasis I:</b> Dominant allele of one gene hides effects of both alleles of another gene	Summer squash: color (see Fig. 3.17a)	9	3	3	1	12:3:1
<b>Dominant epistasis II:</b> Dominant allele of one gene hides effects of dominant allele of other gene	Chicken feathers: color (see Fig. 3.18a)	9	3	3	1	13:3
<b>Redundancy:</b> Only one dominant allele of either of two genes is necessary to produce phenotype	Maize: leaf development (see Fig. 3.19b)	9	3	3	1	15:1

## 5. Several genes can contribute to the same trait

- **Heterogeneous traits** are traits determined by a number of different genes. A mutation at any one of these genes can give rise to the same phenotype.
  - Albinism in plants, animals and humans.
  - Deafness in humans is a heterogeneous trait. About 50 different genes have mutant alleles that can cause deafness.



- Difficult to determine which of many genes mutated in a person with a heterogeneous mutant phenotype.
- Complementation testing can determine if mutations arise from the same or different genes.

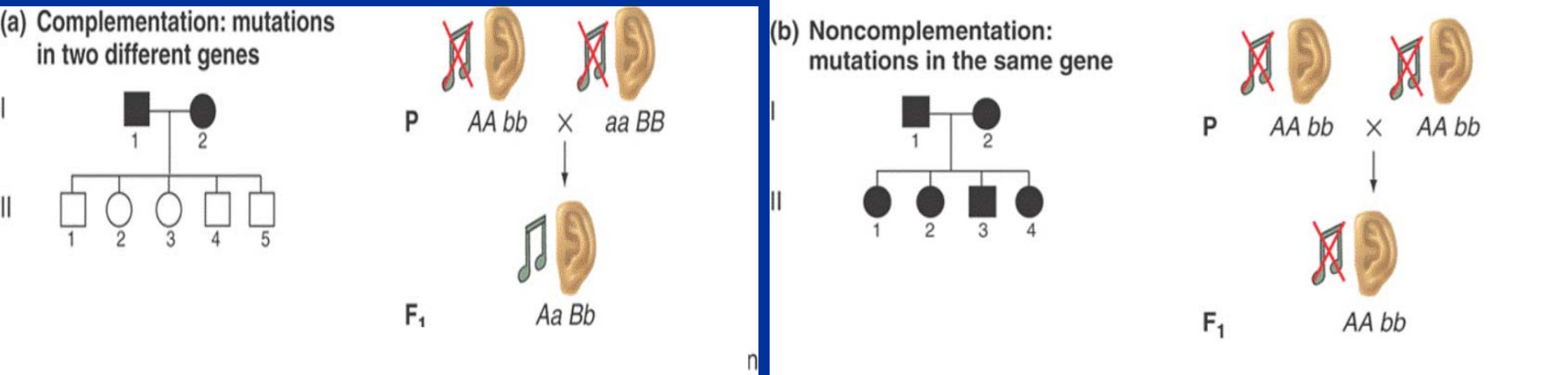


Fig. 3.16



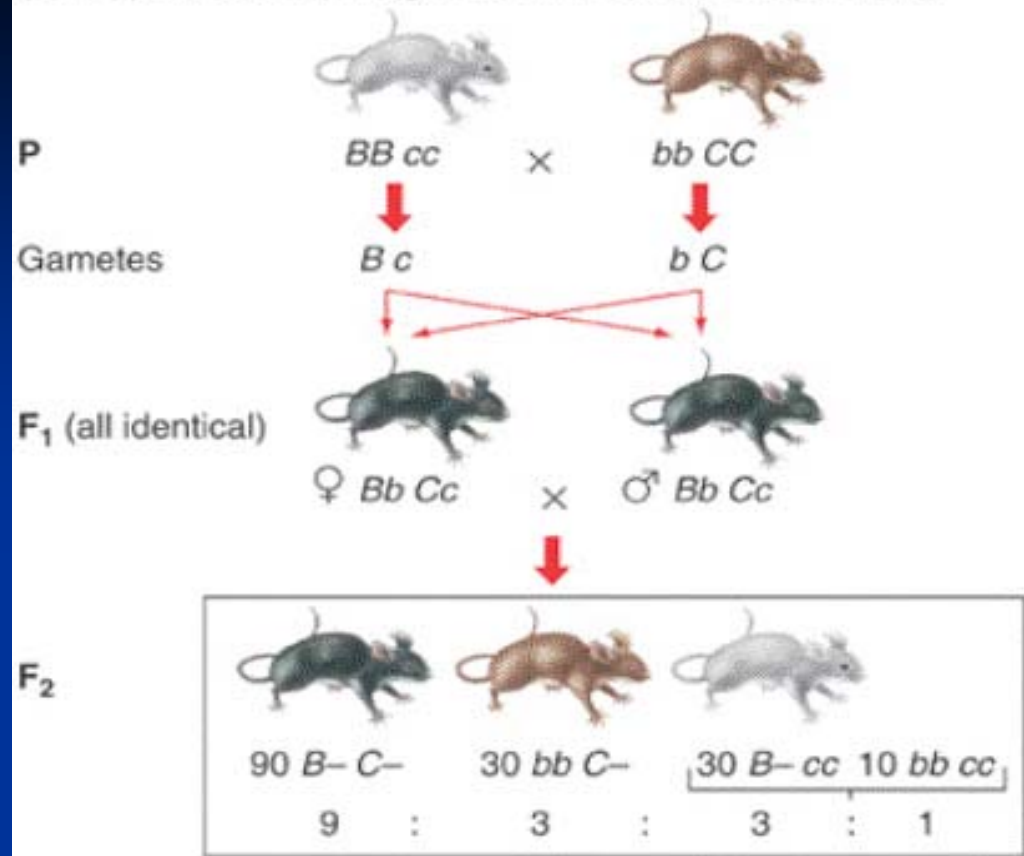
## **Breeding studies help determine how a trait is inherited**

- **How do we know if a trait is caused by one gene or two genes that interact?**
  - **Ratios such as 9:7 or 13:3 can indicate potential gene interaction.**
  - **Further breeding studies can confirm hypotheses.**



# Testing two gene and one gene hypothesis – example from mice

(a) Hypothesis 1 (two genes with recessive epistasis)



If two-gene hypothesis is correct:

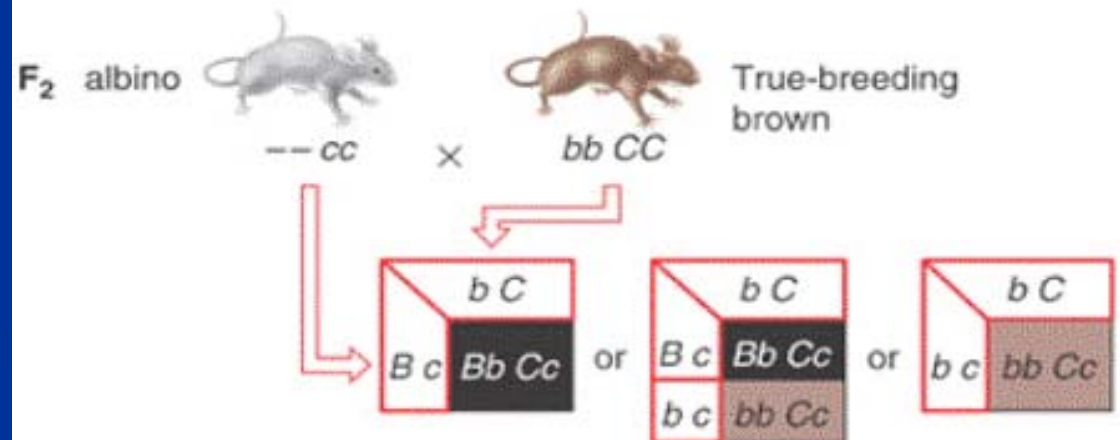
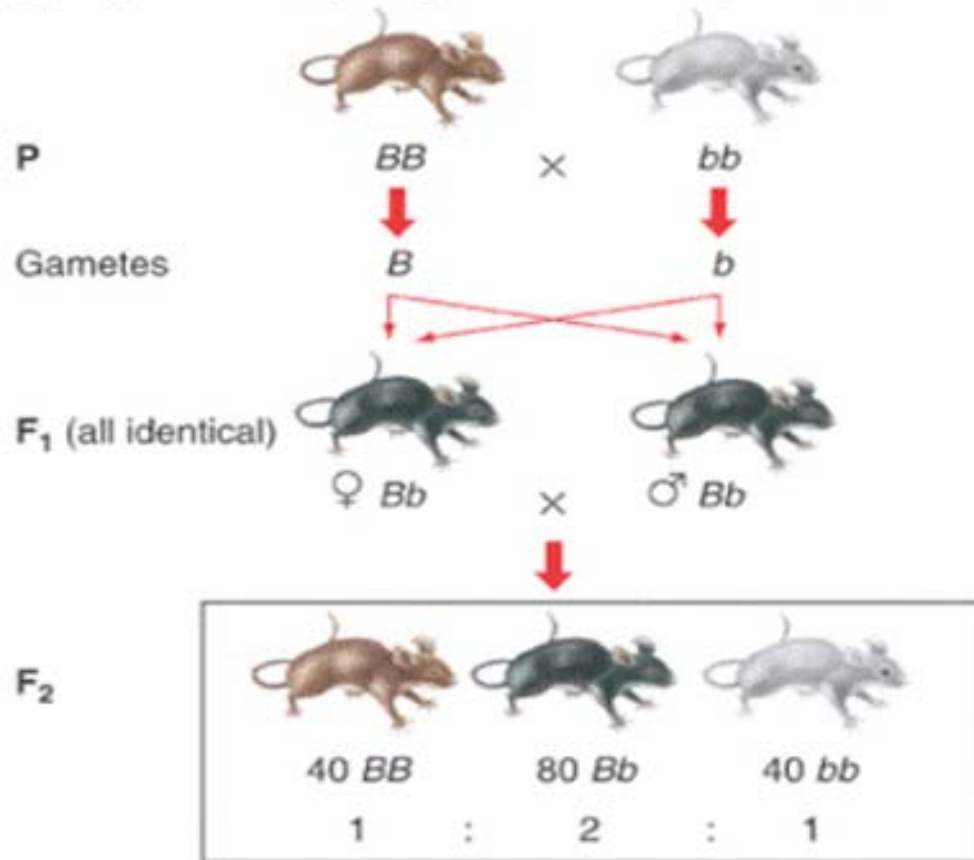


Fig. 3.18a

(b) Hypothesis 2 (one gene with incomplete dominance)



If one-gene hypothesis is correct:

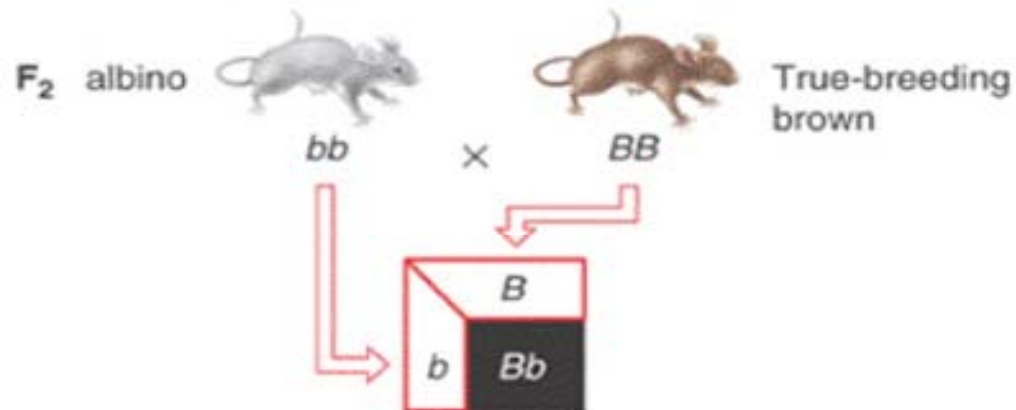


Fig. 3.18b

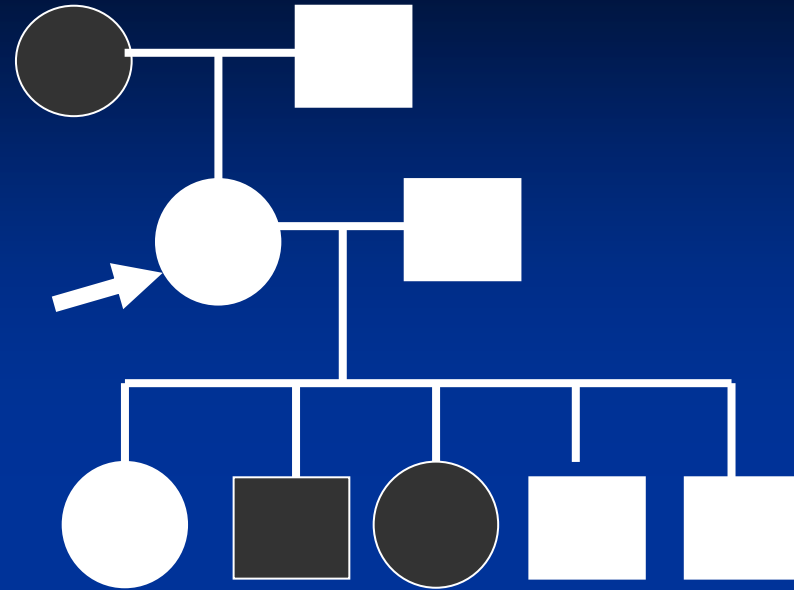
## **Summary of genetic variations on multifactorial traits**

- **Genes can interact to generate new phenotypes.**
- **The dominant alleles of two interacting genes can both be necessary for producing a phenotype.**
- **One gene's alleles can mask the effects of another gene's.**
- **Mutants alleles at one of two or more different genes can result in the same phenotype.**

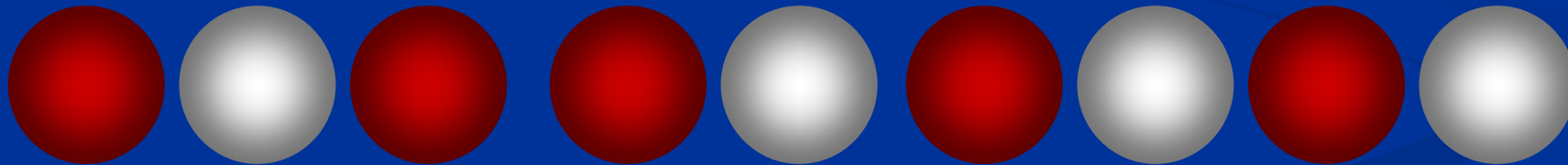
# The same genotype does not always produce the same phenotype

- Phenotype often depends on penetrance and expressivity.
  - **Penetrance (外显率)**: Percentage of individuals with a particular genotype that show the expected phenotype in a population.
    - Penetrance can be complete (100%) or incomplete (retinoblastoma 75%).
  - **Expressivity (表现度)**: Degree or intensity with which a particular genotype is expressed in a phenotype
    - Expressivity can be variable or unvarying.

**Incomplete penetrance**

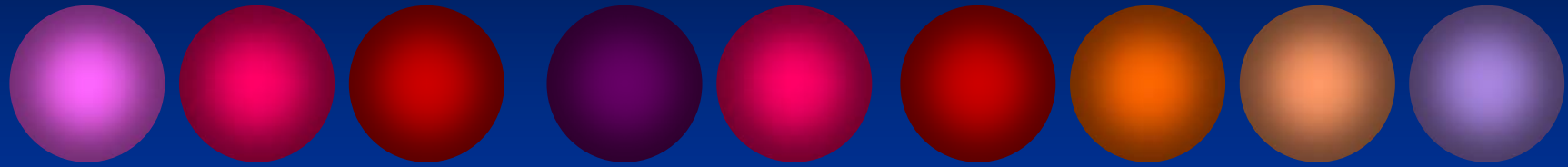


**Phenotypic expression**  
(each circle represents an individual)



**Variable penetrance**

## Variable expressivity



## Color spots in beagle



- **Modifier genes** have subtle, secondary effect on a phenotype from a major gene.
- **Environment can affect the phenotypic expression of a genotype.**
  - **Temperature**
    - Coat color in Siamese cats is lighter in its extremities.
    - **Conditional lethals**
      - *shibire* gene in fly is lethal at temperatures above 29°C.



# Temperature can affect coat color

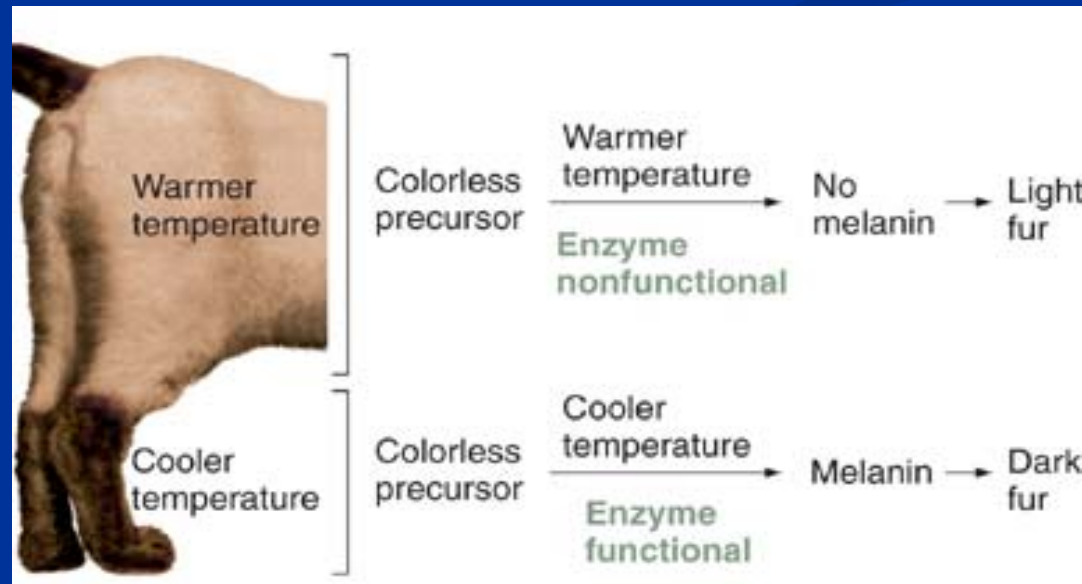
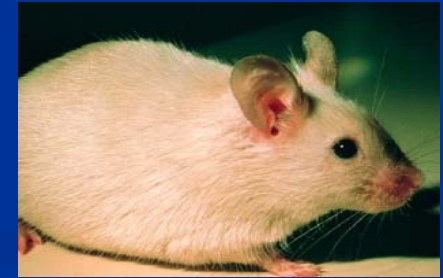
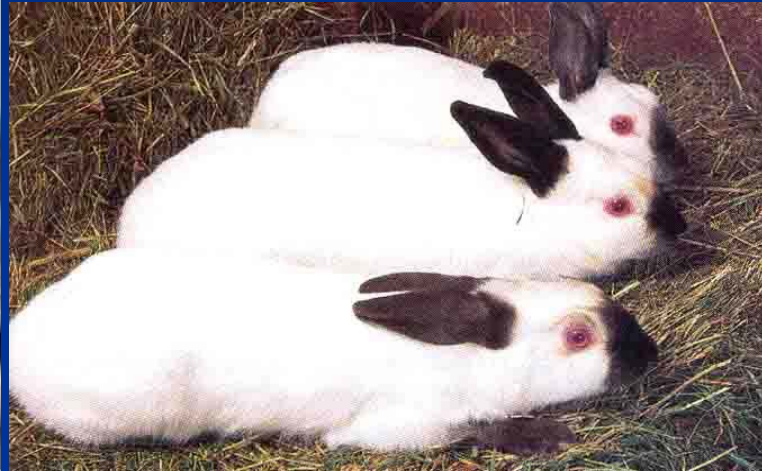


Fig. 3.20

- **Environment can affect the phenotypic expression of a genotype.**
  - **Temperature**
    - Coat color in Siamese cats is lighter in its extremities.
    - Conditional lethals
      - *shibire* gene in fly is lethal at temperatures above 29°C.
  - **Phenocopy (表型模拟):** Change in phenotype arising from environmental agents that mimics the effect of a mutation at a gene.
    - In early 1960s, the drug thalidomide disrupts limb development in normal fetuses.
    - Diet, exercise, smoking contribute to certain heart and lung diseases.

## Continuous variation can be explained by Mendelian analysis

- **Discontinuous traits:** Phenotype is clear cut without variation.
- **Continuous traits (连续性状):** Phenotype shows continuous variation.



# Height is a continuous trait in humans



Fig. 3.21a

# Skin color is a continuous trait

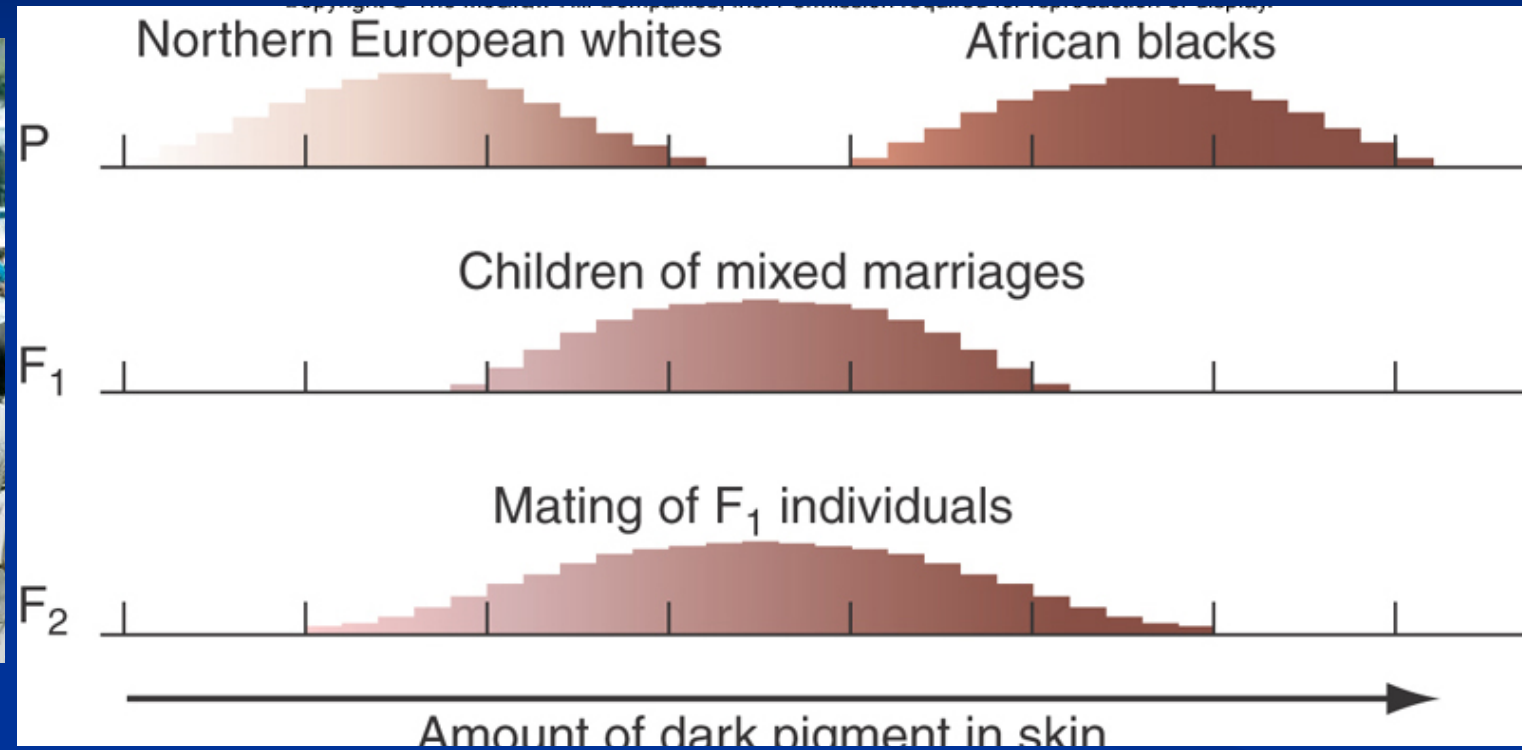


Fig. 3.21b

- **Continuous traits:** Phenotype shows continuous variation.
  - Continuous traits are also called **quantitative traits** by geneticists and are usually polygenic.
  - Continuous traits such as height in humans are determined by segregating alleles of many genes interacting with one another and the environment.
  - The more genes that contribute to a trait, the greater the number of possible phenotypic classes and the greater the similarity to continuous variation.