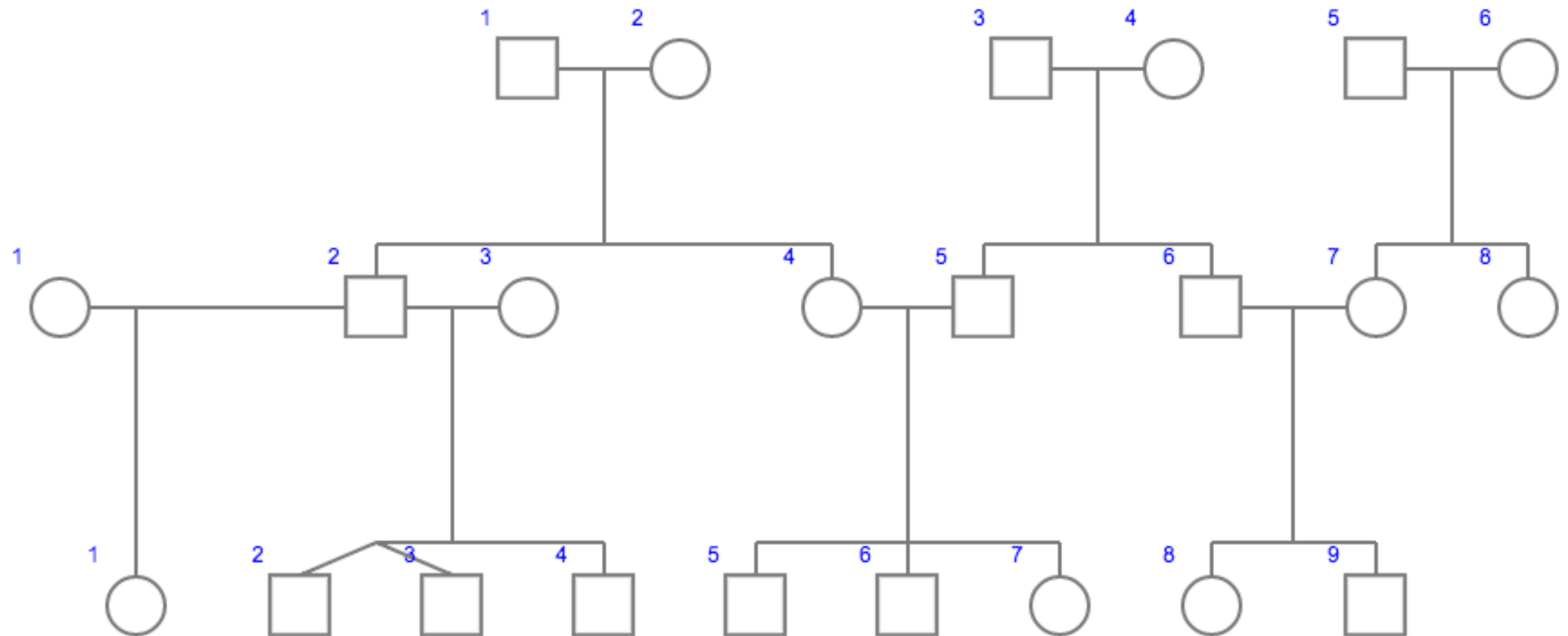
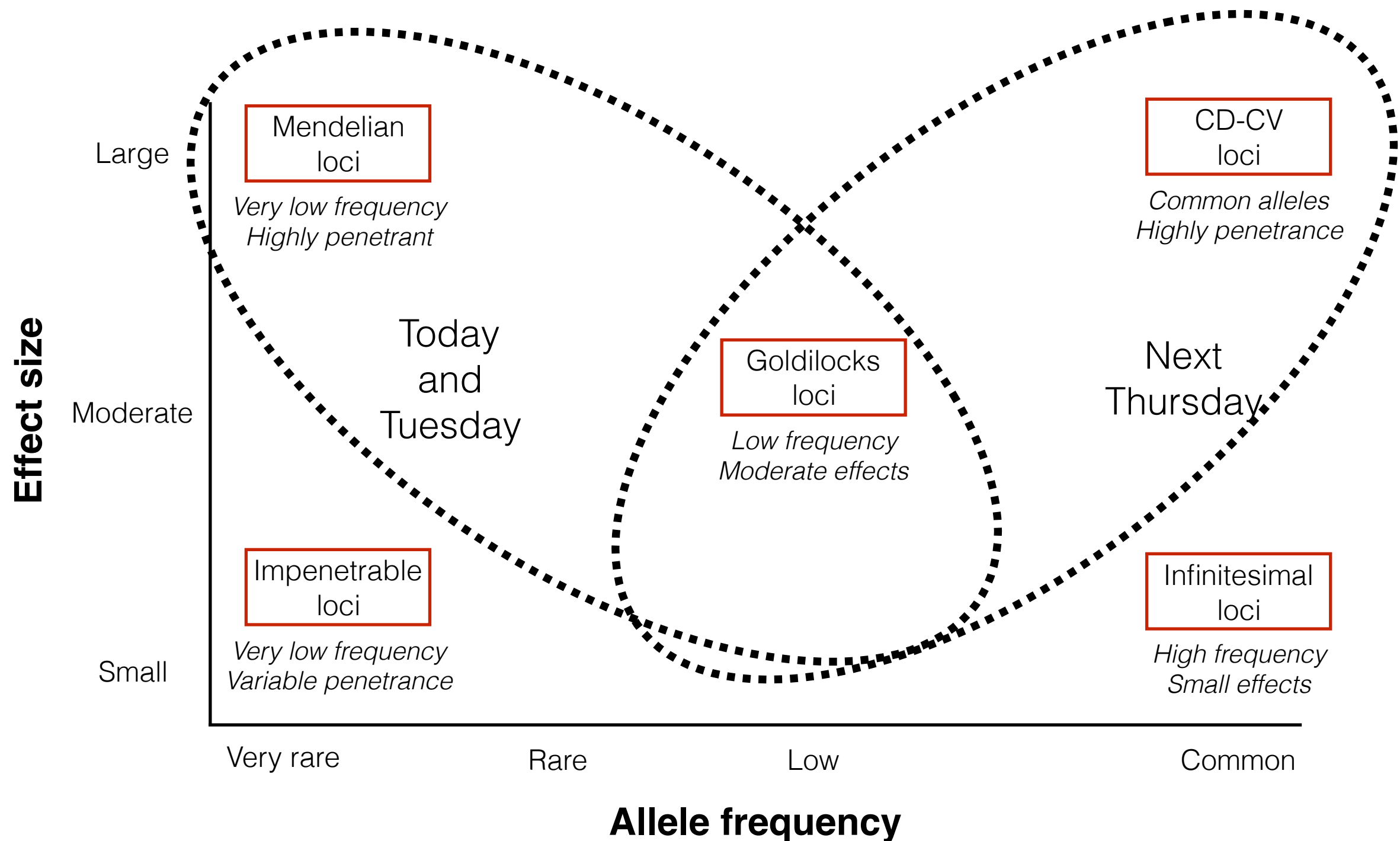


# Bio393: Genetic Analysis

Family-based analysis, Modes of inheritance, Phase

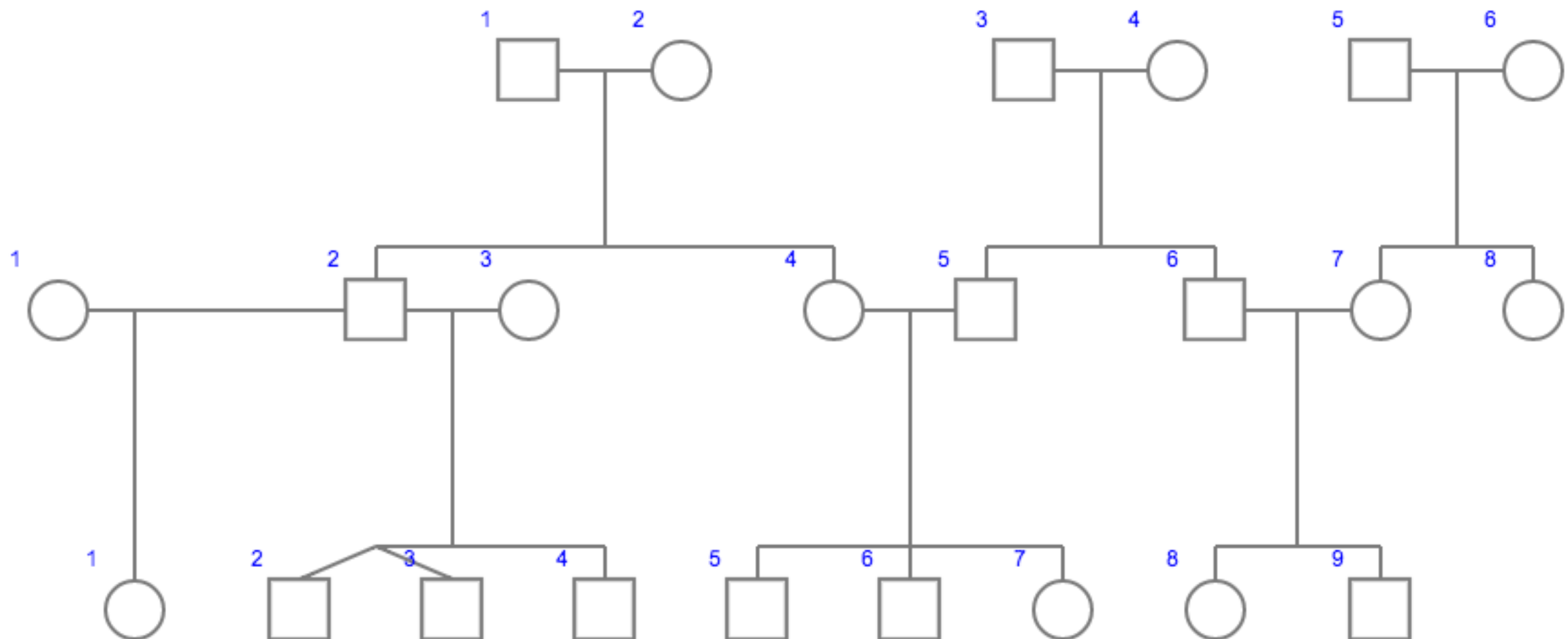


# The spectrum of how variation contributes to disease



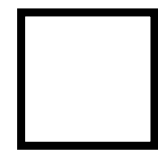
Linkage mapping studies or family-based mapping studies

# Why do we study inheritance in families?

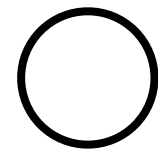


Correlating genetic variants with disease tells us the disease gene is near that variant (or is that variant)

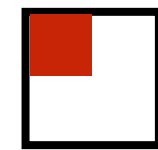
# Human pedigree analysis allows us to follow traits in families



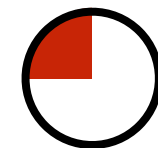
Male



Female



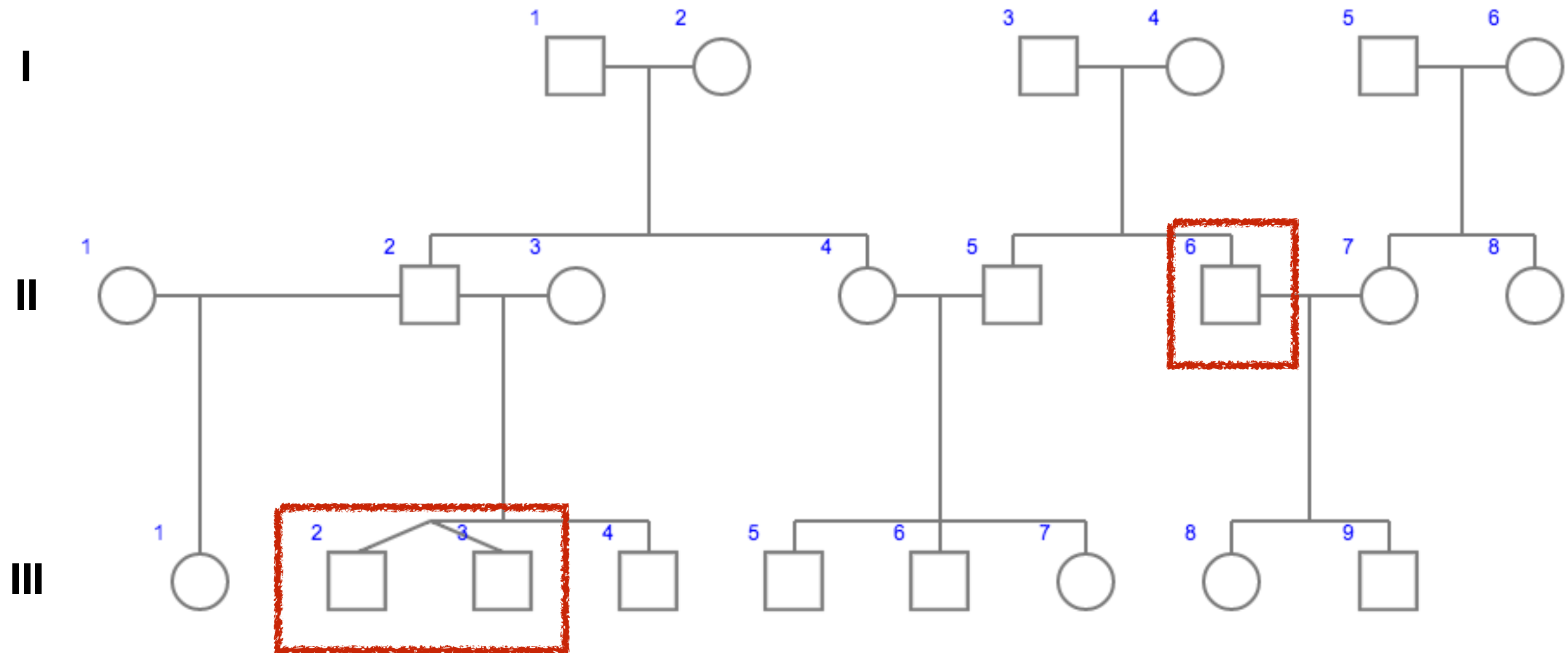
Affected male



Affected female

Remember that humans are diploid.

# Human pedigree analysis allows us to follow traits in families

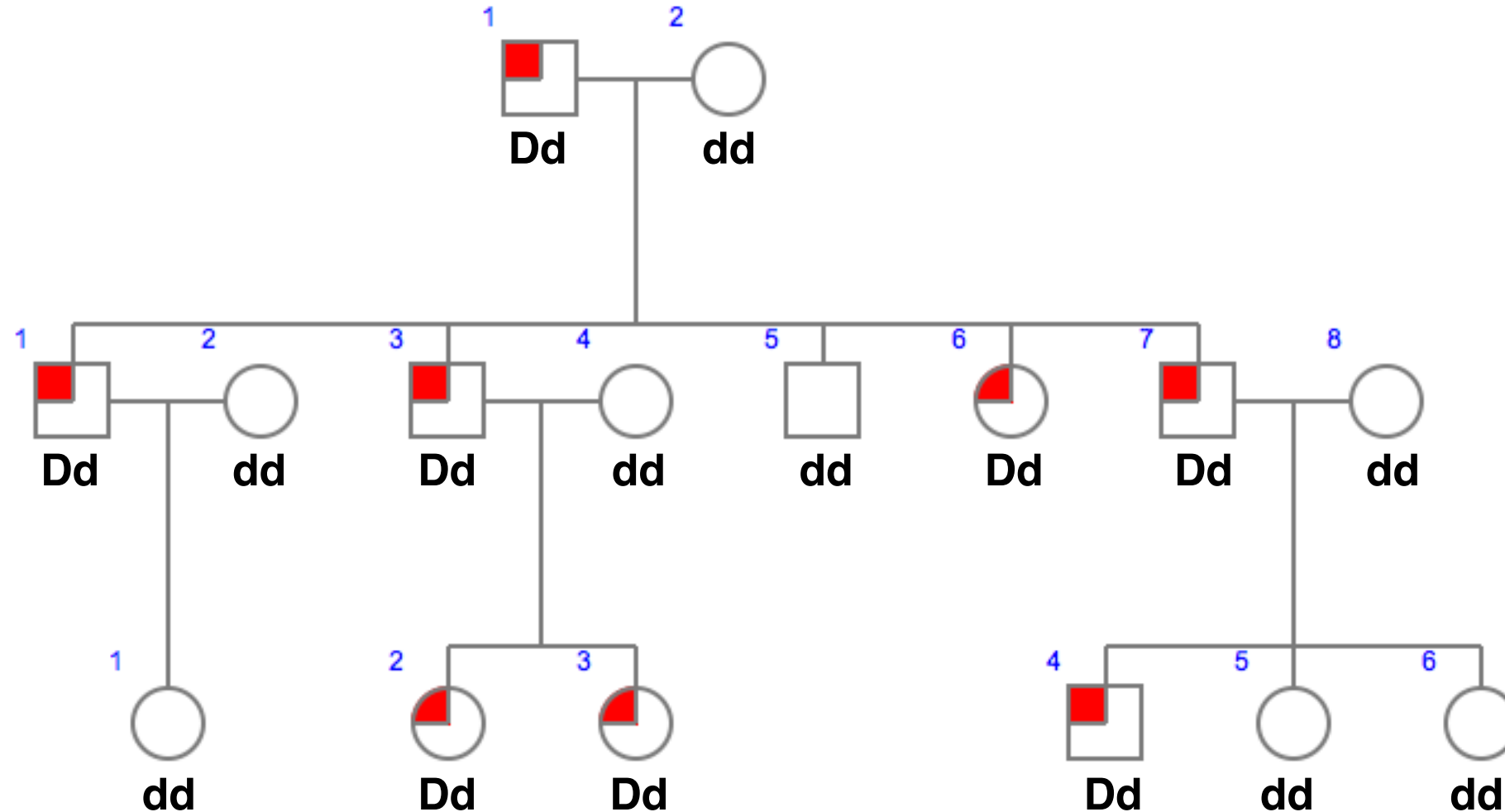


Individuals are numbered from left to right

Generations are numbered from top to bottom in Roman numerals

Most diseases are rare, individuals breeding into families are usually unaffected

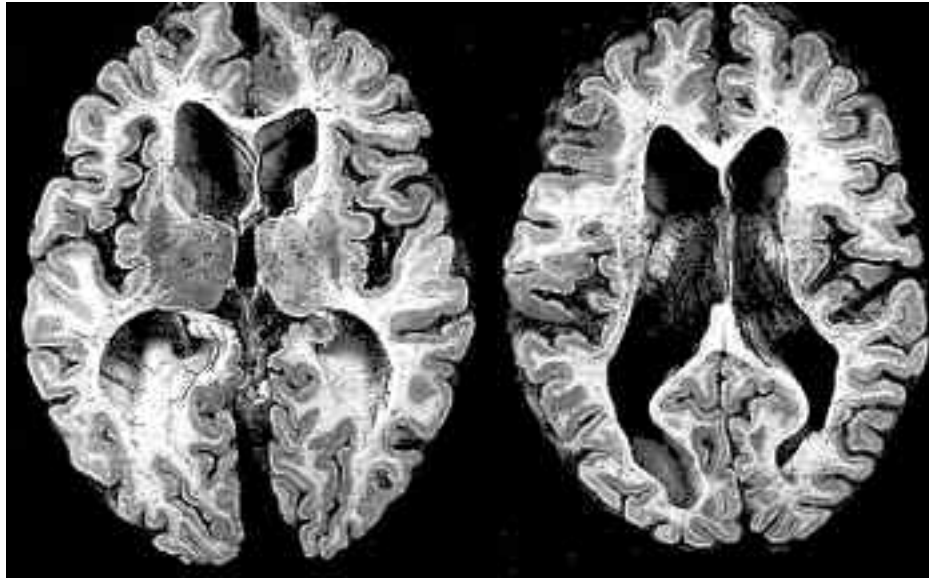
# Modes of inheritance



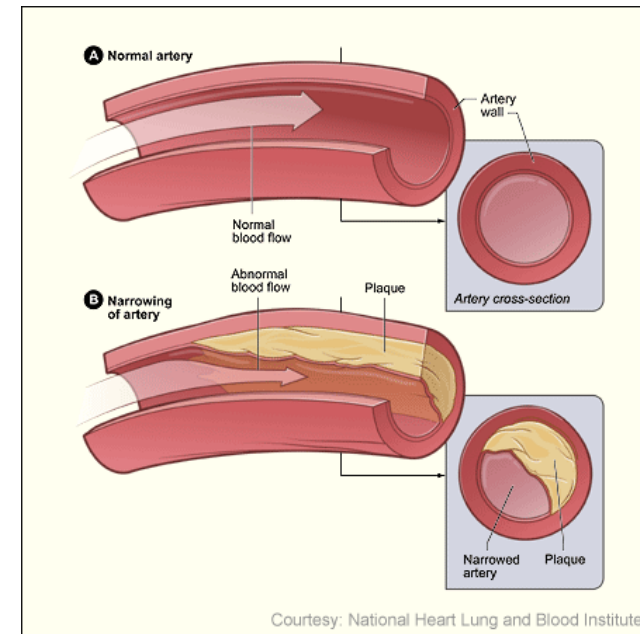
- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

## Autosomal dominant

# Examples of human autosomal dominant disorders



Huntington's Disease  
chr. 4



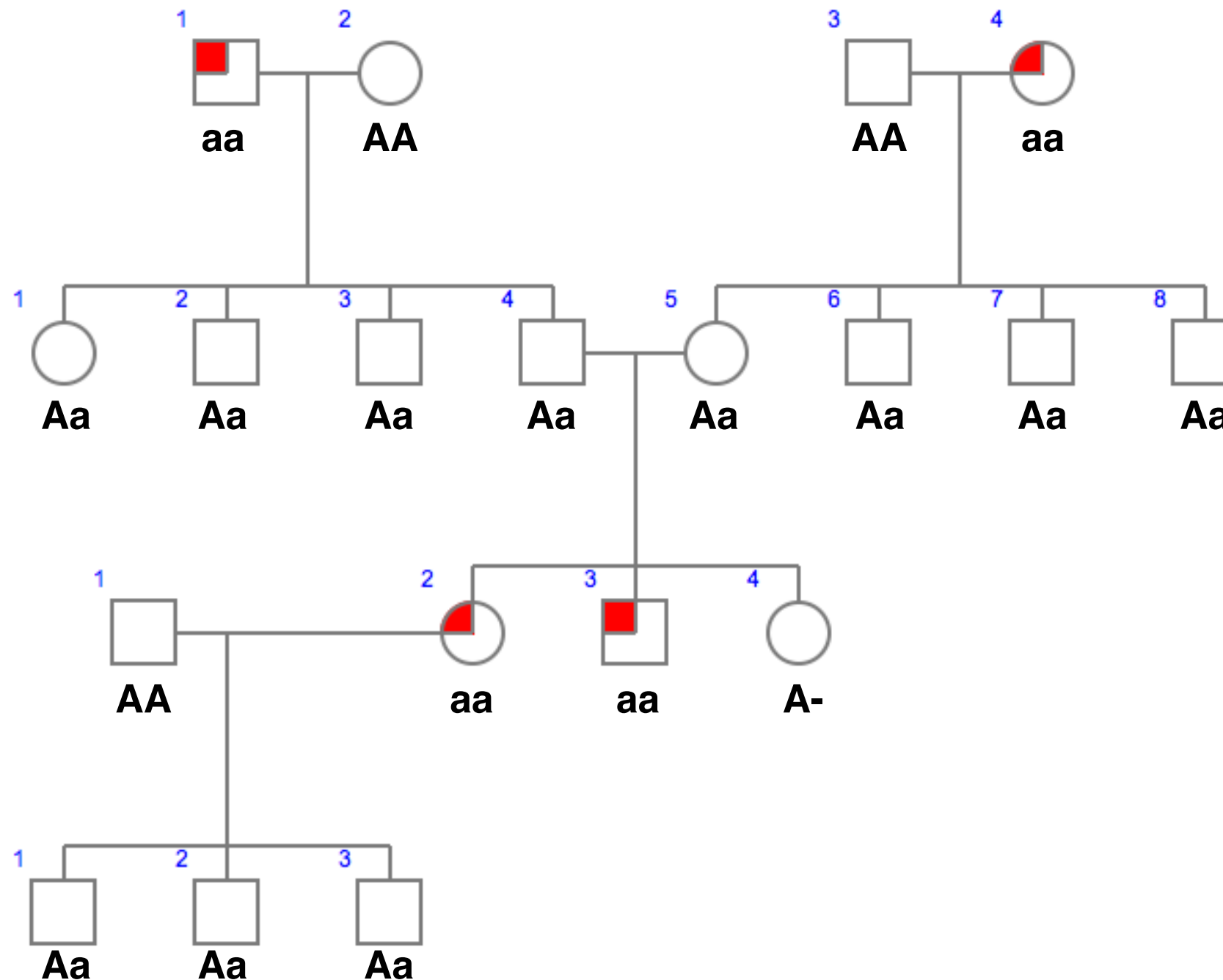
Familial Hypercholesterolemia  
chr. 19

Caused by loss-of-function or gain-of-function?

Most affected individuals are heterozygotes

What is the chance that a child is affected?

# Modes of inheritance

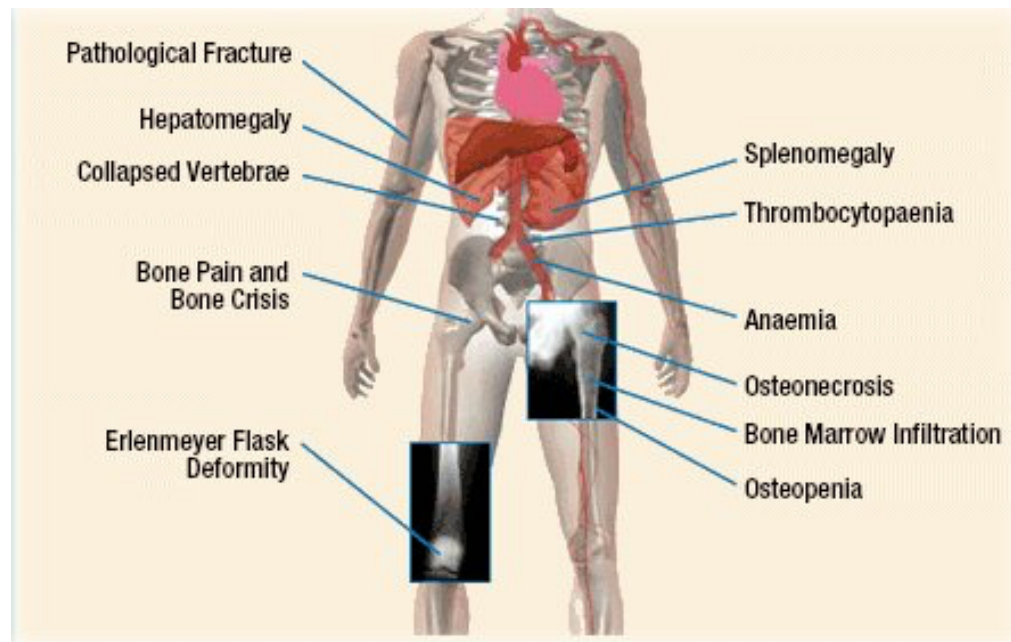


- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

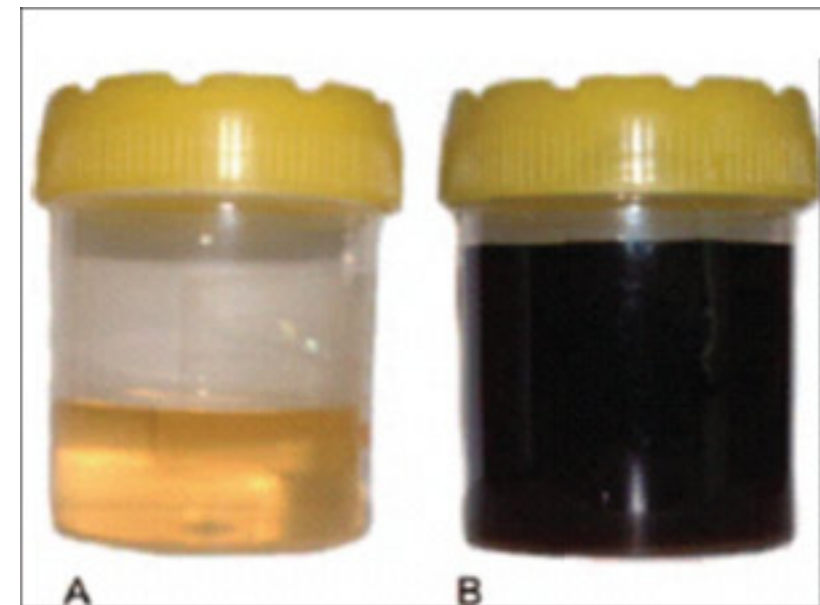
## Autosomal recessive



# Examples of human autosomal recessive disorders



Gaucher's Disease  
chr. 1

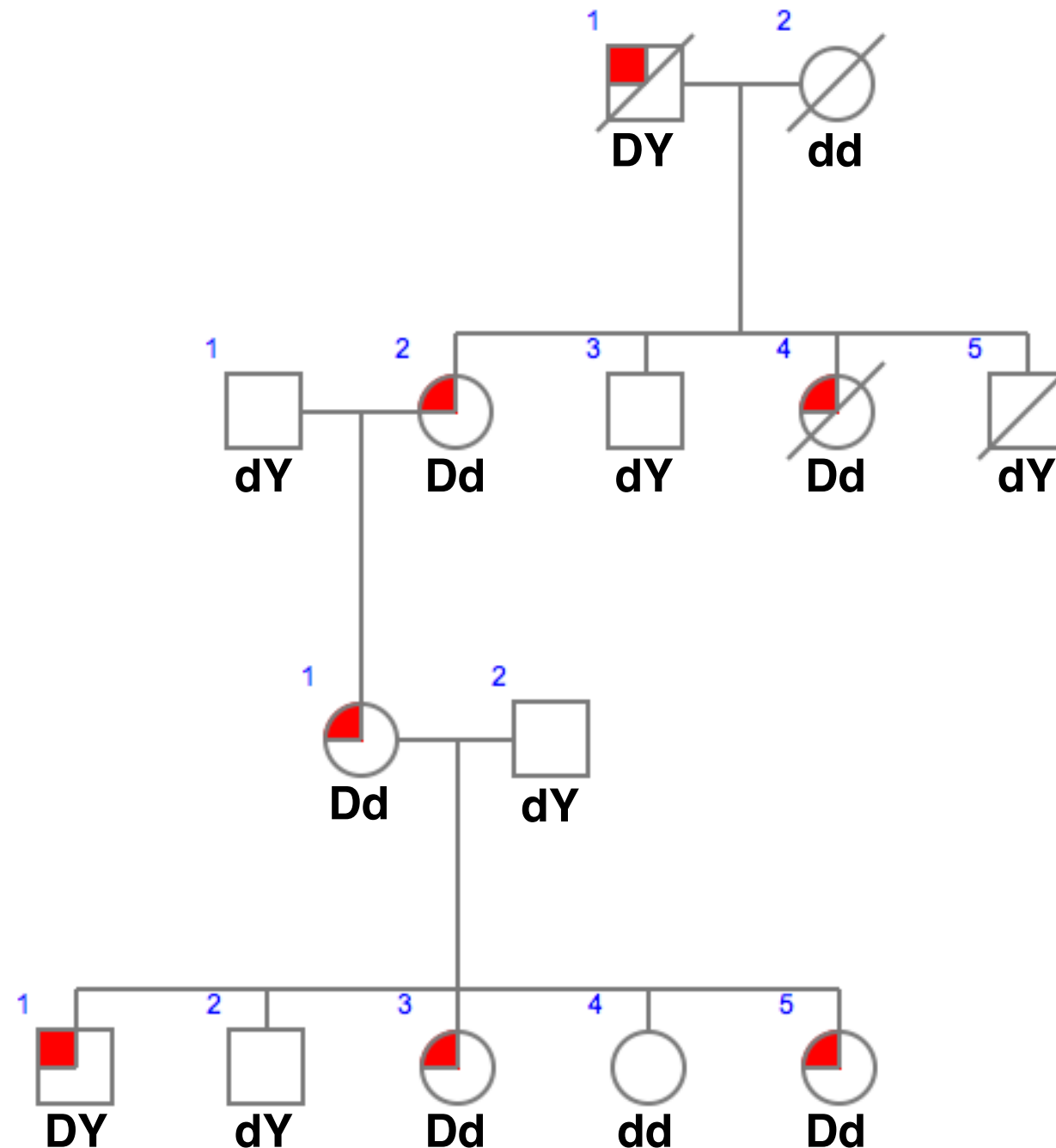


Maple Syrup Urine Disease  
chr. 1, 6, or 19

Caused by loss-of-function or gain-of-function?

All affected individuals are homozygotes

# Modes of inheritance



- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

## X-linked dominant

# Examples of human X-linked dominant disorders



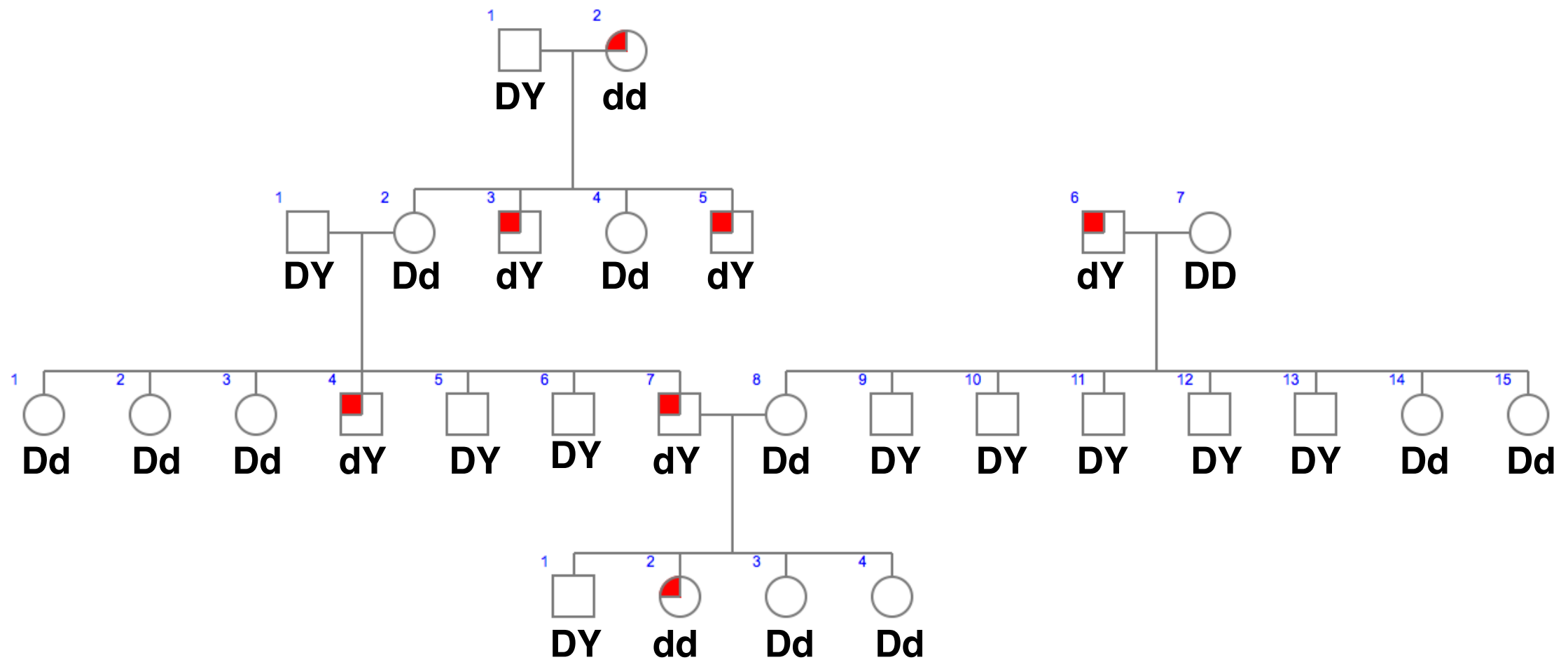
Rett syndrome



Fragile X syndrome

All daughters of affected fathers are affected

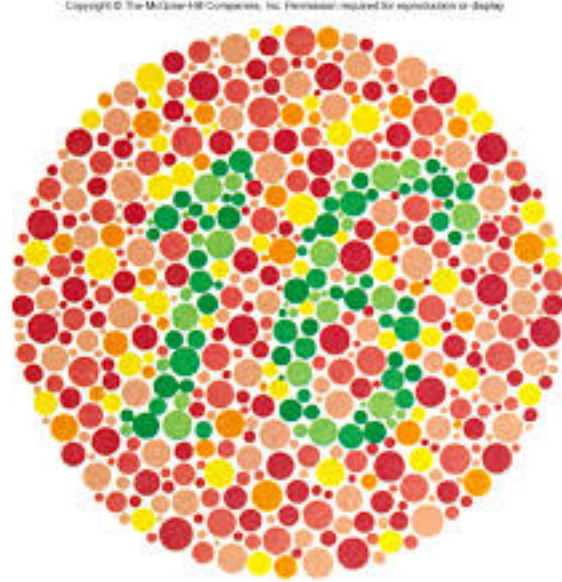
# Modes of inheritance



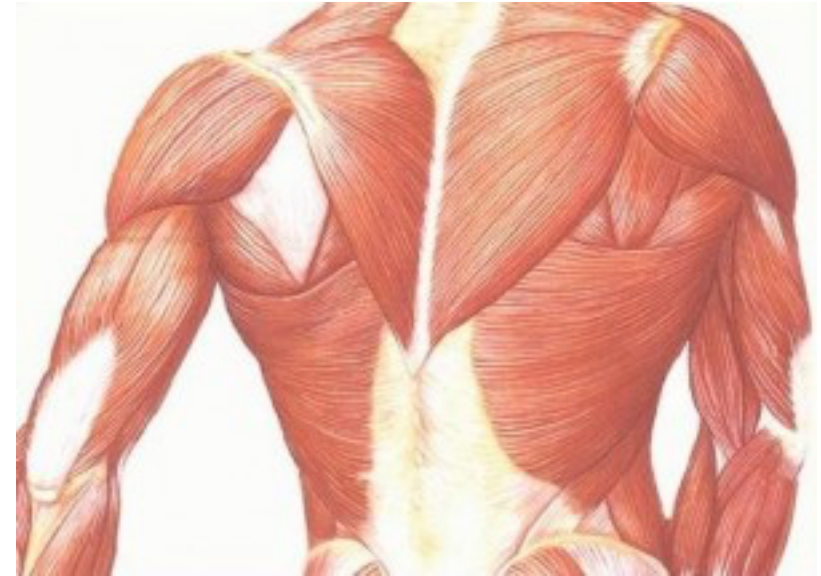
- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

## X-linked recessive

# Examples of human X-linked recessive disorders



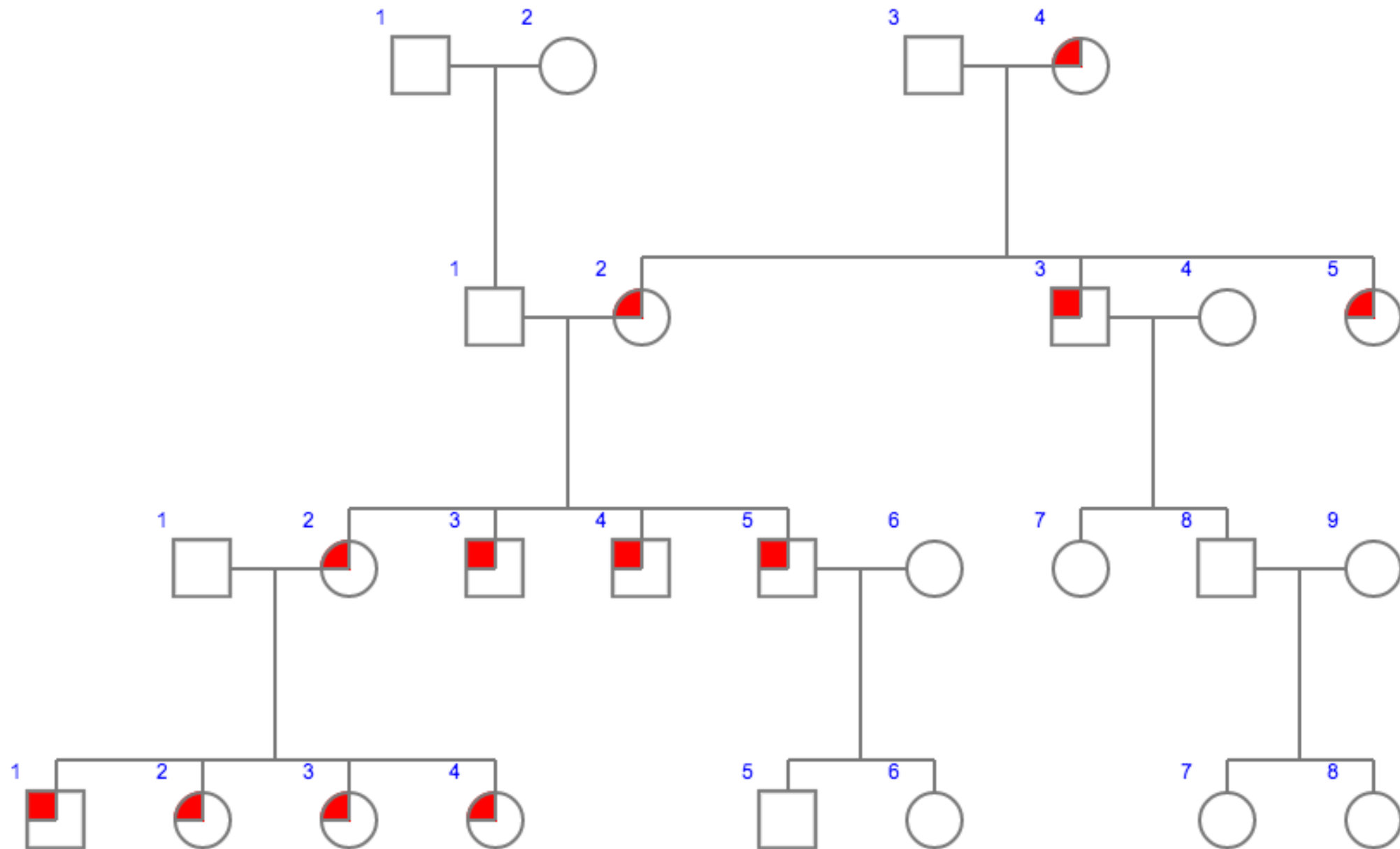
Red-green color blindness



Duchenne muscular dystrophy

All sons of affected mothers are affected

# Modes of inheritance



- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

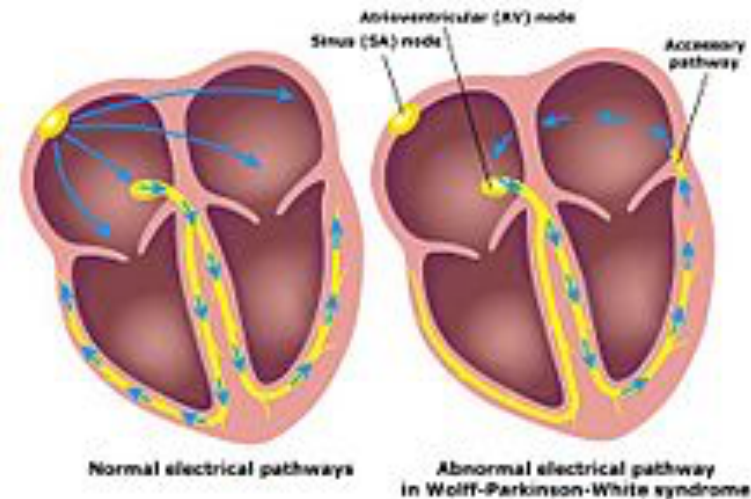
## Cytoplasmic inheritance



# Examples of human cytoplasmic inheritance disorders



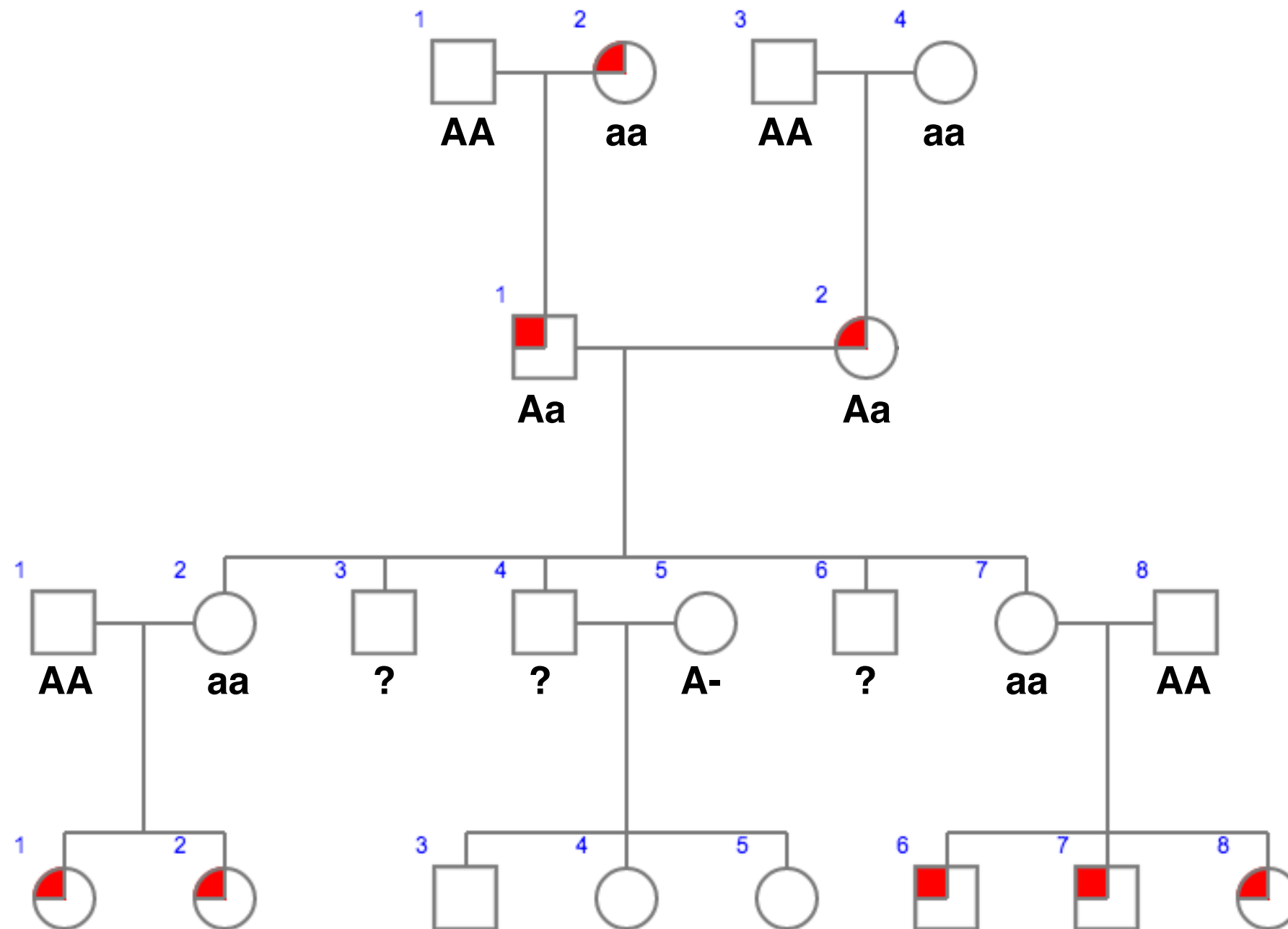
Mitochondrial  
myopathy



Wolff-Parkinson-White  
syndrome

All children of affected mothers are affected

# Modes of inheritance

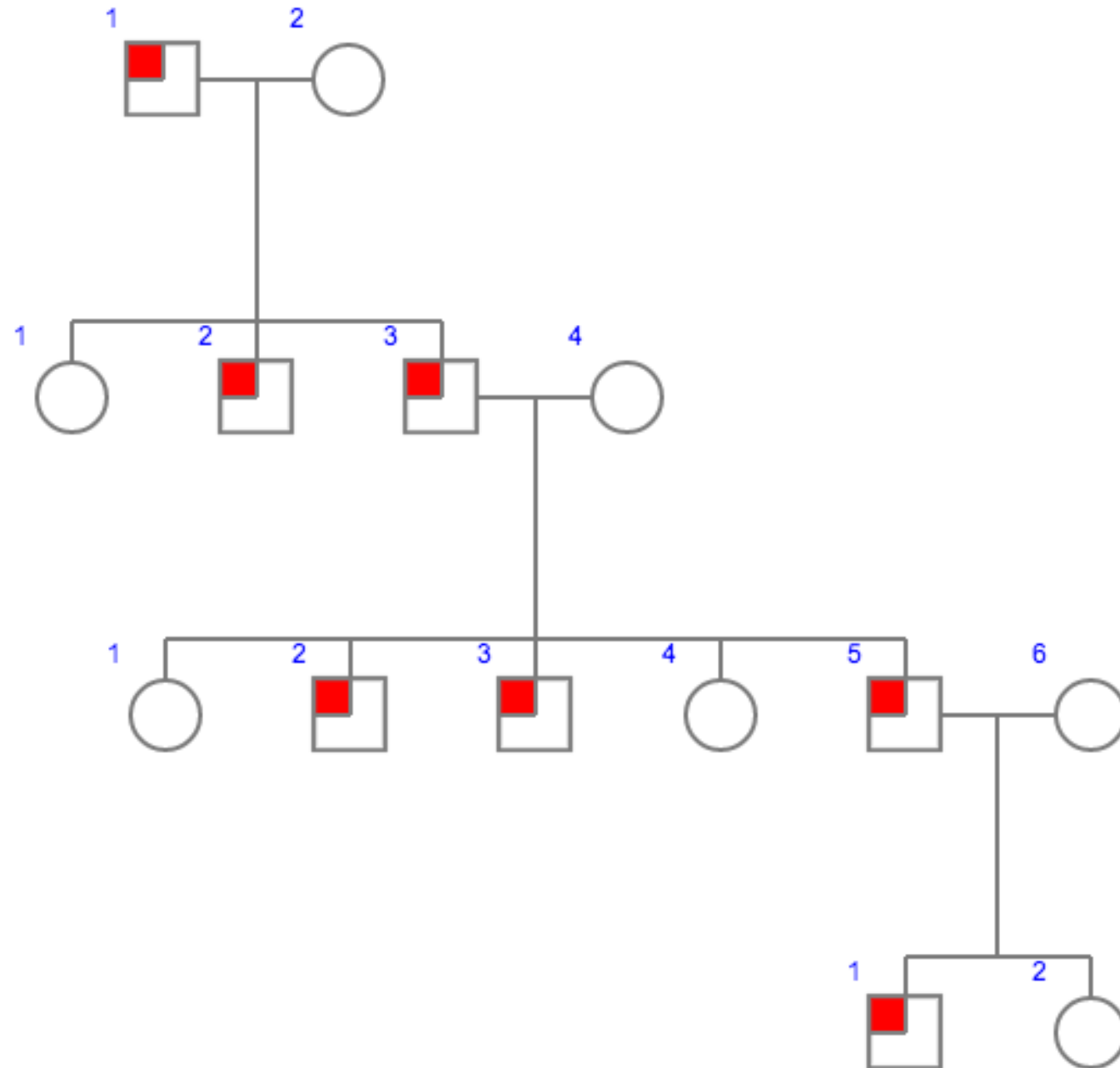


- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

## Recessive maternal-effect inheritance



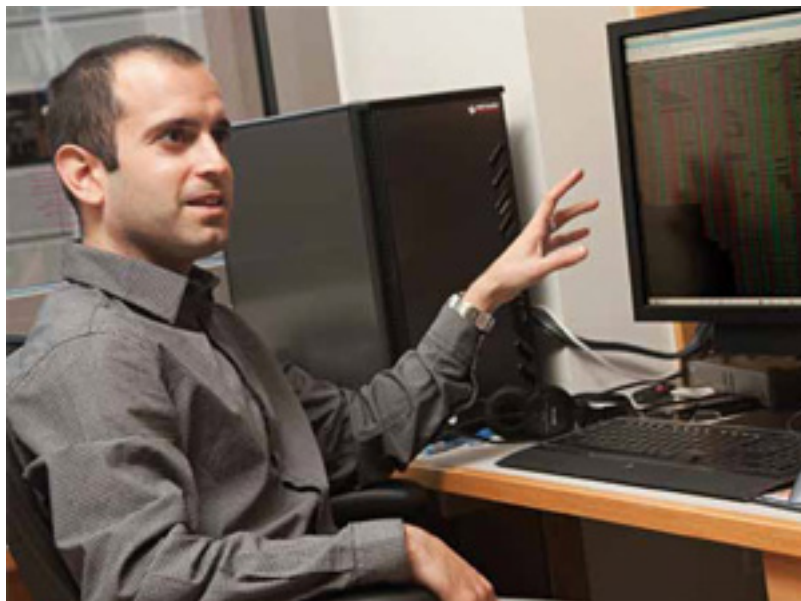
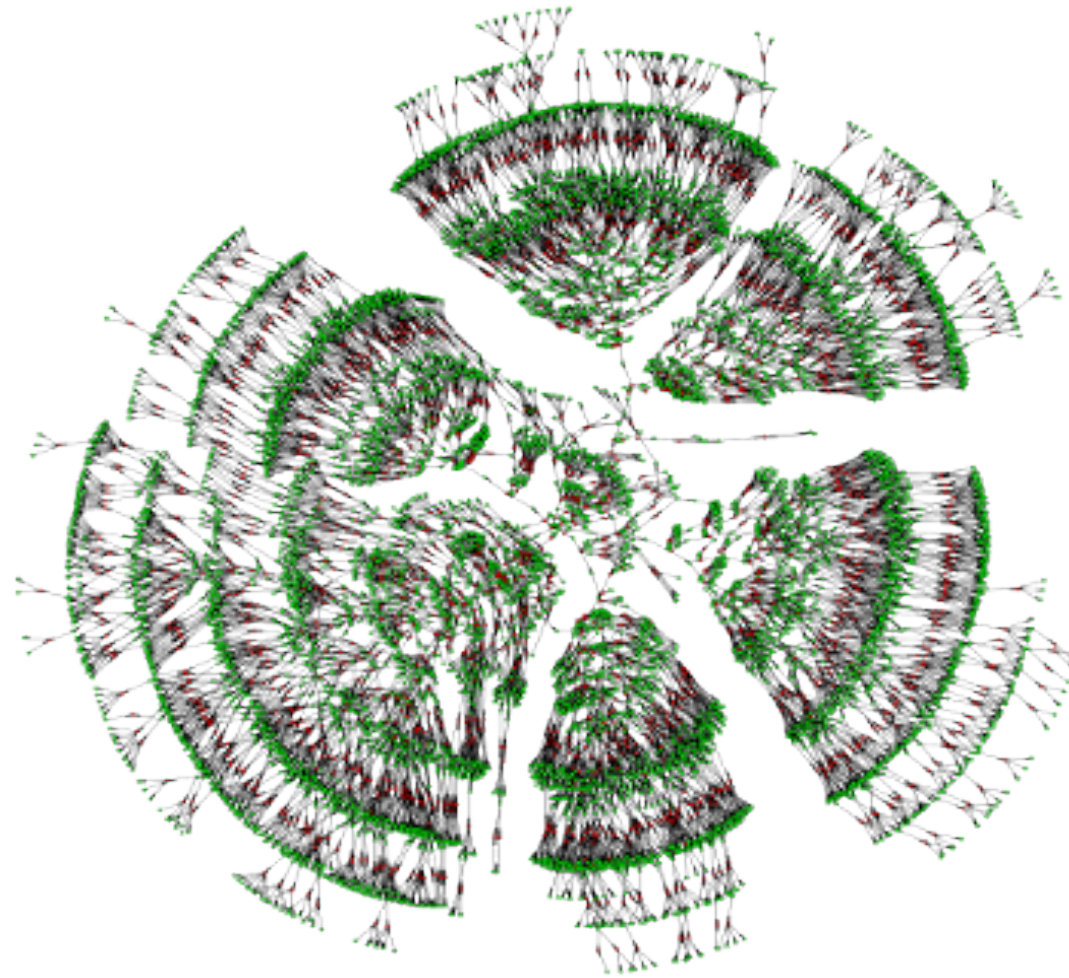
# Modes of inheritance



- How many individuals are affected?
- In each generation?
- Are males preferentially affected from affected mothers?
- Are females preferentially affected from affected fathers?

## Y-linked inheritance

# Some pedigrees can contain millions of individuals

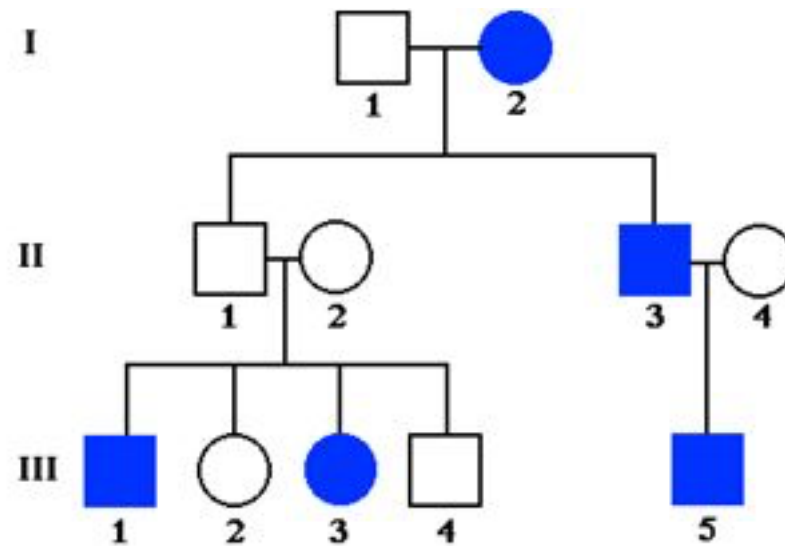


Yaniv Erlich

Ancestry websites offer rich family data



# Remember all of the genetics we've learned so far



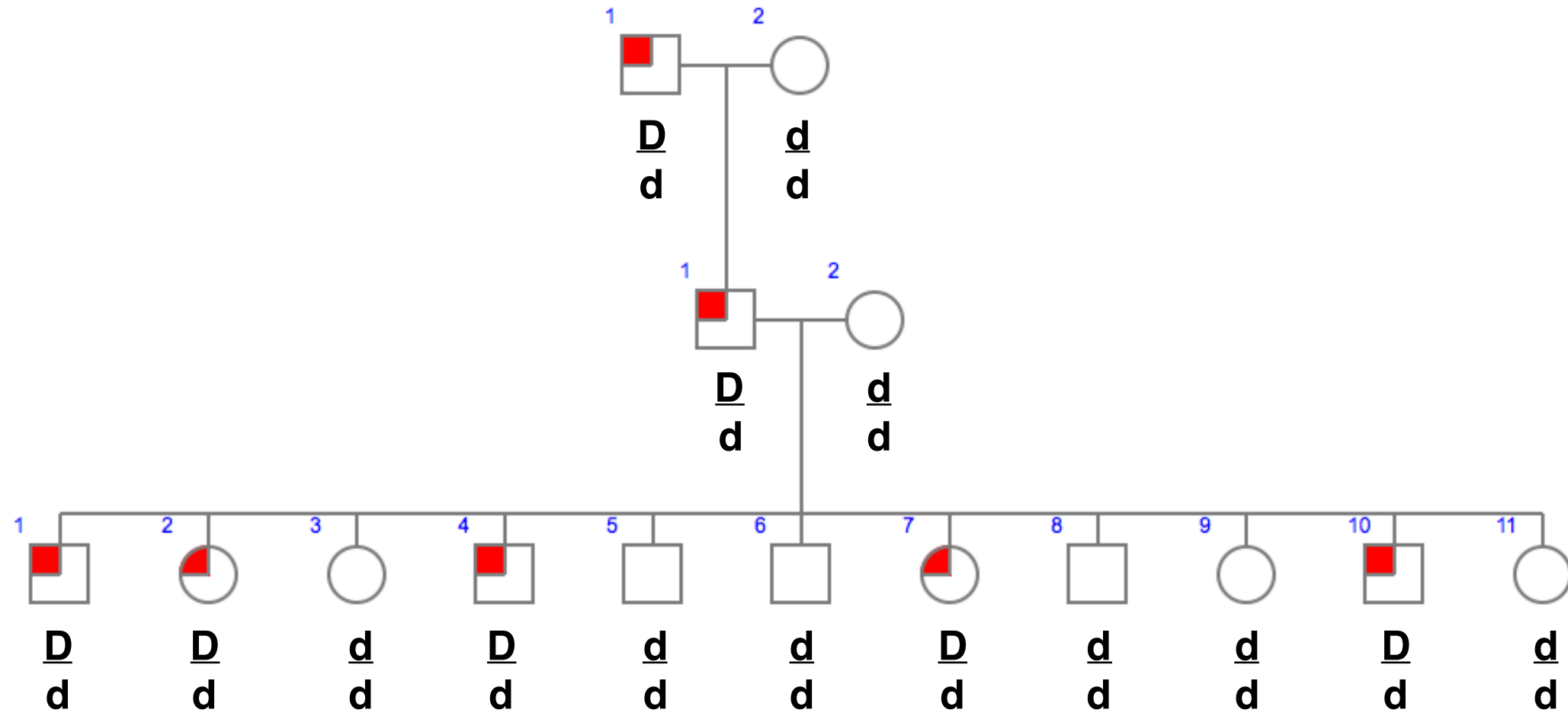
Incomplete penetrance

Non-complementation

Haploinsufficiency

Suppression and enhancement

**We want to be able to find a marker linked with disease to identify the disease gene**



Autosomal dominant

# Genetic variants are used as markers to track disease

Single nucleotide  
variants (SNVs)


Reference ATGTGCAGACGTAGACGTA

Alternative ATGTGCAGACTTAGACGTA

Insertion-deletion  
variants (indels)

Reference ATGTGCAGACGTAGACGTA

Alternative ATGTGCAGACGTAGACGTA

  
Addition of 126 bp

Copy-number  
variants (CNVs)

Reference Diploid (2 copies)

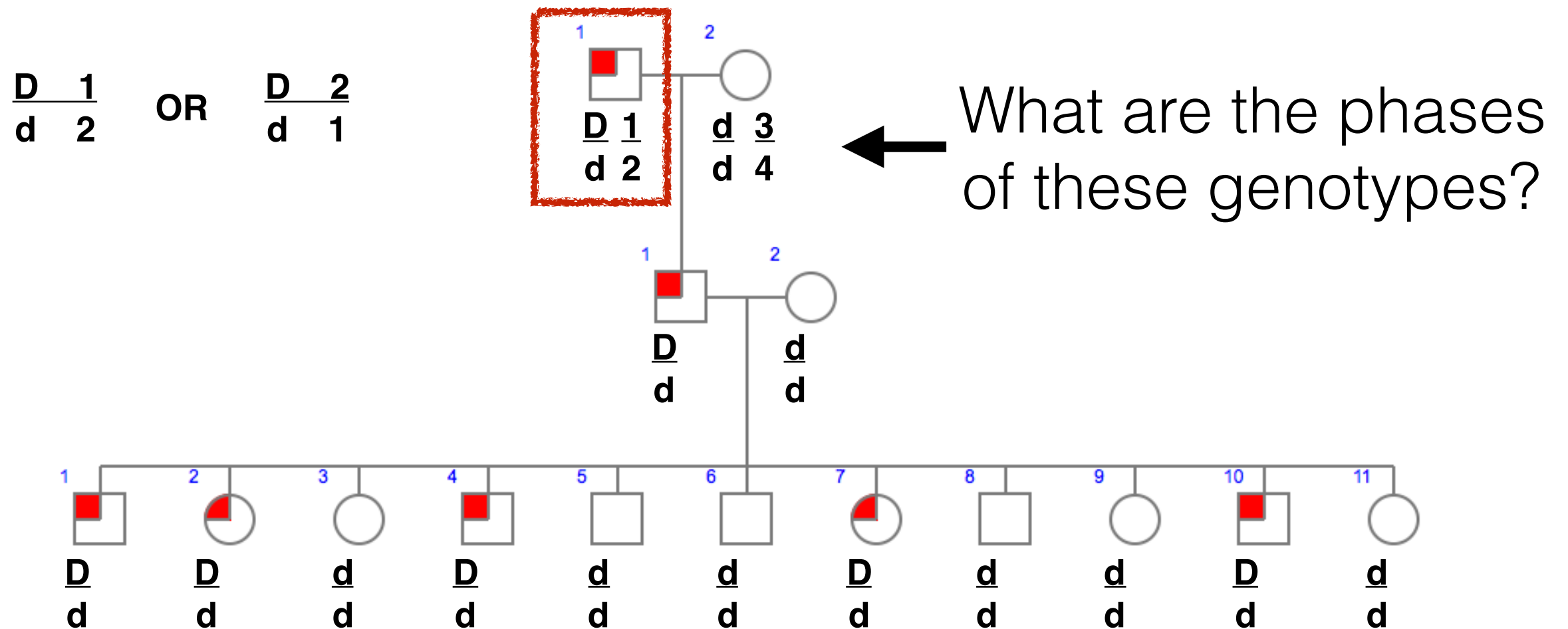
Alternative More (or fewer) than 2 copies

Microsatellites  
or short tandem repeats (STRs)

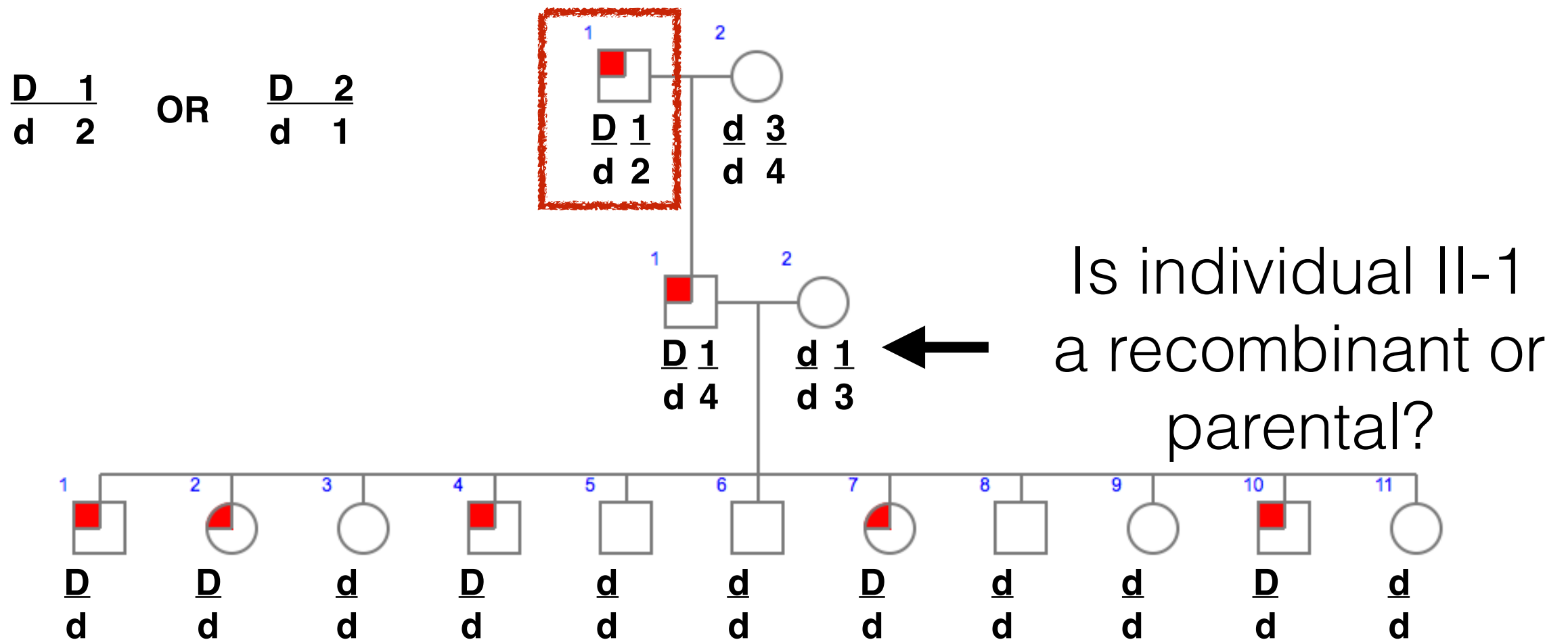
Reference ATGTGCAGCAGCAGCAGCGTA

Alternative ATGTGCAGCAGCGTAGTGACT

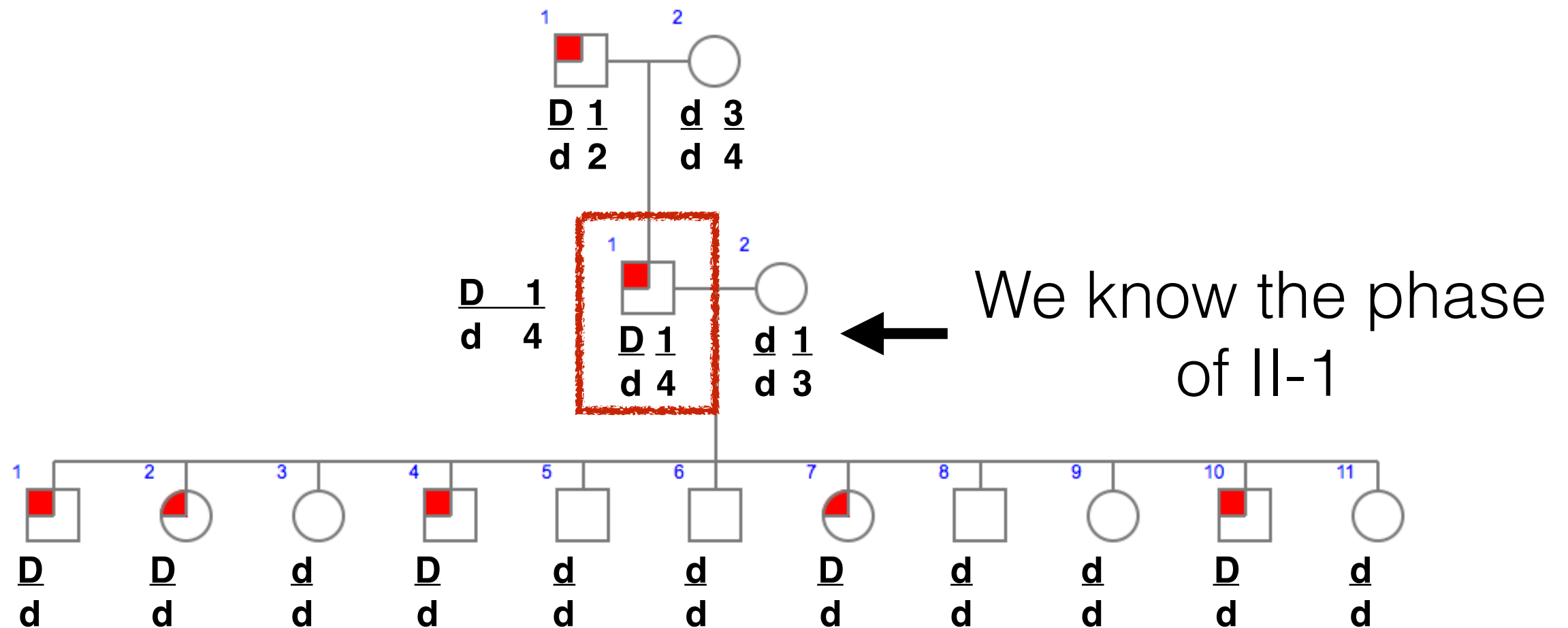
# Linkage to genetic markers tells us where disease genes are



# Linkage to genetic markers tells us where disease genes are

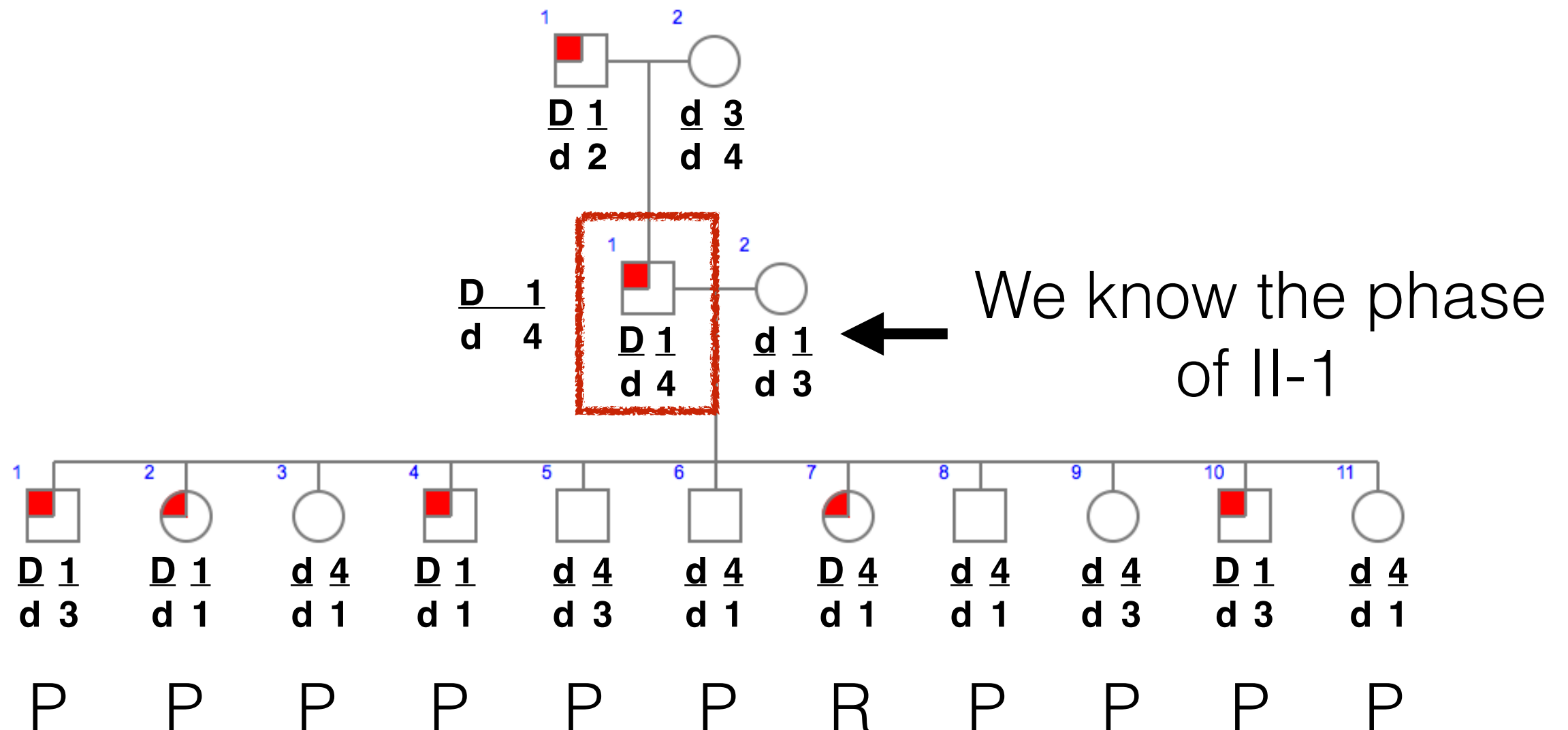


# Linkage to genetic markers tells us where disease genes are

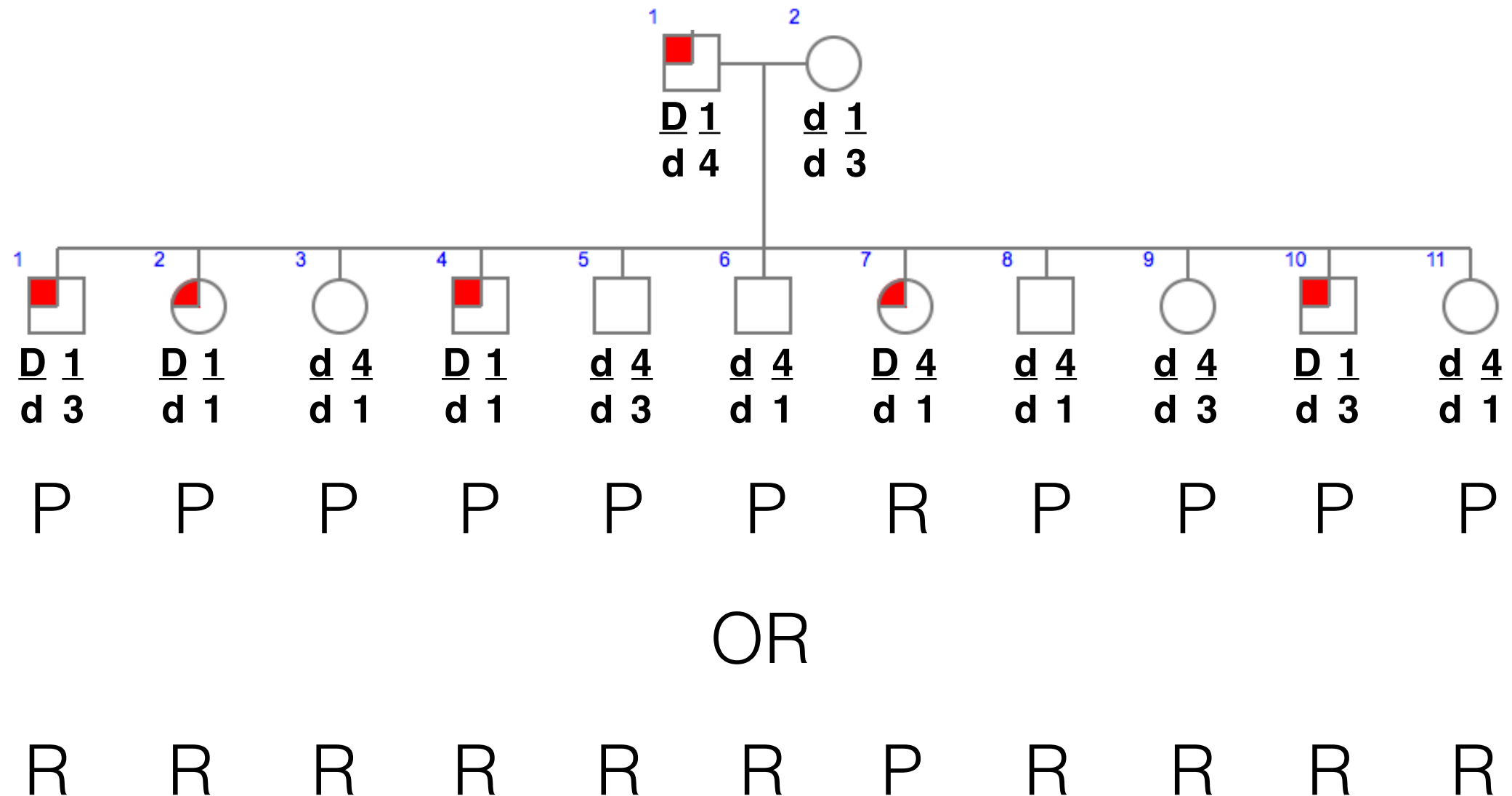




# Linkage to genetic markers tells us where disease genes are



# Linkage to genetic markers tells us where disease genes are



Sometimes, we don't know the phase of the parent, and both possibilities of phase are equally likely