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# Cytogenetics Chromosomal Genetics

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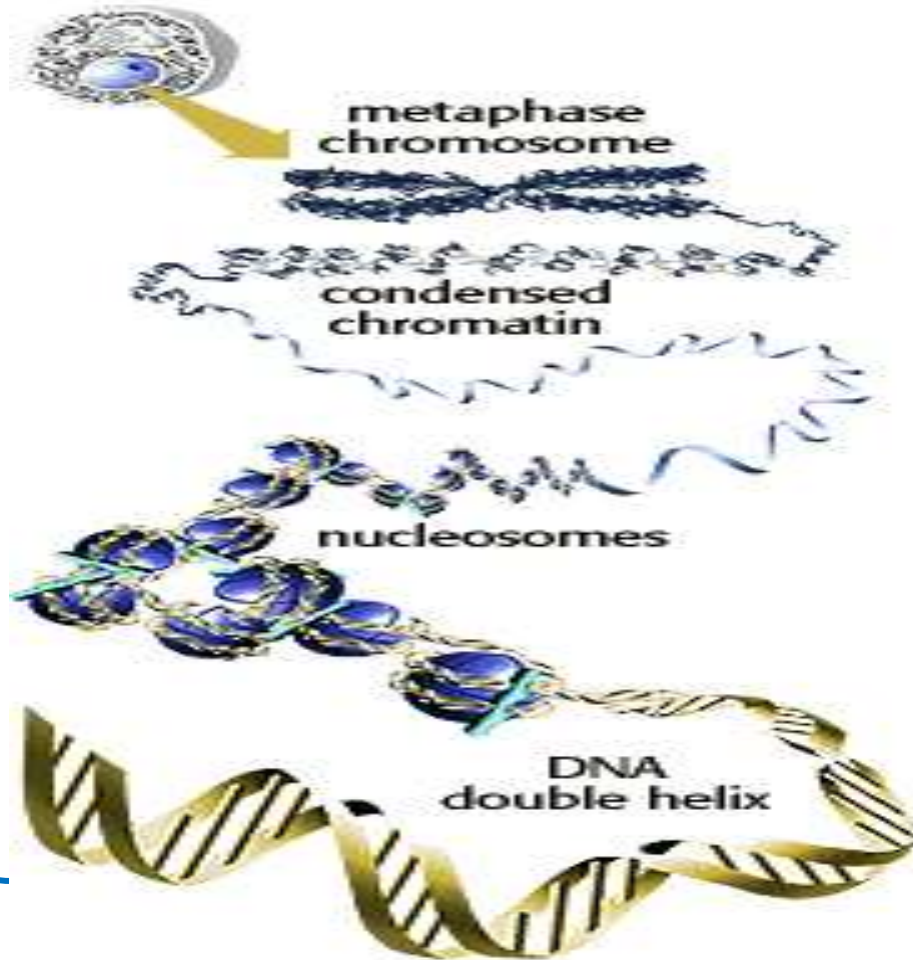
Geneva, Switzerland

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Training Course in Sexual and Reproductive Health Research  
Geneva 2011

Cytogenetics is the branch of genetics that correlates the structure, number, and behaviour of chromosomes with heredity and diseases

**DNA packs tightly into metaphase chromosomes**



Conventional cytogenetics

Molecular cytogenetics

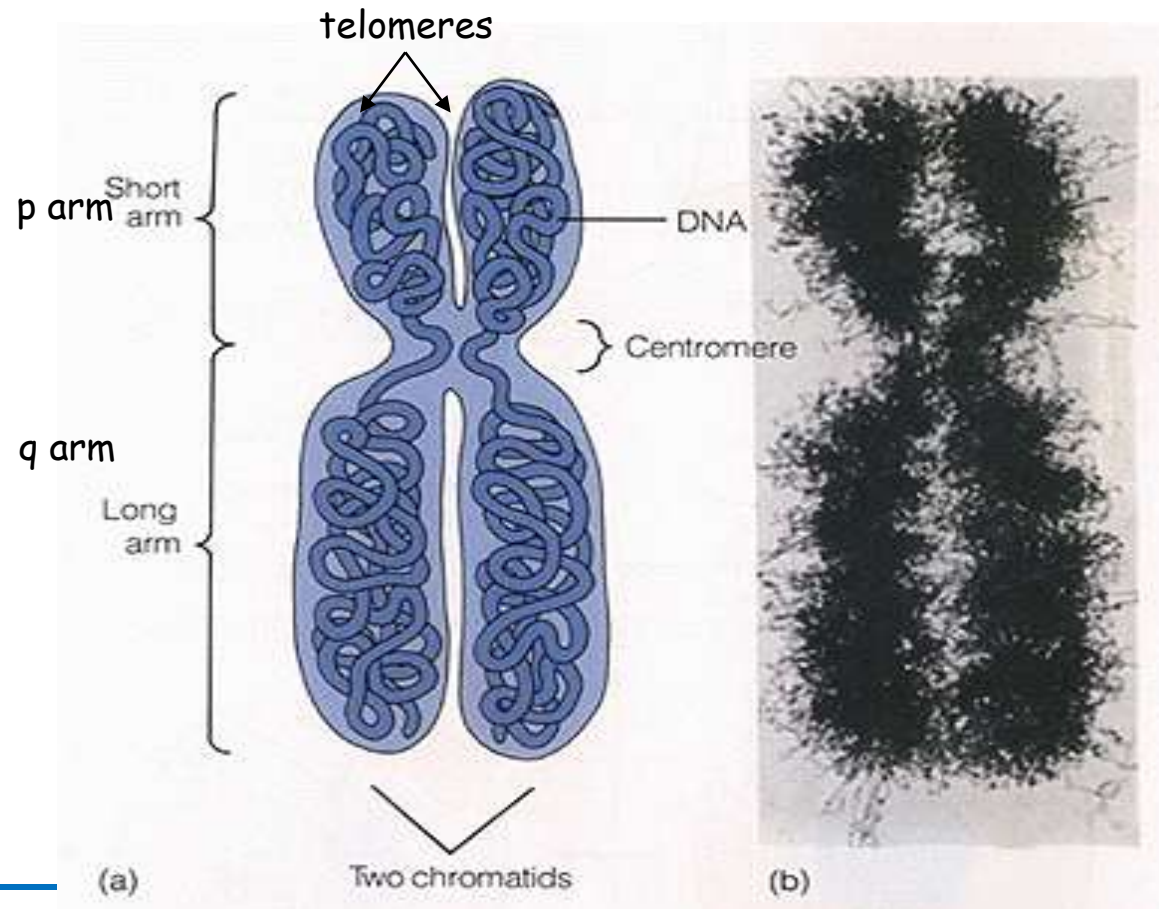
Molecular Biology

# I. Karyotype

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- Definition
- Chromosomal Banding
- Resolution limits
- Nomenclature

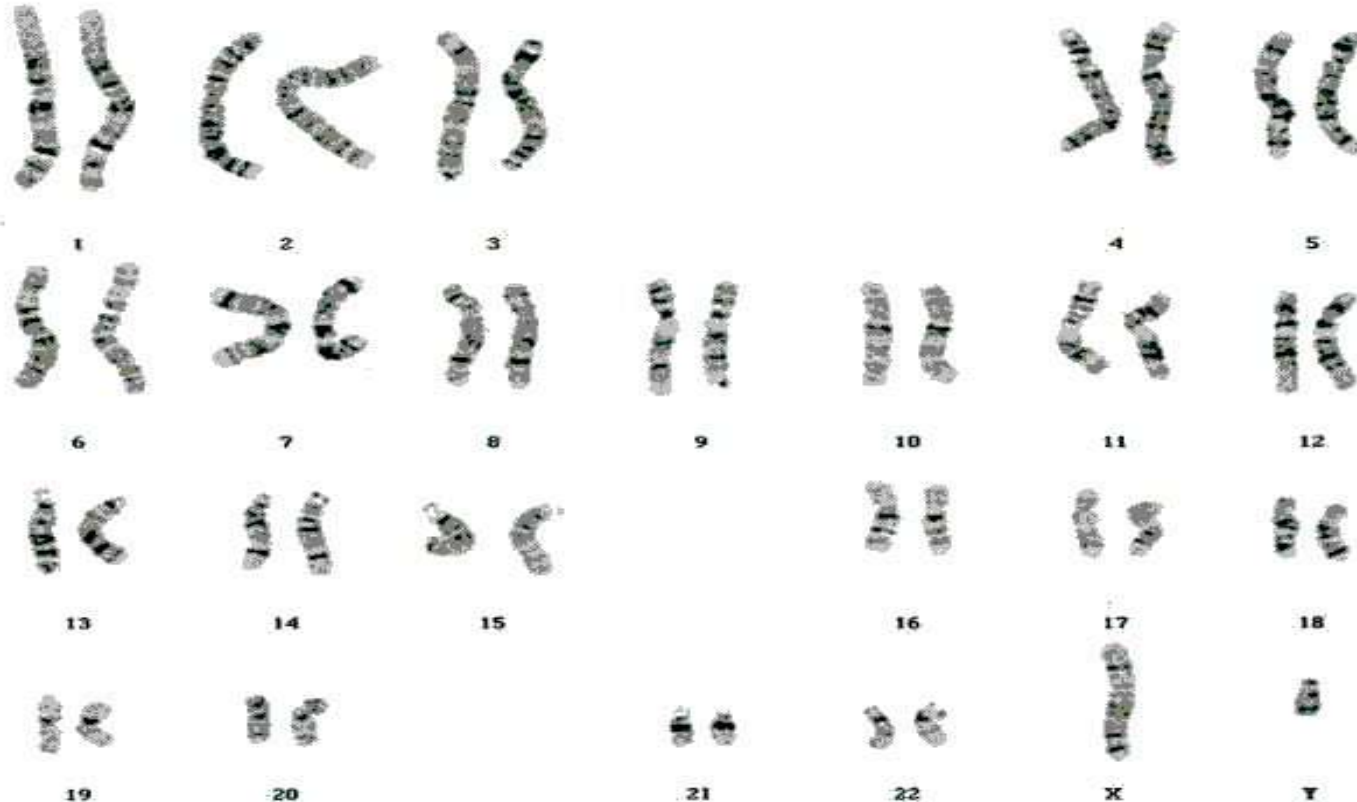
# The metaphasic chromosome



# G-banded Human Karyotype

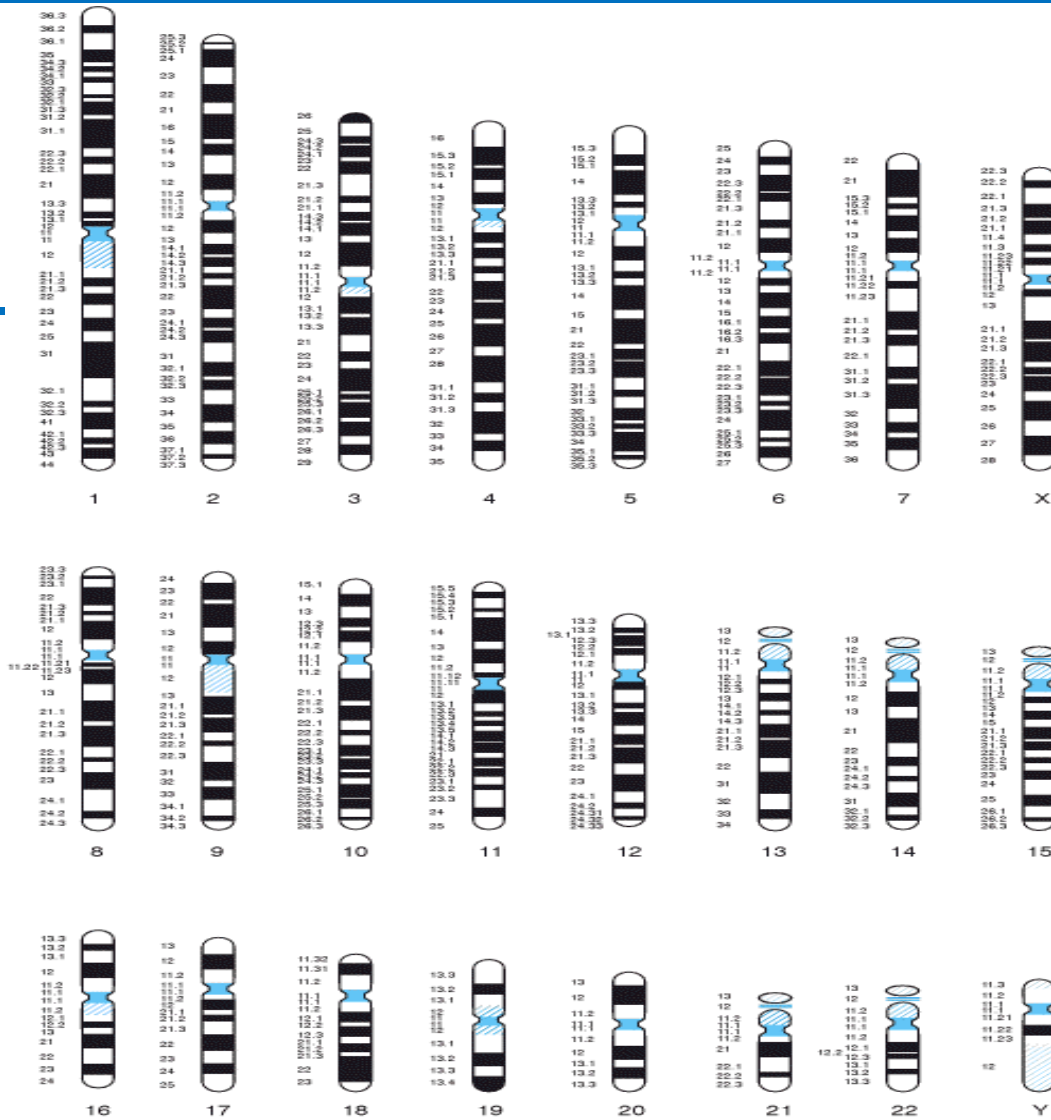
Tjio & Levan 1956

HOPITAL CANTONAL UNIVERSITAIRE GENEVE DIVISION DE GENETIQUE MEDICALE



**Karyotype:** The characterization of the chromosomal complement of an individual's cell, including number, form, and size of the chromosomes.

A photomicrograph of chromosomes arranged according to a standard classification.



Key:  
 ■ Centromere  
 ■ rDNA  
 ■ Noncentromeric heterochromatin

A chromosome banding pattern is comprised of alternating light and dark stripes, or bands, that appear along its length after being stained with a dye. A unique banding pattern is used to identify each chromosome

# Chromosome banding techniques and staining

- Giemsa has become the most commonly used stain in cytogenetic analysis. Most G-banding techniques require pretreating the chromosomes with a proteolytic enzyme such as trypsin. G-banding preferentially stains the regions of DNA that are rich in adenine and thymine.
- R-banding involves pretreating cells with a hot salt solution that denatures DNA that is rich in adenine and thymine. The chromosomes are then stained with Giemsa.
- C-banding stains areas of heterochromatin, which are tightly packed and contain repetitive DNA.
- NOR-staining, where NOR is an abbreviation for "nucleolar organizing region," refers to a silver staining method that identifies genes for ribosomal RNA.

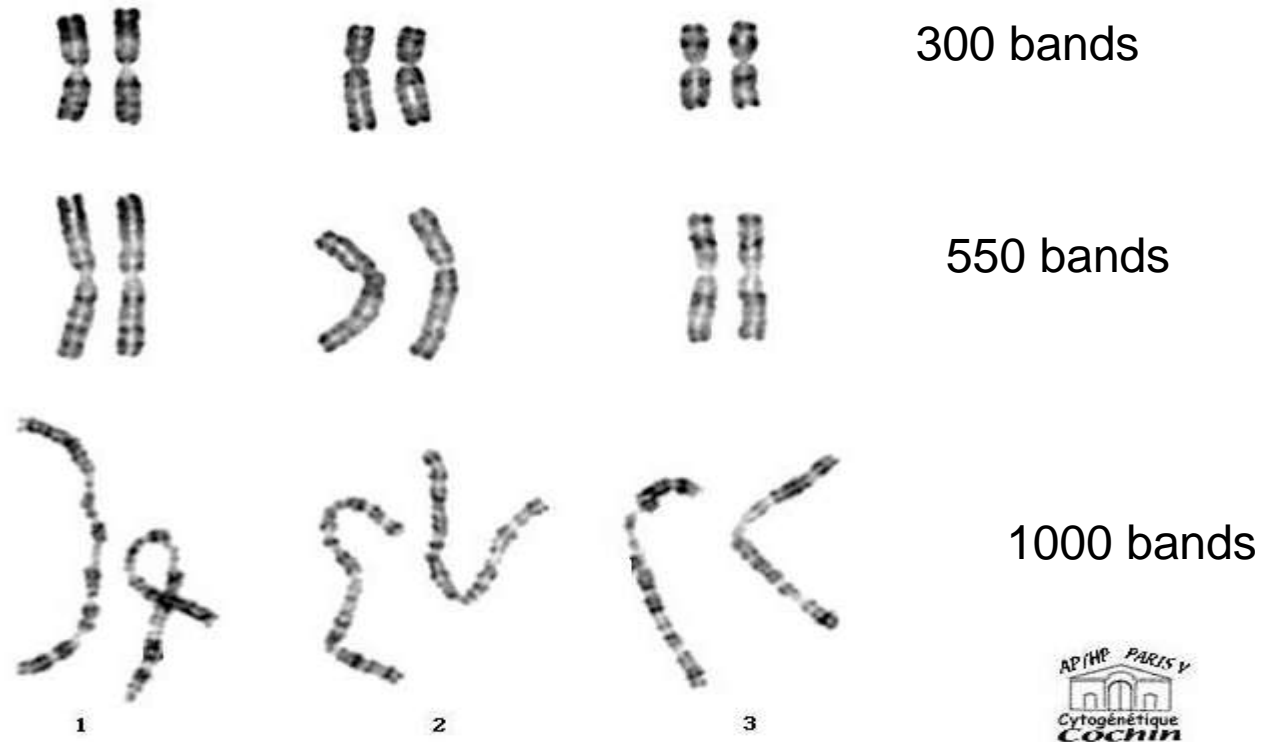






# Limits of resolution

## Metaphase Chromosomes at different levels of resolution



Depending on the length of the chromosomes, the karyotype has a limit of resolution, indicated par the count of bands for a haploid genome

# Nomenclature

International System for human Cytogenetic Nomenclature (ISCN) 2009

## **In designating a particular band,**

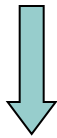
- ⇒ chromosome number
- ⇒ Arm symbol
- ⇒ Region number
- ⇒ Band number

## **Description of chromosome abnormalities**

- ⇒ Total number of chromosomes including sex chromosomes
- ⇒ Sex chromosome constitution
- ⇒ Numerical abnormalities
- ⇒ For example a female Down syndrome or trisomy 21 is written as 47,XX,+21
- ⇒ Structural changes are designated by letters, for example 'dup' for duplication  
Such as 46,XY,dup(1)(q22q25 ) (duplication of a segment in long arm of chromosome 1, q, in region 2 between bands 22 and 25.

# Chromosomes can be studied in any nucleated body cell in an individual

Peripheral blood



Lymphocyte culture  
3 days

Blood sample is taken



Chromosome



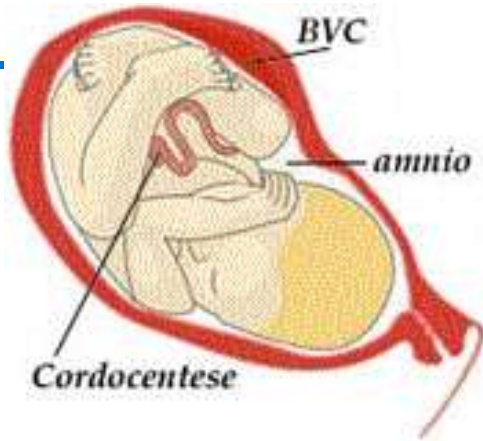
Skin biopsy



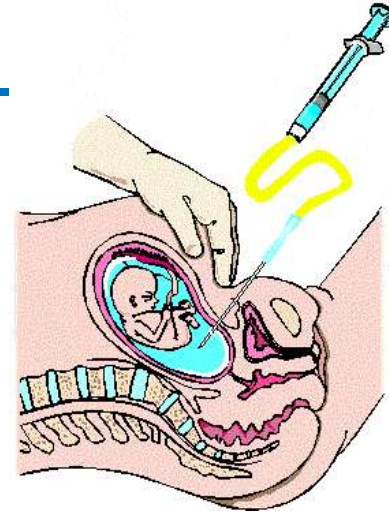
culture of fibroblasts  
15 -21 days



# Prenatal tests to study fetal chromosomes



**Choriocentesis**  
**(Chorion villus biopsy)**  
Risk of abortion 2-3%

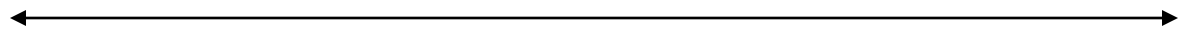


**Amniocentesis**  
Risk of abortion 1%

**Choriocentesis**



**Amniocentesis**



**GW 10 11 12 13 14 15 16 17 18 19 20 21 22**

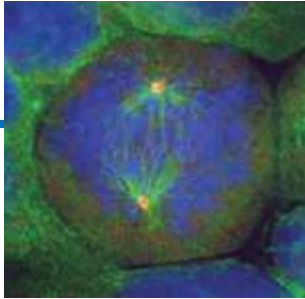


**Cordocentesis**

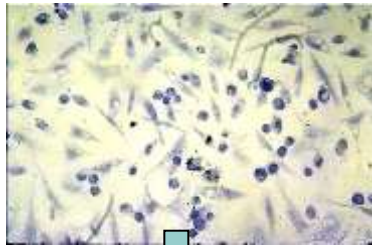
(GW: gestational weeks)

(Blood from umbilical artery)

# Chromosome preparation



Addition of colchicine inhibits formation of mitotic spindle



Hypotonic solution to disperse chromosomes

Fixation of chromosomes on a slide

Staining of chromosomes



## II. Chromosome abnormalities

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- Statistics
- Meiosis
  - Description
  - Crossing over, recombination
- Errors of meiosis I
- Errors of meiosis II
- Promoted factors

# Chromosome abnormalities

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1. Constitutional : exist at birth. These are usually present in all tissues, if present only in some tissues, it is called mosaicism and it means that the abnormality occurred in the mitotic divisions that follow zygote formation
2. Acquired: occur during the life of a healthy individual and are confined to one tissue as seen in tumour cells



**Constitutional  
Chromosome  
abnormalities**

**Acquired  
chromosome  
abnormalities**

**Exist at birth**

**Present  
in all tissues**

**Mosaicism in some tissues  
caused by Postzygotic mitotic  
abnormality**

**occur during life  
in a healthy individual**

**Confined in a tissue  
Tumors**

# Frequencies of chromosome abnormalities

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- 2% of sperms have Chromosomal abnormalities
- 20% of ova have Chromosomal abnormalities
- So among 100 conceptions, there are 25% chromosome abnormalities

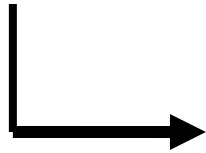
# Frequencies of chromosome abnormalities

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- In every 100 pregnancies, there occurs 15 spontaneous miscarriages, 50% of which have chromosome abnormalities
- Among 160 births, one baby is born with a chromosome abnormality

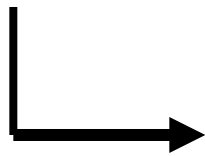
2% of sperms have Chromosomal abnormalities  
20% of ova have Chromosomal abnormalities

100 conceptions



25 Chromosomal abnormalities

100 Pregnancies

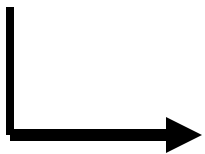


15 miscarriages



50% Chromosomal abnormalities

160 Births



1 child  
With a Chromosomal abnormality

# Meiosis

- Is the process of reductional division in which a diploid cell  $2N = 46$  (2 x sets of chromosomes) is reduced to a haploid cell  $(N) = 23$  (1 set of chromosomes)
- It comprises MI (meiosis I) and MII (meiosis II)
- Meiosis always results in the formation of gametes (ova and sperms)

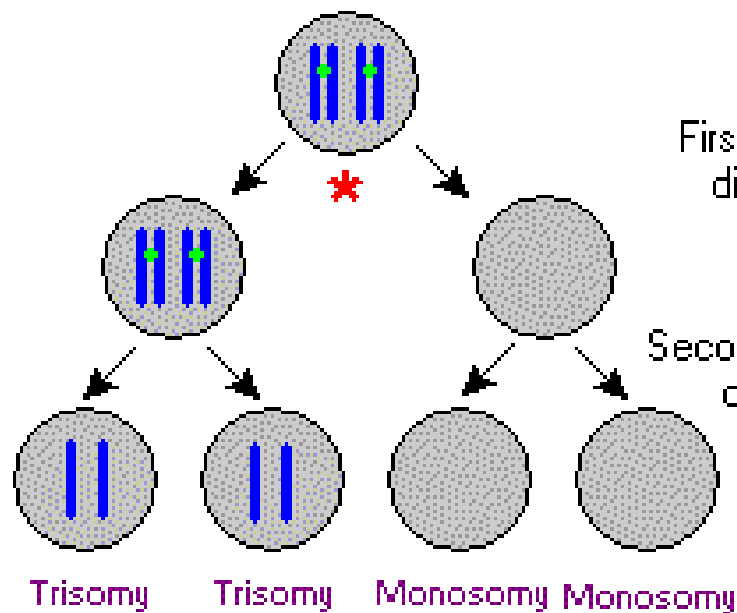
# Non-disjunction in meiosis

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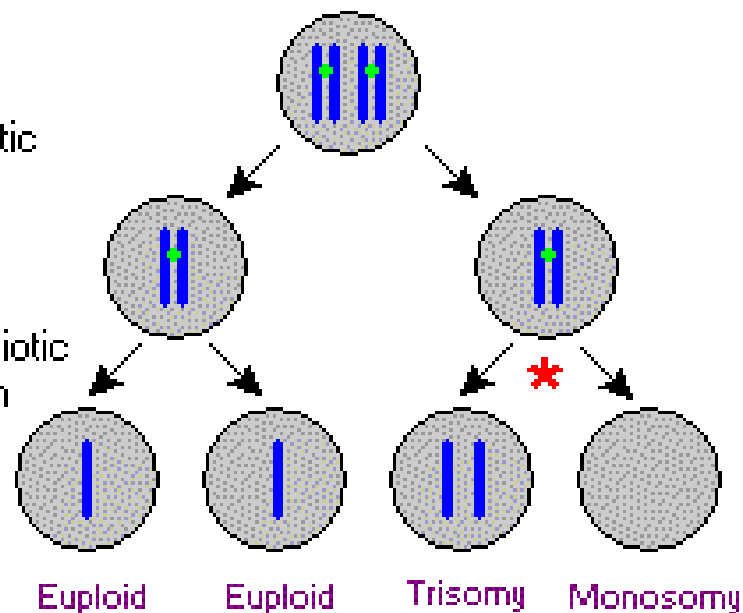
- This is an abnormal division where one daughter cell gets an extra chromosome (24) and the other daughter cell gets one chromosome less than normal (22).
- It can happen in MI or MII.
- Fertilisation with a normal gamete gives either a trisomic zygote ( $24+23=47$ ) or a monosomic zygote ( $22+23=45$ )

# Mechanism. Meiotic nondisjunction

## Nondisjunction in meiosis I

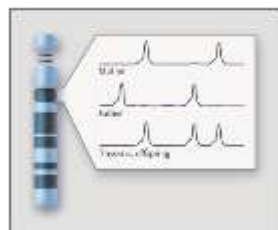


## Nondisjunction in meiosis II



Electrophoresis profiles

offspring after fertilization with another normal gamete







Maternal non disjunction

Known risk factors

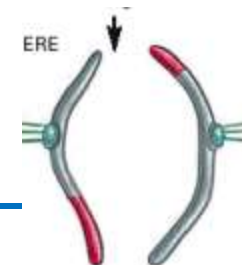
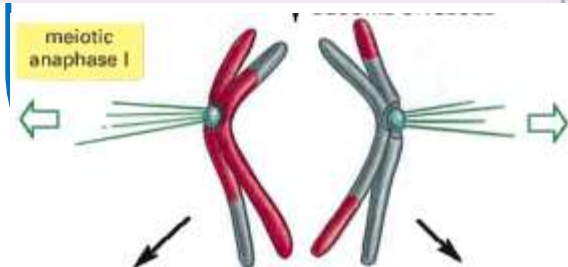
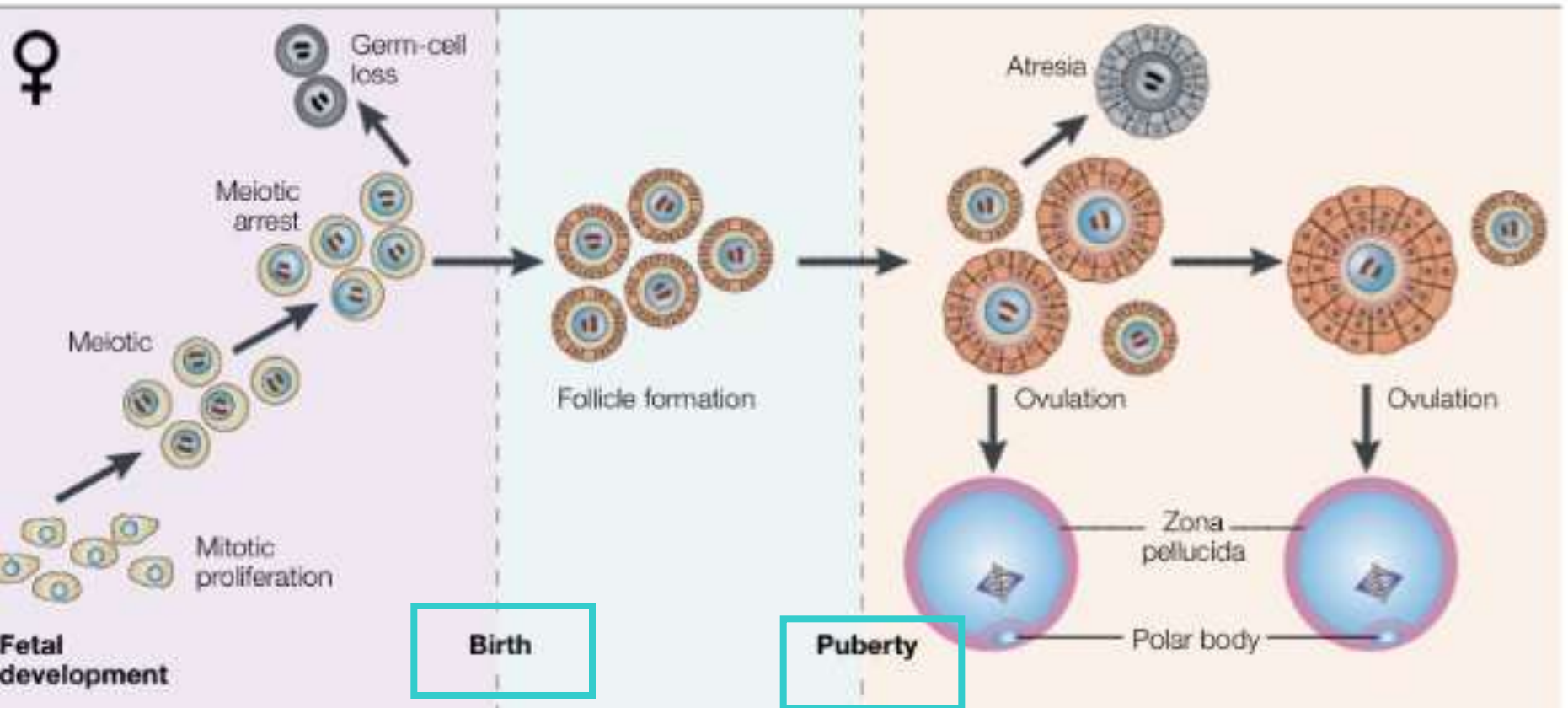
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## Period of gametogenesis in the female

meiosis starts at intrauterine life with ovulation starting at puberty. Each month one ovum is produced and 1000 follicles become undergo atresia

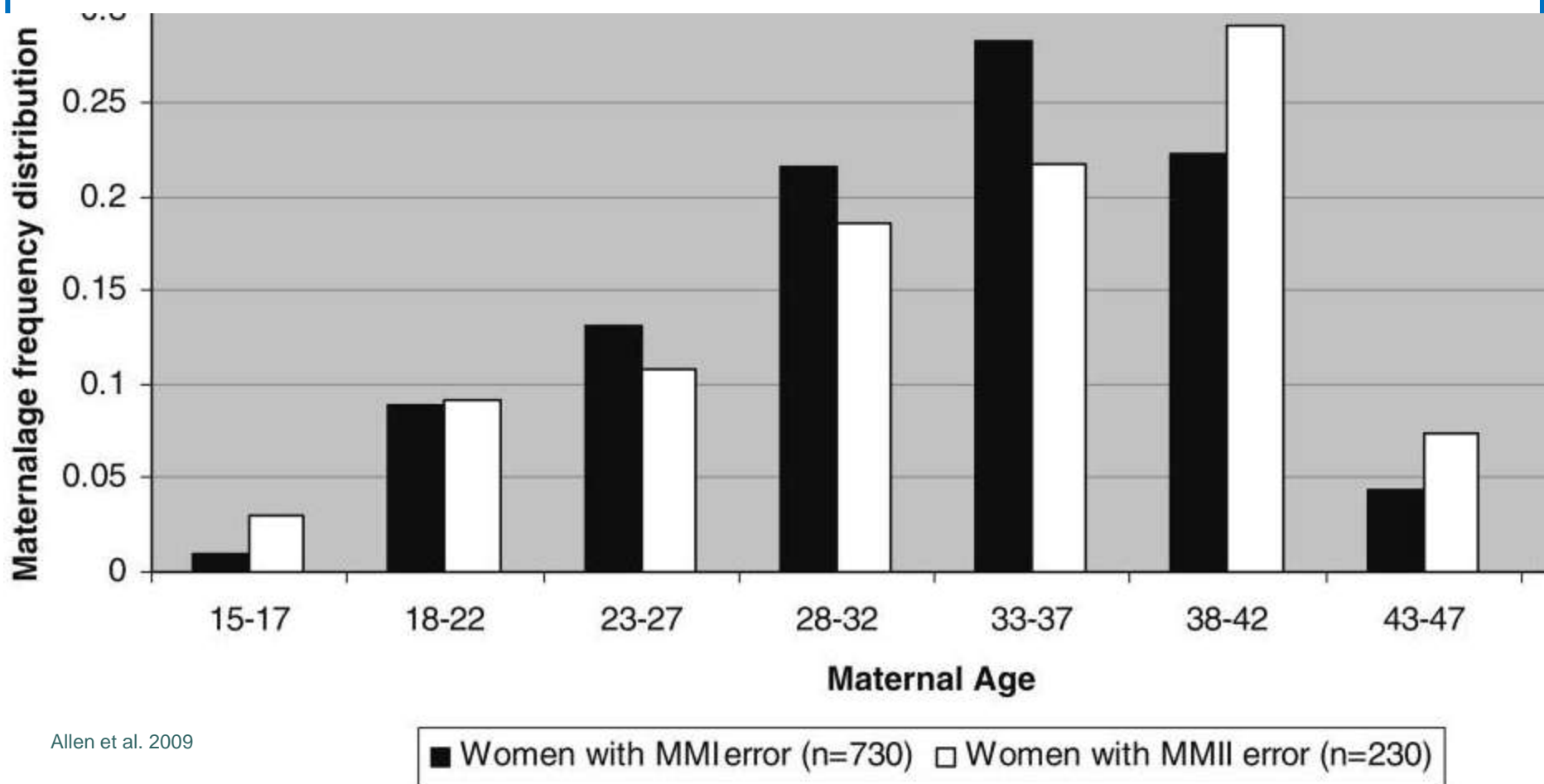


# Known predisposing causes for non-disjunction in the female

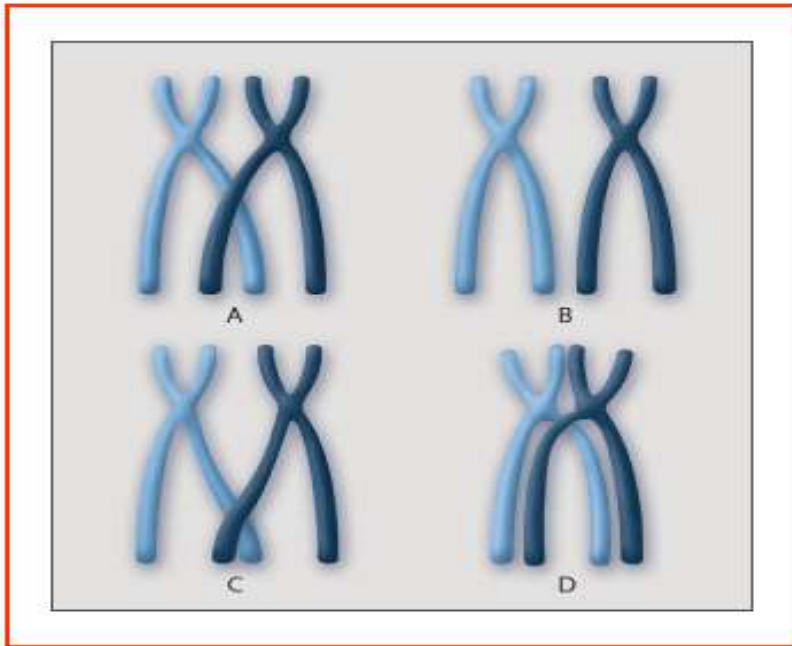
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- Advanced maternal age
- Sites and rate of meiotic recombination (crossing over or chiasma formation)
- Genetic factors
- Mosaicism with trisomic cells in ovaries

# Advanced Maternal Age



# Recombination and non disjunction

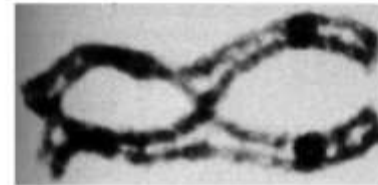


- Normal
  - 1 chiasma/chromosome A
- Trisomy 21 MMI,
  - 45% achiasma B
  - 41% 1 telomeric chiasma C
- Trisomy 21 MMII
  - Pericentromeric Chiasma D



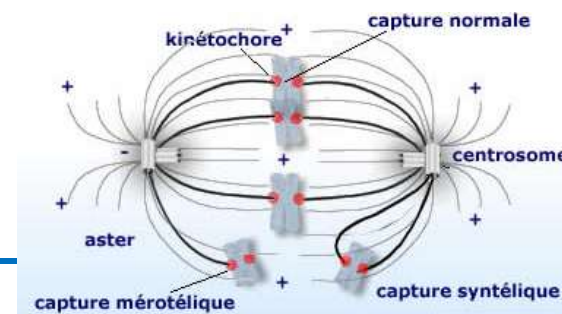
# Two-hit model of non disjunction

- Establishment of "susceptible " exchange in the fetal oocyte



**NinIntena**

- Age dependant abnormal processing



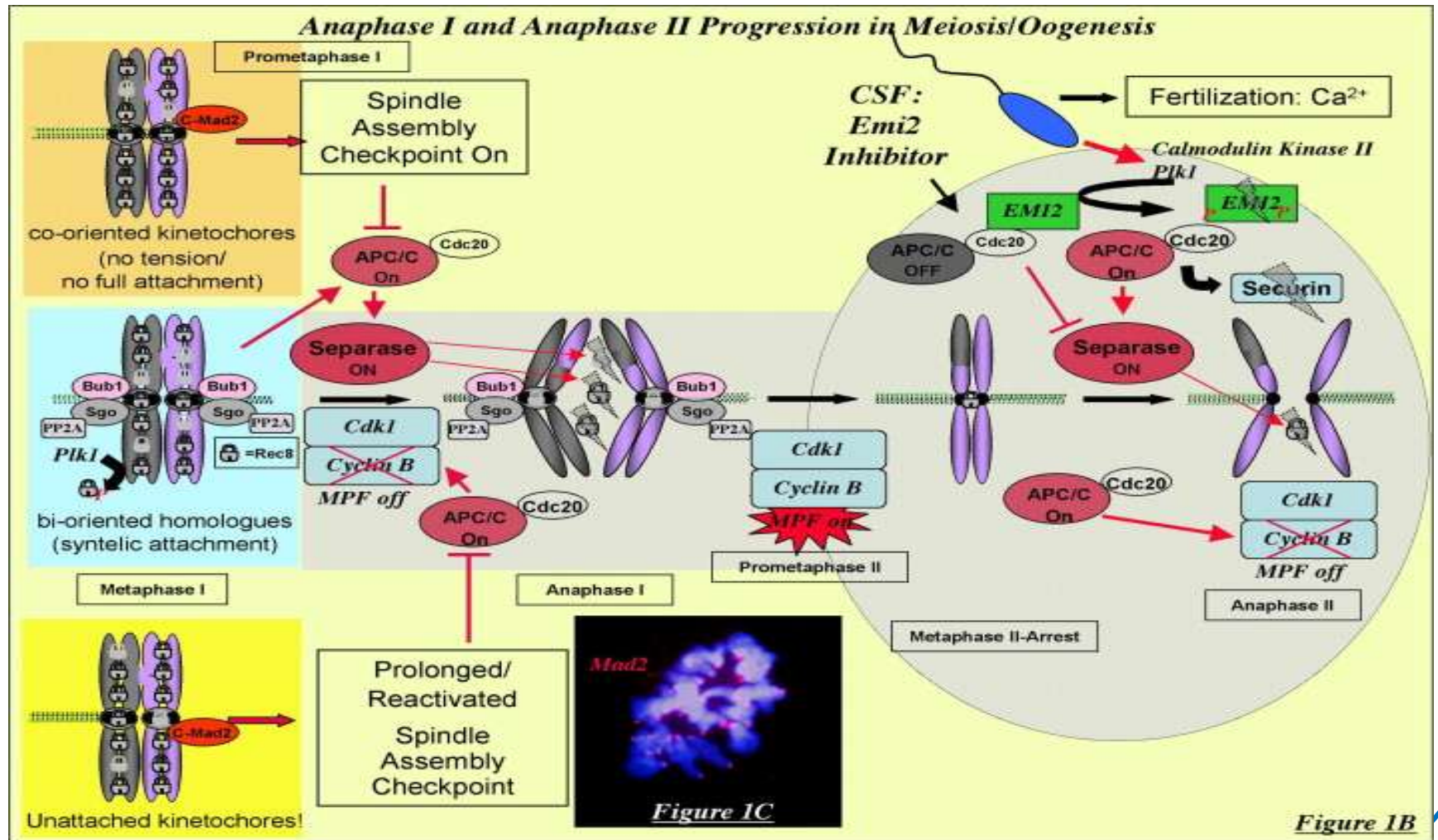
# Genetic factors

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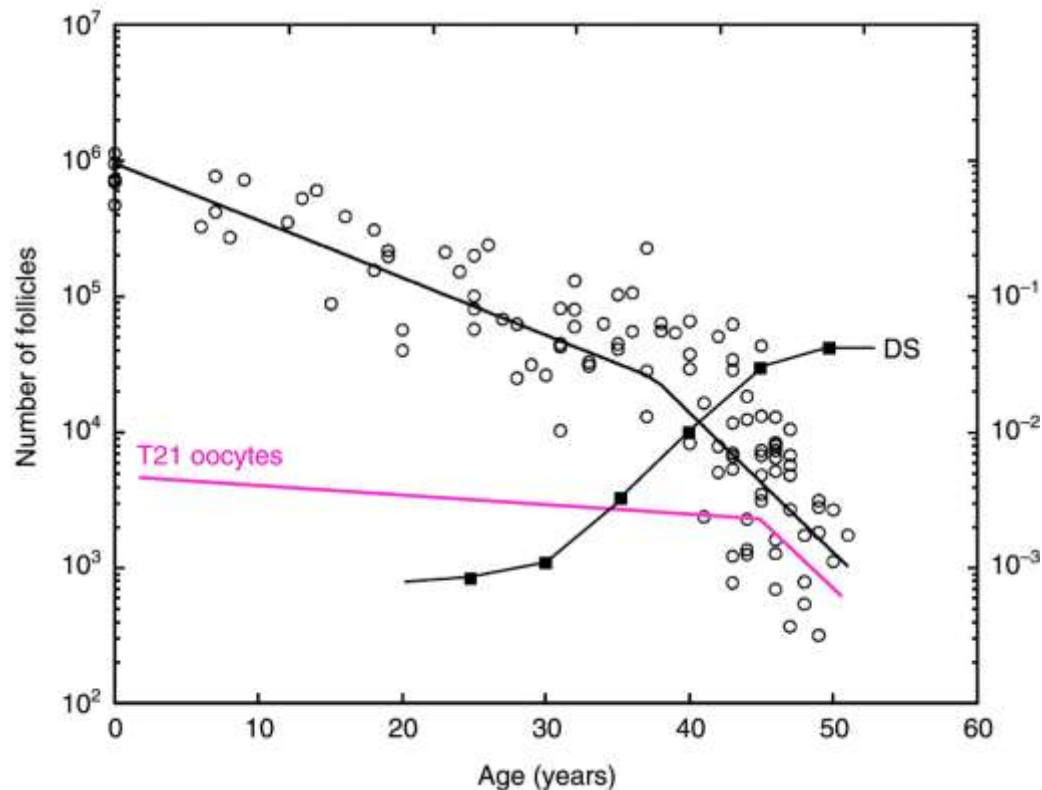
- Homologous chromosomes pairing
- Assembly of the synaptonemal complex
- Chiasmata formation
- Sister chromosome cohesion
- Spindle formation
- etc...



# Mutations in the genes that function during meiosis may play a role in causing non-disjunction



# Germinal mosaicism: the gonads have some cells with trisomy 21 and so some gametes are trisomic



0.54% mosaicism observed by Hultén *et al.* (2008).

accumulation of trisomy 21 oocytes in the ovarian reserve of older women

A microscopic image of sperm cells, showing their characteristic heads and tails. The cells are arranged in a somewhat circular pattern, with some heads pointing towards the center and others towards the periphery. The background is a light, grainy texture.

**Paternal non disjunction  
datas**

# Where did non disjunction causing trisomic Down syndrome occur?

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- Maternal MI 69%
- Maternal MII 21.5%
- Paternal MI 2%
- Paternal MII 3.5%
- Post zygotic 4%

**Table 1.** Origin of nondisjunction in human trisomy 21 by DNA polymorphism analysis

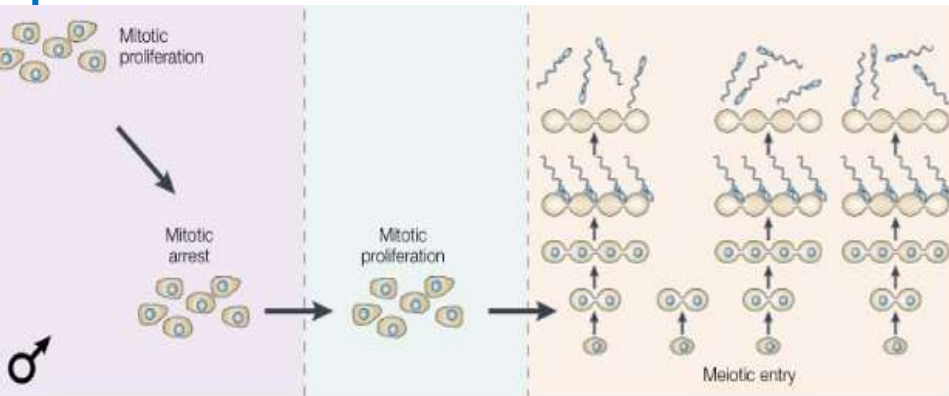
Origin <sup>a</sup>	Number of cases	%	Meiotic recombination
Maternal	732	90.7%	
MI	556	68.9%	Reduced
MII	176	21.8%	Increased
Paternal	44	5.5%	
MI	17	2.1%	Reduced
MII	27	3.3%	
Mitotic	31	3.8%	
“Maternal”	17	2.1%	
“Paternal”	14	1.7%	

<sup>a</sup> MI = meiosis I, MII = meiosis II, “Maternal” and “Paternal” refer to the parental origin of the chromosome that was duplicated by postzygotic nondisjunction. Data from Antonarakis et al. (1993), Lamb et al. (1996), Savage et al. (1998).



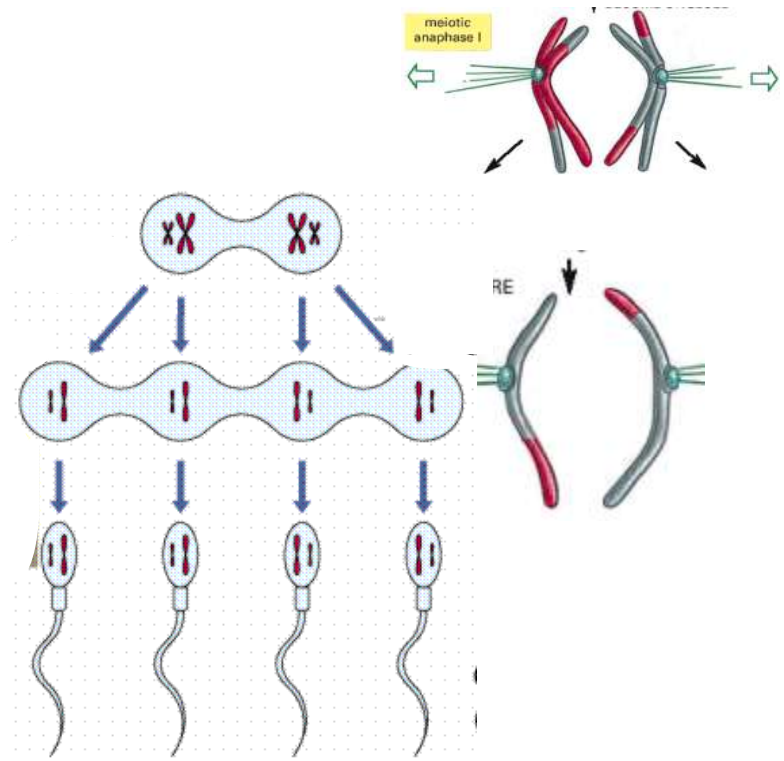
# Period of gametogenesis in the male

## Meiosis starts at puberty



birth

puberty



# Paternal age

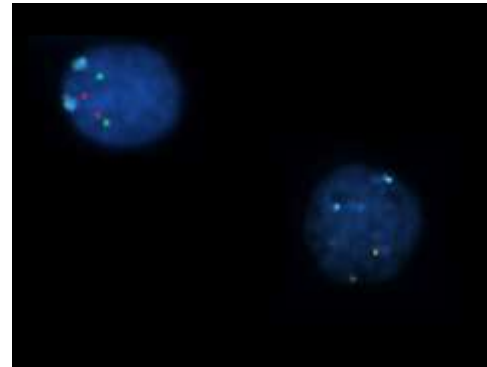
**Table 2.** Mean parental ages by origin of non-disjunction in population-based newborn studies

Origin <sup>a</sup>	n	Maternal age	Paternal age
Maternal			
MI	145	30.1	32.0
MII	50	31.2	33.3
Paternal			
MI+MII	16	25.6	29.9
Mitotic	12	28.2	30.5

<sup>a</sup> MI = meiosis I, MII = meiosis II. Data from Mikkelsen et al. (1995) and Yoon et al. (1996).

# Germinal mosaicism

- FISH to determine testicular T21 mosaicism in four male fetuses showed that male 21 trisomy germinal mosaicism is very low compared to female ovarian T21 mosaicism



Hultén MA et al;2010



# Chromosomal abnormalities

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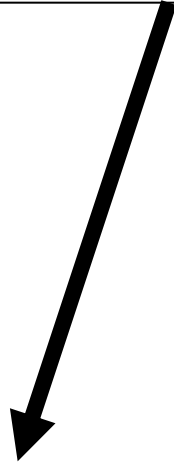
- Numerical
  - Unbalanced
  - Autosomal
  - Sex chromosomes
- Structural
  - Unbalanced vs balanced
  - Transmission

# Consequences of chromosomal abnormalities

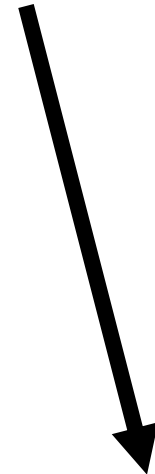
```
graph TD; A[Consequences of chromosomal abnormalities] --> B[Depends on presence or absence of unbalanced chromosome constitution]; B --> C[Unbalanced Phenotypic consequences]; B --> D[Balanced Normal phenotype];
```



Depends on presence or absence of unbalanced chromosome constitution



**Unbalanced  
Phenotypic consequences**



**Balanced  
Normal phenotype**

# Chromosomal abnormalities

Numerical

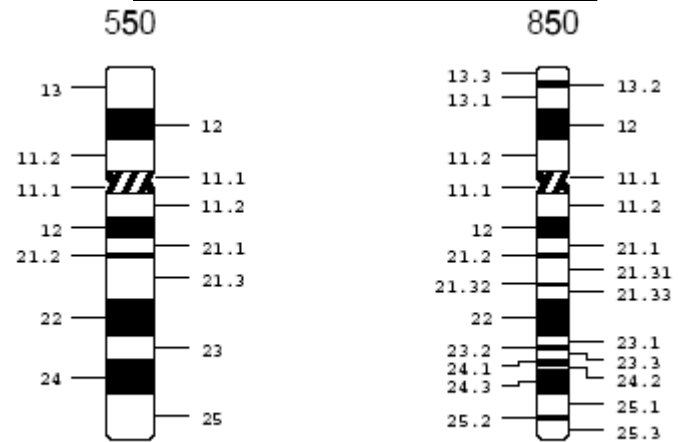


21

Always unbalanced

Abnormal Phenotype

Structural



Unbalanced or Balanced

Normal phenotype

# Numerical Anomalies (Aneuploidies)

Extra  
Chromosomes

+1 Trisomy	47
+2 Tetrasomy	48
+3 Pentasomy	49
+23 Triploidy	69
+46 Tetraploidy	92

Deficient  
Chromosome

-1 Monosomy 45

Chromosome's  
Segment

Partial  
Trisomy



# Viabile aneuploidies

## Autosomes

extra or deficient chromosome material



- Mental Retardation
- Dysmorphy
- +/- Internal Malformations
- +/- Growth Retardation

# Chromosome syndromes

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Down's syndrome  
Trisomy 21



Edward's syndrome  
Trisomy 18



Patau's syndrome  
Trisomy 13

# Malformations (examples)



- Congenital heart defects
- Renal abnormalities
- Brain abnormalities

# Down syndrome

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**Frequency: 1/800 livebirths**

**In newborn: hypotonia and dysmorphic features**

**Frequently associated malformations :**

- Cardiovascular in 50% of cases
- Digestive: duodenal atresia or stenosis

**Mental retardation :**

- IQ around 50 at 5 years of age.



# Chromosome abnormalities in Down syndrome

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- 95% trisomy 21
- 2.5% translocation of chromosome 21 and another acrocentric chromosome
- 2.5% mosaicism

# Aneuploidies of sex chromosomes

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**Mildly or not dysmorphic**  
**Mild or no mental retardation**

**+/- height**

**Fertility problems**

# Klinefelter syndrome

- No frontal baldness
- Poor beard growth
- Breast development
- Female type pubic hair pattern
- Small testicles
- Long legs

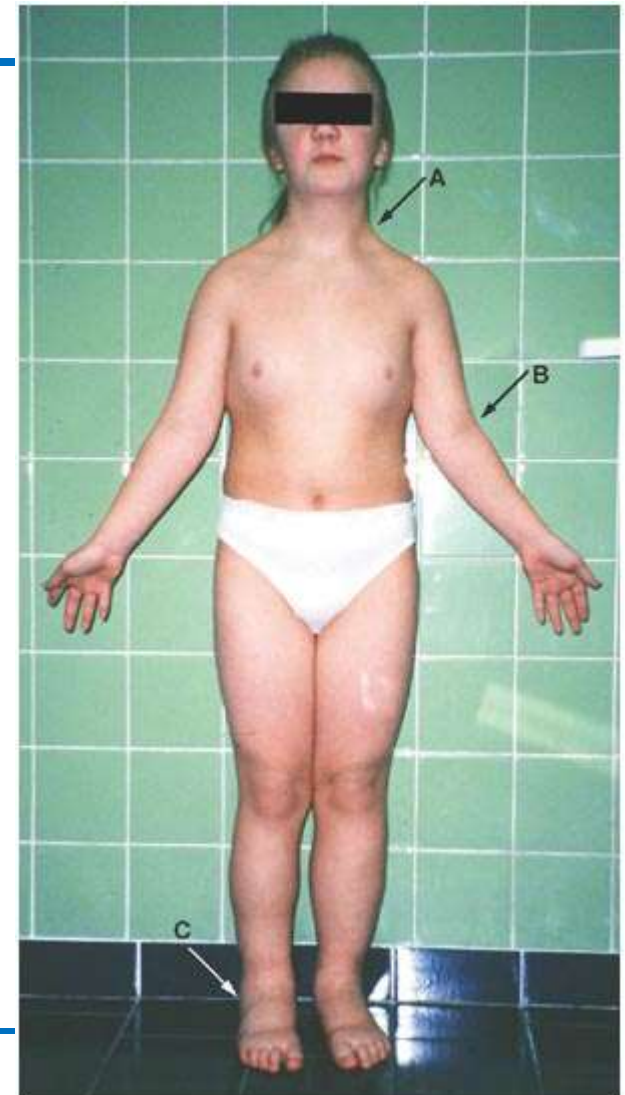
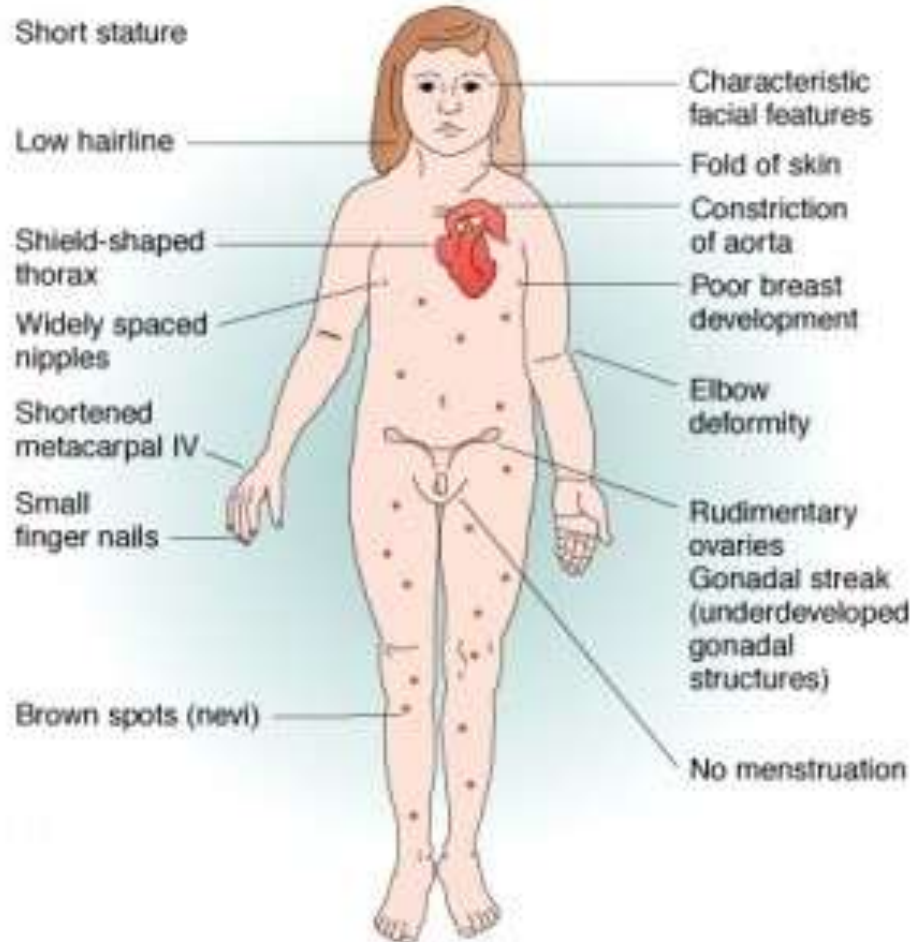


## Cytogenetics :

85 % 47,XXY in all the studied cells

15% *mosaics* 47,XXY/46,XY or 47,XXY/46,XX

# Turner Syndrome



# Cytogenetics Turner syndrome



45,X in 50% of cases, the X chromosome is of maternal origin in 76 % of the cases

45 % of the remaining cases are either numerical variation or structural variation

mosaic : 46,XX/45,X

structural anomalies (could be mosaic) :

ring X : 46,X,r(X)

deletions : del Xp,del(Xq)

isochromosome X : 46,X,i(Xq)



# Structural Balanced Anomalies

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1 Chromosome

**Inversion**  
pericentric  
paracentric

2 Chromosomes

**Translocation**  
reciprocal  
Robertsonian  
**Insertion**

Complex

Structural anomalies

```
graph TD; A[Structural anomalies] --> B[Unbalanced after meiosis]; A --> C[Balanced]; B --> D["Abnormal Gametes  
Partial Anomalies"]; D --> E[Abnormal zygotes]; C --> F[Normal phenotype]
```

Unbalanced after meiosis

Balanced

Abnormal Gametes  
Partial Anomalies

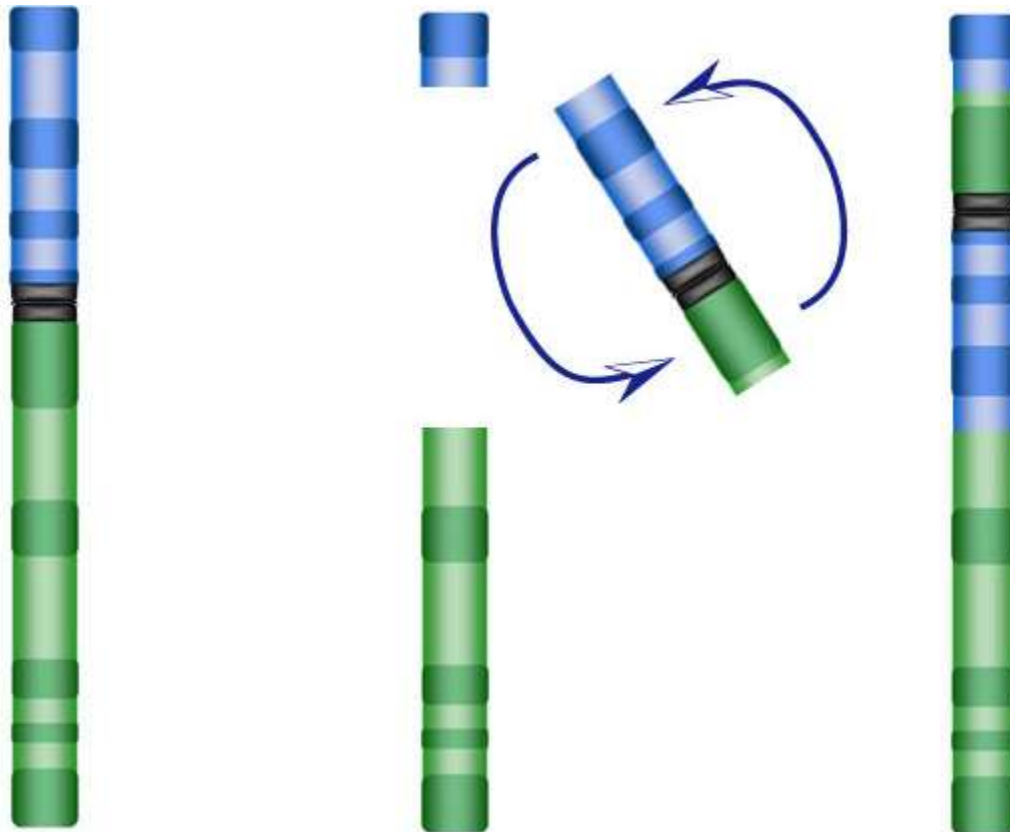
Abnormal zygotes

Normal phenotype

# Pericentric Inversion

1 chromosome  
2 breakpoints

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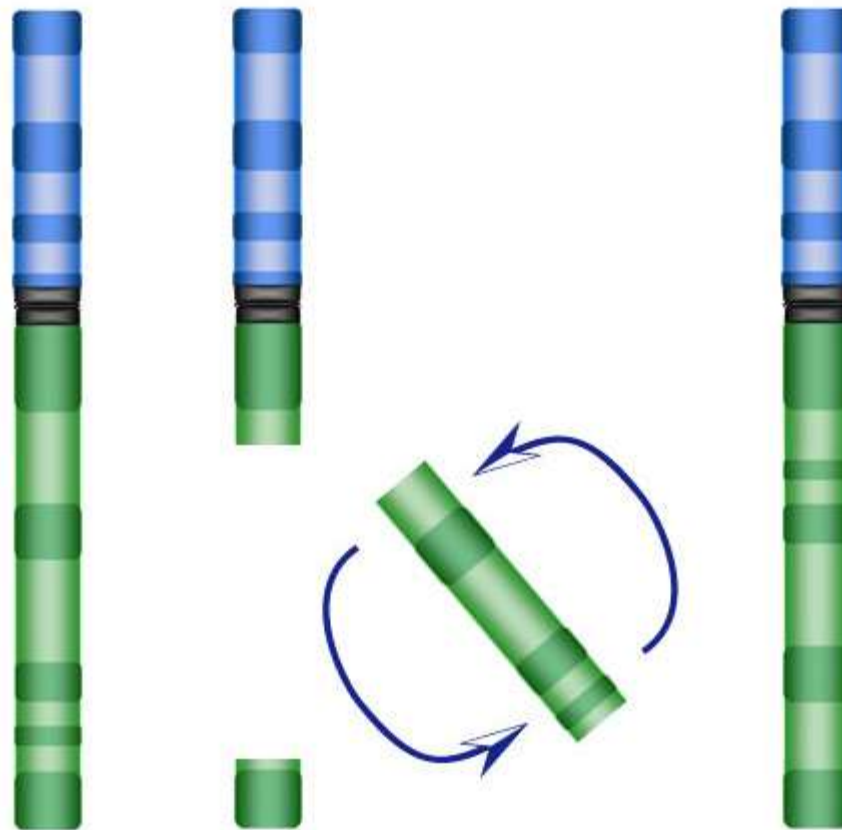
All inversions 1/1000 newborns



# Paracentric inversion

1 chromosome  
2 breakpoints

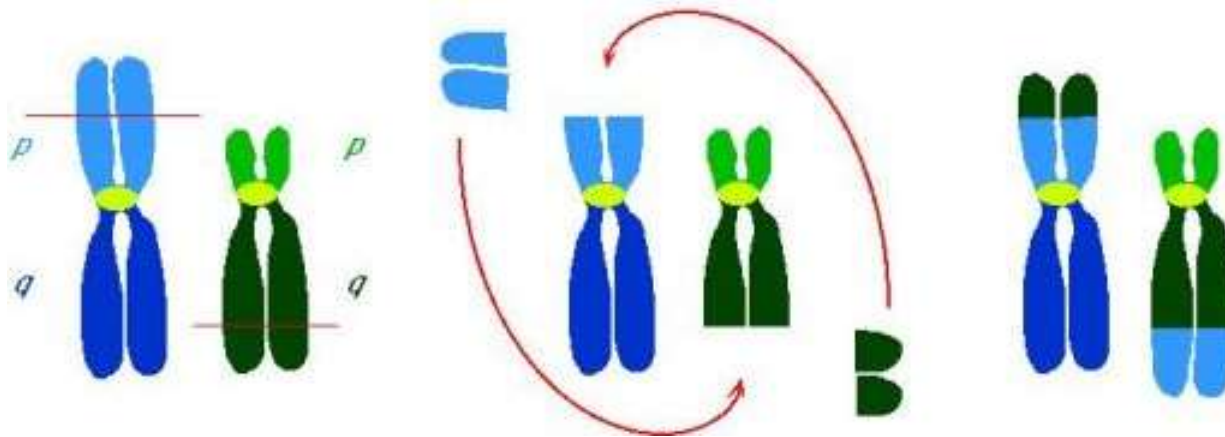
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# Reciprocal translocation

2 chromosomes  
2 breakpoints

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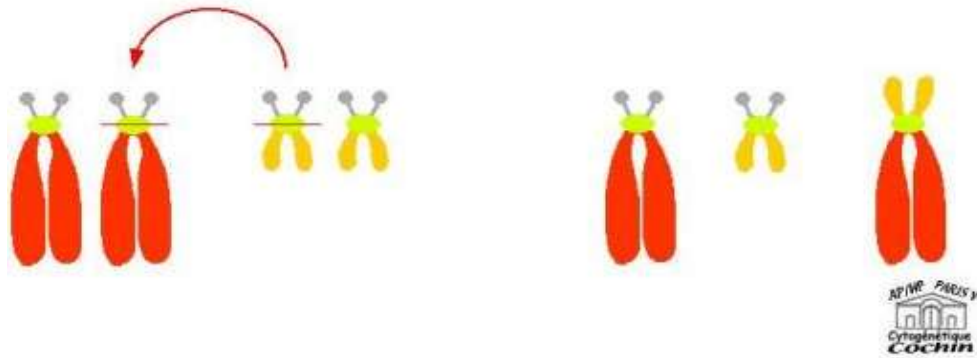


all translocations 1/500 newborns

# Example: translocation between q arm of a chromosome 11 and q arm of a chromosome 22



# Robertsonian Translocation ACROCENTRICS

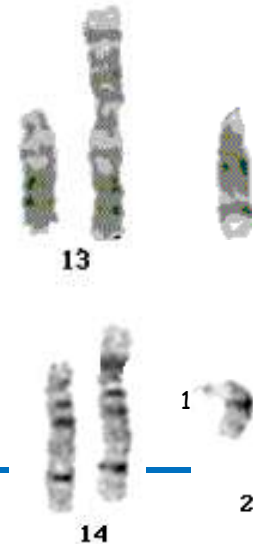


Robertsonian translocations 1/833 newborns

Evans et al.1978

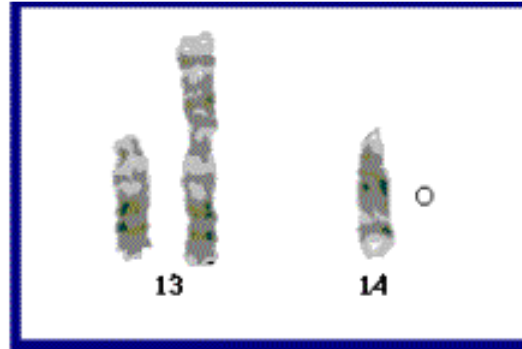
$45,der(13;14)(q10;q10) \Rightarrow 73\%$

$45,der(14;21)(q10;q10) \Rightarrow 10\%$

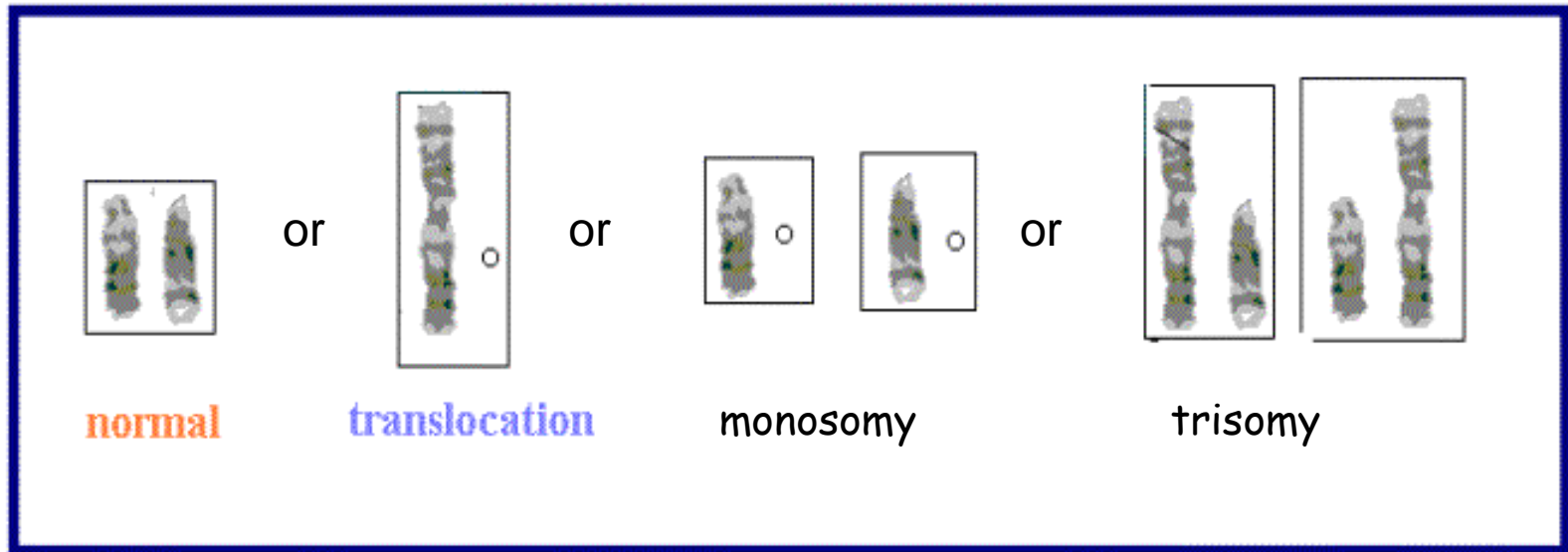


# Meiosis chromosomal segregation of a t(13;14) translocation

Constitutional karyotype

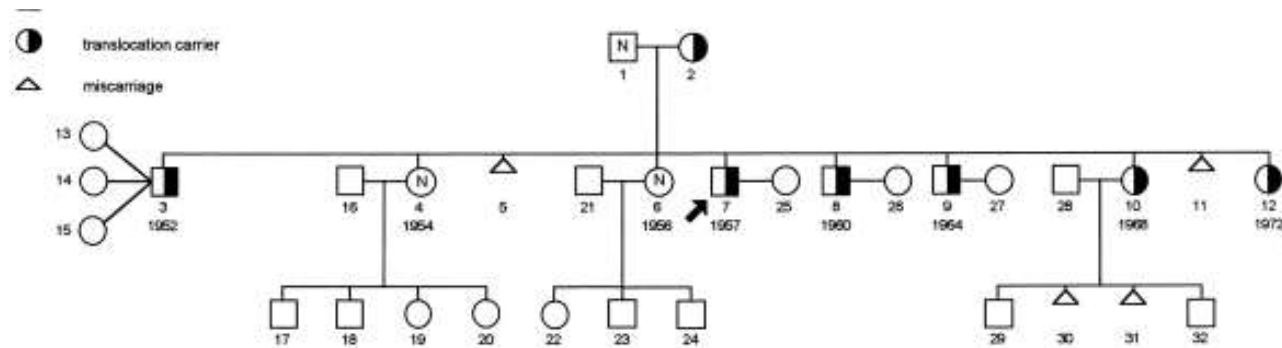


Gametes



# Clinical Consequences of a Translocation

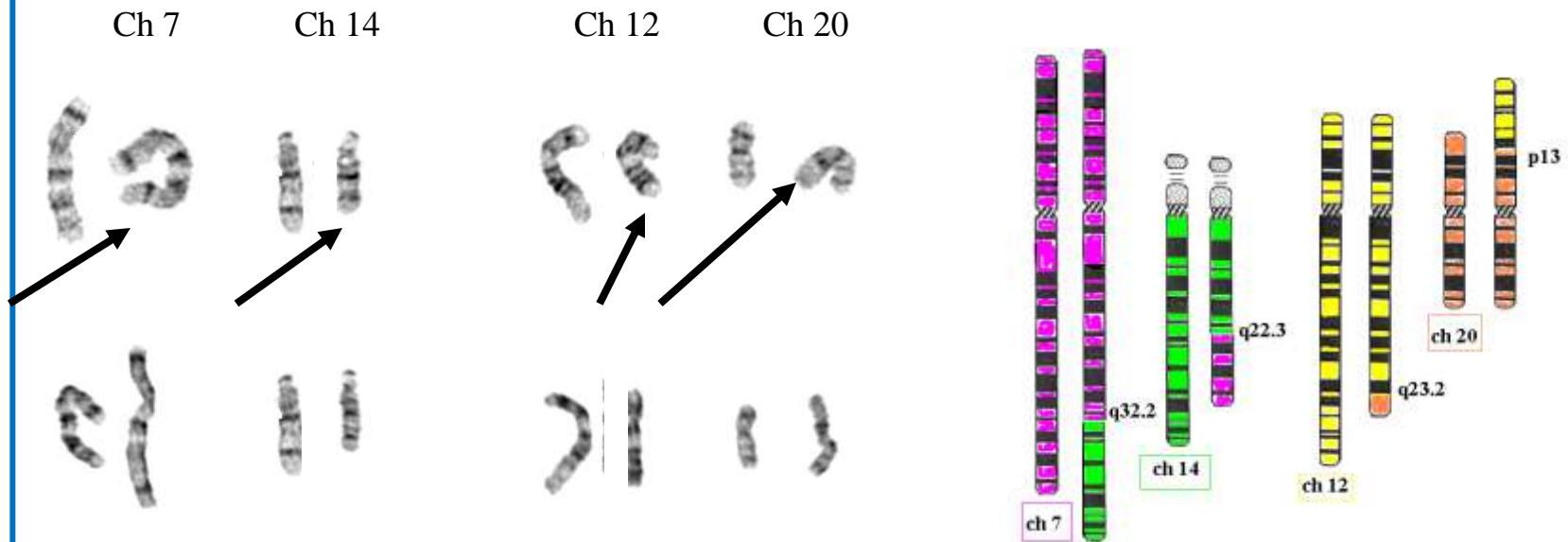
- **Infertility**
- **Miscarriages**



- **Trisomy** by transmission of unbalanced translocation



**Partial Karyotype (GTG banding) of the double translocation  
 $t(7;14)(q32.2;q22.3), t(12;20)(q23.2;p13)$**



*Exemple of complex karyotype with 2 familial translocations*

# Unbalanced Structural Anomalies

1 Chromosome

Deletion  
Duplication  
Ring

2 Chromosomes

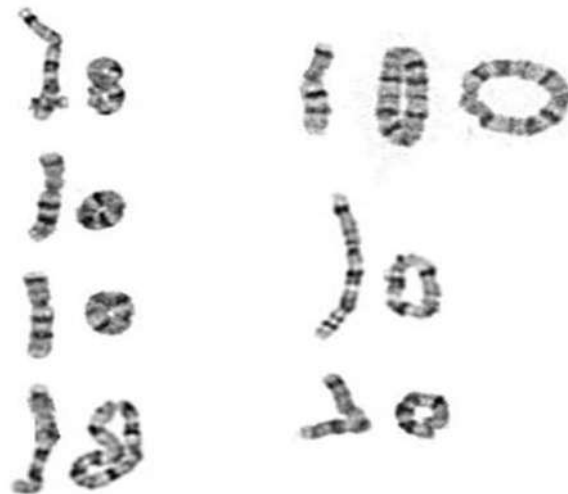
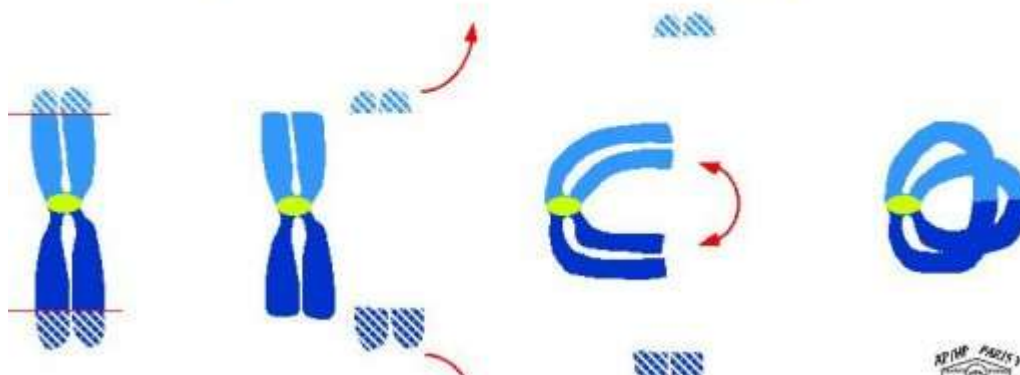
Translocation  
Insertion

Complex

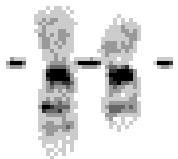
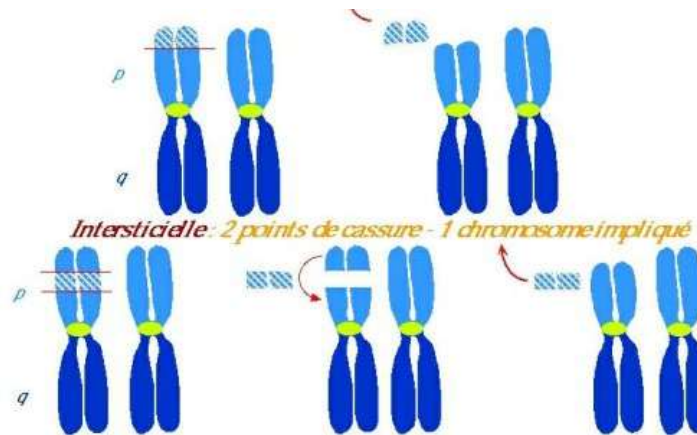


# Ring

1 chromosome  
2 breakpoints

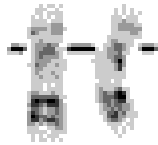


# Deletion



Del(16)(q21)

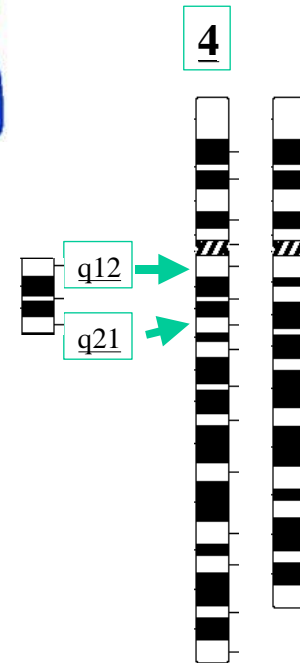
**16**



**17**



**18**



del(4)(q12q21.1)



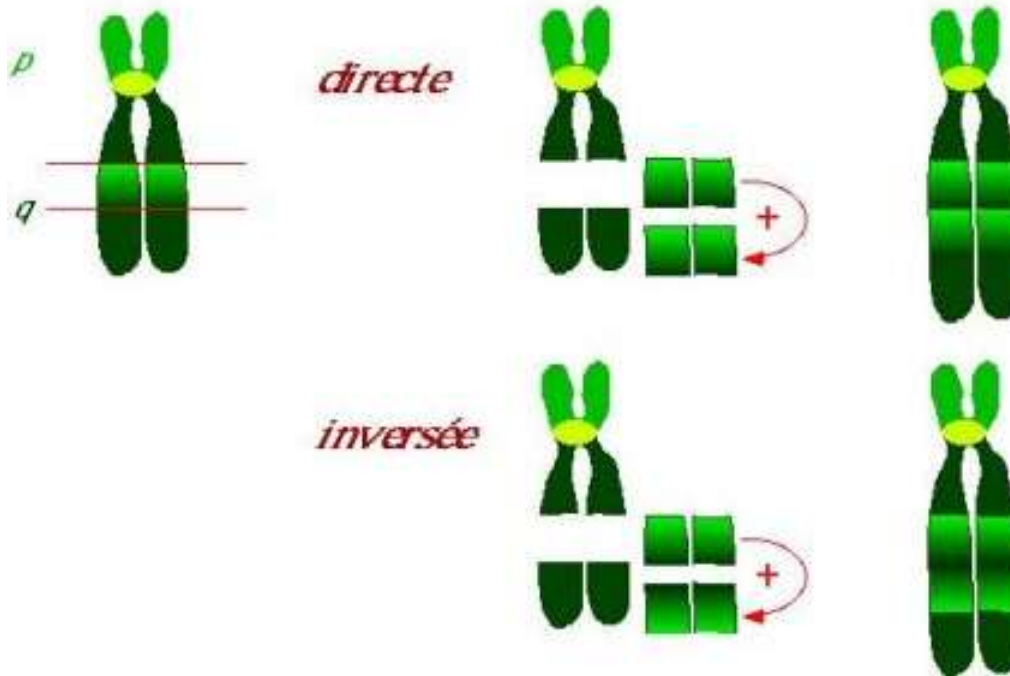
**4**

**Terminal deletion**  
1 chromosome  
1 breakpoint

**Interstitial deletion**  
1 chromosome  
2 breakpoints

# Duplication

1 chromosome  
2 breakpoints



# Conclusions

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- Chromosomes can be studied in any nucleated cell postnatally as well as prenatally from chorion villus samples and amniocytes
- 1/160 newborns has a chromosome abnormality
- The most common syndromes are Down syndrome (trisomy 21) and Klinefelter syndrome (47,XXY)