

Chromosomal Abnormalities

Part 2

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Abnormalities in Chromosome Structure

- Structural abnormalities are rearrangements in chromosomes that can originate either *de novo* (new in proband) or familial (passed through generations)
- Structural abnormalities include:
 - Translocations
 - Inversions
 - Deletions
 - Duplications
- When a structural abnormality is detected, parents should be karyotyped to determine if the abnormality is familial

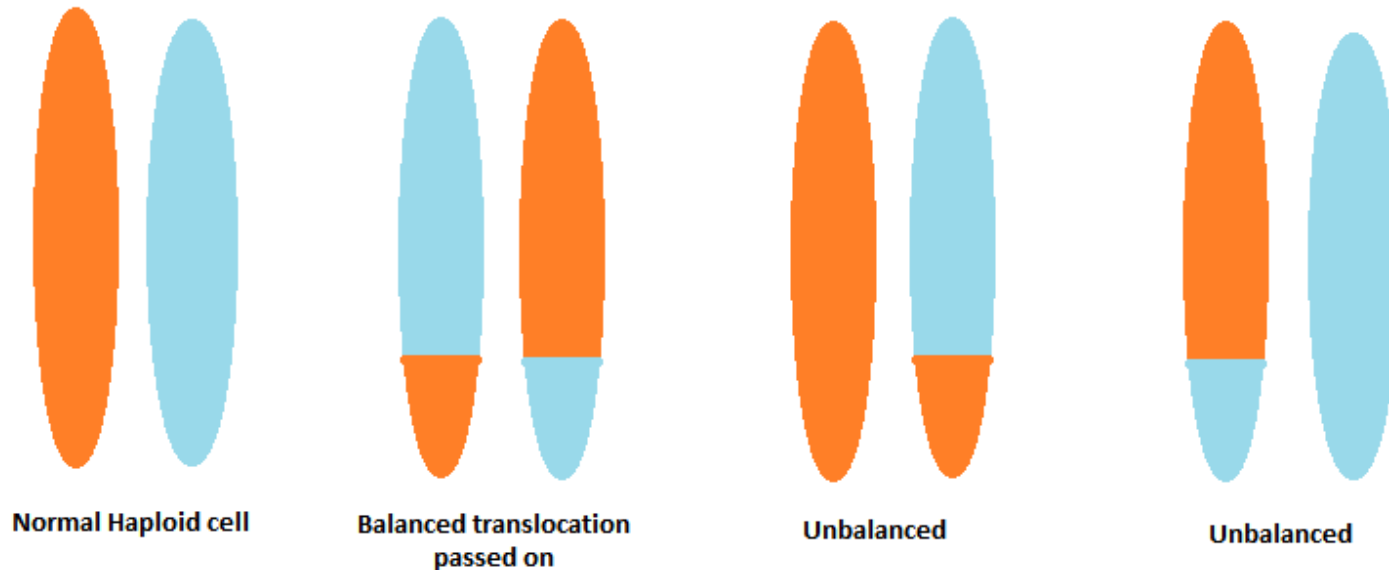
Chromosomal Rearrangements

- Overall present in 1 in 375 newborns
- Result from chromosome breakage, recombination, or exchange, followed by reconstitution in an abnormal combination
- Types of rearrangements: Balanced vs. unbalanced
- Best detected on karyotype or FISH, occasionally microarray if unbalanced

Chromosomal Rearrangements

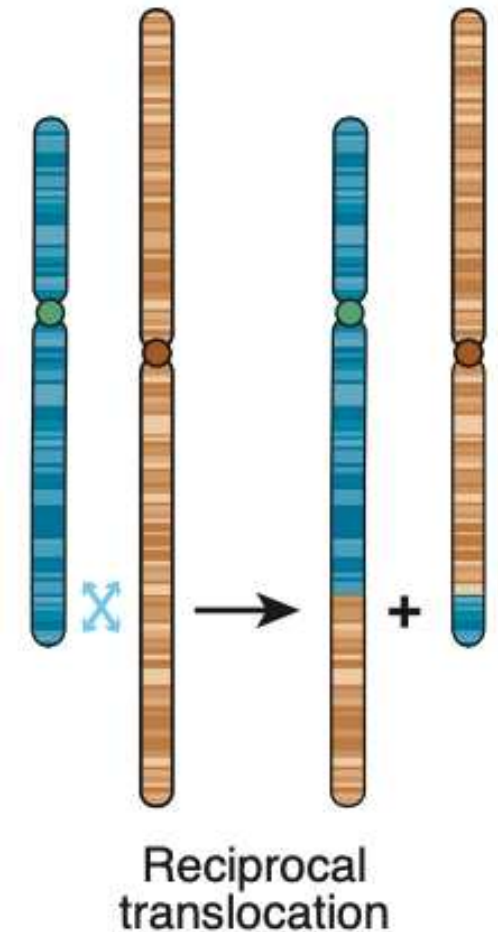
- **Balanced:** all genetic material remains present
 - Typically **normal** phenotype
 - **Increased risk for abnormal offspring**
 - 1 in 500 people

- **Unbalanced:** either gain or loss of material
 - Typically **abnormal** phenotype
 - 1 in 1,600 live births



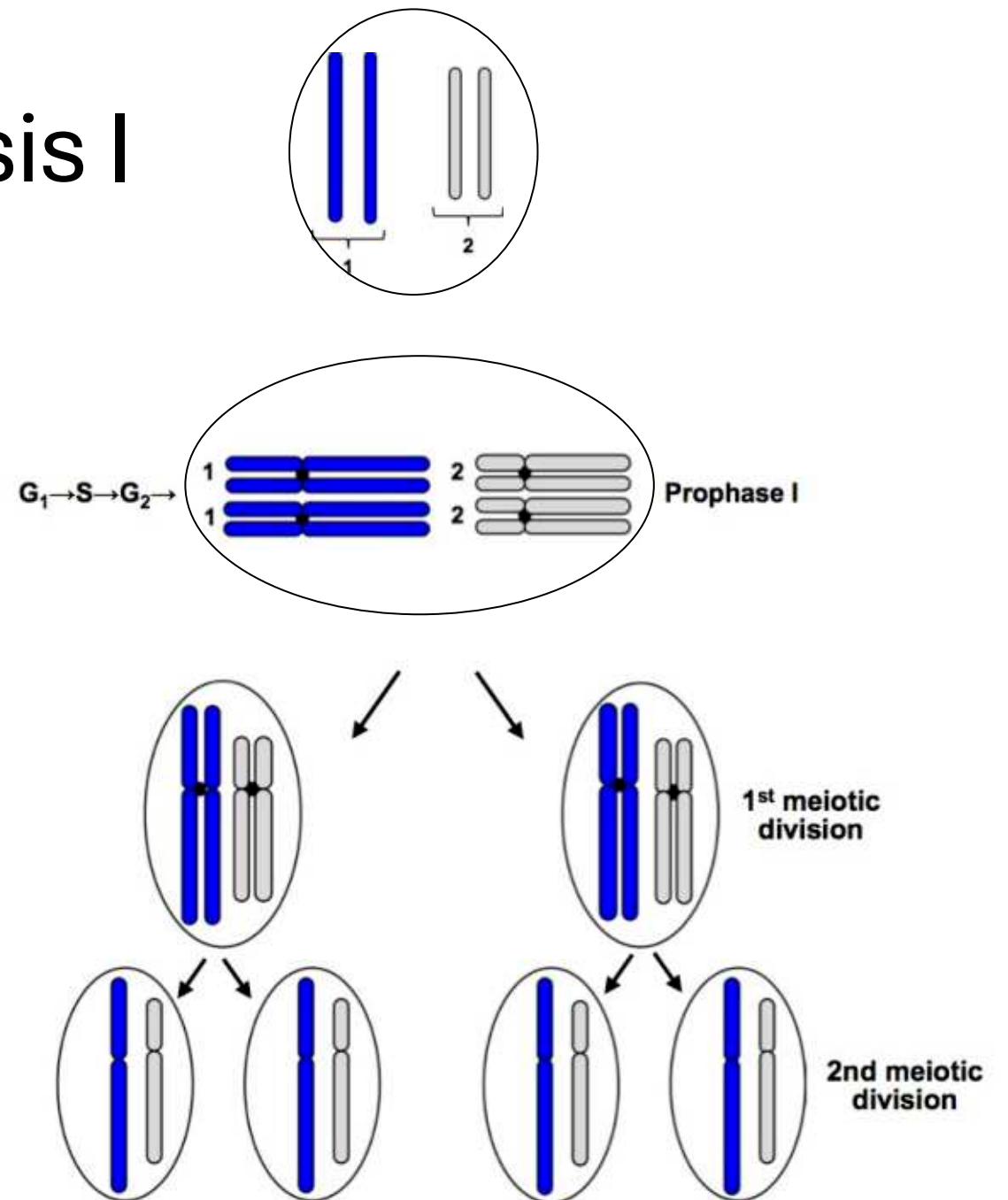
Translocations

- Involves the exchange of chromosome segments between two chromosomes
- Two main types: reciprocal and non-reciprocal
- Reciprocal:
 - Results from breakage or recombination involving nonhomologous chromosomes, with reciprocal exchange of the broken-off or recombined segments
 - Most often leads to a balanced rearrangement
 - Risk for unbalanced gametes and abnormal progeny due to malsegregation during gametogenesis



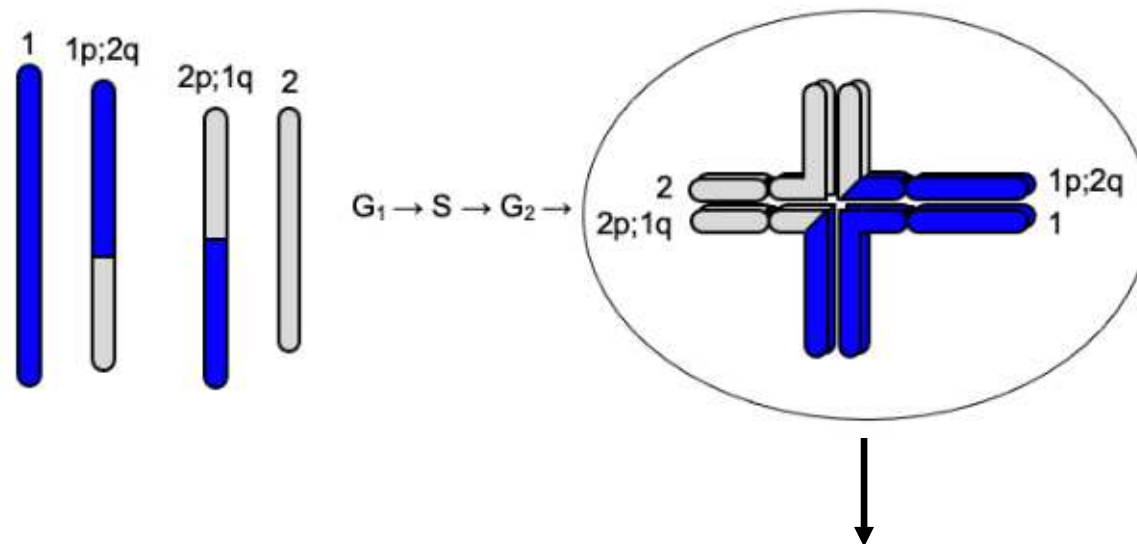
Review of normal meiosis I

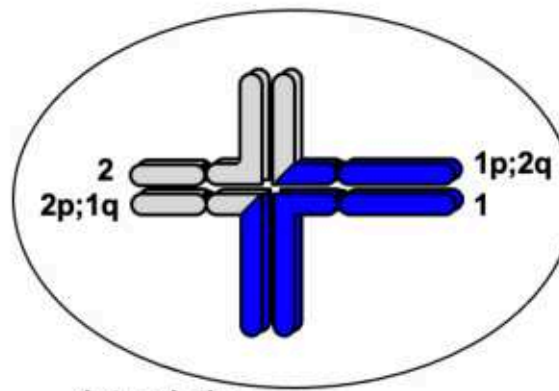
- Segregation of chromosomes
- **Tetrads (bivalent)** form between homologous pairs of chromosomes during prophase I
- Two pairs are shown in this schematic (*crossing over occurs, but not shown here*)



Meiosis I in balanced reciprocal translocations

- Malsegregation of chromosomes
- **Tetrads (quadrivalent)** form to ensure proper alignment of homologous sequences during prophase I

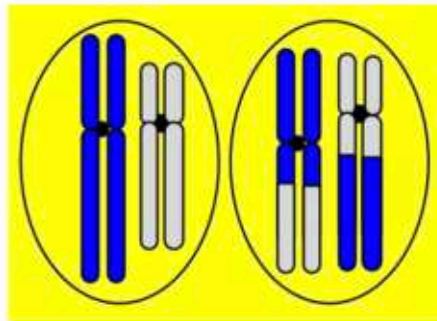




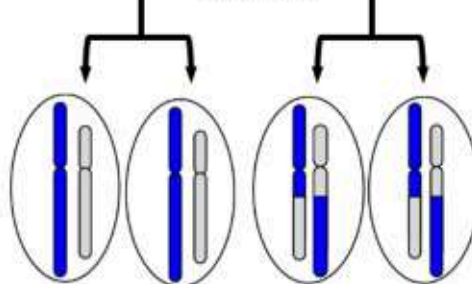
3 possible sets of gametes can be generated depending on how chromosomes segregate during meiosis I into secondary oocytes or secondary spermatocytes

1st meiotic division

Alternate Segregation
(balanced)



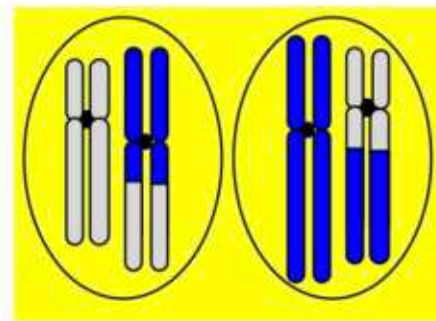
2nd meiotic division



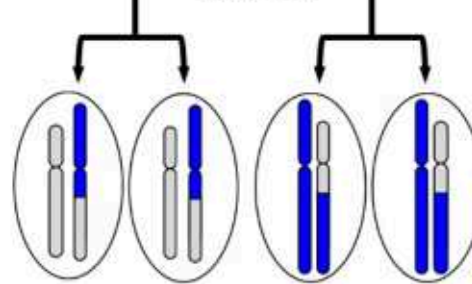
normal

balanced carrier

Adjacent I Segregation
(unbalanced)

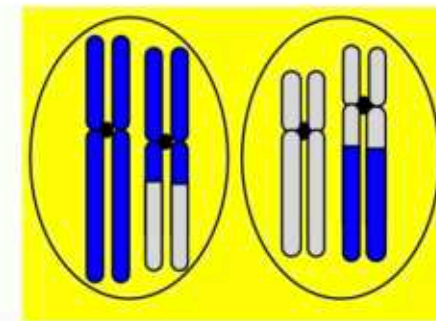


2nd meiotic division

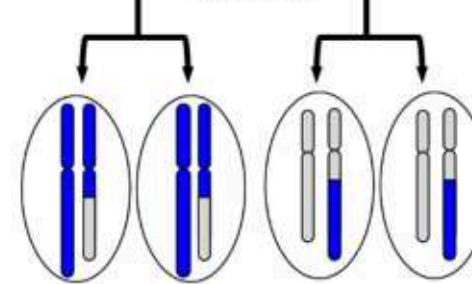


all phenotypically abnormal due to duplications and deletions

Adjacent II Segregation
(unbalanced)

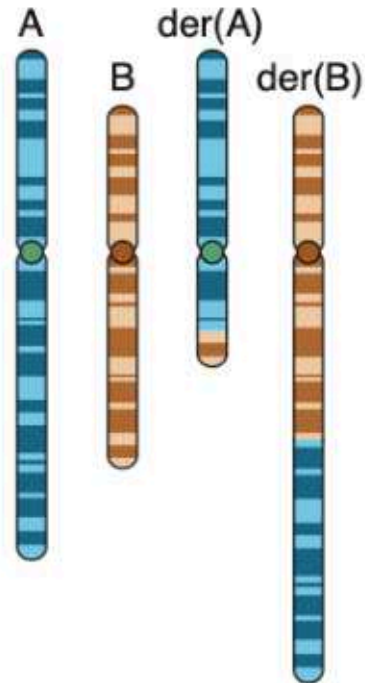


2nd meiotic division

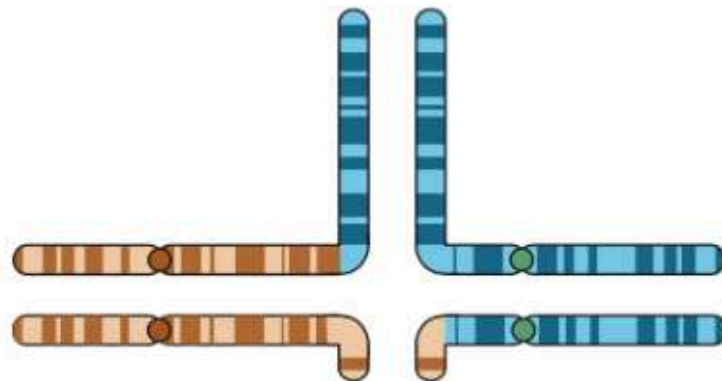


all phenotypically abnormal due to duplications and deletions

A Chromosomes



B Quadrivalent formation in meiosis



C Segregation and gametes

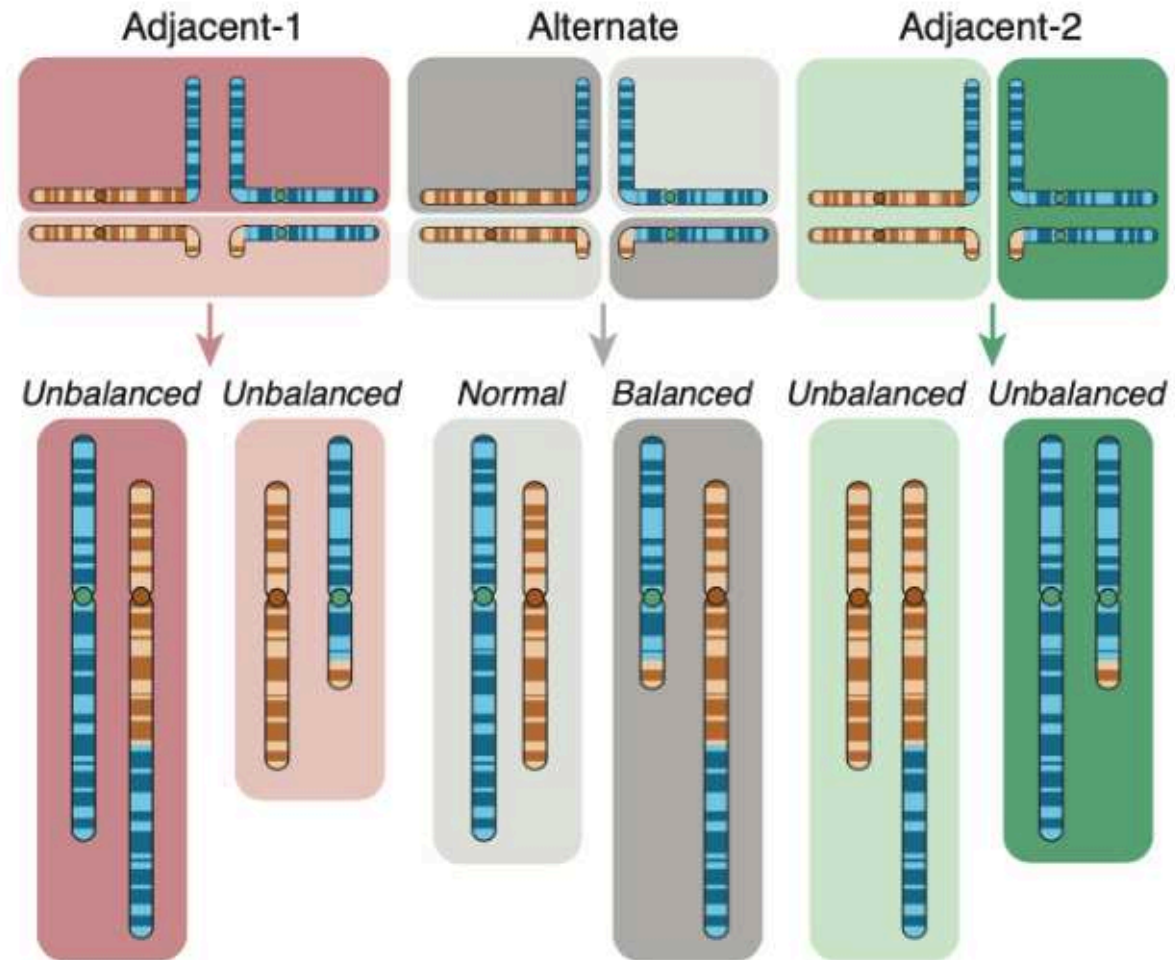
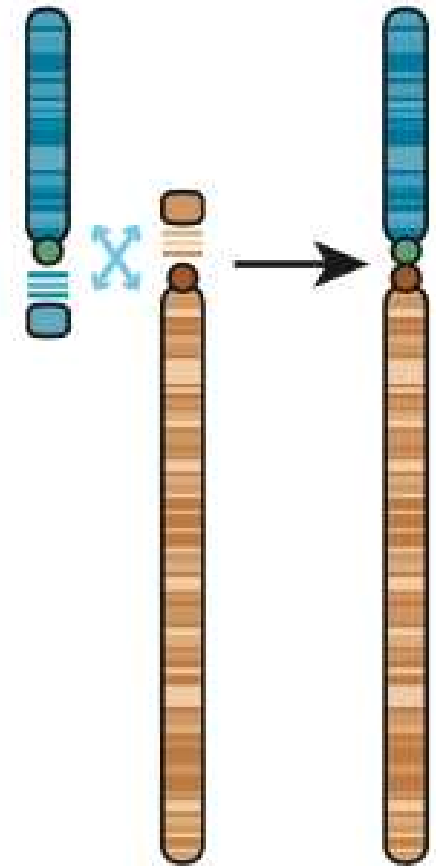


Figure 5-12 A, Diagram illustrating a balanced translocation between two chromosomes, involving a reciprocal exchange between the distal long arms of chromosomes A and B. B, Formation of a quadrivalent in meiosis is necessary to align the homologous segments of the two derivative chromosomes and their normal homologues. C, Patterns of segregation in a carrier of the translocation, leading to either balanced or unbalanced gametes, shown at the bottom. Adjacent-1 segregation (in red, top chromosomes to one gamete, bottom chromosomes to the other) leads only to unbalanced gametes. Adjacent-2 segregation (in green, left chromosomes to one gamete, right chromosomes to the other) also leads only to unbalanced gametes. Only alternate segregation (in gray, upper left/lower right chromosomes to one gamete, lower left/upper right to the other) can lead to balanced gametes.

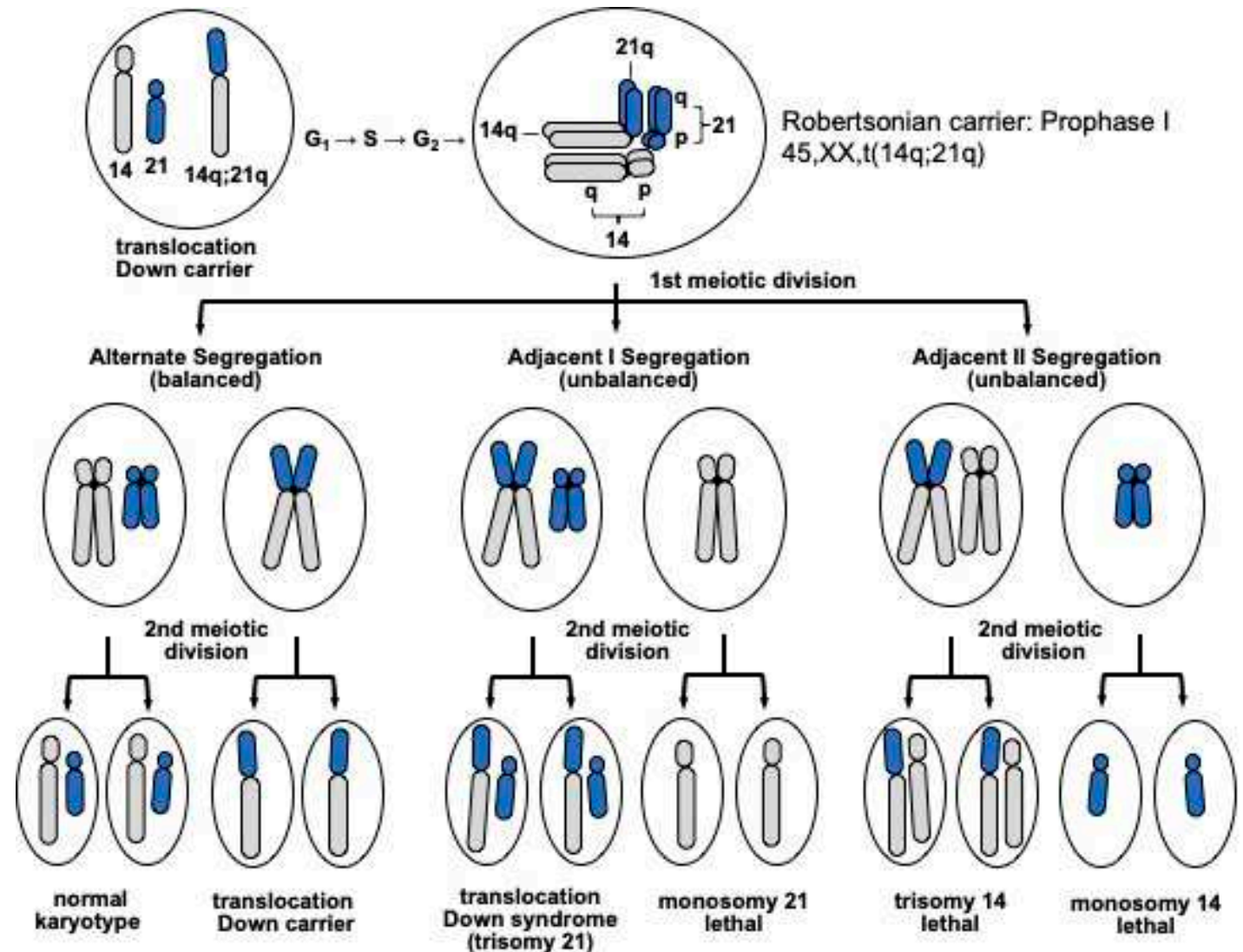
Robertsonian Translocations

- Involve two acrocentric chromosomes (13, 14, 15, 21, 22) that fuse near centromere region with loss of short arms (p)
- Results in karyotype with 45 chromosomes
- “Balanced” as loss of short arm material is not deleterious – normal phenotype
- Risk of abnormal offspring due to malsegregation of chromosomes during gametogenesis or risk for uniparental disomy
- Most common:
 - rob(13;14)(q10;q10) - 1 in 1300 people
 - rob(14;21)(q10;q10)



Robertsonian Translocations: Down Syndrome

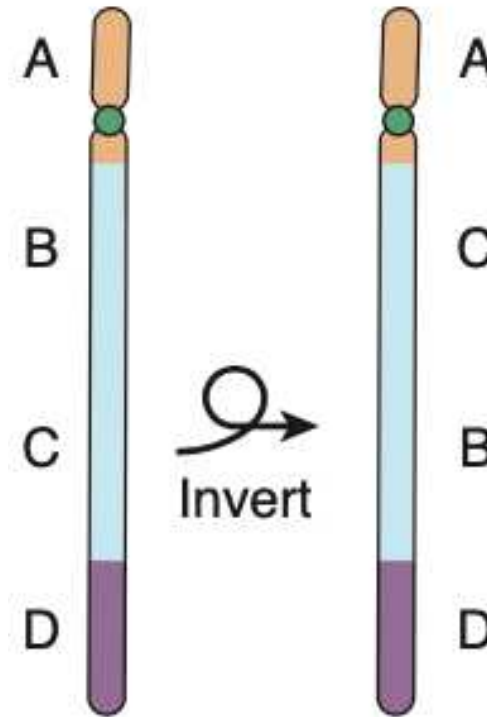
- 5% of Down syndrome patients have 46 chromosomes, one of which is a Robertsonian translocation between 21 and another acrocentric chromosome (most commonly 14 or 22)
- Carriers of Robertsonian translocations involving 21 are at risk for producing a child with translocation Down syndrome



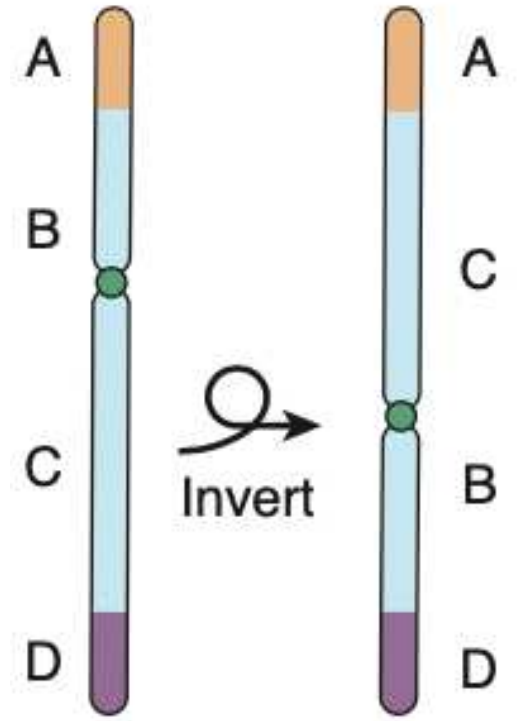
Inversions

- When a single chromosome undergoes two breaks and is reconstituted with the segment between the breaks inverted
- Types:
 - Paracentric – both breaks occur in one arm (**para = beside the centromere**)
 - Pericentric – breaks occur in each arm (**peri = around the centromere**)
- Usually normal phenotype, but risk for abnormal gametes due to loop formation for pairing of homologous segments in meiosis I

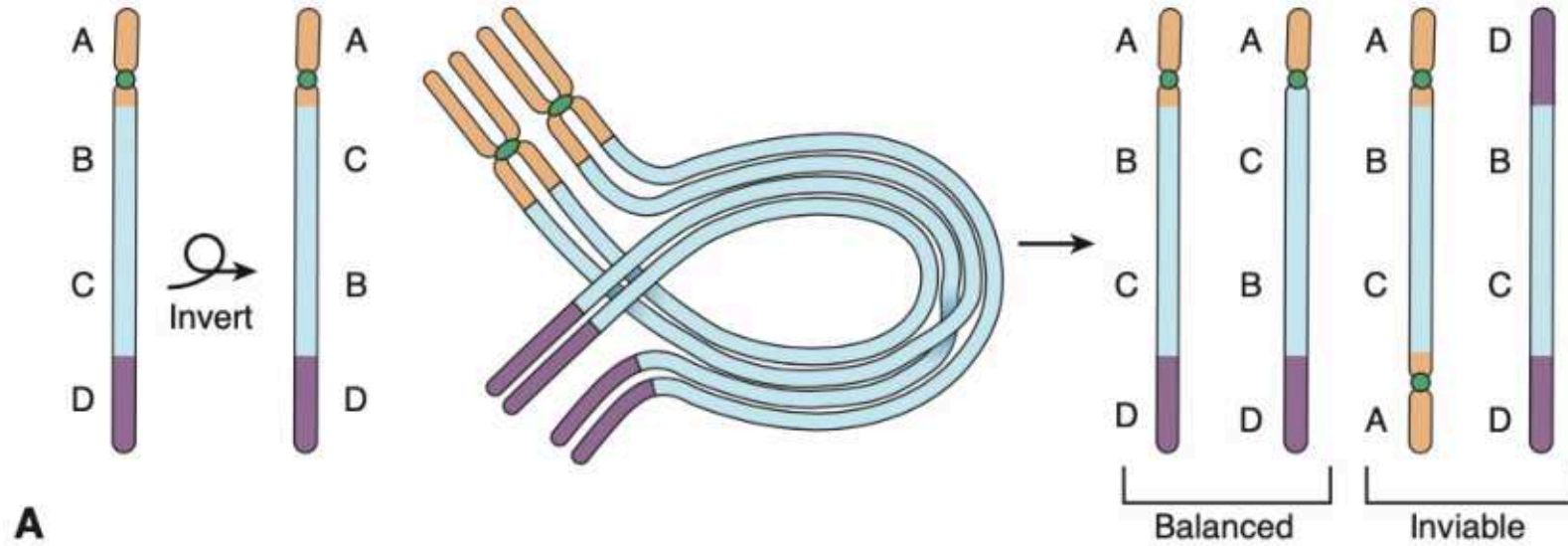
Paracentric



Pericentric

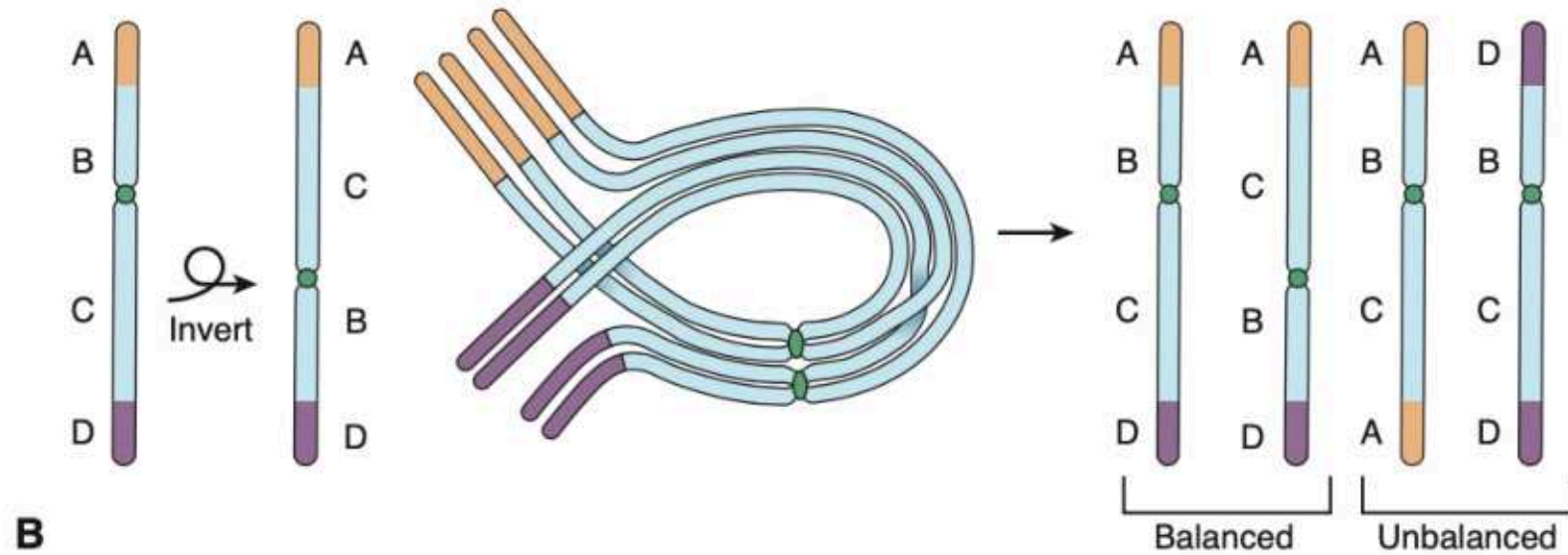


Paracentric



Paracentric:
unbalanced
recombinant
chromosomes are
acentric or dicentric
and typically do not
lead to viable
offspring

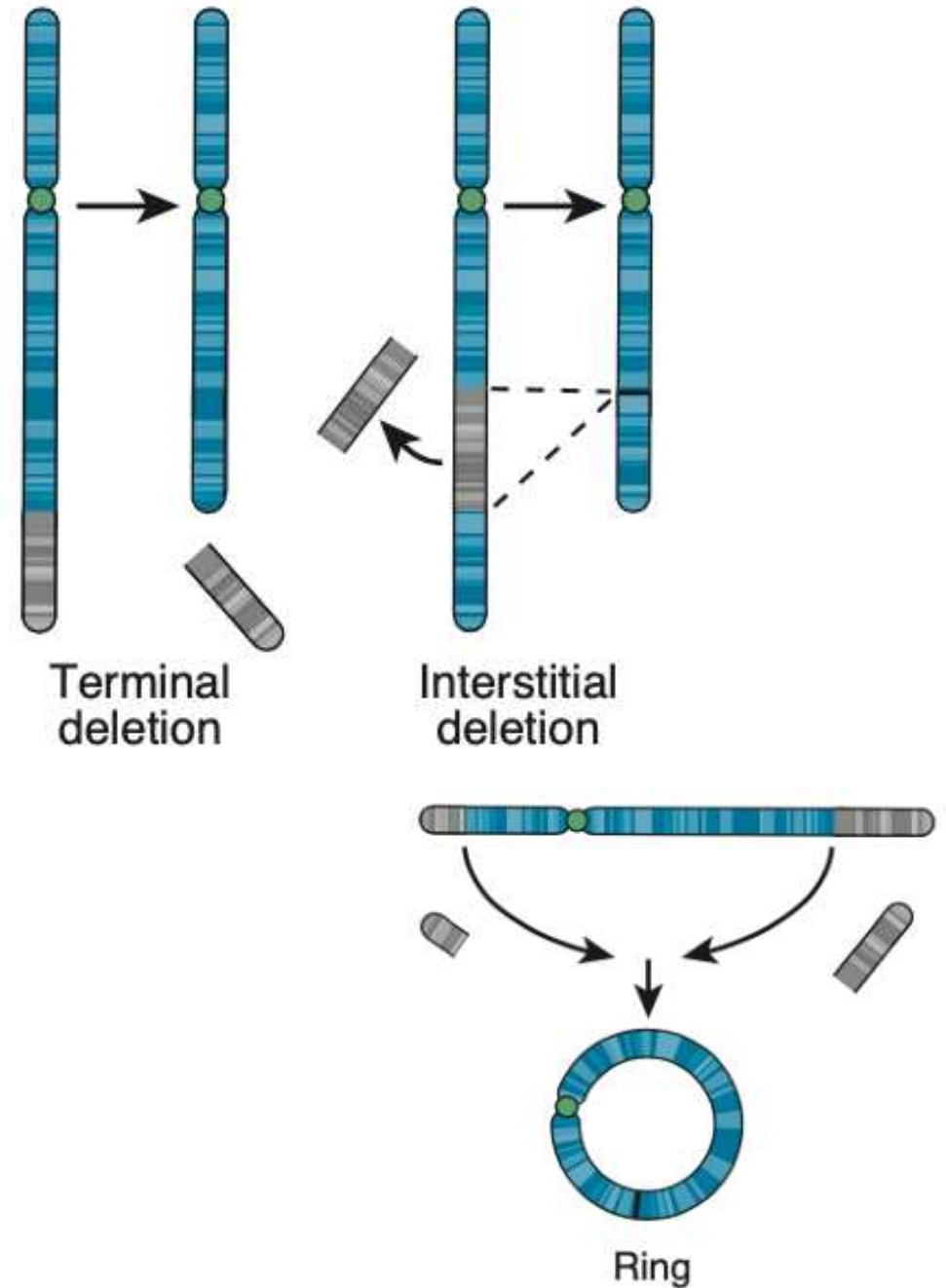
Pericentric



Pericentric:
unbalanced gametes
with both
duplication and
deficiency of
chromosome
segments distal to
the inversion

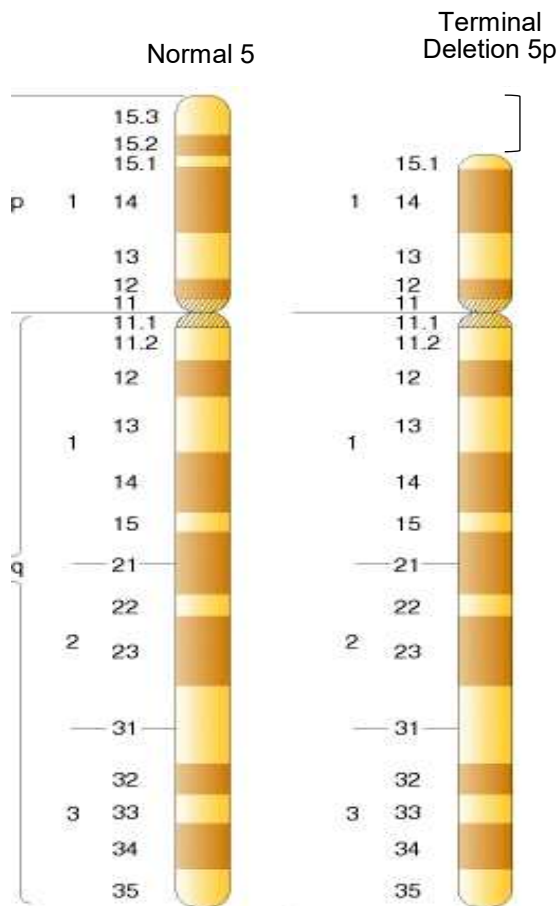
Deletions

- Involve loss of a chromosome segment, resulting in chromosome imbalance
- Monosomic or haploinsufficient
- Severity related to size and number of genes in deleted segment
- Larger, cytogenetically visible deletions detected on karyotype
- Submicroscopic deletions (microdeletions) detected on microarray



Cri du chat Syndrome

Terminal Deletion of 5p – detect on karyotype or microarray



Common Clinical Features

High-pitched cat-like cry

Microcephaly

Hypertelorism

Intellectual disability /
developmental delay

Hypotonia

Round “moon” face



Wolf-Hirschhorn Syndrome

Terminal Deletion of 4p – detect on karyotype or microarray



Common Clinical Features

“Greek warrior helmet” (microcephaly, prominent glabella, broad nasal tip, short philtrum, hypertelorism)

Cardiac defects

Hypotonia

Seizures

Growth restriction / failure to thrive

Intellectual disability

Cleft lip and palate

DiGeorge (22q11.2 deletion) Syndrome

Interstitial microdeletion – detect on microarray



Common Clinical Features

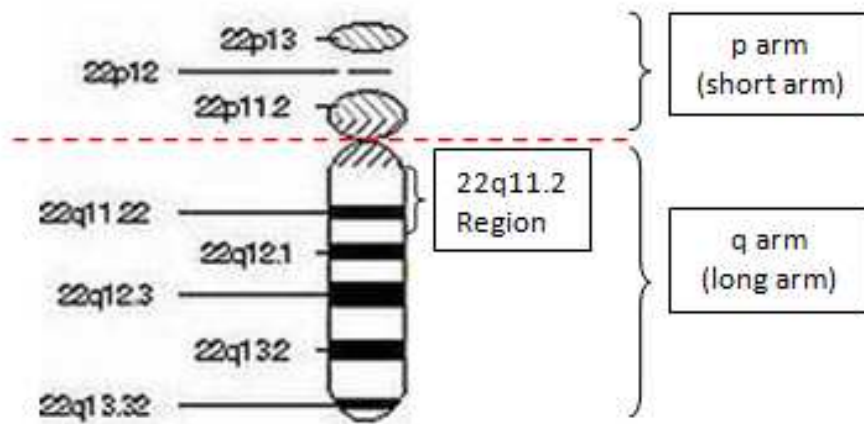
Cardiac defects (conotruncal)

Abnormal facies (hypertelorism, bulbous tip of nose, low set ears)

Thymic hypoplasia (immunodeficiency)

Cleft palate

Hypoparathyroidism + Hypocalcemia



Williams Syndrome

Interstitial microdeletion of 7q11.23 – detect on microarray

Common Features / Buzzwords

Outgoing (cocktail party personality)

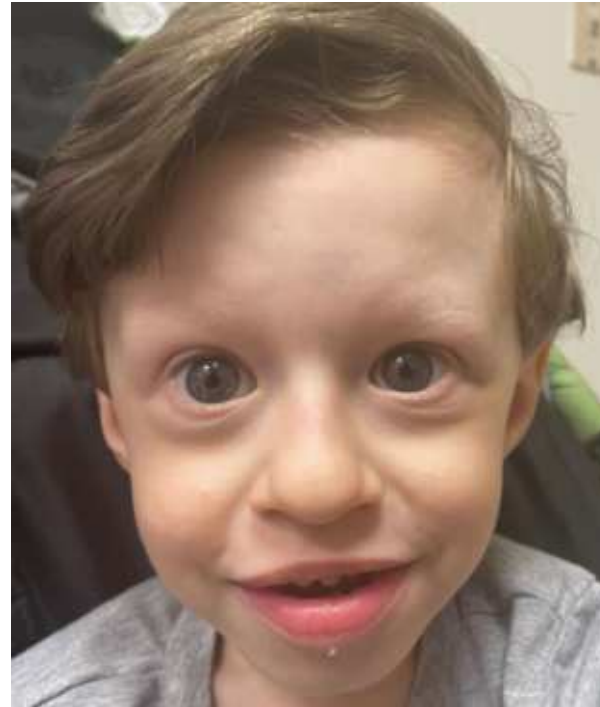
Elfin-like facies

Cardiac defects (supravalvular
aortic stenosis)

Infantile hypercalcemia

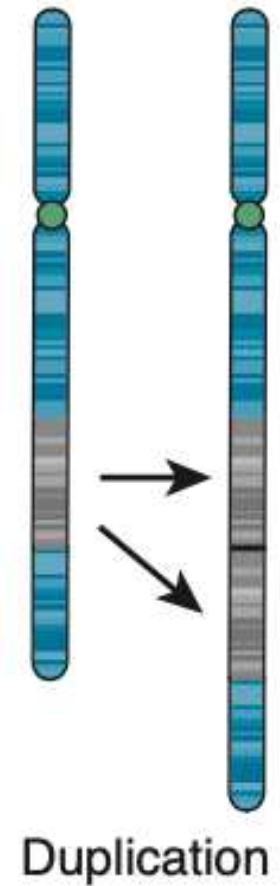
Mild to moderate intellectual disability

Stellate iris pattern



Duplications

- Duplications often occur at loci in the chromosome that contain tandem or inverted repeats.
- Duplications are generated in most cases by:
 - DNA replication errors
 - Unequal crossover during meiosis
- In general, duplications are less detrimental than deletions
- Many small duplications do not have any clinical consequences



Questions?

- Dr. Thompson – thlaur@musc.edu
- Dr. Lancaster - lancastk@musc.edu

Come shadow us in Genetics clinic!