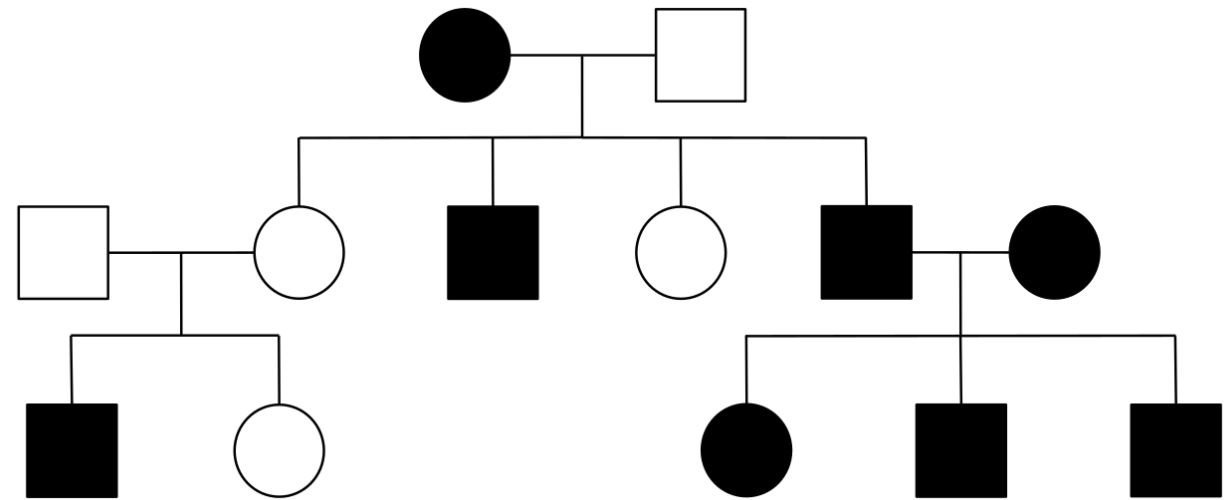


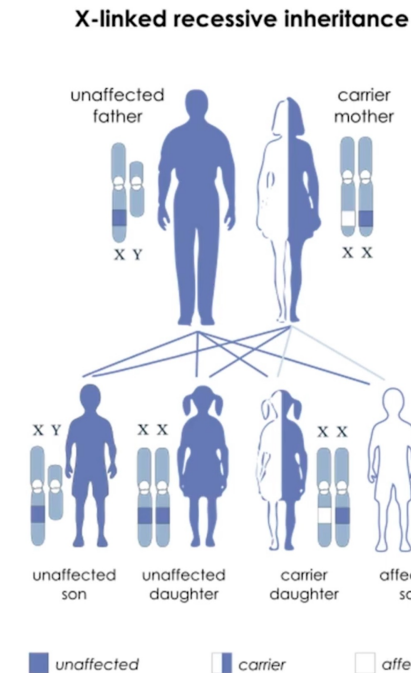
Lesson 16

Pedigrees

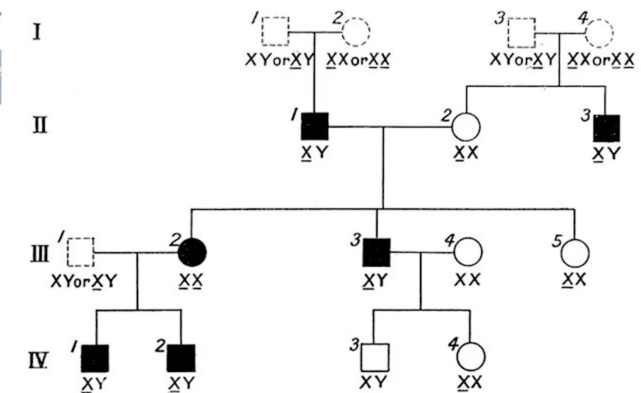


Pedigrees

- Sometimes, the genetics of a particular trait are very complicated
 - There might be many genes involved
 - There might be traits that are associated with sex
 - There might be traits that are influenced in ways that we just don't understand
- **Pedigrees** = another set of tools to understand complex traits



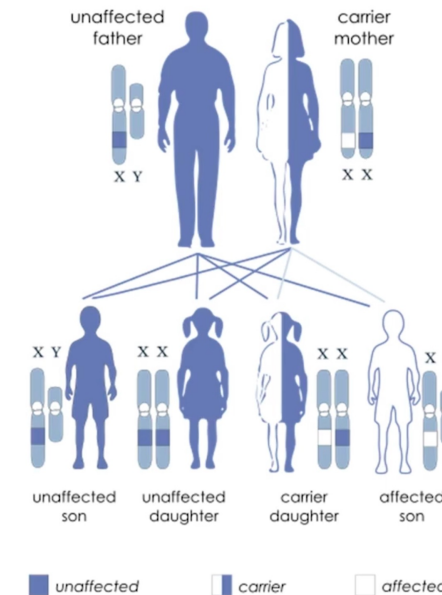
Pedigree analysis
uncovers type of inheritance
suggests gene(s) characteristics



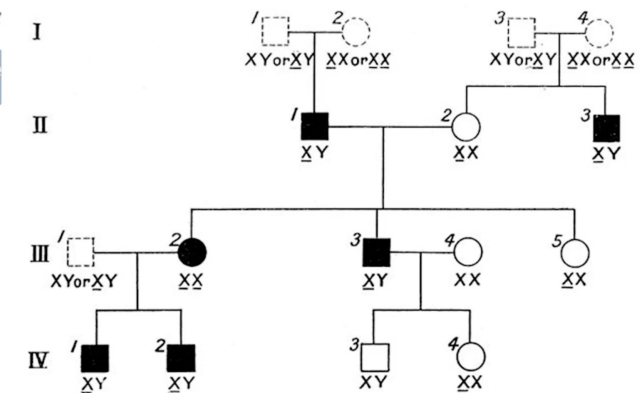
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 - There might be traits that are associated with sex
 - There might be traits that are influenced in ways that we just don't understand
- Pedigrees = another set of tools to understand complex traits
- Pedigrees **track traits through generations**, helping us understand inheritance patterns
- They are used in genetic counseling, medicine, breeding programs, and forensic science
- Pedigree analysis **helps diagnose genetic disorders** and **predict inheritance risks**

X-linked recessive inheritance



Pedigree analysis
uncovers type of inheritance
suggests gene(s) characteristics

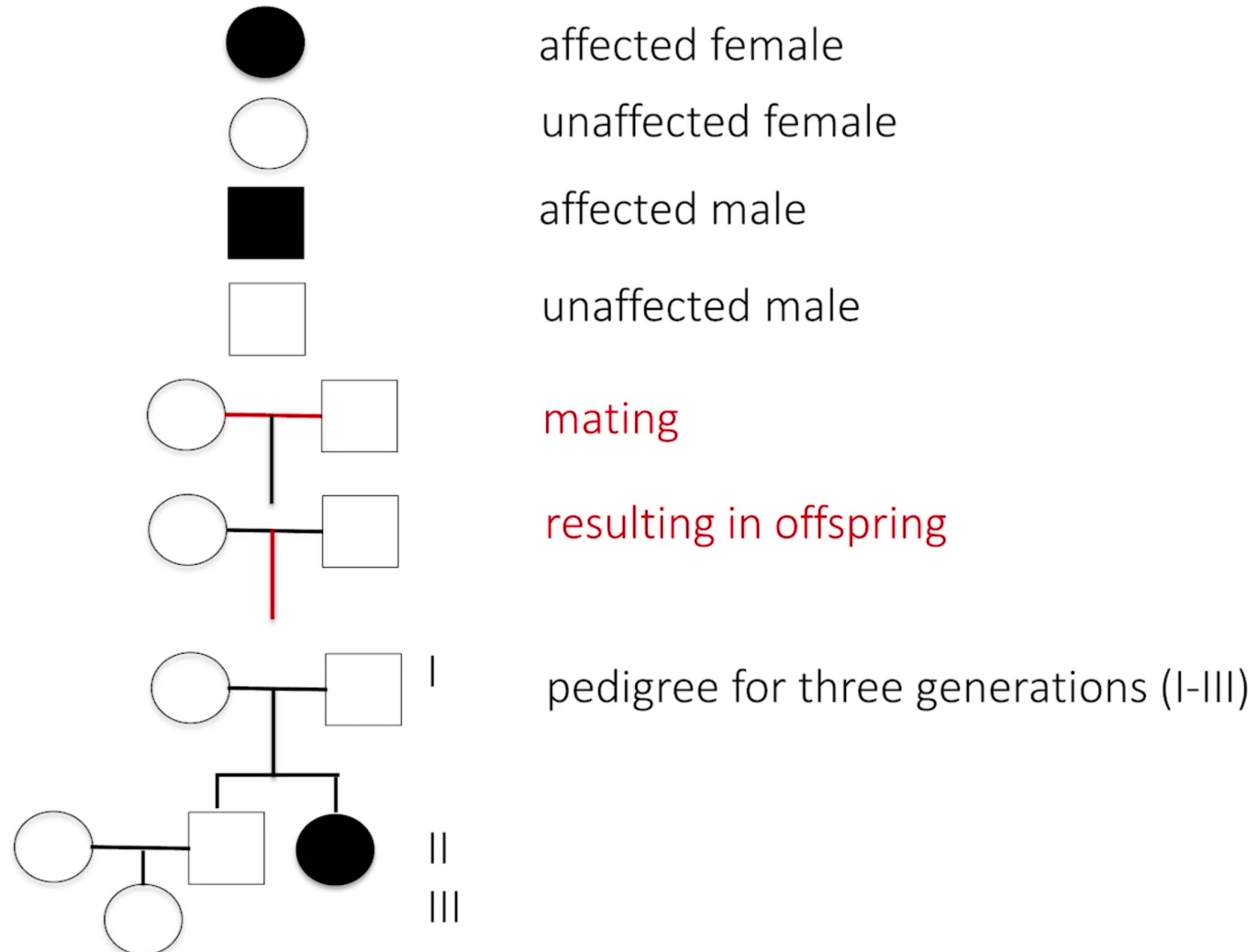


Pedigrees

- **Pedigrees = genetics from family history**
- Conventions in pedigree writing:
 - ○ female
 - □ male
 - ● ■ the trait you are looking at is present (*e.g.*, a disease, brown eyes, etc.)

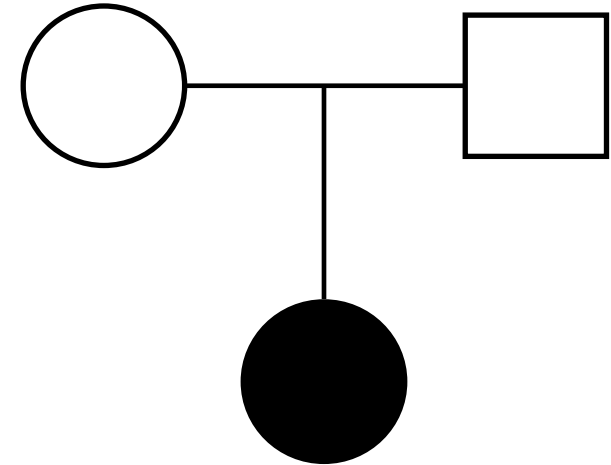
More pedigree nomenclature

Pedigree nomenclature



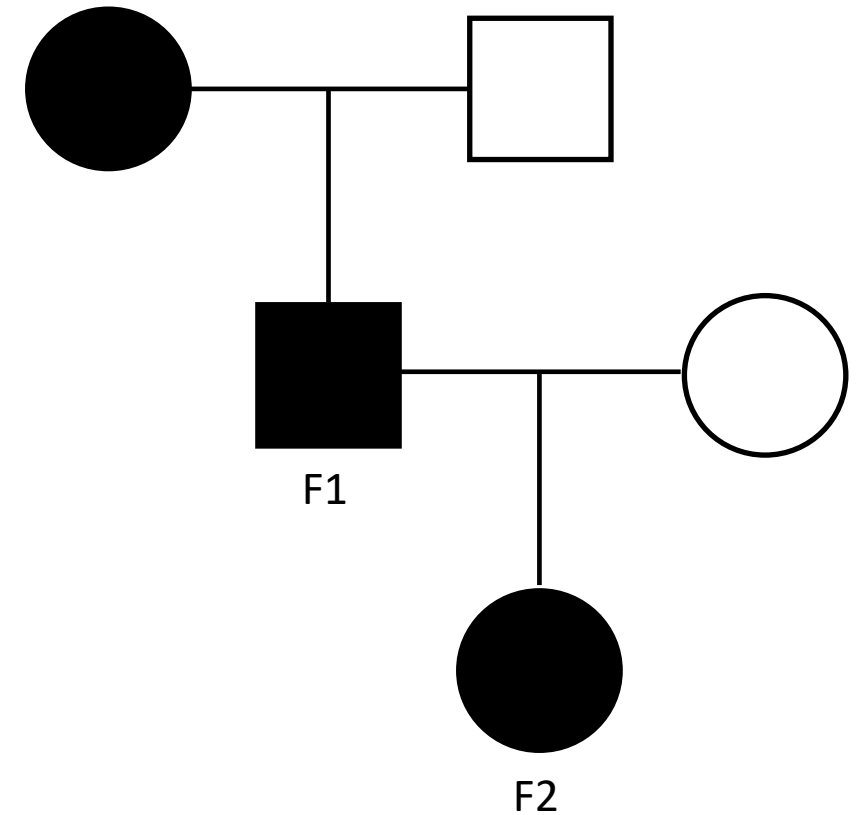
Pedigrees - 1

- This pedigree (*e.g.*, disease):
 - Two healthy parents
 - One affected daughter
- **This is a characteristic of a recessive trait**



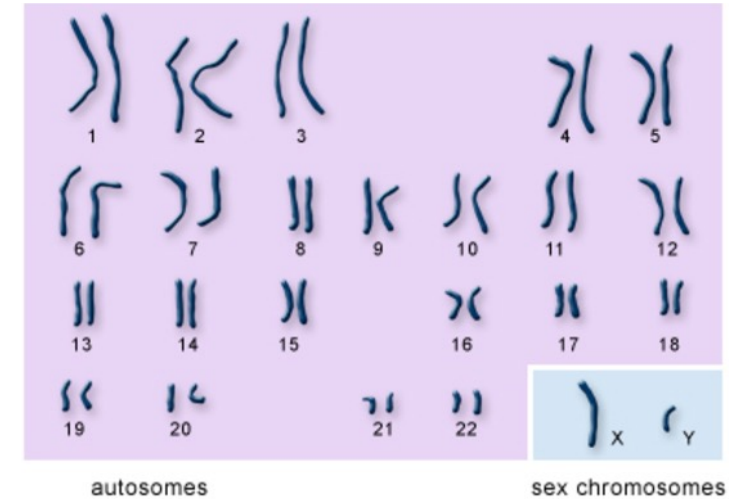
Pedigrees - 2

- This pedigree (*e.g.*, disease):
 - One affected parent and one unaffected parent (*e.g.*, father)
 - One affected son (F1)
 - The affected son mates an unaffected female
 - One affected daughter (F2)
- **The pedigree pattern where every affected offspring has an affected parent is characteristic of a dominant trait**



Chromosomes and pedigrees

- Eukaryote cells have two types of chrs:
 - Autosomes = 22 paired chrs (2 chrs 1, 2 chrs2, etc....)
 - Sex chrs*:
 - XX (paired) → female * simplified
 - XY (unpaired) → male



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 - Autosomal dominant
- } Males and females affected equally

Chromosomes and pedigrees

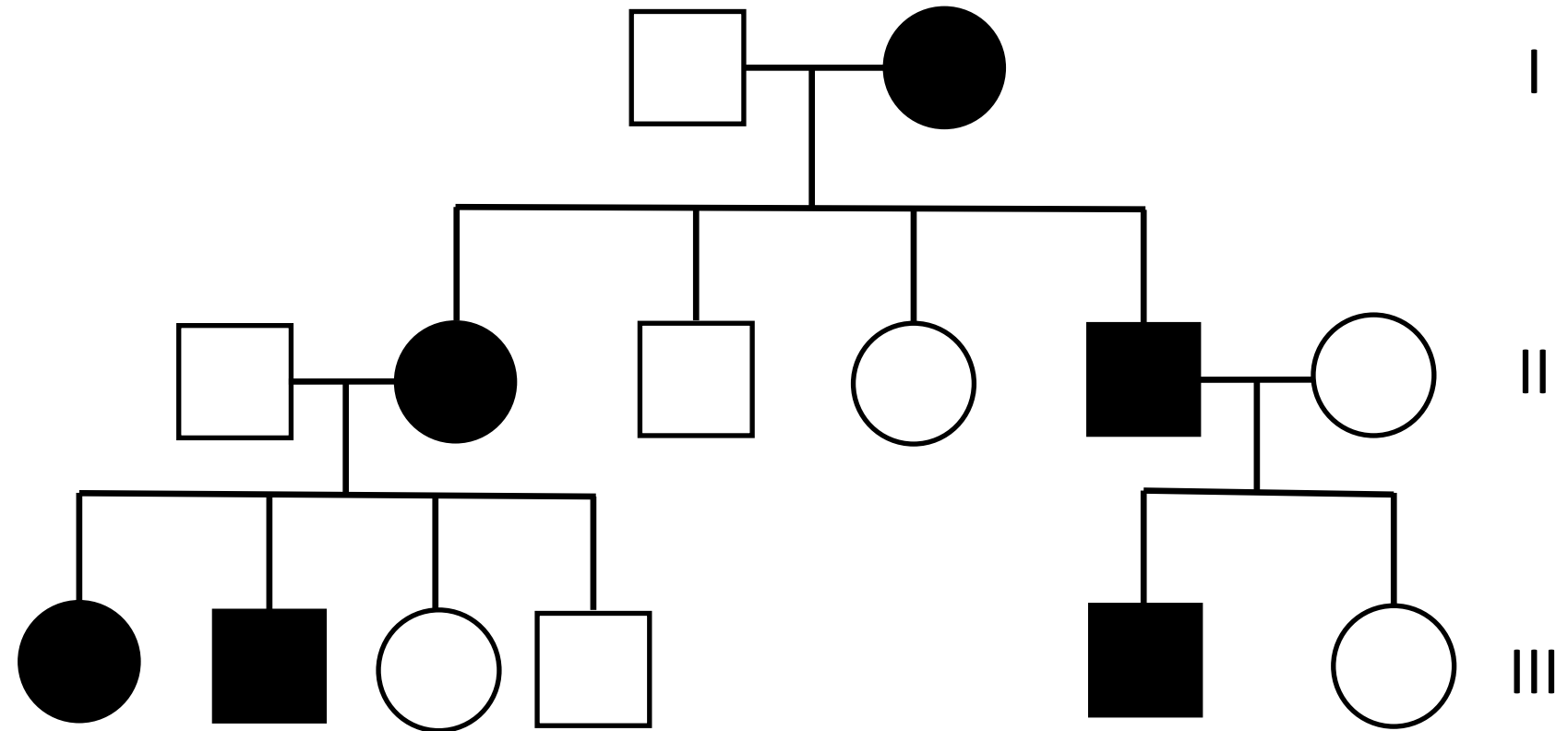
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- **X-linked recessive → Males affected more than females**
 - there is only one X (the other allele is Y) and is affected → the trait will show itself)
 - It is X-linked recessive because this trait on the X allele is always transmitted to the SONS (XY) from the mother

Pedigree for autosomal dominant trait

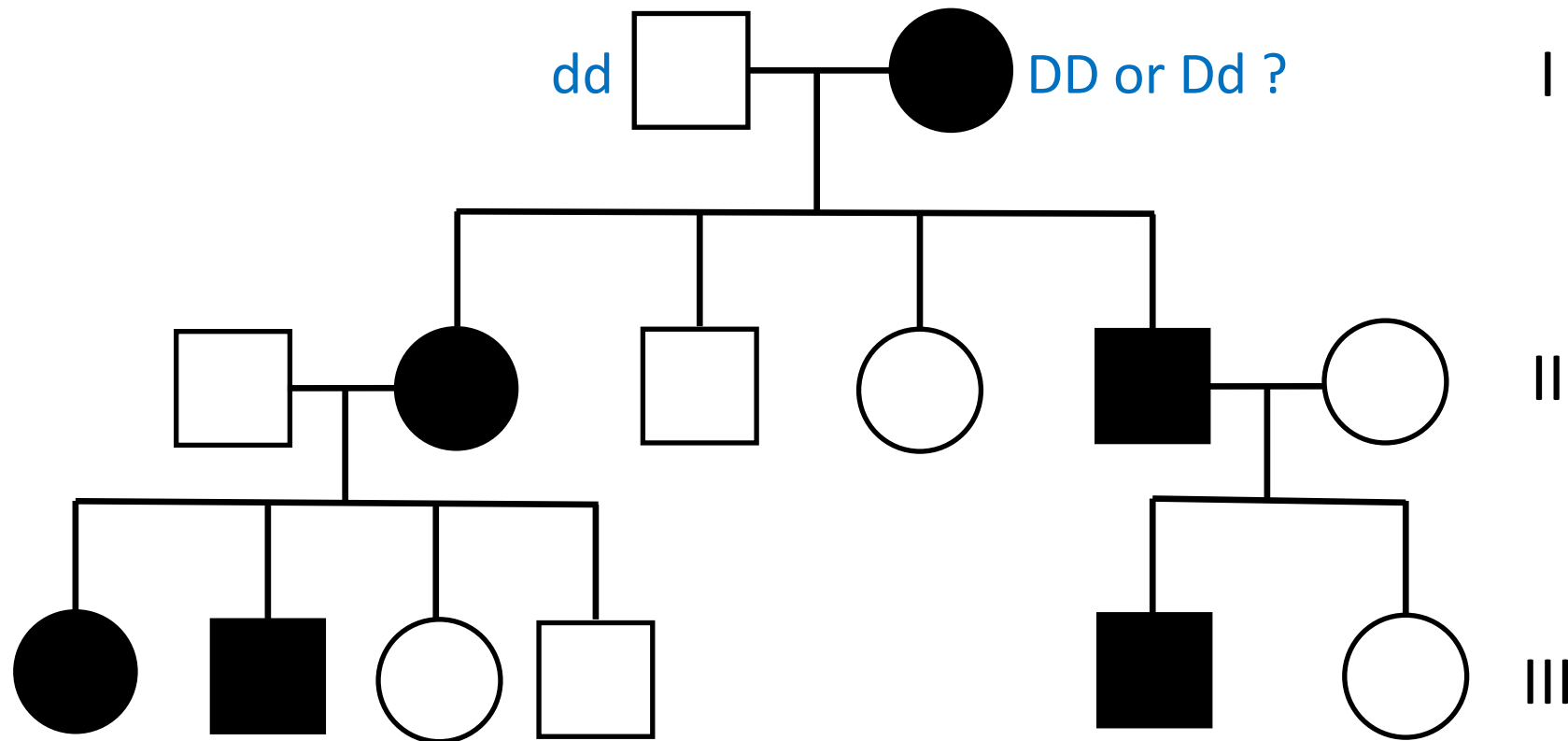
Clues

Autosomal: both males and females are affected



Dominant: every affected child has one affected parent



Pedigree for autosomal dominant trait (with genotypes)

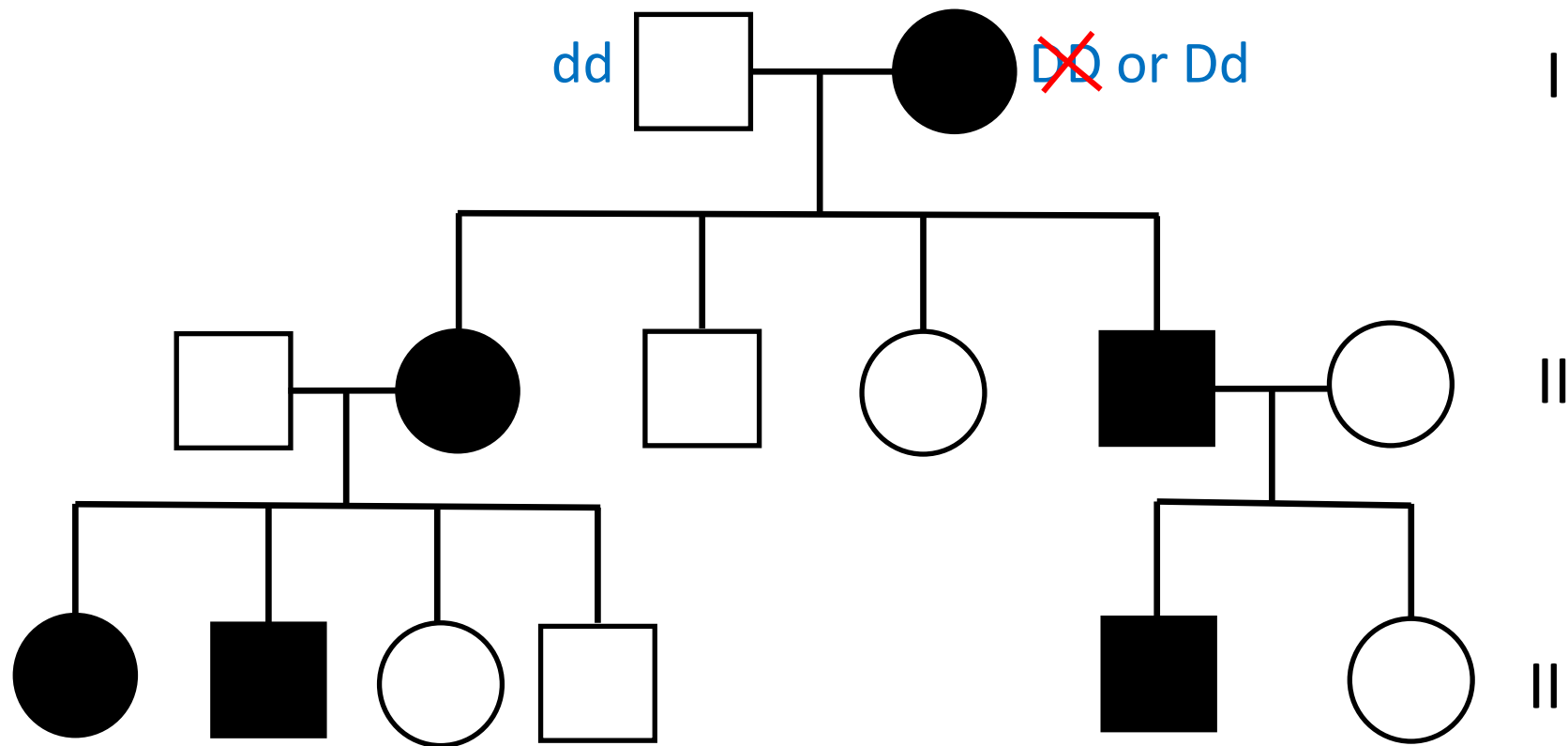


Punnett squares for gen II (gen 1 = parents)

| | | | |
|---|--|---------|----|
| | | Gametes | |
|  G a m e t e s |  d | D | D |
| | d | Dd | Dd |
| | d | Dd | Dd |

100% of II would be affected (all have the dominant allele D)

Pedigree for autosomal dominant trait (with genotypes)



Punnett squares for gen II (gen I = parents)

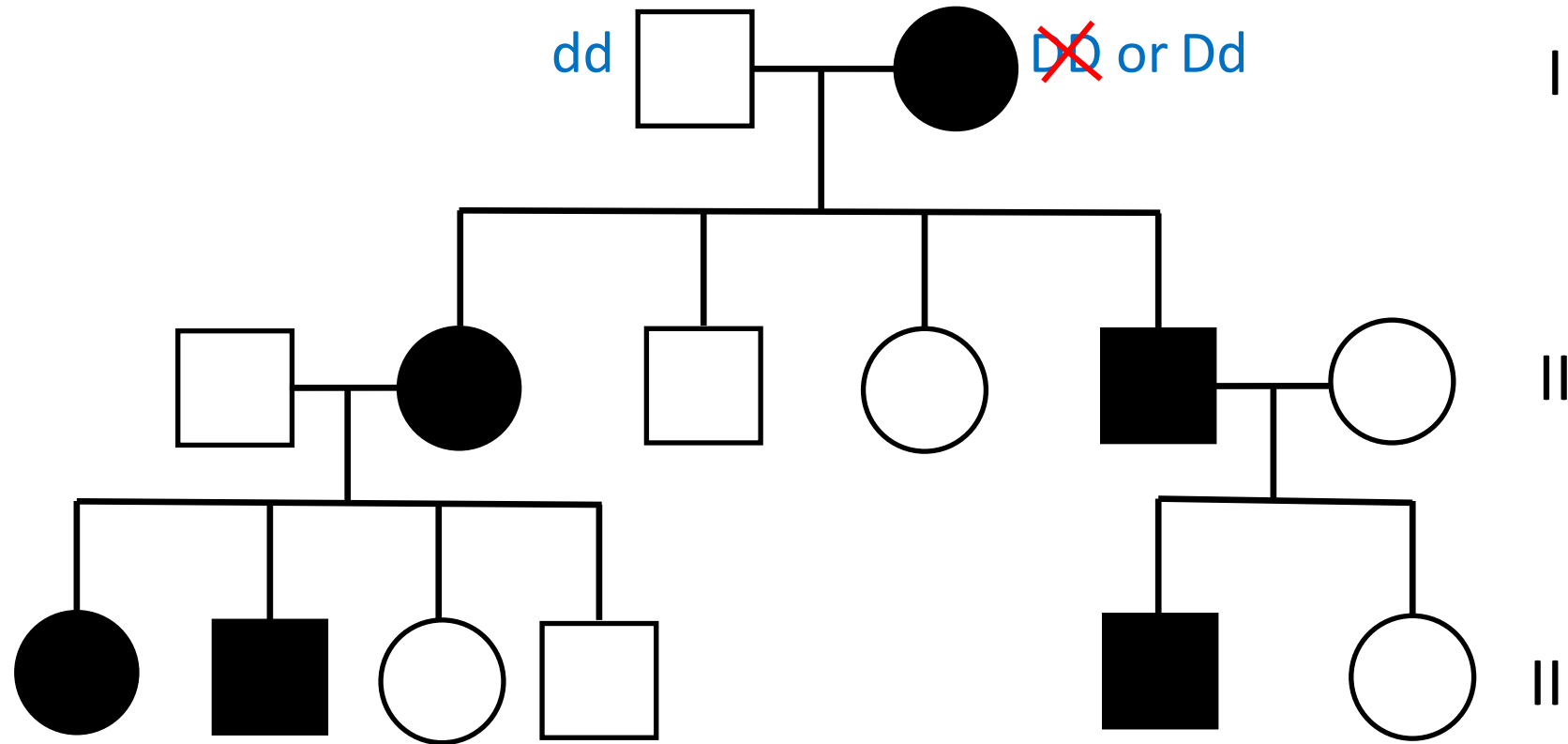
| | | | |
|---------------------------------|---|---------|----|
| | | Gametes | |
| | | D | D |
| G a m e t e s | d | Dd | Dd |
| | d | Dd | Dd |

100% of F1 would be affected (all have the dominant allele D)

| | | | |
|---------------------------------|---|---------|----|
| | | Gametes | |
| | | D | d |
| G a m e t e s | d | Dd | dd |
| | d | Dd | dd |

50% of II would be affected (half have the dominant allele D)

Pedigree for autosomal dominant trait (with genotypes)



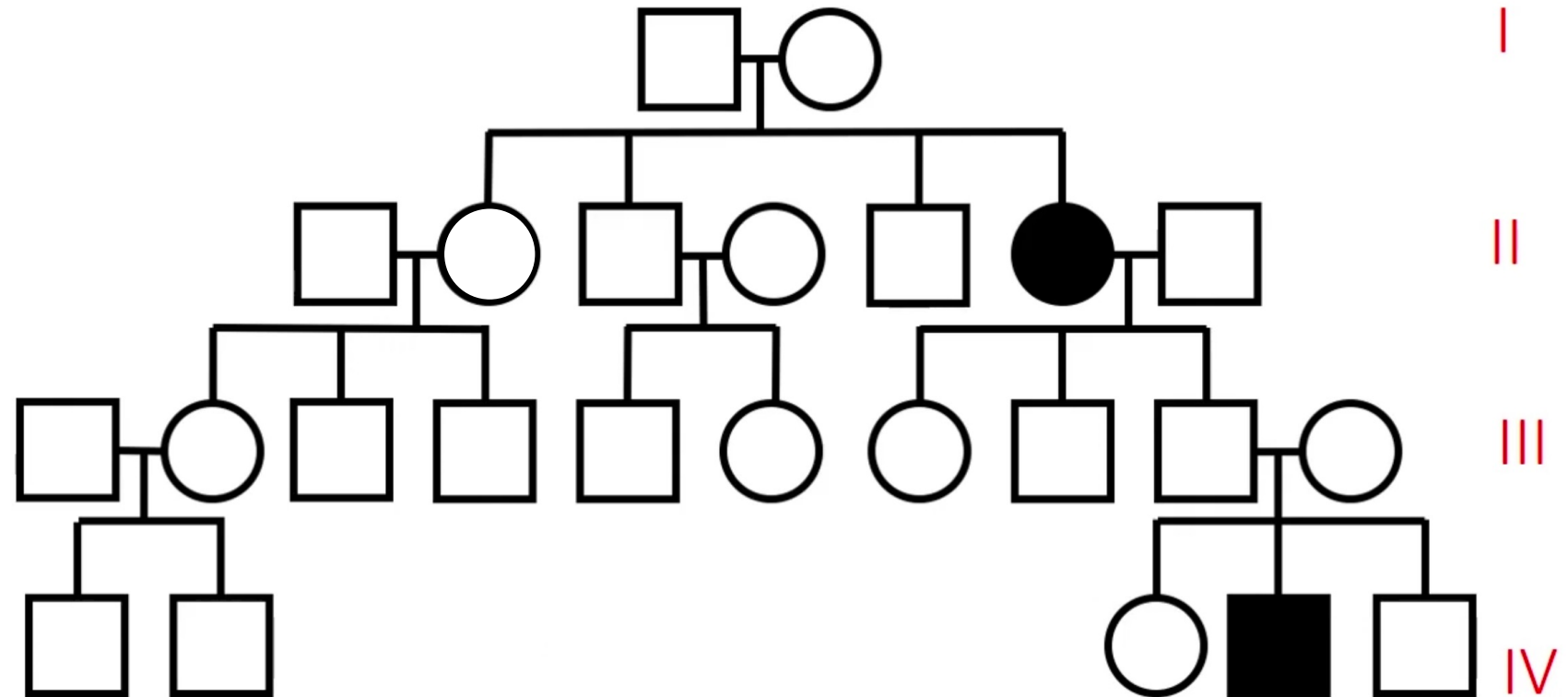
Note: you may not always get ratios identical to predicted*

Pedigree for autosomal recessive trait

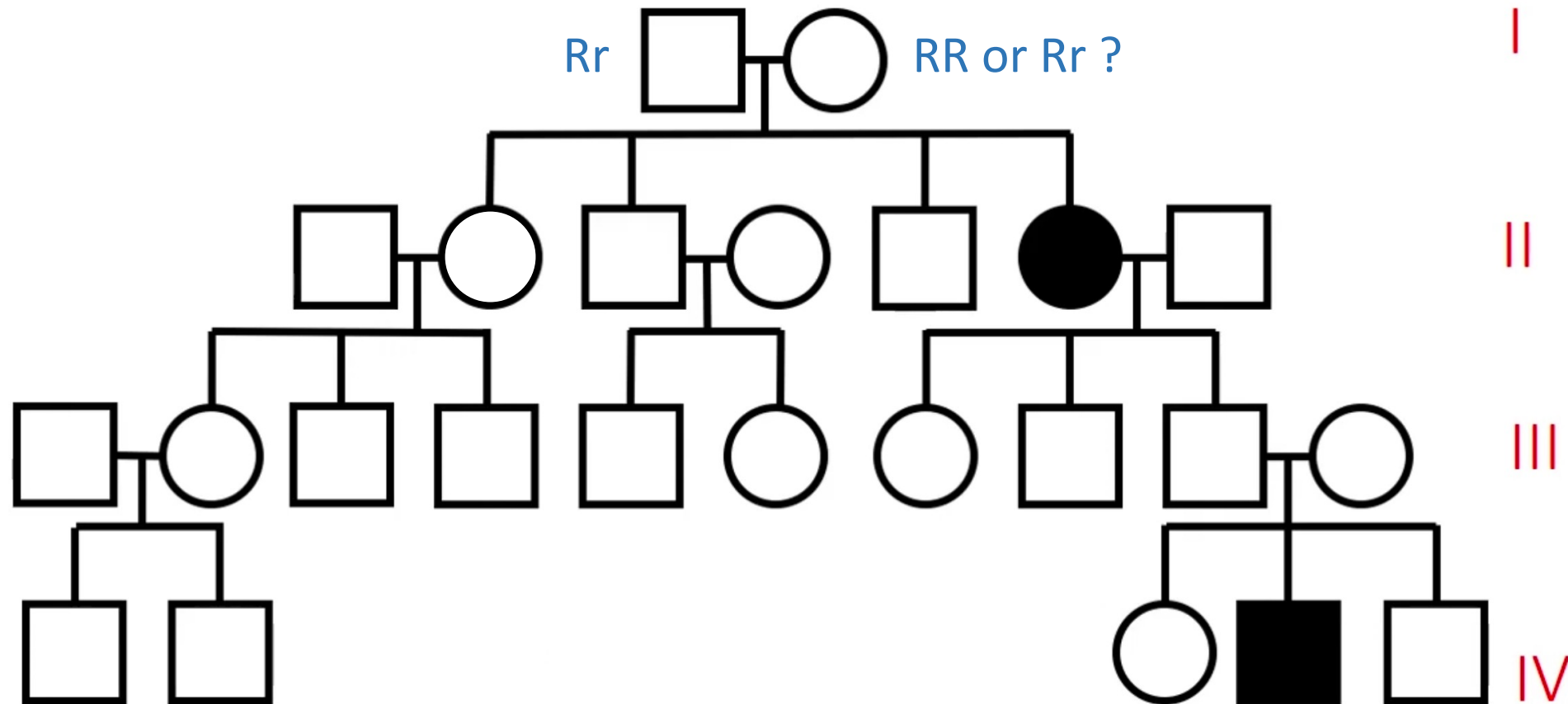
Clues

Autosomal: both males and females are affected

Recessive: affected
child(ren) from
unaffected parents
not many affected
offspring





Pedigree for autosomal recessive trait (with genotypes)



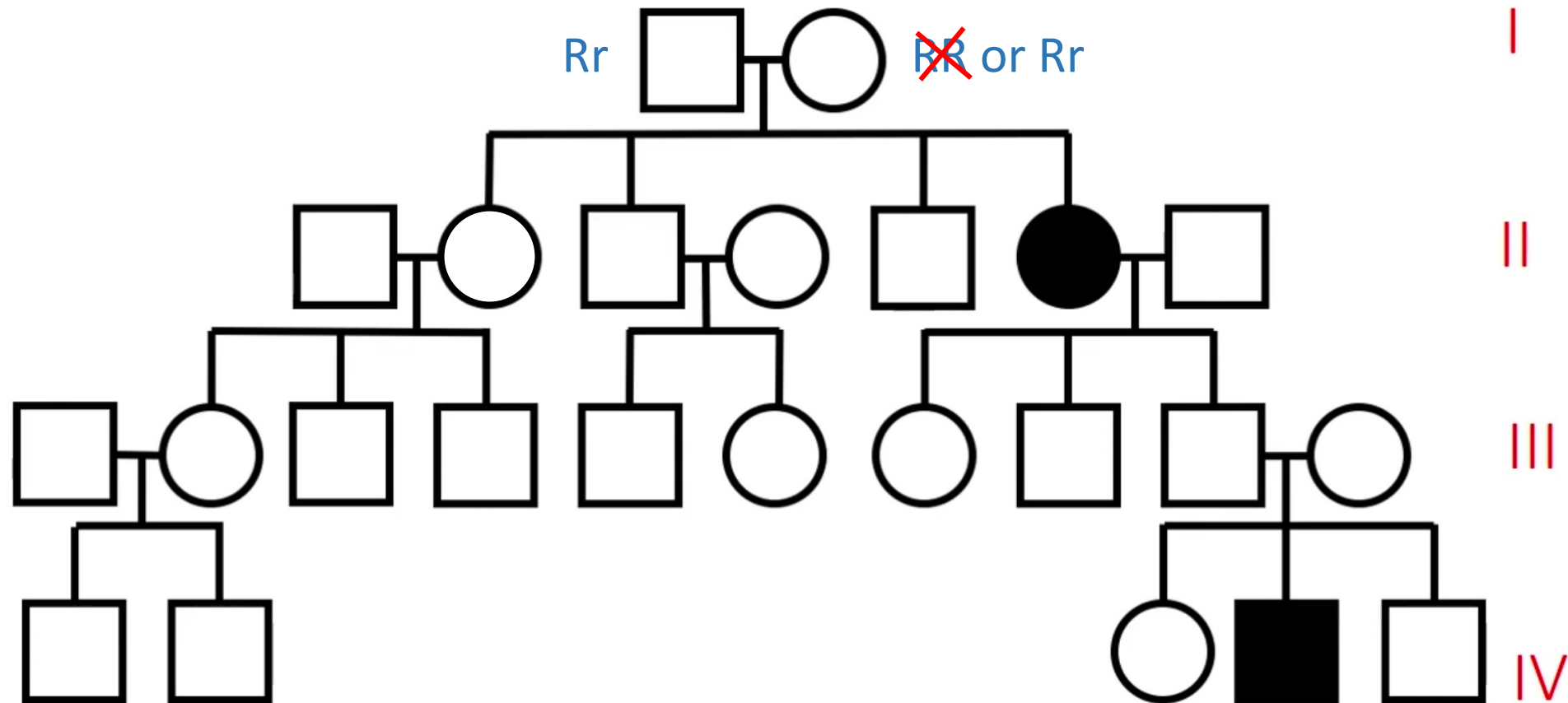
Remember: it is the recessive allele r that expresses the disease

Punnett squares for gen II (gen 1 = parents)



| | | | |
|---|--|---------|----|
| | | Gametes | |
|  G a m e t e s |  R | RR | RR |
| | r | Rr | Rr |

0% of II would be affected (none has genotype rr)



Pedigree for autosomal recessive trait (with genotypes)



Punnett squares for gen II (gen 1 = parents)

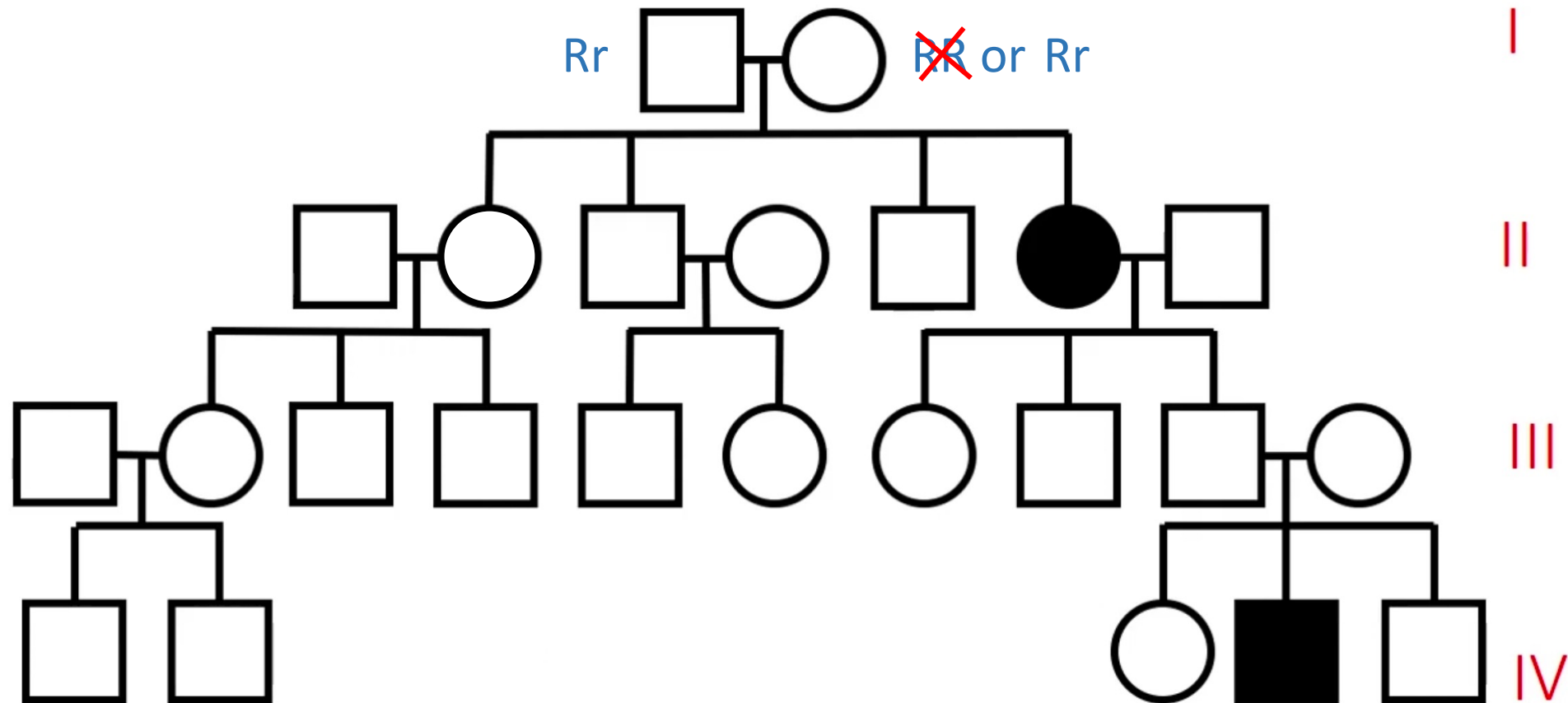
| | | | |
|---|---|---------|----|
| | | Gametes | |
|  G a m e t e s |  | R | R |
| | R | RR | RR |
| | r | Rr | Rr |

0% of II would be affected (none has genotype rr)

| | | | |
|--|---|---------|----|
| | | Gametes | |
|  G a m e t e s |  | R | r |
| | R | RR | Rr |
| | r | Rr | rr |

25% of II (1:3) would be affected (only 1 has rr)

Pedigree for autosomal recessive trait (with genotypes)



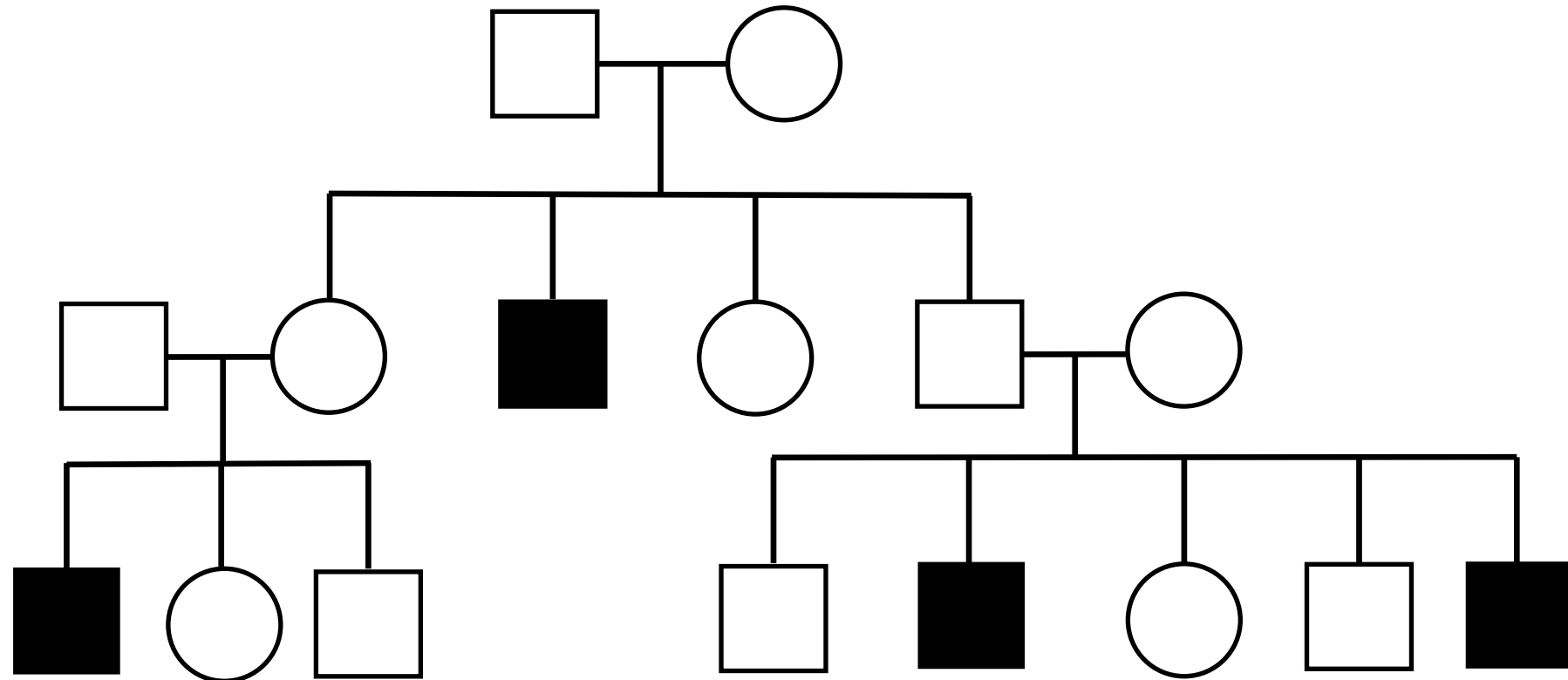
Note: you may not always get ratios identical to predicted*

Pedigree for X-linked recessive trait

Clues

X-linked: only males affected

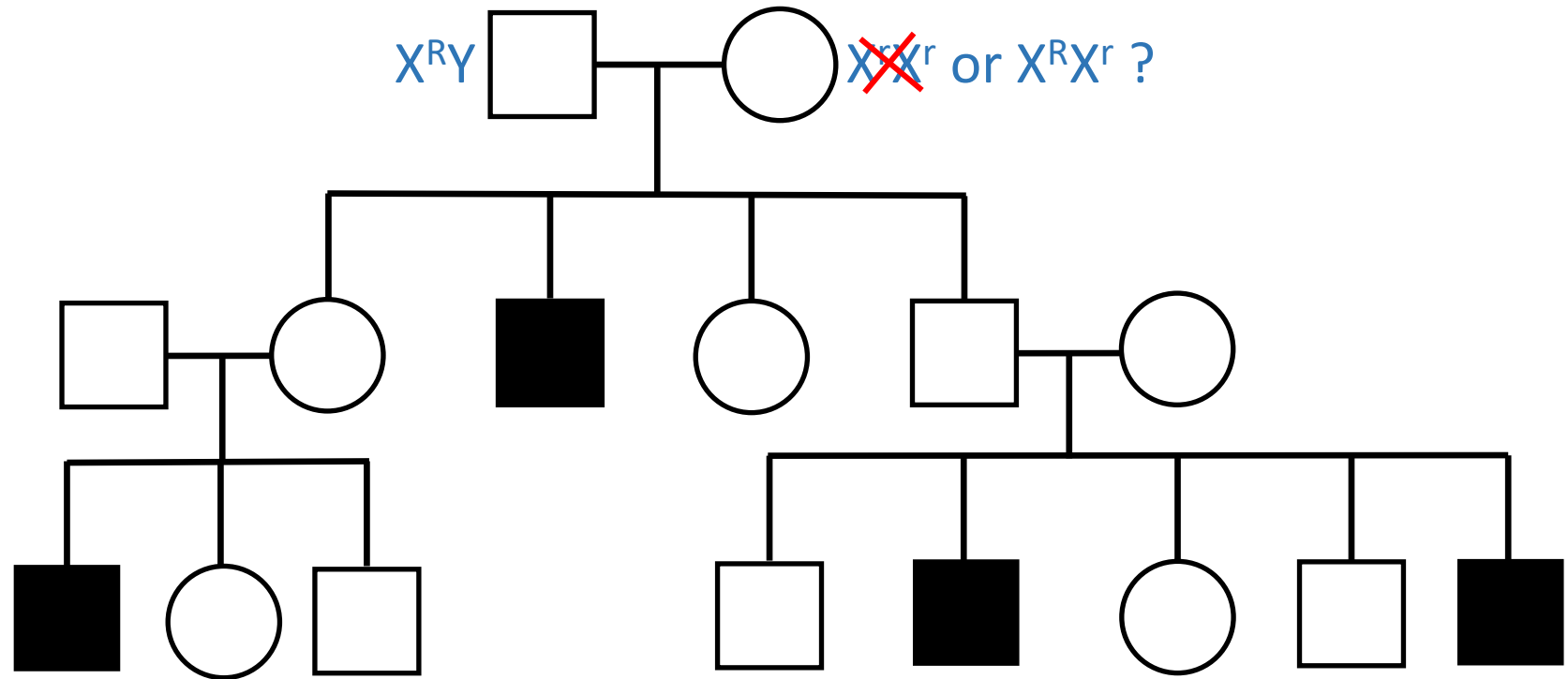
Recessive: affected child(ren) from unaffected parents
not many affected offspring





Pedigree for X-linked recessive trait (with genotypes)

Remember: it is the recessive allele r that expresses the disease

X^rX^r can be immediately excluded otherwise the mother would have the trait

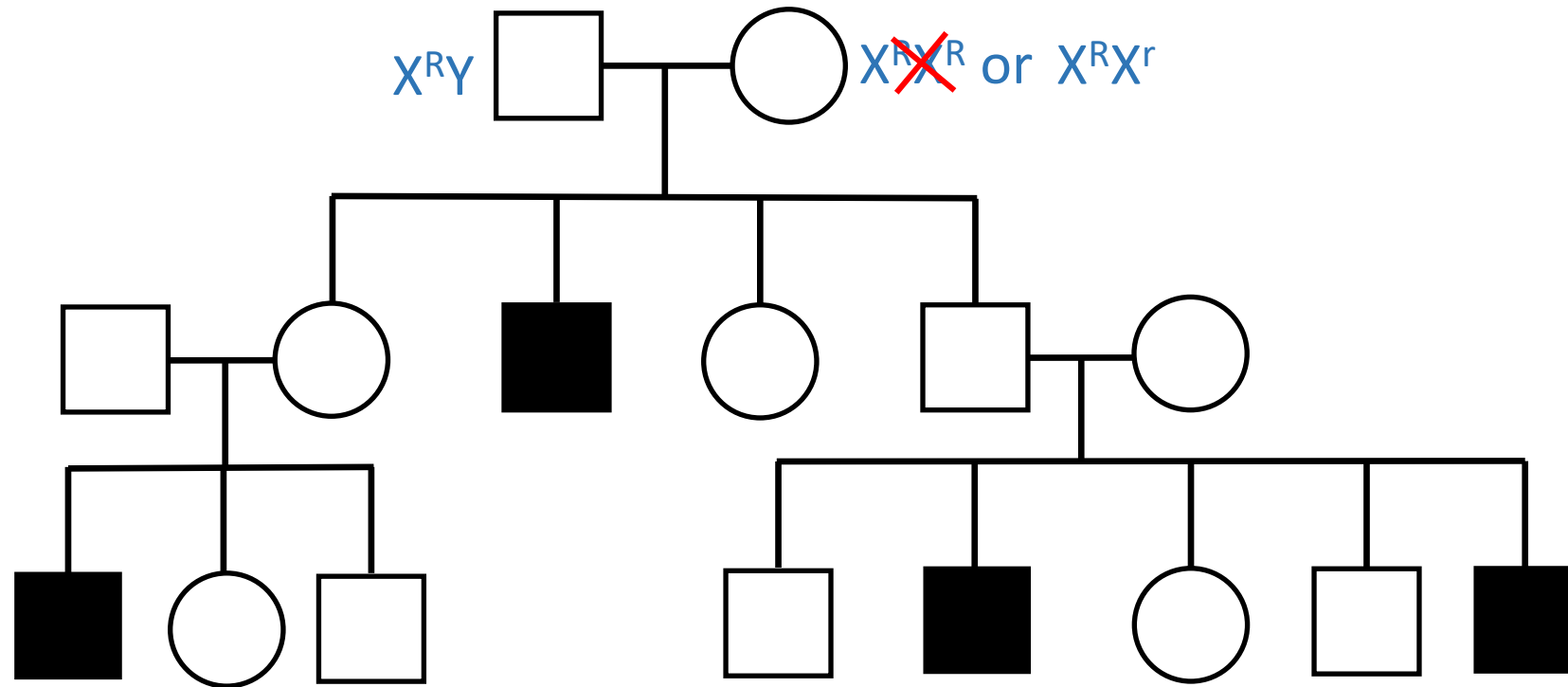


Punnett squares for gen II (gen 1 = parents)

| | | | |
|--|---|-----------|-----------|
| | | Gametes | |
|  G a m e t e s |  | X^R | X^r |
| | X^R | $X^R X^R$ | $X^R X^r$ |
| | Y | $X^R Y$ | $X^r Y$ |

50% of MALE II (1:1) would be affected (has genotype $X^r Y$)

Pedigree for X-linked recessive trait (with genotypes)



Note: you may not always get ratios identical to predicted*

Notes

- The discussed pedigrees are simple and ideal
- You may not always get ratios identical to predicted due to non-Mendelian inheritance patterns, which include (among others):
 - **Codominance and incomplete dominance**
 - **Polygenic traits**: some traits are determined by the combined effect of more than one pair of genes → polygenic (or continuous) traits (an example of this is human stature)
 - **Multiple-allele series**: the ABO blood type system is also an example of a trait that is controlled by more than just a single pair of alleles
 - **Modifying and regulator genes**: modifying genes alter how certain other genes are expressed in the phenotype (for instance, here is a dominant cataract gene which will produce varying degrees of vision impairment depending on the presence of a specific allele for a companion modifying gene)

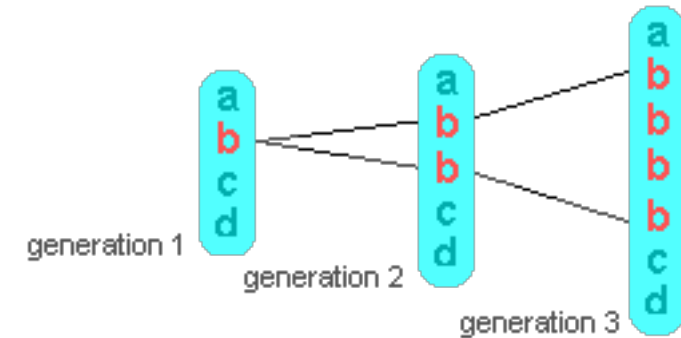
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 - you may inherit the genes that are responsible for type 2 diabetes but never get the disease unless you become greatly overweight, persistently stressed psychologically, or do not get enough sleep on a regular basis;
 - the genes that cause the chronic autoimmune disease, multiple sclerosis may be triggered by the Epstein-Barr virus and possibly other specific environmental stresses

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- **Pleiotropy**: a single gene may be responsible for a variety of traits. The complex of symptoms that are collectively referred to as sickle-cell trait or sickle-cell anemia, is an example.
 - A single gene results in irregularly shaped red blood cells that painfully block blood vessels, cause poor overall physical development, as well as related heart, lung, kidney, and eye problems

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- **Stuttering alleles:** some genetically inherited diseases have more severe symptoms each succeeding generation due to segments of the defective genes being doubled in their transmission to children (typically fatal degenerative nerve disorders)
- You may not have enough offspring at F1 to interpret pedigree
 - You need to analyze F2, F3, ...