

BIO-373  
Genetics & Genomics

**Extension of  
Mendelian Genetics**

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# Why do we need to go beyond Mendelian genetics?

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# Why do we need to go beyond Mendelian genetics?

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**Many phenotypes cannot be explained by Mendel's laws**

→ Co-dominance, incomplete dominance, epistasis, multiple alleles, lethal alleles, genomic imprinting...

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- 1 Incomplete dominance
  - 2 Codominance
  - 3 Multiple alleles of a gene
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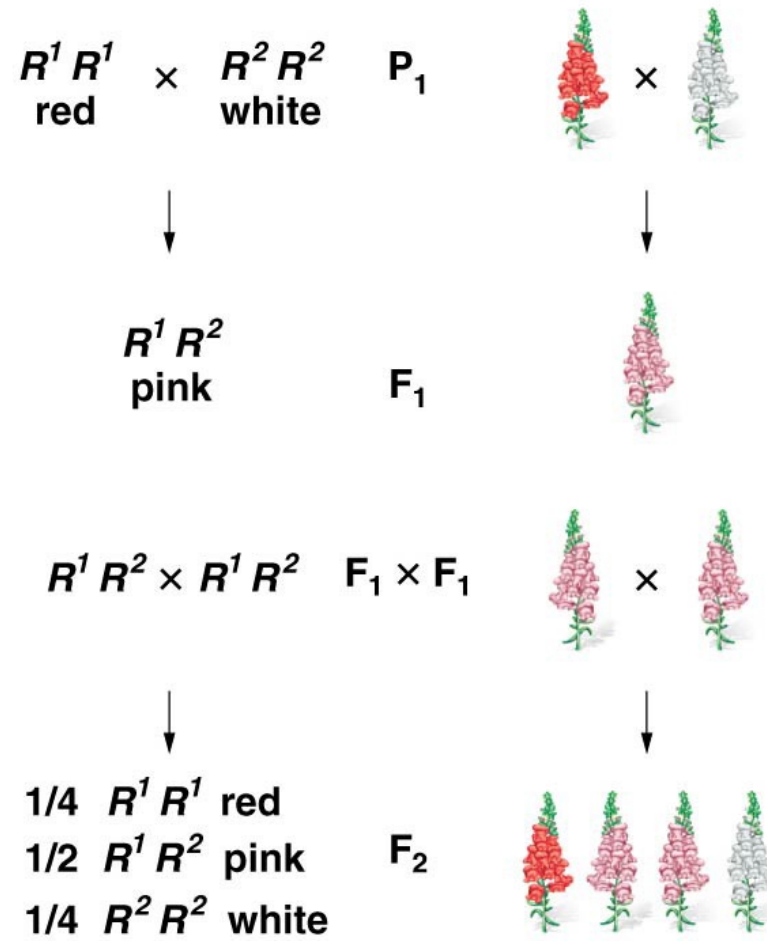
# **1. Incomplete (or partial) dominance**

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# Incomplete (or partial) dominance

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- Neither allele is dominant → intermediate phenotype
  - **Each genotype has its own phenotype**
  - Botanical example: snapdragon
  - Cross between red and white flowers
    - $F_1$  offspring: pink flowers
    - $F_2$  generation: 1/4 red, 1/2 pink, 1/4 white
    - **Phenotypic and genotypic ratios are the same**
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## **2. Codominance**

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# Codominance

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- No dominance or recessiveness
  - No incomplete or blending
  - Joint expression of both alleles in a heterozygote
  - Possible when 2 alleles have distinct phenotypic expression
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# Example of codominance in humans

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- **MN Blood Group**
- M and N are two distinct forms of a glycoprotein that can be expressed on red blood cells
- Alleles  $L^M$  and  $L^N$

## Genotype

$L^M L^M$

$L^M L^N$

$L^N L^N$

## Phenotype

M

MN

N

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## **3. Multiple alleles of a gene**

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# Multiple alleles

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- Mutations can modify a gene in various ways → 3 or more alleles of the same gene might exist
  - The resulting mode of inheritance is unique
  - This can only be studied in populations and not in single individuals (who always carry two alleles per genetic locus)
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# ABO blood groups

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- ABO blood group is controlled by a single gene (*ABO* on chromosome 9)
  - *ABO* is an enzyme that modifies antigens on the surface of red blood cells (RBCs)
  - *ABO* blood group is important for blood transfusion and organ transplantation
  - Three possible alleles of a single *ABO* gene (*I* for isoagglutinogen, another term for antigen)
    - *I<sup>A</sup>* → produces an enzyme that results in **A antigen** on RBCs
    - *I<sup>B</sup>* → produces an enzyme that results in **B antigen** on RBCs
    - *i* → produces an inactive enzyme and results in the **O blood group**
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Genotype	Antigen	Phenotype
$I^A I^A$	A	A
$I^A i$	A	
$I^B I^B$	B	B
$I^B i$	B	
$I^A I^B$	A, B	AB
$ii$	Neither	O

**$I^A$  and  $I^B$  are dominant to  $i$**

**$I^A$  and  $I^B$  are codominant to each other**

# ABO blood groups

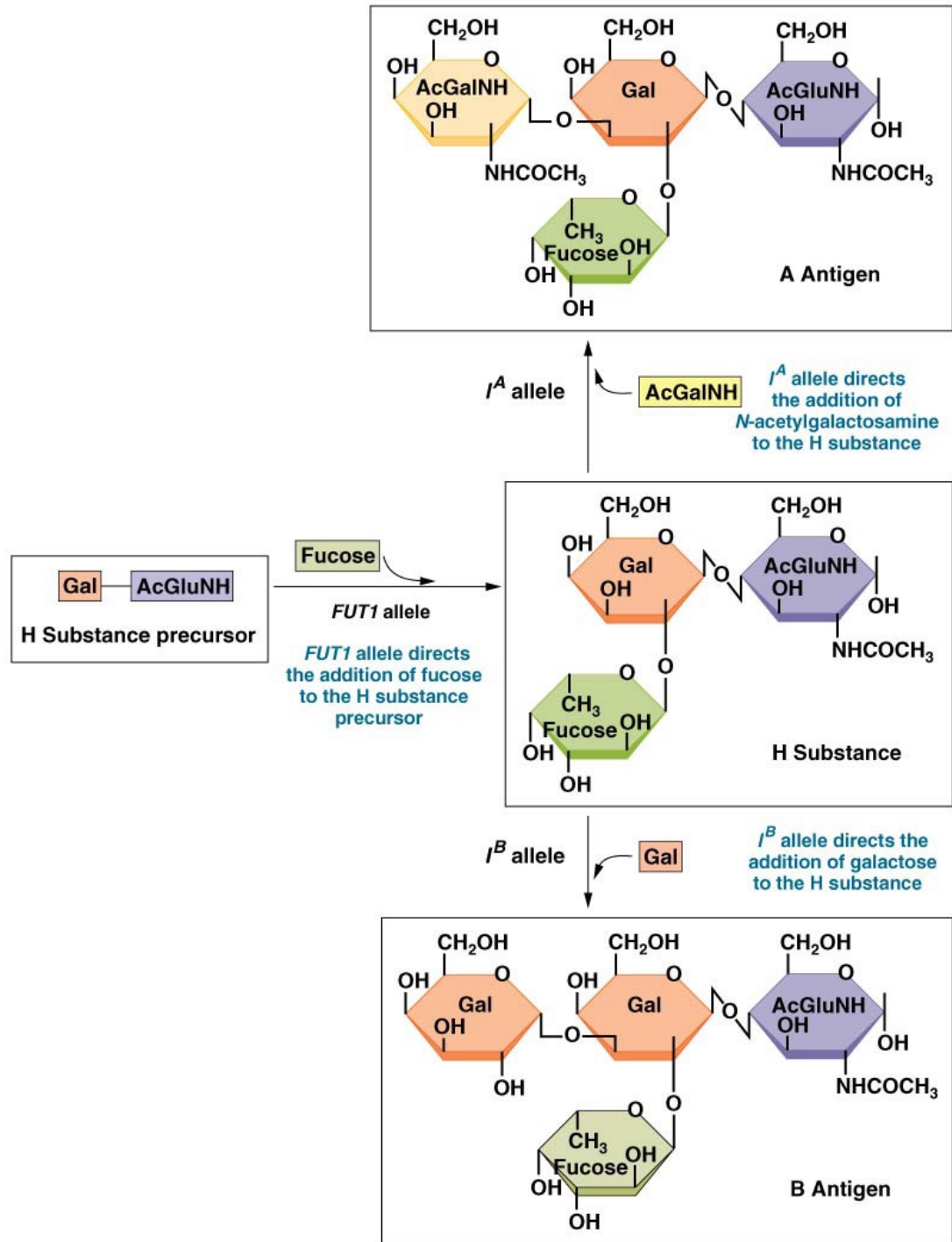
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- **A and B antigens**

- Carbohydrate groups (N-Acetylgalactosamine for A antigen or galactose for B antigen) bound to H substance on red blood cells

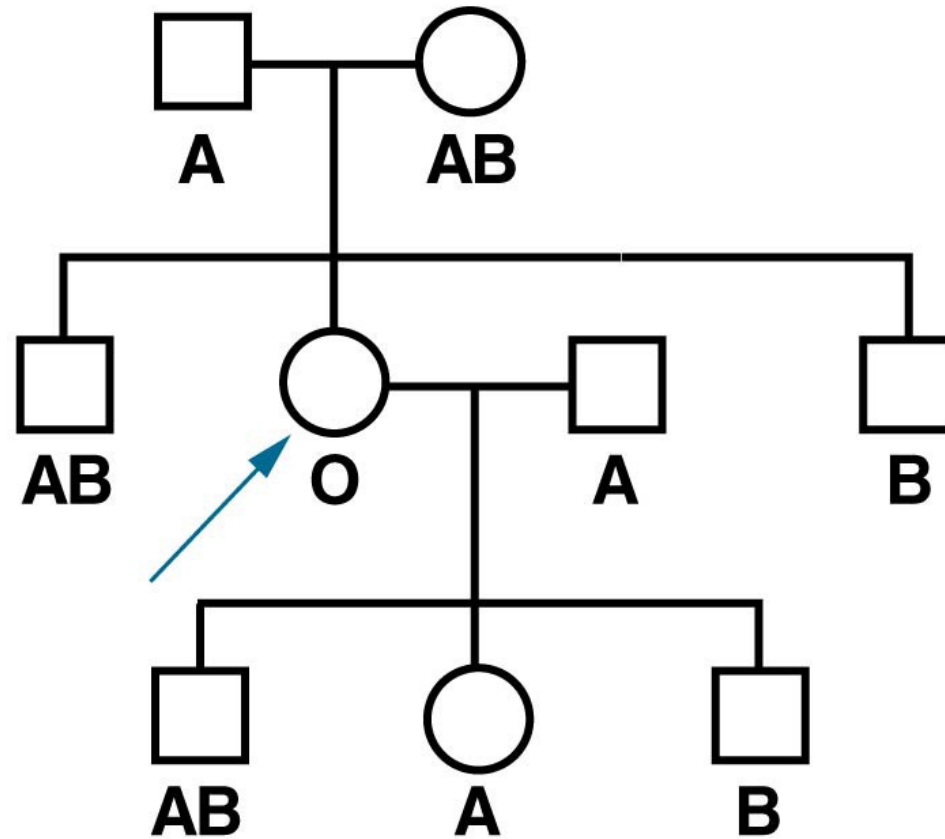
- **H antigen (or also termed H substance)**

- H antigen is the precursor to ABO antigens
  - Basic sugar layer at the surface of red blood cells
  - O blood types (*ii*) only have the H antigen (O stands for “Ohne” in German)
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# A more complex case

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# Bombay phenotype

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- Type O female, yet...
    - Mother has the AB antigen
    - Mother gave  $I^B$  allele to two other children
  
  - Female found to be homozygous for a **FUT1** (fucosyl transferase) mutation
    - Prevents her from producing H antigen
    - No base substrate to make A or B antigens
    - Results functionally in type O
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# Bombay phenotype

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- One gene (*FUT1*) is masking the phenotypic effect of another one (*ABO*)
  - Very rare allele
    - 1:10'000 in India
    - 1:100'000 in Europe
  - Individuals with Bombay phenotype can only receive blood transfusion from Bombay donors
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## **4. Lethal alleles and essential genes**

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# Essential genes and lethal alleles

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- Absolutely required for survival
  - May contain lethal alleles
  - **Dominant lethal allele:** the presence of a single copy is incompatible with life
  - **Recessive lethal allele:** tolerated in heterozygous state
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# Dominant lethal allele in humans

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- Huntington disease
    - Characterized by progressive degeneration of nervous system, dementia, and death
    - Due to allele *H* (abnormal number of CAG trinucleotide repeat in *HTT* gene)
    - Autosomal dominant = Hh genotype
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# Recessive lethal allele in mice

- **Agouti (*Asip*) gene:** responsible for coat color
  - Paracrine antagonist of MC1R on melanocytes
  - Agouti allele *A* (*eumelanin*, black)
  - Mutant *yellow* allele  $A^Y$  (phenomelanin, yellow)

Crosses				
(A) agouti	×	agouti	→	all agouti
(B) yellow	×	yellow	→	2/3 yellow: 1/3 agouti
(C) agouti	×	yellow	→	1/2 yellow: 1/2 agouti

**Cross A**

P<sub>1</sub> **AA** × **AA**  
agouti agouti



F<sub>1</sub> **AA**  
agouti



all agouti  
(All survive)

agouti mouse



**Cross B**

P<sub>1</sub> **AA<sup>Y</sup>** × **AA<sup>Y</sup>**  
yellow yellow



F<sub>1</sub> **AA** **AA<sup>Y</sup>**  
agouti yellow  
**A<sup>Y</sup>A** **A<sup>Y</sup>A<sup>Y</sup>**  
yellow lethal



2/3 yellow  
1/3 agouti  
(Survivors)

yellow mouse



**Cross C**

P<sub>1</sub> **AA** × **AA<sup>Y</sup>**  
agouti yellow



F<sub>1</sub> **AA** **AA<sup>Y</sup>**  
agouti yellow



1/2 agouti  
1/2 yellow  
(All survive)

# Recessive lethal allele in mice

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- **Mutant allele ( $A^Y$ )**
    - Behaves dominantly to normal allele to control coat color
    - Behaves as homozygous recessive lethal allele
  - Genotype  $A^Y A^Y$  does not survive
  - Why is the  $A^Y$  allele lethal?
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# Recessive lethal allele in mice

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- **Mutant allele ( $A^Y$ )**
    - Behaves dominantly to normal allele to control coat color
    - Behaves as homozygous recessive lethal allele
  - Genotype  $A^Y A^Y$  does not survive
  - Why is the  $A^Y$  allele lethal?
    - The  $A^Y$  mutation is a large deletion that removes the *Raly* gene yet retains the *Raly* promoter
    - The *Raly* promoter drives expression of *Asip* (yellow coat color and other effects like obesity)
    - $A^Y A^Y$  is lethal due to complete loss of the *Raly* gene (essential for embryonic development)
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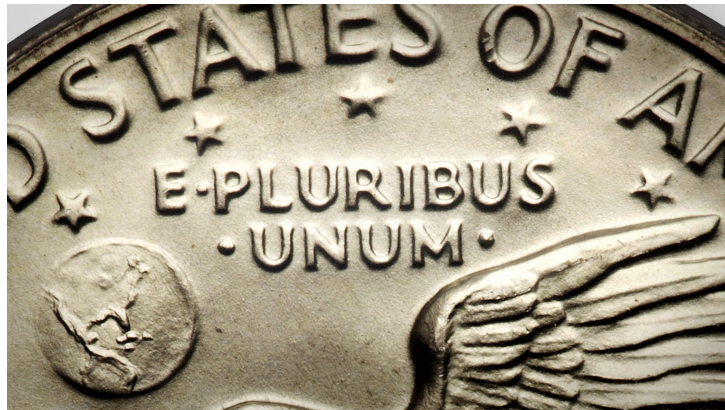
## **5. Impact of multiple genes on a phenotype**

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# E pluribus unum (“out of many, one”)

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- Most phenotype result from the influence of multiple genes
- **Genic interaction:** several genes influence a trait; doesn't imply direct interaction
- **Epistatic interaction:** direct interaction between alleles of different genes



# Example of genic interaction

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- **Hereditary deafness**

- Many genes are involved in ear formation
  - These genes interact to produce a common phenotype
  - Many mutations can interrupt development and lead to hereditary deafness
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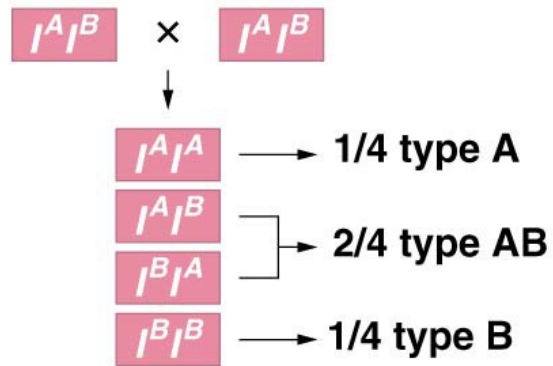
# Example of epistatic interaction

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- **Back to Bombay...**
    - First locus masks expression of second locus
    - Homozygous mutation in *FUT1* gene masks expression of  $I^A$  and  $I^B$  alleles
    - A and B antigen forms only when individual has at least one wild-type allele
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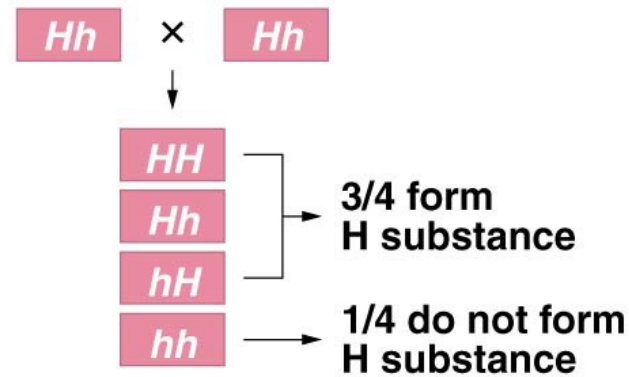
$$I^A I^B Hh \times I^A I^B Hh$$

Consideration of blood types



Genotypes      Phenotypes

Consideration of H substance



Genotypes      Phenotypes

### Consideration of both gene pairs together

Of all offspring	Of all offspring	Final probabilities
↓	↓	↓
1/4 type A	3/4 form H substance	→ 3/16 type A
	1/4 do not form H substance	→ 1/16 type O
2/4 type AB	3/4 form H substance	→ 6/16 type AB
	1/4 do not form H substance	→ 2/16 type O
1/4 type B	3/4 form H substance	→ 3/16 type B
	1/4 do not form H substance	→ 1/16 type O

**Final phenotypic ratio = 3/16 A: 6/16 AB: 3/16 B: 4/16 O**

# Impact of epistasis on phenotypic ratio

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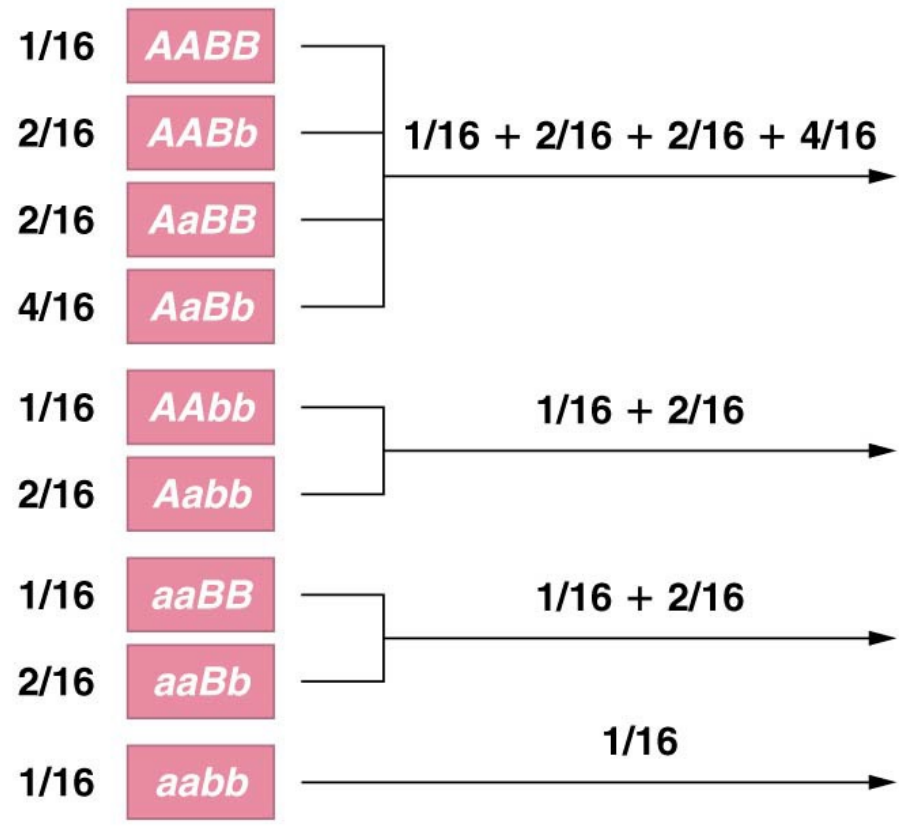
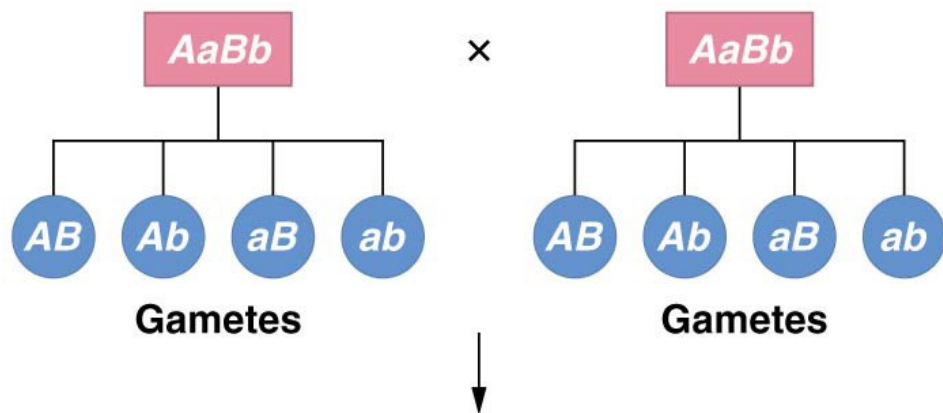
- Epistatic interactions modify the phenotypic ratios expected under Mendel's transmission laws.
  - Dihybrid cross: 9:3:3:1
  - Effect possible on one or more of four phenotypic categories
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# Digenic epistatic ratios:

## gene interactions that modify mendelian ratios

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- **No interaction (simple dominance): 9:3:3:1**
  - **Dominant epistasis: 12:3:1**  
A dominant allele (A-) at one locus masks the other locus.
  - **Recessive epistasis: 9:3:4**  
Homozygous recessive (aa) at one locus masks the other locus.
  - **Duplicate recessive epistasis: 9:7**  
Need at least one dominant allele at both loci (A-B-) for the main phenotype.
  - **Duplicate interaction: 9:6:1**  
Both dominants together give one phenotype, either single dominant gives a second; double recessive gives a third.
  - **Duplicate dominant epistasis: 15:1**  
A dominant allele at either loci gives the same phenotype.
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Dihybrid ratio	Modified ratios				
$9/16 A-B-$	9/16	12/16	7/16	9/16	9/16
$3/16 A-bb$	3/16			6/16	15/16
$3/16 aaB-$	4/16	3/16	1/16	1/16	
$1/16 aabb$		1/16		1/16	1/16

9:3:3:1

# Dominant epistasis

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- Dominant allele at one locus masks an allele at second locus
- Summer squash fruit colour
  - Determined by 2 loci, A and B
  - Dominant allele A = **white** fruit
    - Regardless of second loci allele
  - Absence of A allele (*aa*)
    - Genotypes *BB*, *Bb* = **yellow** fruit
    - Genotype *bb* = **green** fruit



# Dominant epistasis

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$$F_1: AaBb \times AaBb$$

↓

F <sub>2</sub> Ratio	Genotype	Phenotype	Final Phenotypic Ratio
9/16	<i>A-B-</i>	white	12/16 white
3/16	<i>A-bb</i>	white	
3/16	<i>aaB-</i>	yellow	3/16 yellow
1/16	<i>aabb</i>	green	1/16 green

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# Recessive epistasis

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- Homozygous recessive allele at one locus masks an allele at second locus
  - Mouse coat color (again...):
    - *A* allele (dominant): agouti phenotype
    - *B* allele: black pigment
    - *bb* genotype: no black pigment, even if *A* or *a* alleles present → mouse is albino
  - *bb* genotype masks the expression of the *A* allele: **recessive epistasis**
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# Recessive epistasis

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$F_1: AaBb \times AaBb$

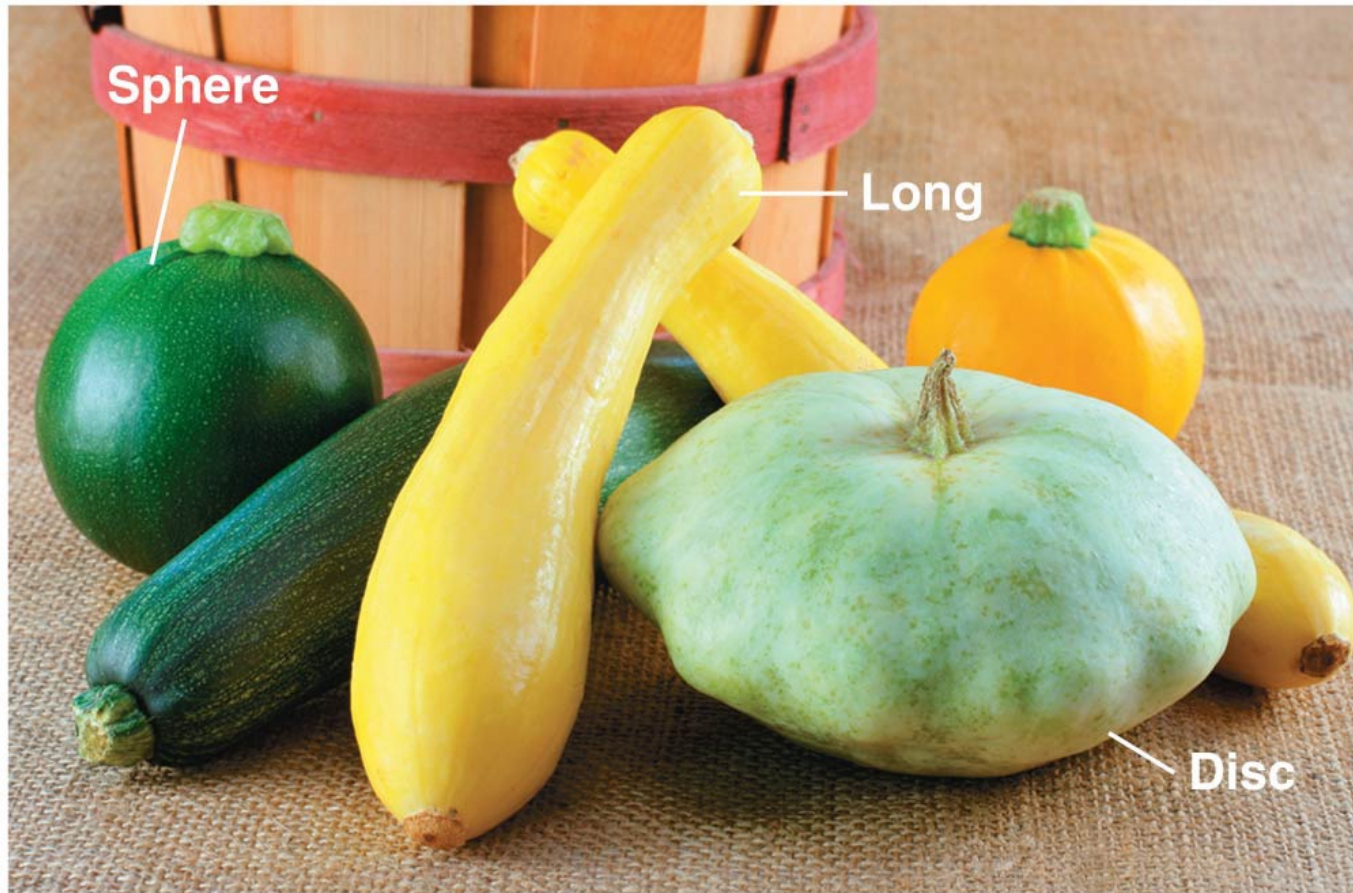


<b>F<sub>2</sub> Ratio</b>	<b>Genotype</b>	<b>Phenotype</b>	<b>Final Phenotypic Ratio</b>
9/16	$A-B-$	agouti	9/16 agouti
3/16	$A-bb$	albino	
3/16	$aaB-$	black	3/16 black
1/16	$aabb$	albino	4/16 albino

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# Duplicate interaction: epistasis resulting in a novel phenotype

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# Duplicate interaction: epistasis resulting in a novel phenotype

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- Two loci contribute to the same trait and dominant alleles have an additive effect
  - Shape of summer squash: **disc-shaped** fruit ( $AABB$ ) crossed with **long** fruit ( $aabb$ )
  - $F_1$ : all **disc-shaped** fruit
  - $F_2$ : includes the two parental phenotypes and a new **spherical** variant
    - Dominant alleles at 2 loci → disc-shaped
    - Dominant allele at 1 locus → spherical
    - Only recessive alleles → long
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# Duplicate interaction: epistasis resulting in a novel phenotypes

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$$F_1: AaBb \times AaBb$$

disc      disc

↓

F <sub>2</sub> Ratio	Genotype	Phenotype	Final Phenotypic Ratio
9/16	<i>A-B-</i>	disc	9/16 disc
3/16	<i>A-bb</i>	sphere	6/16 sphere
3/16	<i>aaB-</i>	sphere	
1/16	<i>aabb</i>	long	1/16 long

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## **6. Pleiotropy**

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# Pleiotropy

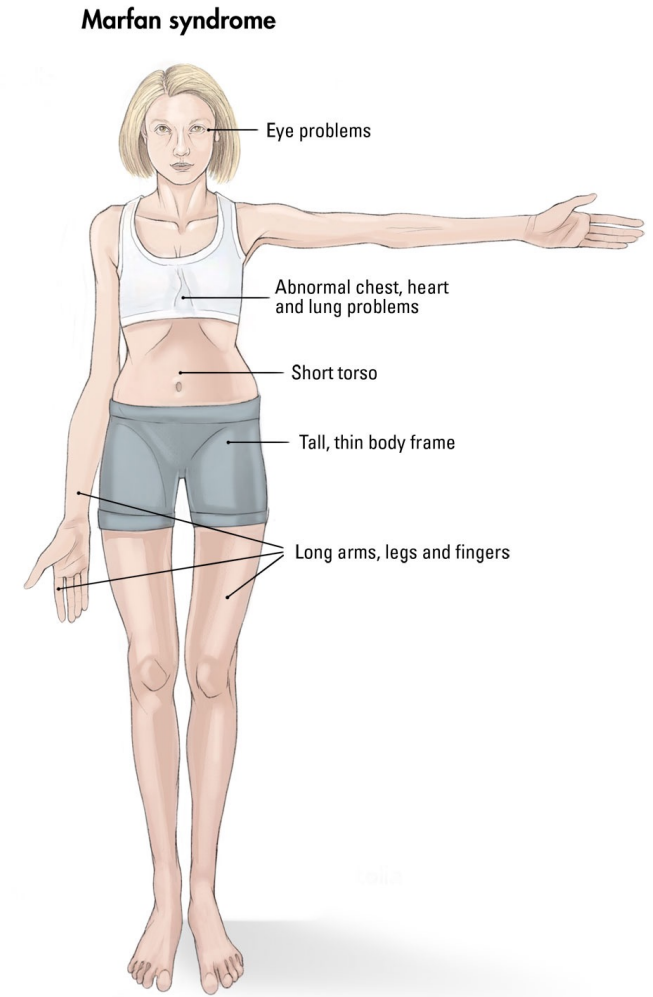
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- We just saw that several genes can influence a single phenotype
  - The opposite situation, where a single gene has multiple phenotypic effects, is called **pleiotropy**
  - Extremely frequent
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# Example of pleiotropy in humans

## ■ Marfan syndrome

- Autosomal dominant allele in fibrillin gene, encoding a protein important for connective tissues
- Multiple phenotypic consequences:
  - Lens dislocation
  - Aortic aneurism
  - Extension of long bones
  - ...



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## **7. Gene x environment interactions**

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# Phenotype = direct expression of genotype?

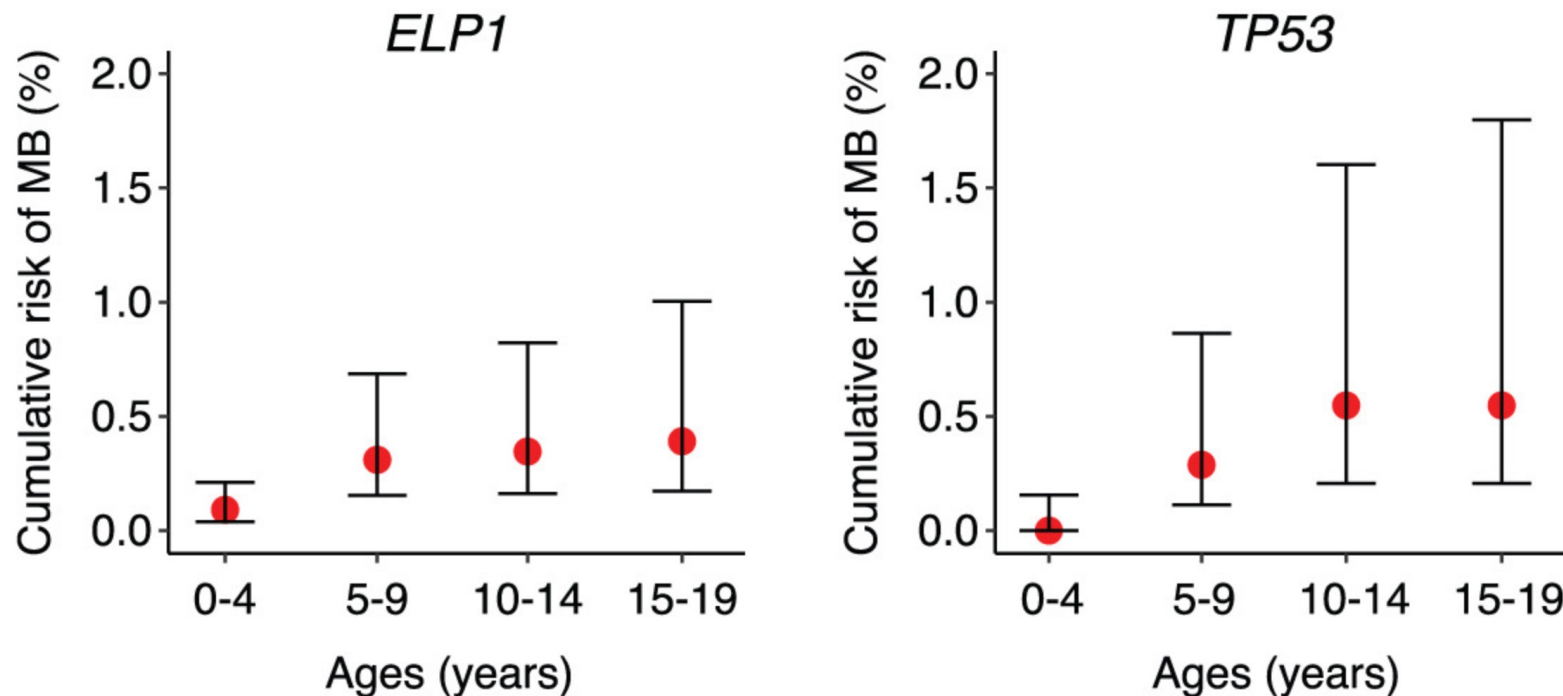
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- **A phenotype usually results from multiple influences**
    - Core gene
    - Other genotypes (genic and epistatic influences)
    - Environment
  - Often difficult to disentangle the respective influences of genes and environment ⇔ long-standing debate between “innate” and “acquired”
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# Penetrance and expressivity

## ■ Penetrance

- Percentage of expression of the mutant genotype in a population
- Incomplete, age-dependent, and incomplete, age-dependent penetrance
- Rare mutations in *ELP1* or *TP53* predispose to medulloblastoma, a rare childhood brain tumor: incomplete, age-dependent penetrance



# Penetrance and expressivity

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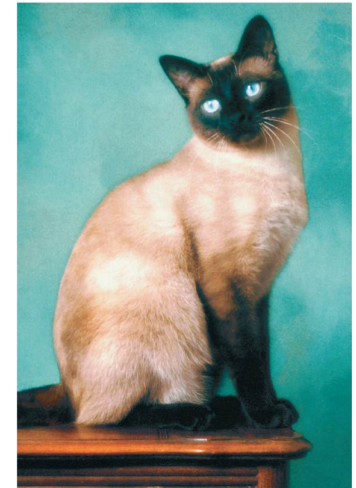
- **Expressivity**

- Range of expression (degree of severity) of mutant phenotype (eg, mild vs severe)
  - Result of genetic background differences and/or environmental effects
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# Causes of variable expressivity

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- **Temperature effects**
  - Evening primrose
    - Red flowers at 23°C
    - White flowers at 18°C
  - Siamese cats and Himalayan rabbits
    - Darker fur on cooler areas of body (tail, feet, ears)
    - Enzymes responsible for pigment formation lose catalytic function at higher temperature



# Causes of variable expressivity

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- **Temperature-sensitive mutations**
    - Known in viruses, bacteria, fungi, and *Drosophila*
    - Mutant allele expresses mutant phenotype at one temperature, wild-type phenotype at another
    - Broadly used in viral genetics: temperature-sensitive mutations are easily induced and isolated in viruses
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# Causes of variable expressivity

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## ■ Nutritional effects

- Some alleles prevent the synthesis of essential nutrients in microbes, they are called **auxotrophs**
  - Phenotype expression can be modulated by diet, which is very useful for *in vitro* studies
  - In humans, diseases are caused by alleles that stops the metabolization of some substances:
    - **Phenylketonuria**: loss of enzyme to metabolize amino acid phenylalanine, severe problems unless low-Phe diet
    - **Galactosemia, lactose intolerance, ...**
-