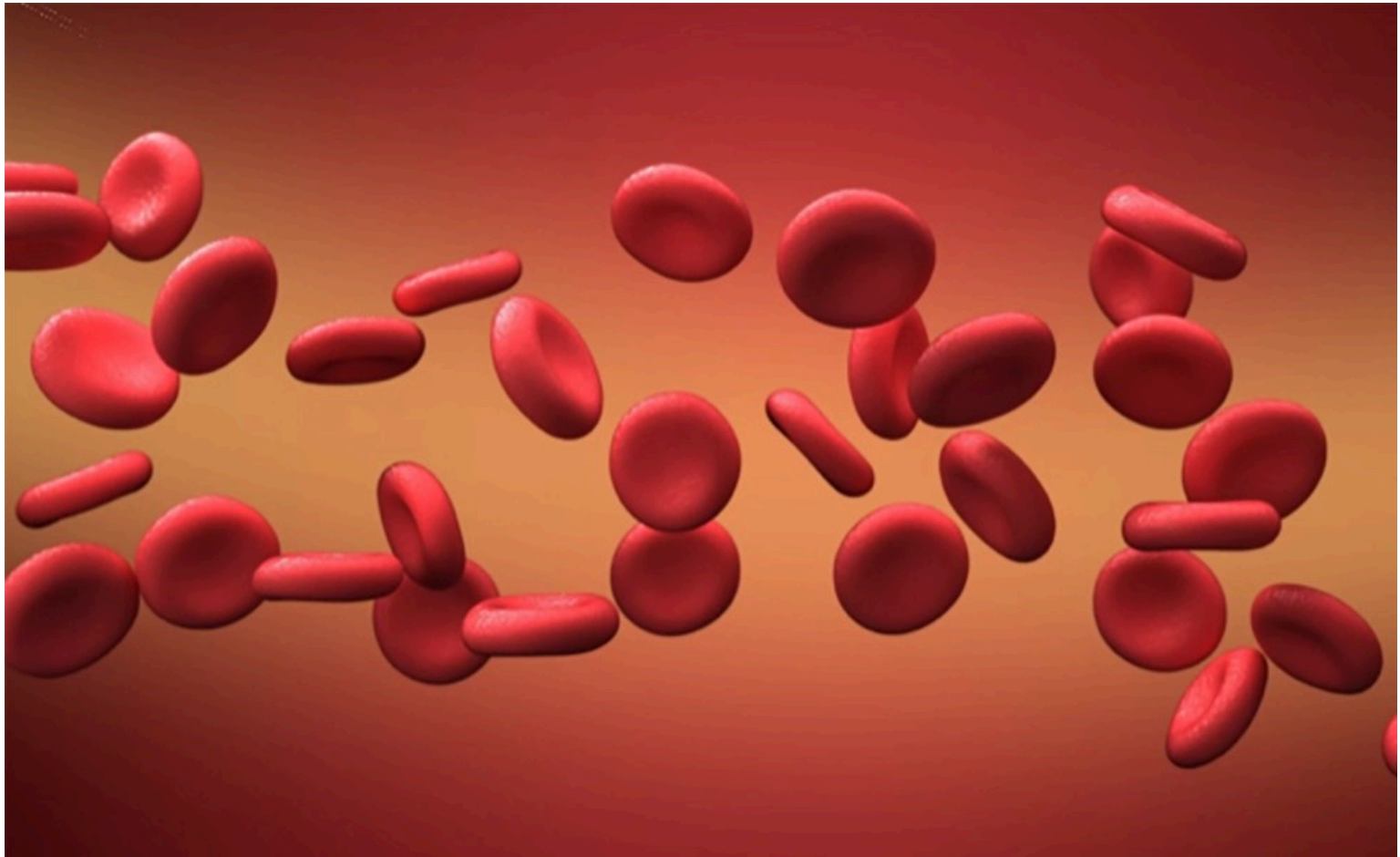


BIOL2107, Fall '23

Lecture 13



# Extensions to Mendelian Inheritance

Incomplete dominance

Codominance

Multiple Alleles

Epistasis

Several Genes

Lethal Alleles

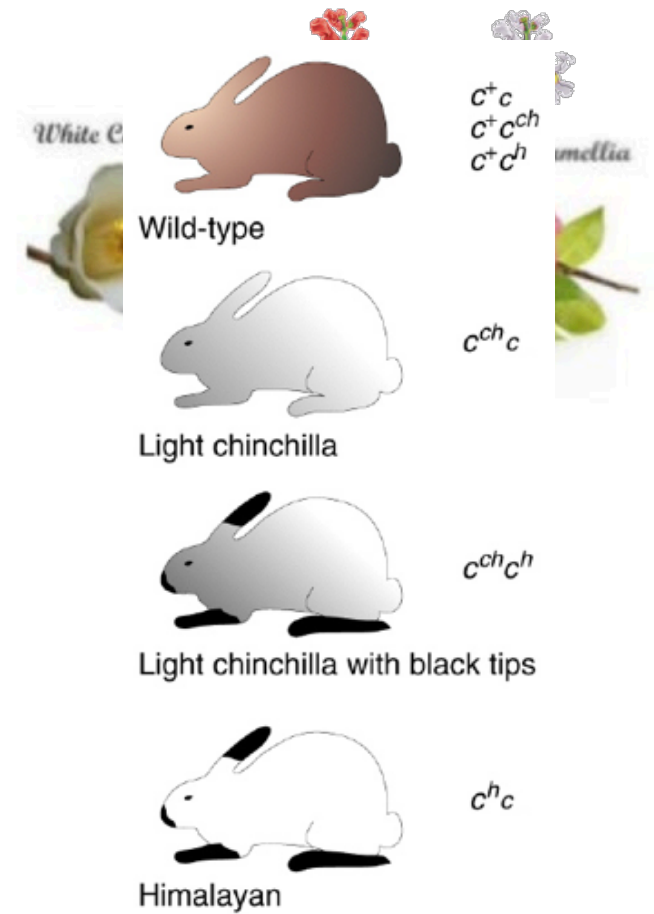
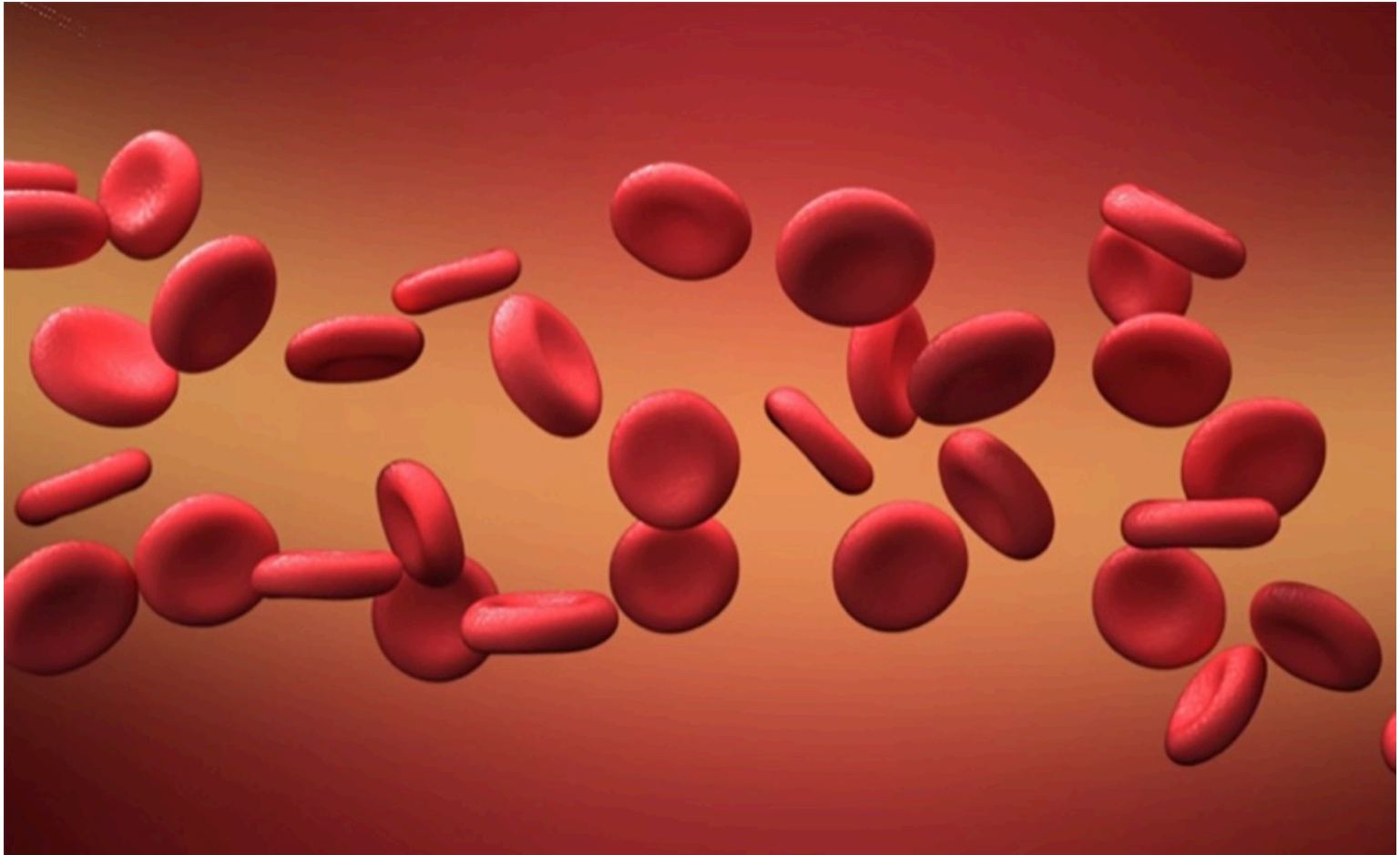


Figure 4.4 Phenotypes of different combinations of *c* alleles in rabbits. The alleles form a series, with the wild-type allele,  $c^+$ , dominant over all the other alleles and the null allele,  $c$  (albino), recessive to all the other alleles; one hypomorphic allele,  $c^{ch}$  (chinchilla), is partially dominant over the other,  $c^h$  (Himalayan). Copyright 2000 John Wiley and Sons, Inc.

# Multiple Alleles





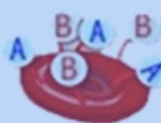




**An example of “co-dominant” alleles  
in humans**

## **The ABO Blood Group System**

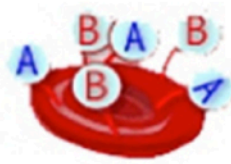


Antigens: molecules, usually on the outside of a cell, that provoke an immune response

## Potential Donors

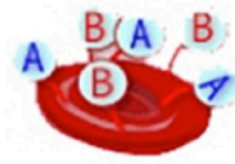
| Blood Type | Antibodies Produced   |  |  |  |  |
|------------|---|--|---|---|---|
| A          |    | +  | -   | -   | +   |
| B          |    | -  | +   | -   | +   |
| AB         | None  | +  | +   | +   | +   |
| O          |  | -  | -   | -   | +   |

## The ABO Blood Group System



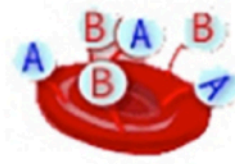
| Blood Type | Genotypes          | ABO Enzymes Present | RBC Antigens Present | Serum Antibodies |
|------------|--------------------|---------------------|----------------------|------------------|
| "A"        | $I^A I^A, I^A I^O$ | "H", "A"            | A, H                 | anti-B           |
| "B"        | $I^B I^B, I^B I^O$ | "H", "B"            | B, H                 | anti-A           |
| "AB"       | $I^A I^B$          | "H", "A", "B"       | A, B, H              | none             |
| "O"        | $I^O I^O$          | "H"                 | H                    | anti-A, anti-B   |

## The ABO Blood Group System



| Blood Type | Genotypes          | ABO Enzymes Present | RBC Antigens Present | Serum Antibodies |
|------------|--------------------|---------------------|----------------------|------------------|
| "A"        | $I^A I^A, I^A I^O$ | "H", "A"            | A, H                 | anti-B           |
| "B"        | $I^B I^B, I^B I^O$ | "H", "B"            | B, H                 | anti-A           |
| "AB"       | $I^A I^B$          | "H", "A", "B"       | A, B, H              | none             |
| "O"        | $I^O I^O$          | "H"                 | H                    | anti-A, anti-B   |

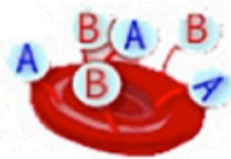
## The ABO Blood Group System



| Blood Type | Genotypes          | ABO Enzymes Present | RBC Antigens Present | Serum Antibodies |
|------------|--------------------|---------------------|----------------------|------------------|
| "A"        | $I^A I^A, I^A I^O$ | "H", "A"            | A, H                 | anti-B           |
| "B"        | $I^B I^B, I^B I^O$ | "H", "B"            | B, H                 | anti-A           |
| "AB"       | $I^A I^B$          | "H", "A", "B"       | A, B, H              | none             |
| "O"        | $I^O I^O$          | "H"                 | H                    | anti-A, anti-B   |



## The ABO Blood Group System



| Blood Type | Genotypes          | ABO Enzymes Present | RBC Antigens Present | Serum Antibodies |
|------------|--------------------|---------------------|----------------------|------------------|
| "A"        | $I^A I^A, I^A I^O$ | "H", "A"            | A, H                 | anti-B           |
| "B"        | $I^B I^B, I^B I^O$ | "H", "B"            | B, H                 | anti-A           |
| "AB"       | $I^A I^B$          | "H", "A", "B"       | A, B, H              | none             |
| "O"        | $I^O I^O$          | "H"                 | H                    | anti-A, anti-B   |

# **Extensions to Mendelian Inheritance**

**Incomplete dominance**

**Codominance**

**Multiple Alleles**

**Epistasis**

**Several Interactive genes**

**Lethal Alleles**



(A)

## Epistasis

Two Genes  
E and B

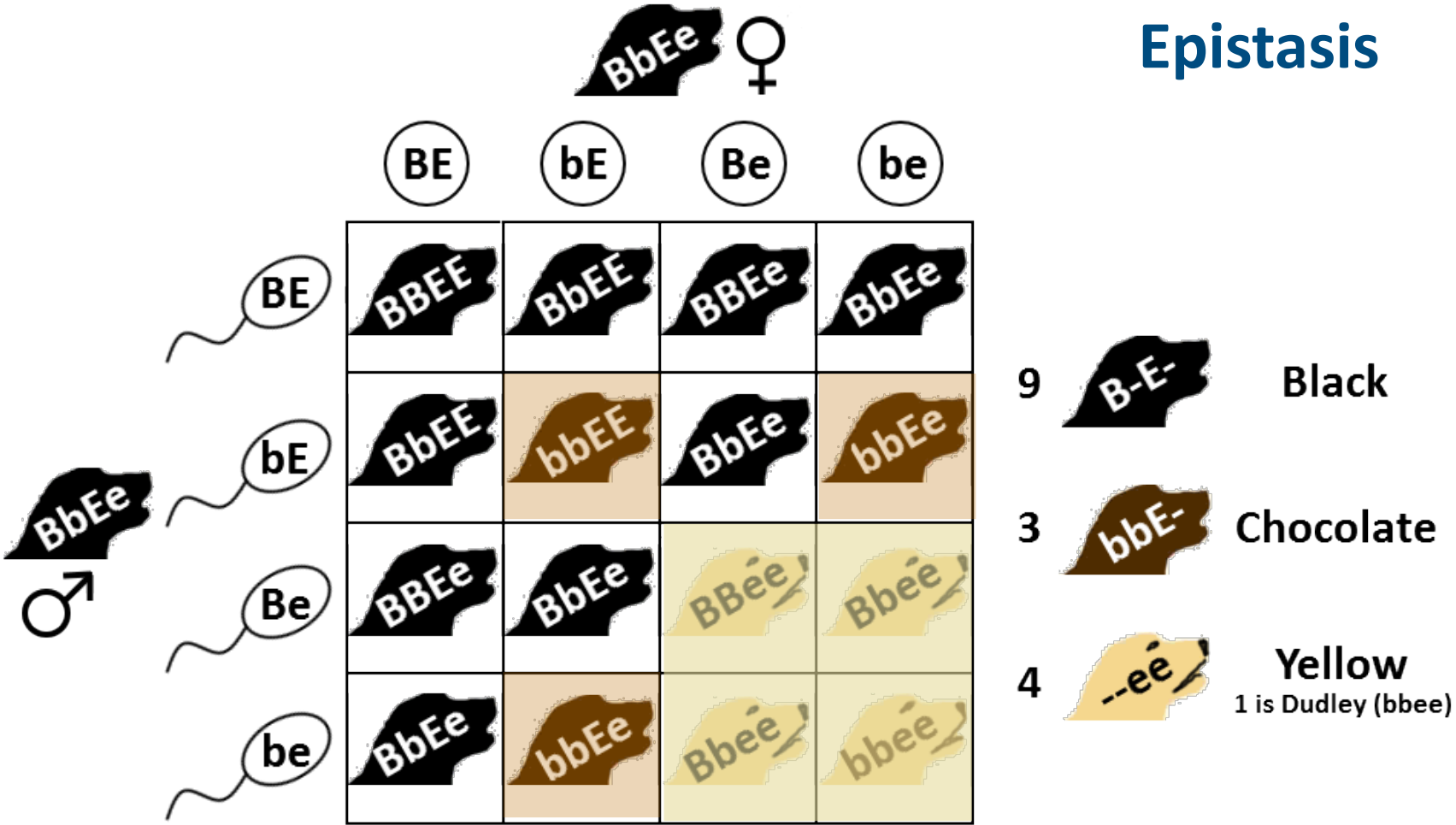


(B)



(C)

# Epistasis



# **Extensions to Mendelian Inheritance**

**Incomplete dominance**

**Codominance**

**Multiple Alleles**

**Epistasis**

**Several Interactive genes**

**Lethal Alleles**

# Several Interactive genes



Agouti... rodents

## Several Interactive genes



# Several Interactive genes



A Parental cross between a **AA, BB** and a **aa, bb**, results in **Aa, Bb** heterozygote.

The **A allele** determines a banded pattern, called **agouti**.

The recessive **a** allele results in **unbanded hairs**.

The genotypes **AA** or **Aa** are, therefore, **agouti**.

The genotypes **BB** or **Bb** result in fur colour that is solidly **black**, whereas **bb** denotes brown fur colour ..



# Several Interactive genes



A 3rd gene at the **C locus**, which is an entirely different locus from either **A** or **B**-determines if any colouration occurs at all.

The genotypes **CC** and **Cc** allow colour to show through, whereas the double recessive **cc** is **albino**, which does not allow any colour to show through, as the **cc genotype** blocks ALL pigment production -note the eyes are now pink.

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb** (where the **C** gene is present as **CC** or **Cc** and does not interfere) would be:

**9 agouti fur**, **3 cinnamon fur (brown, agouti)**, **3 black fur** and **1 brown fur**.

The corresponding **genotypes** are:

**9 agouti fur** (1 BB, AA + 2 Bb, AA + 4 Bb, Aa): **3 cinnamon** (1 bb, AA + 2 bb, Aa):

**3 black fur** (1 BB, aa + 2 Bb, aa) and **1 brown fur** (1 bb, aa).

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb** (where the **C** gene is present as **CC** or **Cc** and does not interfere) would be:

**9 agouti fur**, **3 cinnamon fur (brown, agouti)**, **3 black fur** and **1 brown fur**.

The corresponding **genotypes** are:

**9 agouti fur** (1 BB, AA + 2 Bb, AA + 4 Bb, Aa) **3 cinnamon** (1 bb, AA + 2 bb, Aa):

**3 black fur** (1 BB, aa + 2 Bb, aa) and **1 brown fur** (1 bb, aa).

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb** (where the **C** gene is present as **CC** or **Cc** and does not interfere) would be:

**9 agouti fur**, **3 cinnamon fur (brown, agouti)**, **3 black fur** and **1 brown fur**.

The corresponding **genotypes** are:

**9 agouti fur** (1 BB, AA + 2 Bb, AA + 4 Bb, Aa): **3 cinnamon** (1 bb, AA + 2 bb, Aa):

**3 black fur** (1 BB, aa + 2 Bb, aa) and **1 brown fur** (1 bb, aa).

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb** (where the **C** gene is present as **CC** or **Cc** and does not interfere) would be:

**9 agouti fur**, **3 cinnamon fur (brown, agouti)**, **3 black fur** and **1 brown fur**.

The corresponding **genotypes** are:

**9 agouti fur** (1 BB, AA + 2 Bb, AA + 4 Bb, Aa): **3 cinnamon** (1 bb, AA + 2 bb, Aa):

**3 black fur** (1 BB, aa + 2 Bb, aa) and 1 brown fur (1 bb, aa).

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb** (where the **C** gene is present as **CC** or **Cc** and does not interfere) would be:

**9 agouti fur**, **3 cinnamon fur (brown, agouti)**, **3 black fur** and **1 brown fur**.

The corresponding **genotypes** are:

**9 agouti fur** (1 BB, AA + 2 Bb, AA + 4 Bb, Aa): **3 cinnamon** (1 bb, AA + 2 bb, Aa):

**3 black fur** (1 BB, aa + 2 Bb, aa) and **1 brown fur** (1 bb, aa).

# Several Interactive genes



An **F2 phenotypic ratio** of an initial parental cross between an **AA, BB** and an **aa, bb**

Note that when the **C gene** is present as **cc** **ALL** mice are **ALBINO** -pink eyes

Which provides another example of **EPISTASIS**

# **Extensions to Mendelian Inheritance**

**Incomplete dominance**

**Codominance**

**Multiple Alleles**

**Epistasis**

**Several Interactive genes**

**Lethal Alleles**



# Lethal Alleles



Agouti... rodents

# Lethal Alleles

$Y'$  is often designated  $A^y$  which is dominant over  $y$  or the regular agouti allele

mono hybrid cross of  $Y'y$  heterozygotes...  
2 (+0): 1 ratio

eg. of **pleiotropy**

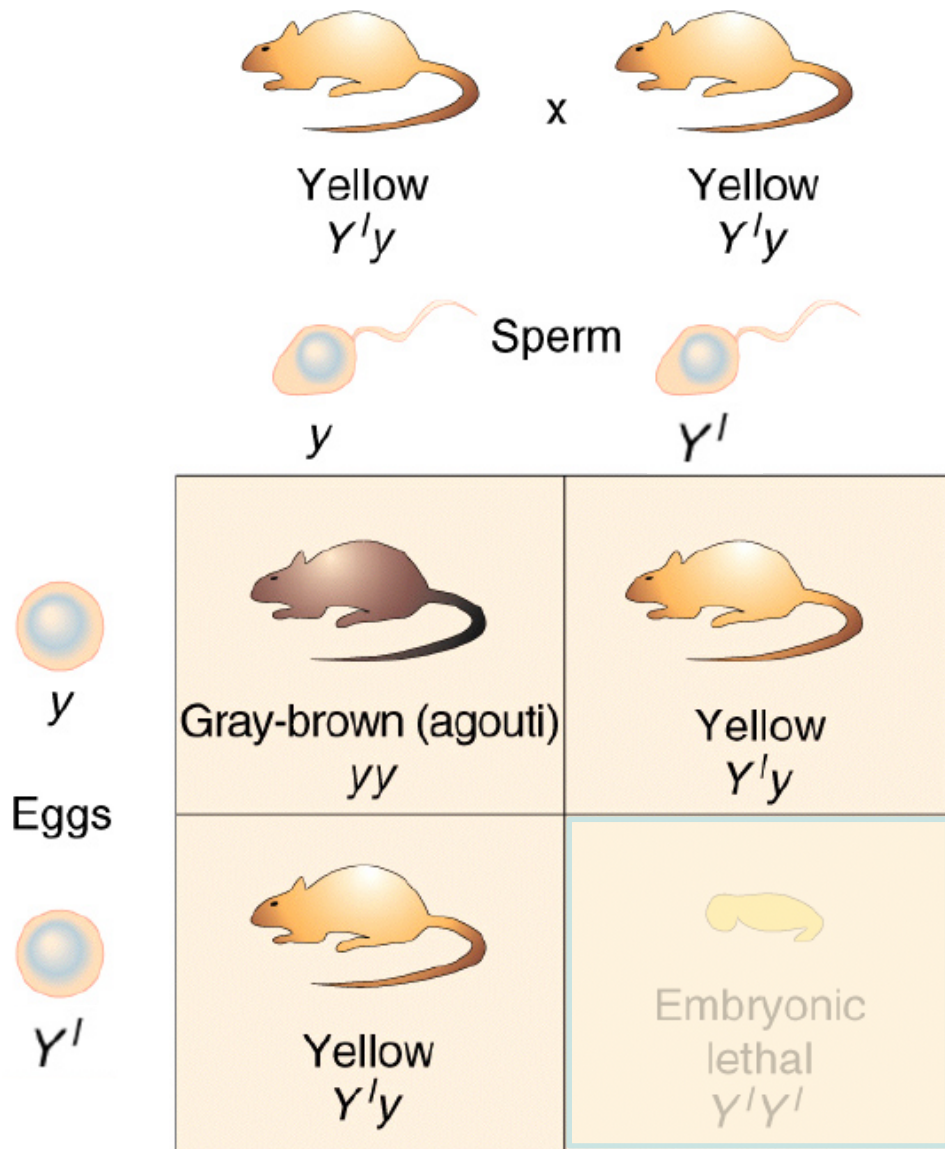


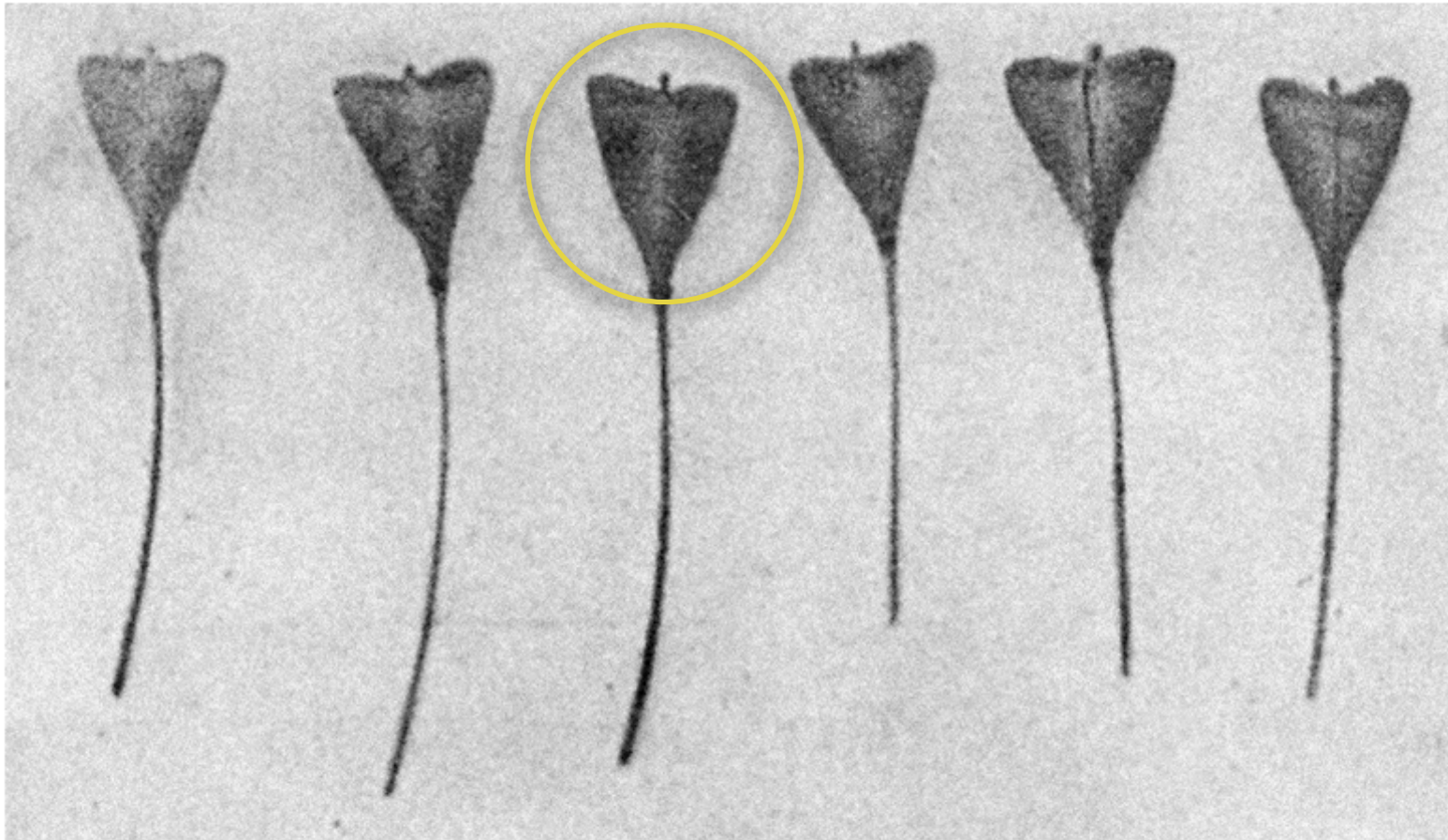
Figure 4.7  $Y'$ , the yellow-lethal mutation in mice: a dominant visible that is also a recessive lethal. A cross between carriers of this mutation produces yellow heterozygotes and gray-brown (agouti) homozygotes in a ratio of 2:1. The yellow homozygotes die as embryos.

## Duplicate Genes

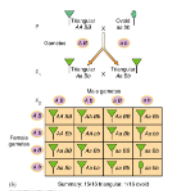


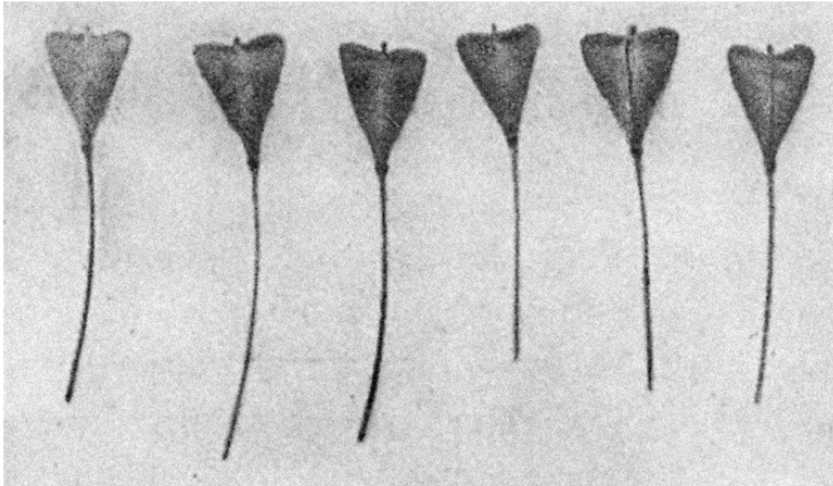
Shepherd's Purse

# Duplicate Genes



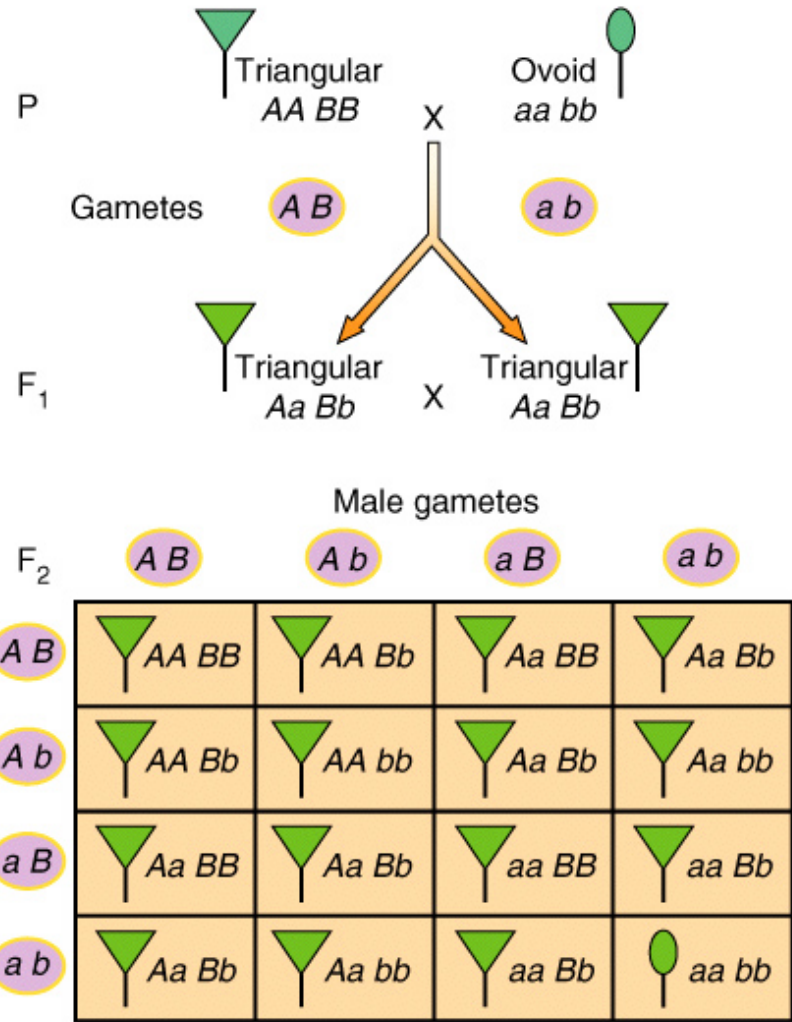
Courtesy New York Public Library





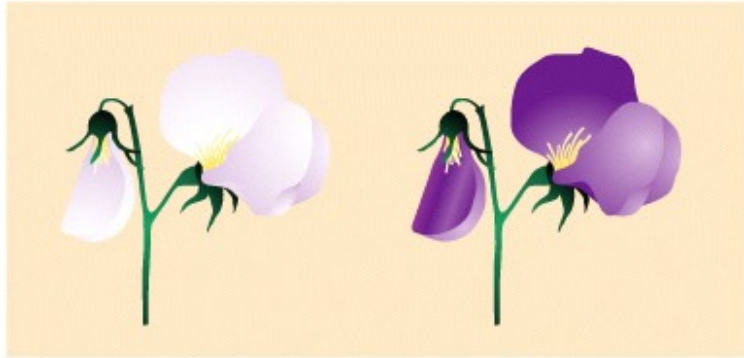
Courtesy New York Public Library

# Duplicate Genes

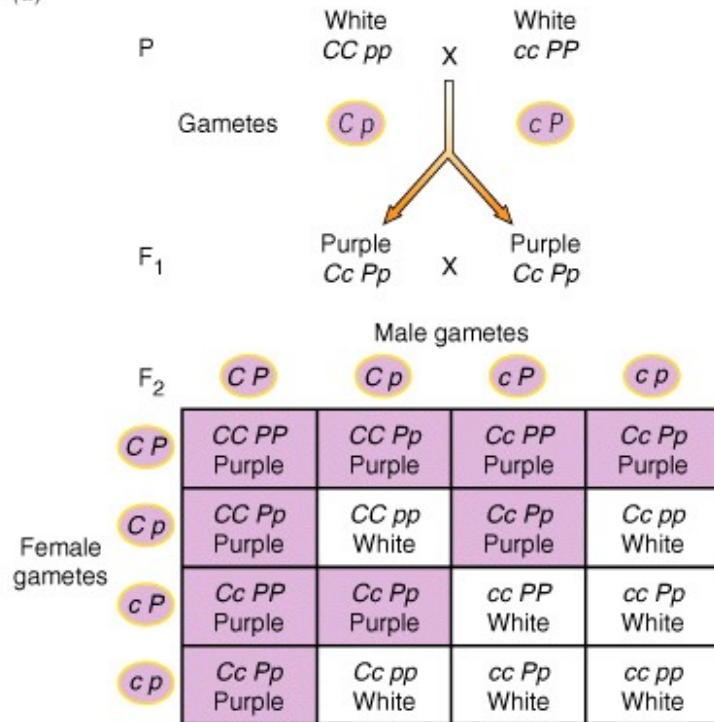


(b)

Summary: 15/16 triangular, 1/16 ovoid



(a)



(b)

Summary: 9/16 purple, 7/16 white

## Complementary Genes

Need BOTH interactive genes to be either in the **Homozygous dominant** ( $CC$ ,  $PP$ ) or **Heterozygous** ( $Cc$ ,  $Pp$ ) for one of the phenotypes to show through....

# Multiple or "Poly" Genes

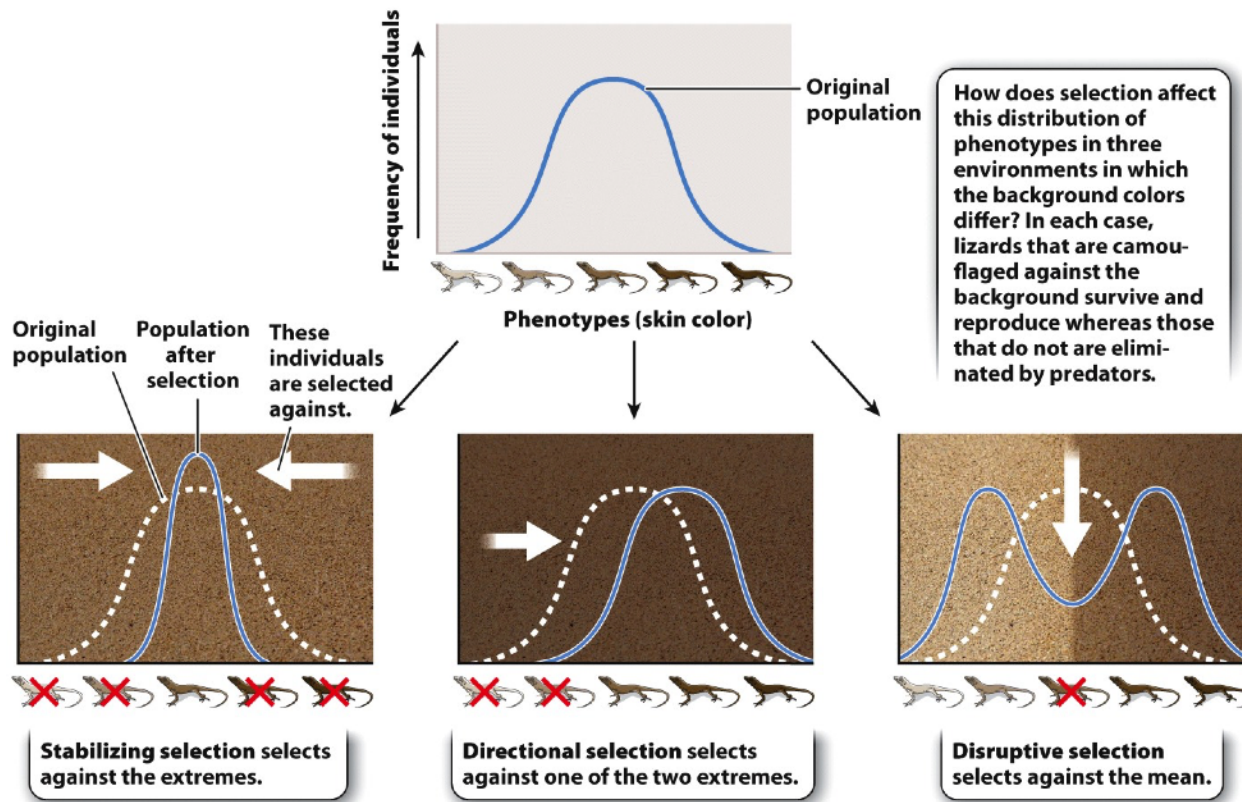
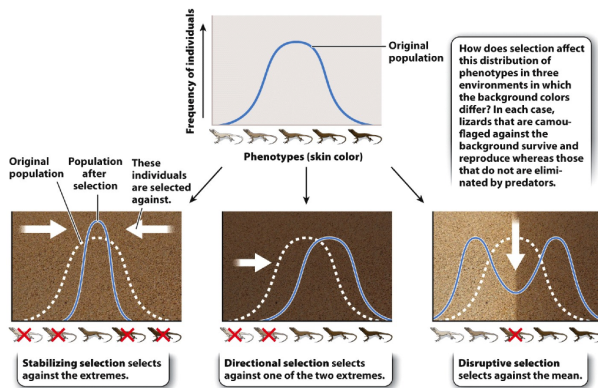


Figure 21.9  
 Biology: How Life Works  
 © 2014 W. H. Freeman and Company



# Multiple or “Poly” Genes



>50 genes heavily involved directly in structural height integrity in humans



## Number of genes linked to height revealed by study

*Date:* October 5, 2014

*Source:* Boston Children's Hospital

*Summary:* The largest genome-wide association study to date, involving more than 300 institutions and more than 250,000 subjects, roughly doubles the number of known gene regions influencing height to more than 400. The study provides a better glimpse at the biology of height and offers a model for investigating traits and diseases caused by many common gene changes acting together.

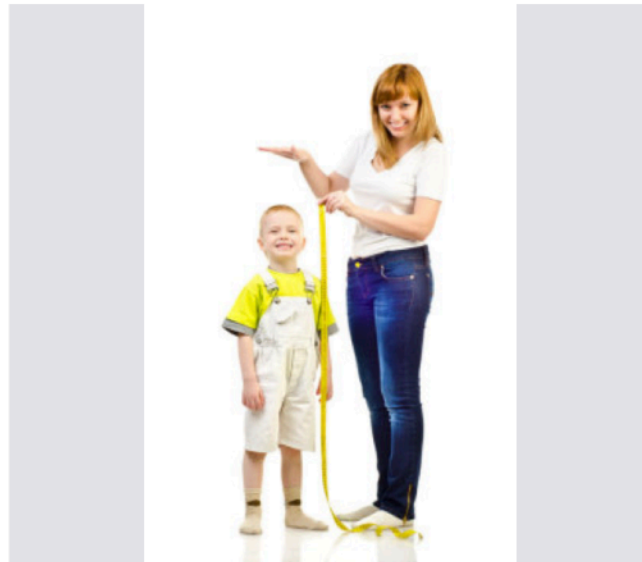
Share: [f](#) [t](#) [G+](#) [p](#) [in](#) [✉](#)

### RELATED TOPICS

#### Health & Medicine

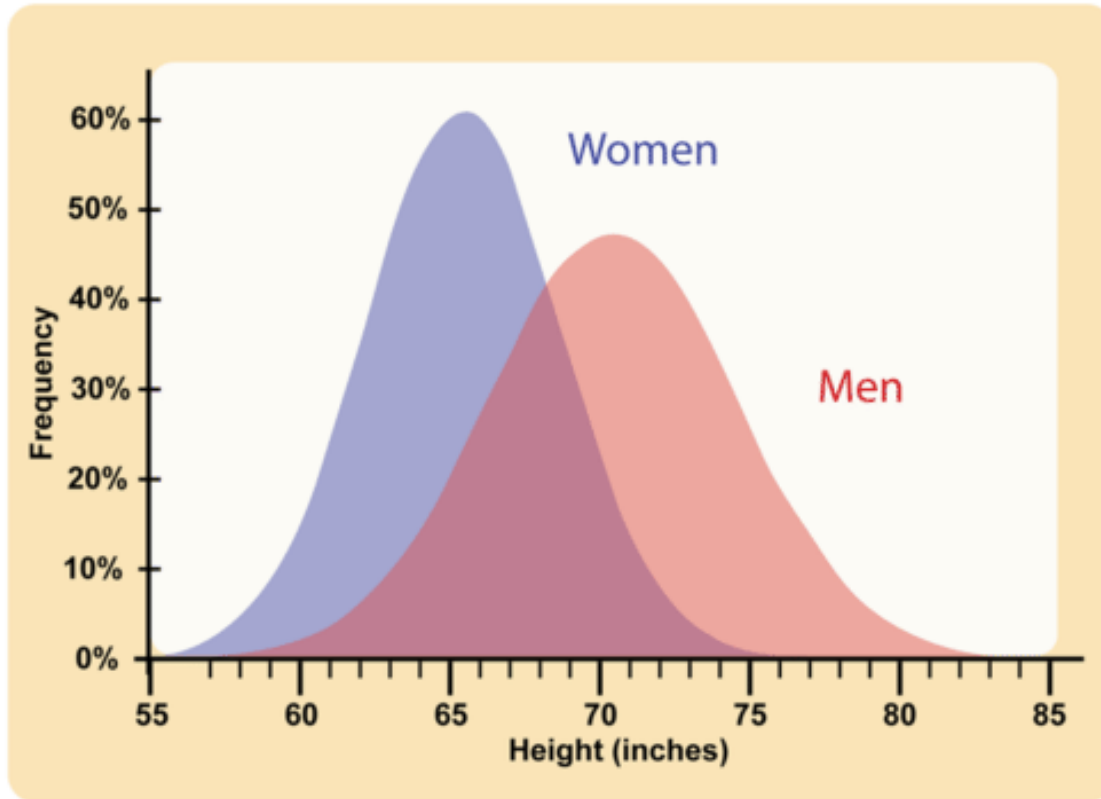
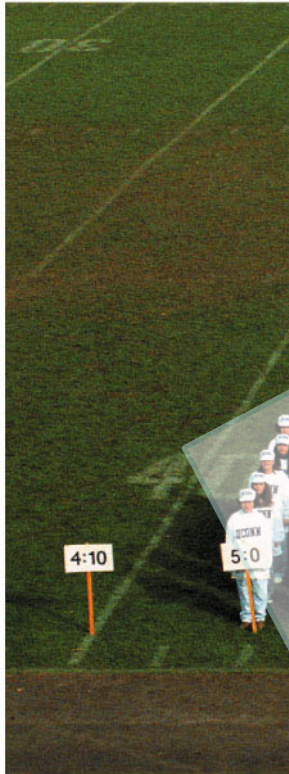
- > [Genes](#)
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- > [Hormone Disorders](#)
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### FULL STORY

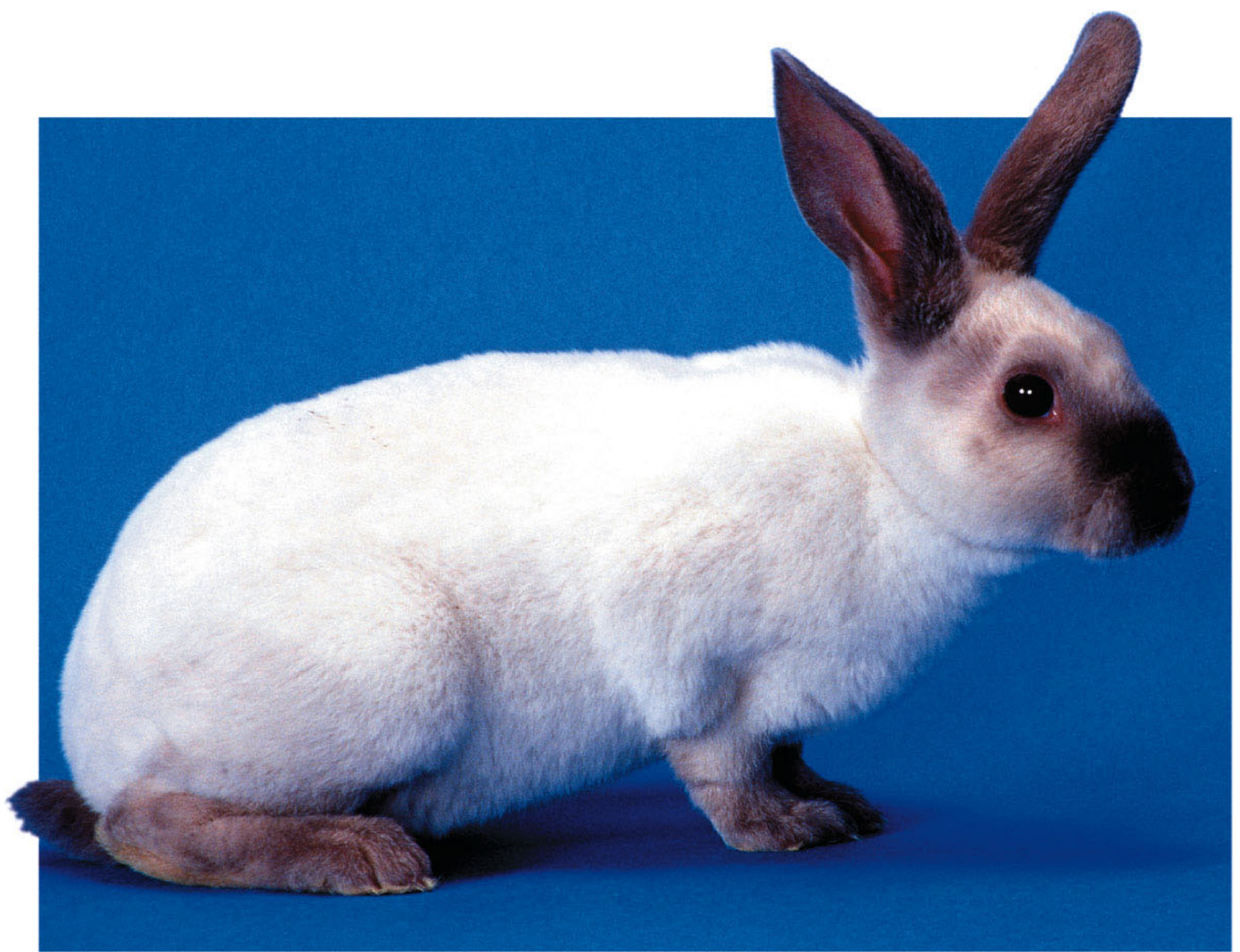


"We can now explain about 20 percent of the heritability of height, up from about 12 percent where we were before," says co-first author Tonu Esko, PhD, of Boston Children's Hospital, the Broad Institute and the University of Tartu (Estonia).

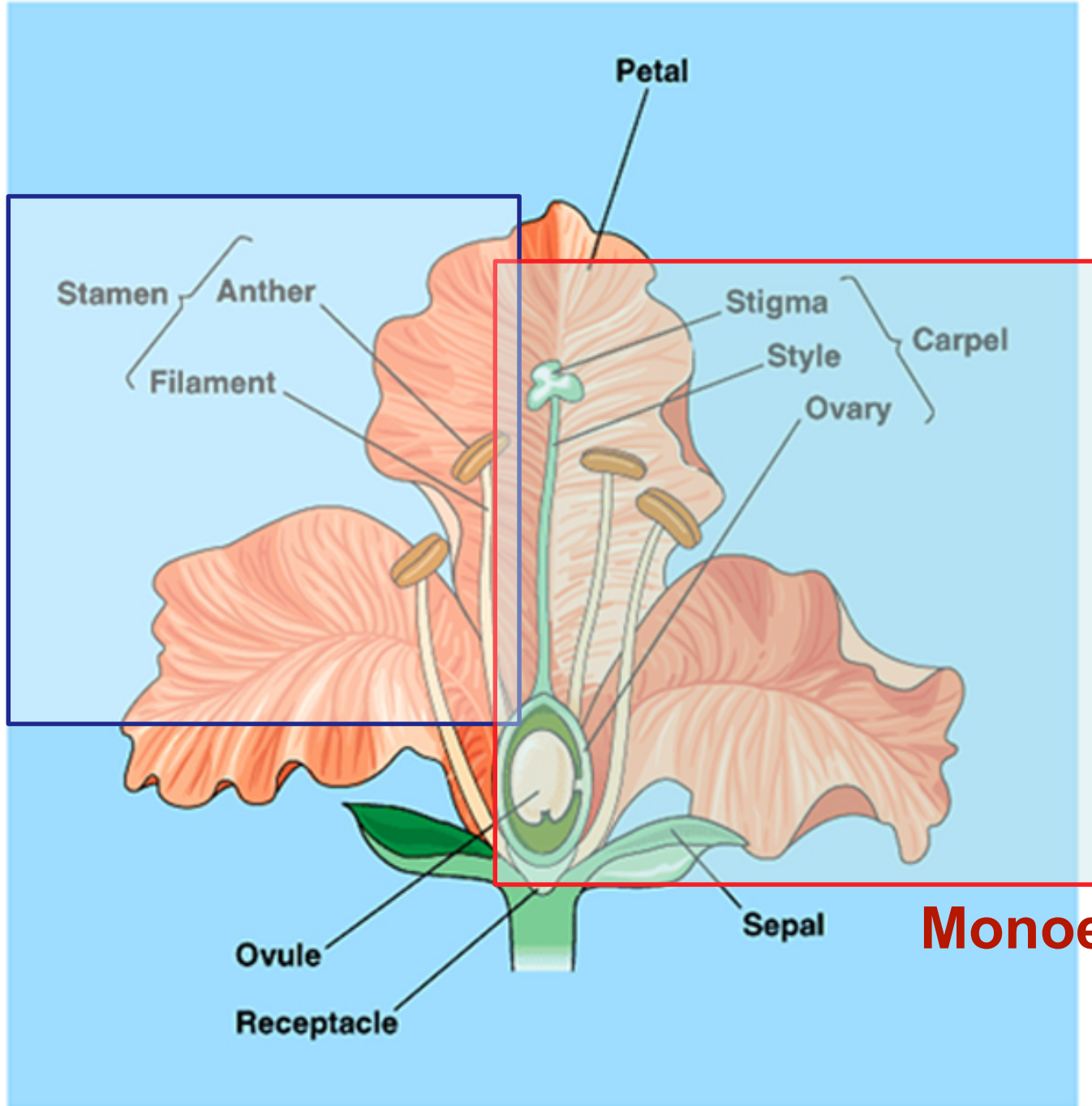
# Polygenes



Some of these assorted genes are also subject to female- and male- specific influences



**Nature vs. Nurture**



**Monoecious**

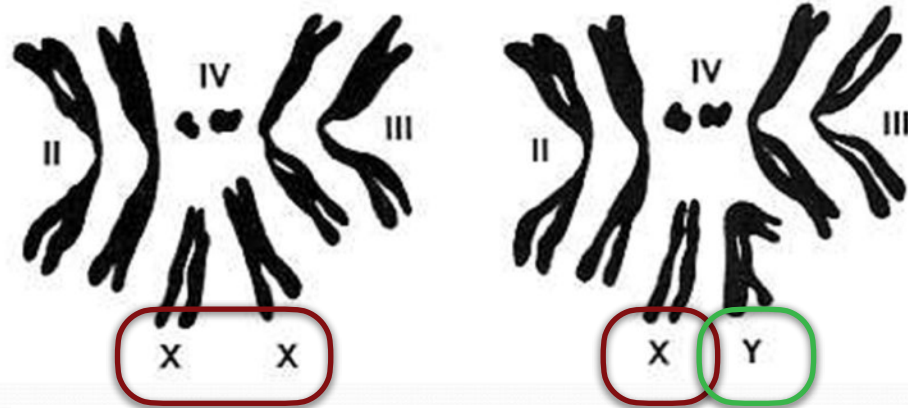


**Dioecious**



Most species of Holly are dioecious, meaning **male** and **female flowers** are on different plants -requiring a male holly plant to pollinate the female, which produce the **red berries**.

**Dioecious**



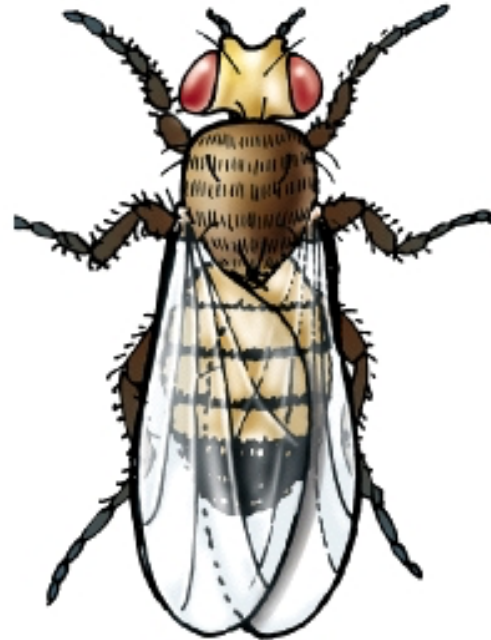
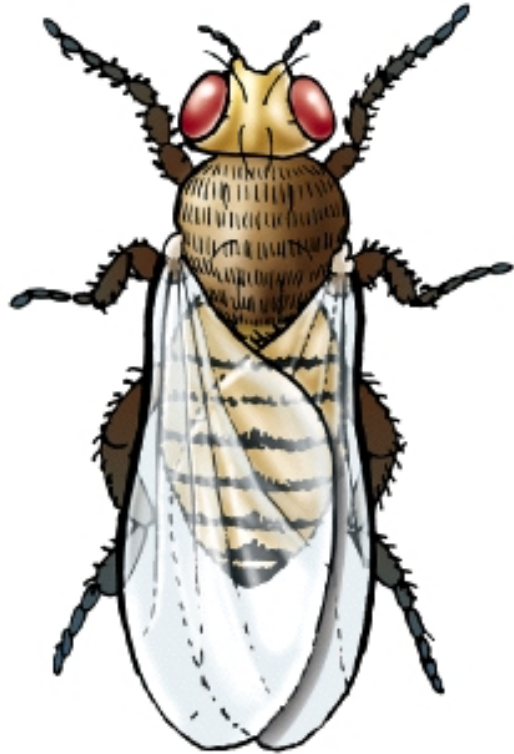
*Drosophila melanogaster* or “fruit fly”



**Dioecious**

*Homo sapiens* or “humans”

XX

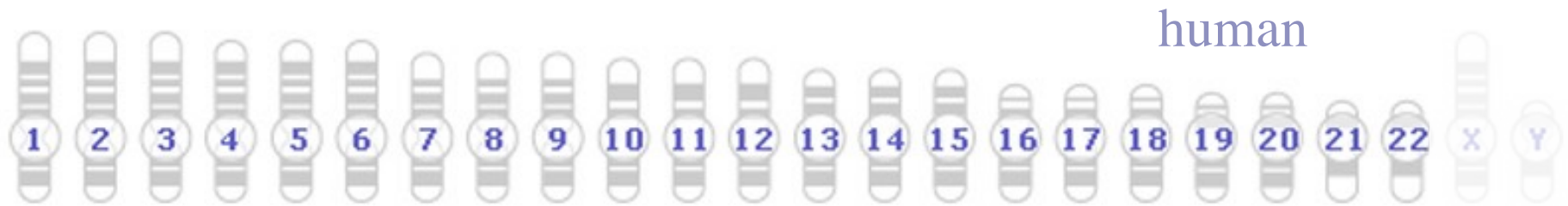


X Y

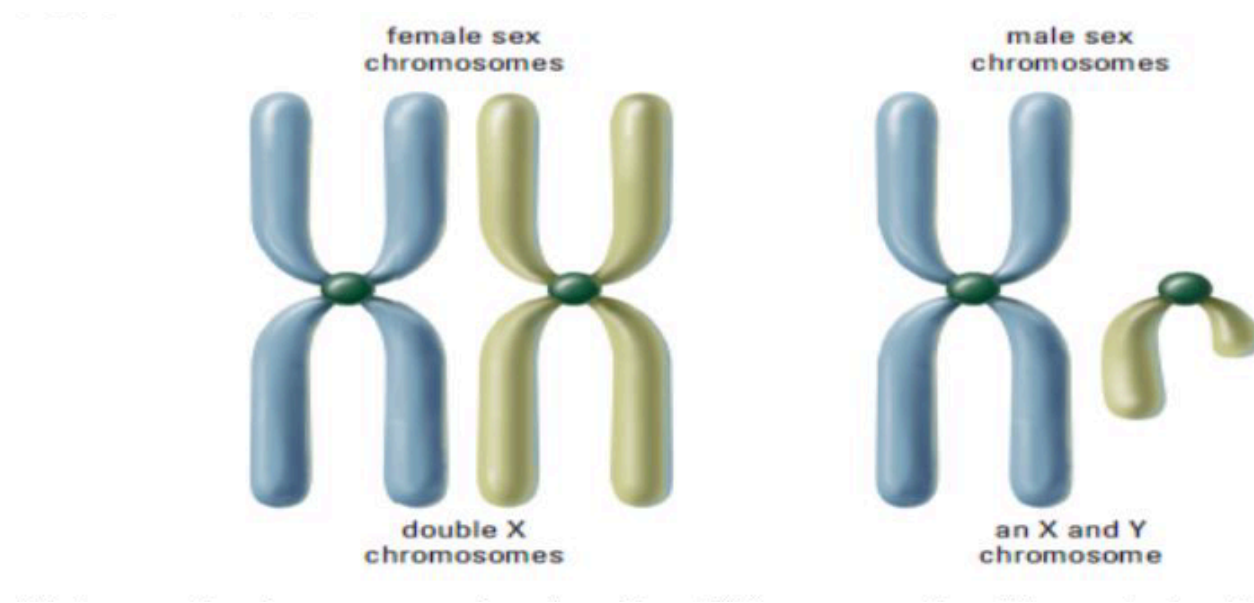


**Dioecious**





**autosomes:** the chromosomes not involved in sex determination



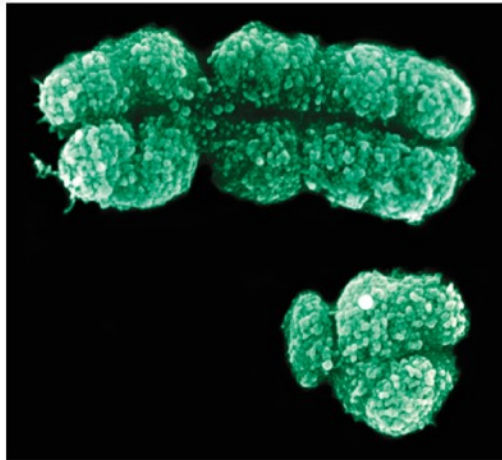
**sex chromosomes:** the pair of chromosomes that have a role in the sex of an individual

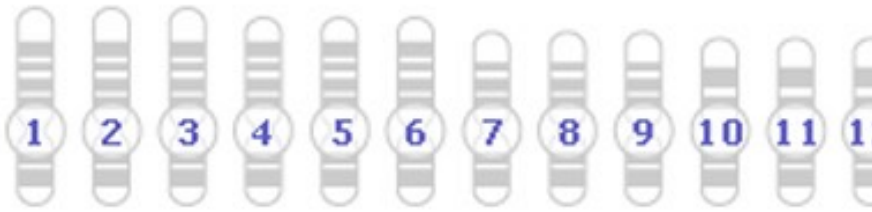
# human



Almost none of the genes in the X chromosome have counterparts in the Y chromosome.

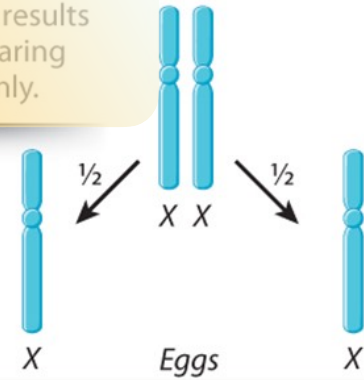
The tips of the arms of the X and Y chromosomes share a small region of homology (red).



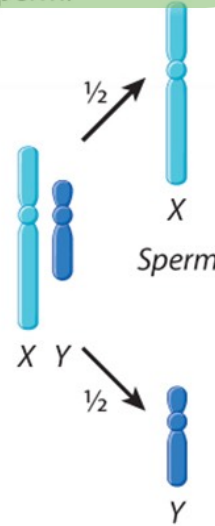


# human

Meiosis in a female results in X-bearing eggs only.

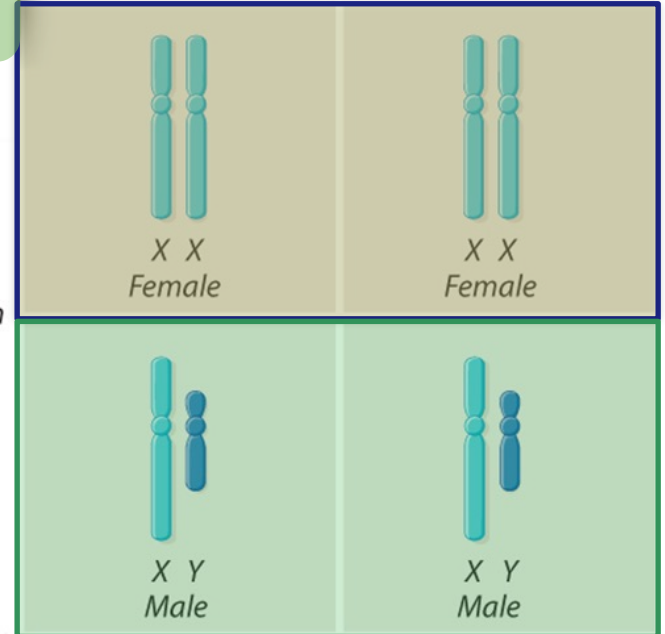
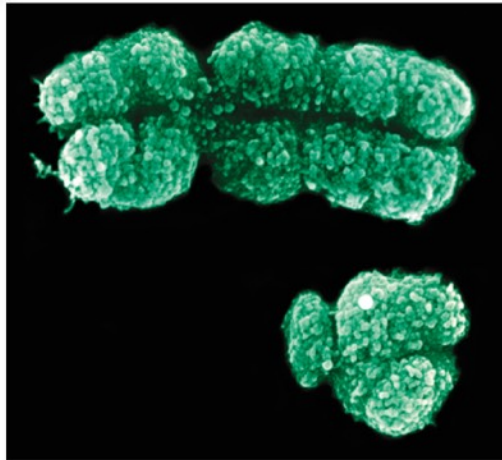
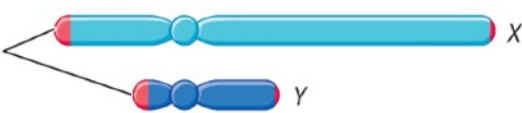


Meiosis in a male results in a 1:1 ratio of X-bearing and Y-bearing sperm.



The tips of the arms of the X and Y chromosomes share a small region of homology (red).

Almost none of the genes in the X chromosome have counterparts in the Y chromosome.



Random fertilization results in an expected ratio of 1/2 XX (female) and 1/2 XY (male) progeny.

# Sex-determining Region Y in Mammals

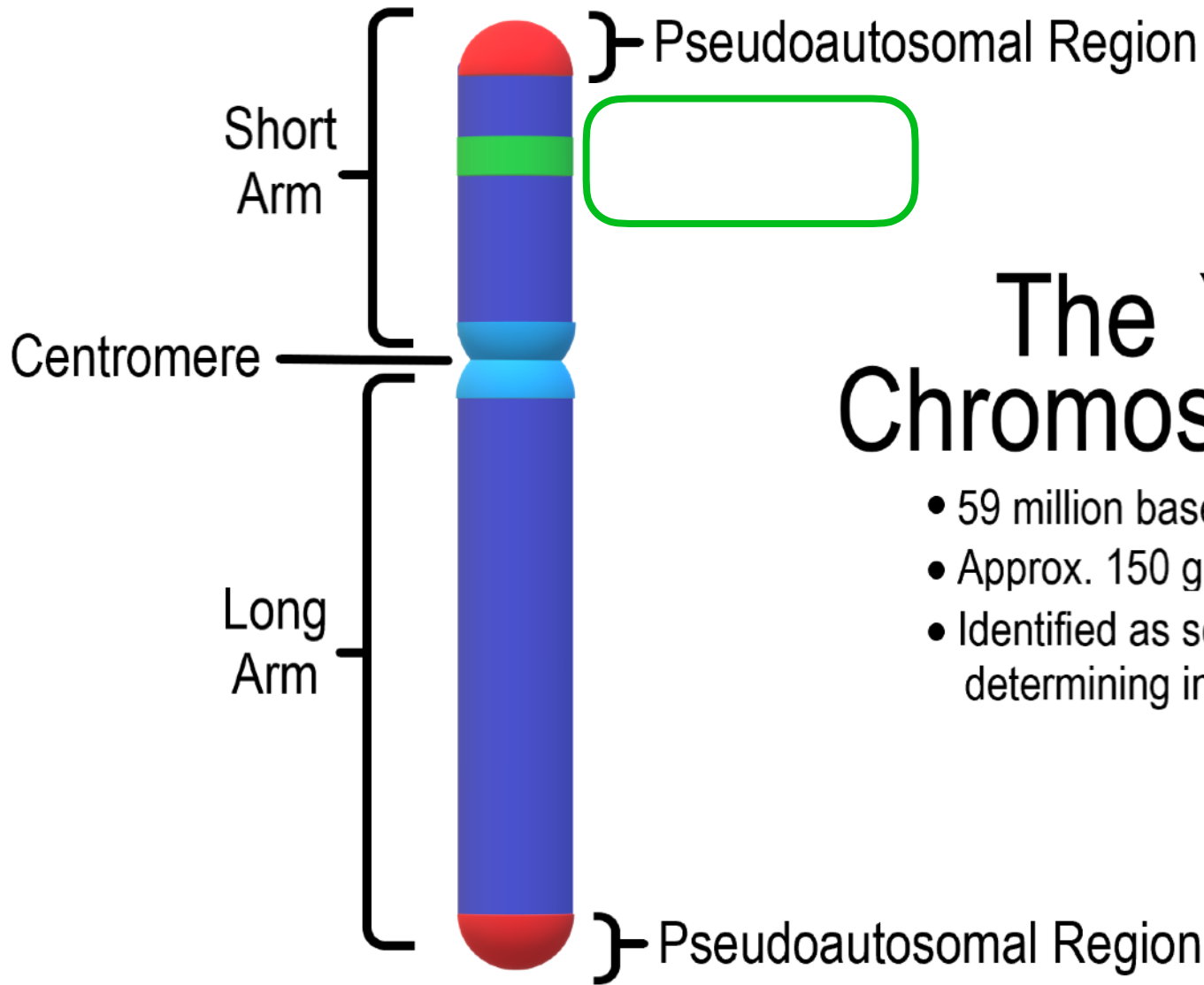
By: *Troy Cox*

Published: 2013-12-31

Keywords: [Sex-Related Gene On Y](#), [Sex Determination Processes](#)

The Sex-determining Region Y (*Sry* in mammals but *SRY* in [humans](#)) is a gene found on Y chromosomes that leads to the development of male phenotypes, such as [testes](#). The *Sry* gene, located on the short branch of the Y chromosome, initiates male embryonic development in the XY [sex determination](#) system. The *Sry* gene follows the central dogma of molecular biology; the DNA encoding the gene is transcribed into messenger RNA, which then produces a single *Sry* protein. The *Sry* protein is also called the testis-determining factor (TDF), a protein that initiates male development in [humans](#), placental mammals, and marsupials. The *Sry* protein is a transcription factor that can bind to regions of testis-specific DNA, bending specific DNA and activating or enhancing its abilities to promote testis formation, marking the first step towards male, rather than female, development in the embryo.

In [humans](#) the first step in the development of an organism's sex is the inheritance of an X chromosome from the mother, and either an X or Y chromosome from the father. Typically, an XX individual develops as a female and an XY individual develops as a male. Studies by [University of Kansas](#) zoologist Clarence Erwin McClung in Lawrence, Kansas at the turn of the twentieth century helped researchers focus on the roles of chromosomes for [sex determination](#). McClung theorized that there were two distinct types of [spermatozoa](#), each of which resulted in different forms of fertilized eggs, leading to either male or female development. Nettie Maria Stevens, a post-doctorate researcher at [Bryn Mawr College](#), located near Philadelphia, Pennsylvania, expanded upon McClung's theory in 1905, observing that [spermatozoa](#) are of two distinct forms, containing either an X or a Y chromosome. Based upon her research on [sex determination](#) in insect species, Stevens concluded that the Y chromosome carries the genetic material that leads to male development.

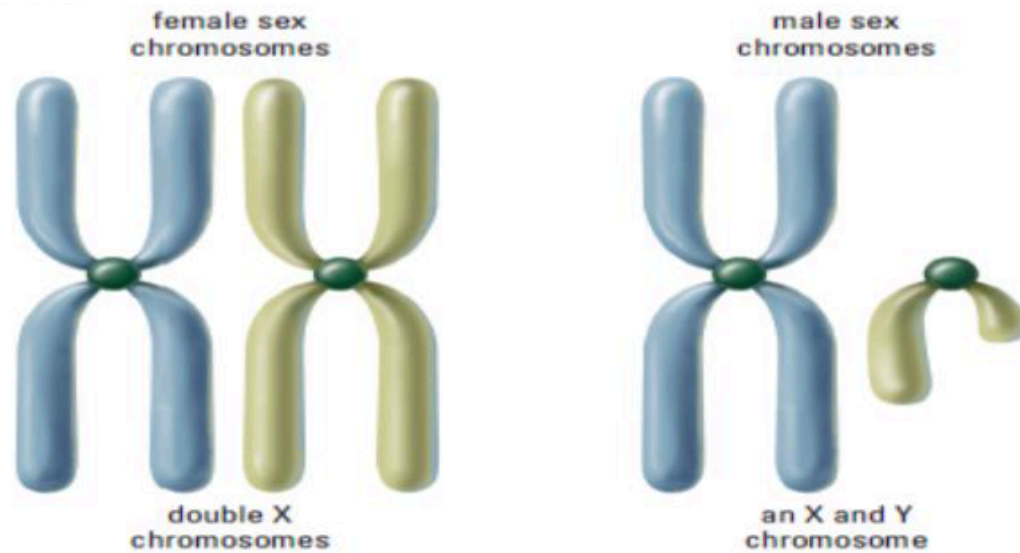


# The Y Chromosome

- 59 million base pairs
- Approx. 150 genes
- Identified as sex determining in 1905

All genes that are present on the X-chromosome,  
demonstrate a genetic phenomenon called...

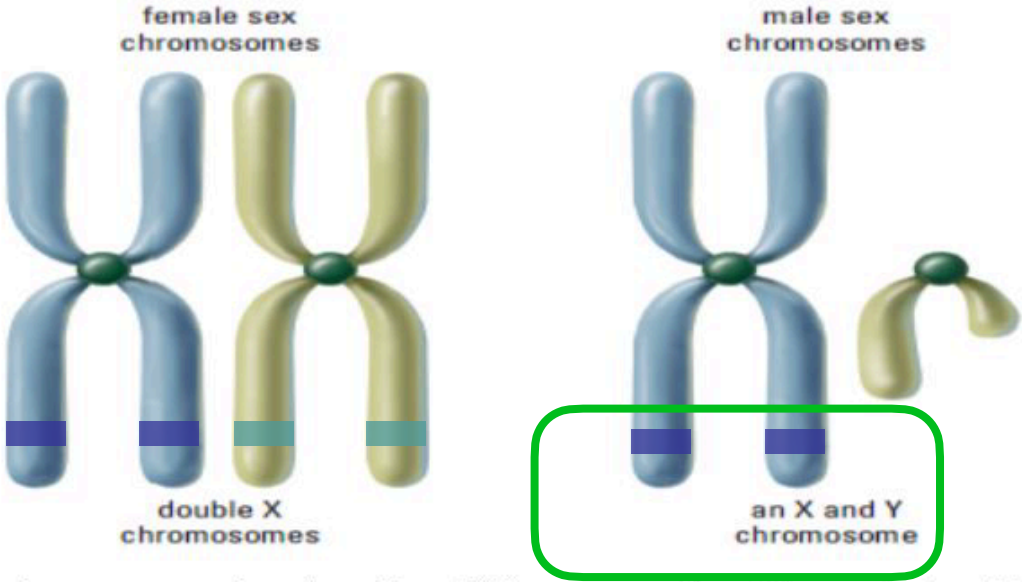
## X-linkage



# Non-Mendelian Inheritance

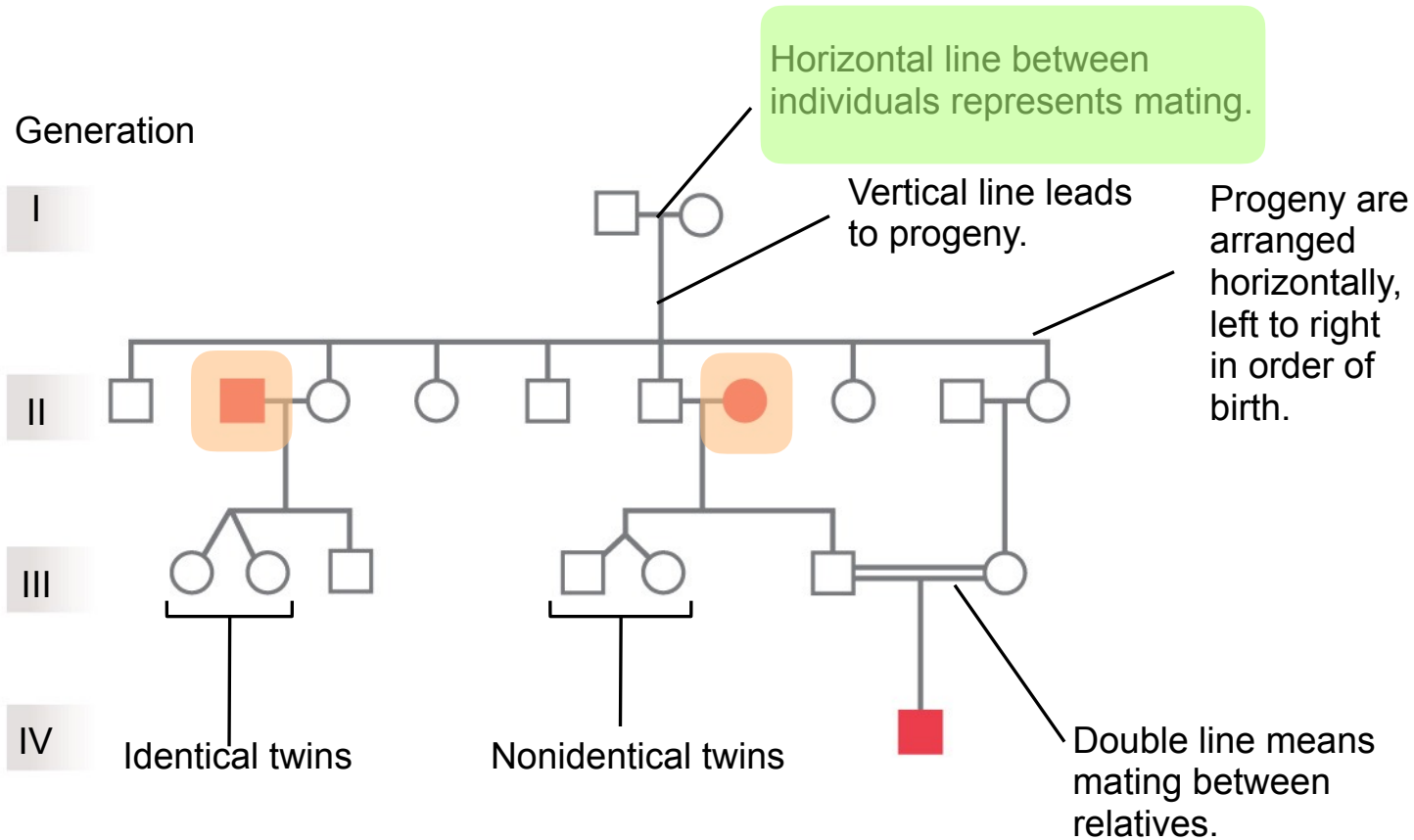


## X-linkage



hemizygous

# Patterns of Inheritance in Humans



## Family Pedigrees

- Female
- Male
- Open symbol means not affected.
- Darkened symbol means affected.

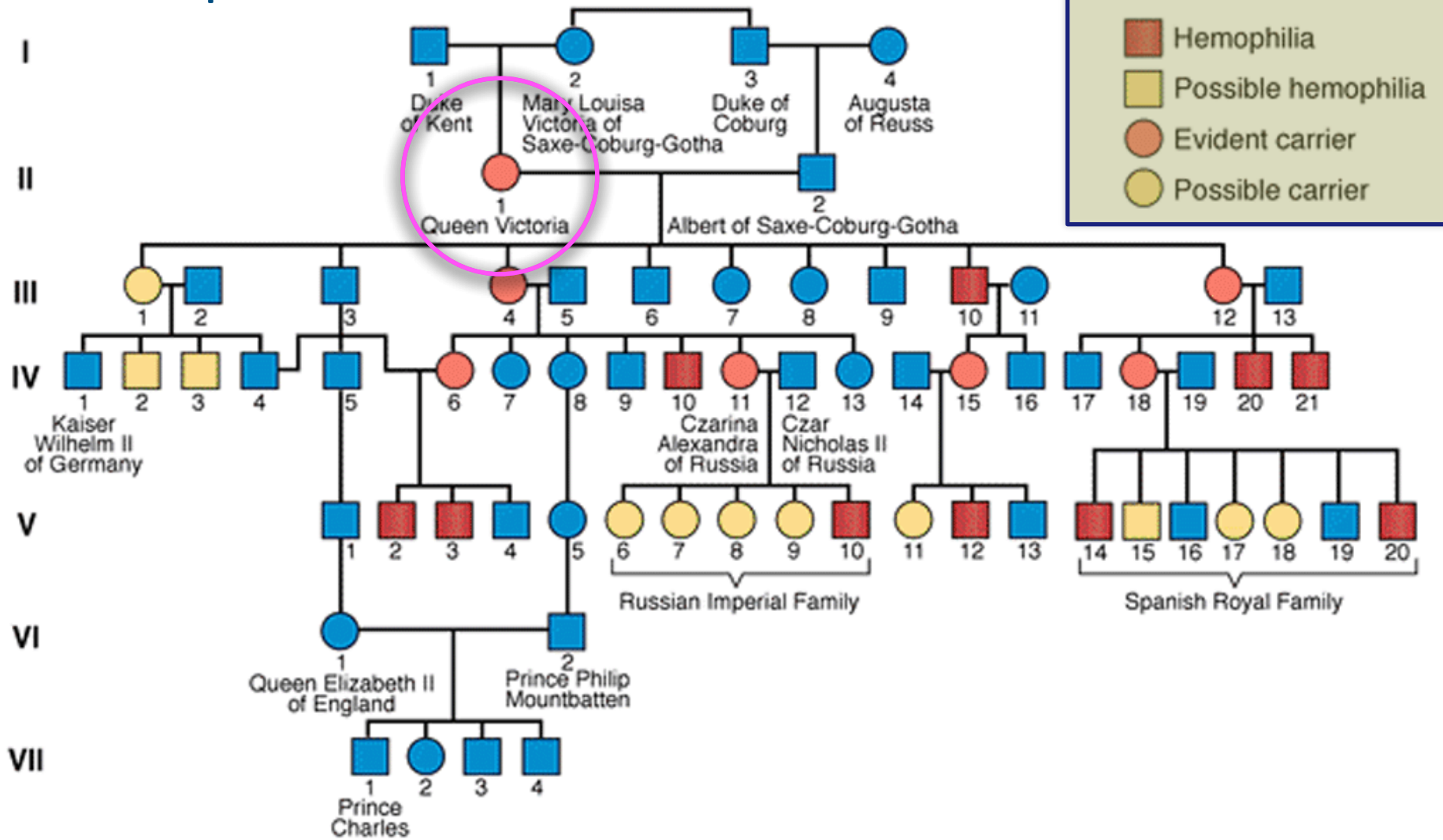


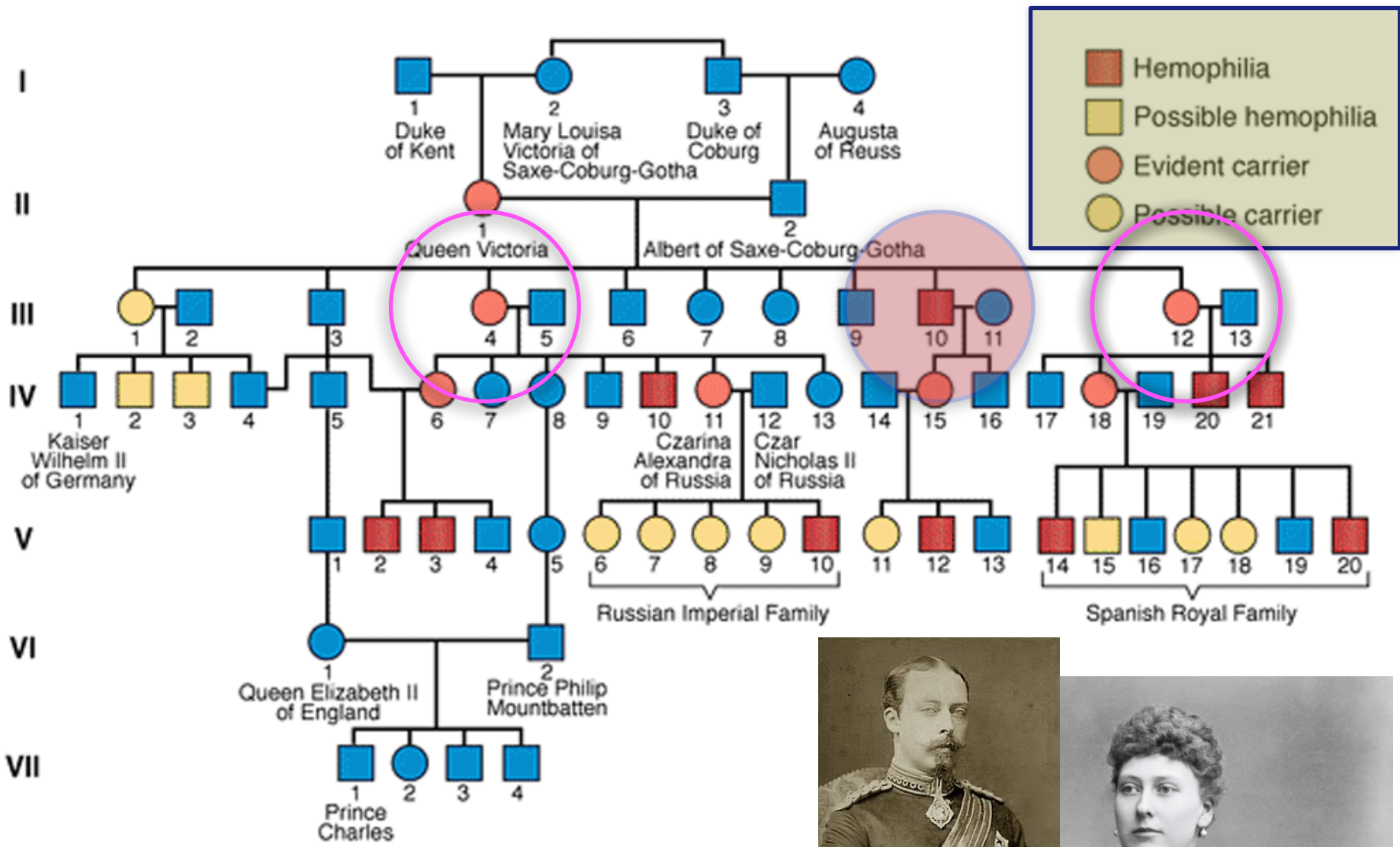
# Haemophilia



Queen Victoria, nine children,  
six of their spouses & twenty three grandchildren

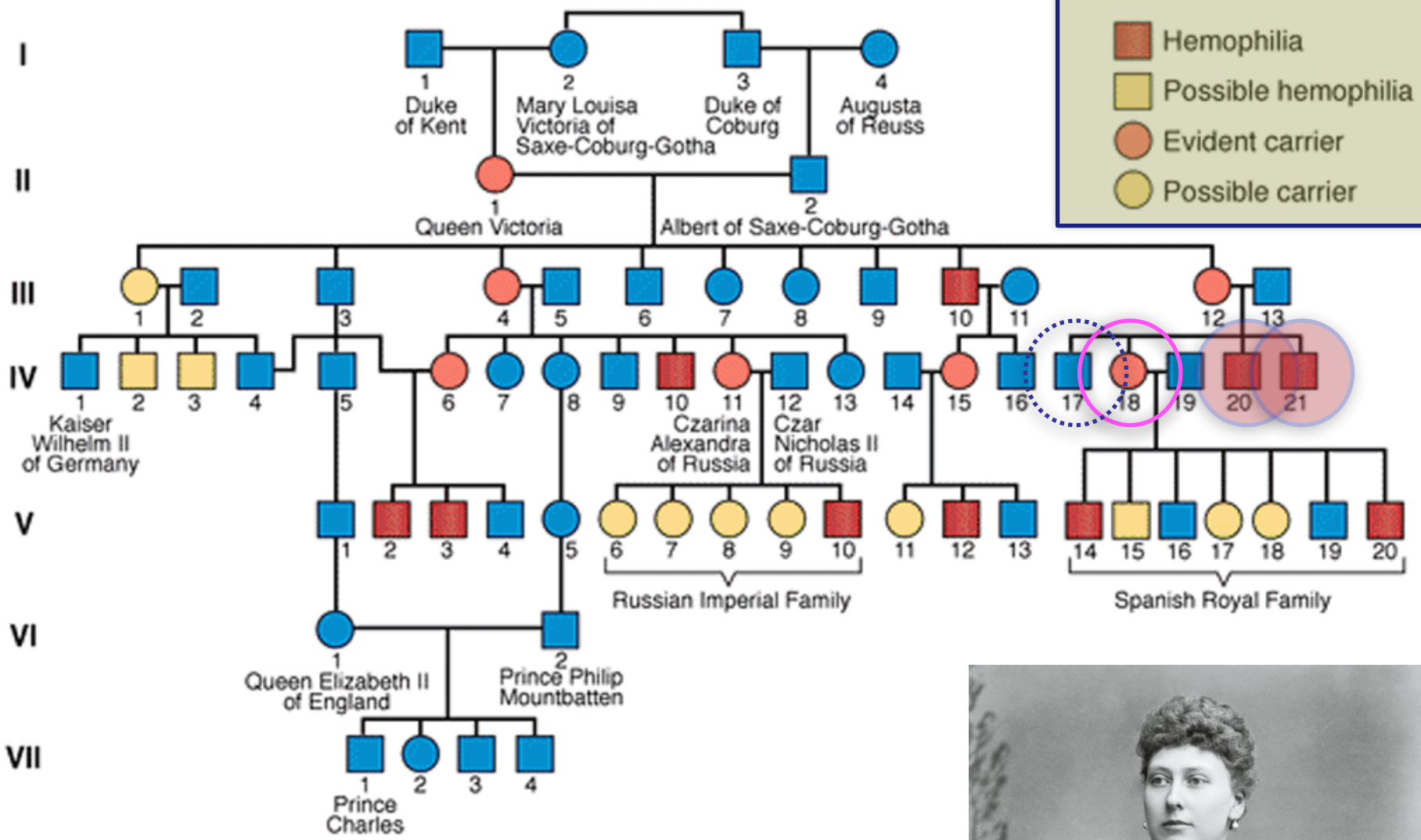
# Haemophilia

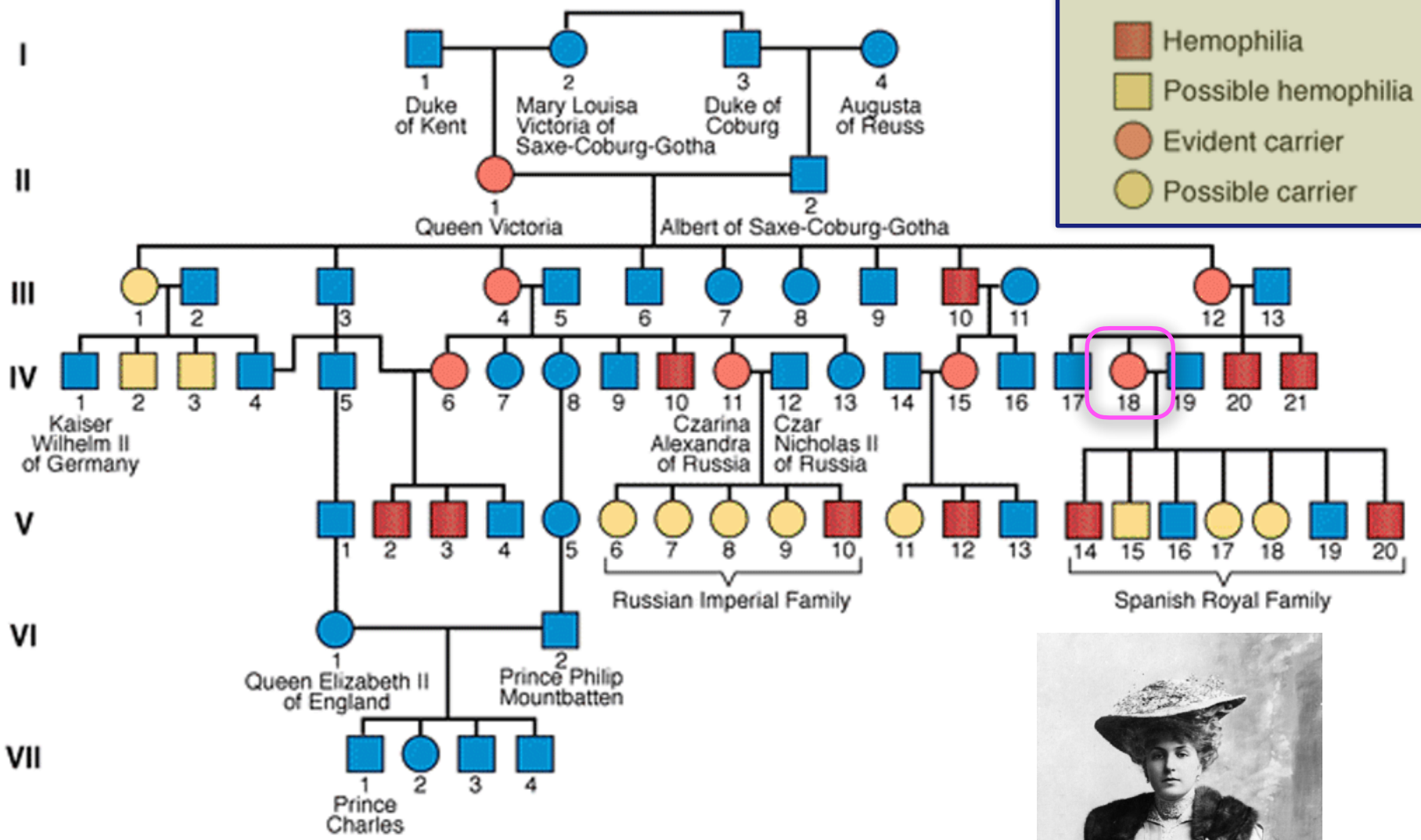


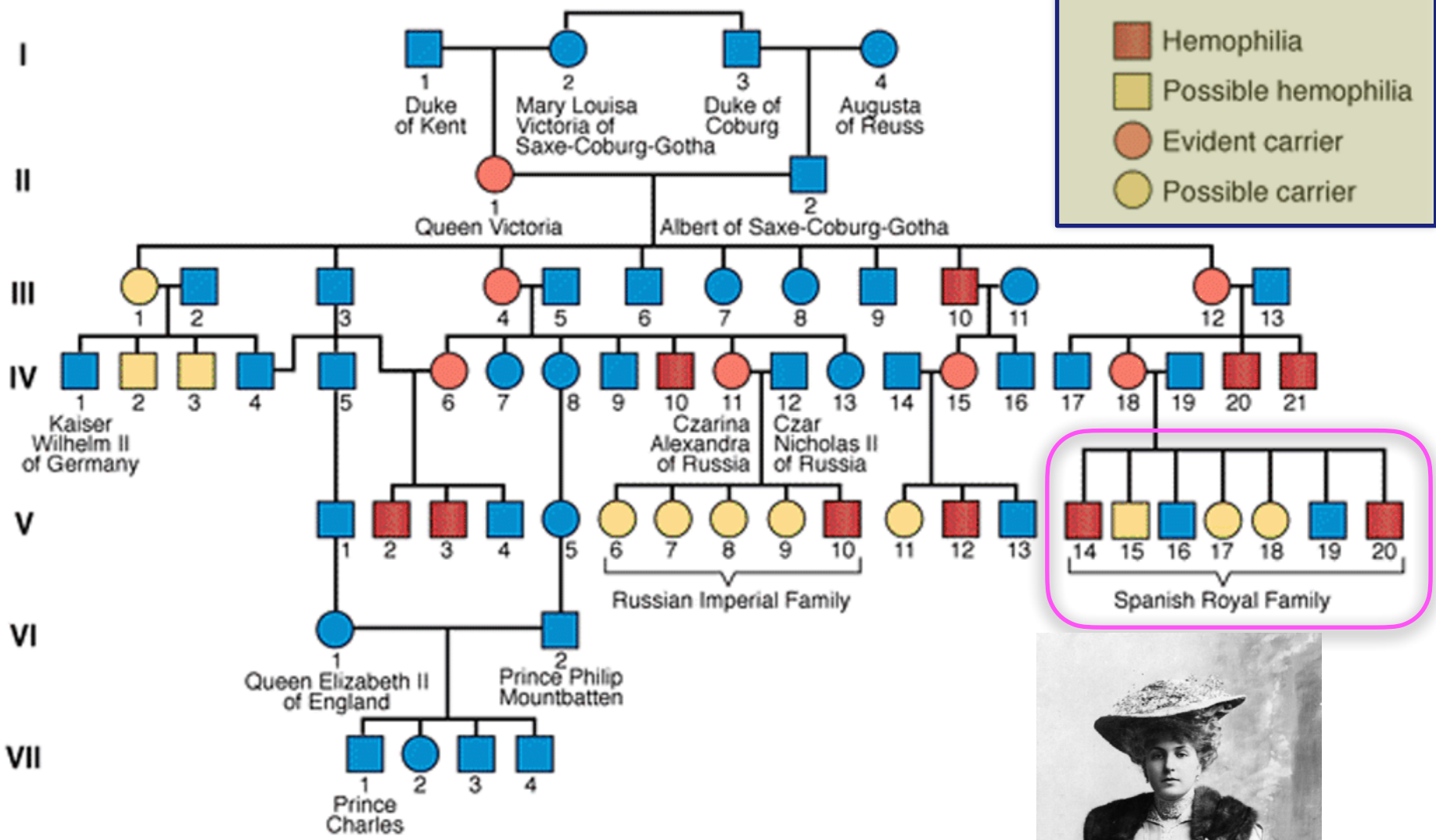


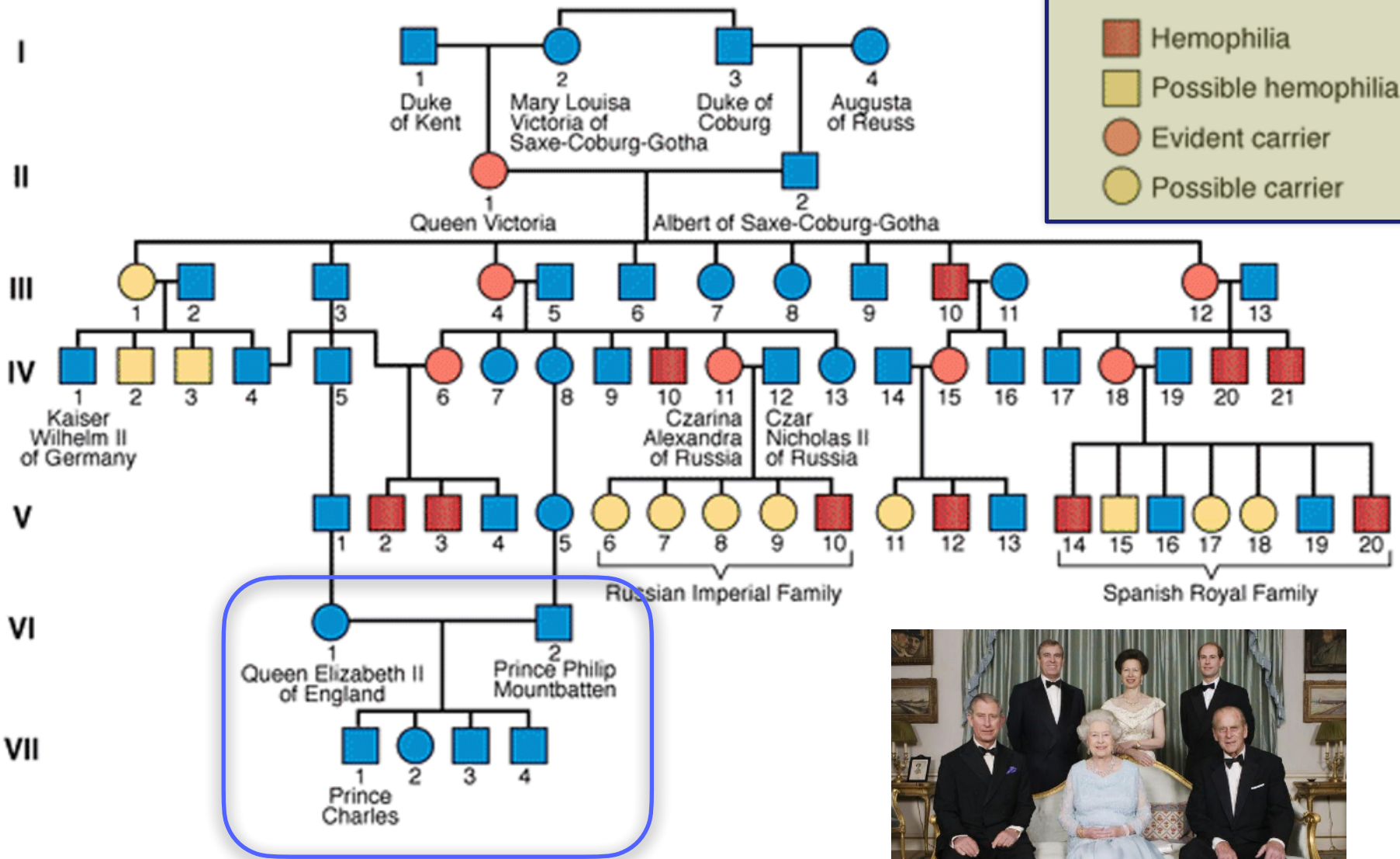
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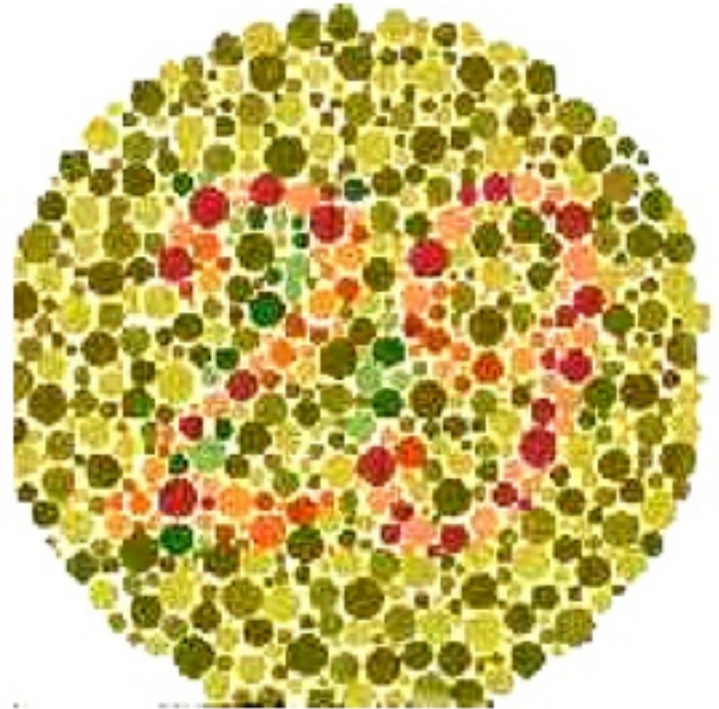
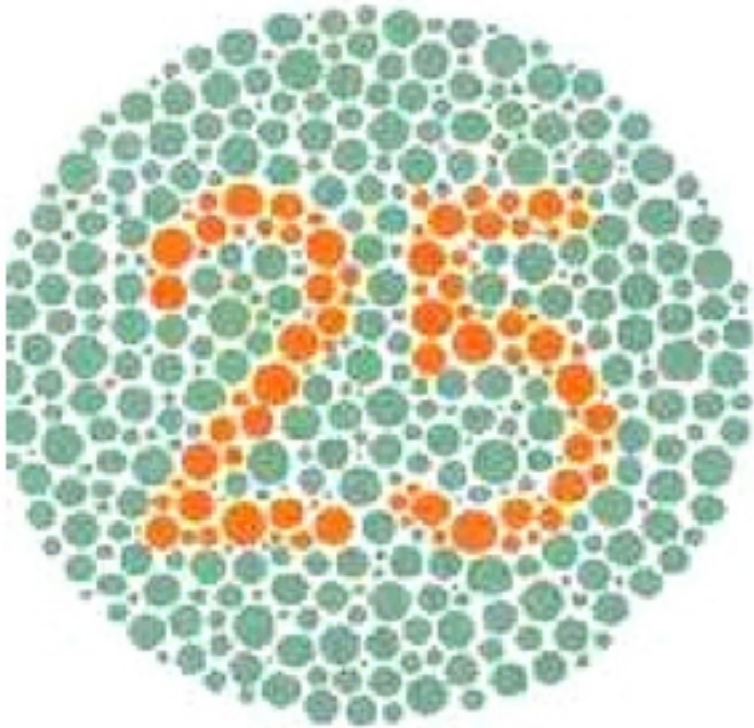


# Red Green Colour blindness

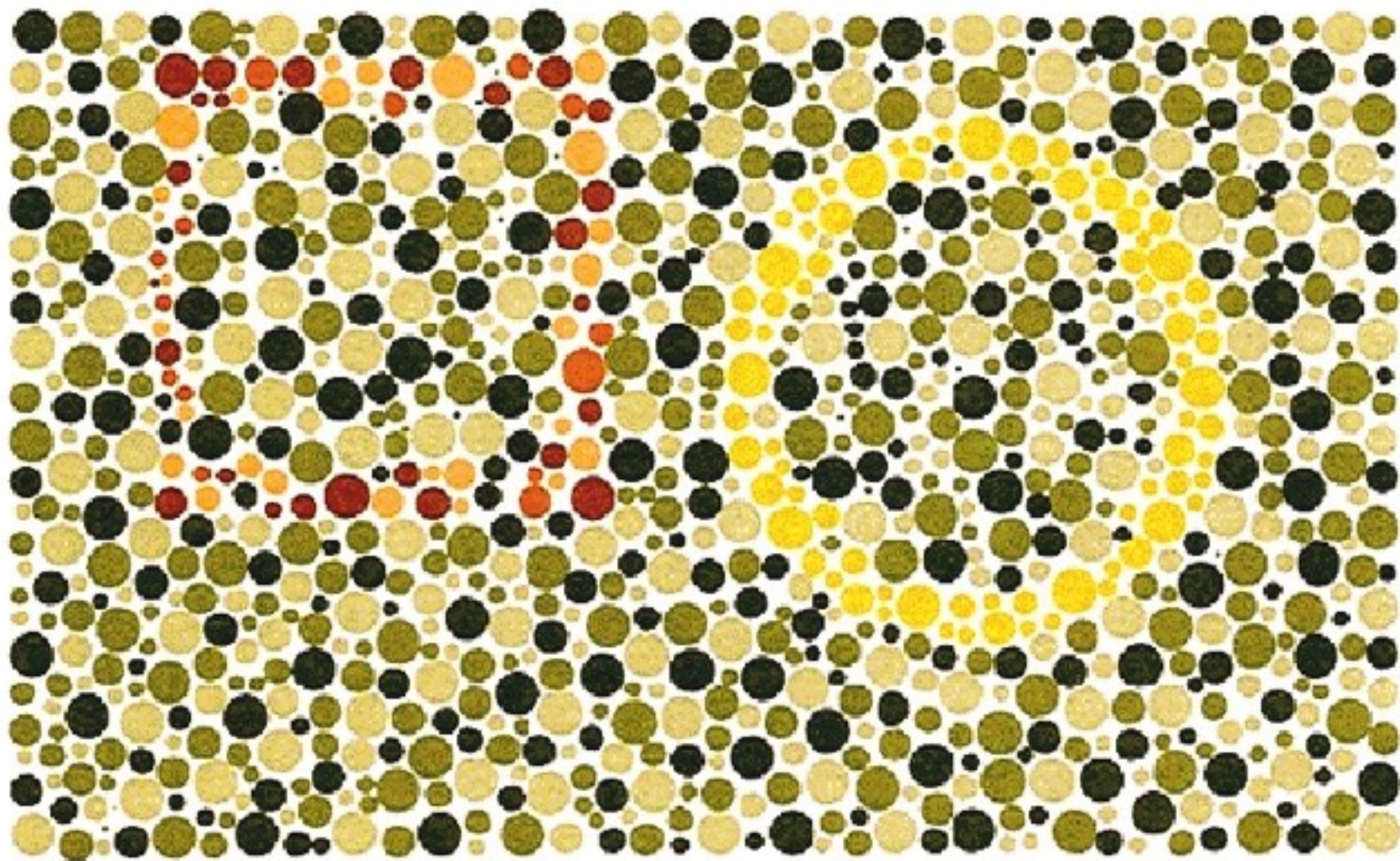


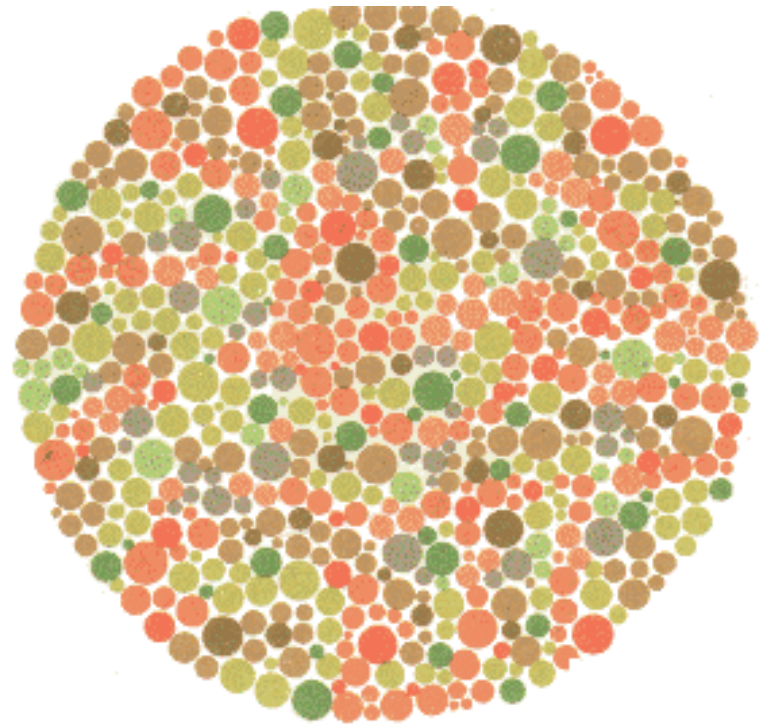
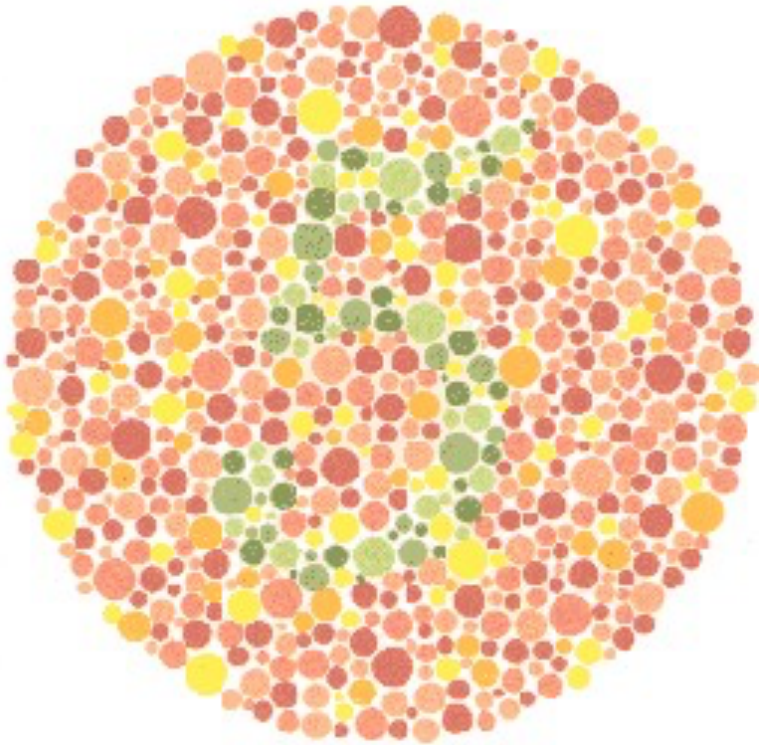




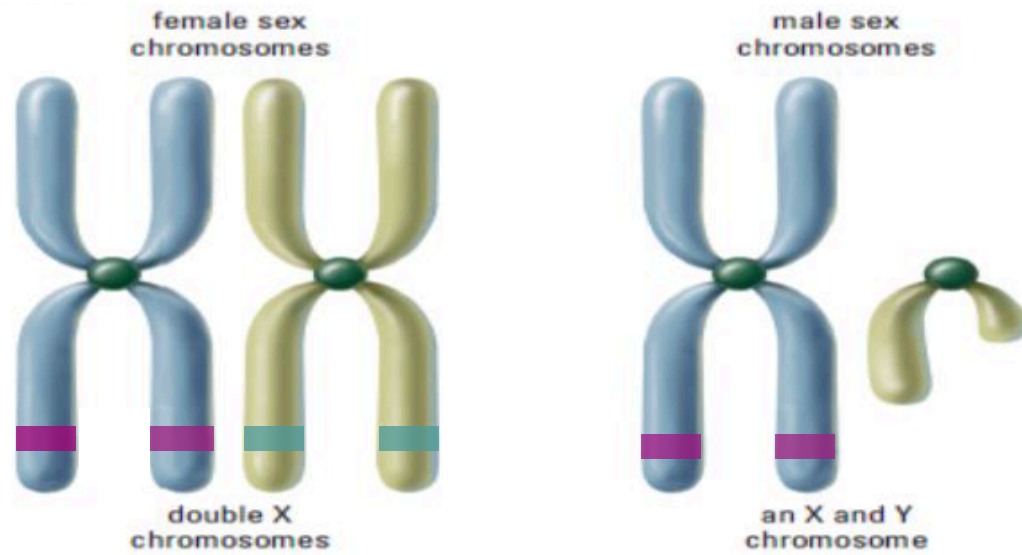


Ishihara colour blindness tests





# X-linkage



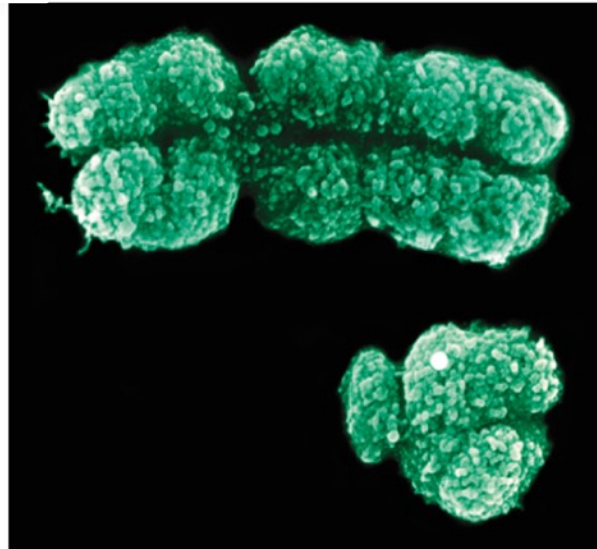
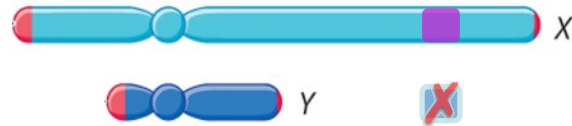
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So while a females can carry an X- linked trait, if it is recessive- the other X chromosome would probably not, and it's expression would DOMINATE giving a WT phenotype.

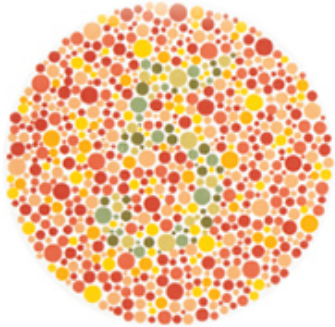
Hence Females can often be carriers of an X- linked trait, but rarely demonstrate the phenotype.

Giving rise to the following inheritable signs for **X- Linkage**



For Males it's a different story, if the X chromosome carries the trait... there is NO compensating X chromosome to help hide the trait, and if it is present it WILL ALWAYS SHOW THROUGH

People with red-green color blindness cannot see the number 5 in this figure.

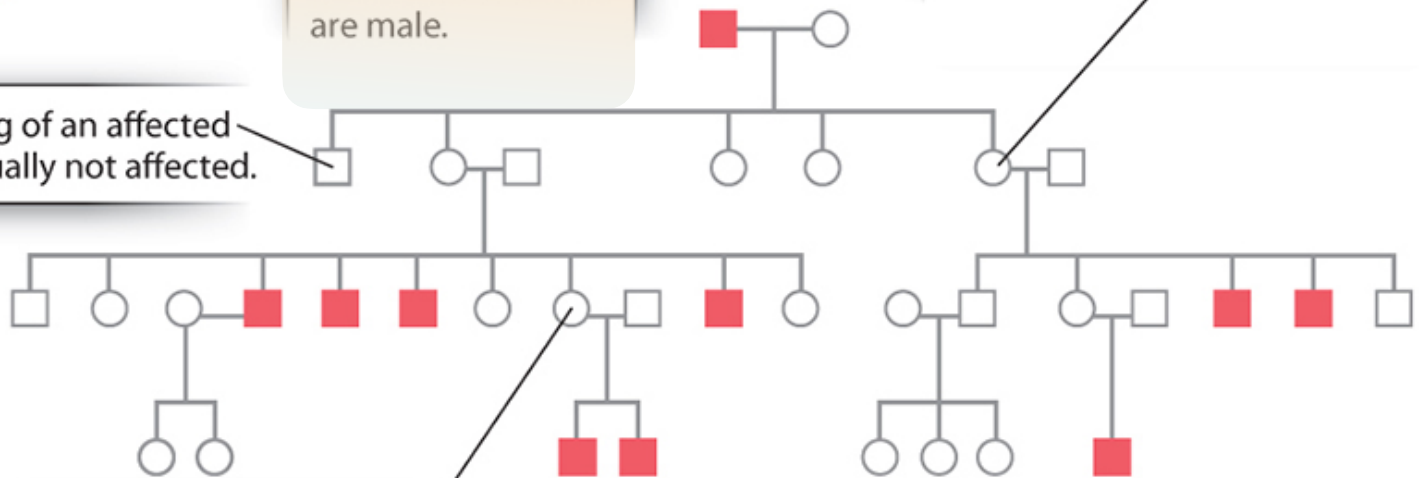


- Unaffected male
- Unaffected female
- Affected male

For a rare X-linked recessive trait, most affected individuals are male.

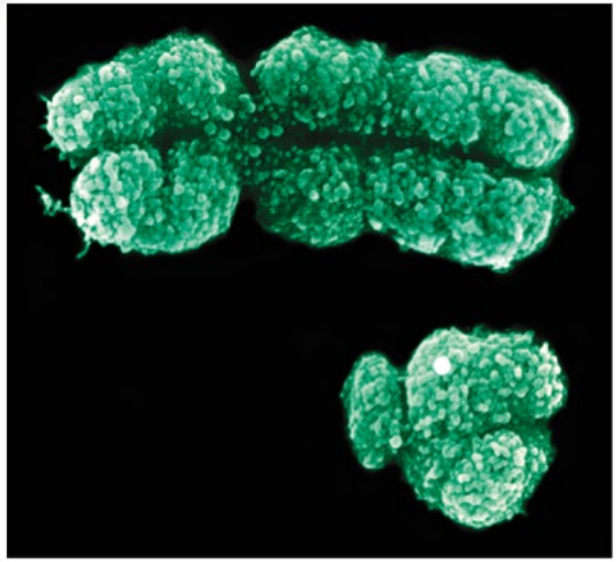
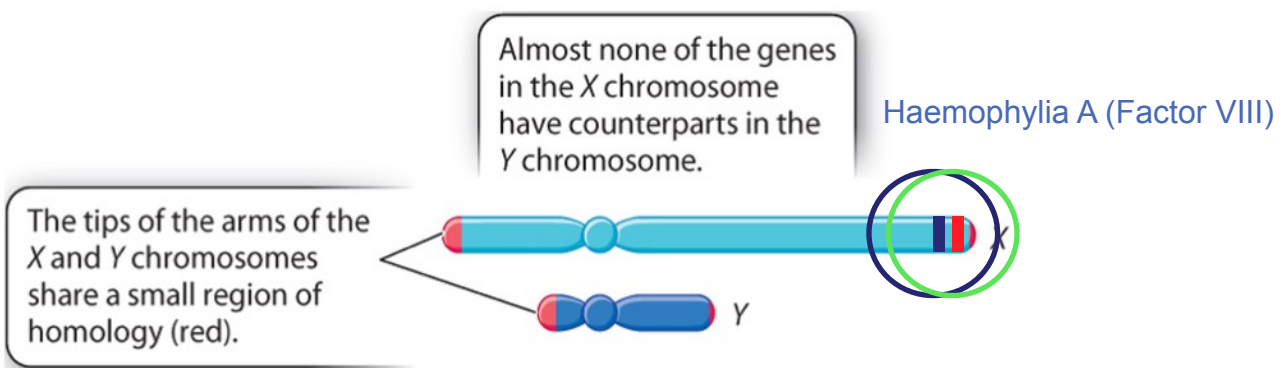
The daughters of affected males can have affected sons.

The offspring of an affected male are usually not affected.



The sisters of an affected male can have affected sons.

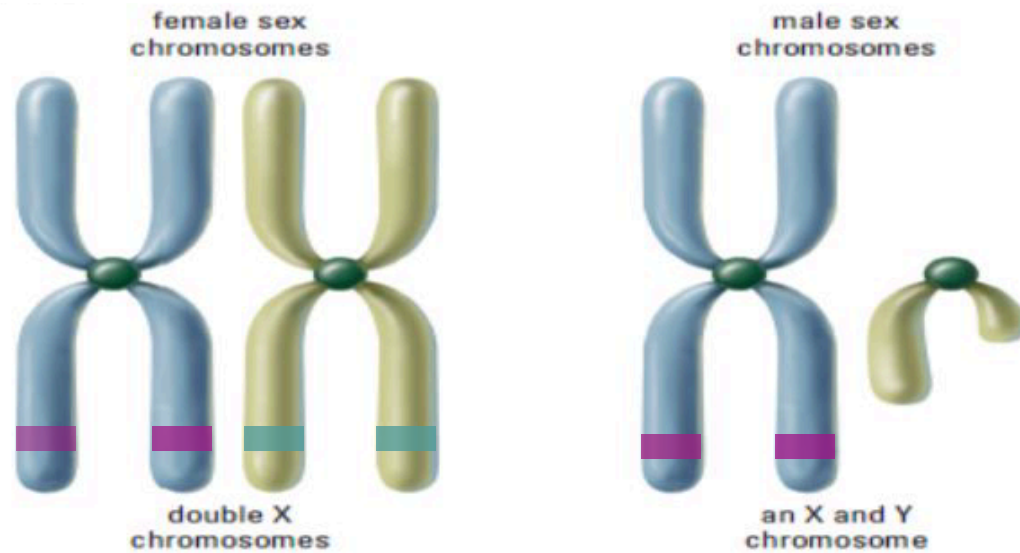


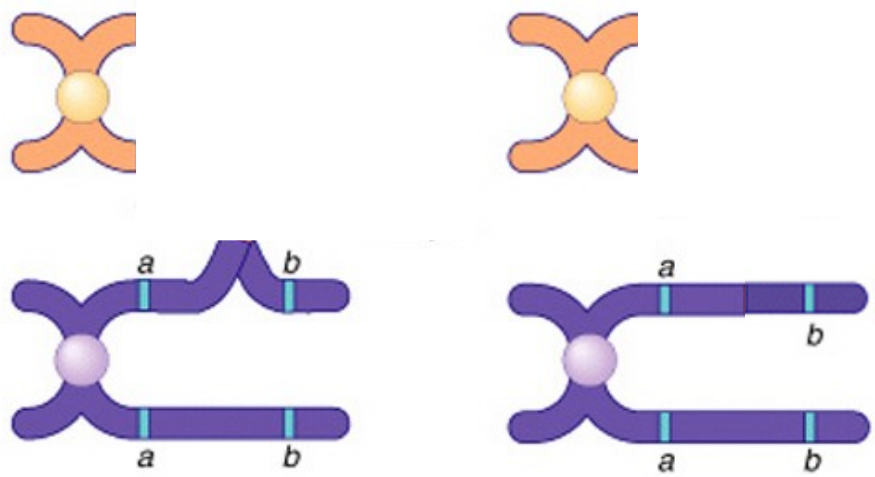




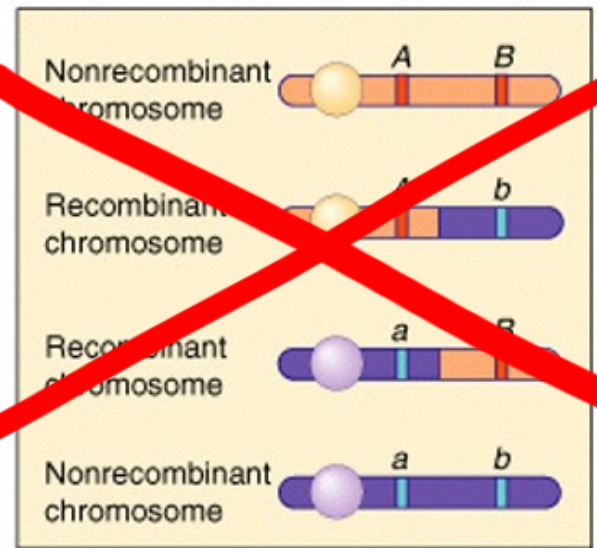
# Non-Mendelian Inheritance

## X-linkage

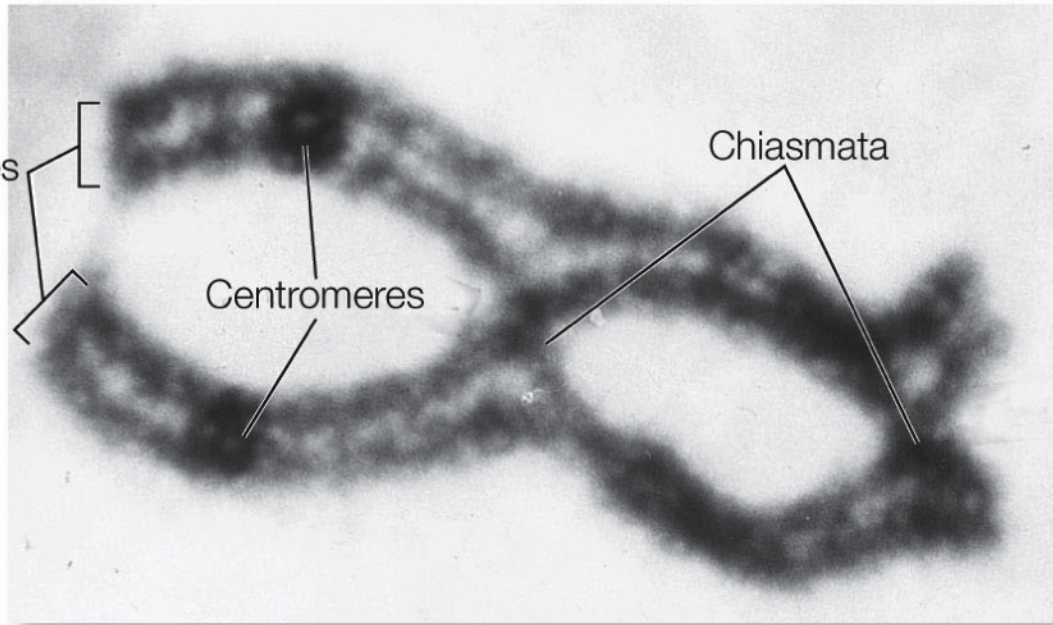




Four products of meiosis



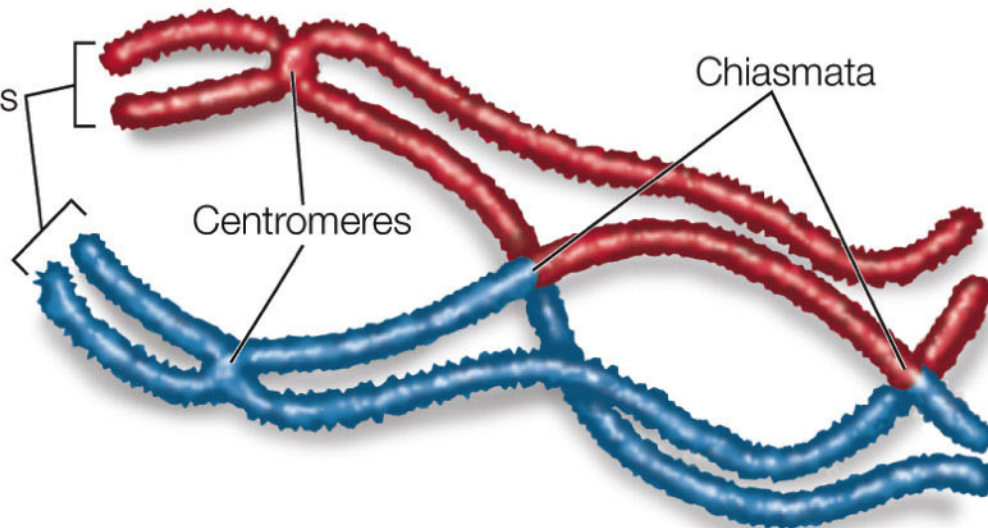
Homologous chromosomes



Centromeres

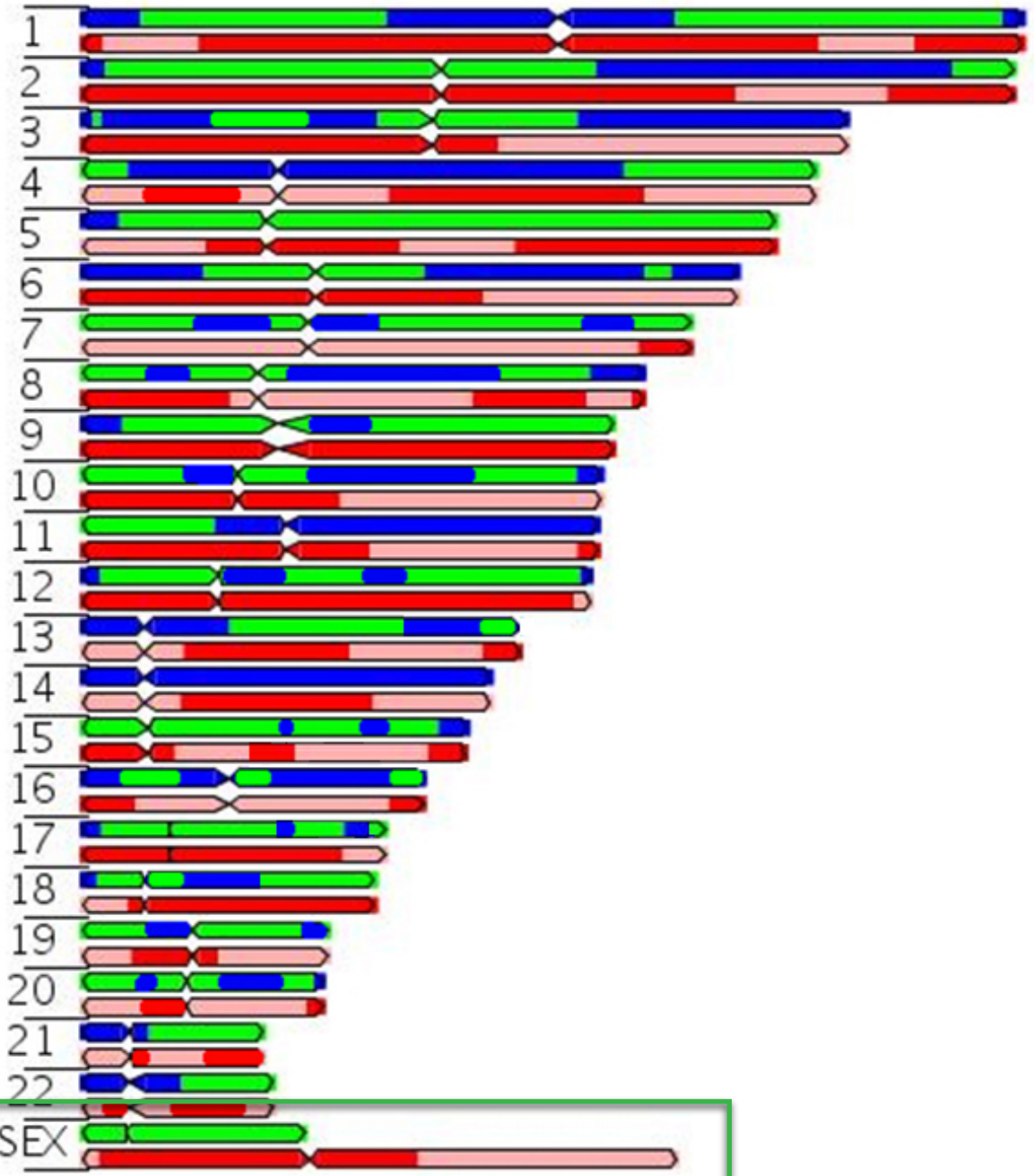
Chiasmata

Homologous chromosomes



Centromeres

Chiasmata



Father



Mother



