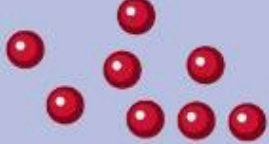
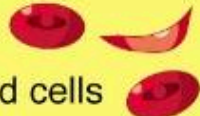
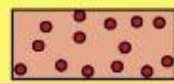
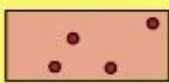
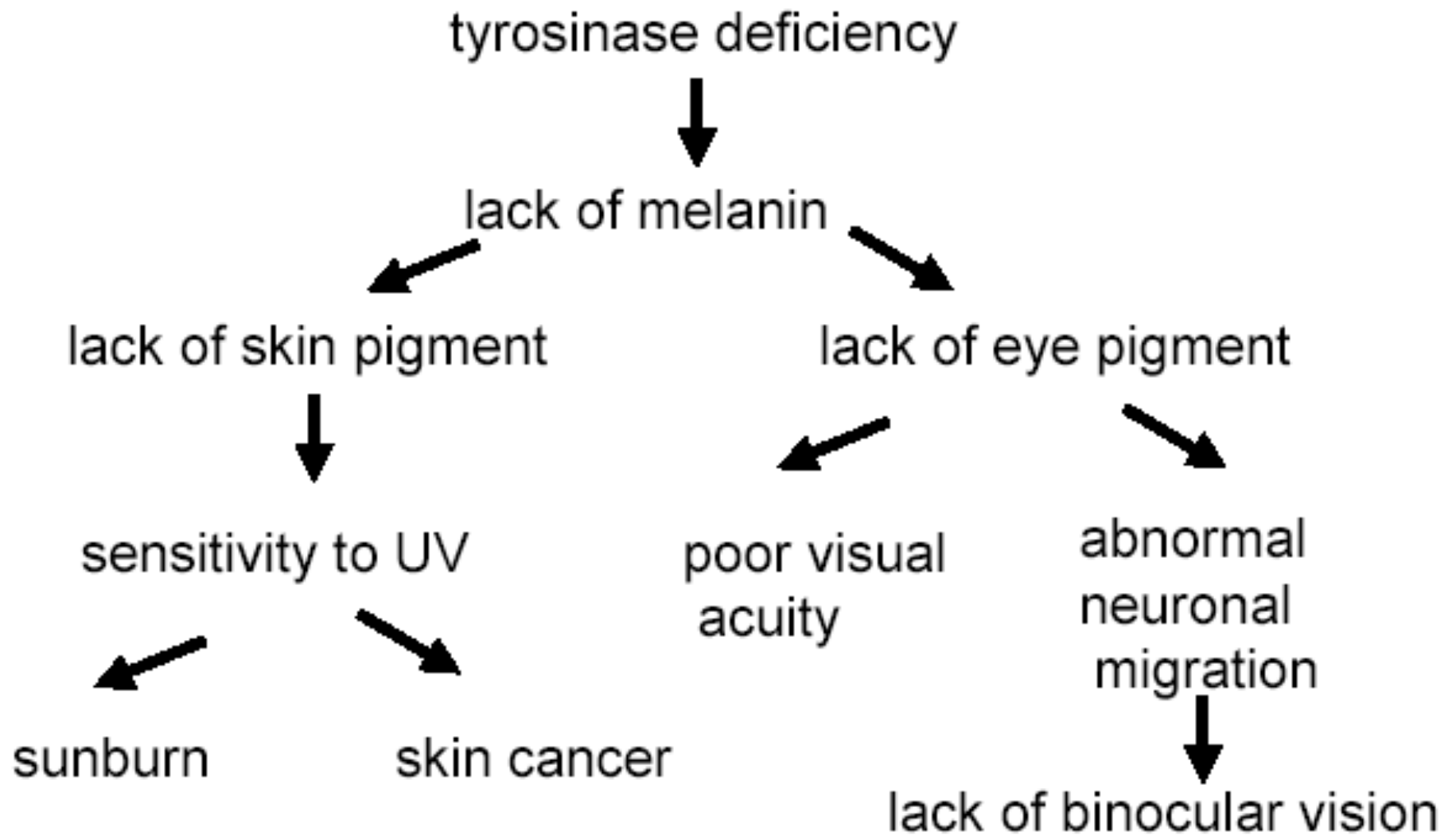


Pleiotropic mutations have effects on several different characters. In this case the dominance relationships depend on the phenotype examined.

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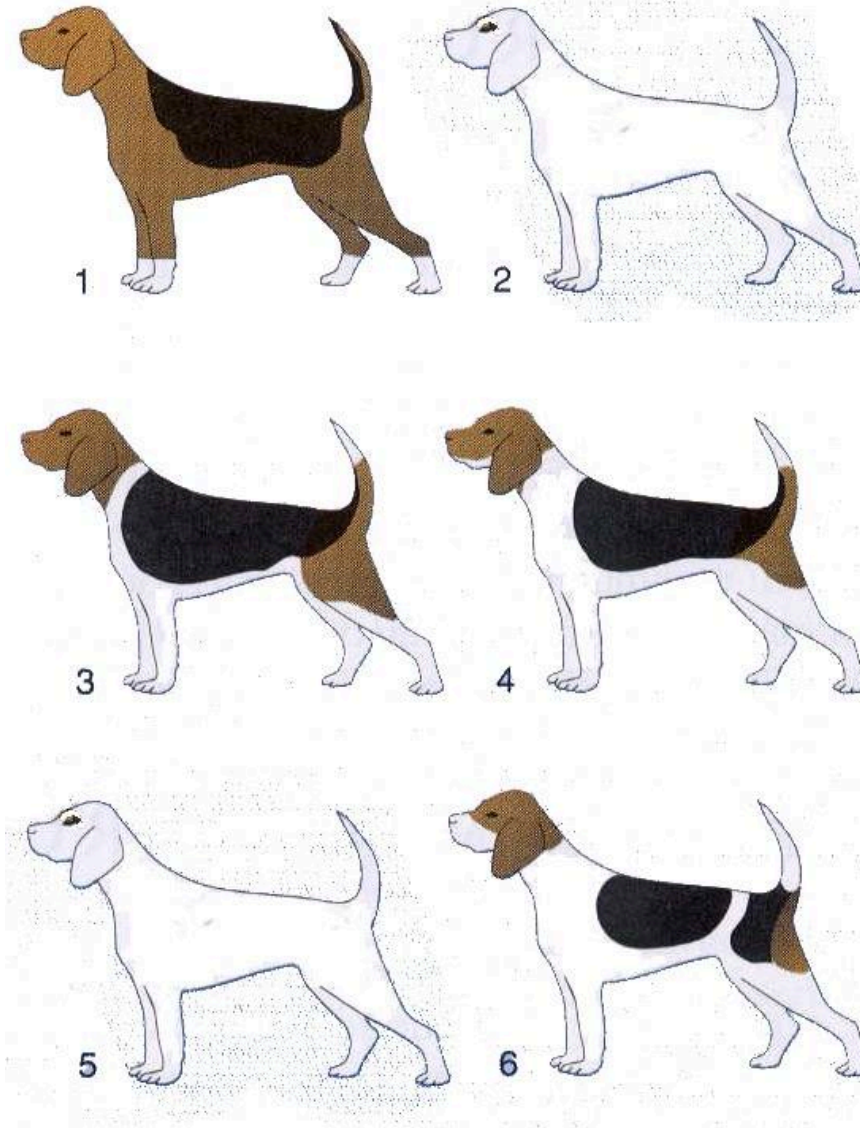
Phenotypes at Different Levels of Analysis	Normal AA	Carrier AS	Diseased SS	Dominance Relations at Each Level of Analysis
β -Globin polypeptide production				A and S are codominant
Red blood cell shape at sea level				A is dominant S is recessive
Red blood cell concentration at sea level	Normal cells 	Normal 	Lower 	
Red blood cell shape at high altitudes				A and S show incomplete dominance
Red blood cell concentration at high altitudes	Normal 	Lower 	Very low, anemia 	
Susceptibility to malaria	 Normal susceptibility	 Resistant	 Resistant	S is dominant A is recessive

PLEIOTROPY



Penetrance

...the frequency at which individuals with a given genotype manifest a specific phenotype.



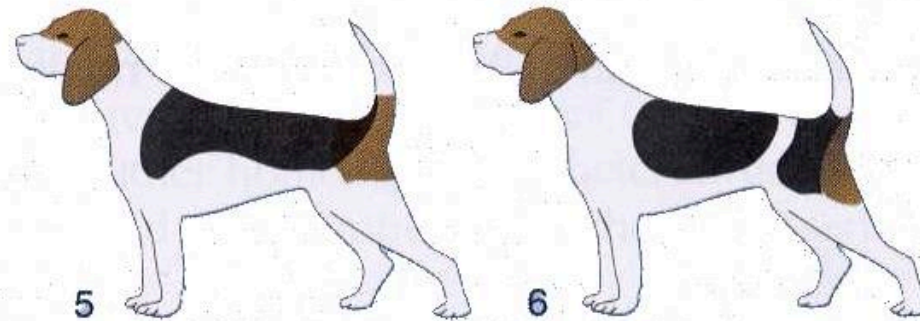
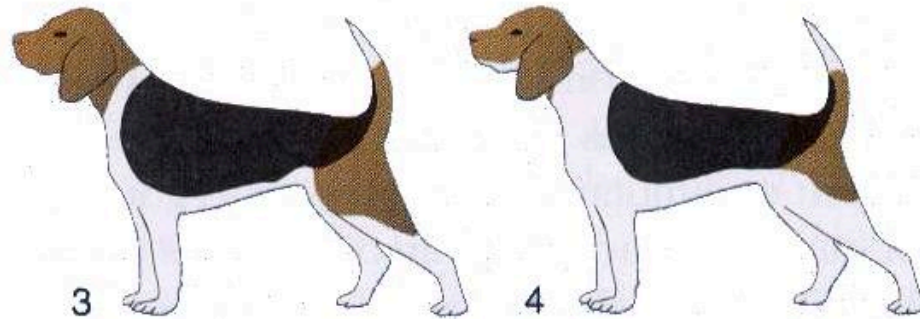
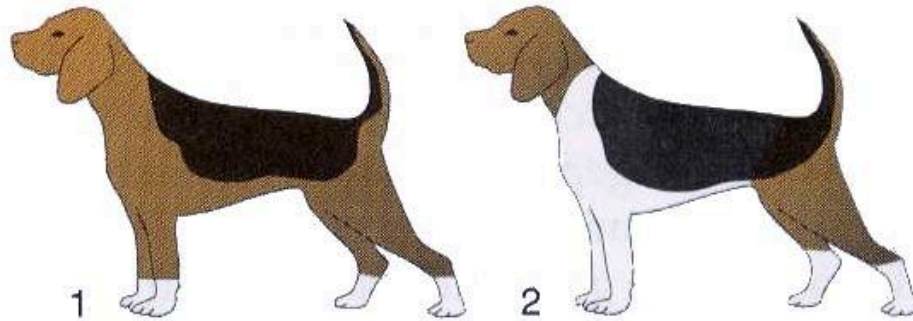
- 4 of 6 dogs, or 66% of the population shows the phenotype, at some level,

- penetrance is usually referred to as a percentage.

all the same genotype

Expressivity

...the degree or range within which a phenotype of a specific genotype is expressed



range of phenotypes

Phenotype strength may depend on the presence or absence of specific alleles at other genetic loci.

all the same genotype

**Himalayan rabbits carry a temperature-sensitive form of tyrosinase
Required for melanin synthesis**



**...Or phenotype may
depend on the
environment**

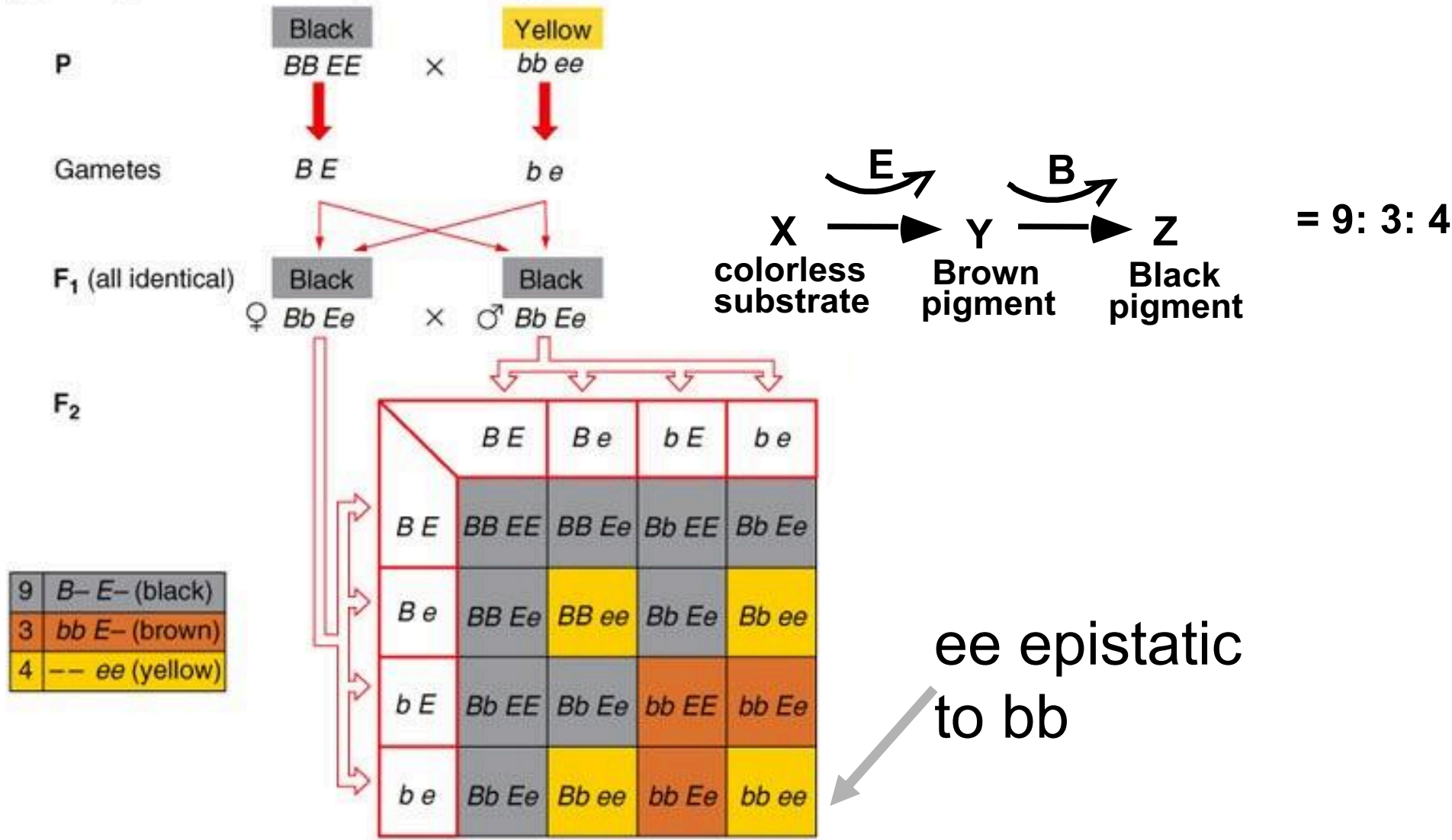
Epistasis and puppies



In epistasis the genotype at one locus masks the ability to determine the genotype at a second locus. Epistasis relationships are often found in biochemical pathways.

Recessive epistasis

Disruption of an early step in the pathway blocks the ability to determine the genotype with respect to genes involved in later steps



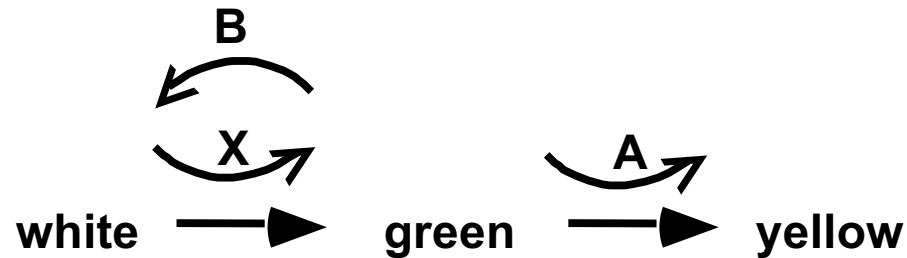
The functions of E and B

- E encodes melanocortin receptor 1 -- promotes production of eumelanin leaving dogs black or chocolate brown; null allele (e) means mc1r is not functional and dogs only produce phaeomelanin (yellow)
- B regulates distribution of melanin in hair shaft. B promotes dense packing of melanin (black) and b promotes less dense packing (chocolate brown).

Dominant epistasis produces ratios of 12:3:1 or 13:3

12:3:1---A dominant allele in one gene, B, masks the contribution of a second gene, A, (blocks color formation) regardless of the A allele. Thus in bb individuals the A- and aa genotypes can express themselves, resulting in a total of three different phenotypes.

Dominant epistasis: one possible mechanism
= 12 (white): 3(yellow): 1(green)

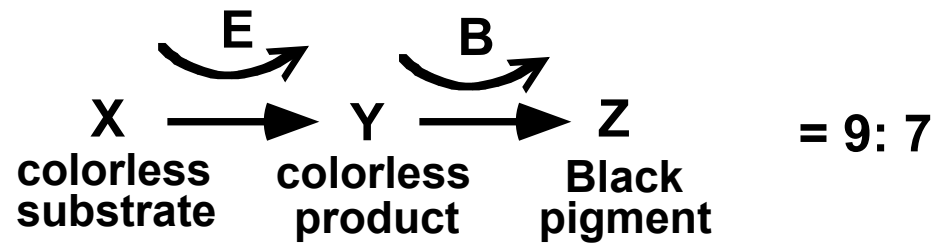


B acts to inhibit the processing of a white precursor into a green product.

In the absence of the B gene product the A product processes the green precursor into a yellow product.

Complementary gene action

Dominant alleles from two different genes must be present to produce a particular phenotype.



	BE	Be	bE	be
BE	BBEE (Black)	BBEe (Black)	BbEE (Black)	BbEe (Black)
Be	BBeE (Black)	BBee (Golden)	BbeE (Black)	Bbee (Golden)
bE	bBEE (Black)	bBEe (Black)	bbEE (Golden)	bbEe (Golden)
be	bBeE (Black)	bBee (Golden)	bbeE (Golden)	bbee (Golden)

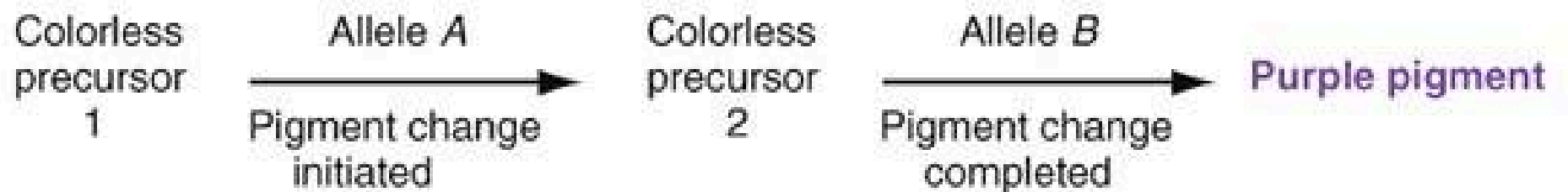
Notice that if we had a way to distinguish colorless product X from colorless product Y this would be an example of recessive epistasis.



**How many genes contribute to a process
(flower color)**



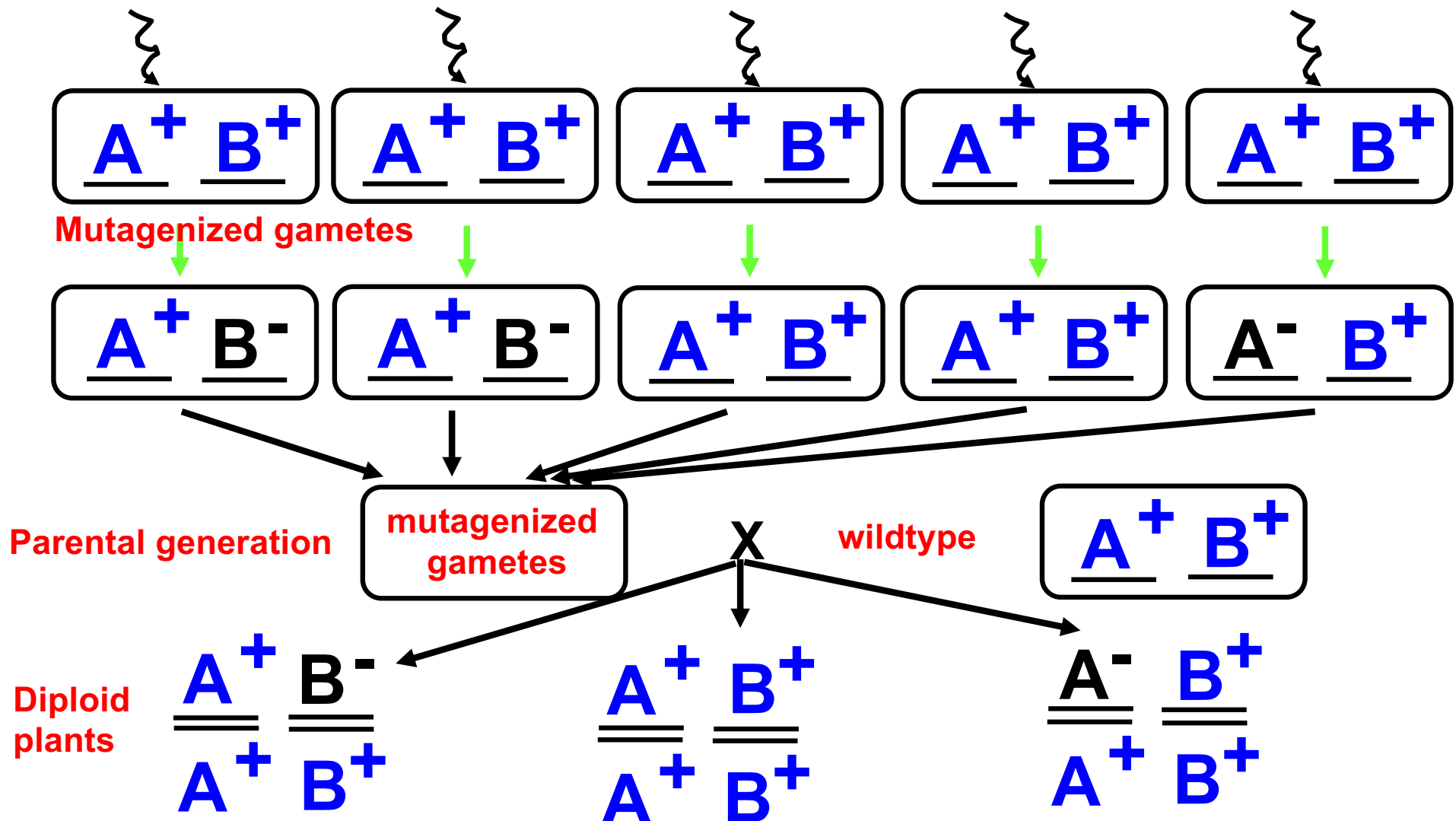
(c) A biochemical model for purple pigment production



Complementation testing 1

How many genes are required to generate a particular phenotype such as flower color ?

1. Carry out a genetic screen for mutations that lack flower color.
2. Carry out complementation testing to determine which mutations effect the same gene.



Complementation testing 2

F1 Diploid plants

$$\begin{array}{c} \underline{\underline{A^-}} \quad \underline{\underline{B^+}} \\ \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \end{array}$$

$$\begin{array}{c} \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \\ \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \end{array}$$

$$\begin{array}{c} \underline{\underline{A^+}} \quad \underline{\underline{B^-}} \\ \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \end{array}$$

Self cross genotypes

$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^-}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^-}} \quad \underline{\underline{B^+}}$
$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^-}} \quad \underline{\underline{B^+}}$
1:	2:	1

$$\begin{array}{c} \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \\ \underline{\underline{A^+}} \quad \underline{\underline{B^+}} \end{array}$$

$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^+}} \quad \underline{\underline{B^-}}$
$\underline{\underline{A^+}} \quad \underline{\underline{B^+}}$	$\underline{\underline{A^+}} \quad \underline{\underline{B^-}}$	$\underline{\underline{A^+}} \quad \underline{\underline{B^-}}$
1:	2:	1

F2

White X white =
True breeding stock

The complementation test

$$\begin{array}{c} \underline{\underline{A^-}} \quad \underline{\underline{B^+}} \\ \underline{\underline{A^-}} \quad \underline{\underline{B^+}} \end{array}$$

X

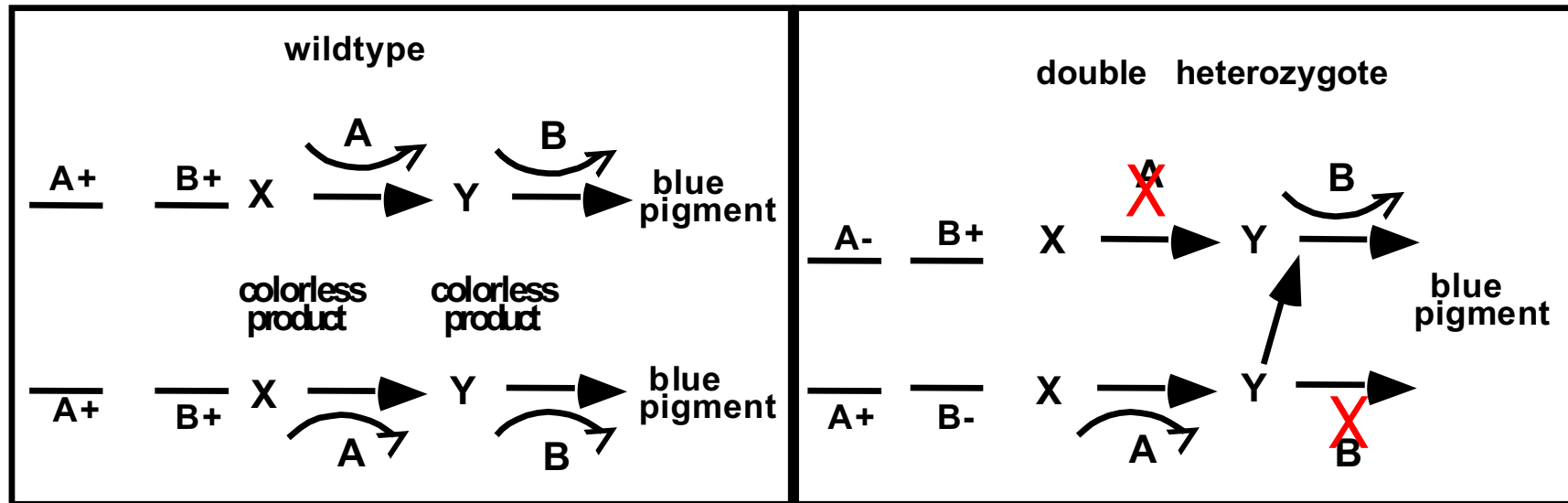
$$\begin{array}{c} \underline{\underline{A^+}} \quad \underline{\underline{B^-}} \\ \underline{\underline{A^+}} \quad \underline{\underline{B^-}} \end{array}$$

$$\begin{array}{c} \underline{\underline{A^+}} \quad \underline{\underline{B^-}} \\ \underline{\underline{A^-}} \quad \underline{\underline{B^+}} \end{array}$$

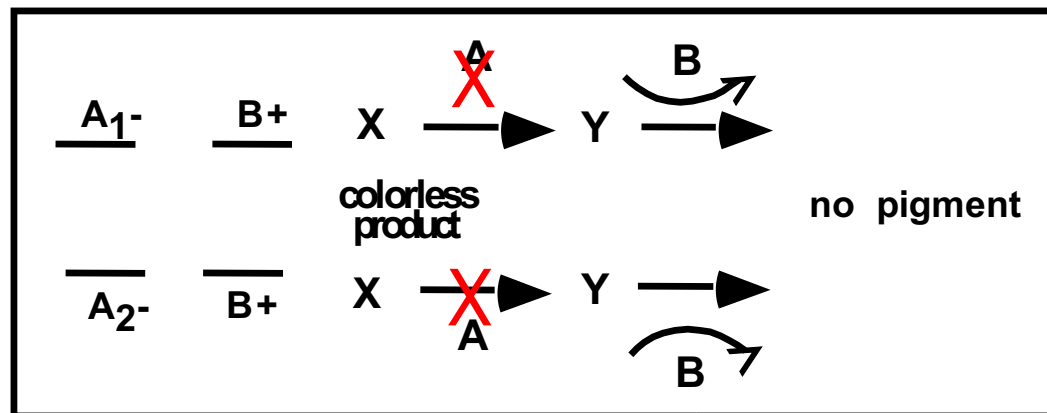
= all blue progeny = complementation
(mutations are in different genes =
complementation groups)

Complementation testing 3

$$\frac{A^-}{A^-} \frac{B^+}{B^+} \times \frac{A^+}{A^+} \frac{B^-}{B^-} = \text{all blue progeny} = \text{complementation}$$

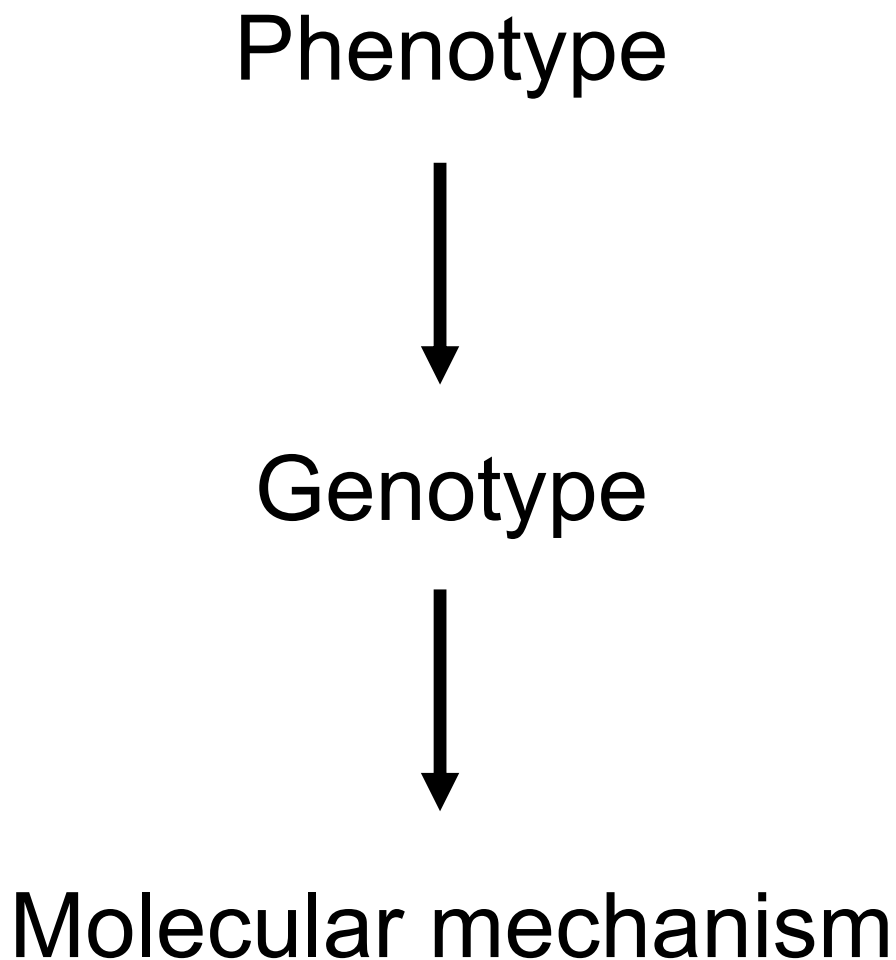


$$\frac{A_1^-}{A_1^-} \frac{B^+}{B^+} \times \frac{A_2^-}{A_2^-} \frac{B^+}{B^+} = \text{all white progeny} = \text{noncomplementation}$$



How many different genes are required for a particular trait?

Genetic screens and complementation testing.



Matings between different white-flowered lines produce purple progeny

↓

Mutations at different genes

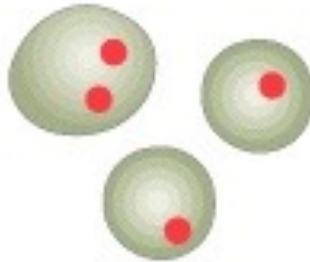
↓

Affected genes encode different proteins that are required for the same biochemical pathway

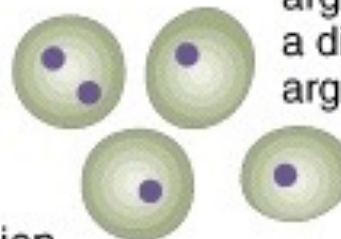
```
graph TD; A[Matings between different white-flowered lines produce purple progeny] --> B[Mutations at different genes]; B --> C[Affected genes encode different proteins that are required for the same biochemical pathway];
```

Complementation can occur during cell fusion.

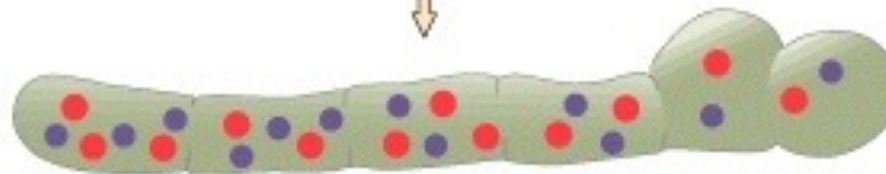
arg-1 cells, defective for one specific enzyme in arginine synthetic pathway



arg-2 cells, defective for a different enzyme in arginine synthetic pathway



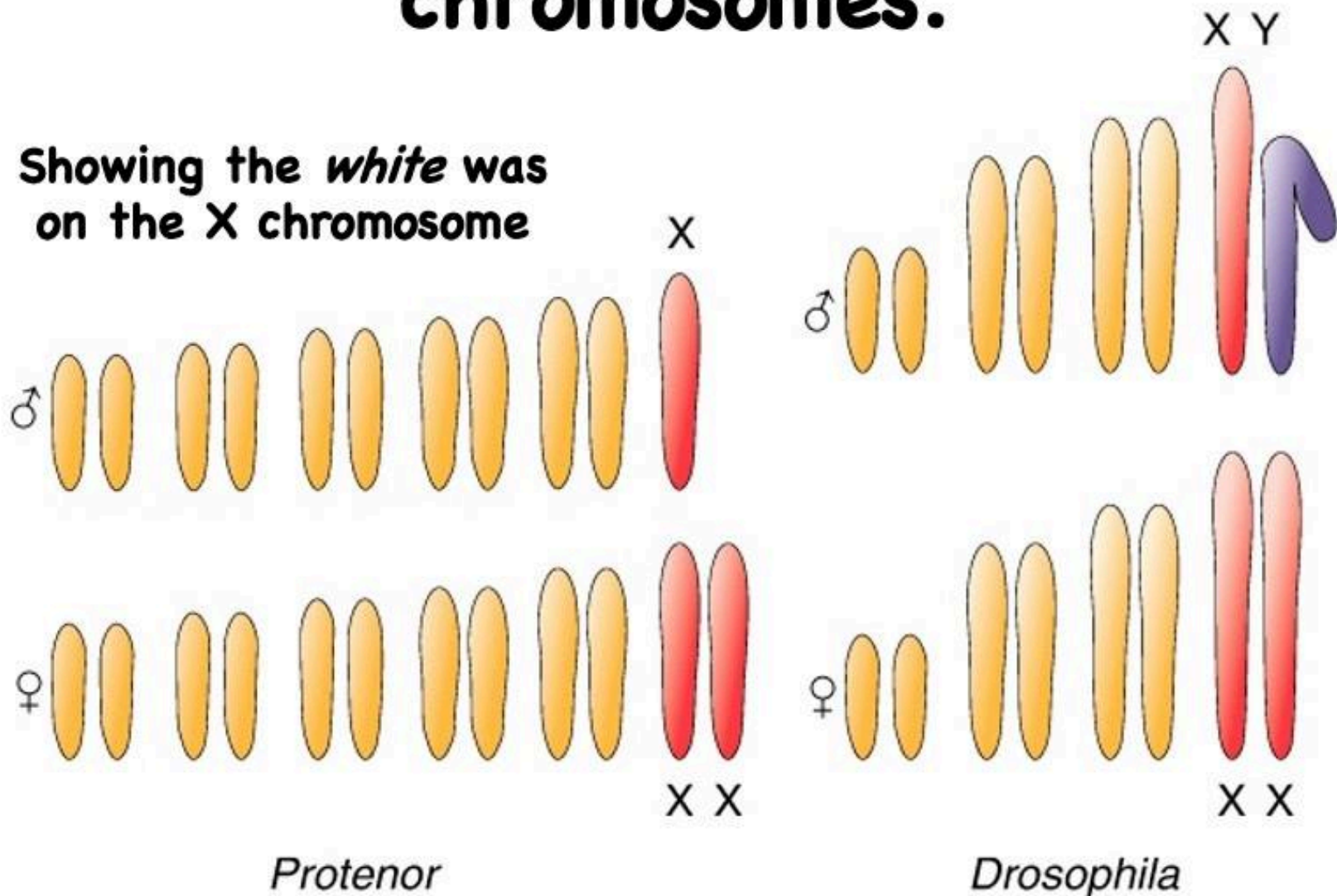
Fusion



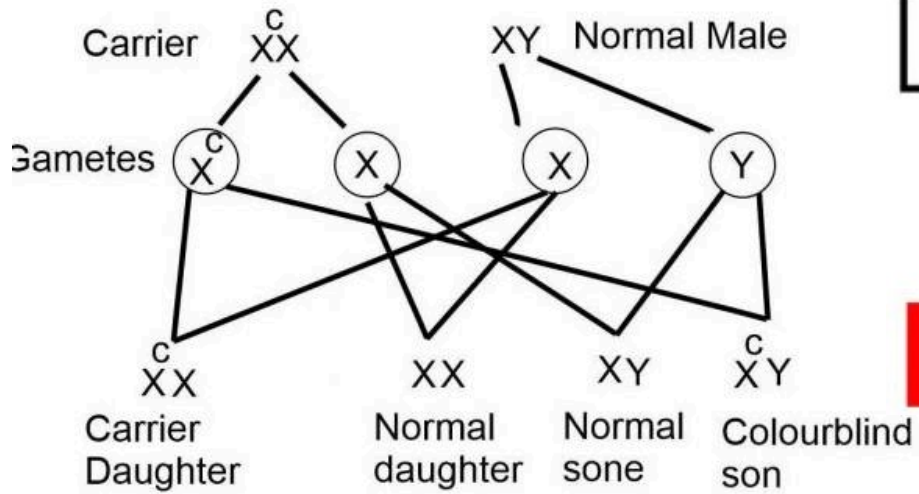
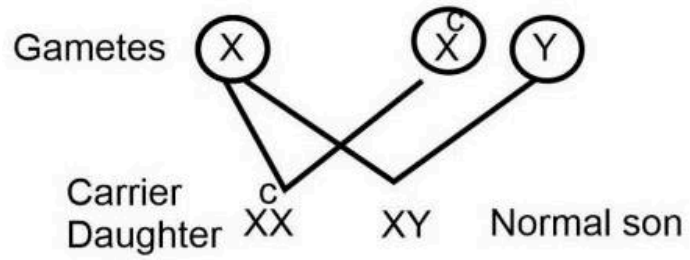
Heterokaryon grows without arginine

Organisms often have a pair of heteromorphous (look different) chromosomes.

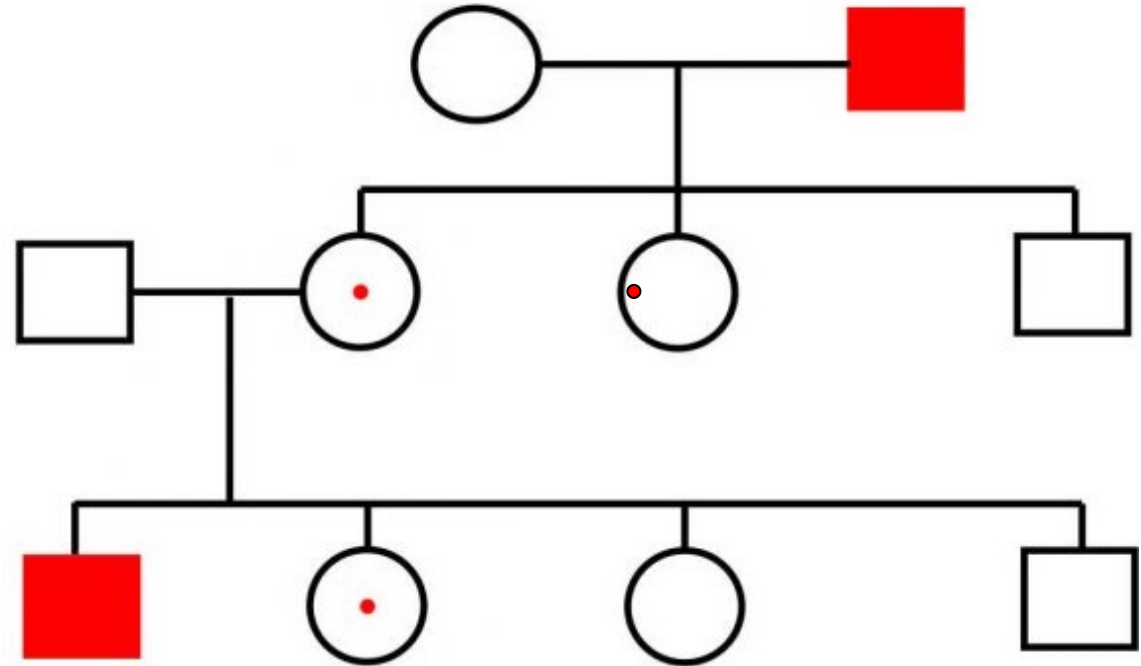
Showing the *white* was on the X chromosome



Parents XX x $\overset{c}{X}Y$
 Female Male (Colourblind)



Color blindness. X-linked recessive



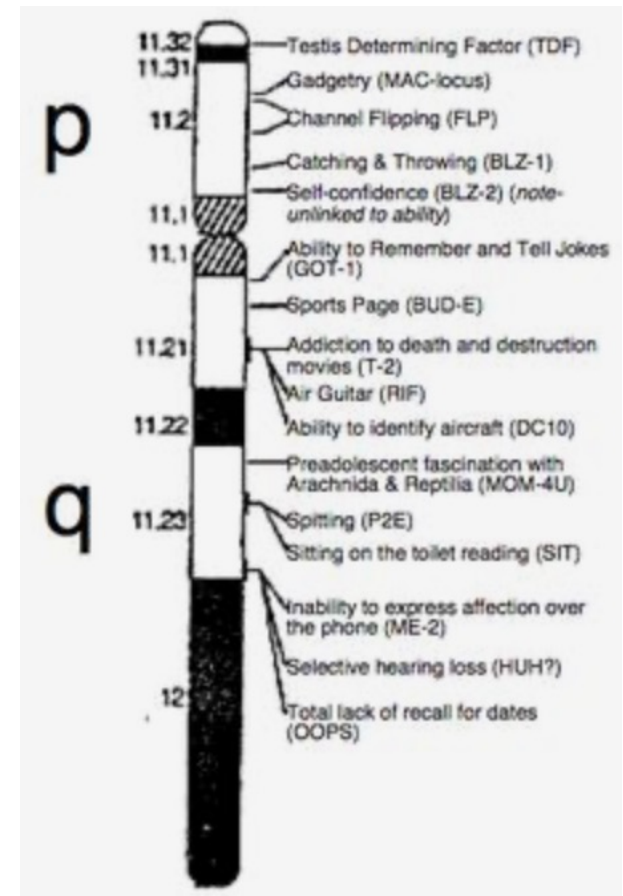
In *Drosophila* (an insect) is not just the number of X chromosomes, but instead the ratio of X chromosomes to the number of autosomal sets (A) that determines sex in the fly. $X:A+0.5$ in the male. $X:A = 1$ in the female

XY AA	male
XX AA	female
XO AA	male (sterile)
XXX AAA	female
XYY AAA	male
XXY AAA	intersex

The mammalian Y chromosome contains very few genes SRY

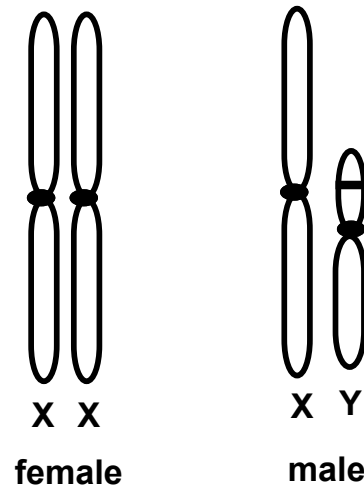
+ Genes needed for sperm development

The Y chromosome is necessary and sufficient for male development!



Sex determination in mammals

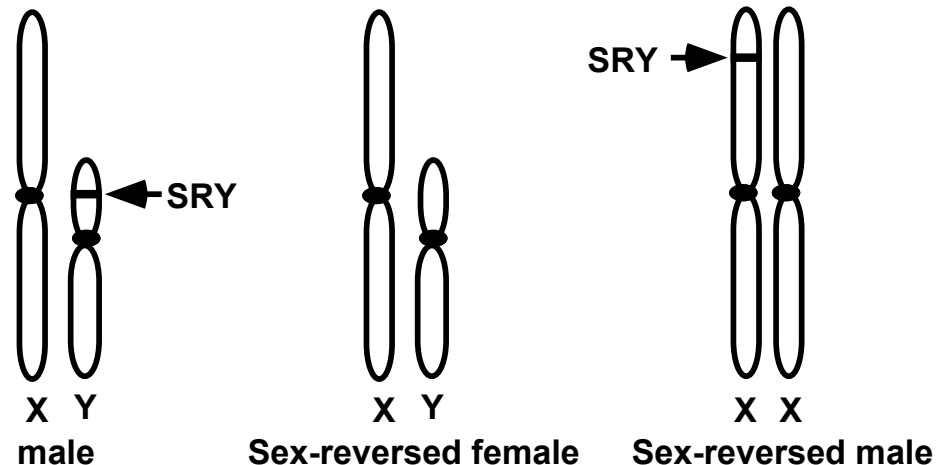
The Y chromosome and the SRY locus



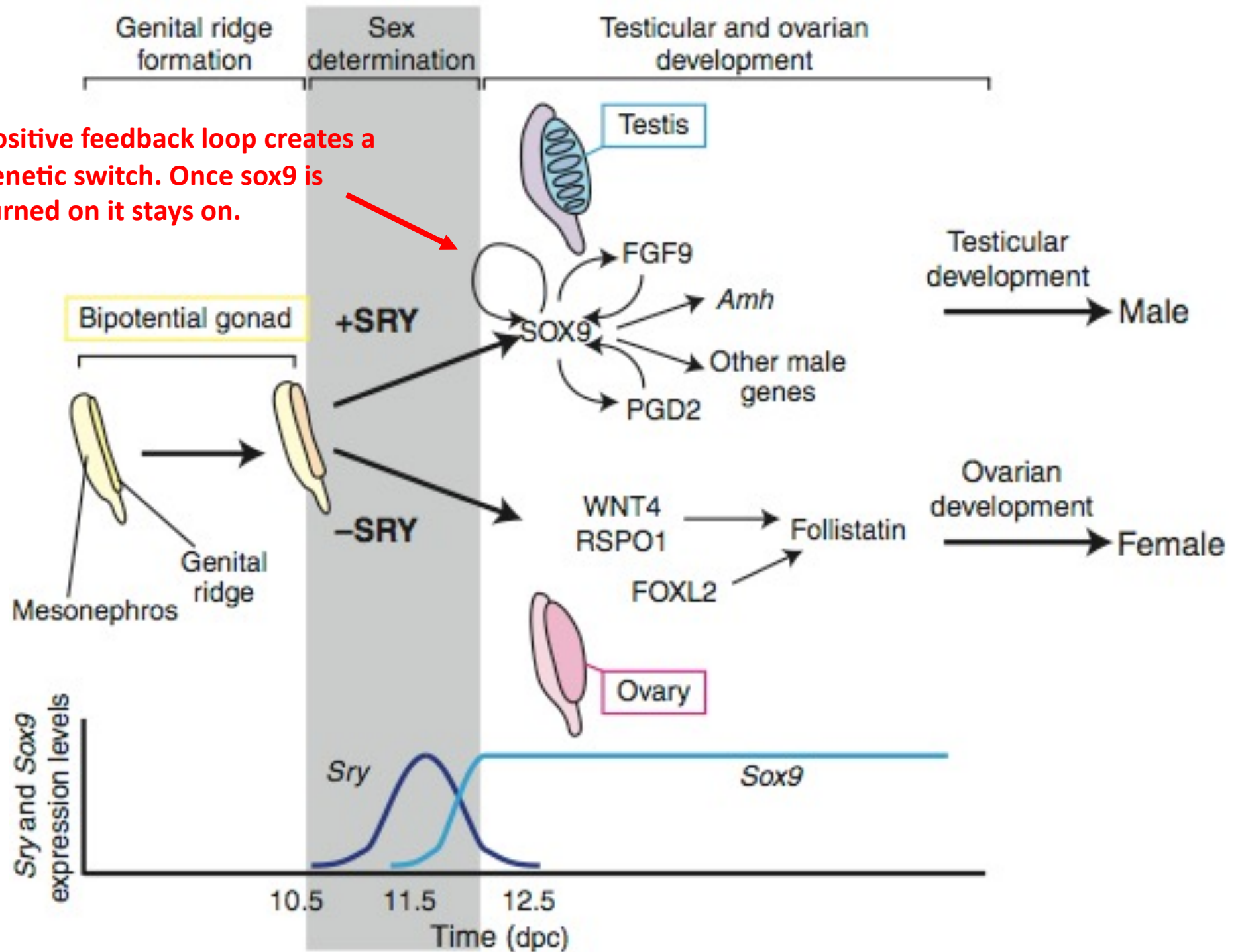
XX	female
XY	male
XXX	female (normal)
XXY	male (Klinefelter's syndrome)
XO	female (Turner's syndrome)

The SRY locus, present on the Y chromosome, determines sex in mammals.

1. XX individuals that show sex reversal (they are phenotypically male) have a small fragment of the Y chromosome moved to the X chromosome.
2. XY individuals that lack the SRY region of the Y chromosome are phenotypically female.
3. Generation of mice that are XX, but that have the SRY gene introduced onto their X chromosome are phenotypically male.



Positive feedback loop creates a genetic switch. Once *sox9* is turned on it stays on.



More than 1% of us have mutations that result in altered biological sex outcome

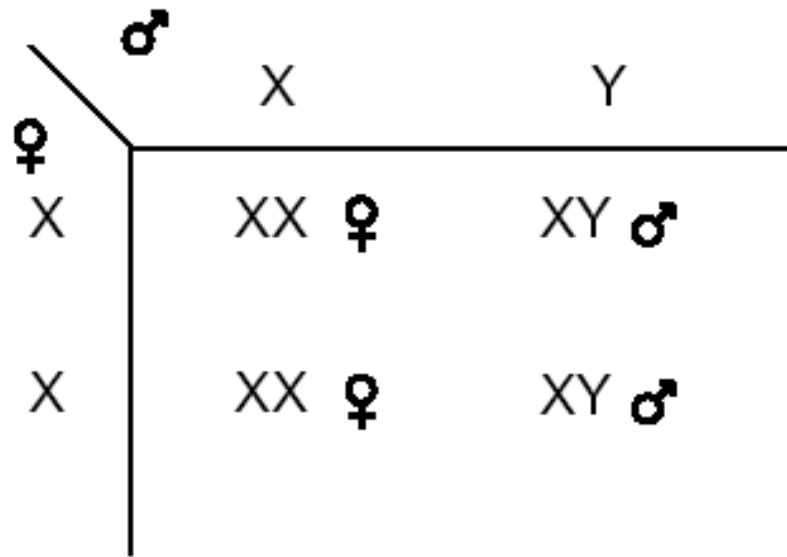
Biological outcome is different than gender

Gender refers to the socially constructed roles, behaviours, expressions and identities of girls, women, boys, men, and gender diverse people.

Fig. 1. Overview of sex determination in mice. Chronological flow of early mouse sex differentiation; the grey area indicates the period of sex determination. During mouse embryogenesis, bi-potential gonads (yellow) arise from the genital ridges by 10.5 dpc. In somatic cells of XY genital ridges, *Sry* expression (shown in dark blue beneath the schematic) starts at 10.5 dpc, reaches a peak at 11.5 dpc and then wanes by 12.5 dpc. A few hours later, *Sox9* expression (shown in light blue beneath the schematic) is upregulated to induce differentiation of Sertoli cells. *Sox9* expression peaks at 11.5-12.5 dpc, continues to be expressed postnatally and is supported by several positive-feedback loops (including FGF9, prostaglandin D2 and SOX9 itself), and SOX9 subsequently activates many male-specific genes, including *Amh*. At 12.5 dpc, testis cords have formed, and morphological differences between testis (blue) and ovary (pink) are evident. In the absence of SRY, genes such as *Wnt4*, *Rspo1* and *Foxl2* are expressed in a female-specific manner and induce ovarian development, as characterized by the expression of follistatin and many other ovary-specific genes. Abbreviations: *Amh*, anti-Müllerian hormone; dpc, days post coitum; FGF9, fibroblast growth factor 9; FOXL2, forkhead box L2; PGD2, prostaglandin D2; RSPO1, R-spondin 1; SOX9, SRY box containing gene 9; SRY, sex-determining region on the chromosome Y; WNT4, wingless-type MMTV integration site family, member 4.

Many routes to sex determination

1) XX - XY

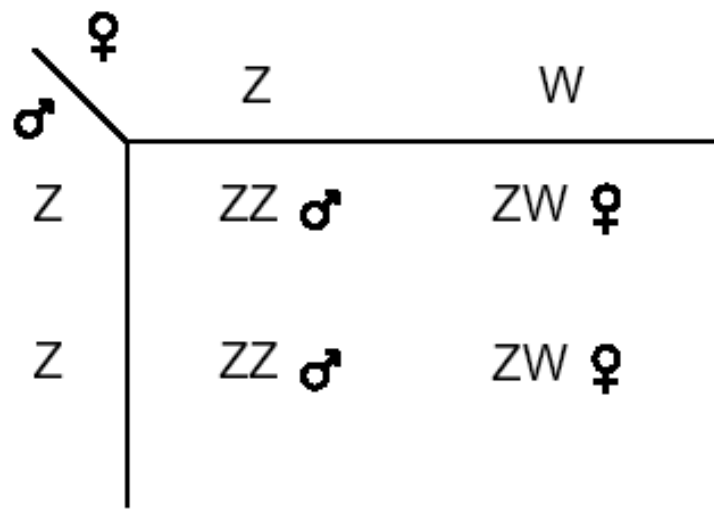


many dioecious angiosperms

many animal species

most vertebrates

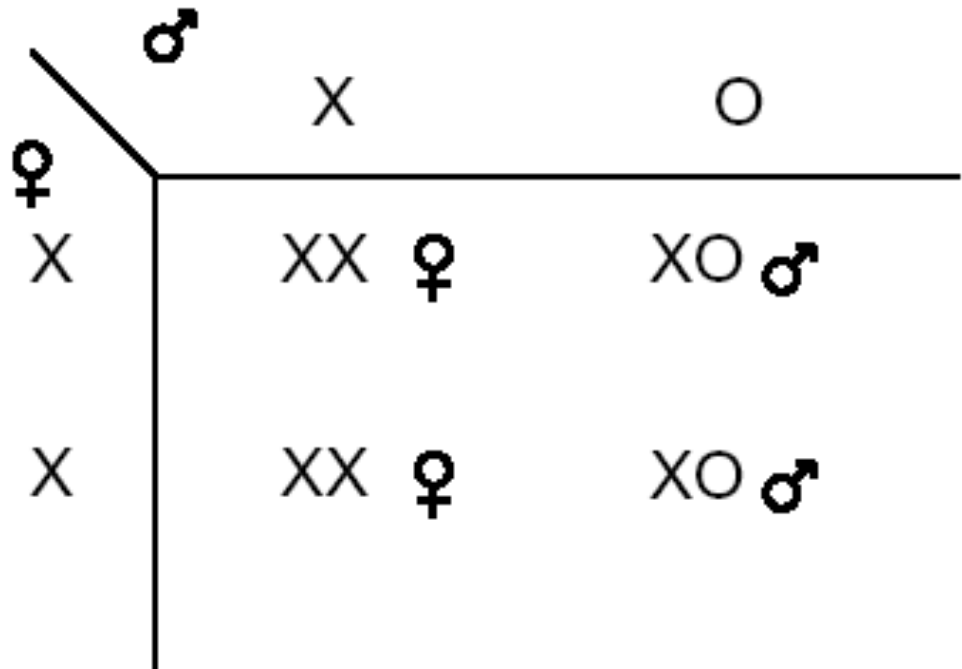
2) ZW - ZZ



some flatworms

- " crustaceans
- " insects especially lepidopterans
- " diptera (some)
- " some fish, amphibians, lizards
- " most birds

3) XX - XO



many insects

5) Arrhenotoky (haplo-diploid sex determination)



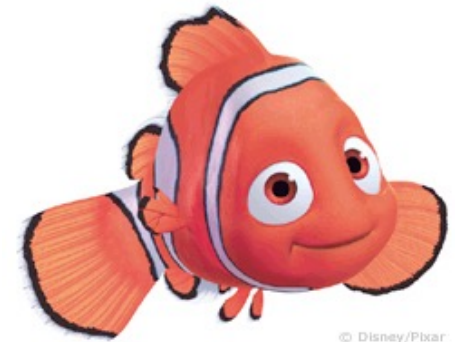
hymenoptera (ants, bees, wasps)
thysanoptera (thrips)
mites and ticks
rotifers

Yet more sex-determination systems

7) Sequential hermaphroditism

(start life as one sex, usually male, and switch later)

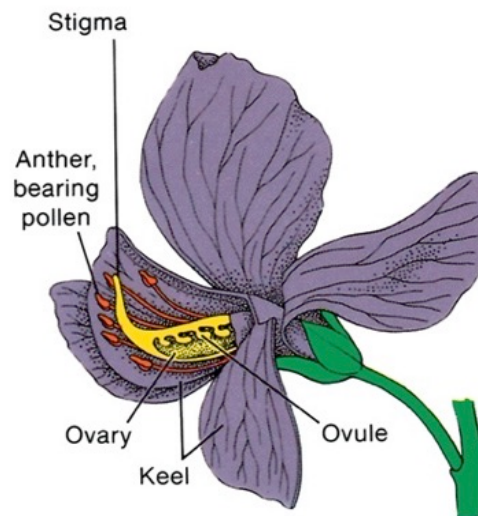
oysters, shrimp, some fish



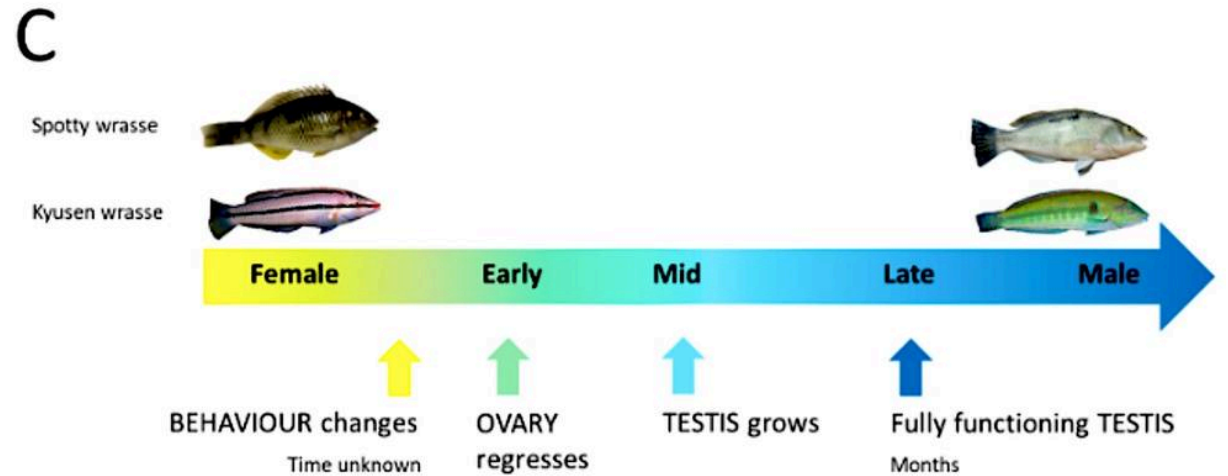
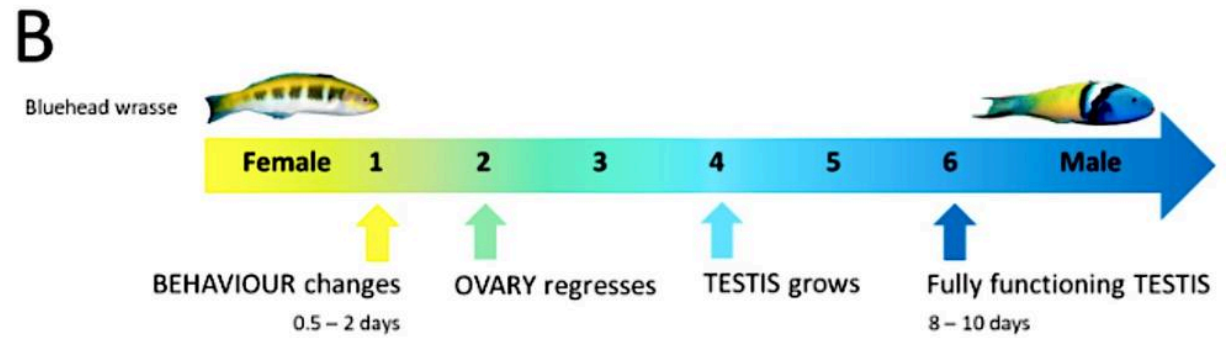
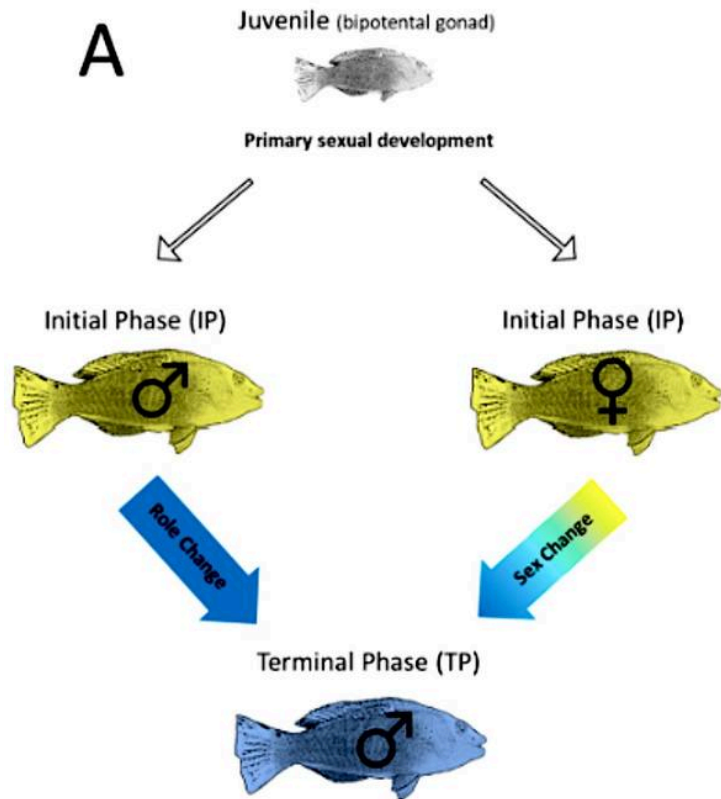
© Disney/Pixar

8) Self-sterility systems

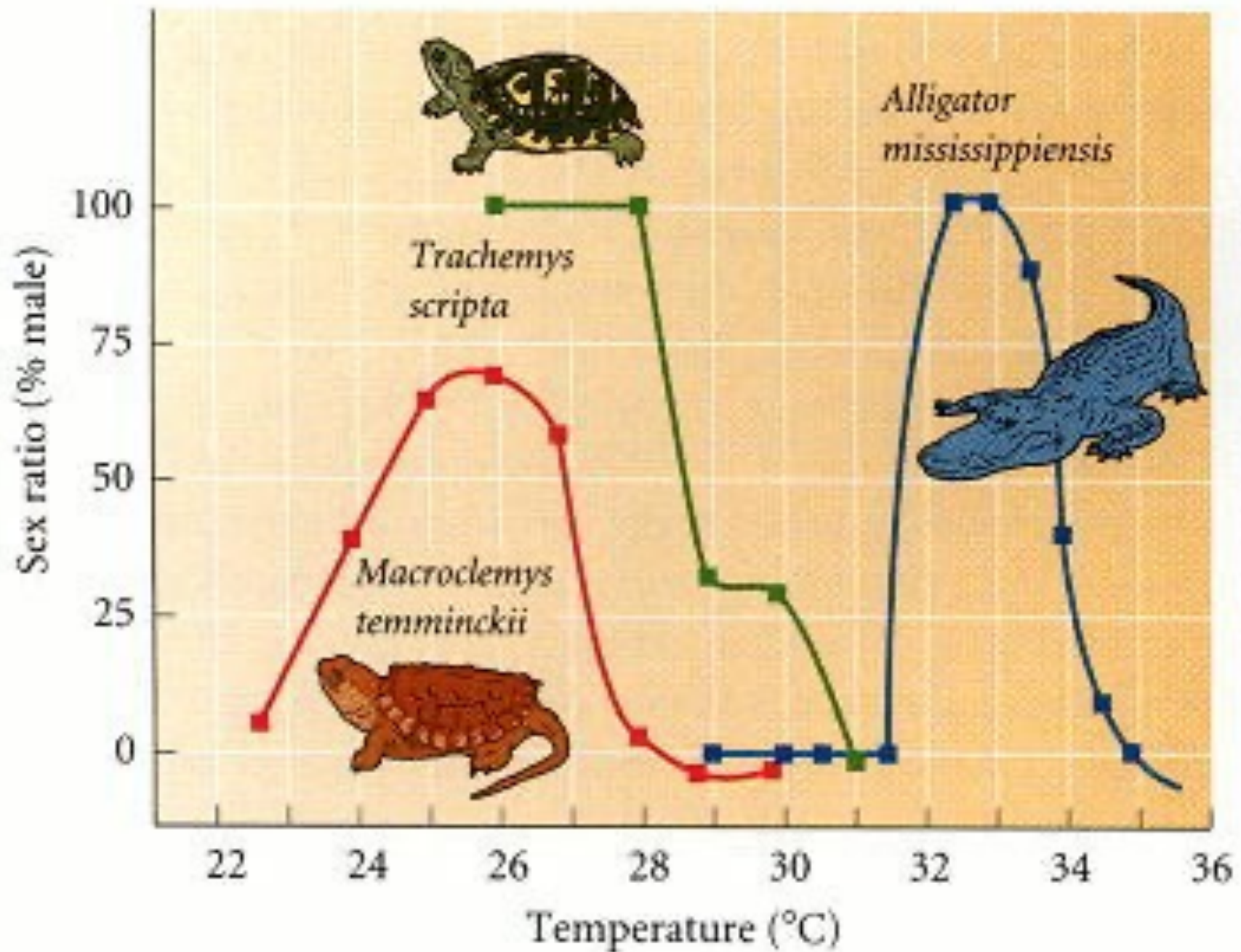
In some angiosperm plants, multiple alleles which allow pollen to succeed only if it does not contain any allele at that locus which is found in the female parent (gametophytic self-incompatibility) or the male does not contain any allele at that locus found in the female (sporophytic self-incompatibility)



Bluehead wrasse sequential sex transformation



Temperature-dependent sex determination



**Some organisms have many mating types. You must mate with
Someone who has a different type than you**



These fairy inkcap mushrooms (*Coprionellus disseminatus*), photographed in London, have 143 mating types. Each is able to reproduce with members of any of the other 142 types.

Stu's Images

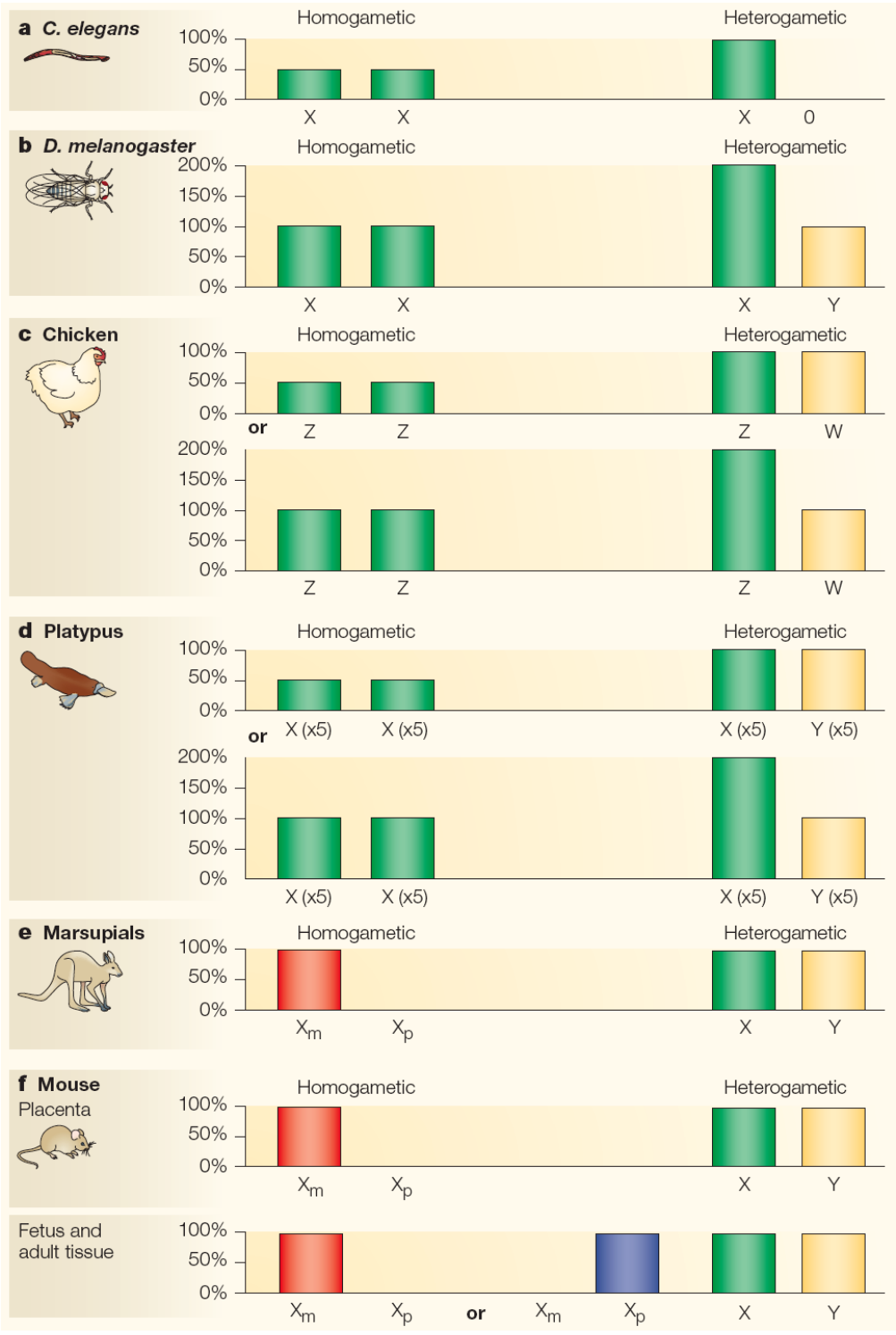
Mammals triploid for an autosomal chromosome usually die, but XXY, XYY, XXX, XXXX and XXXXX survive.

Why?

Dosage compensation

Dosage Compensation

- X chromosomes in females provide twice the genes, as in males,
 - Drosophila: female genes are expressed at 50% of the male levels,
 - Mammals: one X chromosome in females is silenced.



In *Drosophila* dosage compensation occurs through hyperactivation of the male X chromosome

sex	genotype	dose	pigment
Female	$w^a / Df(w)$	1x	1x
Female	w^a / w^a	2x	2x
Female	$w^a / w^a / Dp(w^a)$	3x	3x
Male	$w^a / -$	1x	2x
Male	$w^a / - / Dp(w^a)$	2x	4x

Visualizing dosage compensation in flies using w^a alleles of w

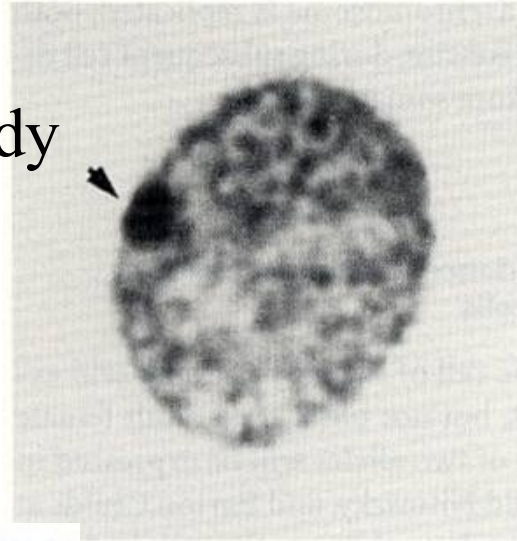
Mammalian X-inactivation: Early studies

- 1953. Two X-linked genes in the mouse, tabby and mottled, showed variegation in heterozygous females.
- 1959. XO mouse females are shown to be viable and fertile.
- 1961. Mary Lyon proposes that dosage compensation in mammals was achieved by transcriptional silencing of one X chromosome in female cells (Gene action in the X-chromosome of the mouse Nature 190: 372-373).
- Beutler et al. Two types of red blood cells in human females heterozygous for G6PDH.

Canadian Cat Scientists Sees it First

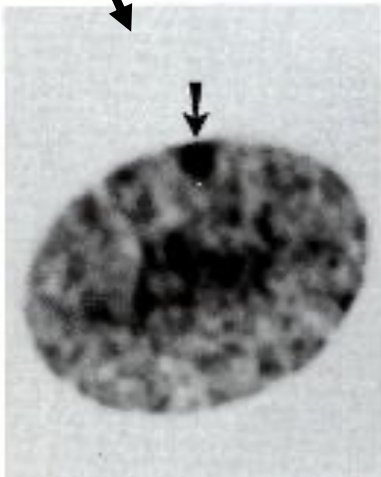
Murray Barr noted that the nuclei of female but not male cats contained a darkly stained element. This is now known as a Barr body.

Barr Body



The number of Barr bodies
varies with the number of
X chromosomes

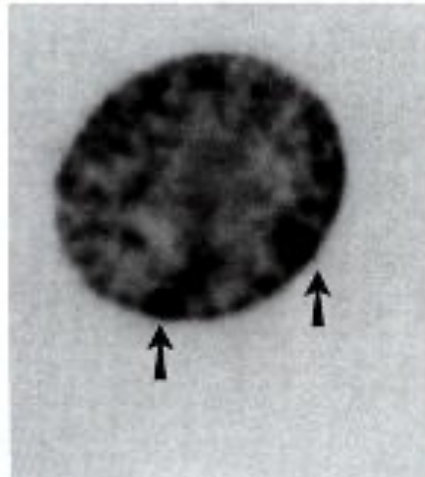
(A)



XX

One Barr body

(B)



XXX

Two Barr bodies

Karyotype

XY

XO

XX

XXX

XXXX

XXXXX

Barr bodies

0

0

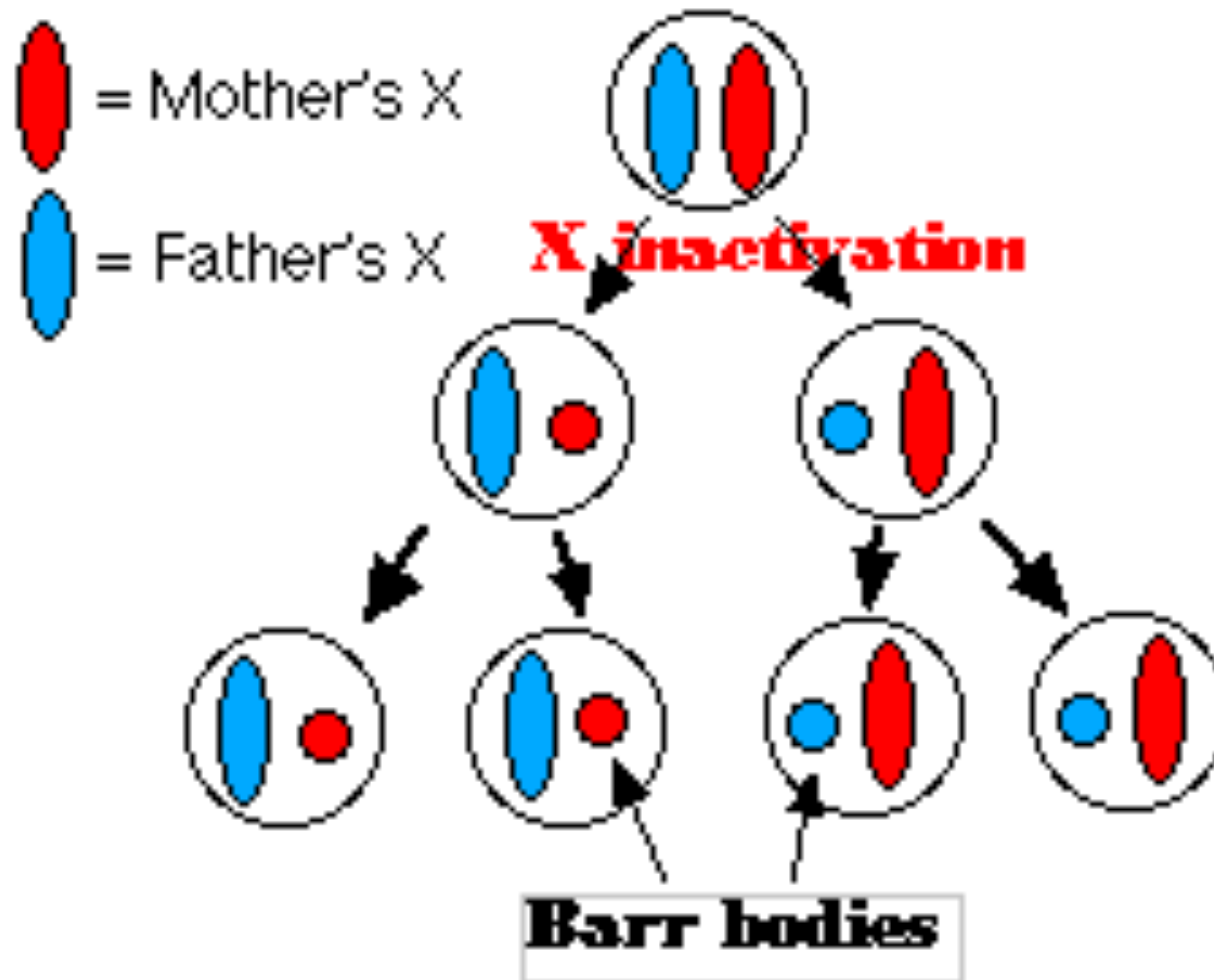
1

2

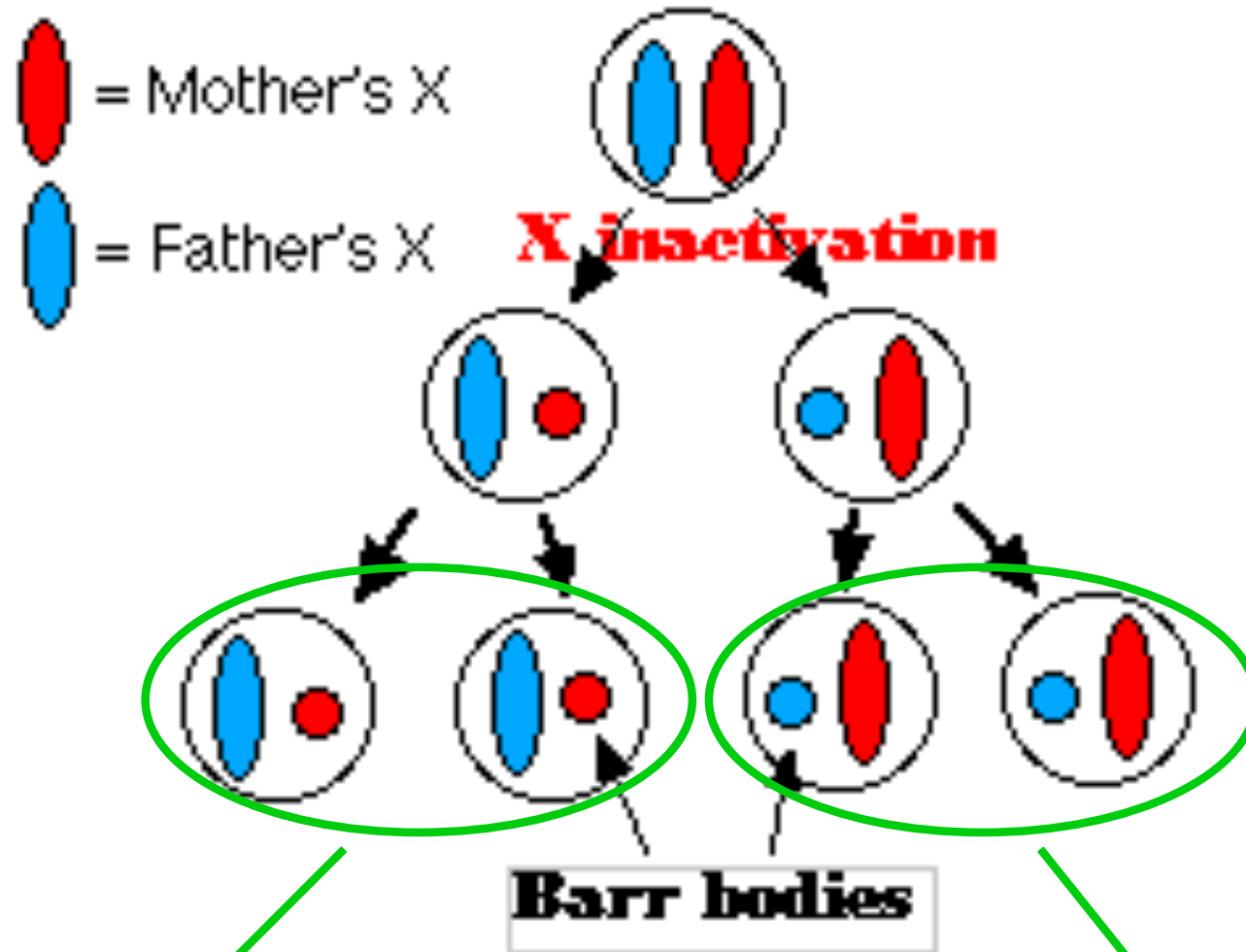
3

4

In mammalian females, early in embryonic development each cell inactivates one X chromosome



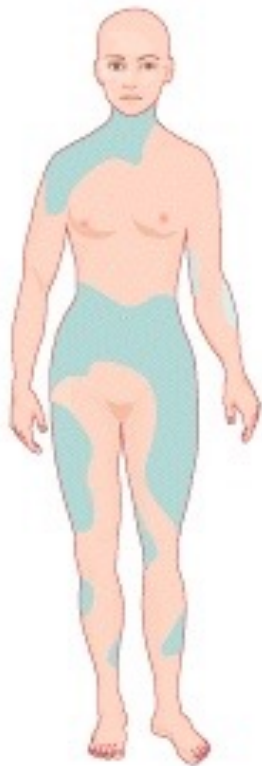
All mammalian females are mosaic!!!



These cells express only paternal X chromosome genes.

These cells express only maternal X chromosome genes.

Certain X-linked inherited traits result in
mosaicism in females.
e.g., anhidrotic ectodermal dysplasia



First
generation



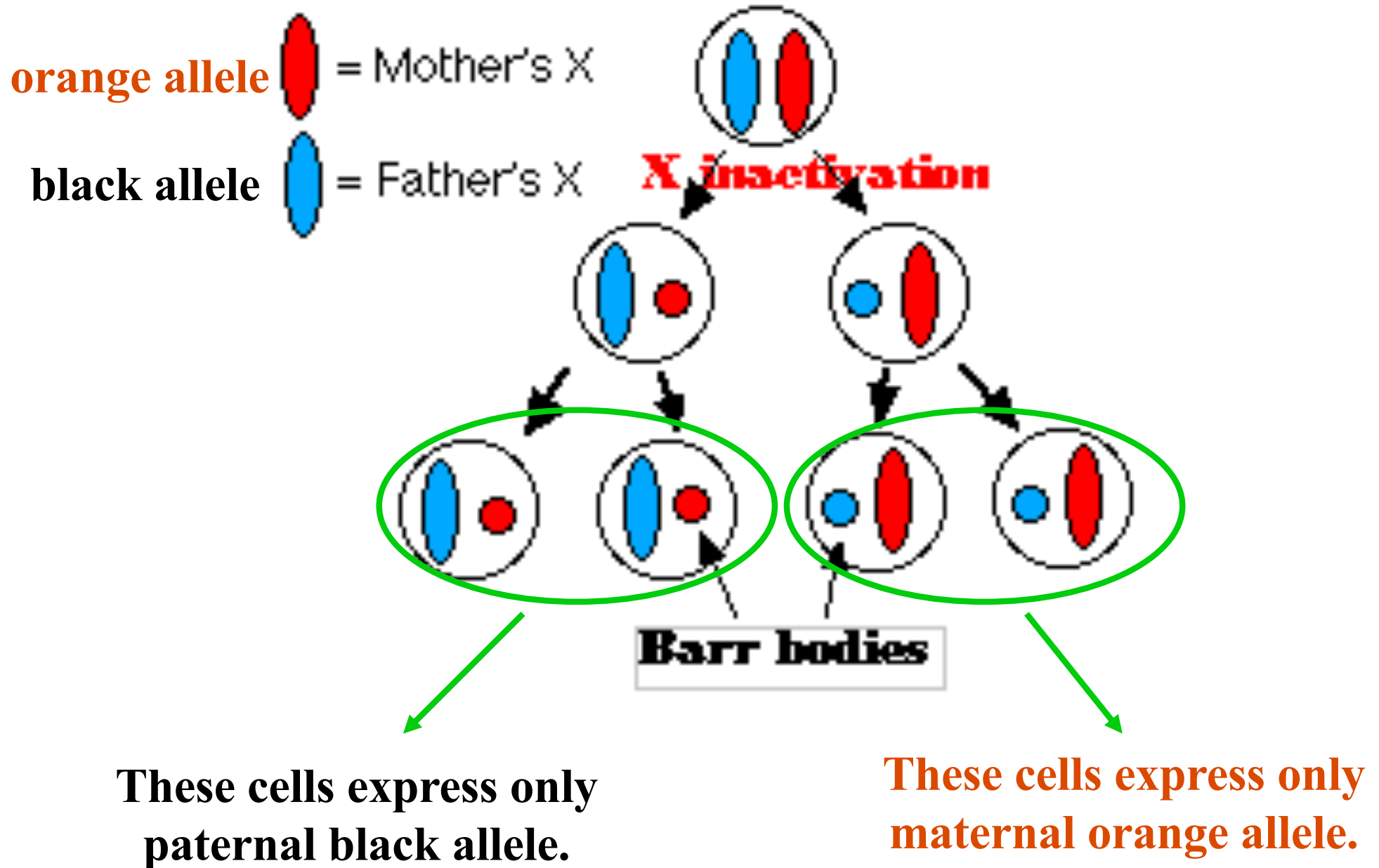
Second
generation



Third
generation



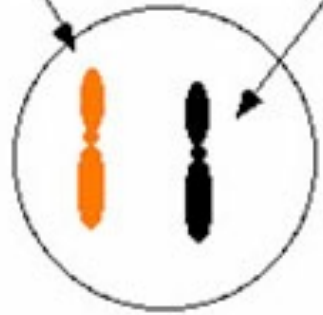
Coat color in Calico and tortoise shell cats



Tortoise shell cat

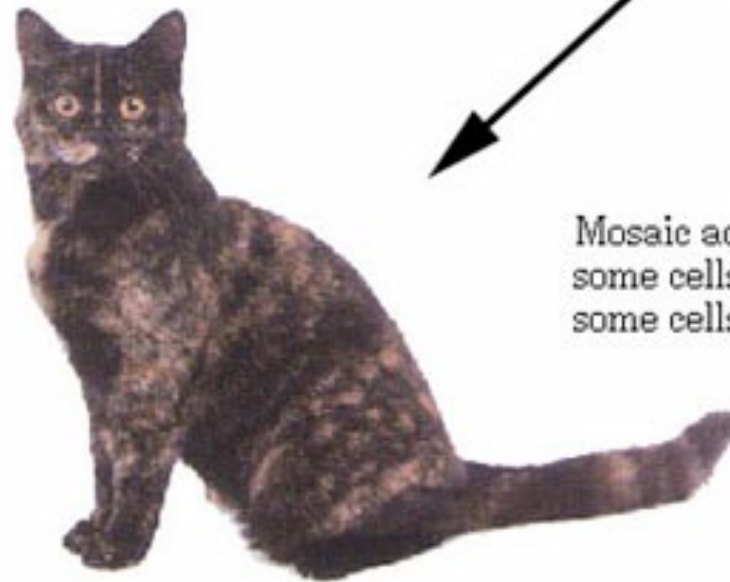
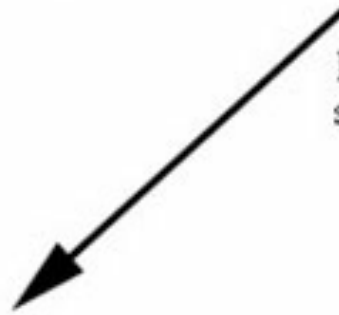
Maternal X chromosome O allele

Paternal X chromosome o allele



Random X inactivation

Portion of embryo
showing clones of cells



Mosaic adult
some cells express O
some cells express o

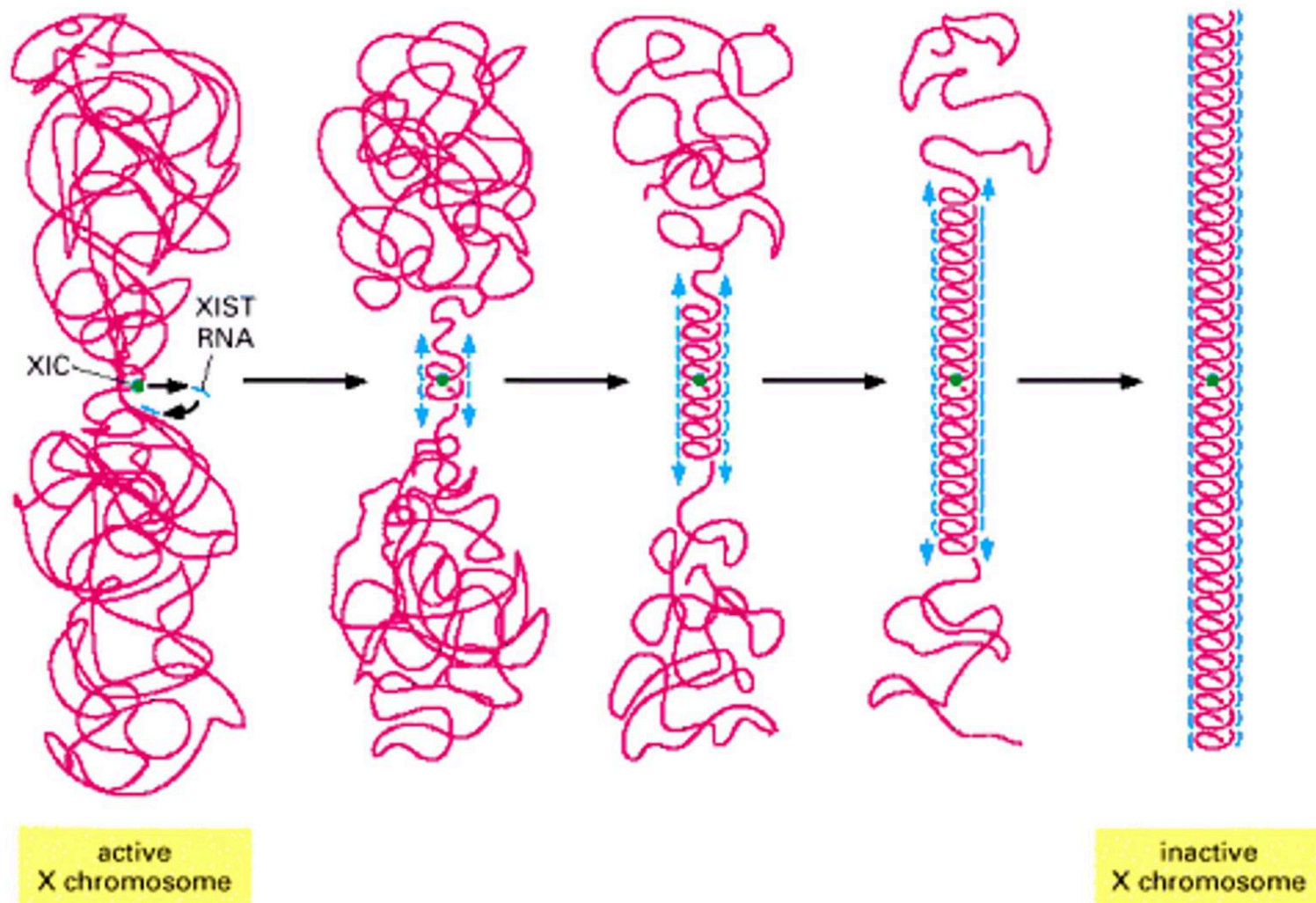
XIST

(X-inactive specific transcript)

- Identified in 1991
- 17 kb spliced, non-coding RNA
- Stable **expression only from inactive X**
- "Paints" inactive X chromosome
- Required to initiate silencing



MAMMALIAN X-CHROMOSOME INACTIVATION



X-chromosome inactivation begins with the synthesis of **XIST** (X-inactivation specific transcript) RNA from the **XIC** (X-inactivation center) locus. The association of XIST RNA with the X chromosome is correlated with the condensation of the chromosome.

After Alberts et al. - Molecular Biology of the Cell 2002



Initial steps of X inactivation occurring at the *Xic* locus

Determines which X remains active

Spread in cis of the inactive state from *Xic* to the entire X

Counting → Choice → Initiation → Spreading → Maintenance

Determines the X chr : Autosome ratio

Not quite all. See next slide

Complete and stable silencing of all X chromosomes genes. The active/inactive states are clonally inherited.

Does X inactivation affect all X chromosome genes?

No!!!!

47, XXX

48, XXXX

47, XXY

45, XO

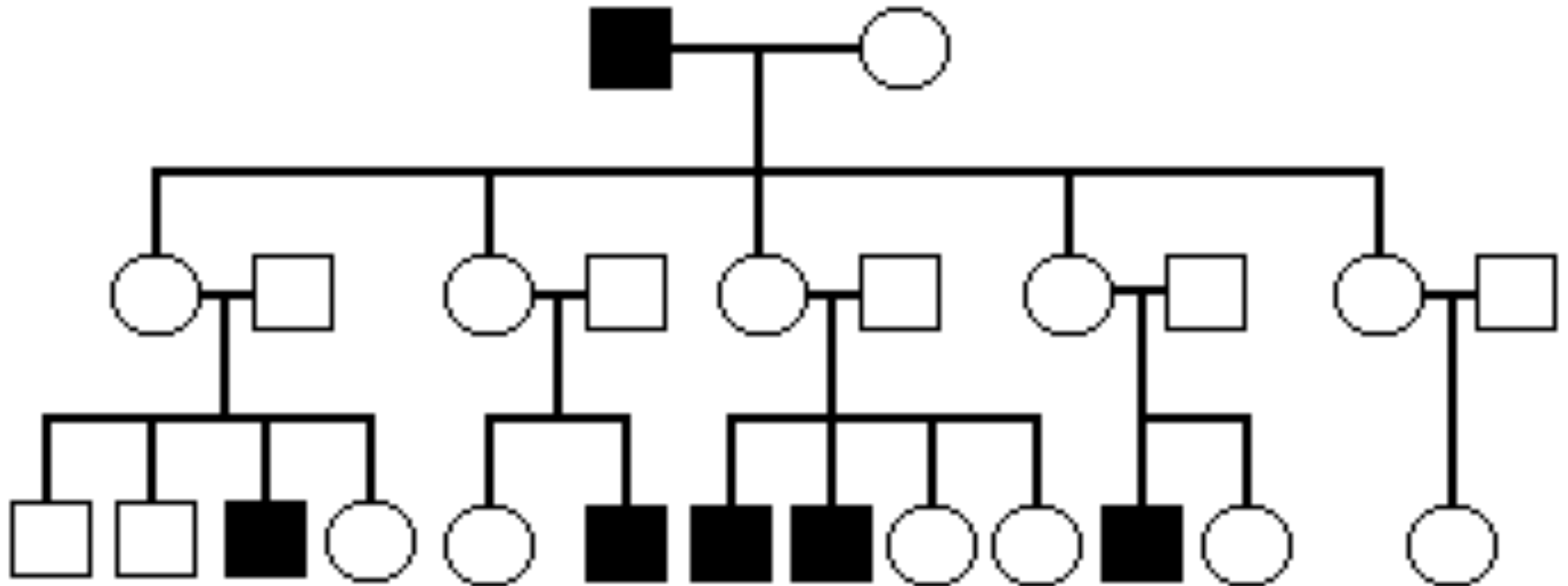
Klinefelter syndrome males

Turner syndrome females

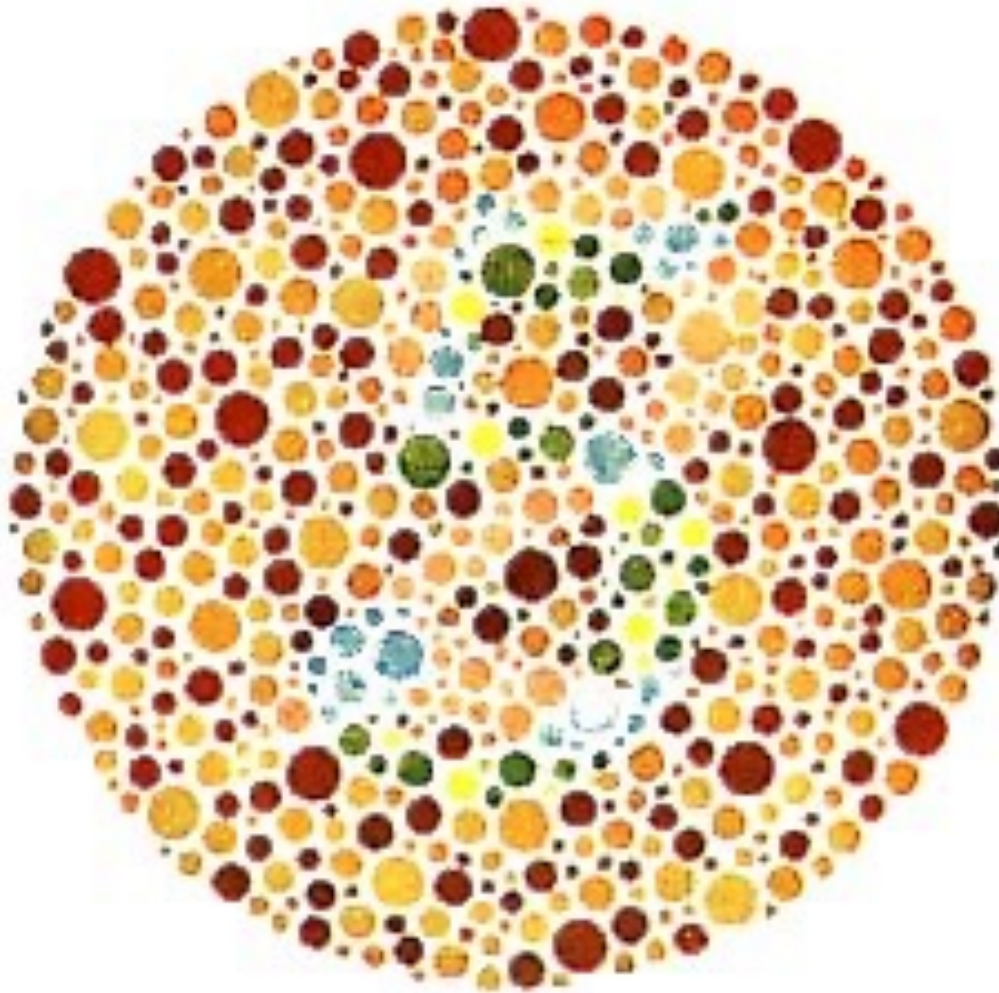
Klinefelter: Some genes in the X chromosome escape X inactivation

Turner: Some genes on the X need to be expressed in two copies

Pedigree of recessive X-linked trait



What do you see?

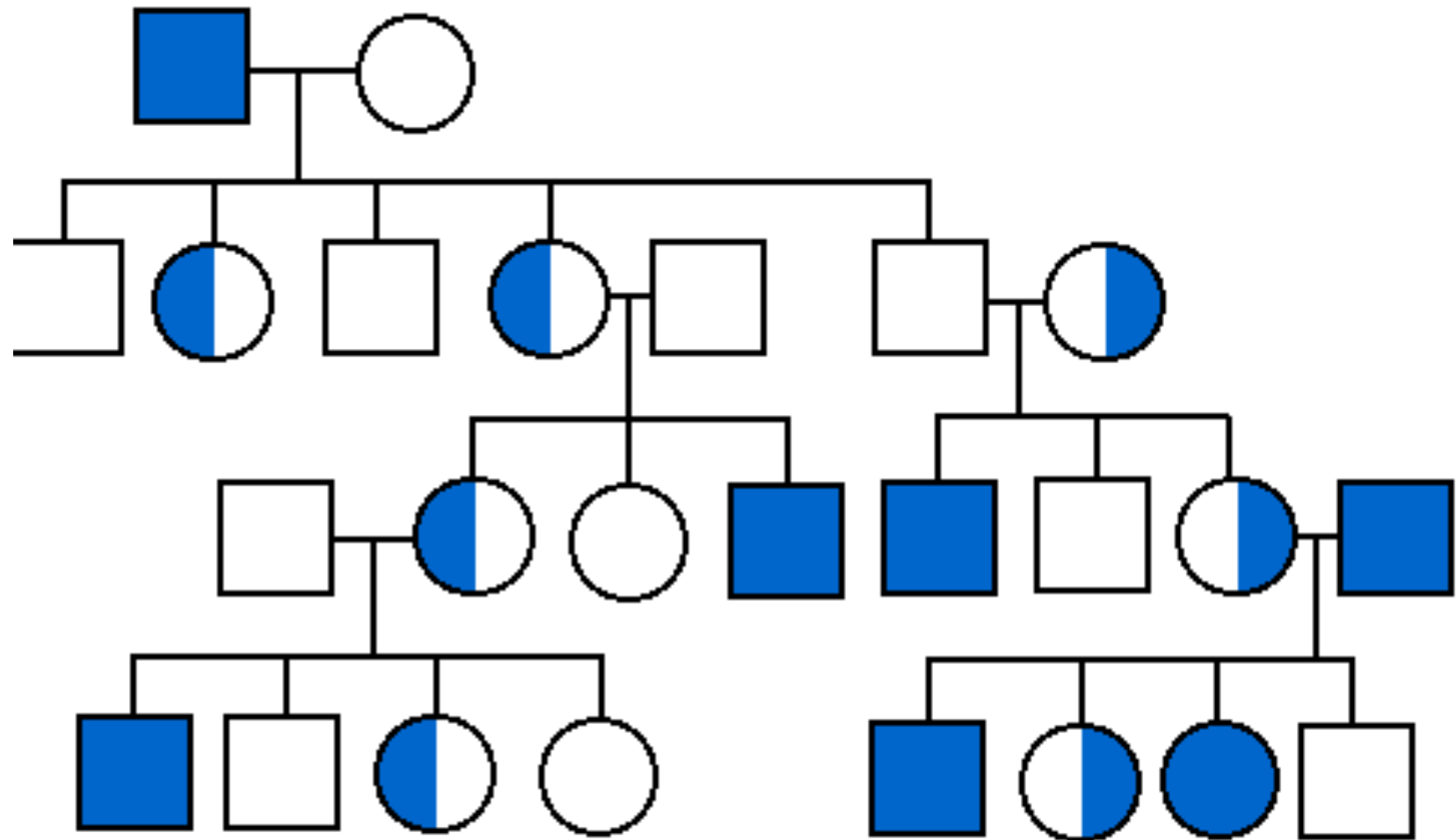


2
colorblind

5
normal vision

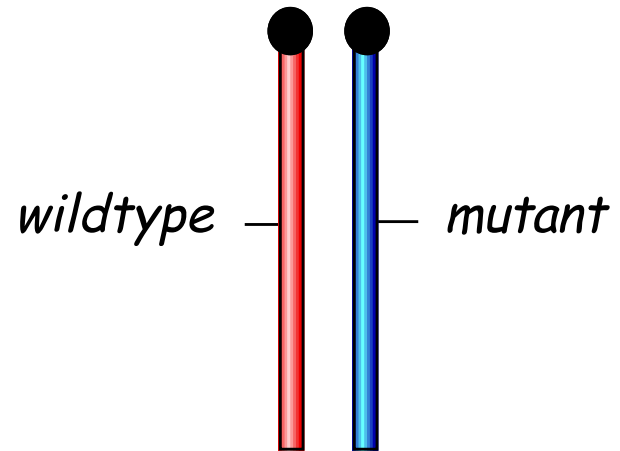
9
negative ctrl

Inheritance of Red-Green Color Blindness: an X-linked Recessive Trait

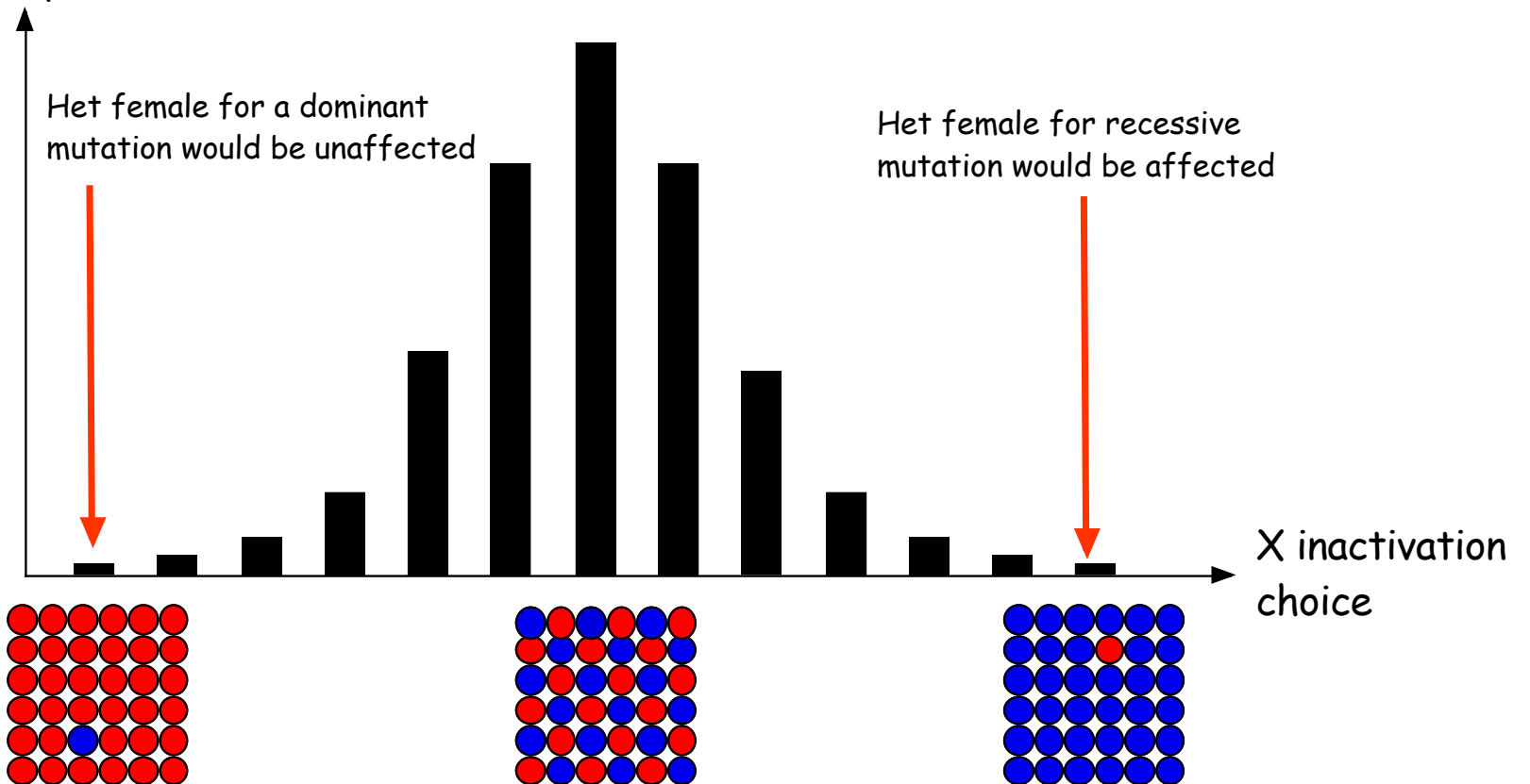


 = Carrier of Trait

↑
colorblind female
How??



% of females with X
inactivation pattern



Doctors Said These Women's Mutated Genes Wouldn't Harm Them

But they were always at risk of developing diseases with potentially severe effects.

By Roxanne Khamsi

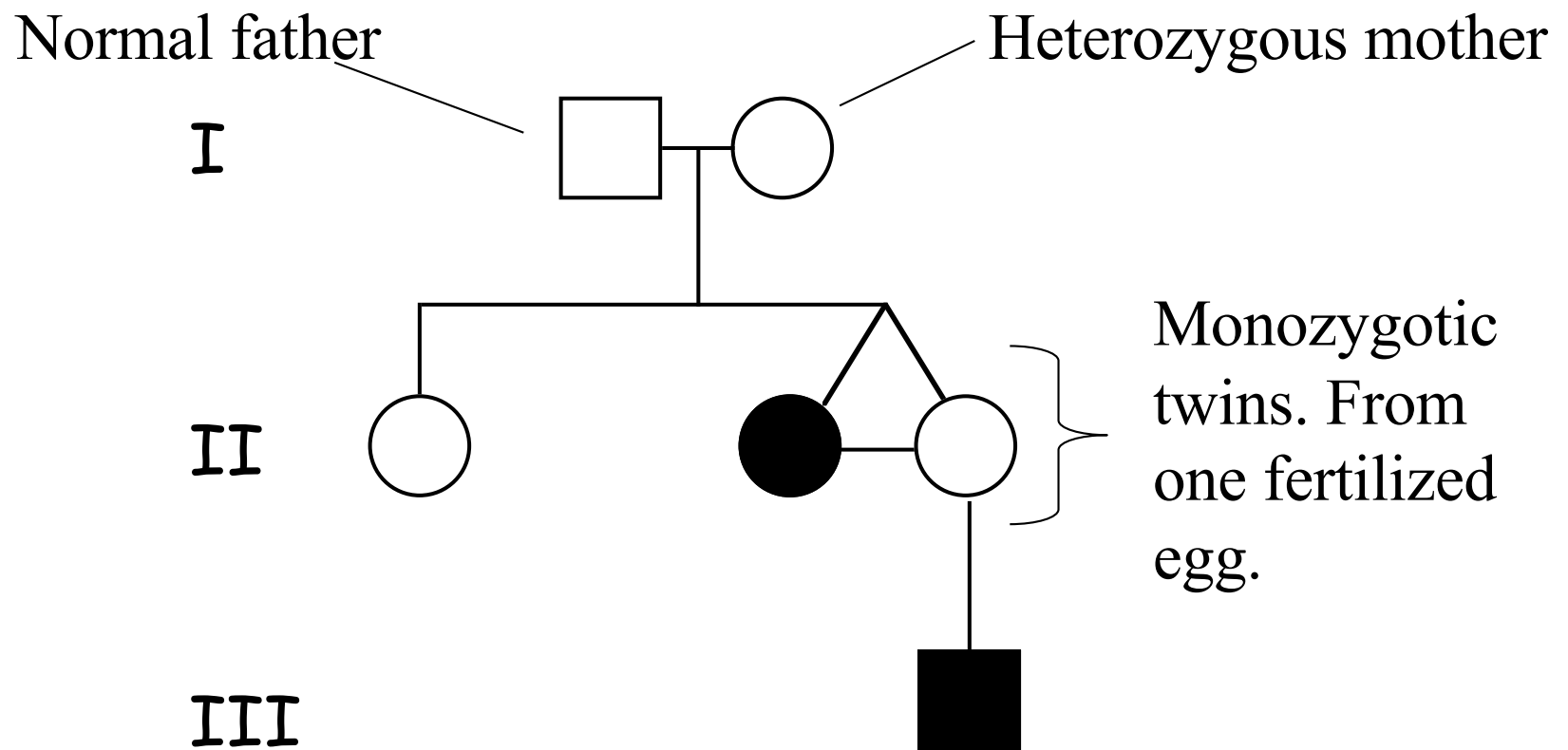


Skewed X inactivation and discordant monozygotic twins

X-linked recessive disease

Fabry disease (α -galactosidase A)

OMIM 301500



Hypophosphatemia Is an X-linked Dominant Trait

