

Introduction to Clinical Cytogenetics

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What is Cytogenetics?

- study of chromosomes; for clinical cytogenetics, it is applying the study of chromosomes to clinical medicine (diagnosis - postnatal, prenatal, cancer; gene mapping)

What types of tissues can be used to study chromosomes?

- for routine chromosome analysis, cells must have a nucleus, be viable at time of collection and capable of undergoing cell division (cells are grown in culture)

- for certain molecular cytogenetic applications, cell division is not a requirement

Why perform clinical cytogenetic testing?

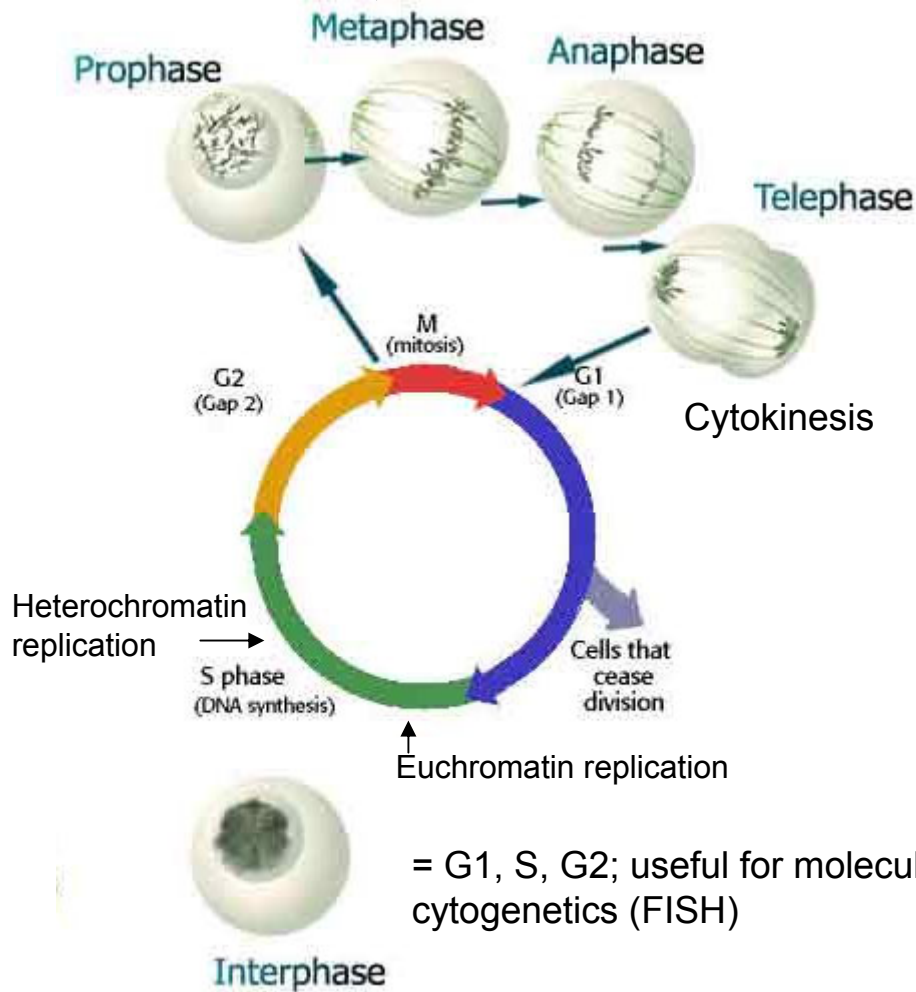
- fertility problems – couples with history of infertility or habitual pregnancy loss; females with amenorrhea
- malformed fetus; stillbirths and neonatal deaths where a cytogenetic basis is likely
- pregnancy in women of advanced age
- family history in first degree relatives
- dysmorphology, problems of early growth and development (FTT, DD, Dysmorphic facies, MR, ambiguous genitalia, multiple malformations including cardiac anomalies
- Neoplasia (can be diagnostic or prognostic)

Frequency of Chromosome Abnormalities

- ½ to 1% of all live births
- 2% of recognized pregnancies in women > 35 y.o.
- 10% of stillbirths
- ½ of recognized spontaneous pregnancy losses
- high in neoplasias (type dependent)

Mitotic Cell Cycle:

Stage examined in routine cytogenetics



- G1- cell grows and performs normal metabolism; organelles duplicate (~10-12 hours)
- S – DNA replication and chromosome duplication (~6-8 hours)
- G2 – cell grows and prepares for mitosis (~2-4 hours)
- M – mitosis – chromosomes are maximally condensed (~ 1hour)
- transitions between phases mark crucial regulatory checkpoints and are often rate limiting



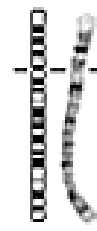
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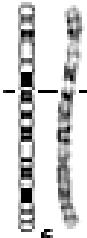
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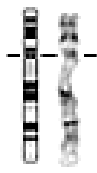
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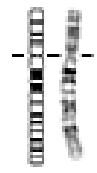
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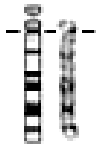
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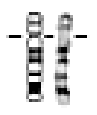
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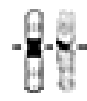
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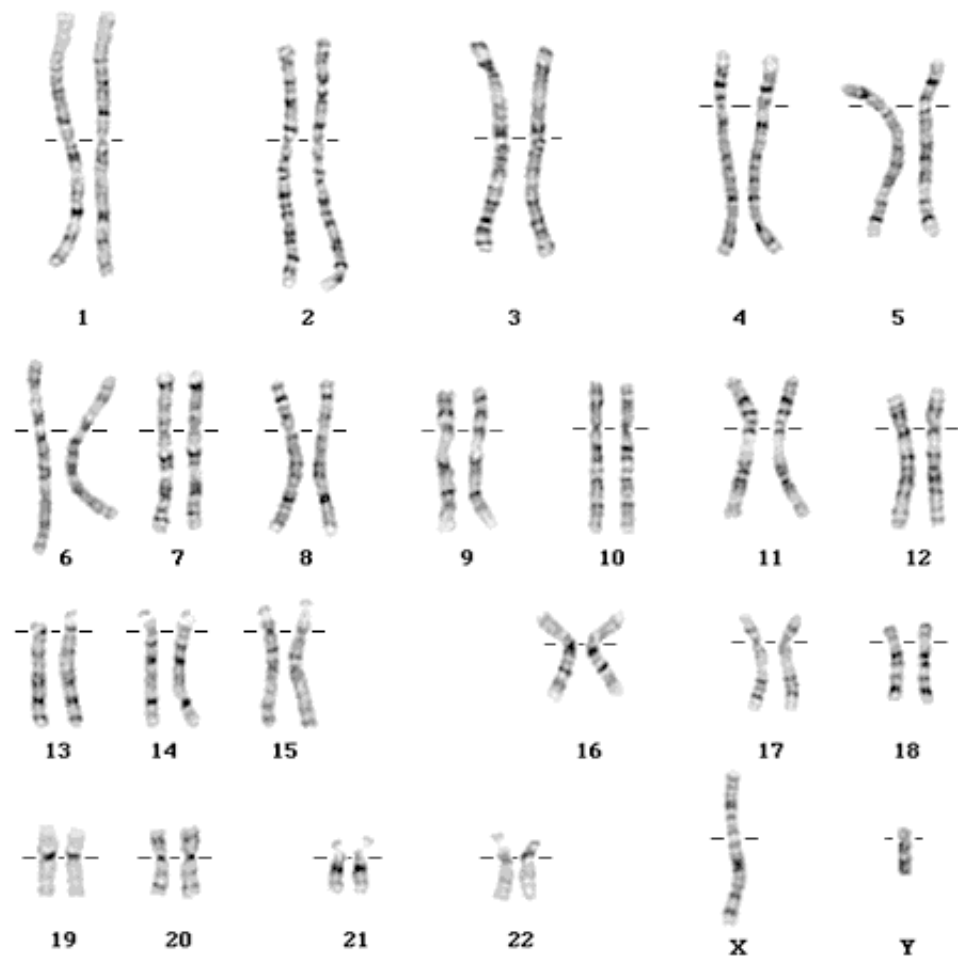
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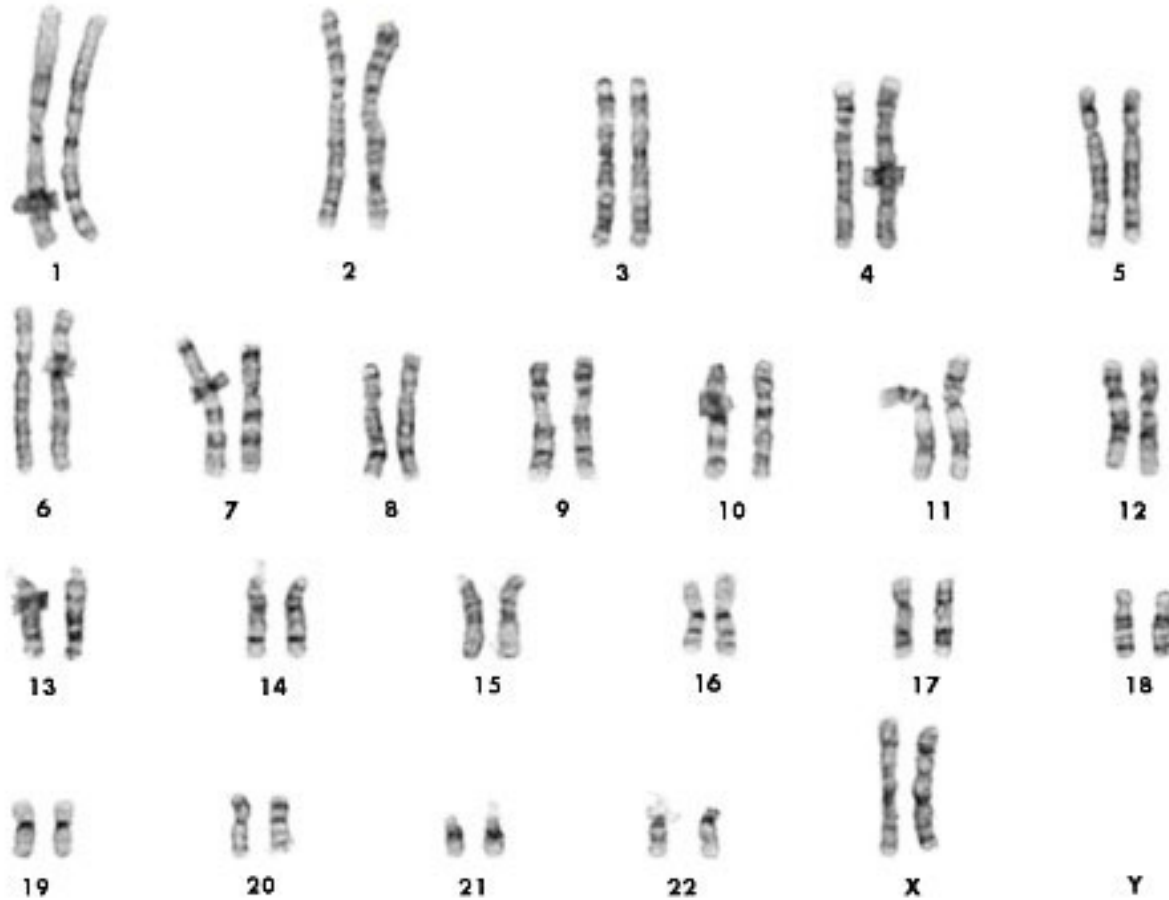
X



Y



Normal Female karyotype



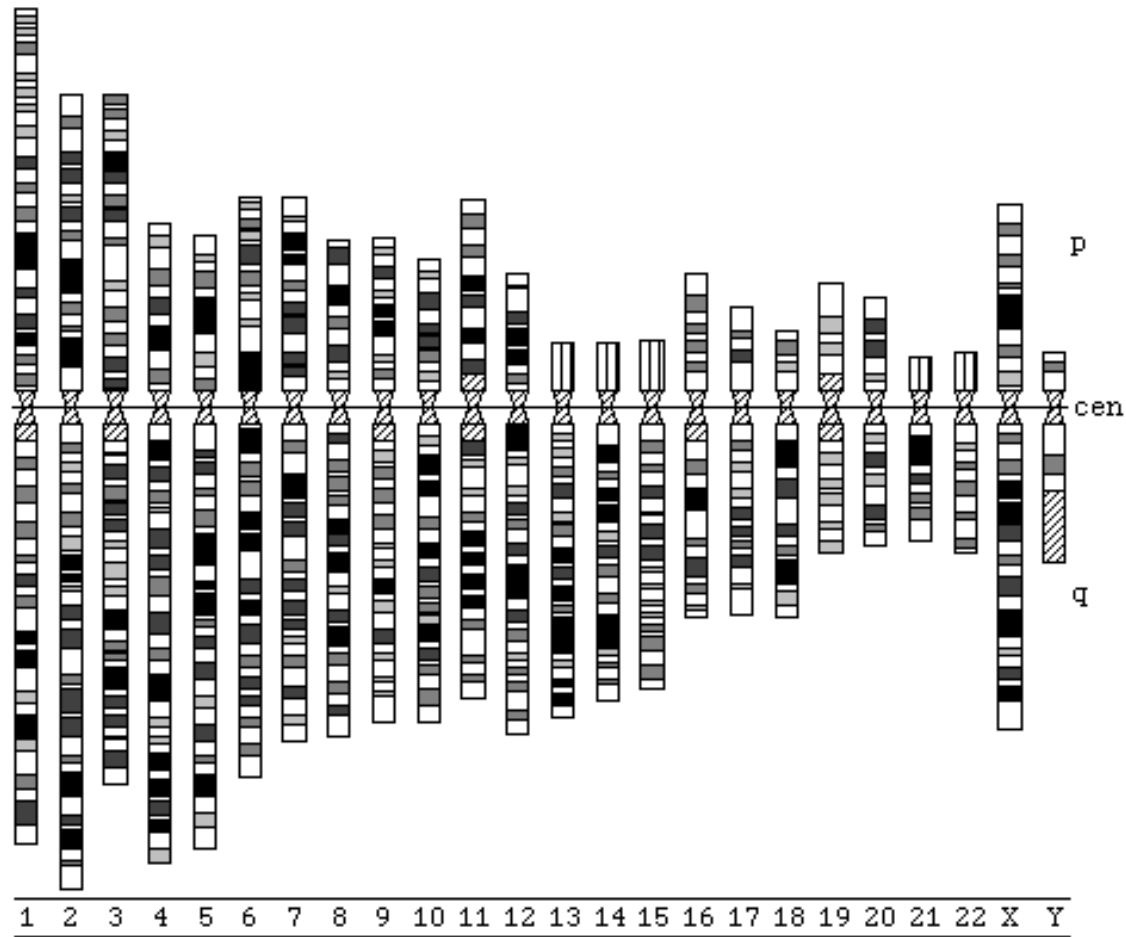
From Children's Mercy Hospital Cytogenetics Laboratory

Normal Male karyotype



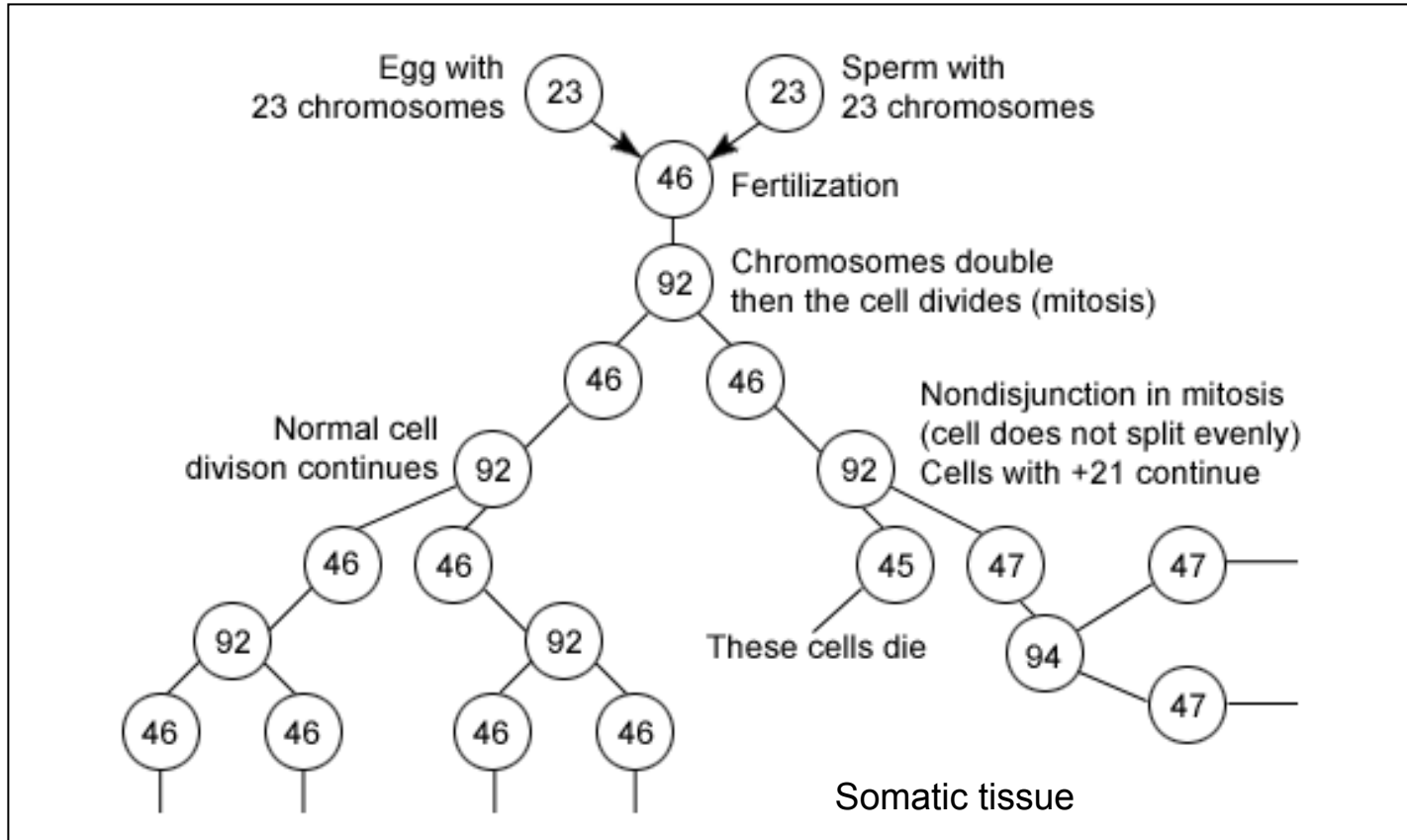
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Ideogram: Human Chromosomes



Chromosomal Mosaicism:

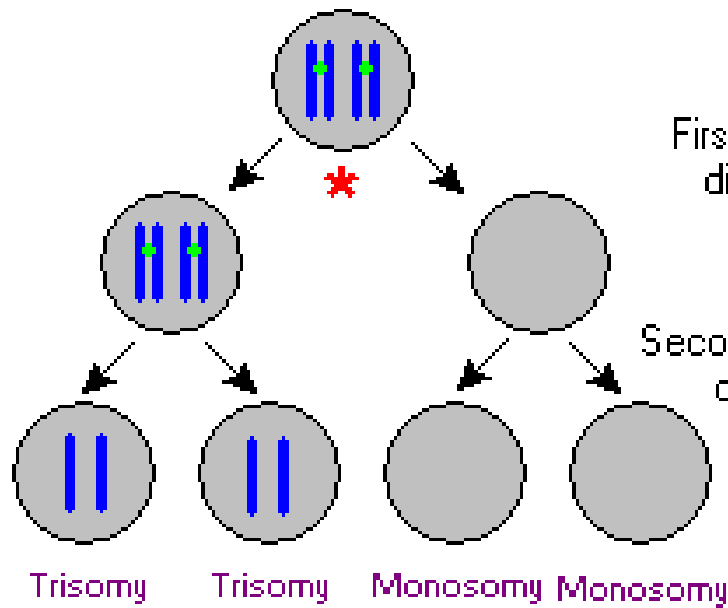
One or more cell lines in an individual derived from a single zygote



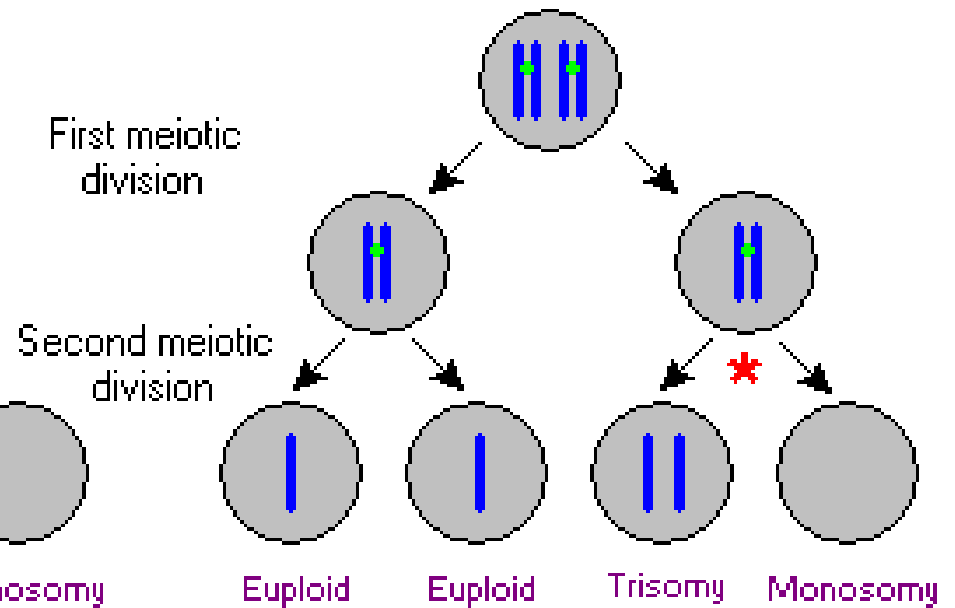
Eg. **47,XX,+21/46,XX or 47,XY,+21/46,XY**

Q: Does 46,XX/46,XY represent a mosaicism?

Nondisjunction in meiosis I

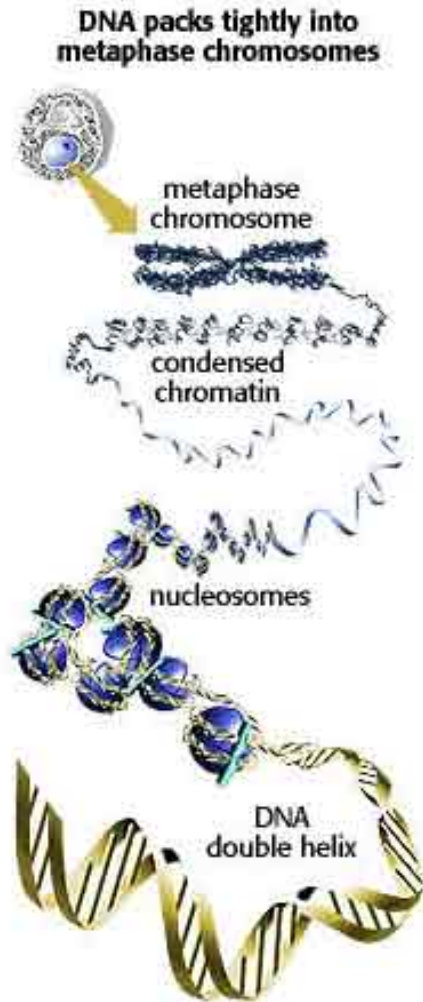


Nondisjunction in meiosis II



Genome of offspring after fertilization with another normal gamete

Chromosome Structure:

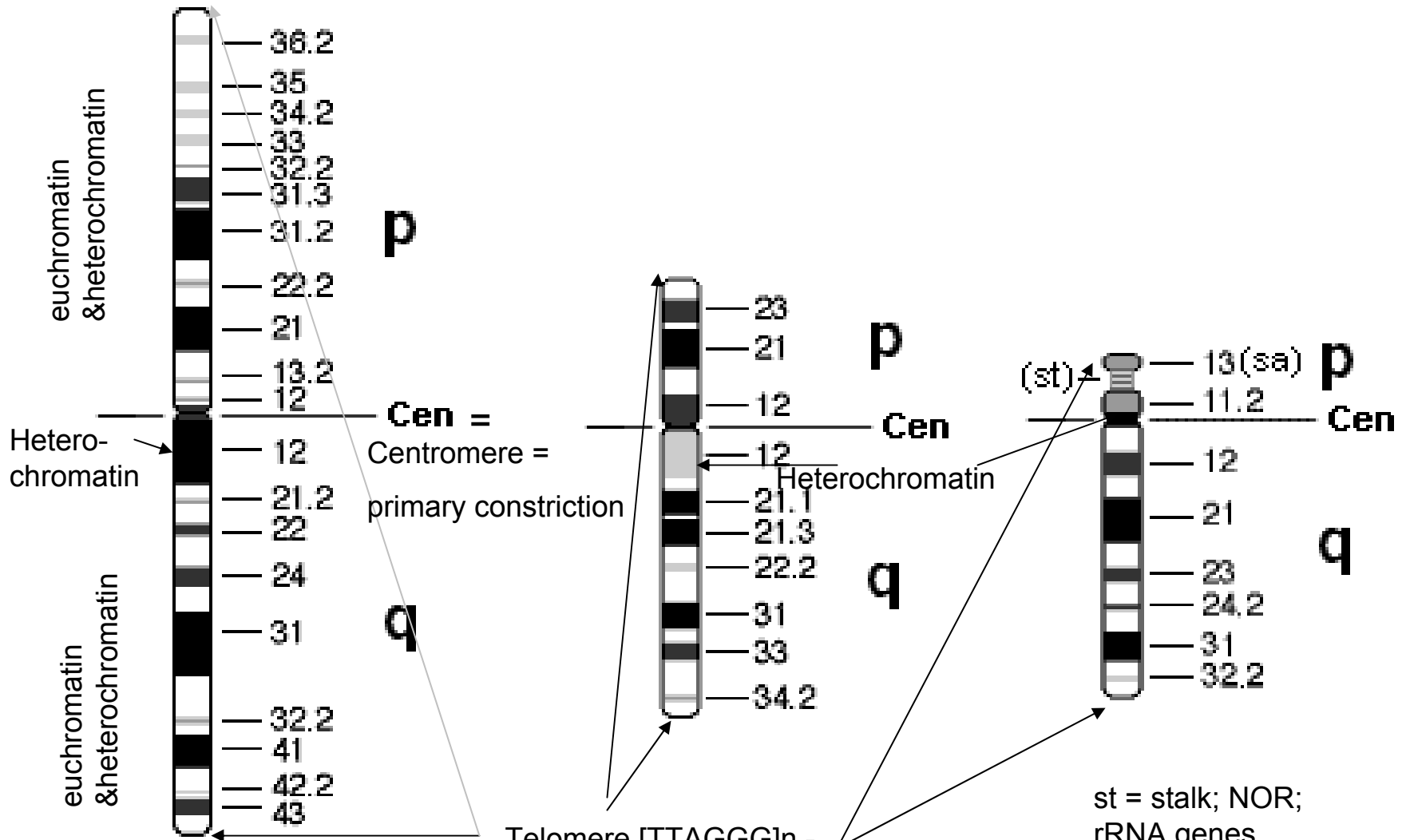


- 3 billion nucleotides per human haploid genome (23 chromosomes)
- chromosome sizes range from 35 million (chrom 22) to ~180 million (chrom 1)
- >2 meters of DNA per diploid nucleus; >10,000 fold compaction
- cell needs to access information in timely and orderly manner
- human genome draft sequence published in Feb'02; completed sequence expected later this year

Metacentric

Submetacentric

Acrocentric



Cen = Centromere = primary constriction

Telomere [TTAGGG]_n - Not chromosome specific; adjacent subtelomeric regions are chromosome specific

st = stalk; NOR; rRNA genes
sa = satellites

Nomenclature:

of chromosomes with centromeres, sex chromosomes, abnormalities listed by chromosome number from 1 through 22 and alphabetically by type of abnormality

Egs. 46,XY and 46,XX

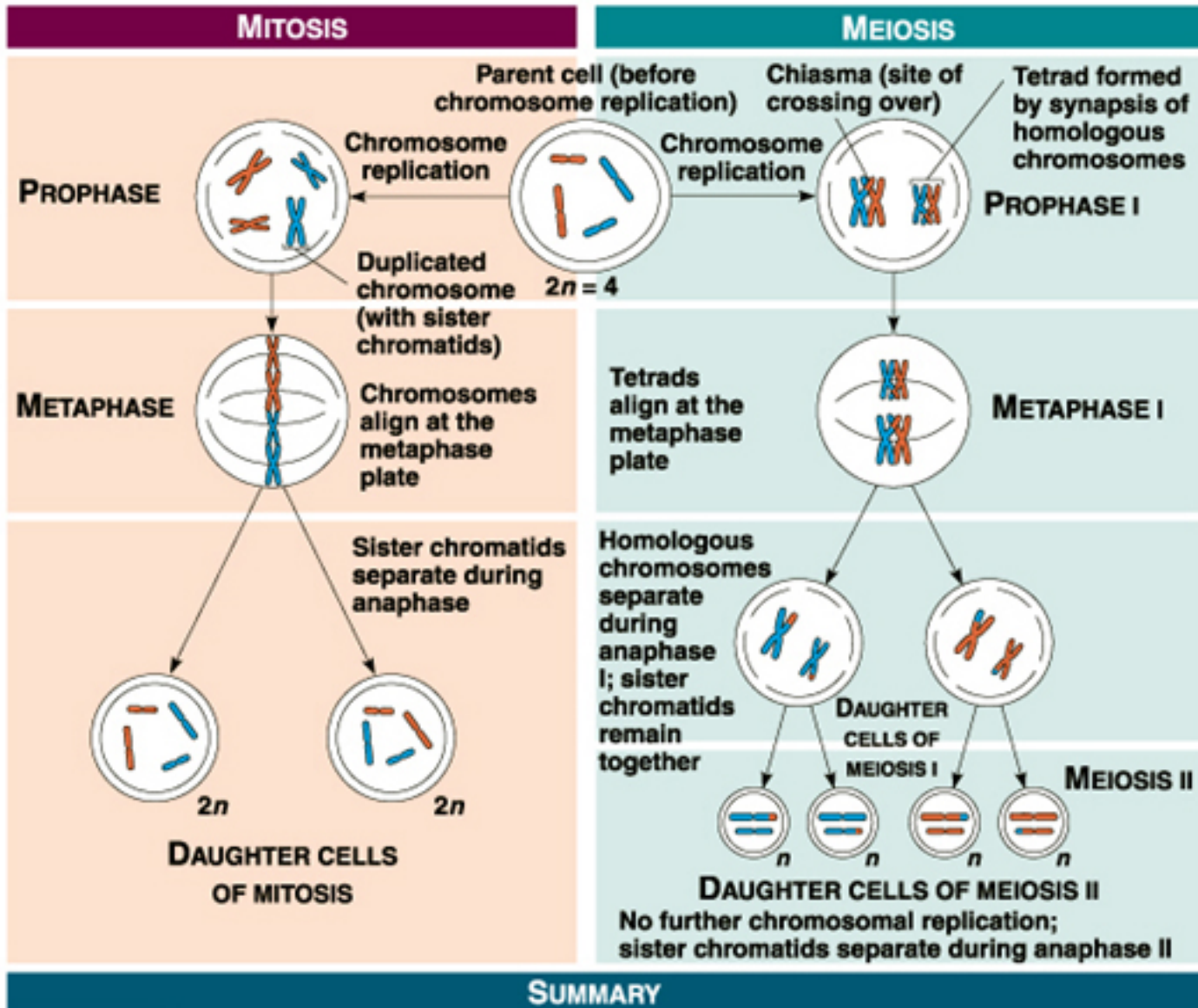
47,XXY and 47,XX,+21 and 45,X

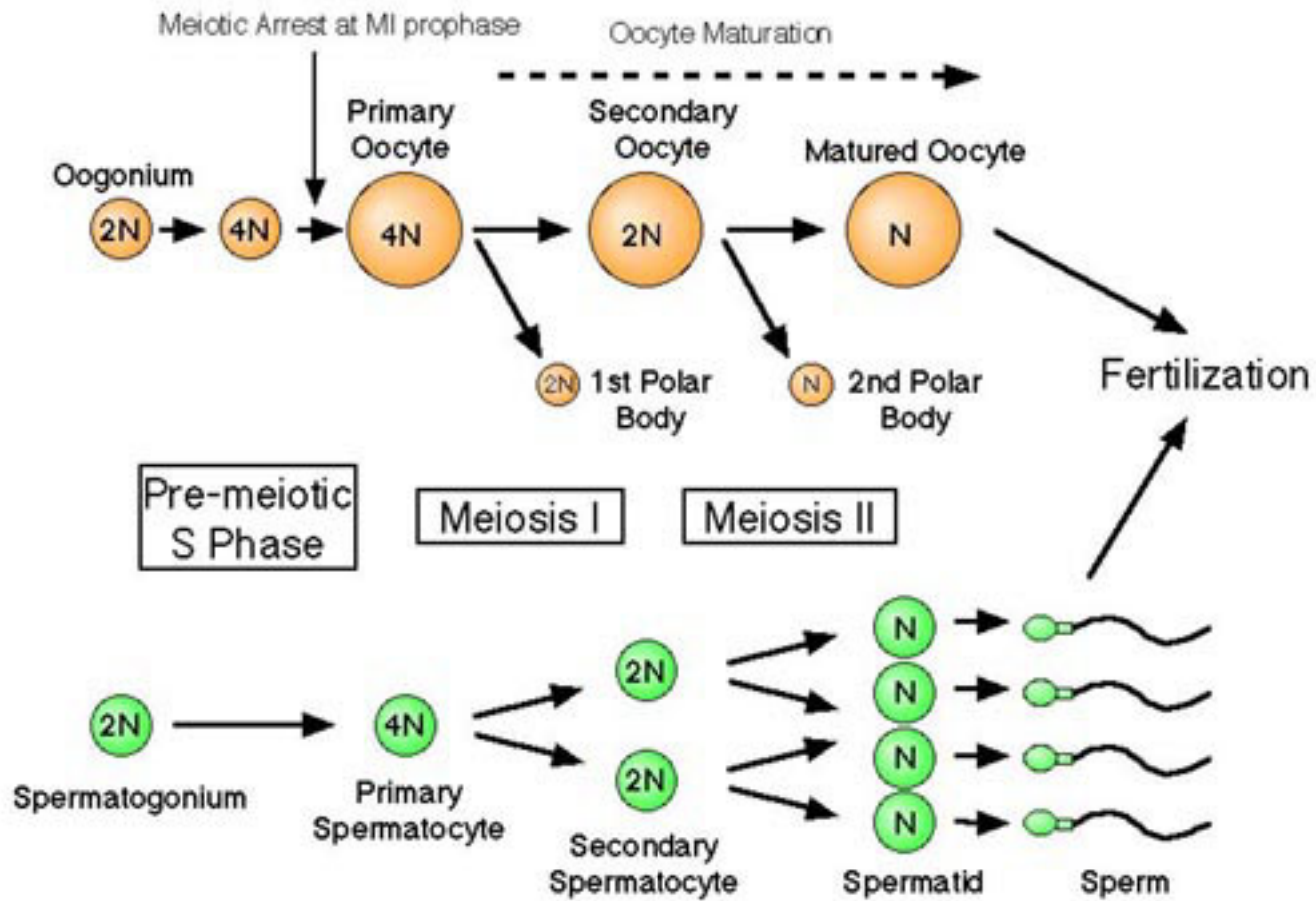
46,XX,del(15)(q11.2q13)

45,XX,t(14;21)(q10;q10)

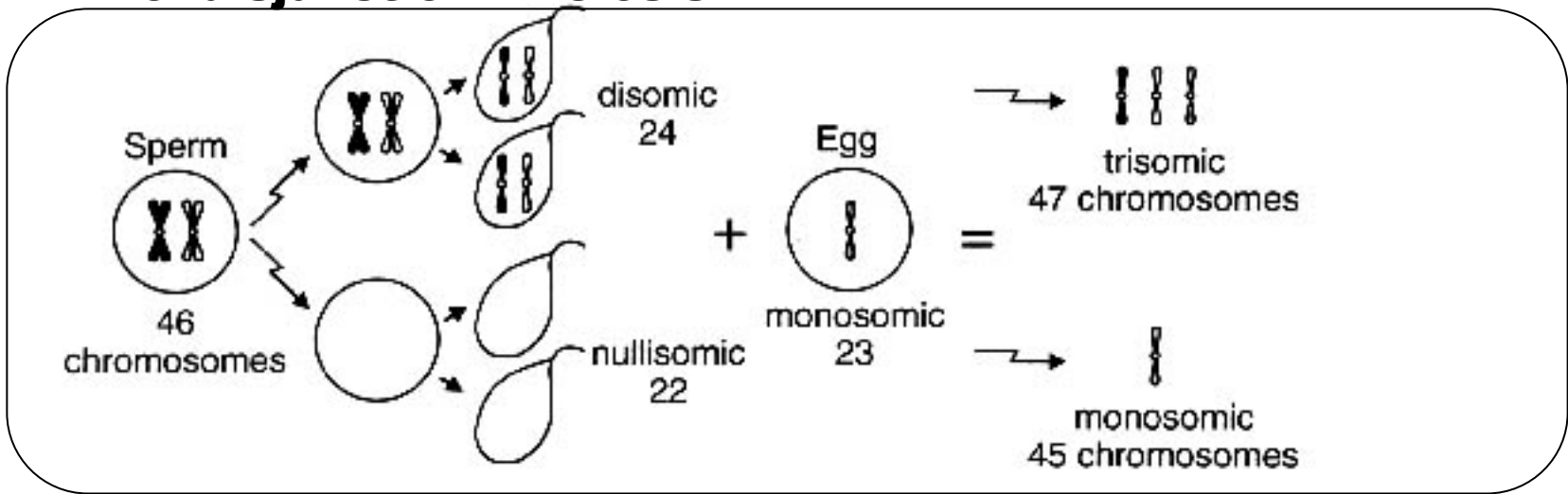
46,XY,t(14;21)(q10;q10),+21

47,XY,t(1;17)(p32;q12),inv(3)(p25),+4,add(6)(q21q23),del(6)(p25)

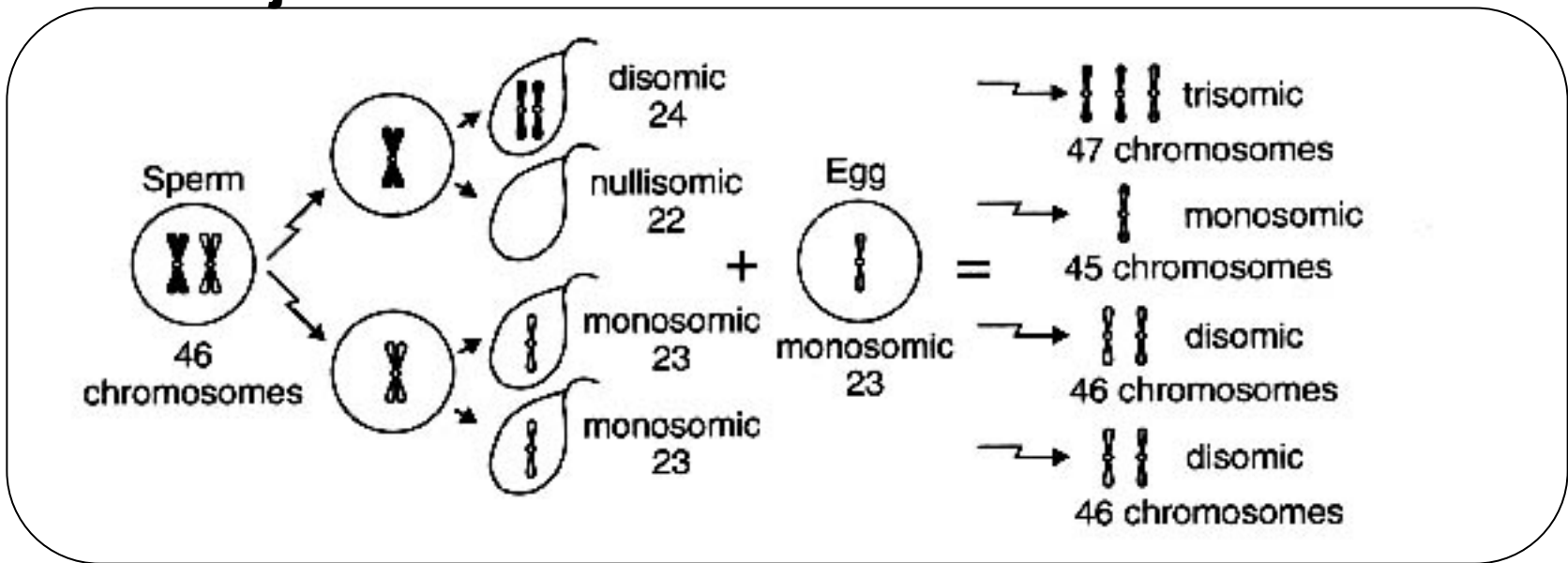




Nondisjunction: Meiosis I



Nondisjunction: Meiosis II

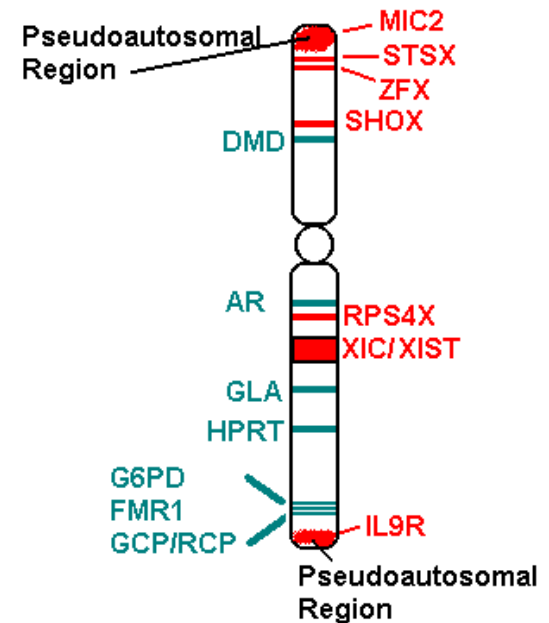


Fertilization

Zygotes

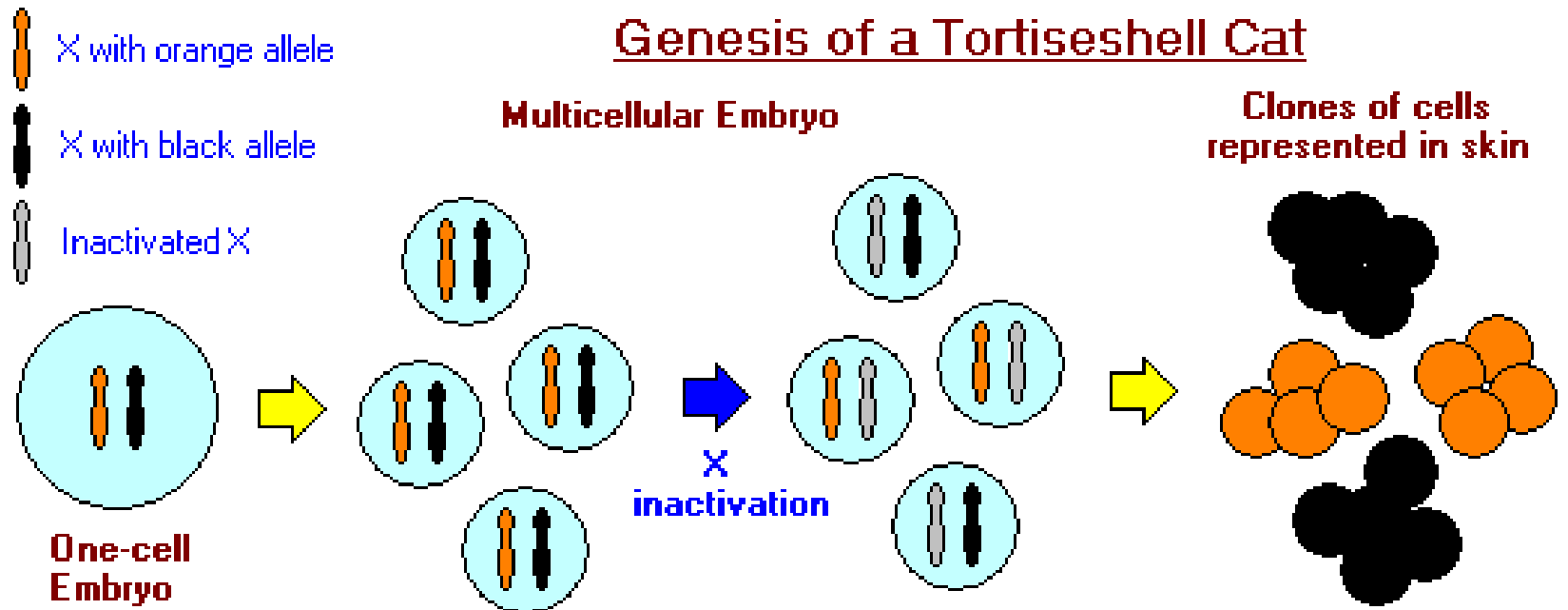
X-Inactivation or Lyonization:

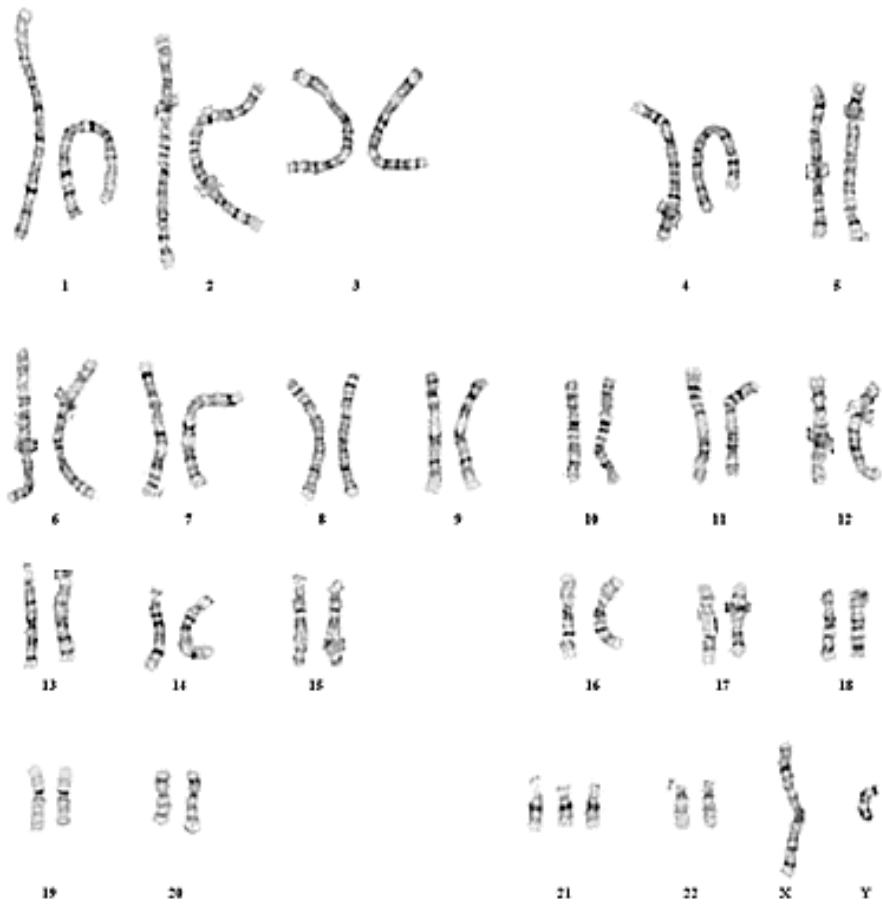
- form of gene dosage compensation between females (XX) and males (XY)
- generally occurs randomly in blastocyst (maternal or paternal)
 - if abnormal X involving t(X;autosome) then abnormal X remains active
 - if abnormal X involves deletion, duplication, etc then abnormal X is inactive
- permanent
- one active X per cell but inactive X does have some genes that remain active
- inactive X is late replicating, forms Barr body and is very compacted
- XIST gene (RNA) initiates inactivation and is expressed from inactive X



Egs. of inactive (left) and active (right) genes on 'inactive' X.

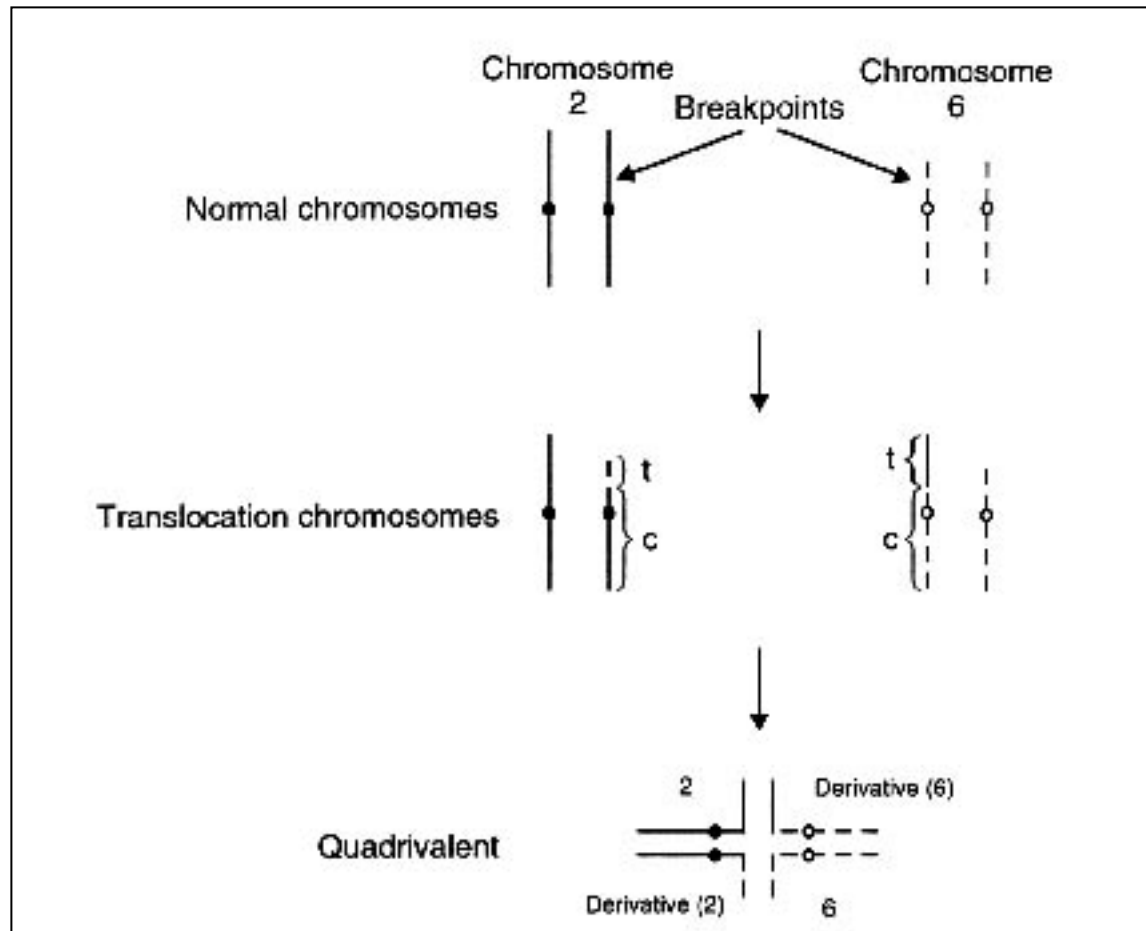
X-Inactivation or Lyonization





47,XY,+21

Meiotic Pairing of a Reciprocal Translocation



Possible Gametes: 2,6 and der(2),der(6) [Alternate segregation – balanced];
 2,der(6) and der(2),6 [Adjacent 1 – unbalanced];
 2,der(2) and der(6),6 [Adjacent 2 – unbalanced]