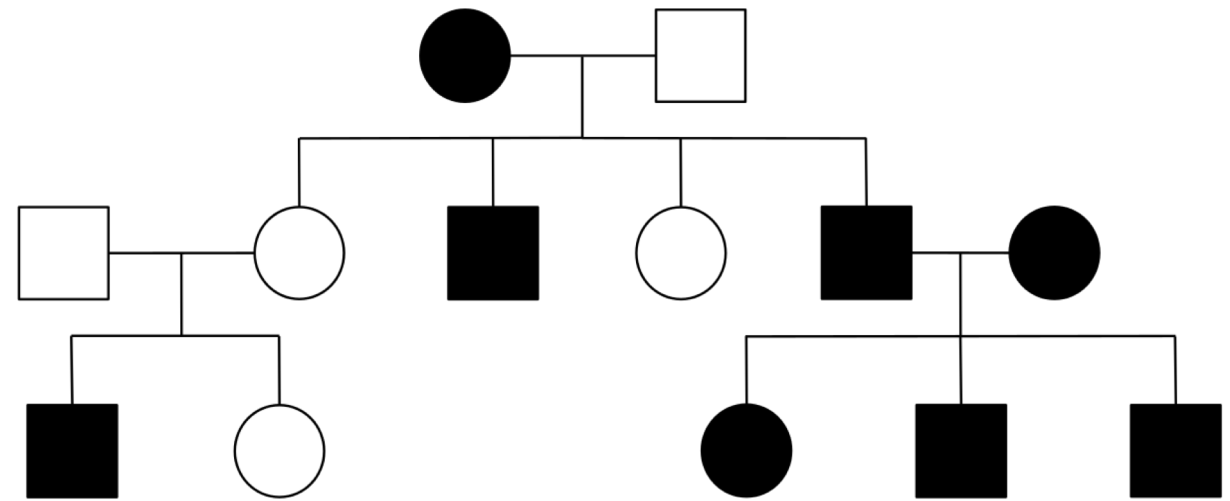


# Lesson 16

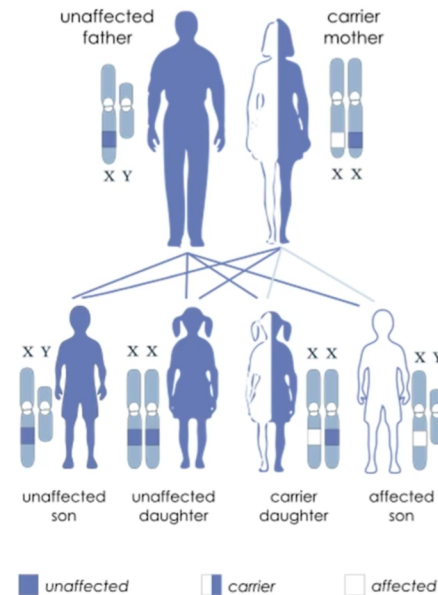
## Pedigrees



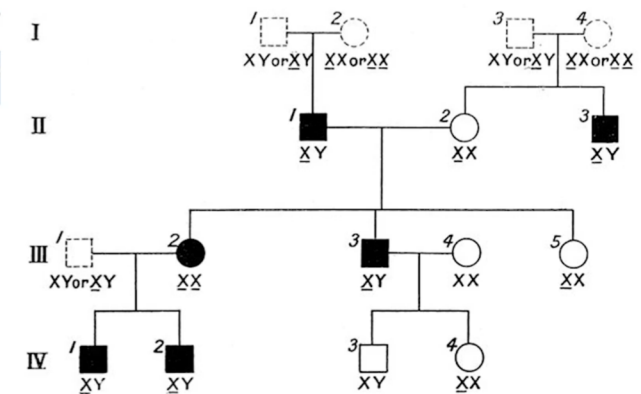
# Pedigrees

- 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
  - Especially important to understand diseases

## X-linked recessive inheritance



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

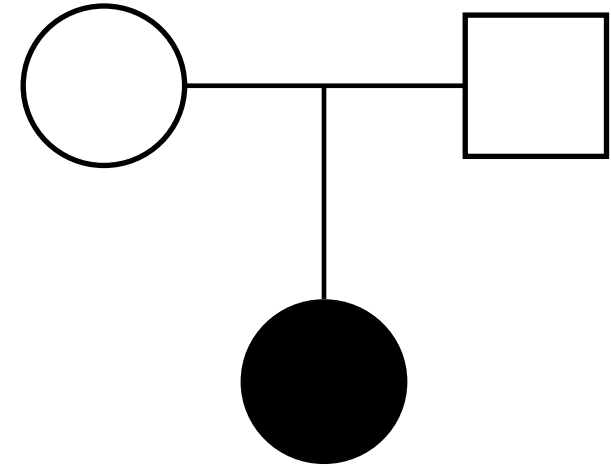


# Pedigrees

- **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.)

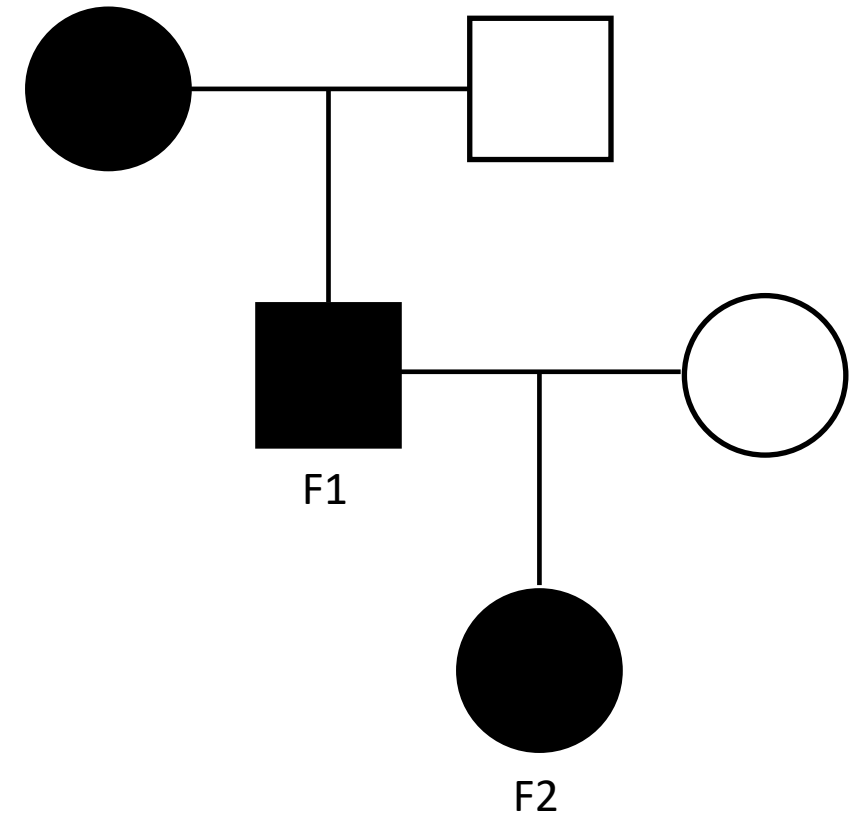
# Pedigrees - 1

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



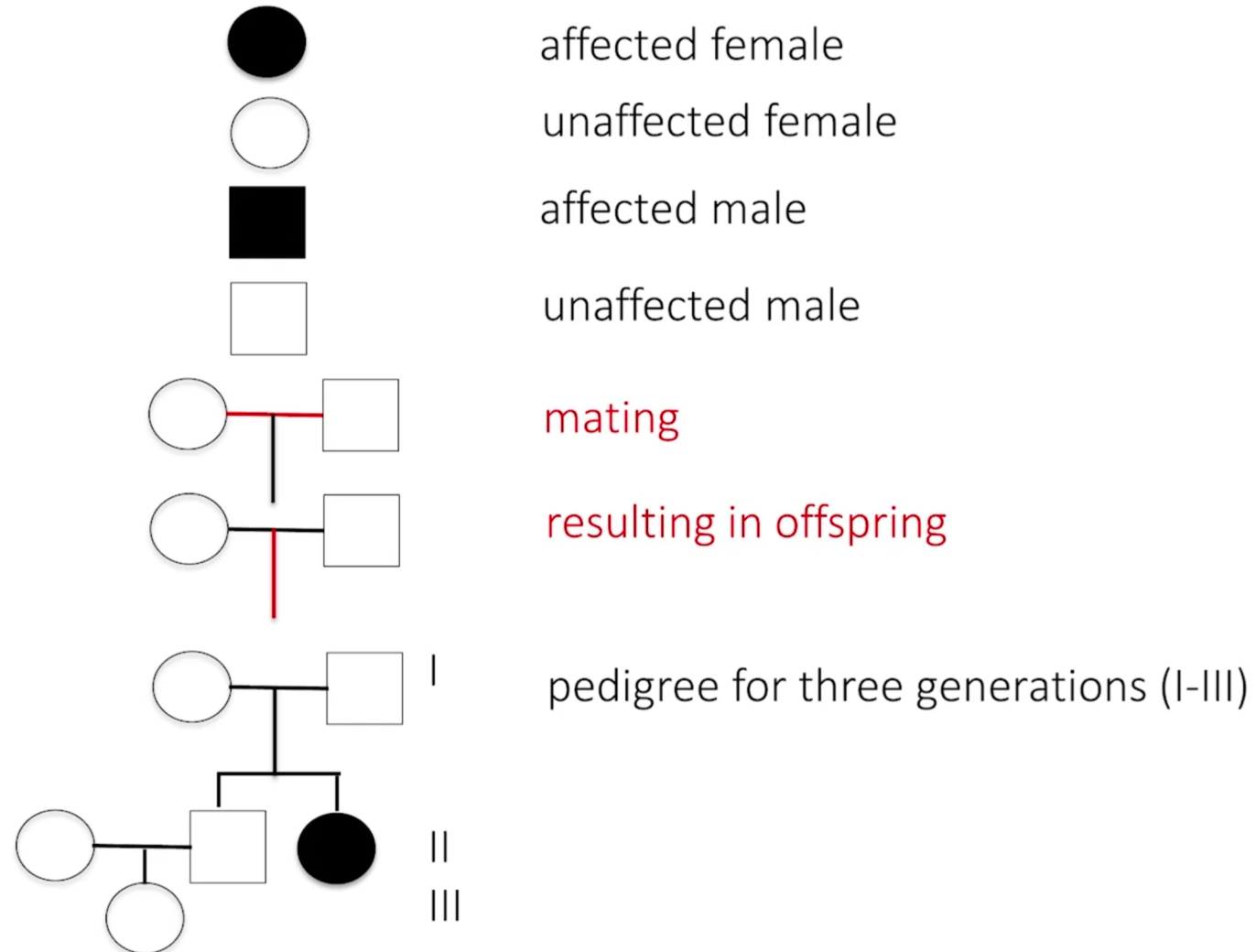
# Pedigrees - 2

- 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**



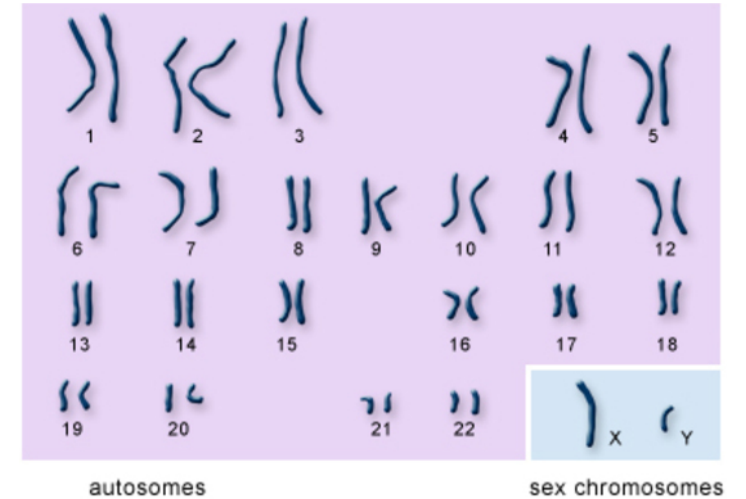
# More pedigree nomenclature

Pedigree nomenclature



# 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



# Chromosomes and pedigrees

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    - XX (paired) → female \* simplified
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- 3 types of pedigree based on autosomes or sex chromosomes:
- Autosomal recessive } Males and females affected equally
- Autosomal dominant }



# Chromosomes and pedigrees

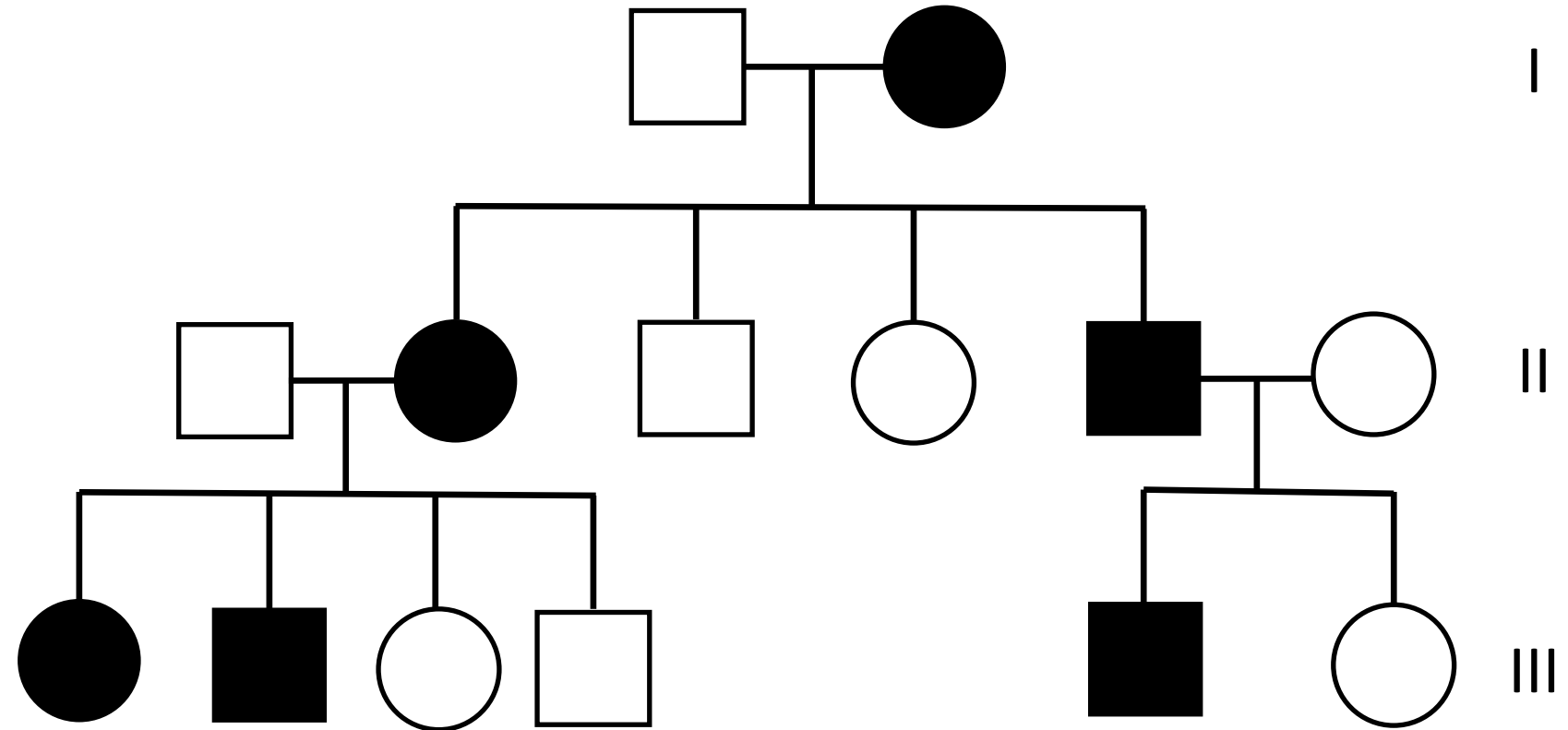
- Eukaryote cells have two types of chrs:
  - Autosomes = 22 paired chrs (2 chrs 1, 2 chrs2, etc....)
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    - XX (paired) → female
    - XY (unpaired) → male \* simplified
- 3 types of of pedigree based on autosomes or sex chromosomes:
- Autosomal recessive
- Autosomal dominant } Males and females affected equally
- **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

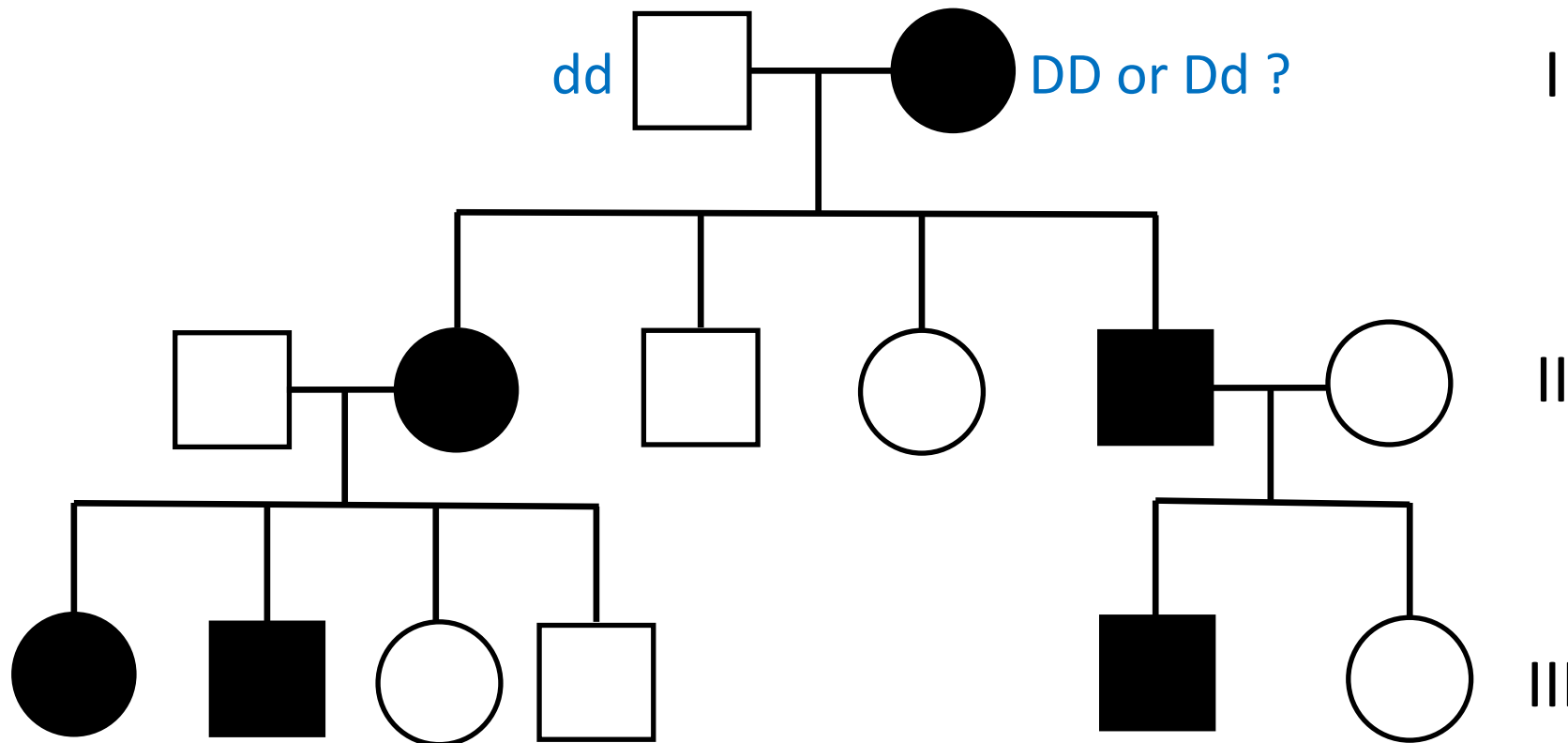
Clues

**Autosomal:** both males and females are affected



**Dominant:** every affected child has one affected parent



# Pedigree for autosomal dominant trait (with genotypes)

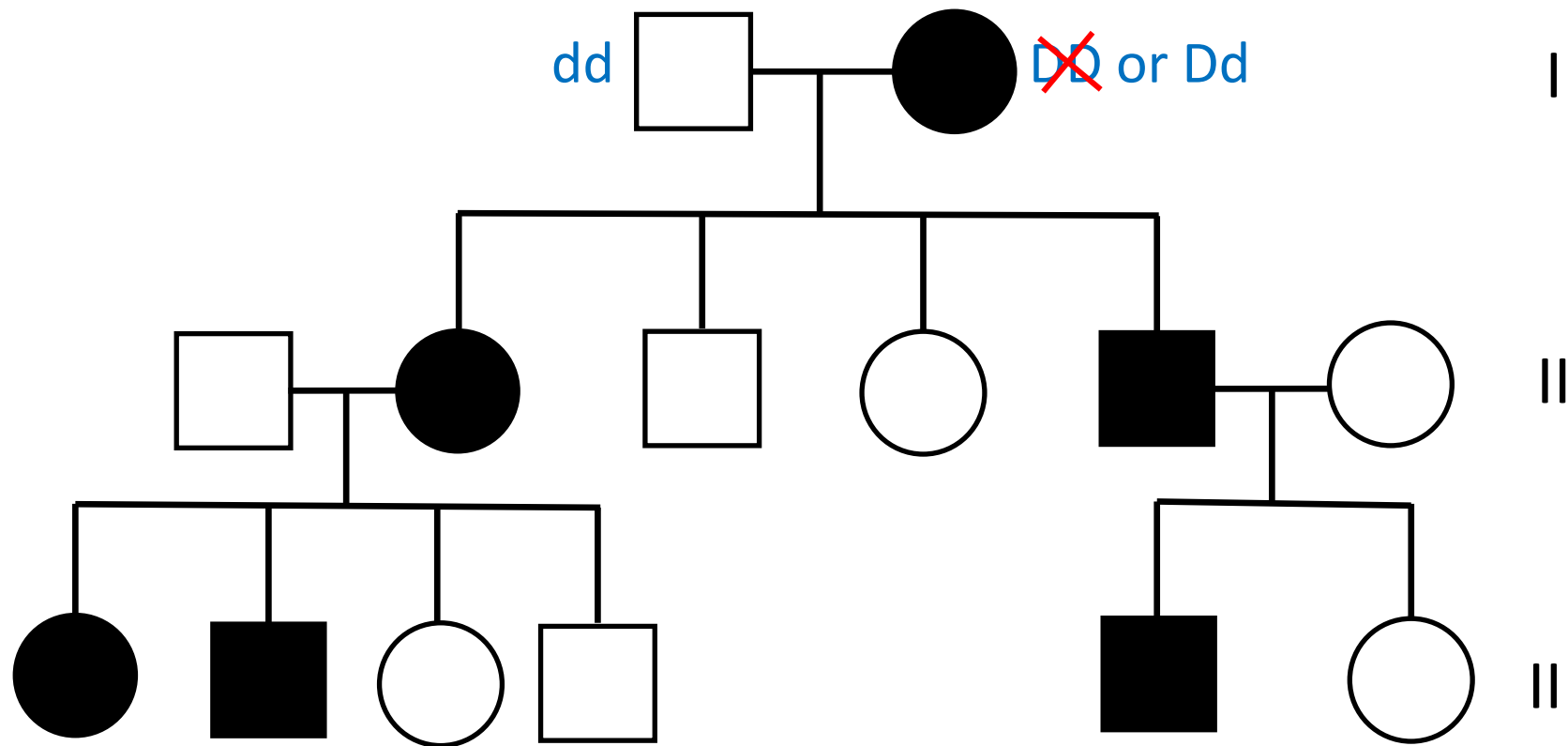


# Punnett squares for gen II (gen 1 = parents)



		Gametes	
		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)



# Pedigree for autosomal dominant trait (with genotypes)



# Punnett squares for gen II (gen I = parents)

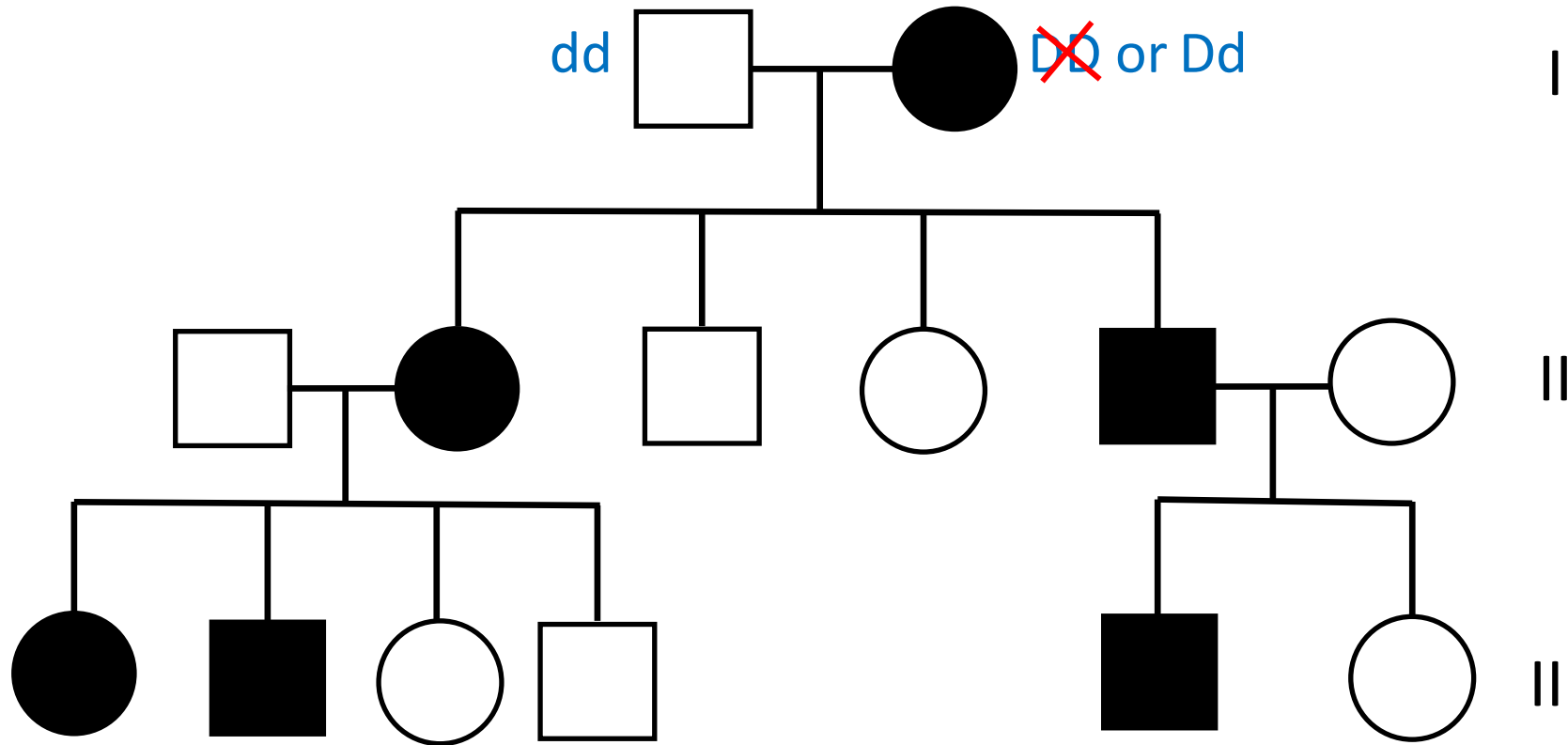
		Gametes	
		D	D
G a m e t e s	d	Dd	Dd
	d	Dd	Dd

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 F1 would be affected (half have the dominant allele D)

# Pedigree for autosomal dominant trait (with genotypes)



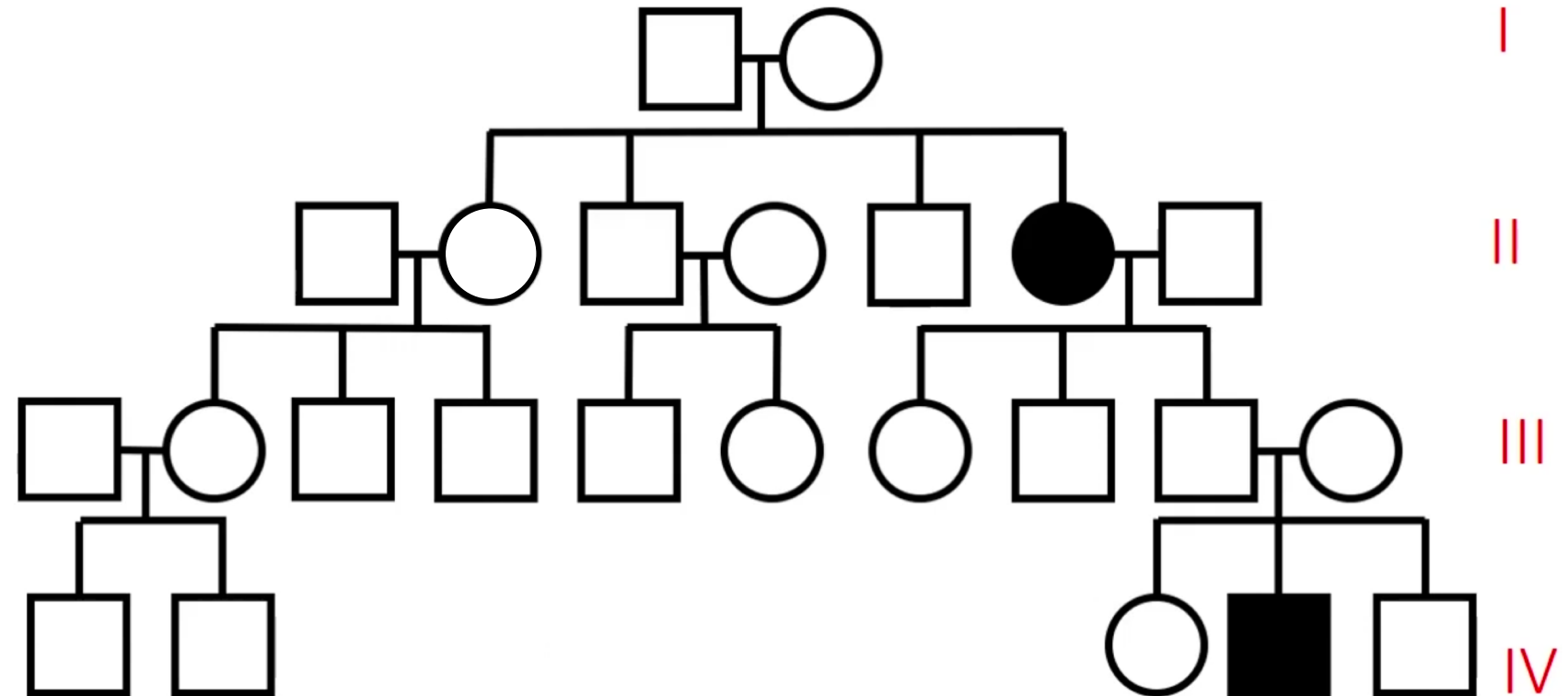
Note: you may not always get ratios identical to predicted due to probability of using specific gametes

# Pedigree for autosomal recessive trait

## Clues

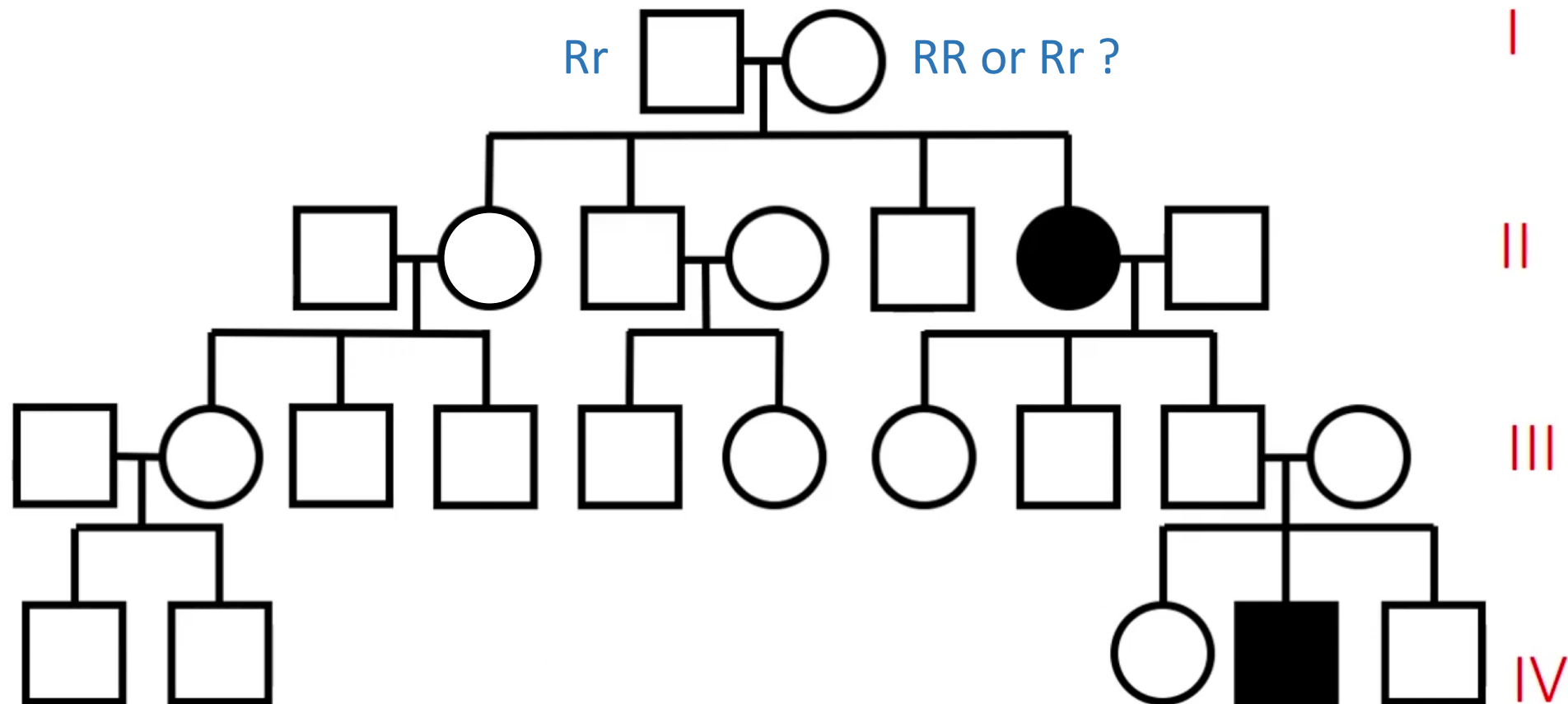
**Autosomal:** both males and females are affected

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







# Pedigree for autosomal recessive trait (with genotypes)



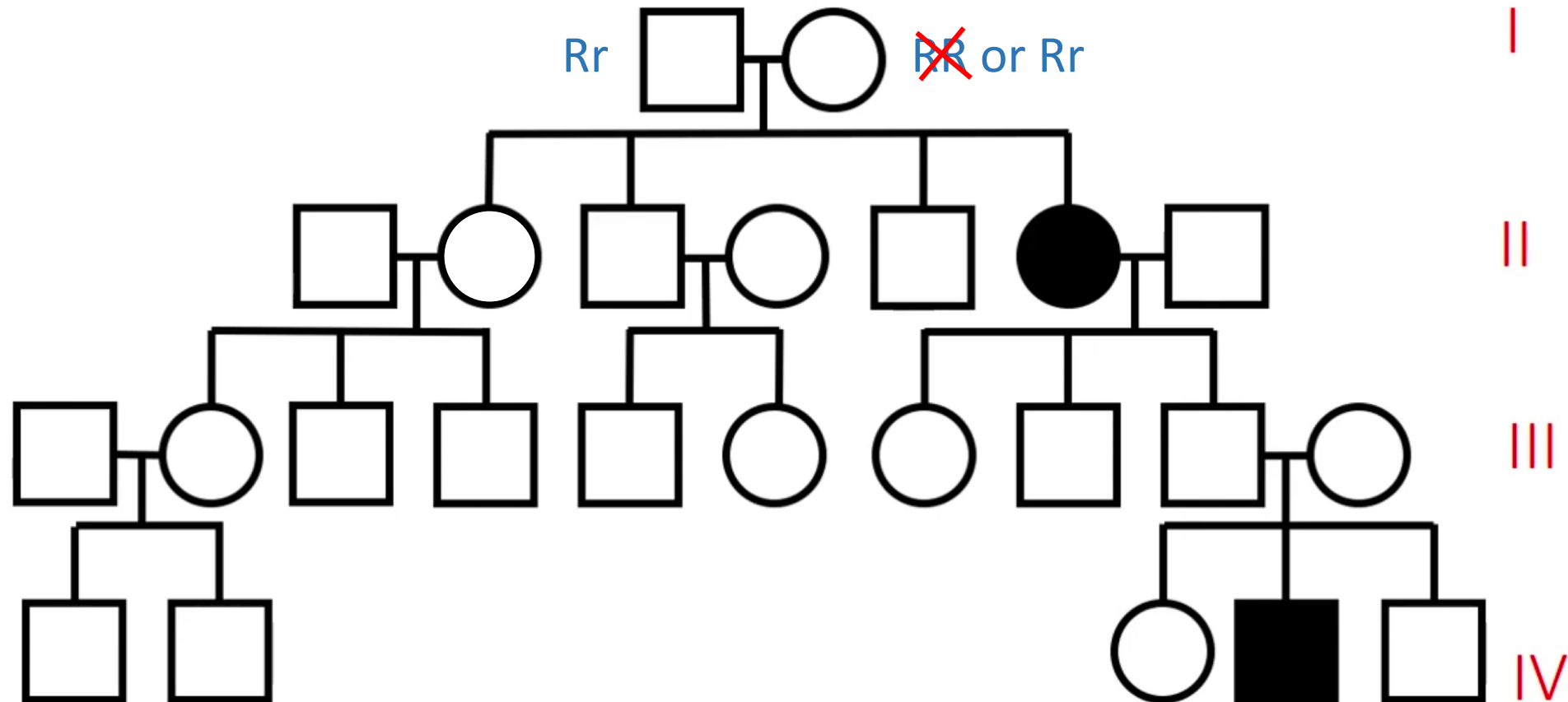
Remember: it is the recessive allele  $r$  that expresses the disease

# Punnett squares for gen II (gen 1 = parents)

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

0% of F1 would be affected (none has genotype rr)

# Pedigree for autosomal recessive trait (with genotypes)



# Punnett squares for gen II (gen 1 = parents)

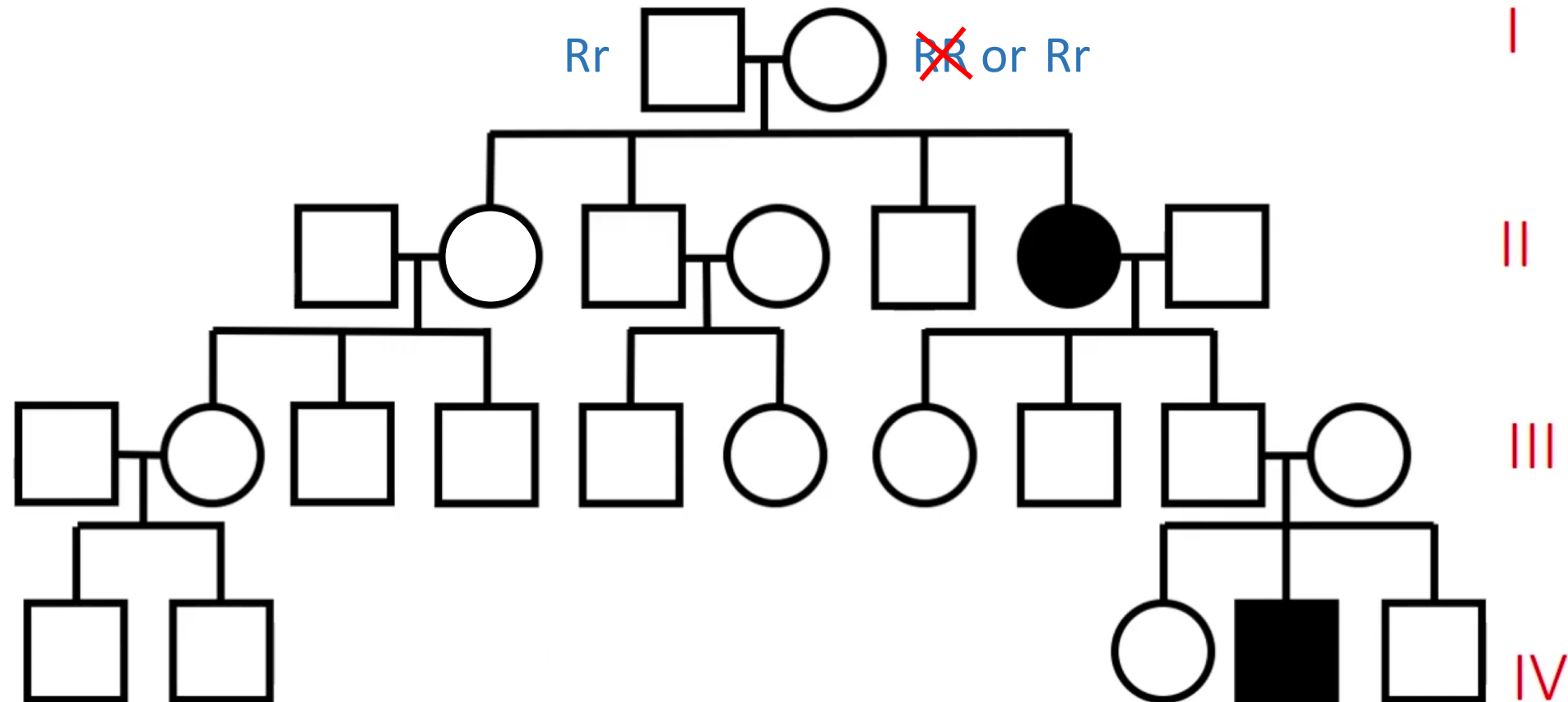
		Gametes	
	○	R	R
□		RR	RR
G a m e t e s	R		
	r	Rr	Rr

0% of F1 would be affected (none has genotype rr)

		Gametes	
	○	R	r
□		RR	Rr
G a m e t e s	R		
	r	Rr	rr

25% of F1 (1:3) would be affected (only 1 has rr)

# Pedigree for autosomal recessive trait (with genotypes)



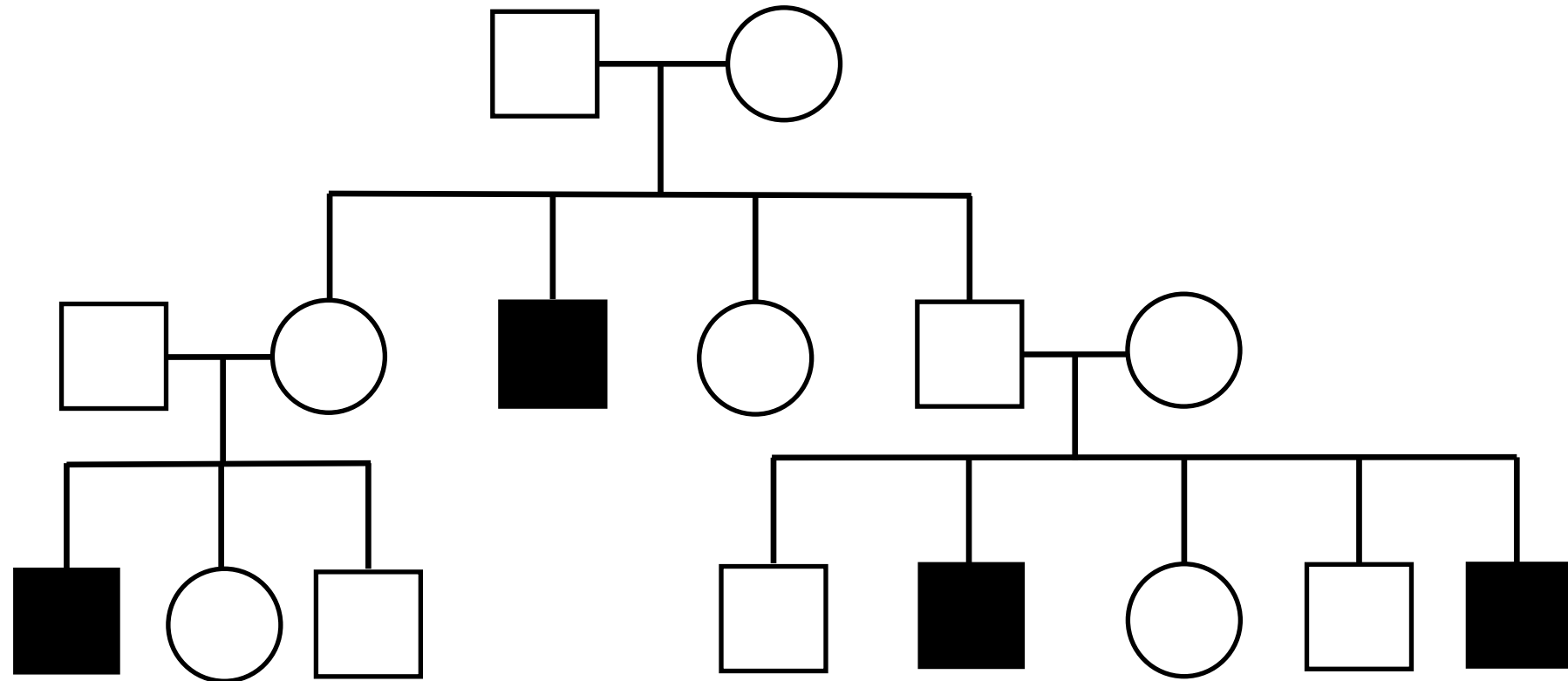
Note: you may not always get ratios identical to predicted due to probability of using specific gametes

# Pedigree for X-linked recessive trait

## Clues

**X-linked:** only males affected

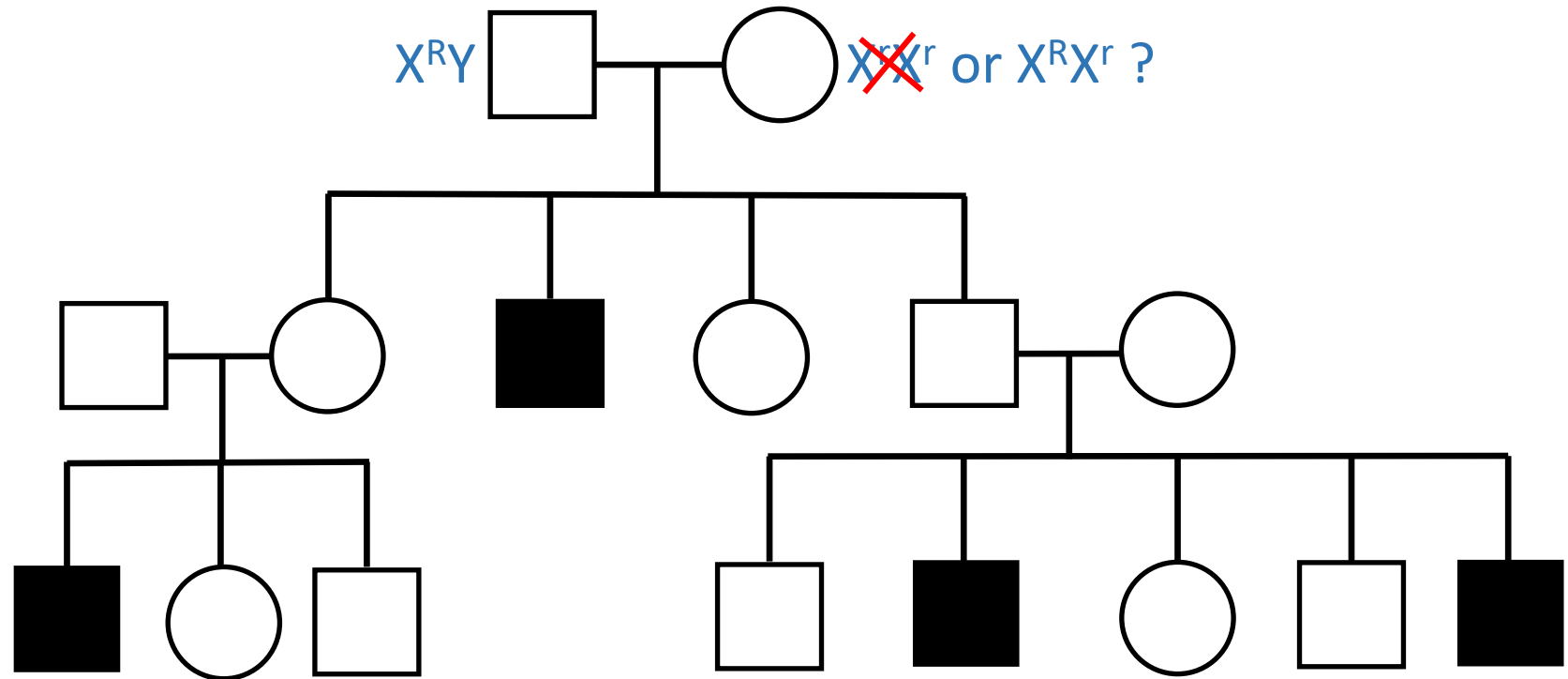
**Recessive:** affected child(ren) from unaffected parents  
not many affected offspring



# 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



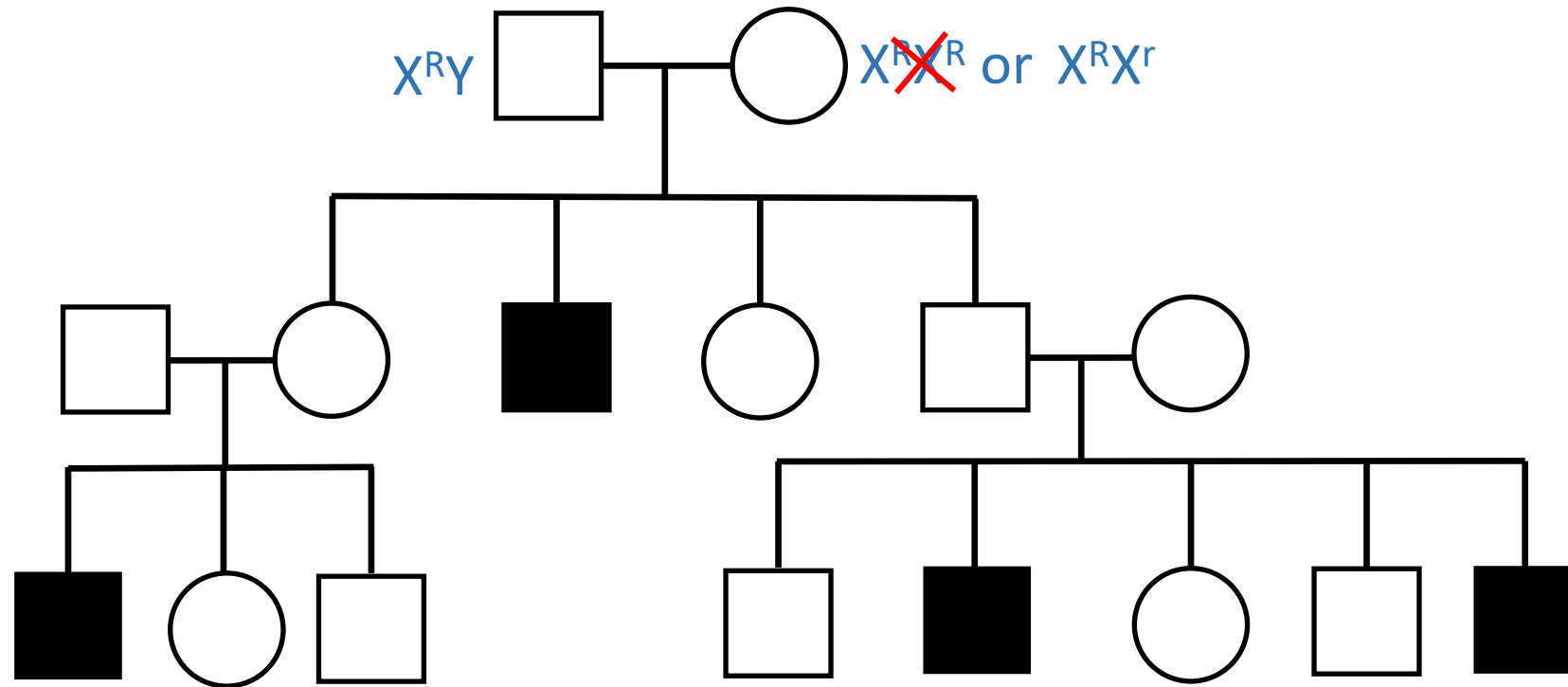
# Punnett squares for gen II (gen 1 = parents)

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

50% of MALE F1 (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 due to probability of using specific gametes

# Notes

- The discussed pedigrees are simple and ideal
- You may not always get ratios identical to predicted due to probability of using specific gametes
- You may not have enough offspring at F1 to interpret pedigree
  - You need to analyze F2, F3, ...

# Pedigrees

- Take assignment 16: **Pedigrees**