

Introduction to the human genome

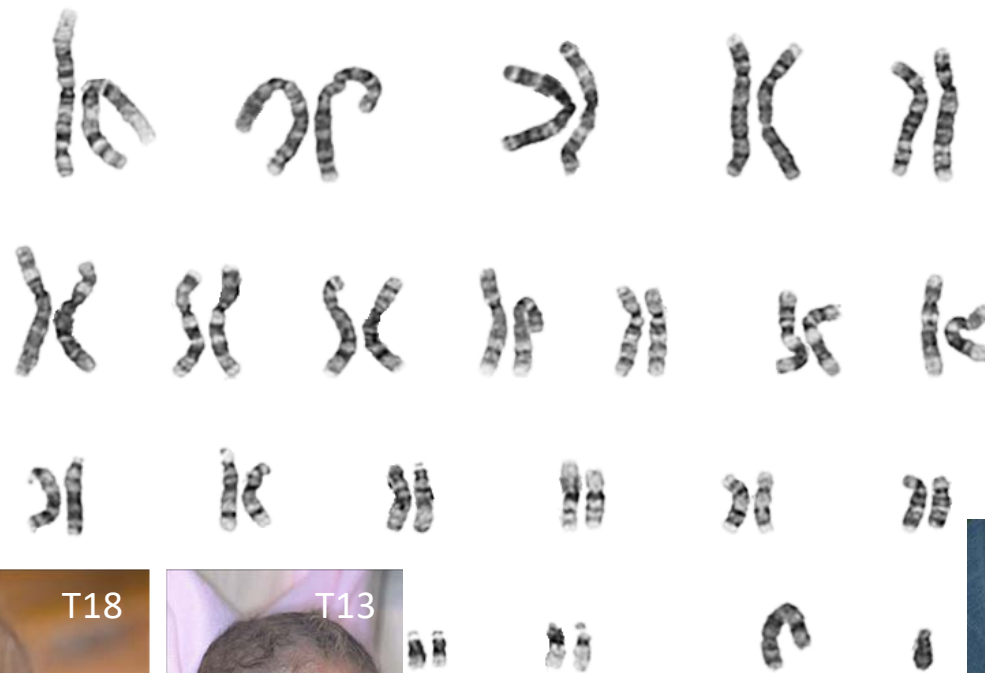
Chapter 2

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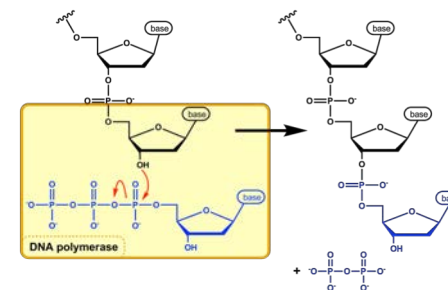


- ✓ The Human Genome Sequence
- ✓ Organization of the Human Genome
 - Single-Copy DNA Sequences
 - Repetitive DNA Sequences
 - Repetitive DNA and Disease
- ✓ Variation in the Human Genome
- ✓ Transmission of the Genome
 - The Cell Cycle
 - Mitosis
 - Meiosis
- ✓ Human Gametogenesis and Fertilization
 - Spermatogenesis
 - Oogenesis
 - Fertilization
- ✓ Medical Relevance of Mitosis and Meiosis

1956 : a normal cell has 46 chromosomes



1975 Sanger sequencing (chain-terminator method)



dATP, dCTP, dGTP, dTTP
+ ddATP

dATP, dCTP, dGTP, dTTP
+ ddCTP

dATP, dCTP, dGTP, dTTP
+ ddGTP

dATP, dCTP, dGTP, dTTP
+ ddTTP

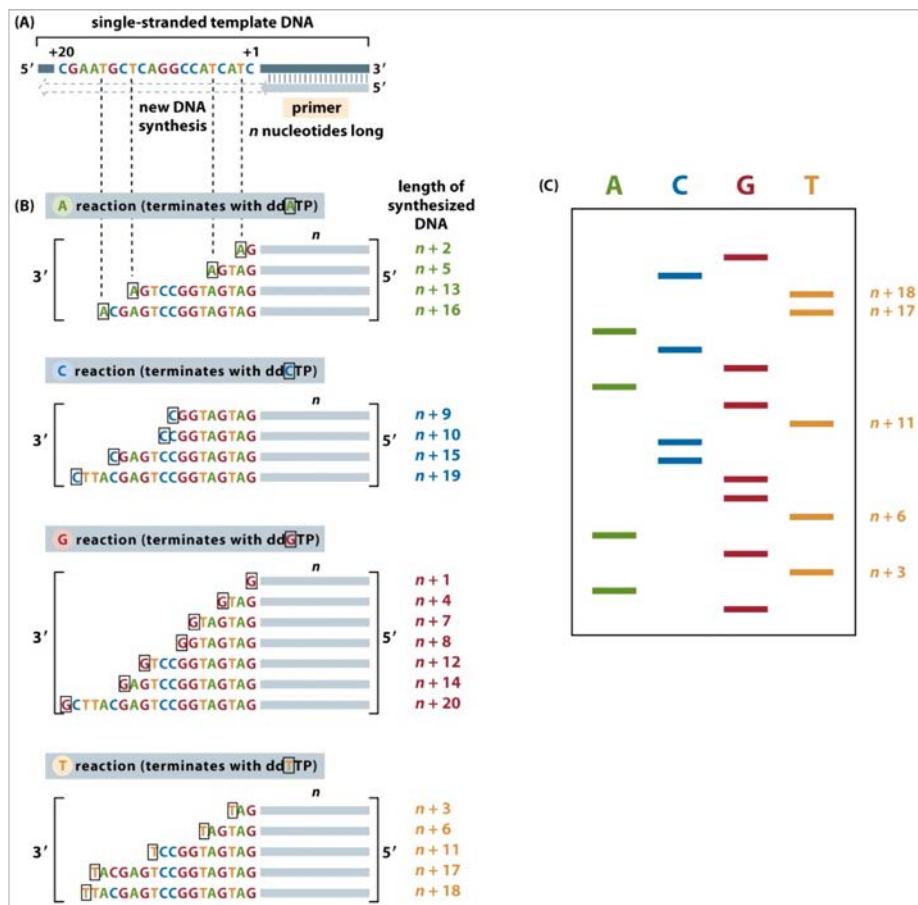


Figure 8.6 Human Molecular Genetics, 4ed. (© Garland Science)

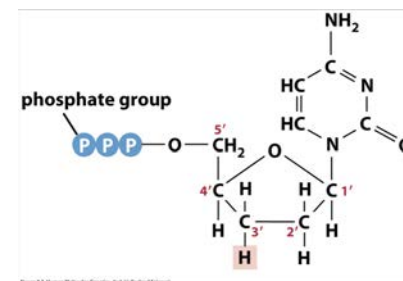


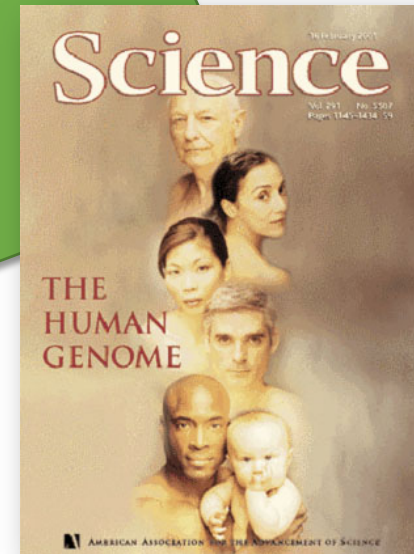
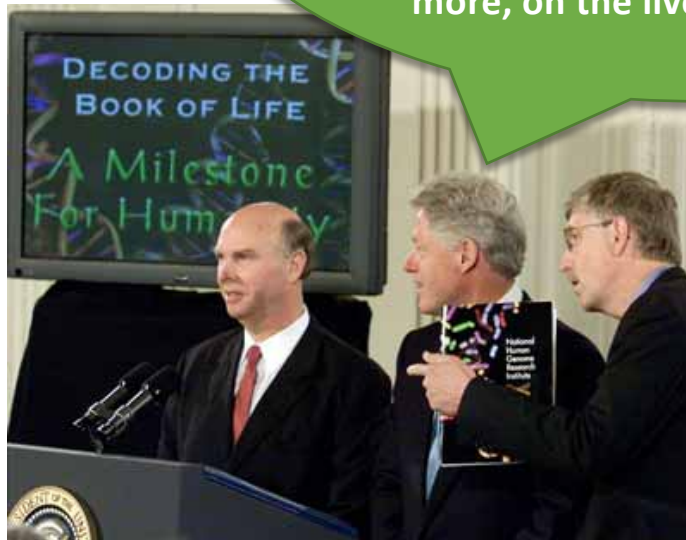
Figure 8.3 Human Molecular Genetics, 4ed. (© Garland Science)

1990 - 2003 : International Human Genome Project

Today we are learning the language in which God created life.

...

[G]enome science will have a real impact on all our lives – and, even more, on the lives of our children



1990 - 2003 : International Human Genome Project



3.000.000.000 bp @ \$ 2.700.000.000

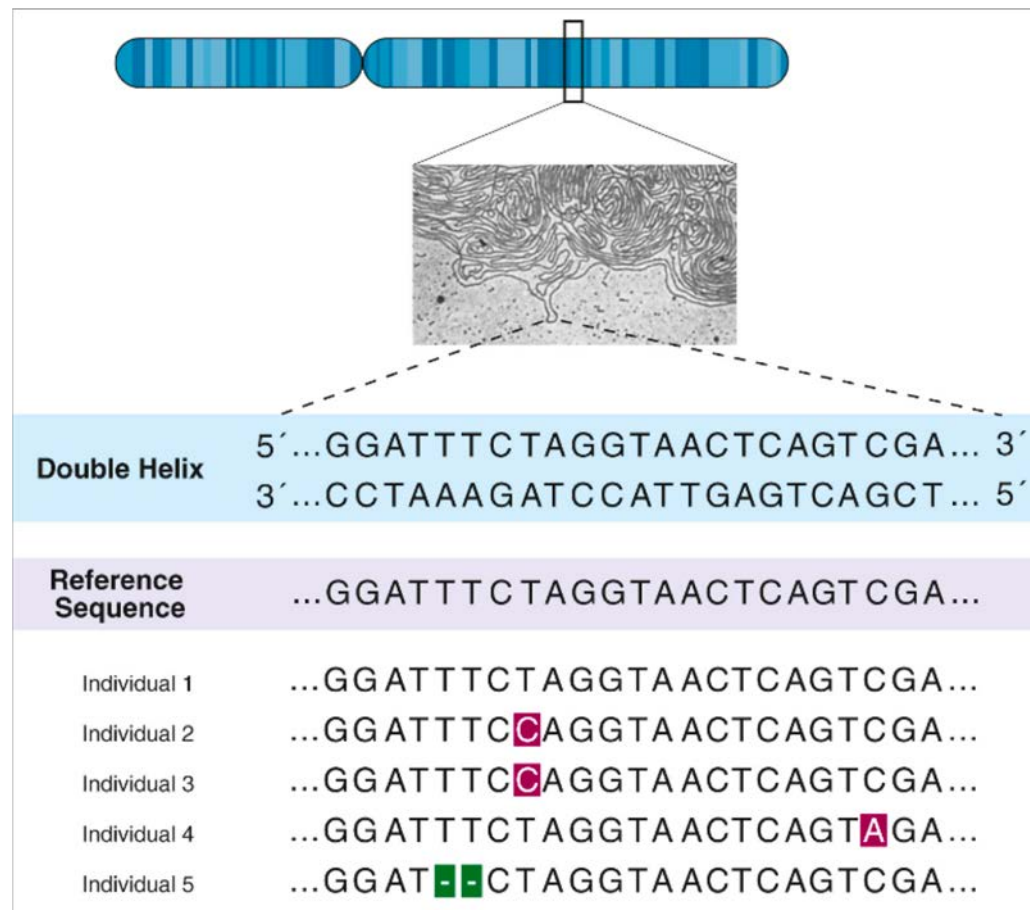
(most advanced appliances: 96 samples, 30.000-60.000 bases per run)

Human reference genome (n>1)



Reading is not understanding

By convention, only sequences from one strand of DNA are presented



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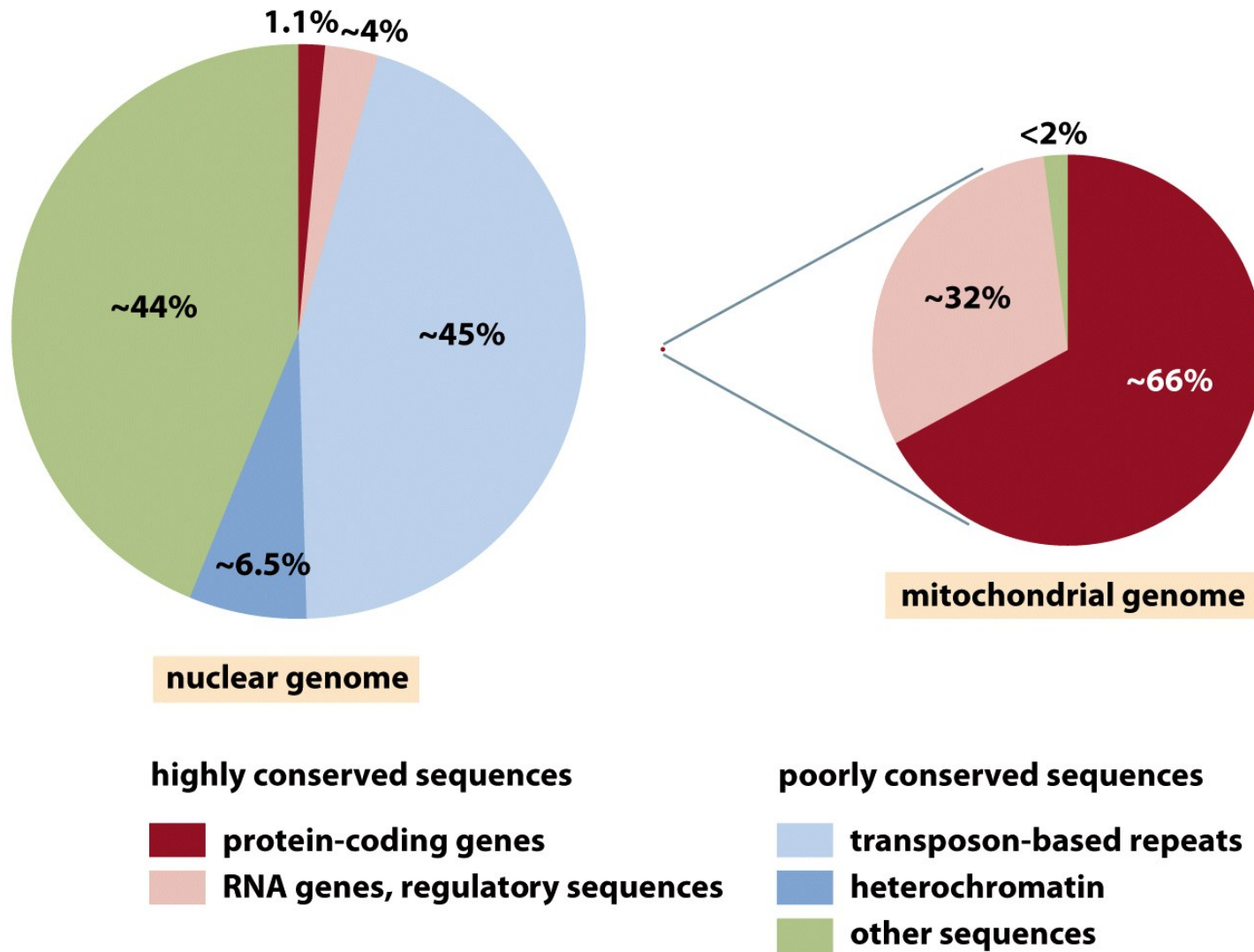
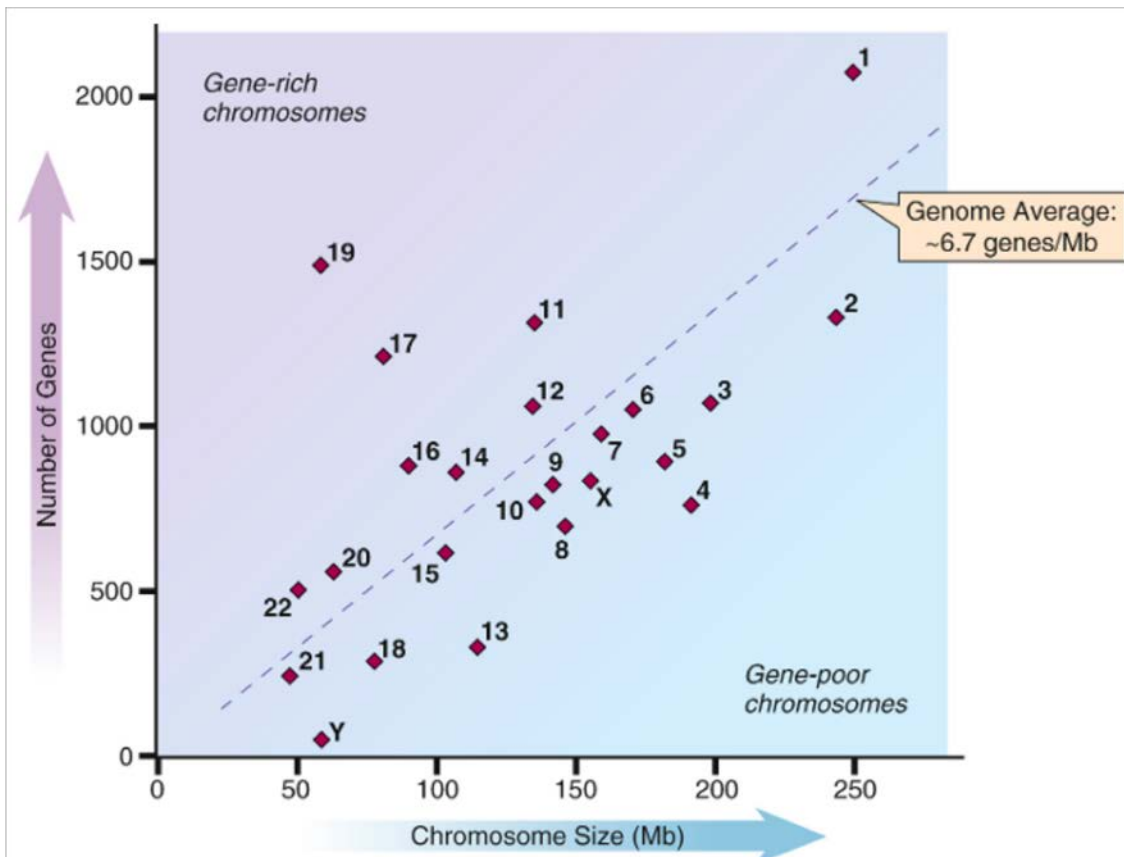
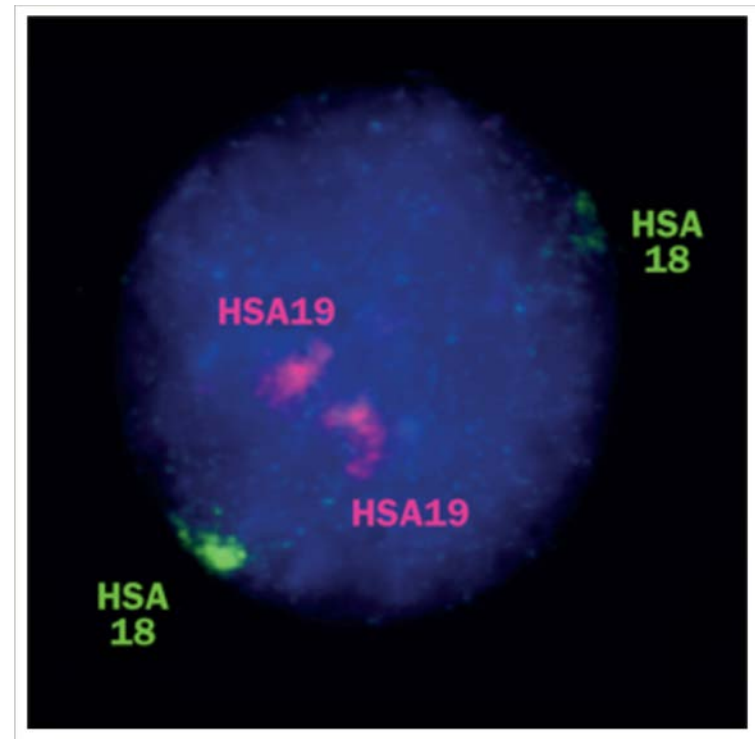


Figure 9.1 Human Molecular Genetics, 4ed. (© Garland Science)

Size and gene content of the 24 human chromosomes



Genetics in Medicine, 8th Edition (Thompson & Thompson)



Human Molecular Genetics, 4th Edition (Tom Strachan & Andrew Read)

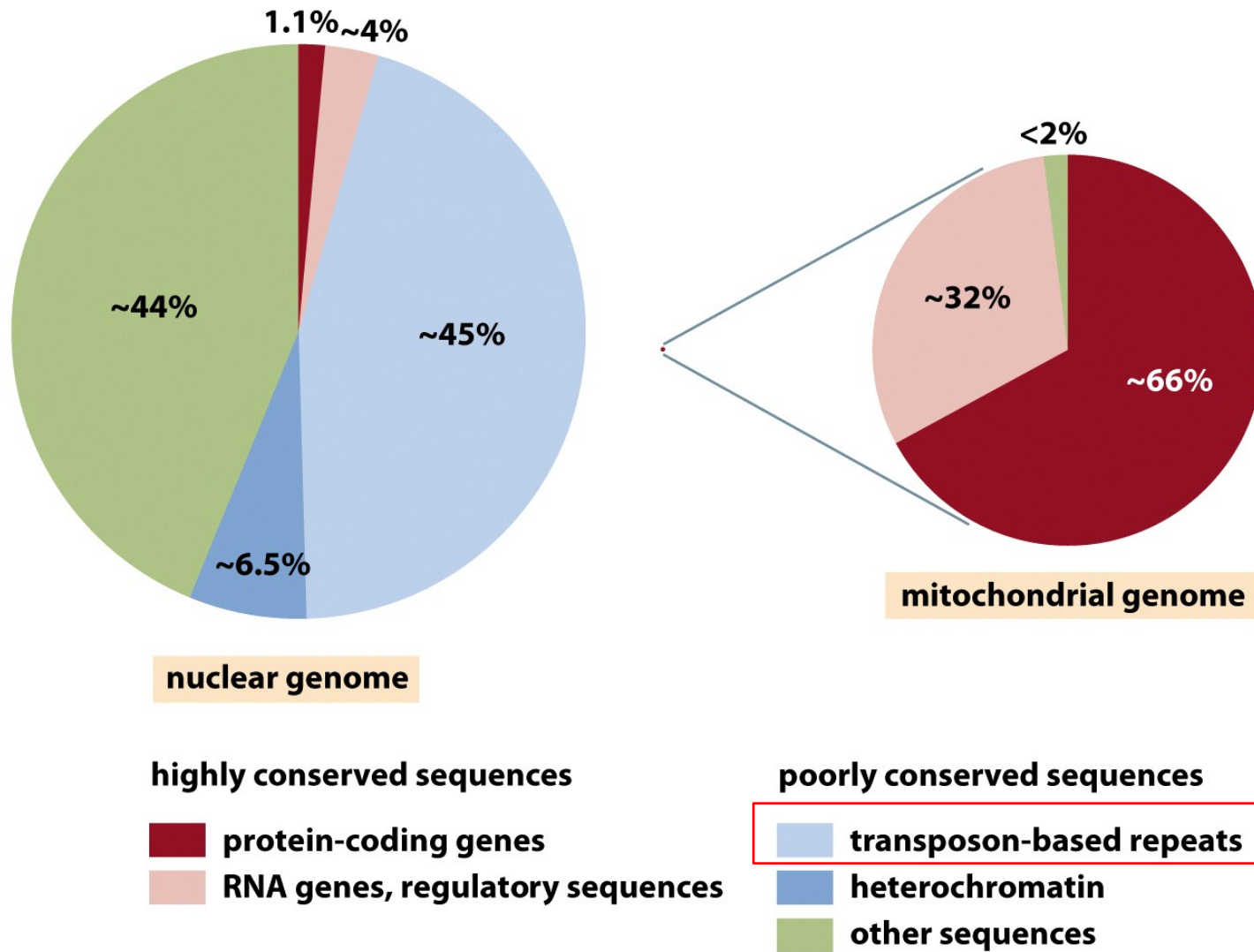
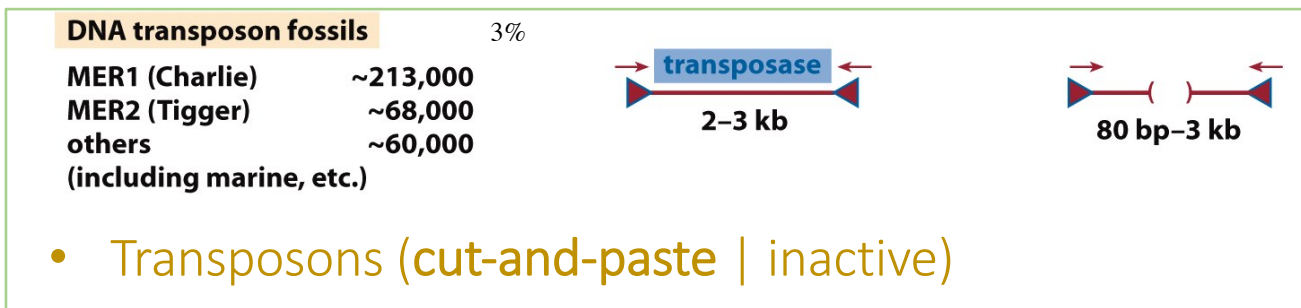
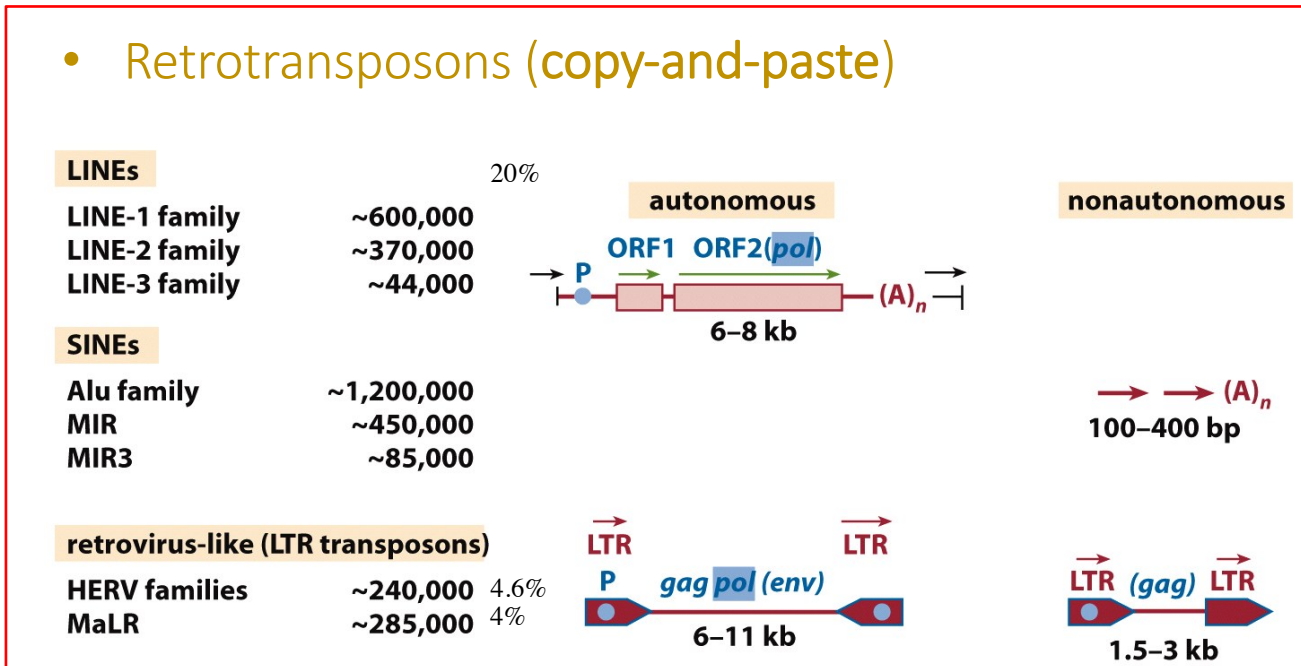


Figure 9.1 Human Molecular Genetics, 4ed. (© Garland Science)

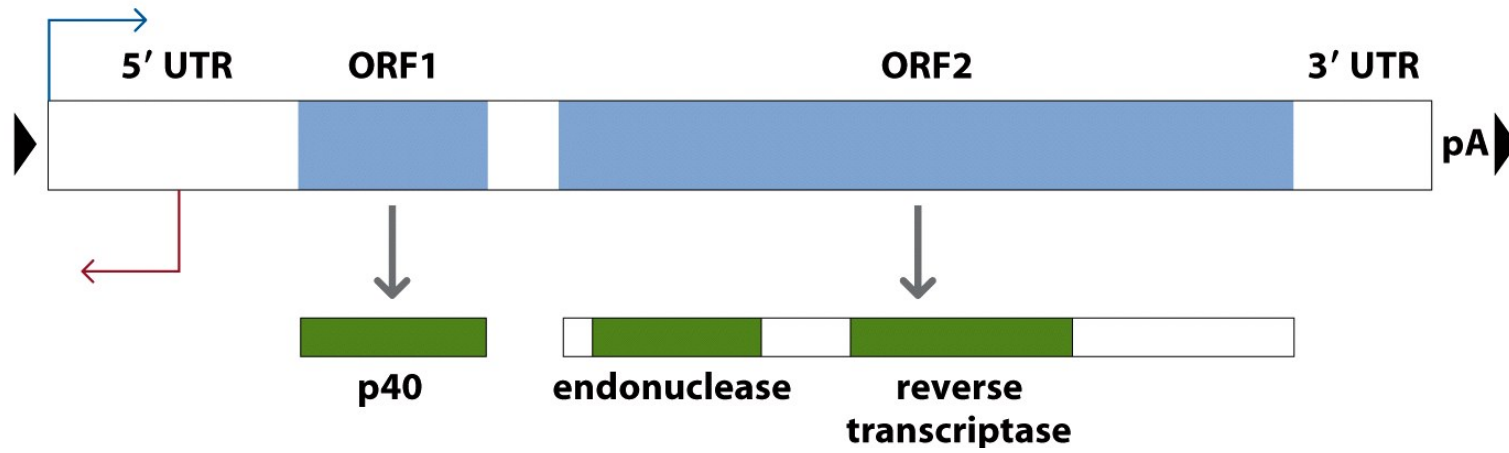
Transposons

- Retrotransposons (copy-and-paste)

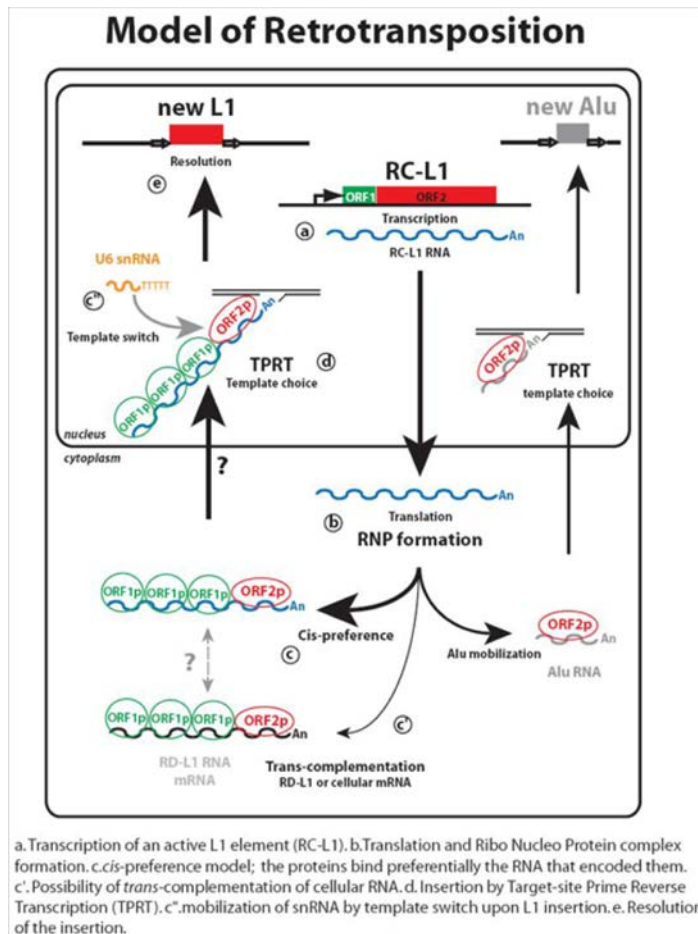


- Transposons (cut-and-paste | inactive)

LINE-1 repeat element



Model of L1- / Alu-retrotransposition



Integration in TTTT|A (preference for AT-rich regions)

1/100 are full lengths (genome-wide average = 900bp)

80-100 full-lengths L1 (n=6000) active

Gene mutation by L1-insertion

LINE-1 products used for retrotransposition of SINEs, mRNAs (-> processed pseudogenes) and retrogenes

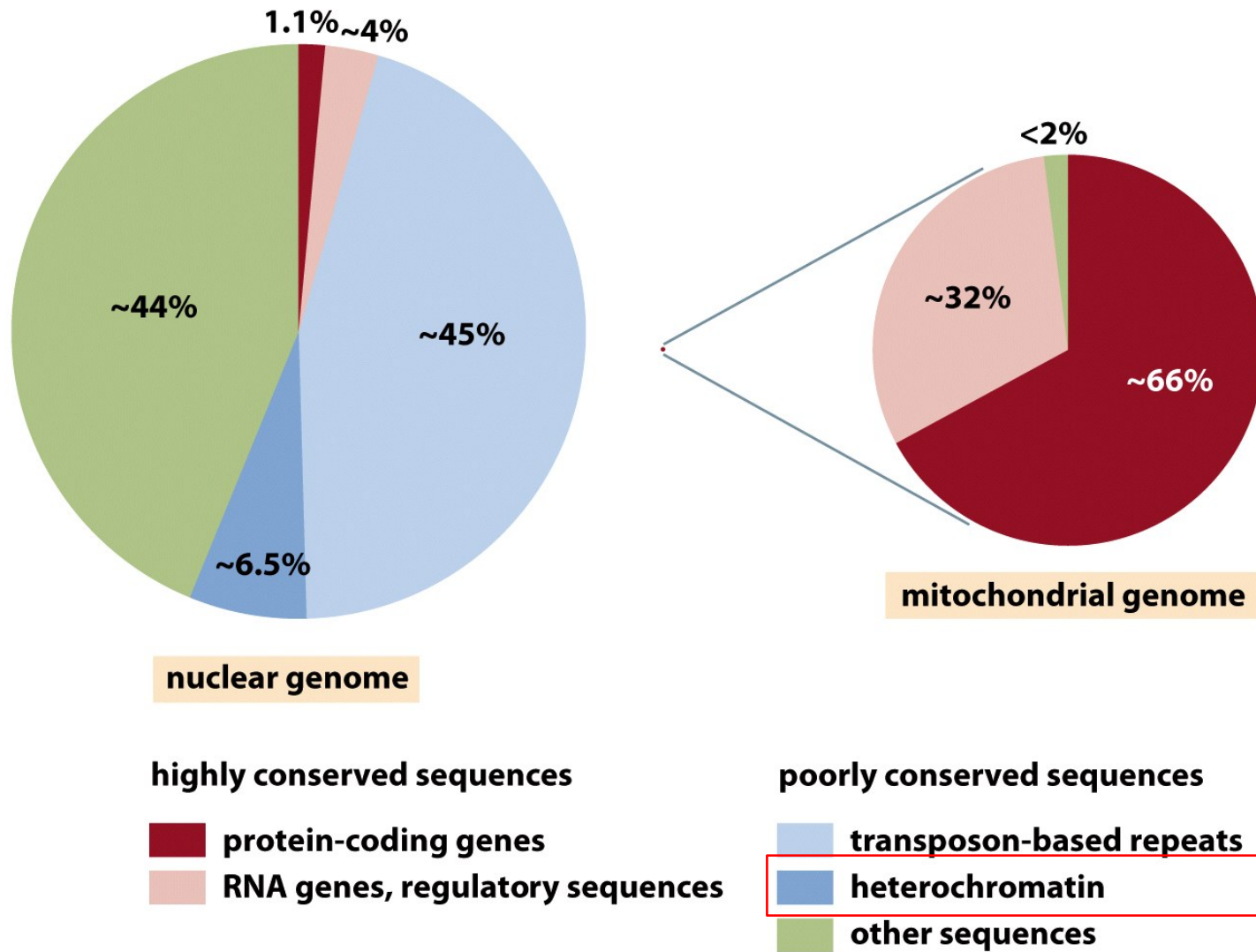
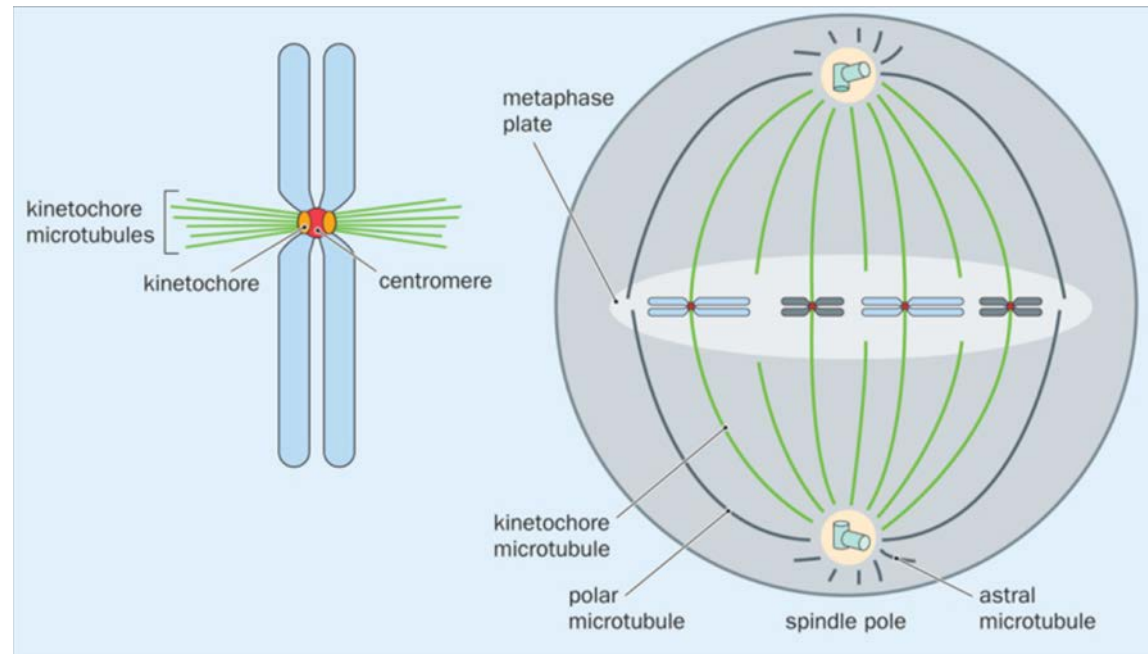
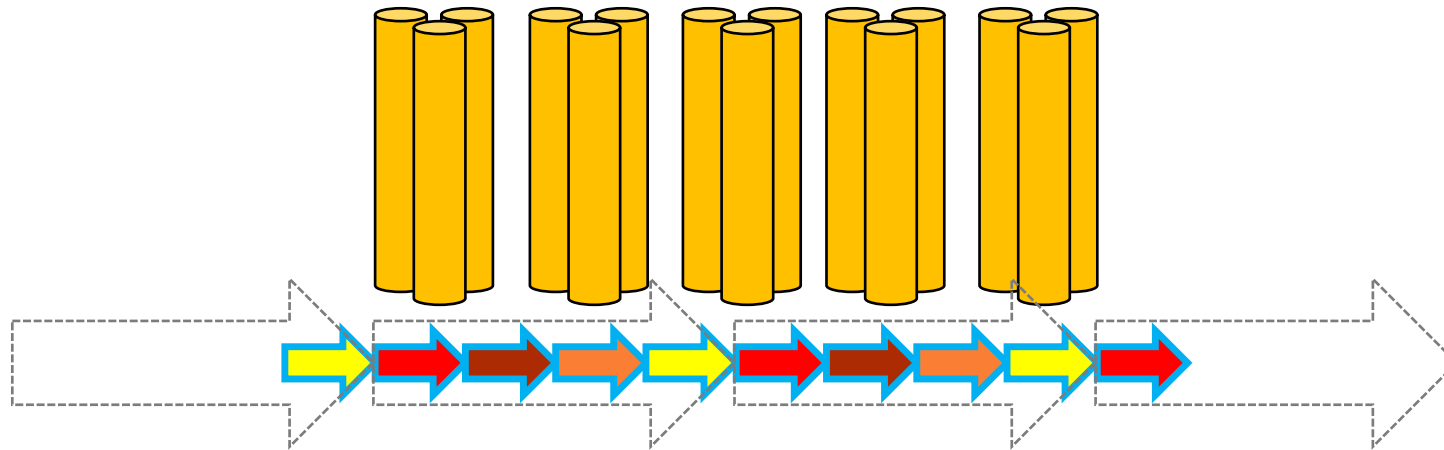


Figure 9.1 Human Molecular Genetics, 4ed. (© Garland Science)

Centromeric and telomeric DNA



Centromeric DNA:
Alpha-satellite or alphoid DNA at normal human chromosomes

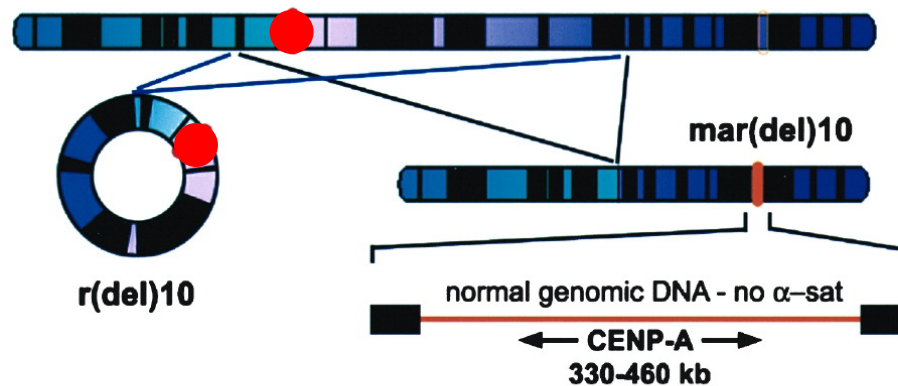
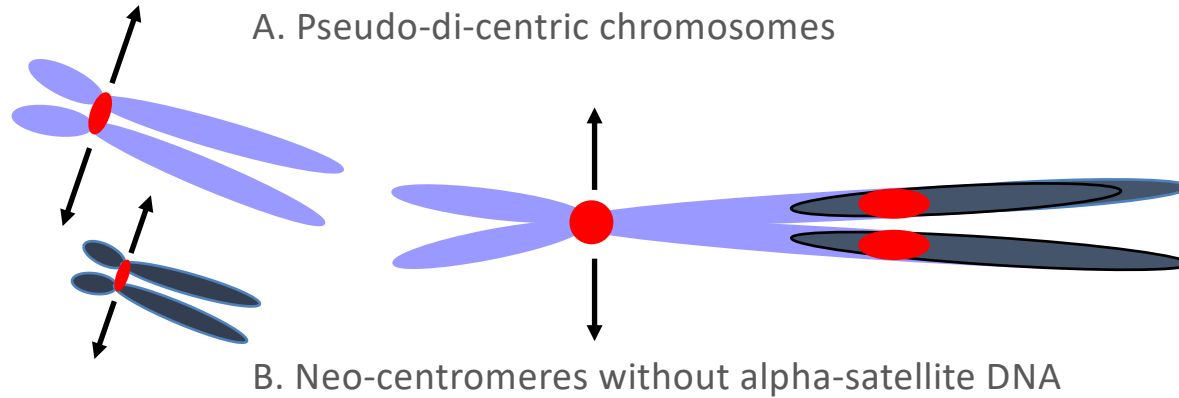


One alphoid higher order repeat can be :

- specific for one chromosome
- occurring on different chromosomes

Different alphoid higher order repeats can be co-existing on the same chromosome

Alpha-satellite DNA is not sufficient nor necessary for centromere function



➡ Centromere function is epigenetically regulated

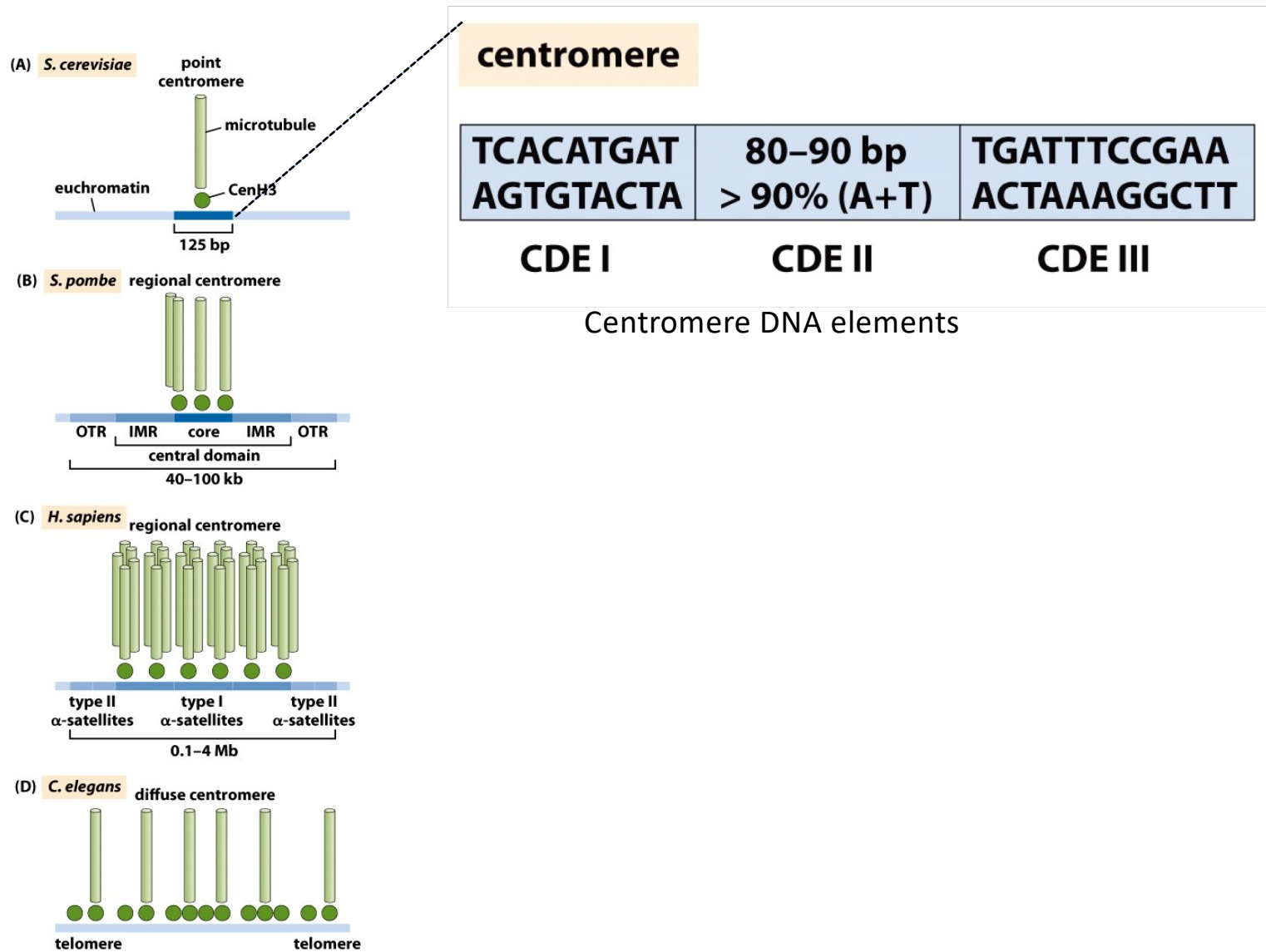
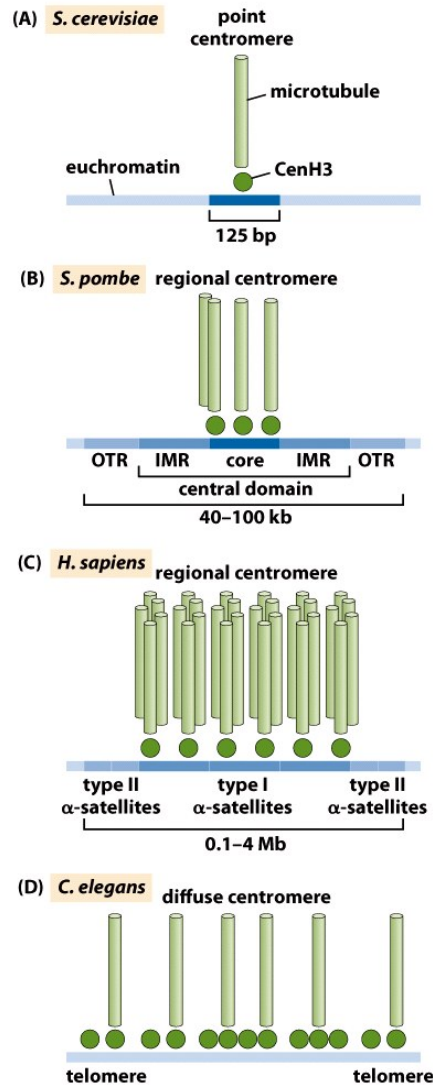


Figure 2.11 Human Molecular Genetics, 4ed. (© Garland Science)

Histone H3 variant CENP-A is the best candidate to carry the epigenetic centromere mark



(c) *H. sapiens*

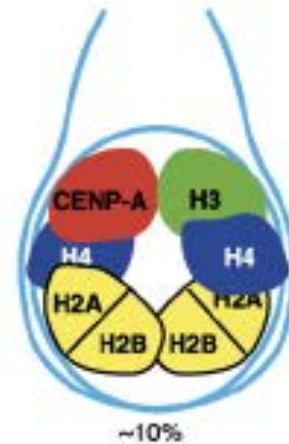
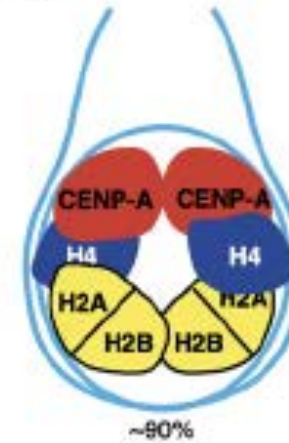
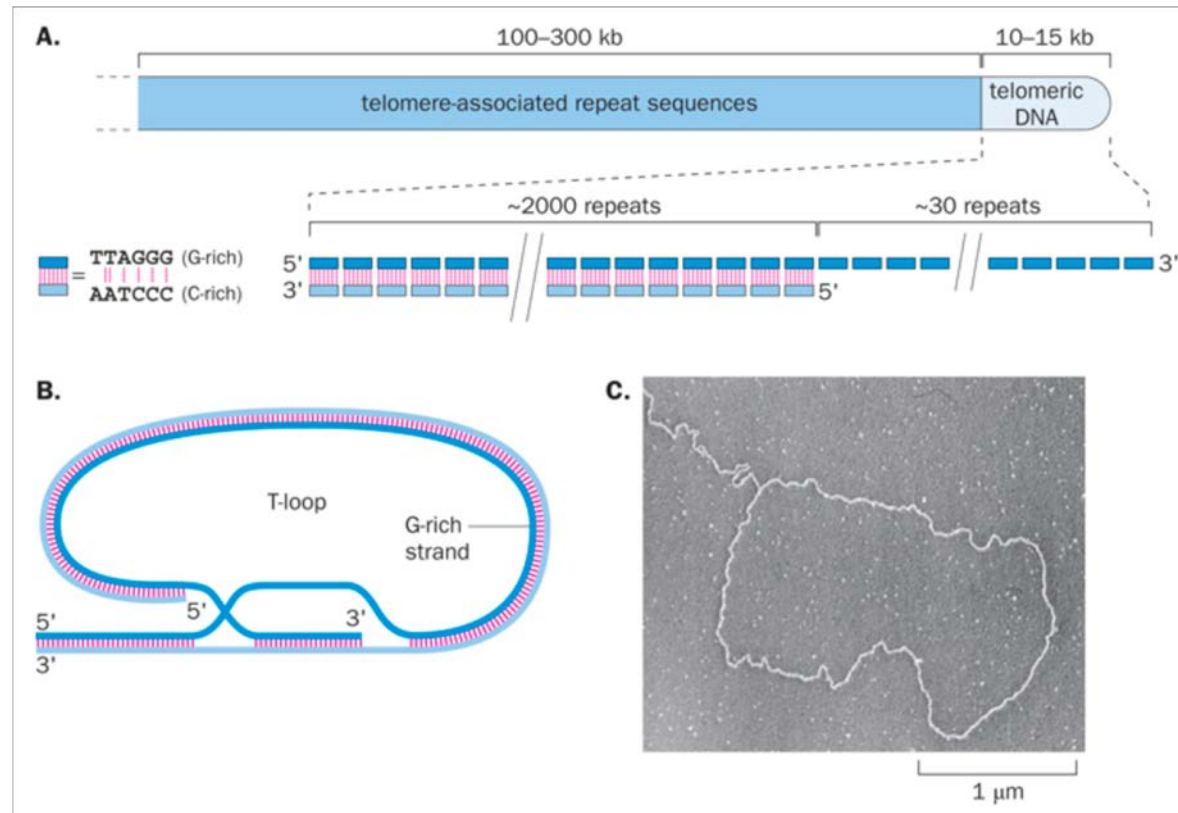


Figure 2.11 Human Molecular Genetics, 4ed. (© Garland Science)

Telomeric DNA

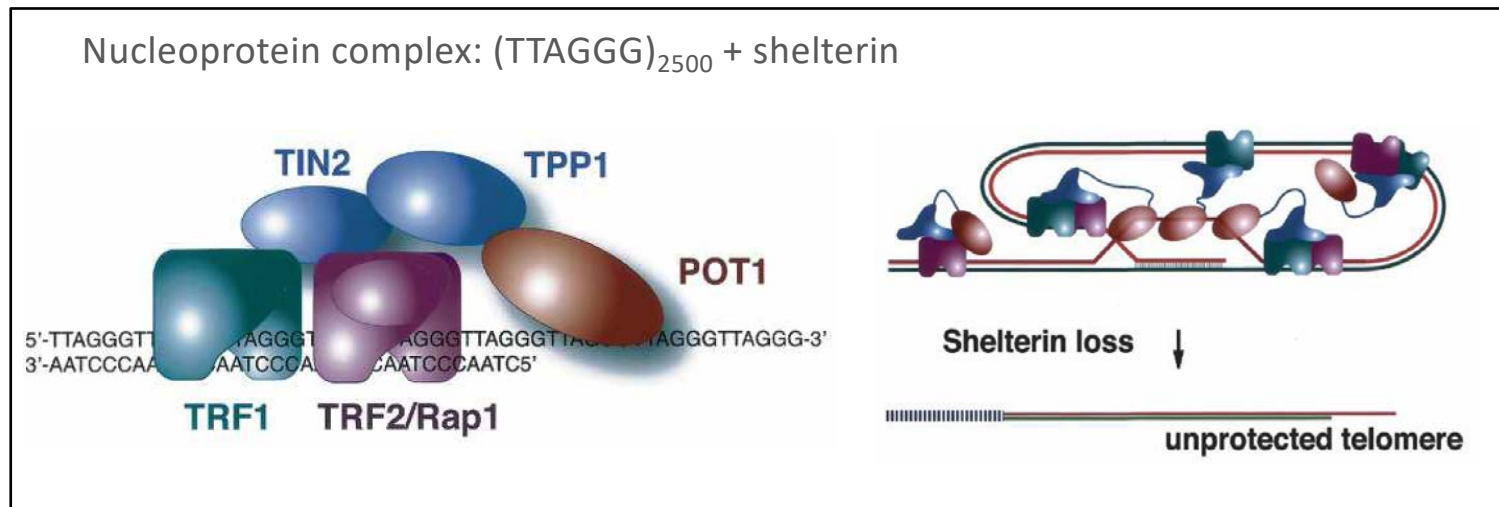


Telomeres:

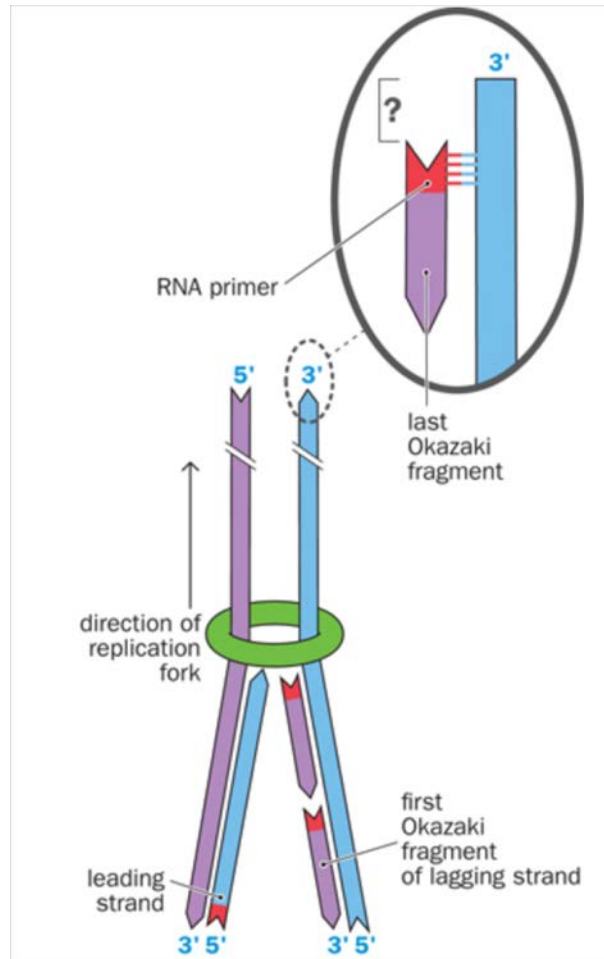
- important for maintaining structural integrity (prevents instability of ends, degradation and fusion with ends of broken chromosomes)
- required for complete replication of chromosome end
- in some cells: interaction with nuclear envelope for positioning of chromosomes in the nucleus

Telomeric DNA

Shelterin - Telosome

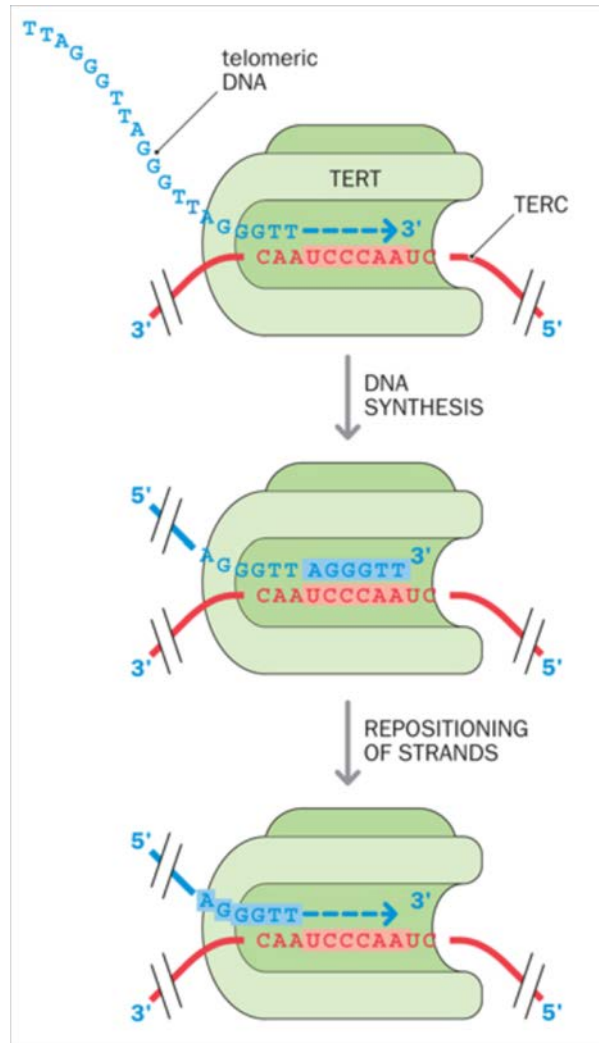


The 'chromosome end-replication' problem: incomplete synthesis



DNA synthesis starts at free 3'-OH
of an RNA primer


Telomerase uses a reverse transcriptase and a non-coding RNA template to make new telomeric DNA repeats



Telomerase:

- Germ cells
- Embryonic cells
- Stem cells
- Cancer cells

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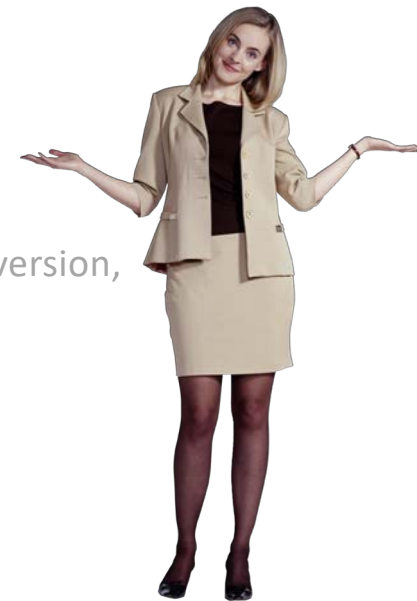
Humane genetic variation

The key to:


- Understanding differences between people
- Identifying genes / variants that play a role in disease and health



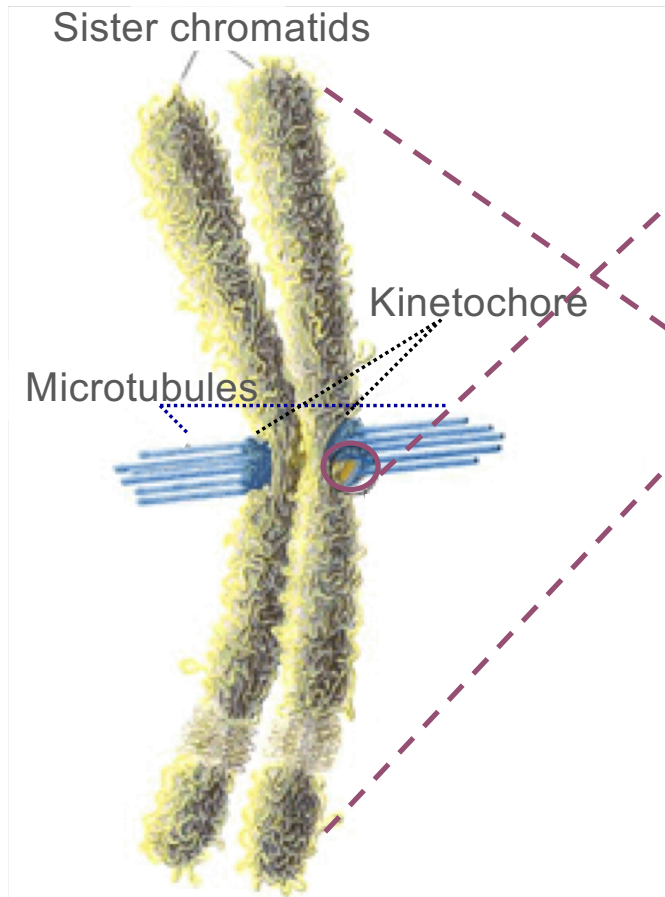
single nucleotide variants
small insertions and deletions
copy number variation
structural variation
(Deletion, Duplication, Amplification, Translocation, Inversion,
Retrotransposition)



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Chromosomes ensure transport and integrity of genetic information



Functional domains

Centromere

- correct segregation (capture microtubules)
- chromosome movements

Telomeres

- protect against degradation, fusion and recombination
- complete end replication
- chromosome movements
- subtelomeric gene expression

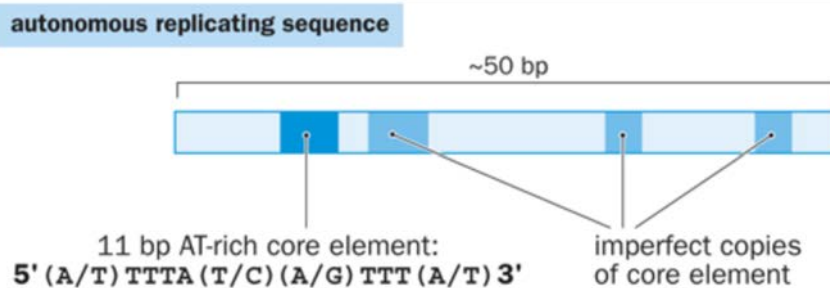
Origins of replication

- replication of the genetic information once per cell cycle

Origins of replication

DNA sequence in cis where proteins bind in preparation for DNA replication

S. cerevisiae



ARS provides a binding site for:
ORC (multi-protein origin of replication complex) + TF

Mammalia

DNA is replicated from multiple initiation sites per chromosome, with an average of **one initiation site per 40-80 kb DNA**

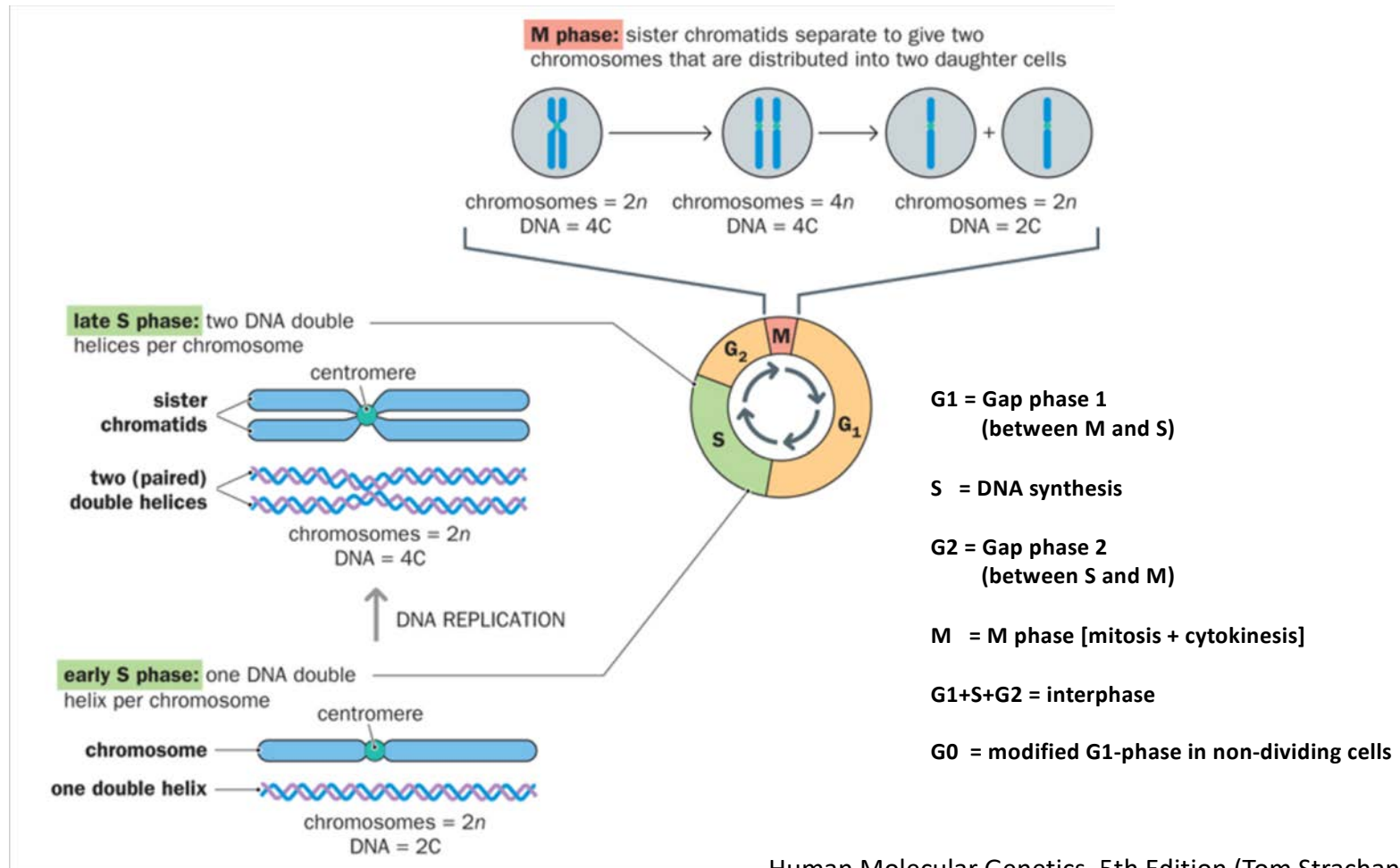
Structural motives can be important: probable replication origins often have guanine-rich DNA sequences with the potential to form G-quadruplexes, a four-stranded DNA structure with Hoogsteen binding between the guanines

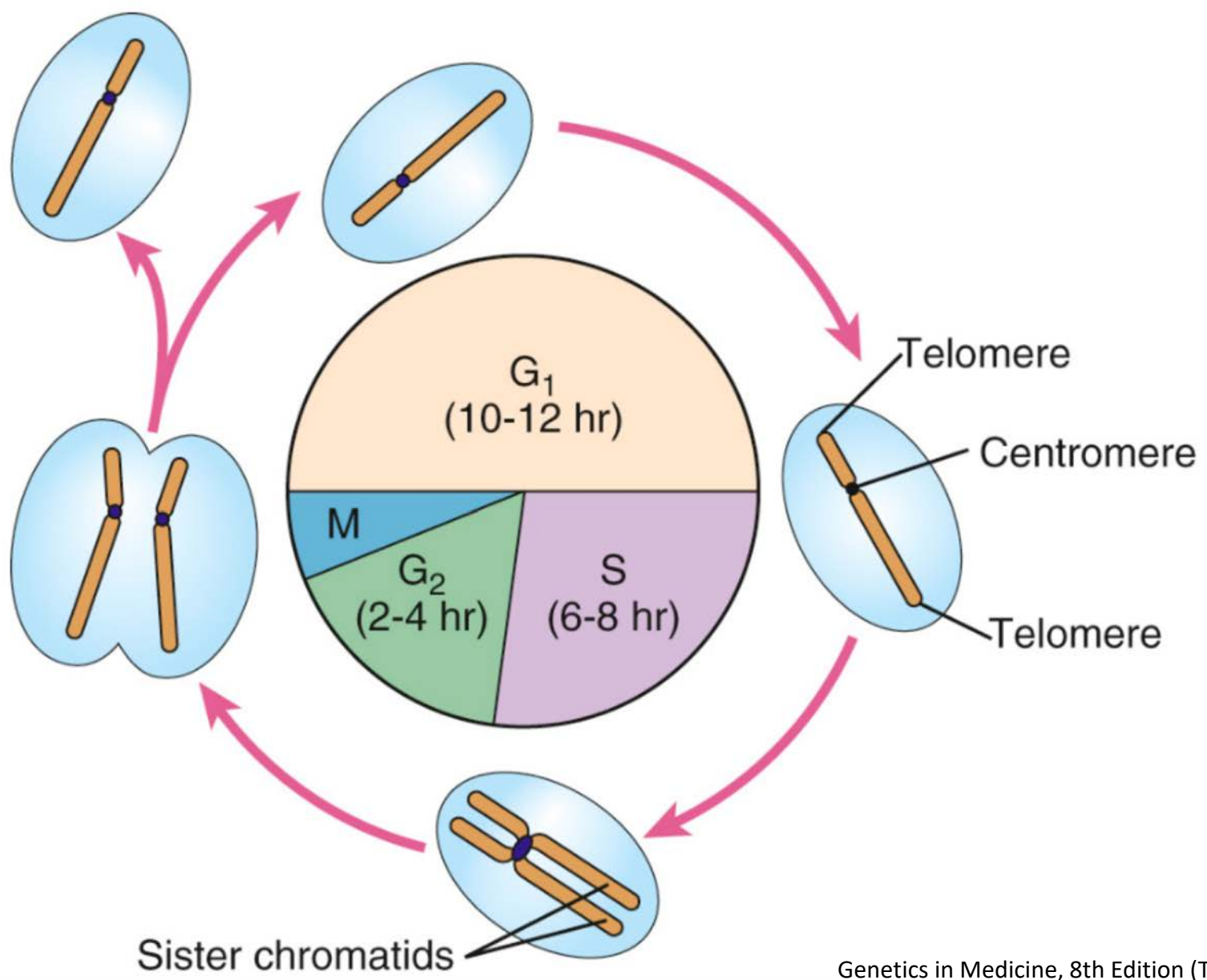
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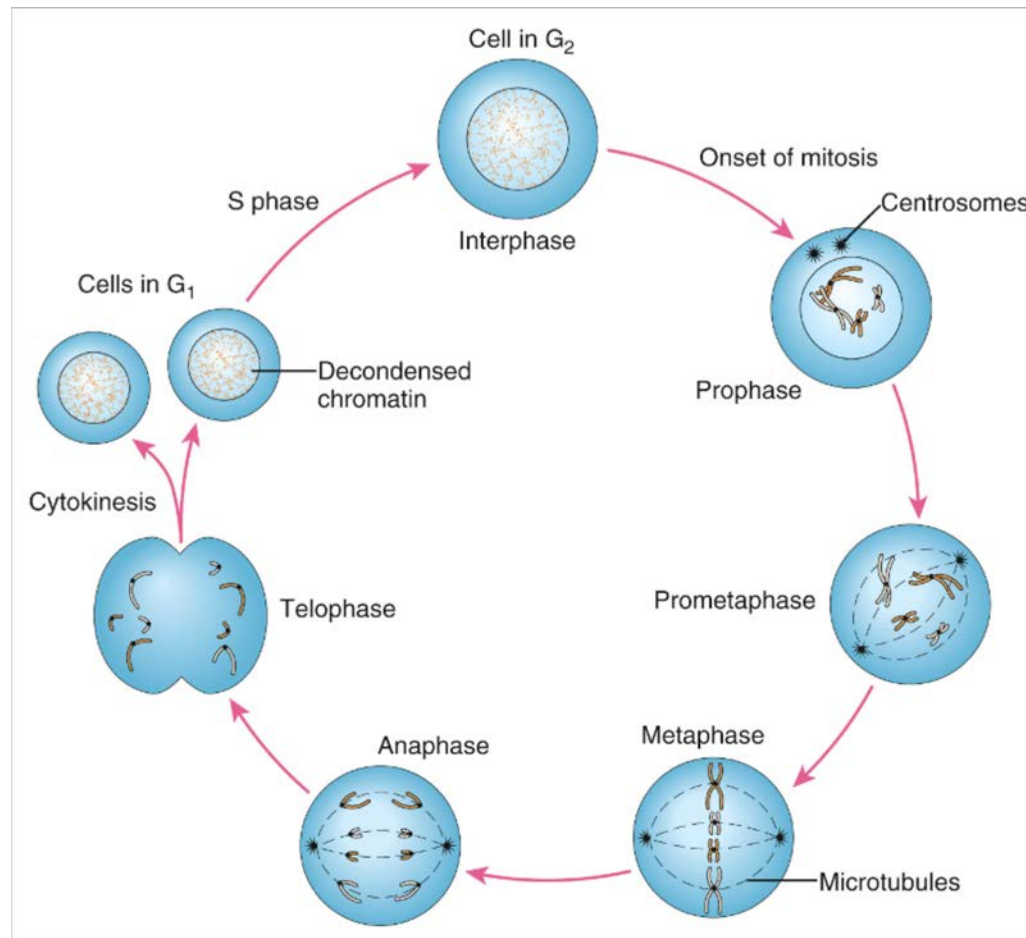


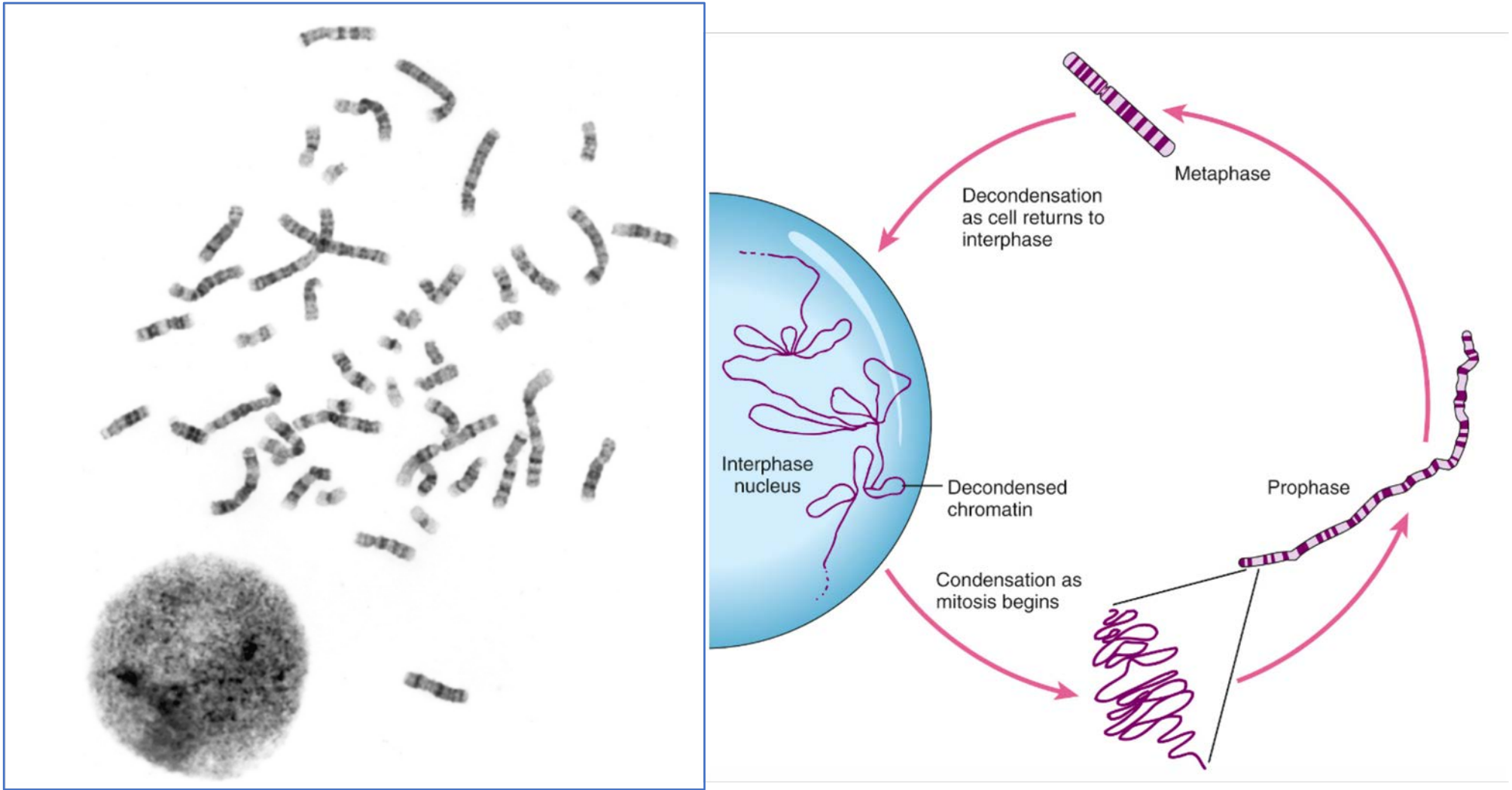
Doubling the number of chromosomes and the DNA content prior to mitosis during the cell cycle



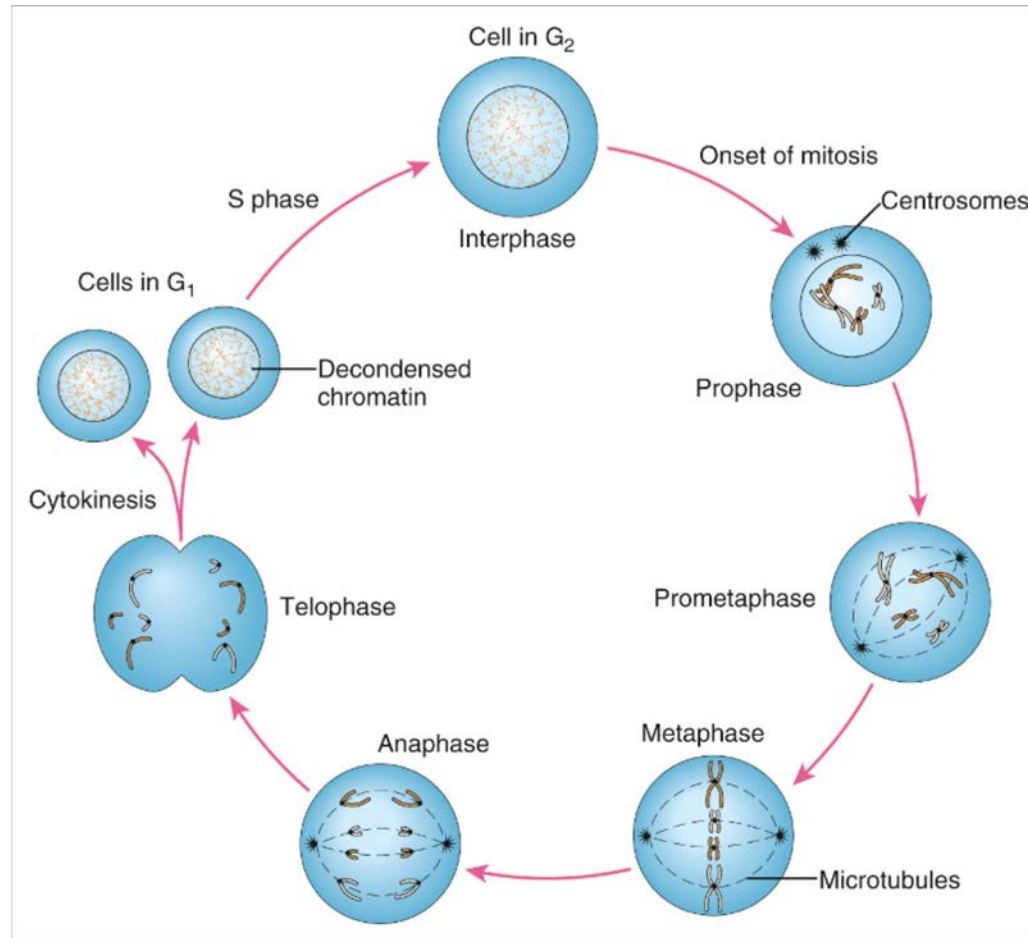


Mitosis (nuclear division) and cytokinesis (cell division)



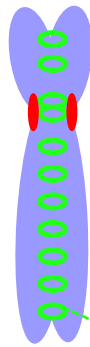


Mitosis (nuclear division) and cytokinesis (cell division)

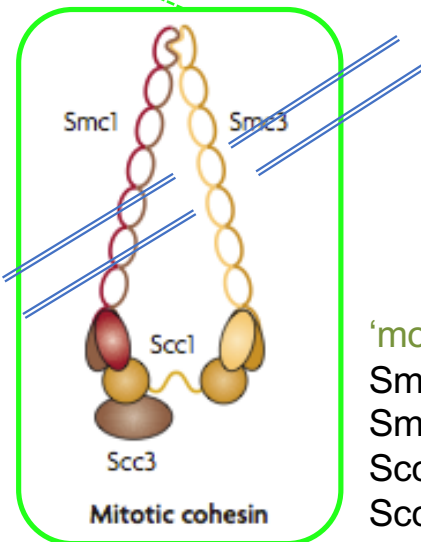


Molecular glue between replicated DNA-molecules = cohesin complex

Prophase



Sister chromatids
of a chromosome



'molecular glue' between replicated DNA-molecules (sister chromatids)

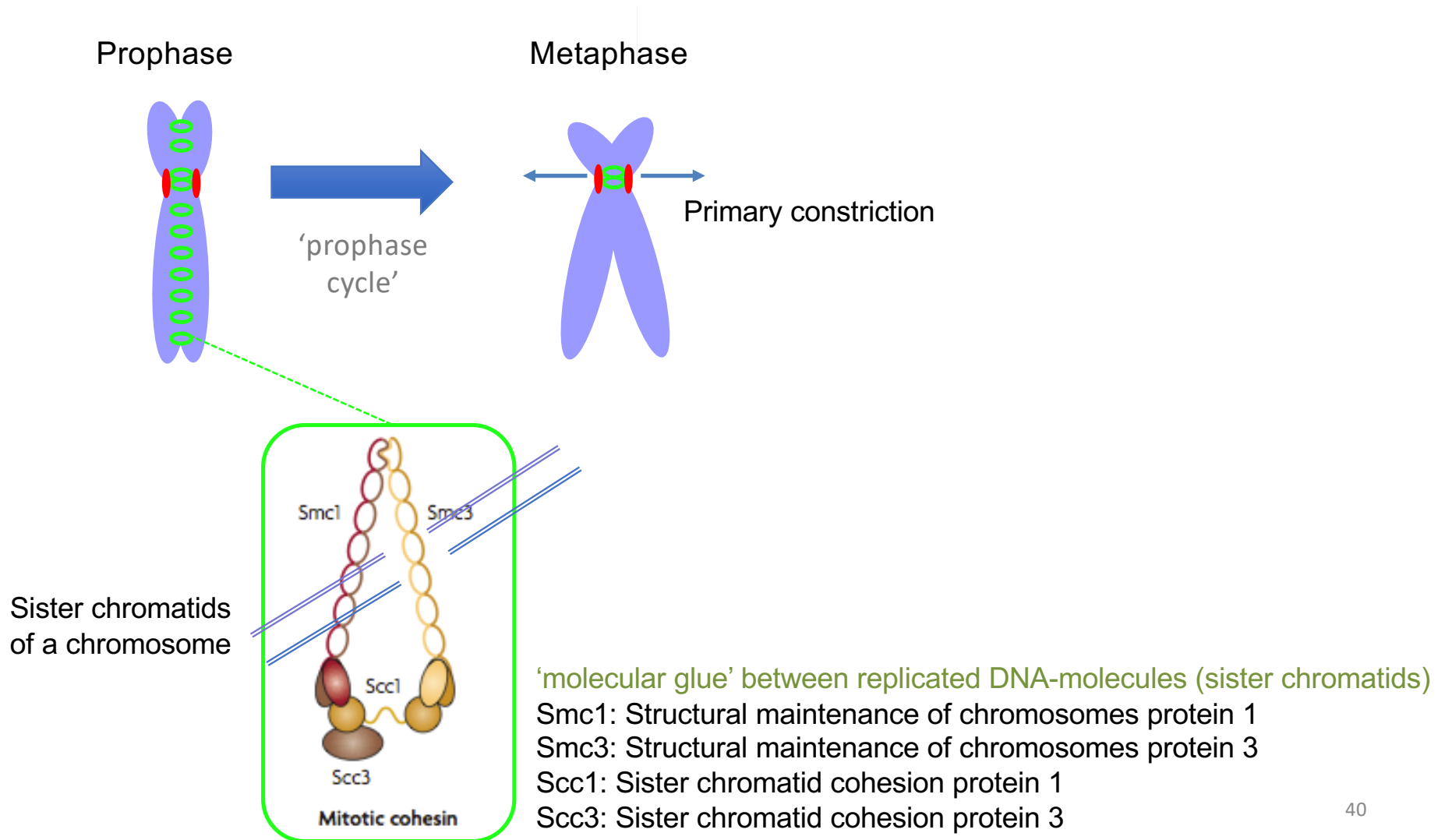
Smc1: Structural maintenance of chromosomes protein 1

Smc3: Structural maintenance of chromosomes protein 3

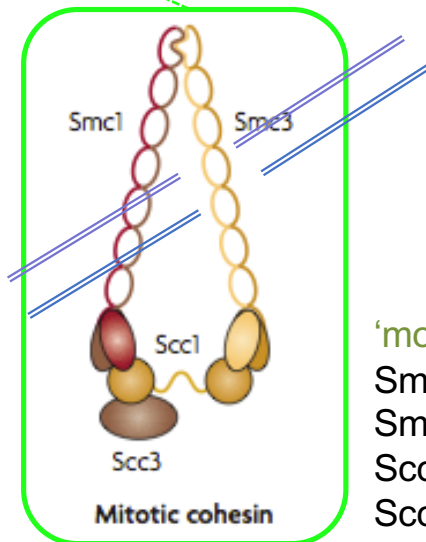
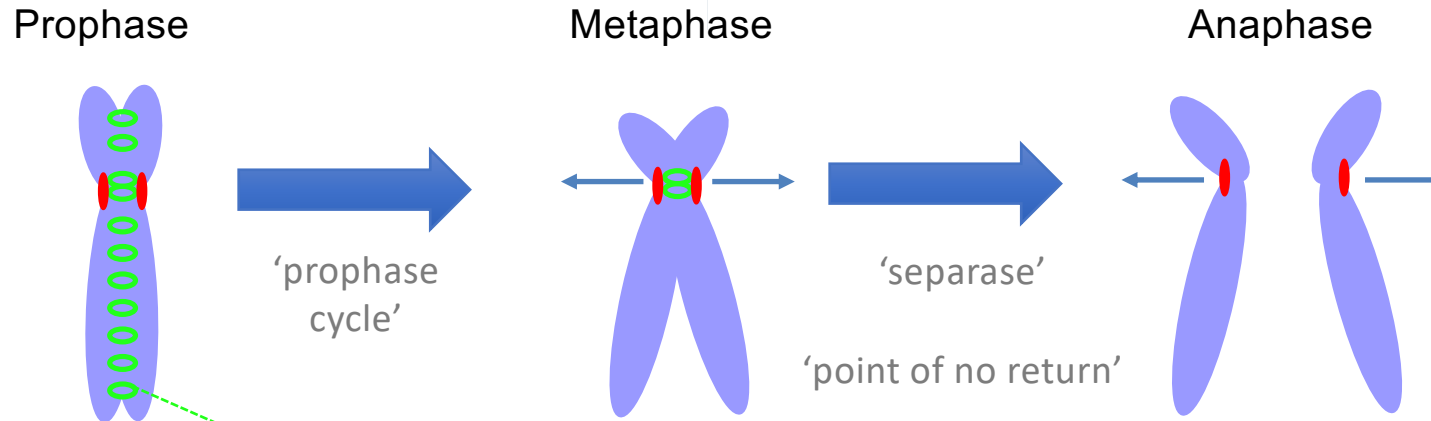
Scc1: Sister chromatid cohesion protein 1

Scc3: Sister chromatid cohesion protein 3

Loss of sister chromatid cohesion during mitosis



Loss of sister chromatid cohesion during mitosis



'molecular glue' between replicated DNA-molecules (sister chromatids)

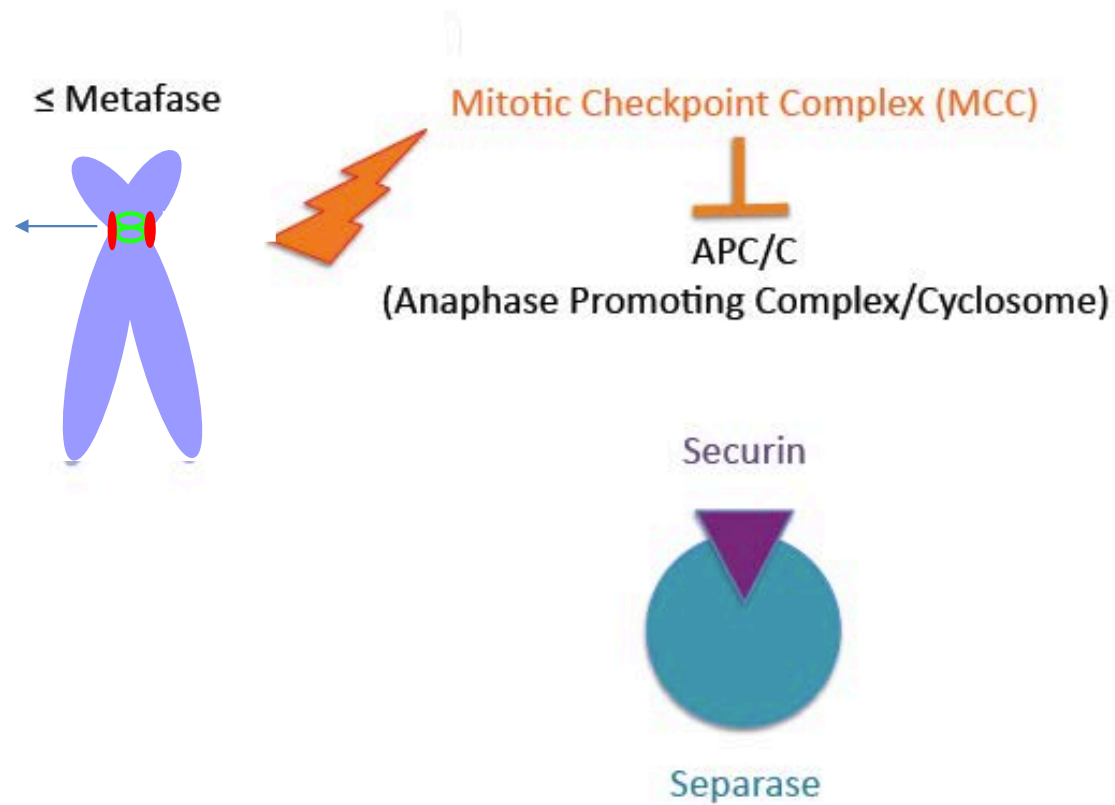
Smc1: Structural maintenance of chromosomes protein 1

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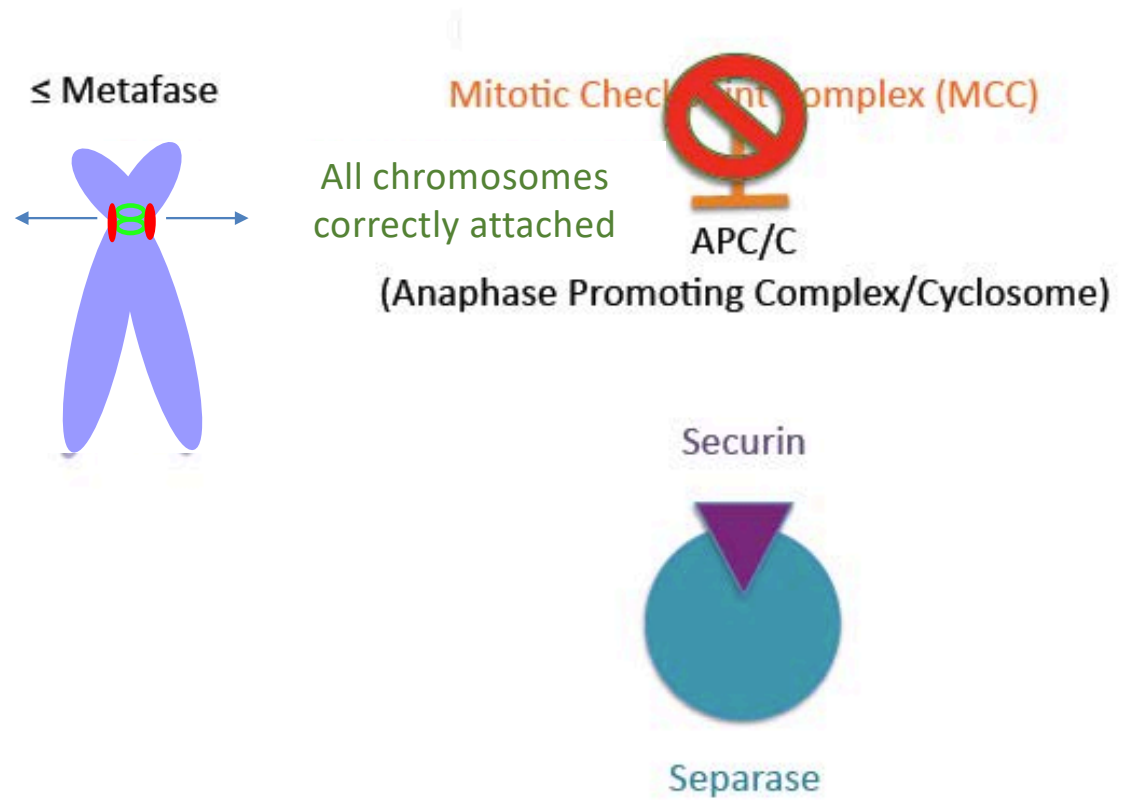
Scc1: Sister chromatid cohesion protein 1

Scc3: Sister chromatid cohesion protein 3

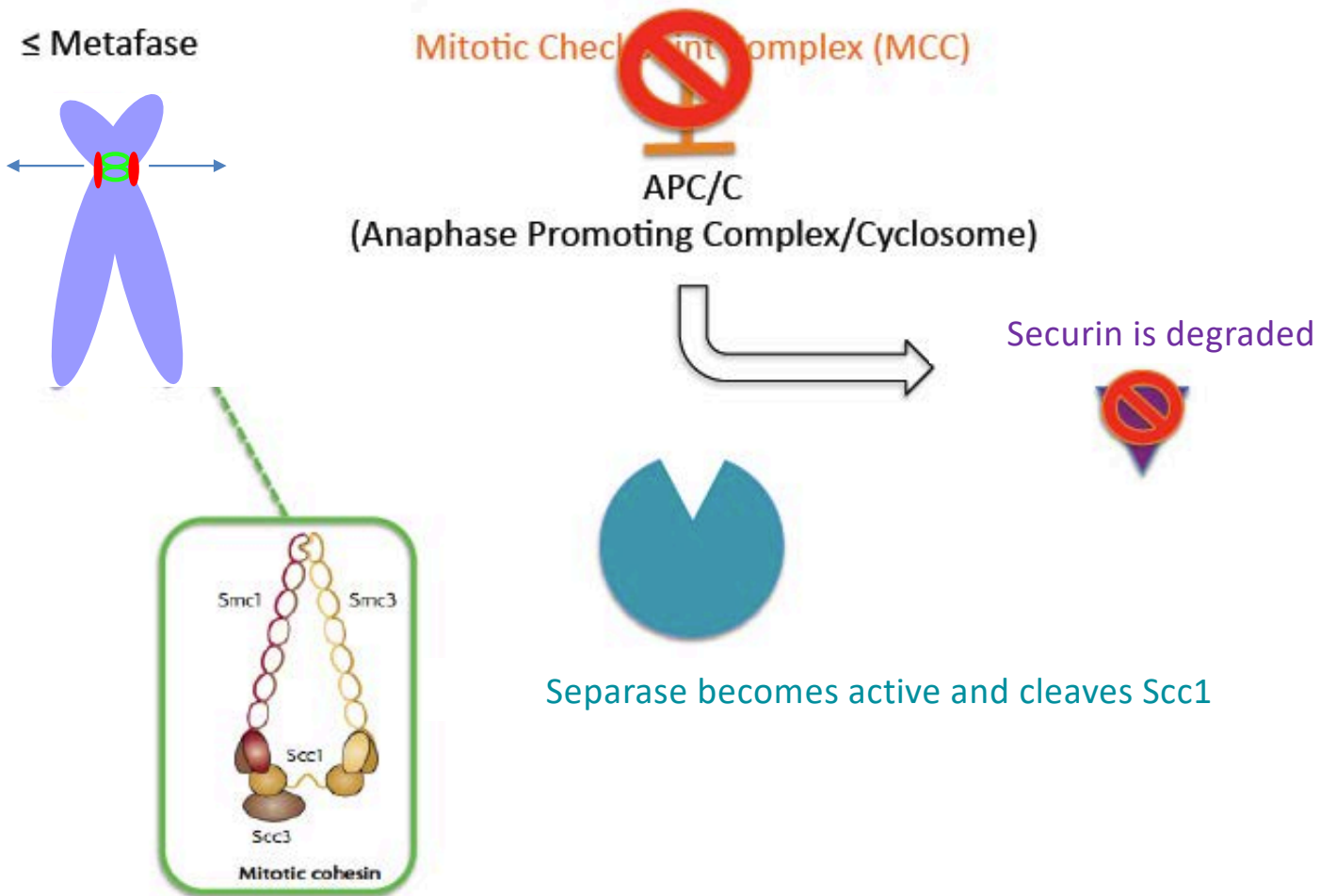
Spindle Assembly Checkpoint



Spindle Assembly Checkpoint

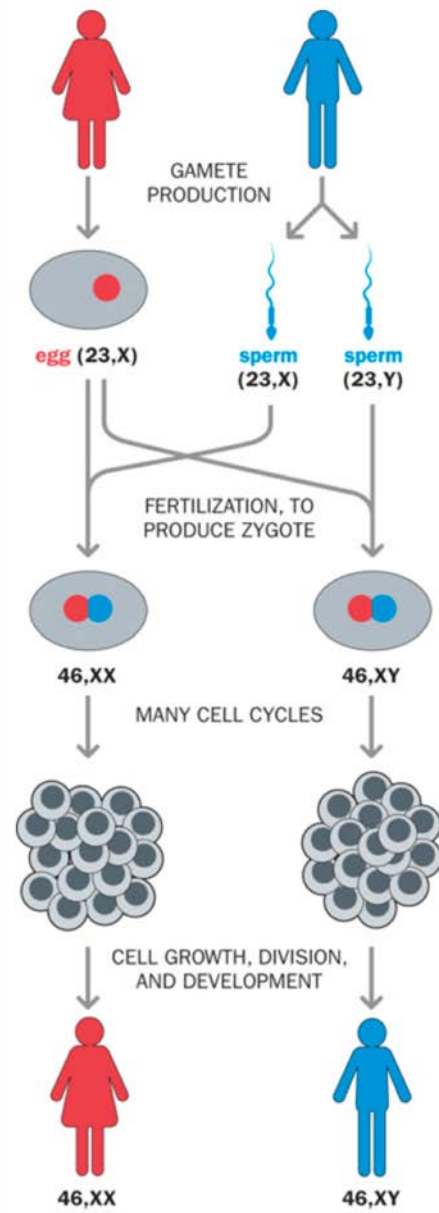


Spindle Assembly Checkpoint



Hoofdstuk 2

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The human life cycle, seen from a chromosomal angle

Meiosis:

1 diploid cell ($2n$)

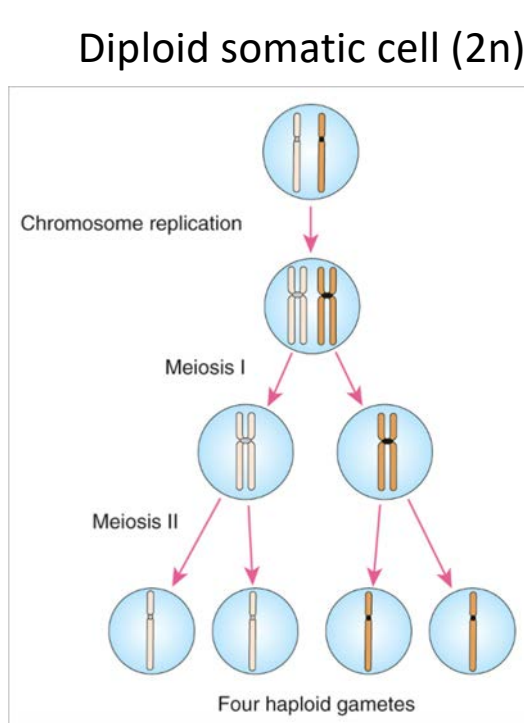
-> man: 4 haploid gametes ($1n$)

-> woman: 1 haploid gamete + polar bodies

Mitosis:

1 diploid cell ($2n$) -> 2 diploid daughter cells ($2n$)

Meiosis: 'to reduce'



Haploid gametes
4 x (n)

Chromosomes replicated once

Crossovers -> genetic diversity

-> correct homologue segregation

meiosis I: reductional division (disjunction)

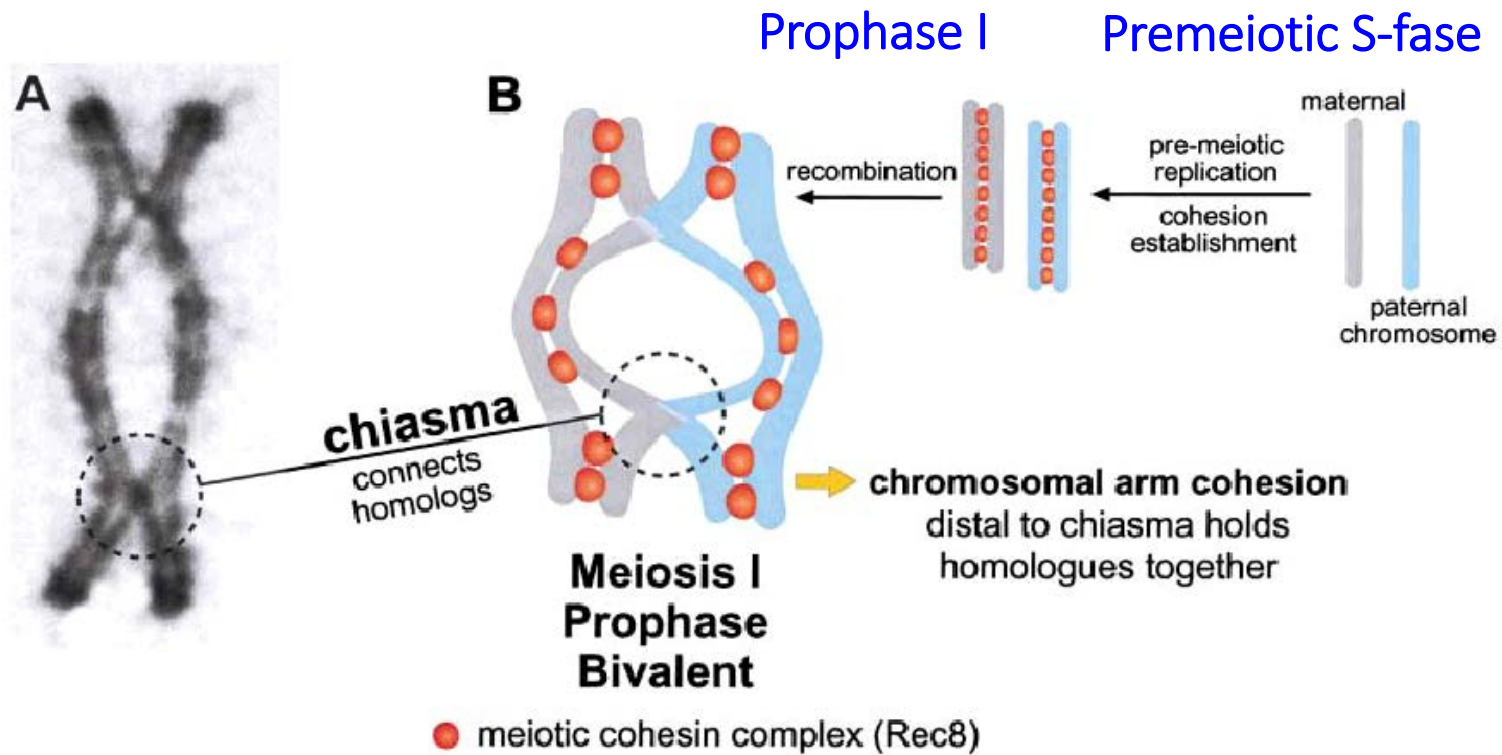
- separation of homologous autosomes

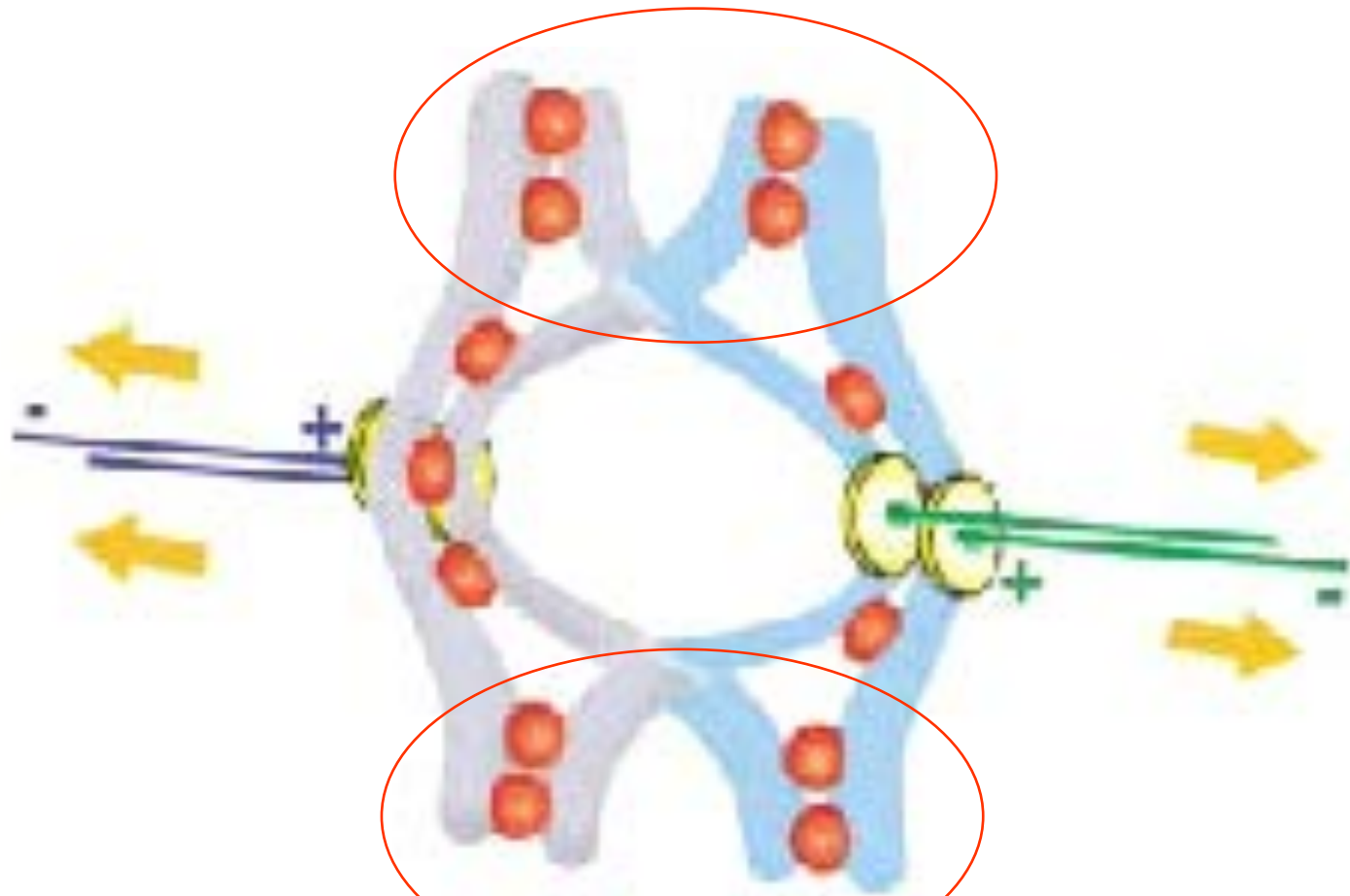
- separation of sex chromosomes

Random assortment

meiosis II: equational division

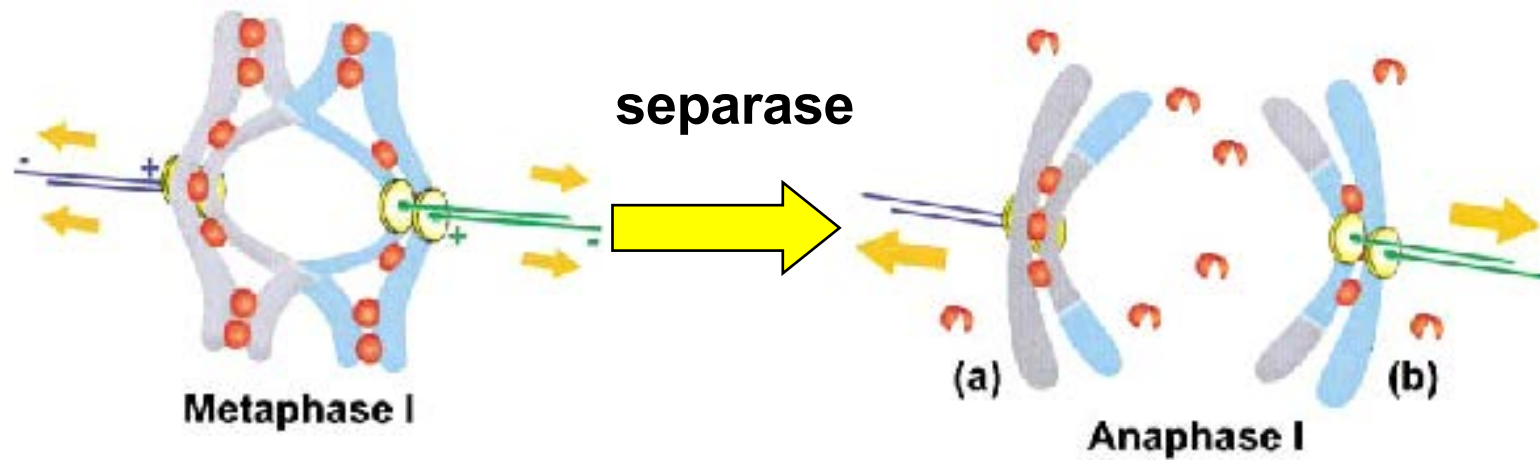
- separation of sister chromatids (~mitosis)





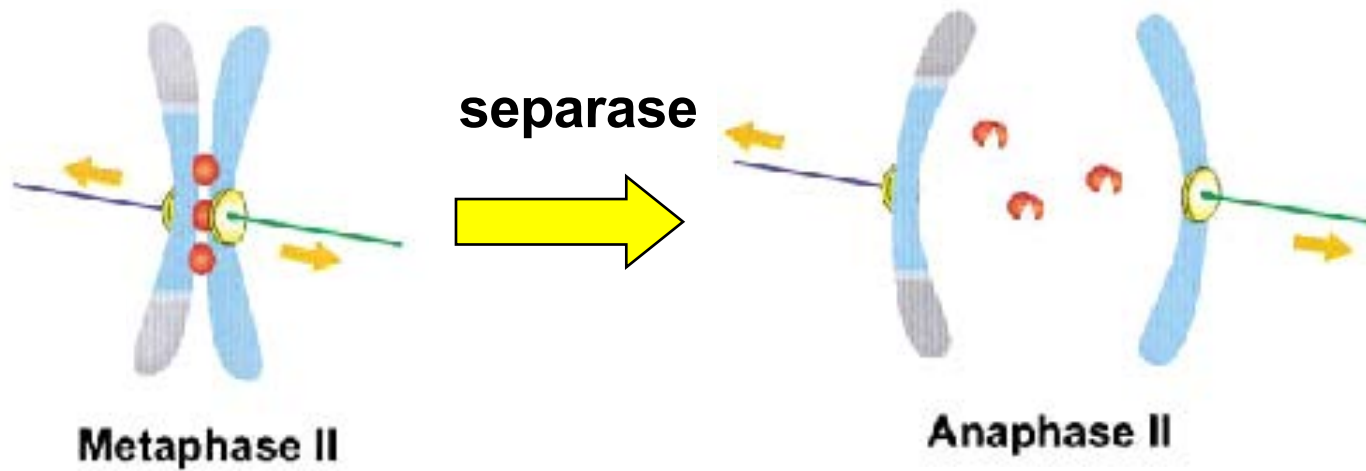
Metaphase I

First meiotic division



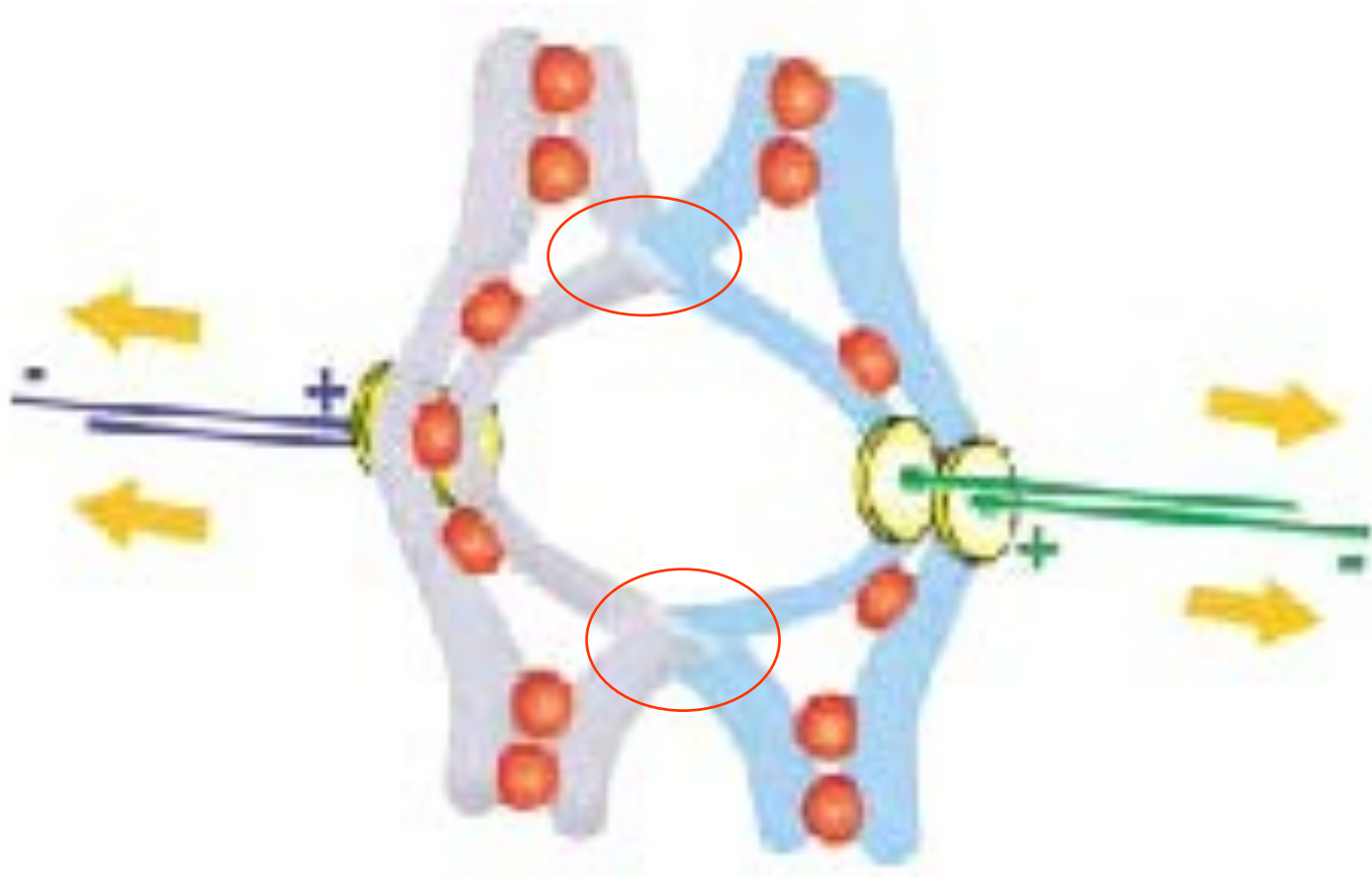
Cohesin rings at the chromosomal arms are opened, but remain intact at the centromeres

Second meiotic division

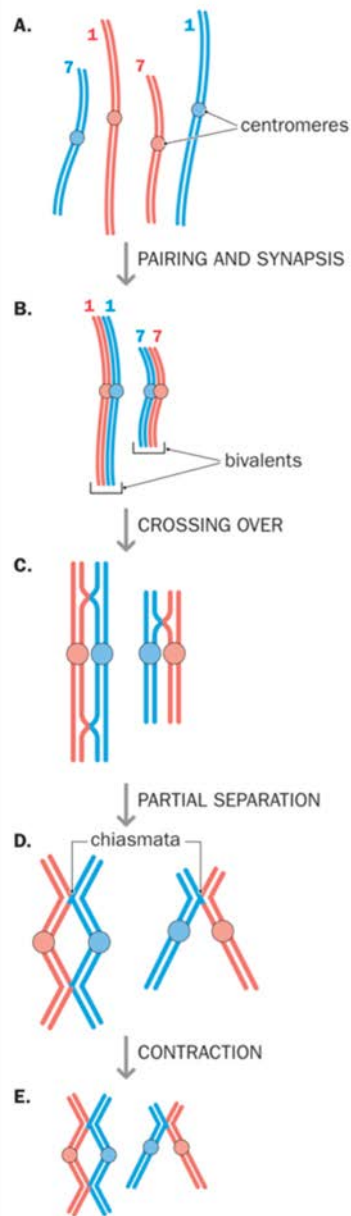


Cohesin rings at centromeres are opened

Formation of chiasma(ta) by homologous recombination



± 55 chiasmata per cell in human male meiosis
± 90 chiasmata per cell in human female meiosis



The five stages during prophase of meiosis I

A: leptotene (chr condensation, chr unpaired, dsDNA breaks [DSB])

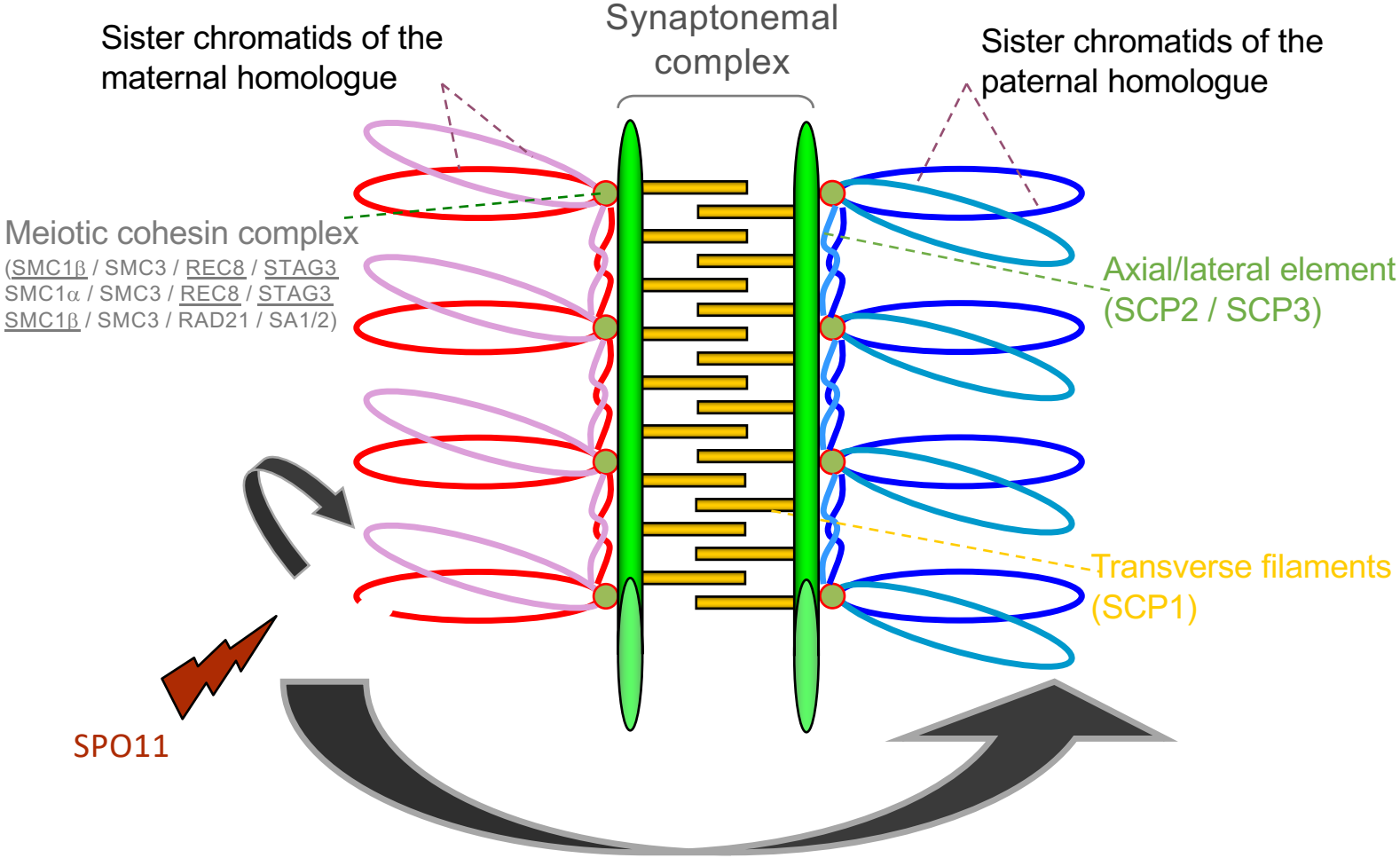
B: zygotene (repair of DSB in progress; pairing of homologues to bivalents; synapsis through synaptonemal complex)

C: pachytene (synapsis complete; crossing-over complete; formation of chiasmata)

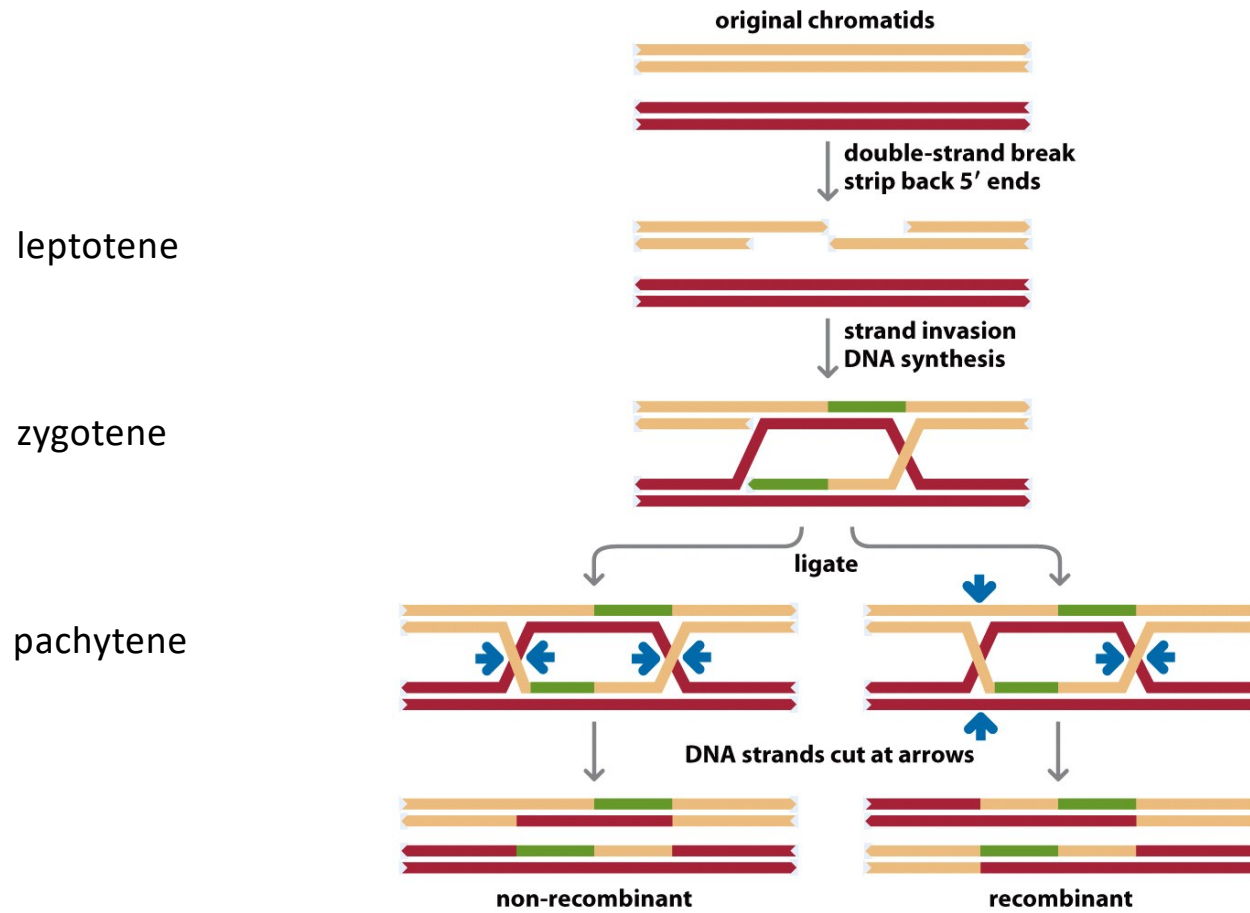
D: diplotene (partial separation of homologues by breakdown of the synaptonemal complex, held together by chiasmata)

E: diakinesis (chromosome condensation and transition to metaphase I)

Between leptotema and pachynema, these double-stranded breaks will be restored as crossovers and non-crossovers. The homologous chromosomes will align, pair and go in synapsis for this purpose (synapsis = formation of a synaptonemal complex between homologous chromosomes).

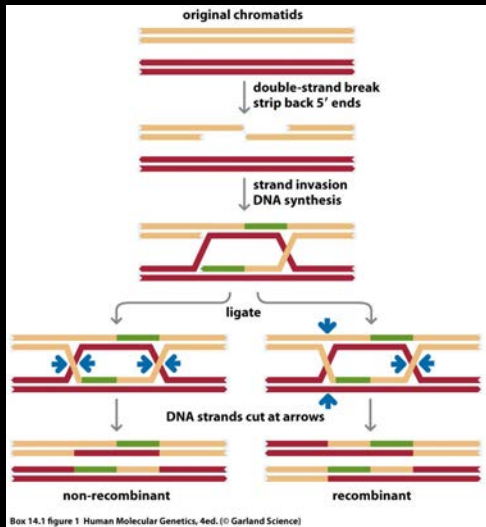


Homologous recombination

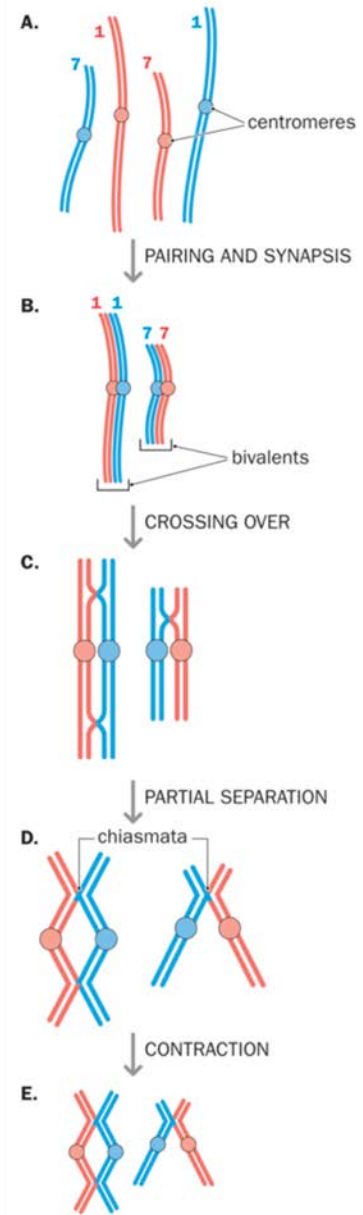
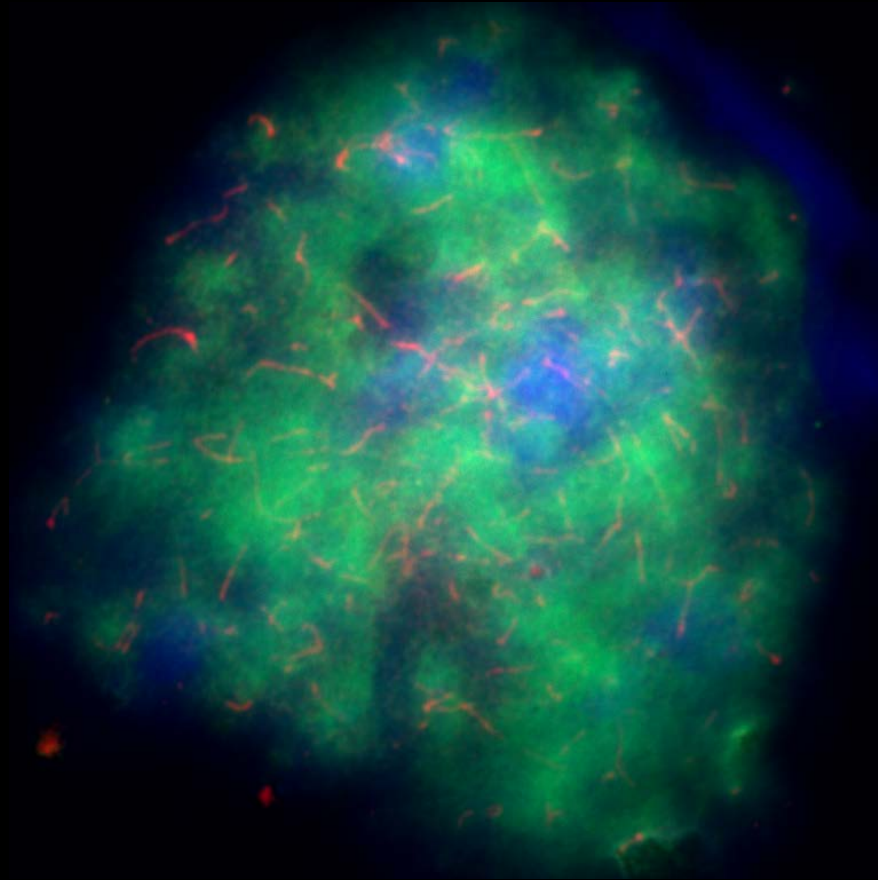


Box 14.1 figure 1 Human Molecular Genetics, 4ed. (© Garland Science)

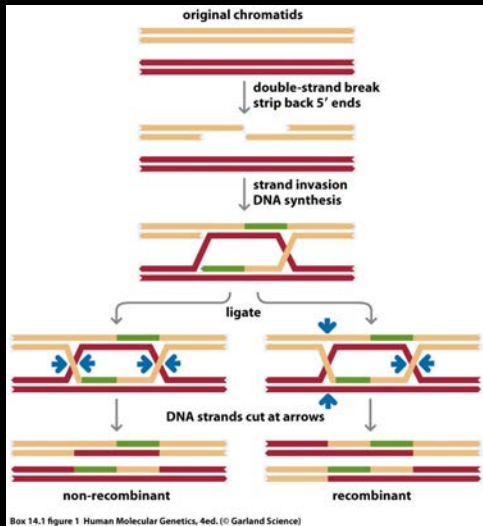
Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (γ H2AX (marker for DSB) - SCP3 - DNA staining)



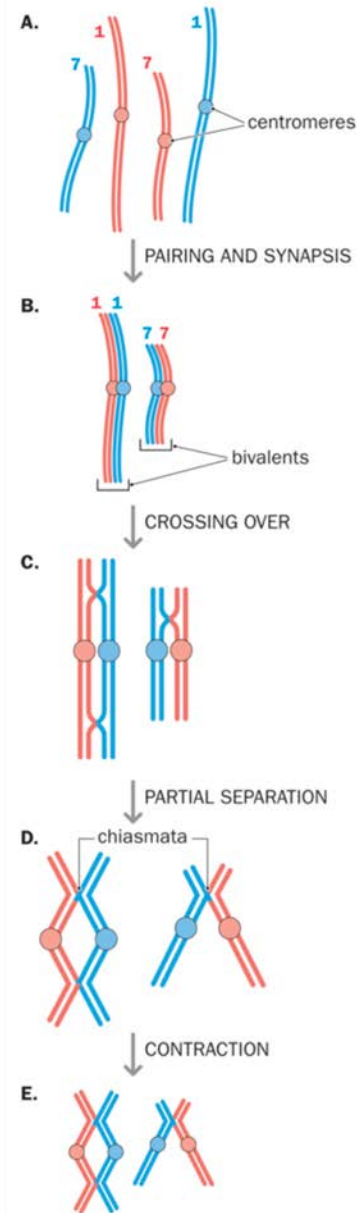
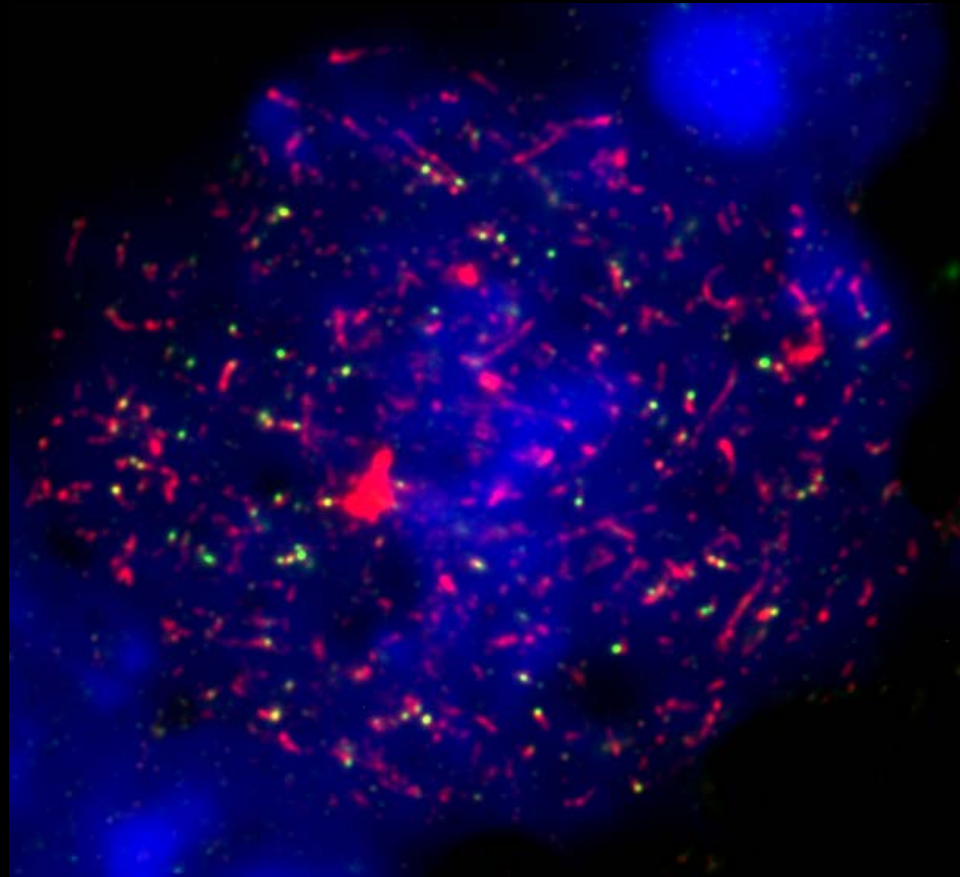
Leptonema



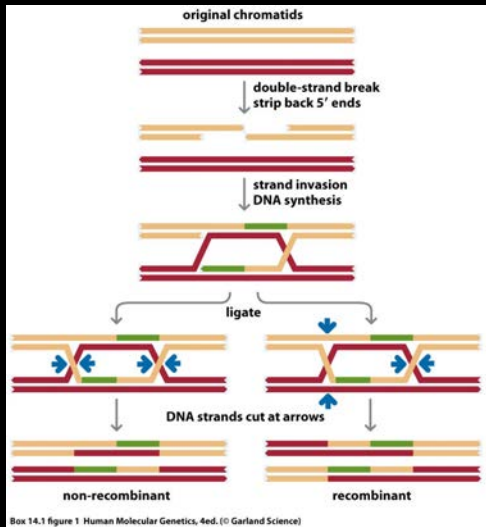
Analysis of prophase I :
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 (Rad51 – SCP3 - DNA staining)



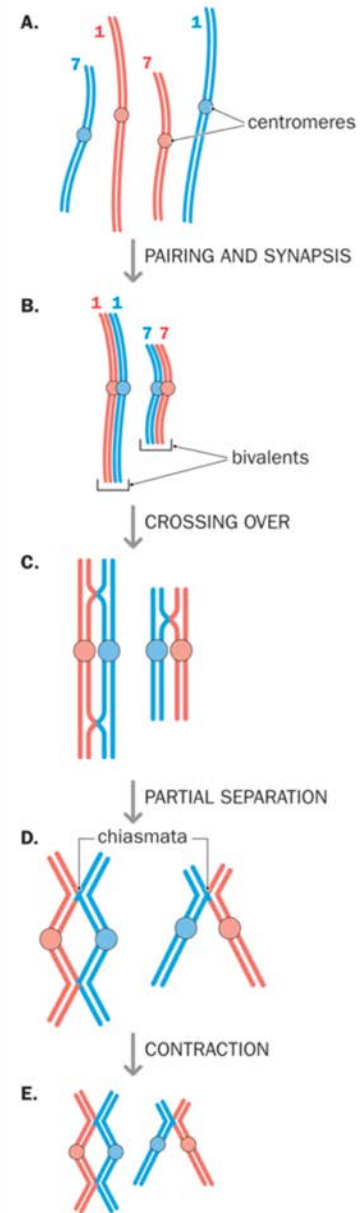
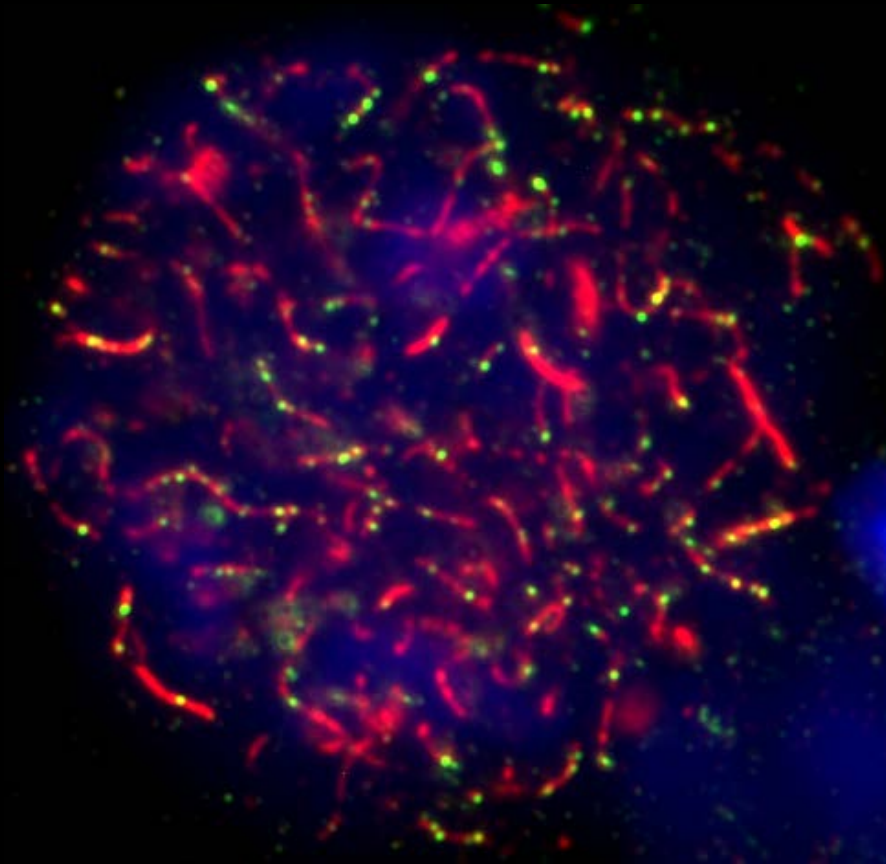
Early leptonema



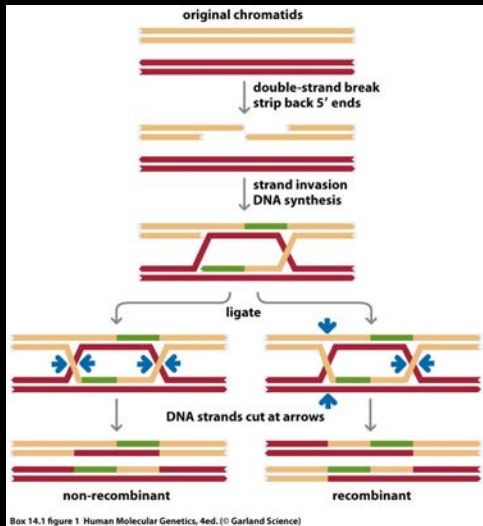
Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (Rad51 – SCP3 - DNA staining)



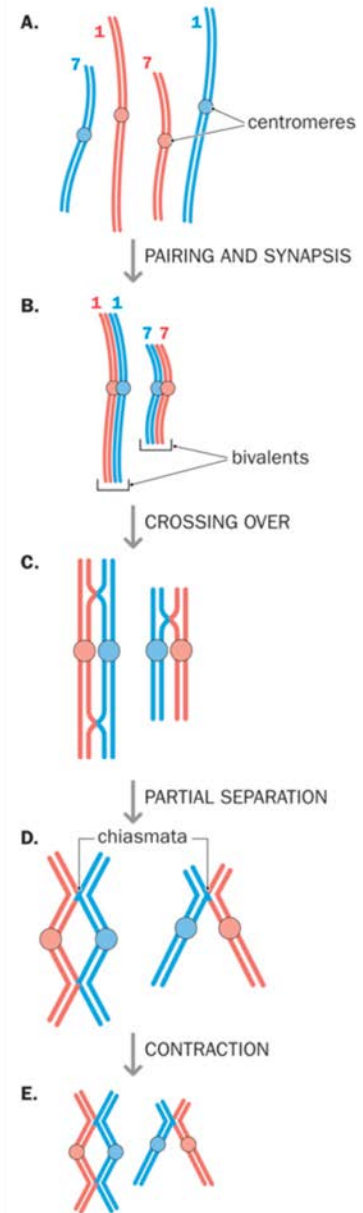
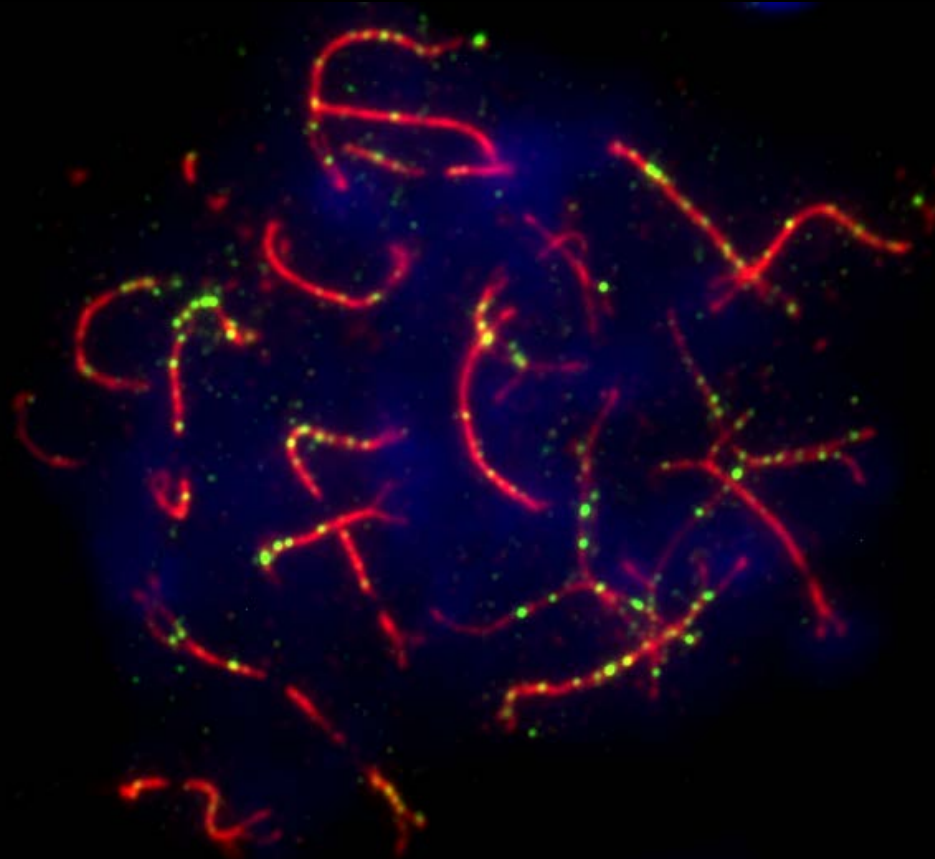
Late leptonema - Early zygonema



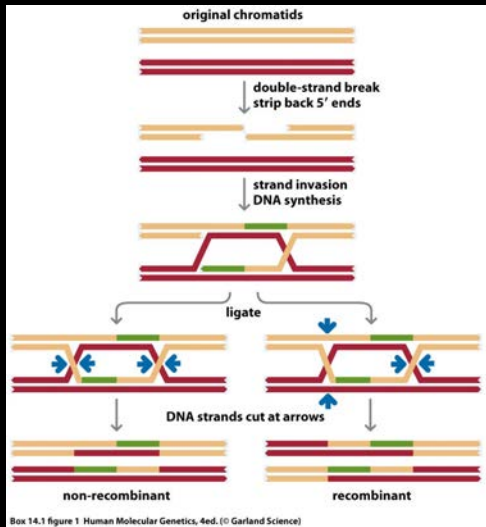
Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (Rad51 – SCP3 - DNA staining)



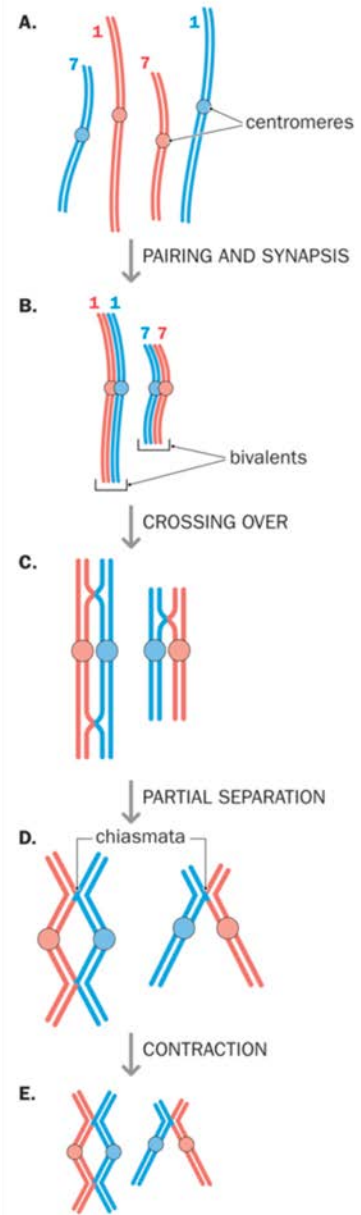
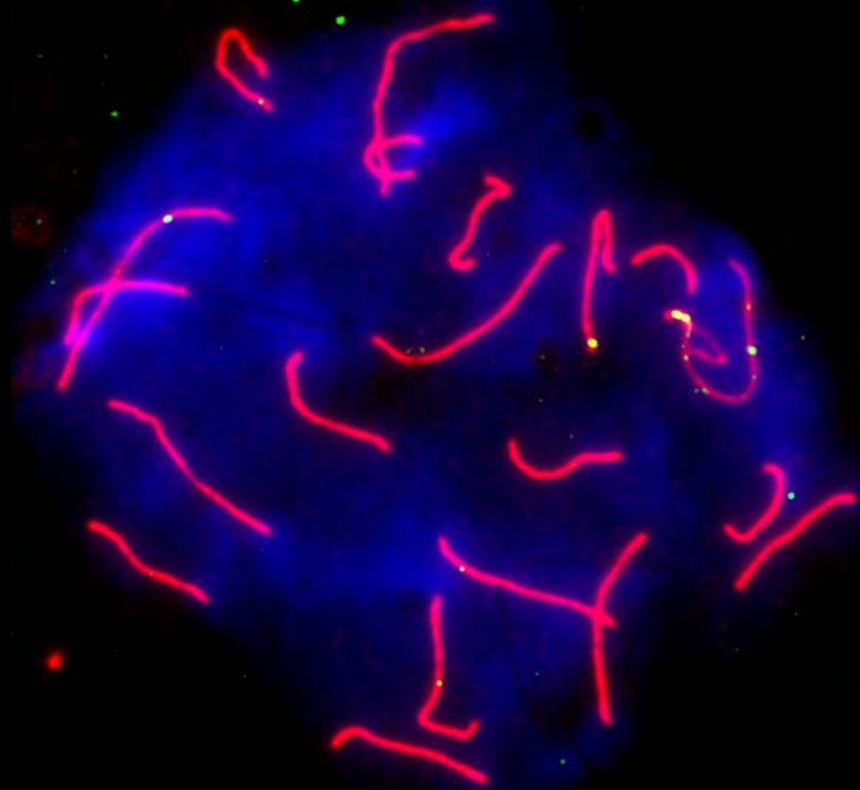
Late zygonema



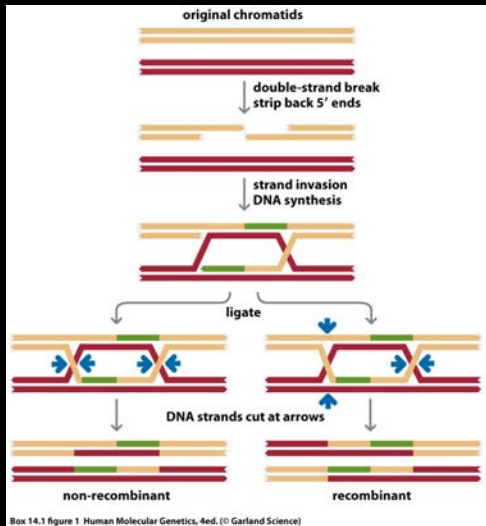
Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (Rad51 – SCP3 - DNA staining)



Early-Pachynema

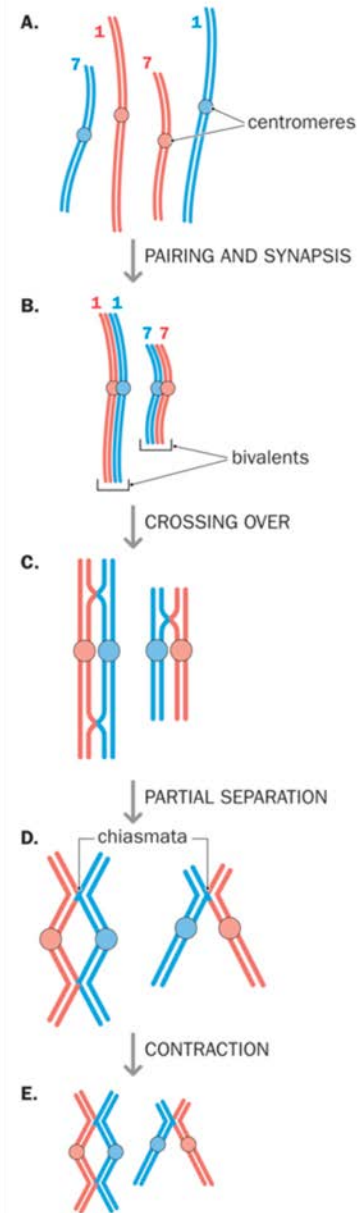
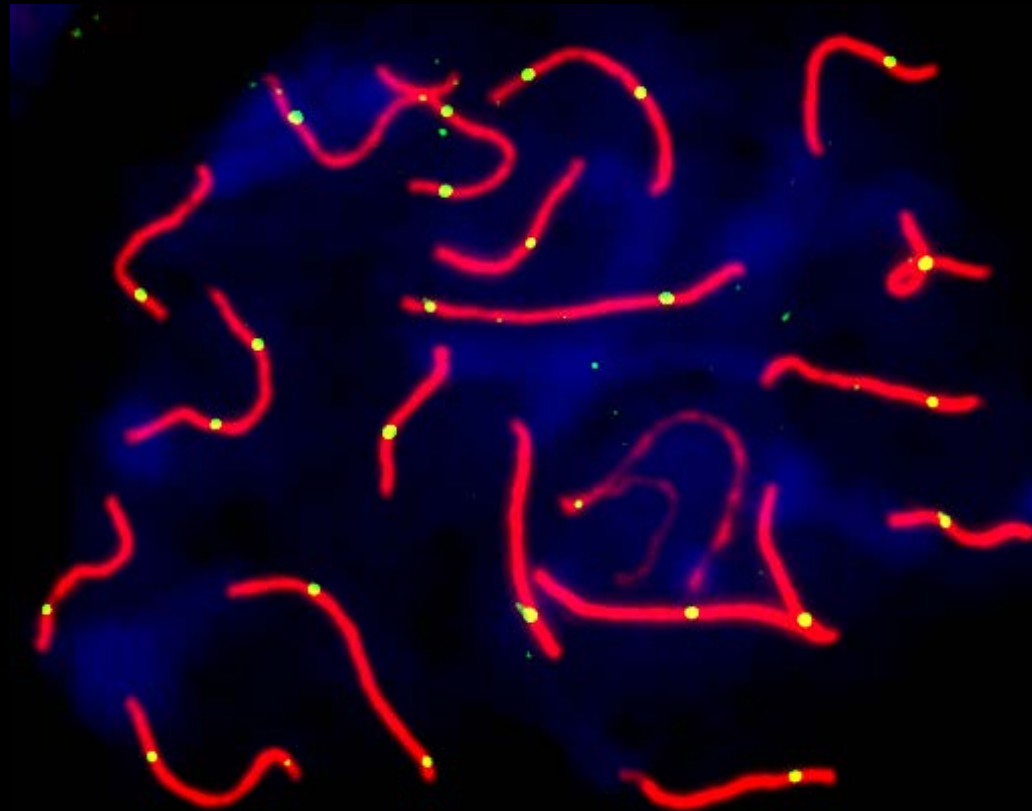


Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (Mlh1 (marker for genetic crossover) - SCP3 - DNA staining)

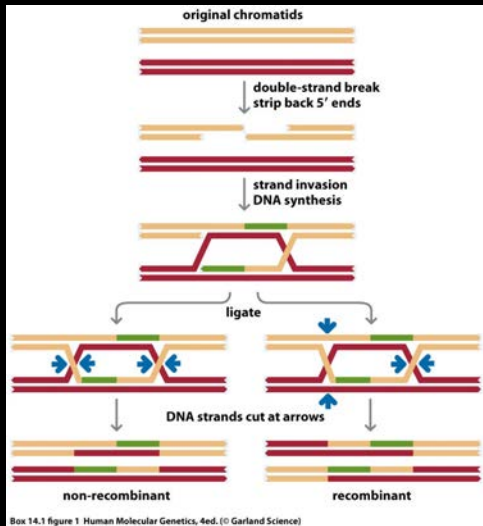


Cross-over interference

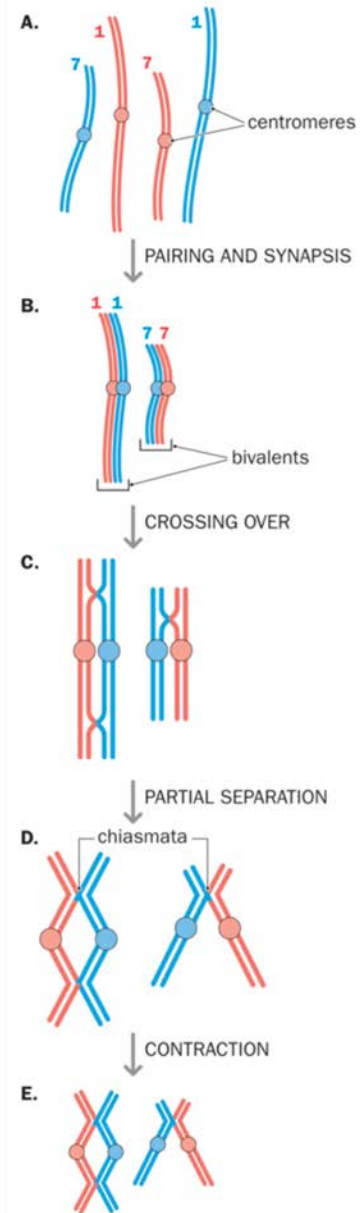
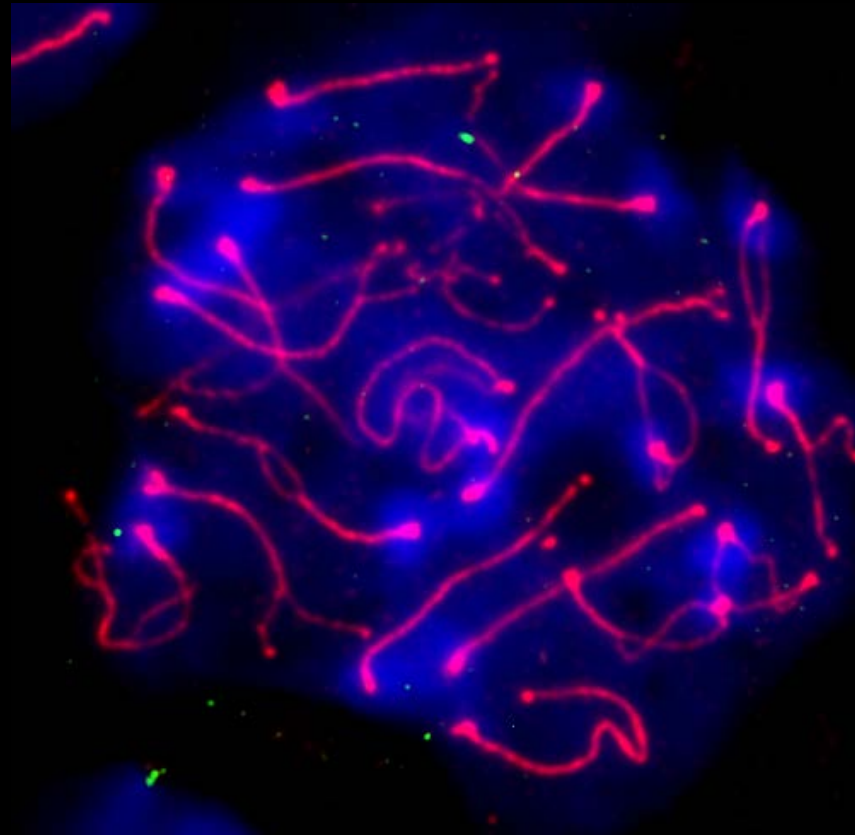
Mid-Pachynema



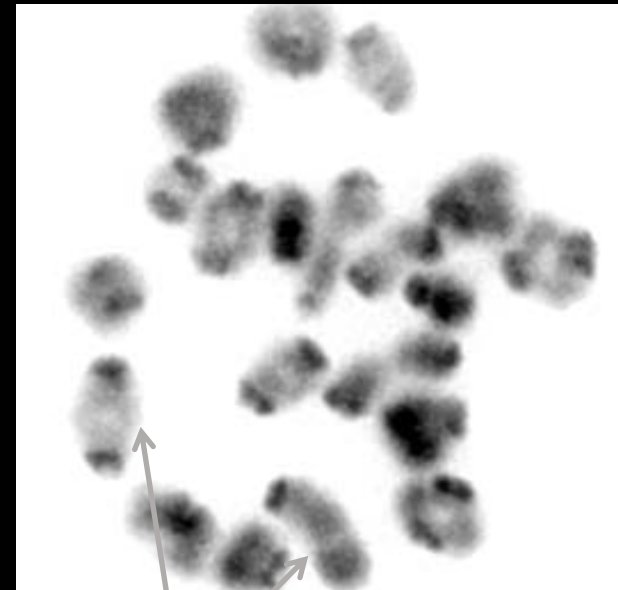
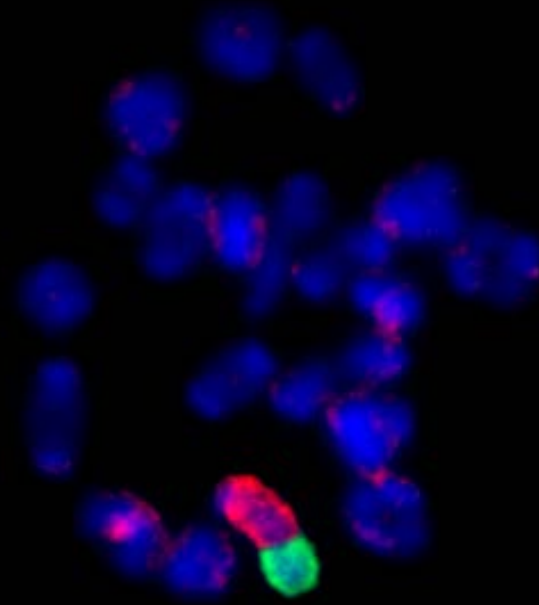
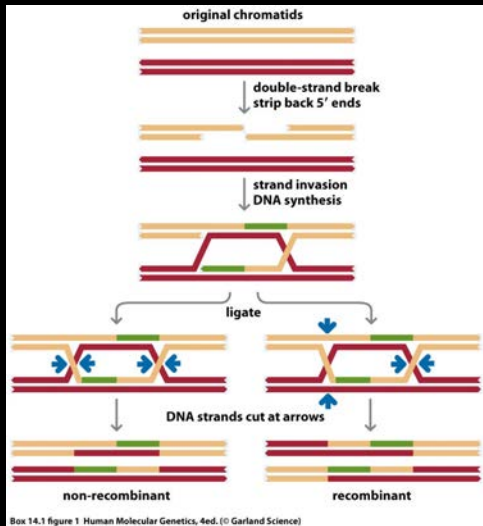
Analysis of prophase I :
 DSB formation and repair process / Homologue pairing and synapsis
 (Rad51 – SCP3 - DNA staining)



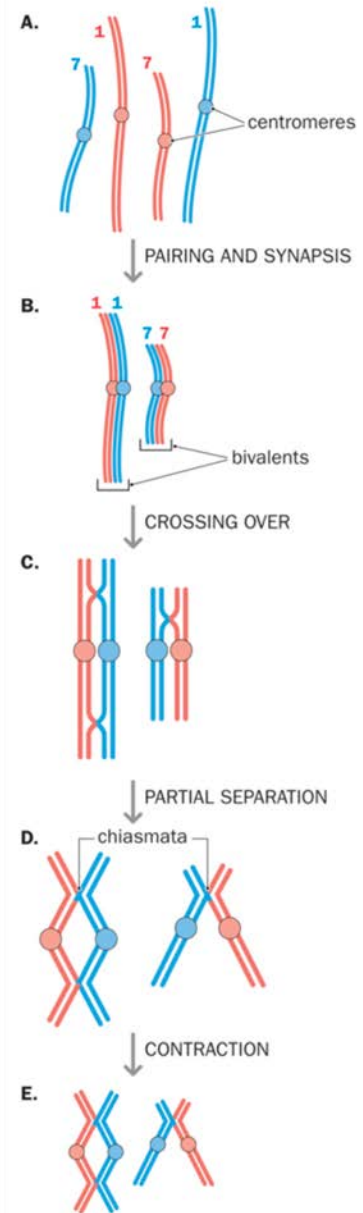
Diplonema

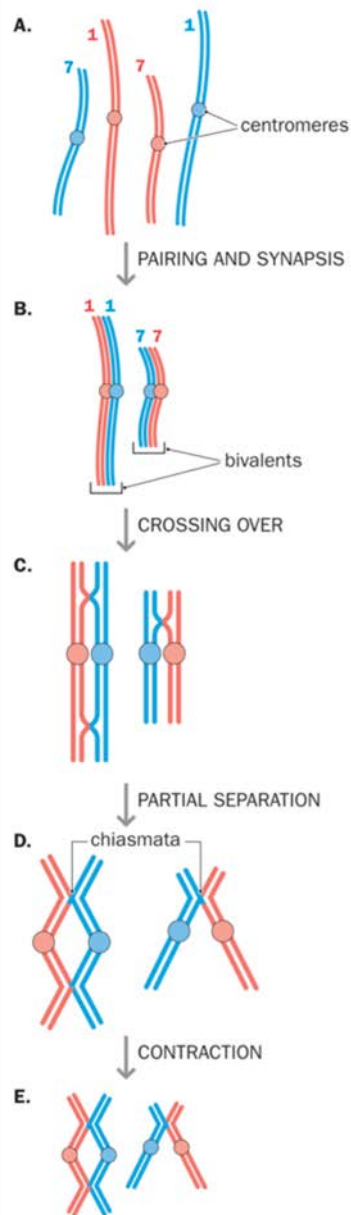


Analysis of metaphase I :
 20 bivalents in normal mouse meiosis
 (FISH Y – X – DNA staining)



Chiasmata





The five stages during prophase of meiosis I

A: leptotene (chr condensation, chr unpaired, dsDNA breaks [DSB])

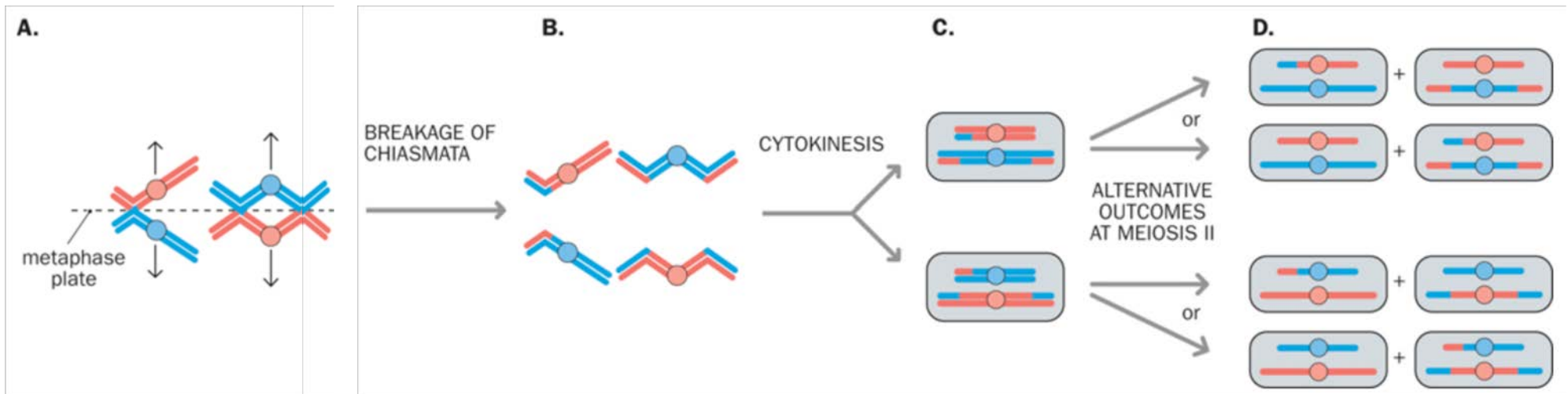
B: zygotene (repair of DSB in progress; pairing of homologues to bivalents; synapsis through synaptonemal complex)

C: pachytene (synapsis complete; crossing-over complete; formation of chiasmata)

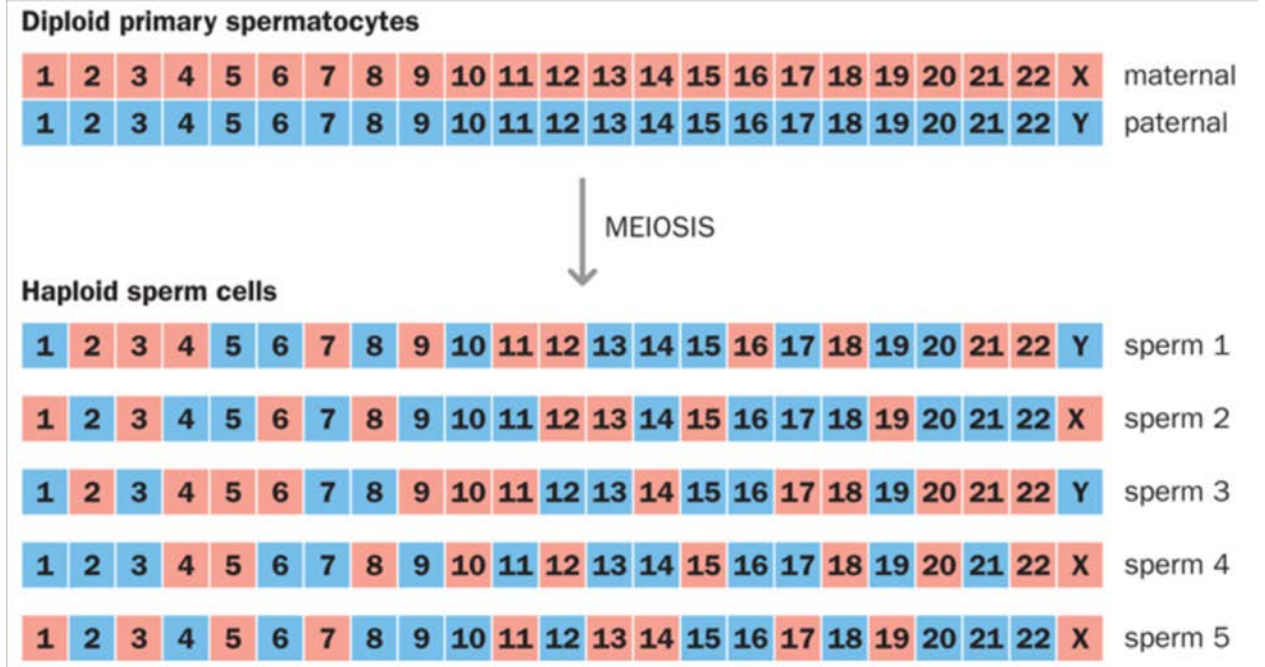
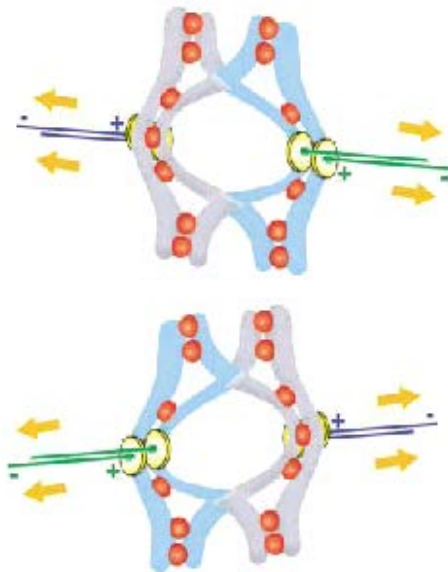
D: diplotene (partial separation of homologues by breakdown of the synaptonemal complex, held together by chiasmata)

E: diakinesis (chromosome condensation and transition to metaphase I)

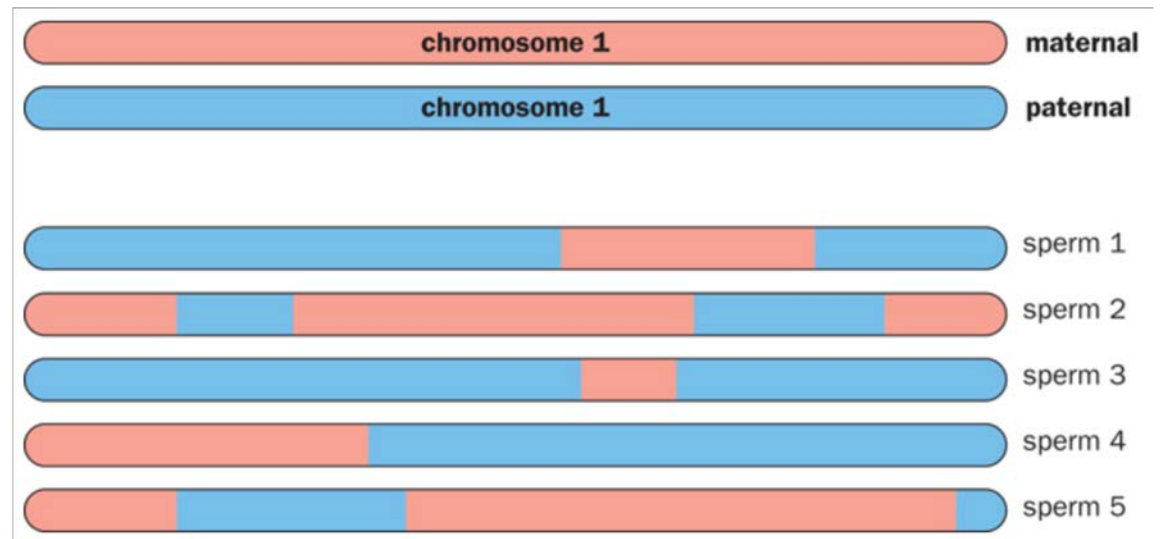
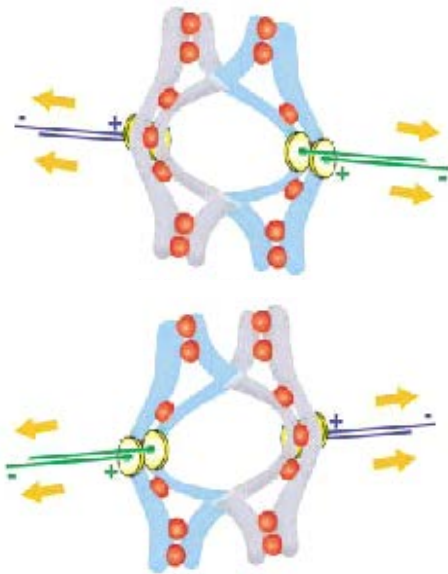
From metaphase I to gametes

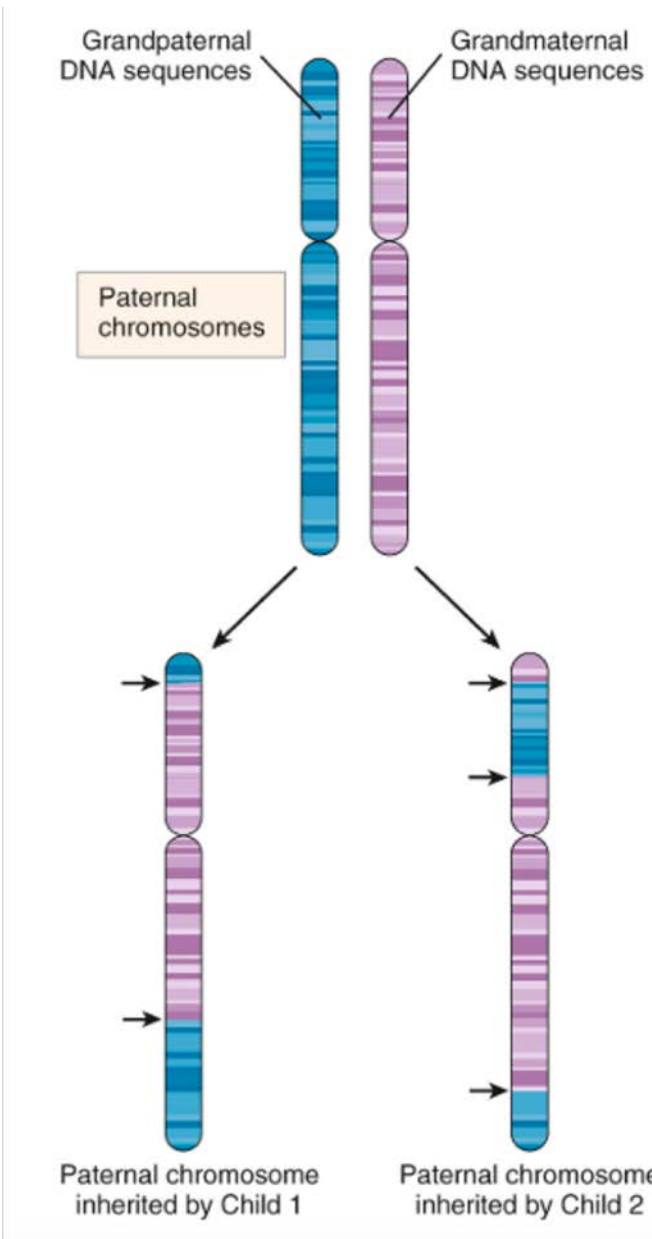


Source of genetic diversity: **Independent assortment** + homologous recombination




Source of genetic diversity: Independent assortment + **homologous recombination**

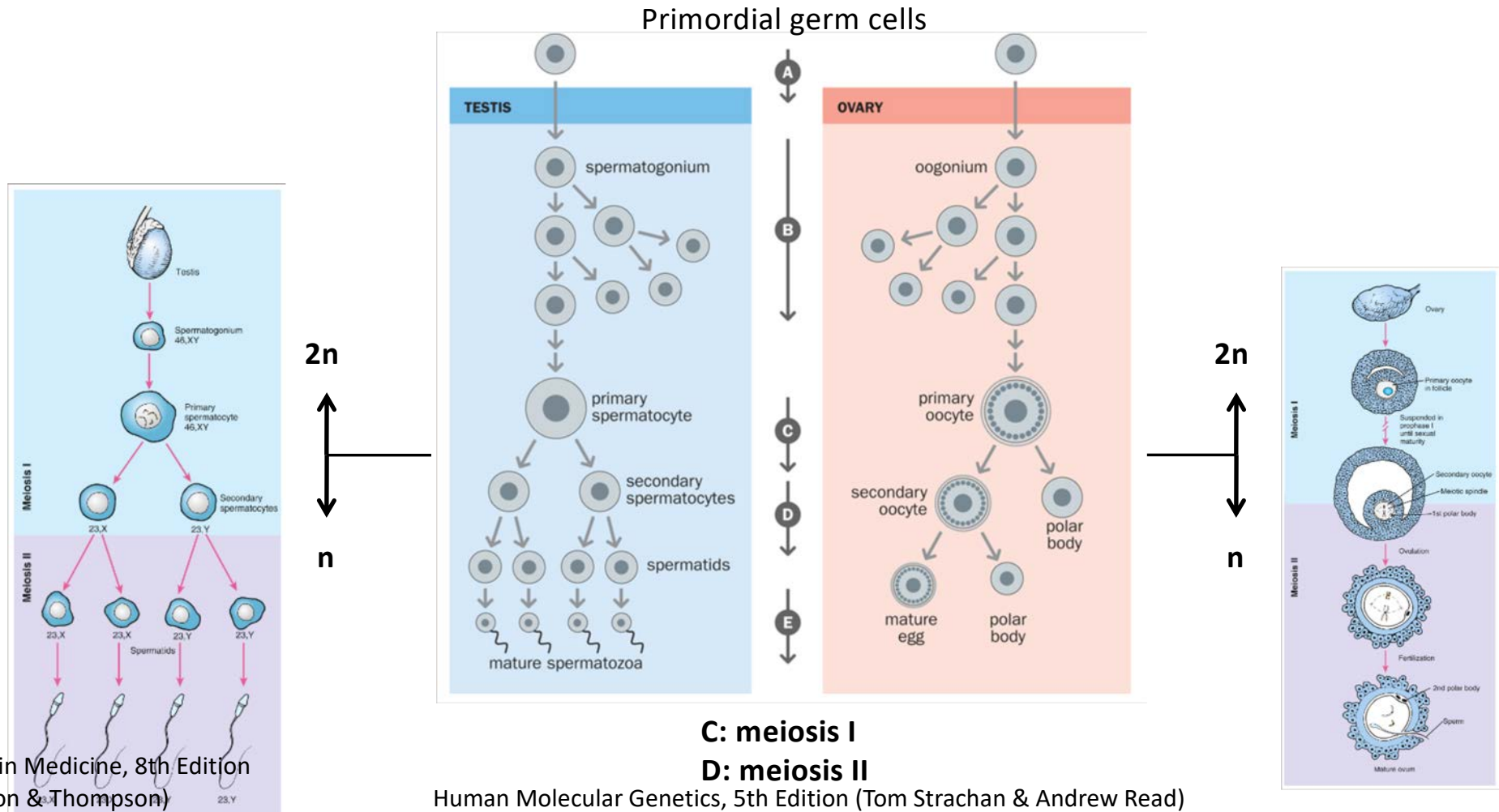




Chapter 2

- ✓ The Human Genome Sequence
- ✓ Organization of the Human Genome
 - Single-Copy DNA Sequences
 - Repetitive DNA Sequences
 - Repetitive DNA and Disease
- ✓ Variation in the Human Genome
- ✓ Transmission of the Genome
 - The Cell Cycle
 - Mitosis
 - Meiosis
-  ✓ Human Gametogenesis and Fertilization
 - Spermatogenesis
 - Oogenesis
 - Fertilization
- ✓ Medical Relevance of Mitosis and Meiosis


General overview of gametogenesis in ovary and testis



Genetics in Medicine, 8th Edition
(Thompson & Thompson)

Human Molecular Genetics, 5th Edition (Tom Strachan & Andrew Read)

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