

Premedical - Biology

Mitosis and Cell cycle

Physiological modes of somatic cell



Proliferation in:

- ontogenesis
- physiological renewal of cells
- reparation and wound healing
- immune response

Aging (senescence)

- limited number of cell division (maximum 50) →

Hayflick's limit

both in vivo and in vitro

- accumulation of mutations
- decreased cytokines response, increased synthesis of inhibitory proteins
- shortening of telomere sequences at the ends of chromosomes

Resting (Quiescent) Cells: G₀

G₀ phase relates
to terminal stages of
differentiation

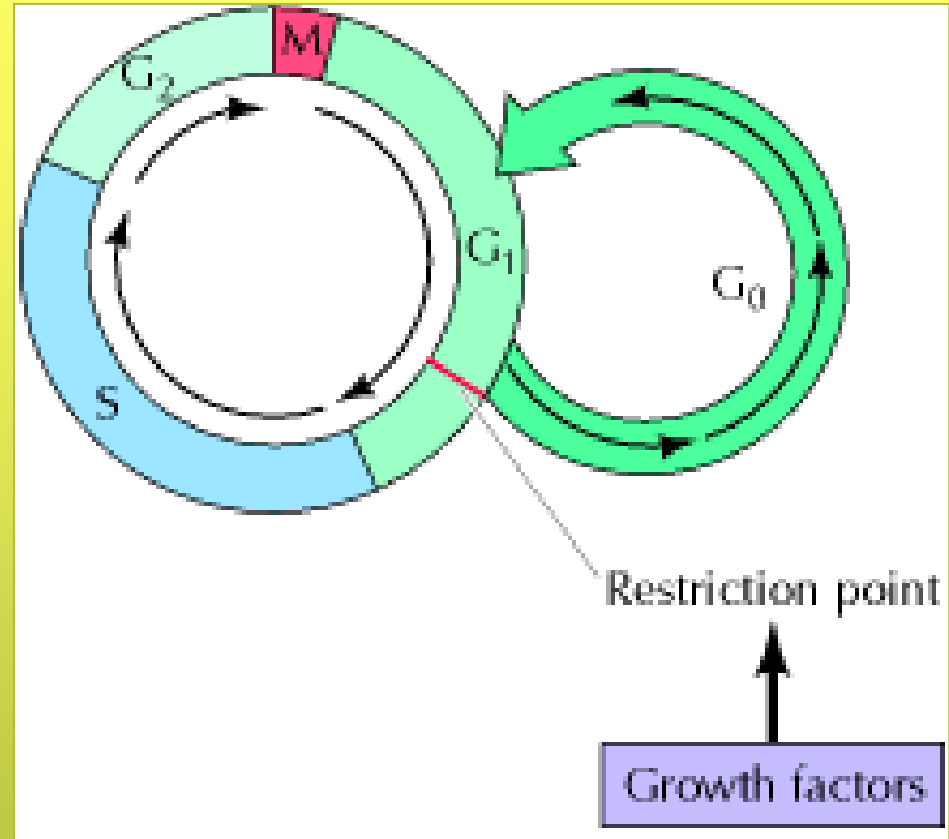
e.g. hepatocytes divide

1x a year;

neurons, myocytes

do not divide;

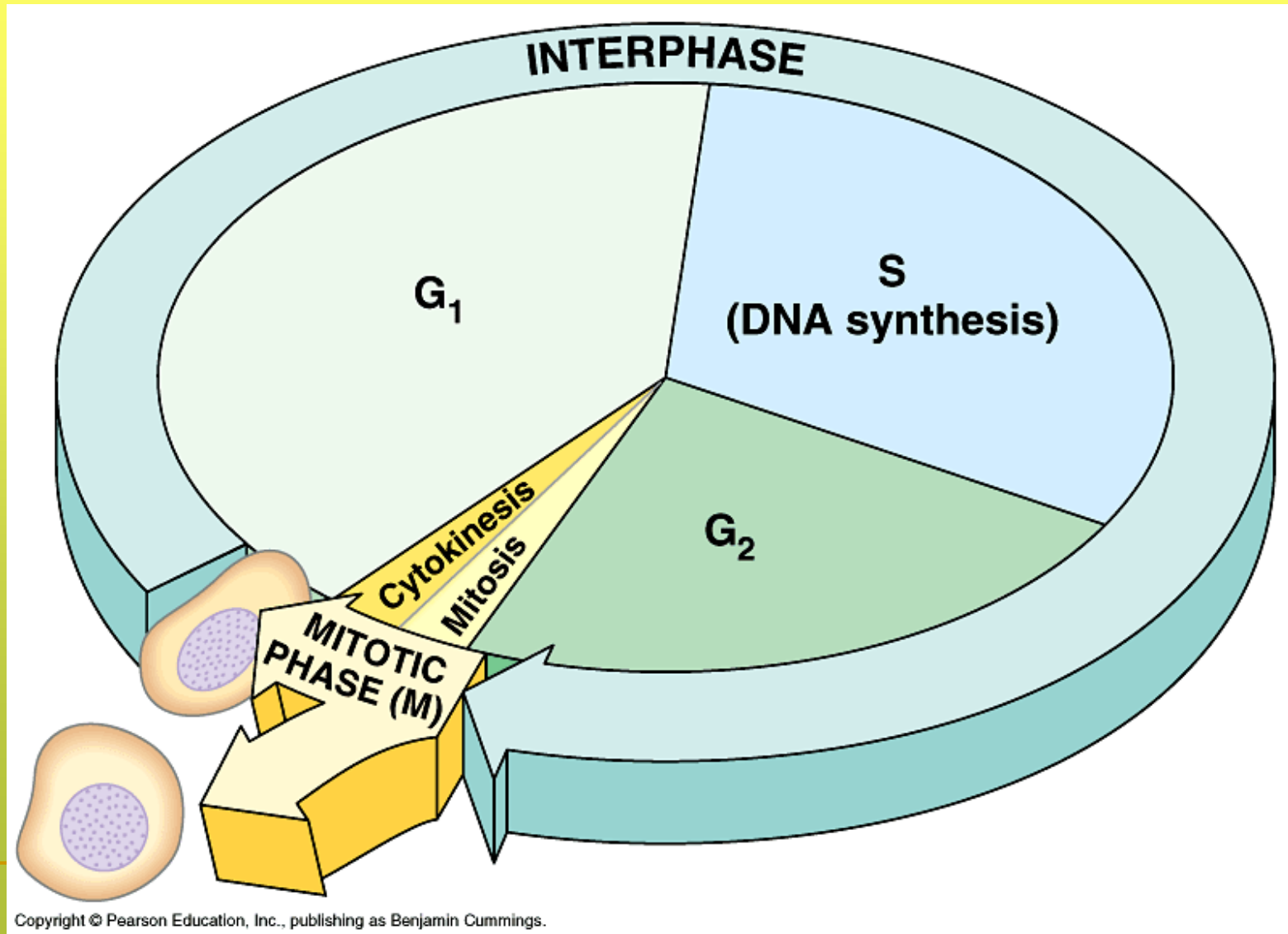
epithelial cells divide 1-2x a day



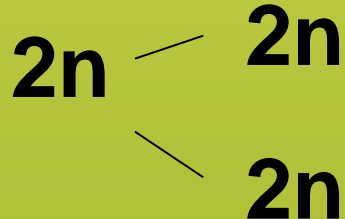
Cell cycle

G ~ Gap/Growth

S ~ DNA sythesis



The cell cycle

- **M phase and interphase**
 - **M phase: Mitosis and cytokinesis**
 - Interphase : G1, S, G2 phase
 - 46 chromosomes, 23 chromosomes from each parent
 - mitosis – **distribution of identical sets of 46 chromosomes to daughter cells**
- 
- The diagram illustrates the process of mitosis. A parent cell, labeled with $2n$, is shown on the left. Two lines diverge from this cell to the right, leading to two daughter cells, each also labeled with $2n$. This represents the equal distribution of genetic material during cell division.

Cell Cycle

- **G1** phase – the longest and the most variable part of the cell cycle
 - growth of the cell
 - completion of organelles (ribosomes, mitochondria, endoplasmic reticulum etc.)
 - RNA and protein synthesis
 - synthesis of nucleotides, preparation for replication

- **S phase** – replication of nuclear DNA
(extranuclear DNA replicates during the whole interphase)
- **G2 phase** – cell growth, protein and RNA synthesis, origin of cell structures
- **M phase:**

Mitosis - division of the nucleus

Cytokinesis – division of the cell

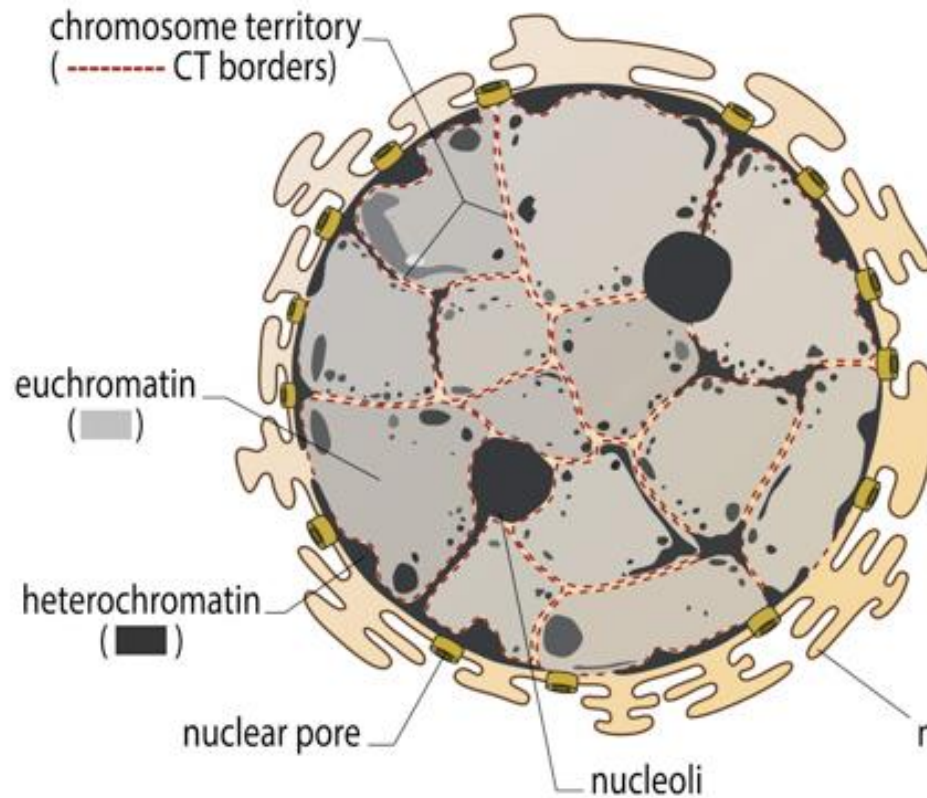
Mitosis – animal cells

Prophase: Chromatin fibers are getting coiled, spiralized.

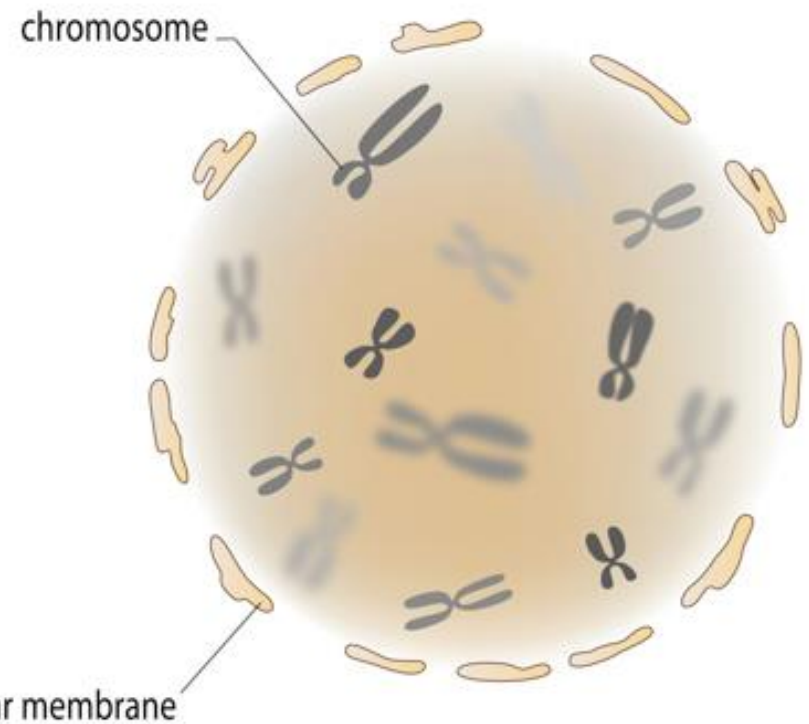
Nucleoli disappear and microtubuli begin to form mitotic spindle

Prometaphase: Nuclear envelope fragments. Mitotic spindle interacts with chromosomes. Chromosomes are getting more condensed.

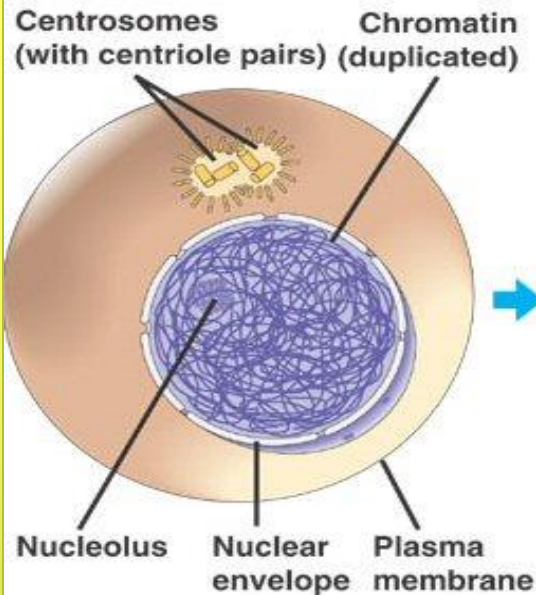
A. Interphase nucleus



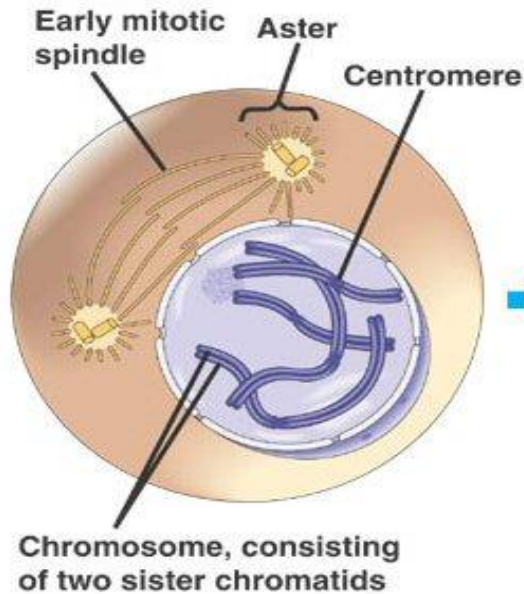
B. Prometaphase nucleus



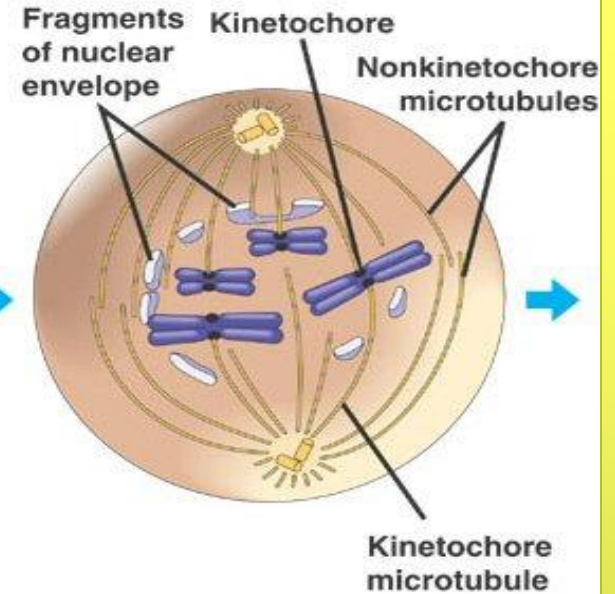
G₂ OF INTERPHASE



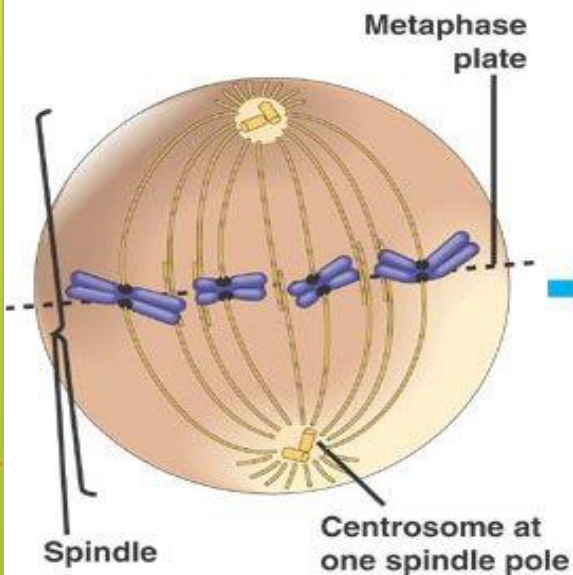
PROPHASE



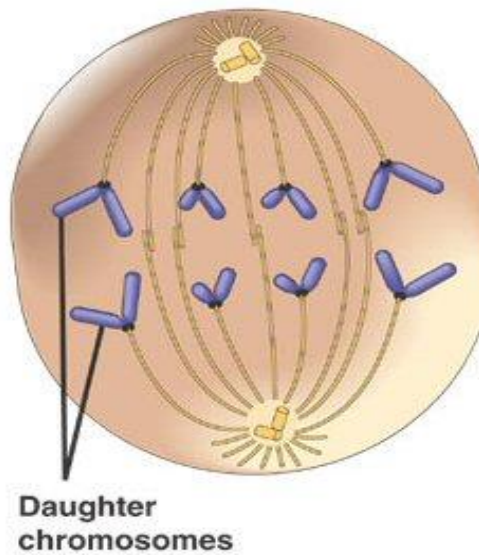
PROMETAPHASE



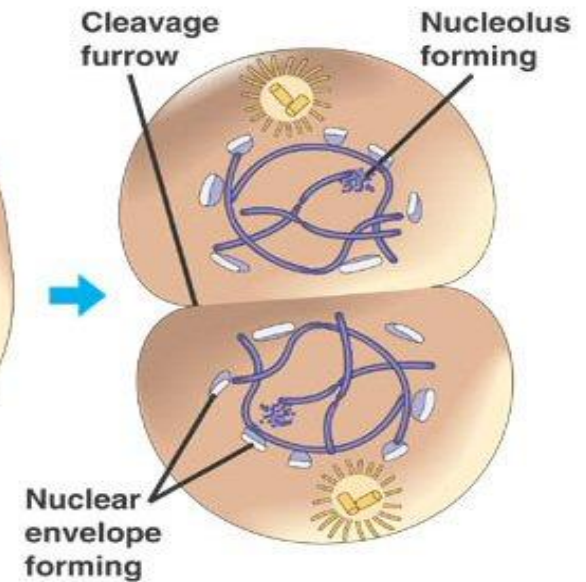
METAPHASE



ANAPHASE



TELOPHASE AND CYTOKINESIS

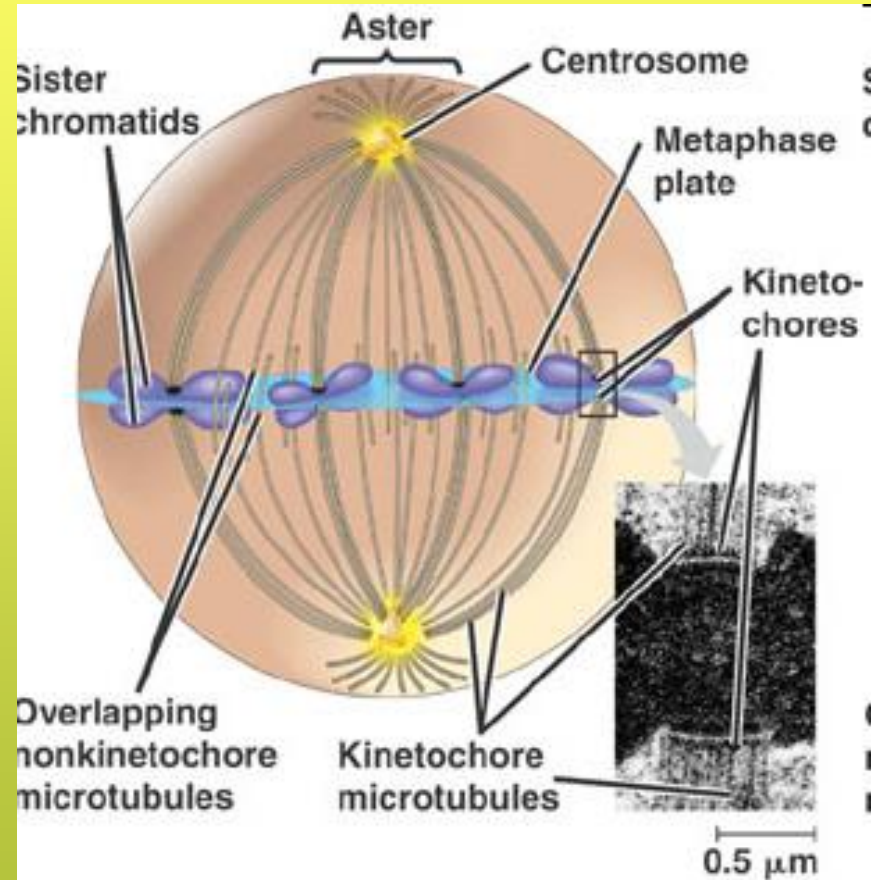





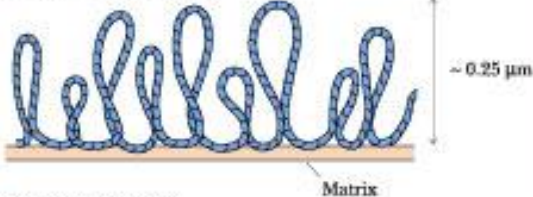


Mitotic spindle

- Fibers made of microtubules, spindle starts from centrioles - 9 sets of triplets of microtubules from subunits tubulin α , β
- **Microtubule organizing center**
- Mitotic spindle elongates by incorporating subunits of protein tubulin
- Microtubule polarity

- Kinetochore microtubules
- Non-kinetochor m. (polar)
- Astral microtubules

kinetochor – proteins and chromosomal DNA at the centromere



	Base pairs per turn	Packing ratio
DNA double helix 	10	1
"Beads on a string" chromatin form 	80	6-7
Solenoid (six nucleosomes per turn) 	1200	~40
Loops (50 turns per loop) 	60,000	680
Miniband (18 loops) 	$\sim 1.1 \times 10^6$	1.2×10^4
Chromosome (stacked minibands) 	18 loops/miniband	1.2×10^4

Nucleosome:

DNA double helix + histone core

Histone core = octamer of two copies of H2A, H2B, H3, H4 histons

Spacer segment between two nucleosomes is free or associated with H1 histone

String of nucleosomes is coiled into **solenoid** (6 nucleosomes in each turn)

Solenoid is packed into loops, attached to **non-histone protein scaffold** (Laemli loops).

Non-histone protein scaffold with loops is coiled into spiral structure of

chromatids

Metaphase: Spindle poles are at opposite positions.

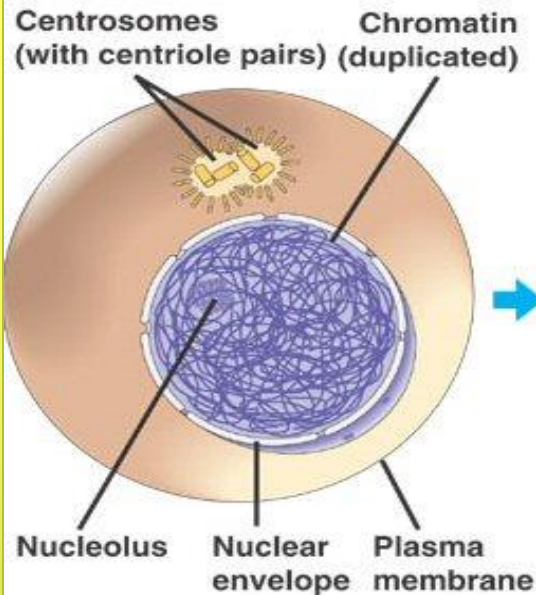
Chromosome are located on the metaphase plate (in equatorial plane). Each chromosome is attached by kinetochore to the mitotic spindle.

Anaphase: Chromatids move to opposite poles of the cell.

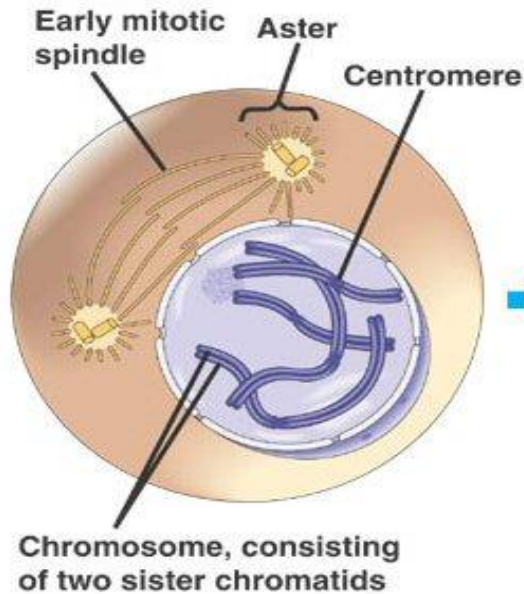
Kinetochore microtubules are getting shorter, the poles move further apart. There are at the end two collection of chromosomes.

Telophase: Non-kinetochore microtubules elongate. Nuclear envelope generates. Cytokinesis starts to run.

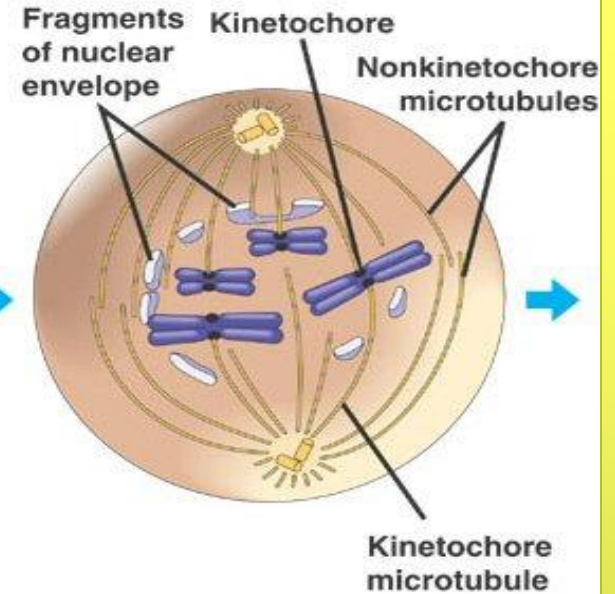
G₂ OF INTERPHASE



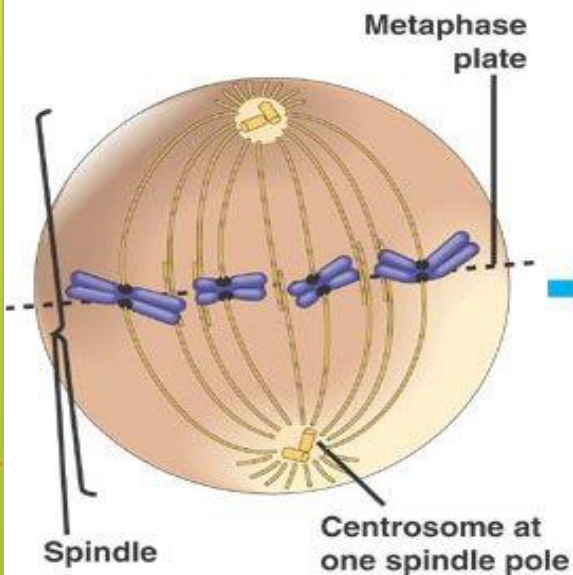
PROPHASE



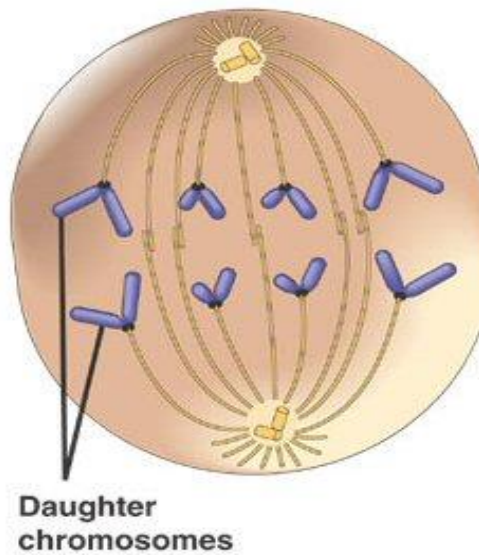
PROMETAPHASE



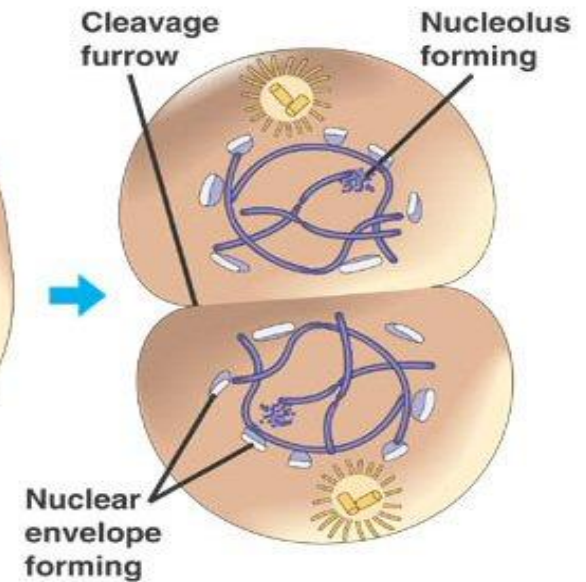
METAPHASE



ANAPHASE



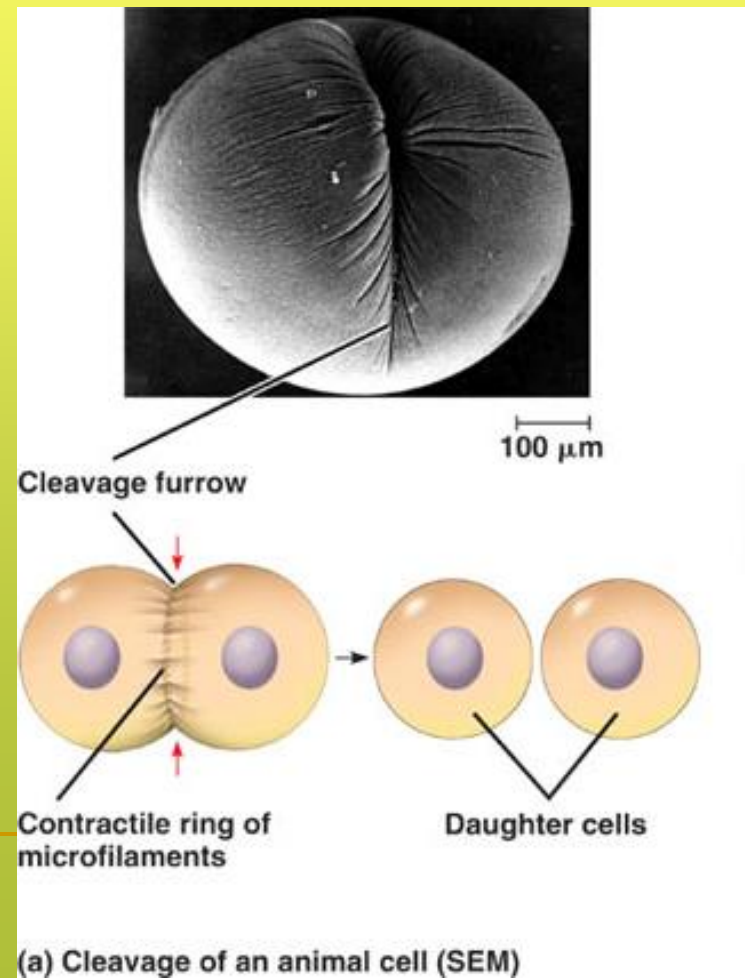
TELOPHASE AND CYTOKINESIS



Cytokinesis

Cleavage

- contractile ring of actin microfilaments
- cell plate in plant cells



Cell cycle

- External signals and internal network of interactions – signalling transduction pathways regulate the cell cycle
 - **Cancer cells have escaped from cells cycle controls**
-

Check points

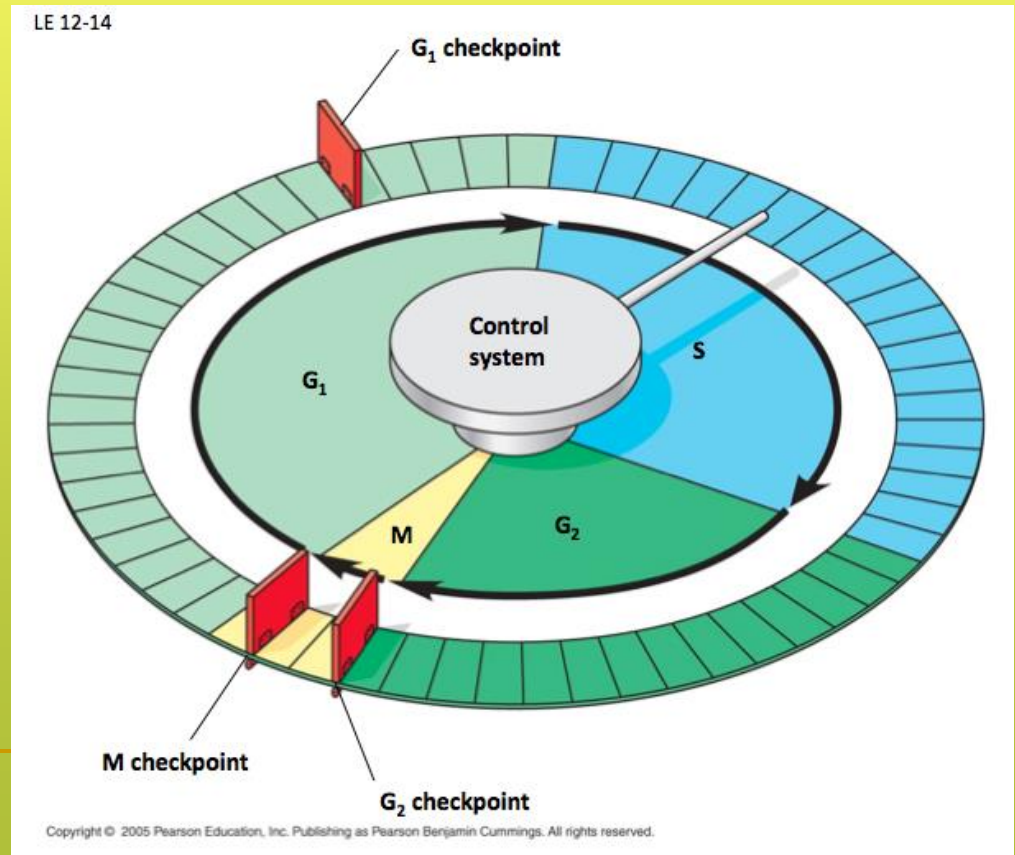
trigger and coordination of key events

Checkpoints are critical points, where signals can stop or go-ahead to the next phase of cell cycle:

G1 checkpoint

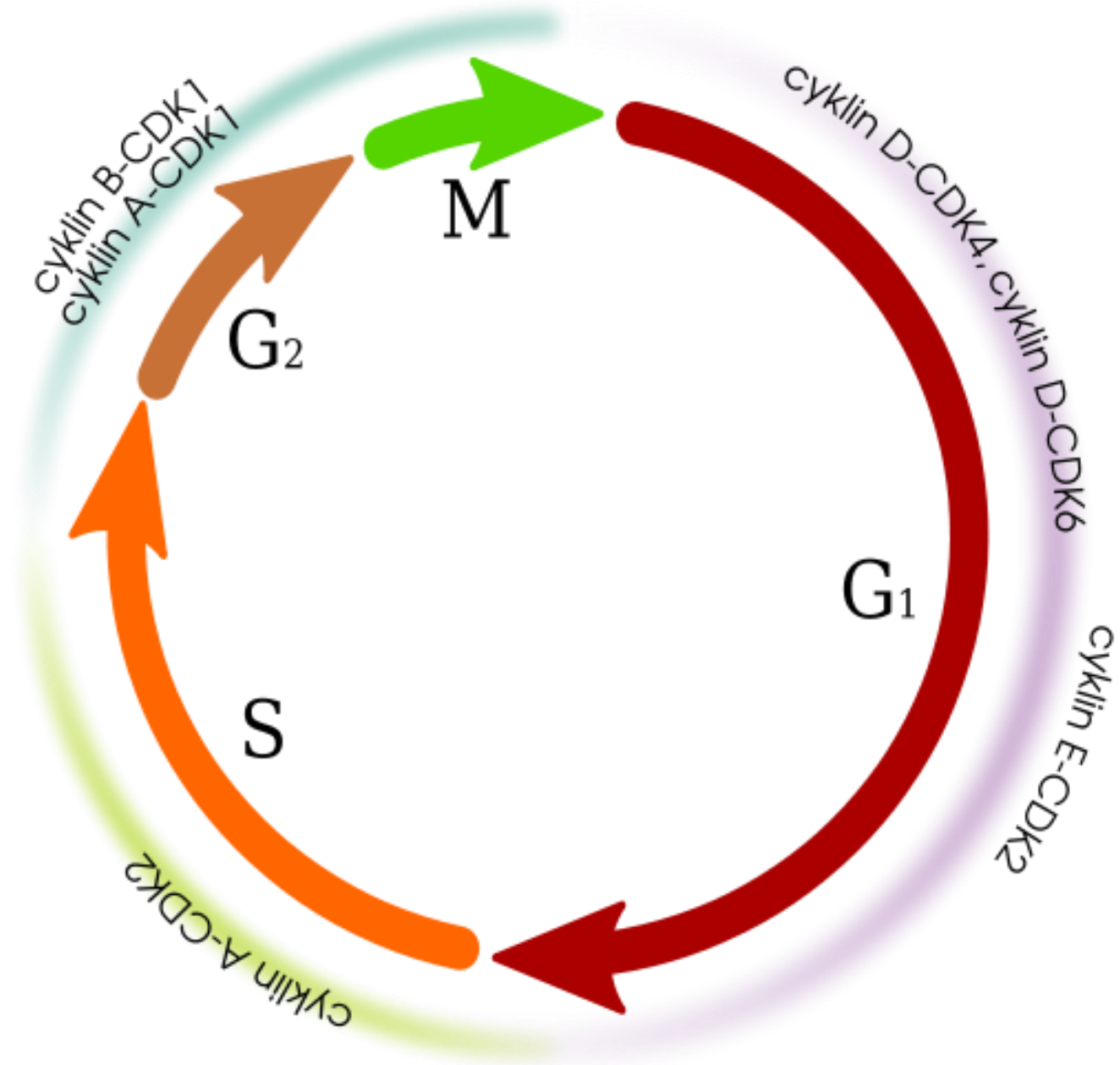
G2 checkpoint

M checkpoint

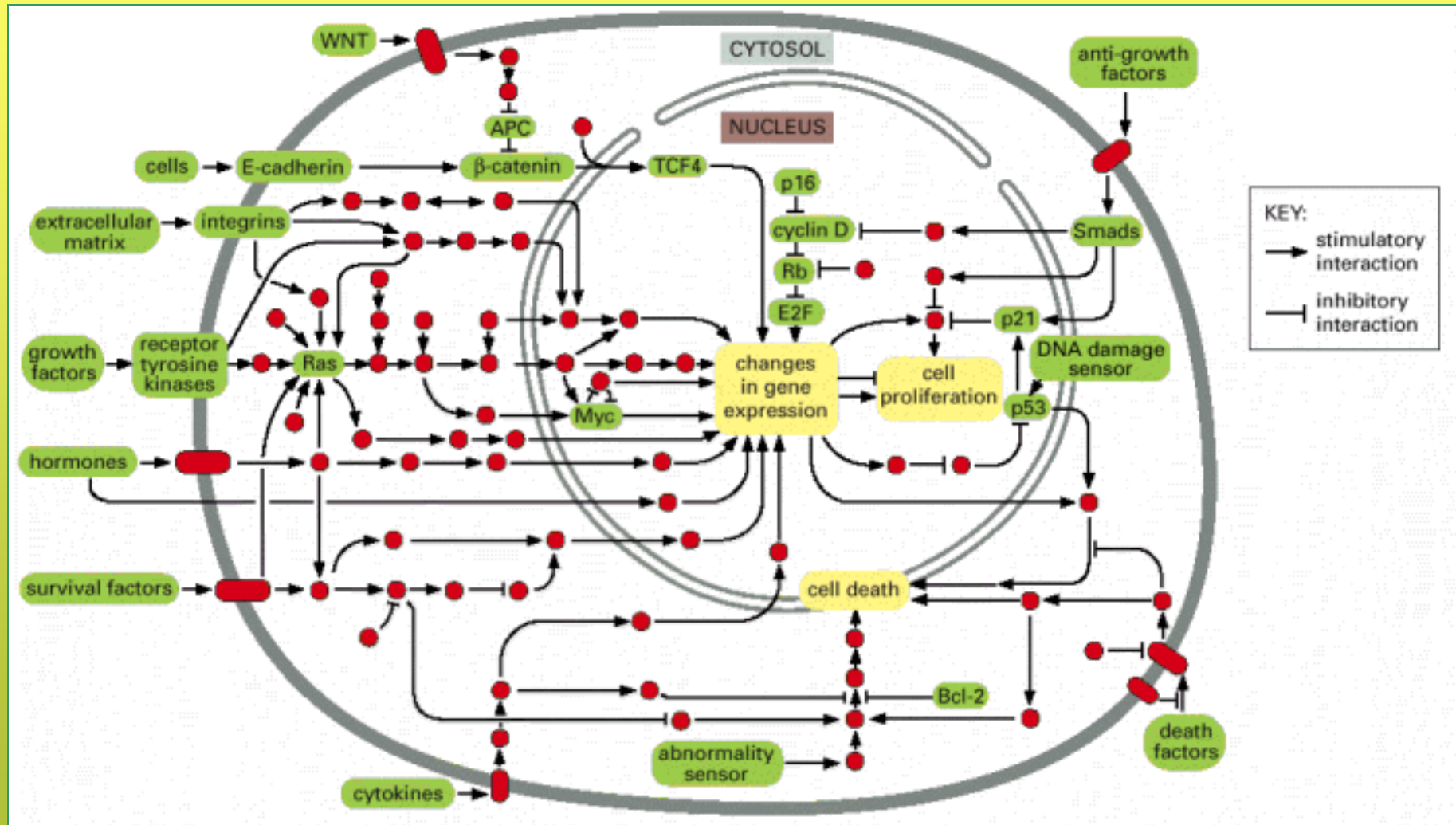


Control system of cell cycle

- **Cyclin** – cyclic accumulation and degradation of proteins during the cell cycle
 - **Cdk – cyclin dependent kinases (CDK)**
 - = enzymes that phosphorylate other proteins in active states
 - = **activation by cyclin**
- complex cyclin / kinase => protein phosphorylation =>
triggers cell cycle phases**



Signal transduction pathways – proliferation, apoptosis



Genes regulating cell cycle:

Protooncogenes

- products **stimulate** cell division
- Genes for **growth factors, receptors, regulatory proteins, Ras proteins**
- mutated forms = **oncogenes** => permanent or increased mitotic activity
(effect of one allele mutated)

Tumor suppressor genes (TSG) „antioncogenes“

- products **inhibit** mitotic division
- effect of **both** alleles mutated
- **Rb1** gene, product RB protein
 - Mutations in retinoblastom and other tumors
- **TP 53** gene, **p53** product – induction of DNA repair or apoptosis = programmed cell death
- mutations in many tumors

Carcinogenesis

Mutator genes – genes for reparation enzymes

Proteins encoded by proto-oncogenes and tumor-suppressor genes are components of cell-signalling pathways.

Multistep model of cancer development

Apoptosis = programmed cell death

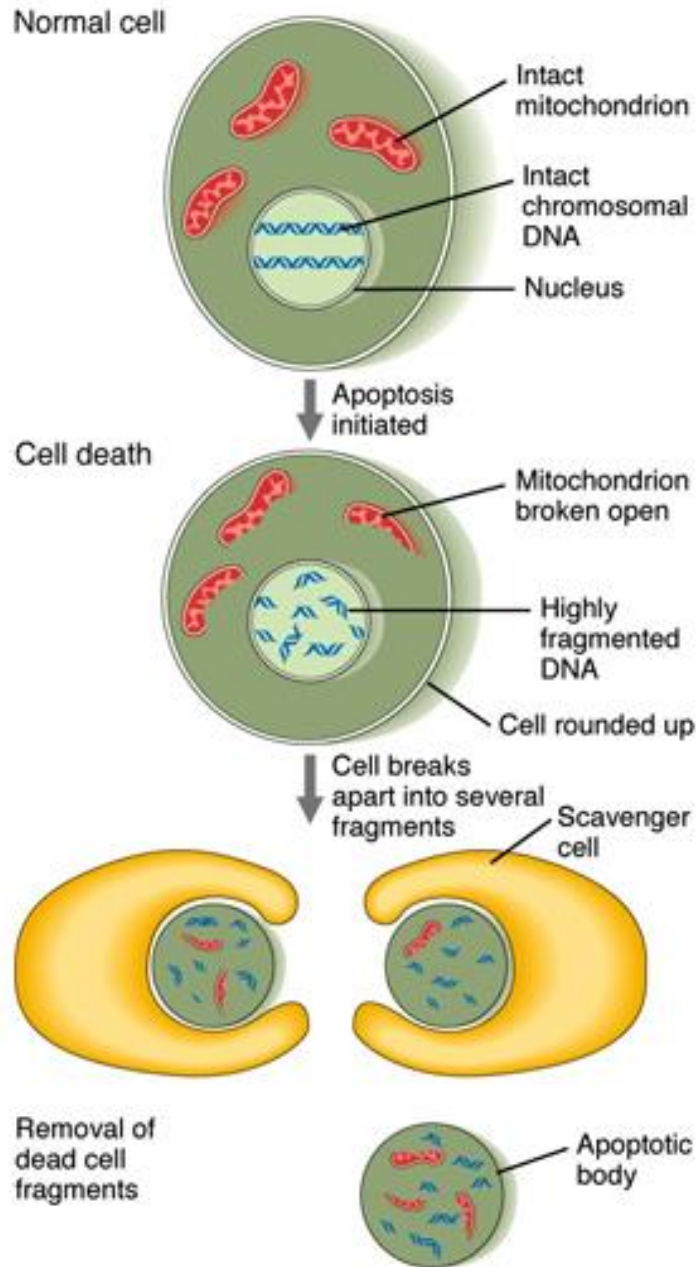
- final stage of aging process in the cell
- elimination of cells, which can not be repaired
- during embryogenesis - reduction of redundant parts
- some diseases
- Purpose: elimination of cells, that accomplished their fate and could become destructive for the organism

Apoptosis:

- without disintegration of both plasma membrane and organelles
- chromatin condensation, surface blebbing, cell fragmentation → **apoptotic bodies**
- phagocytosis without inflammation

Necrosis:

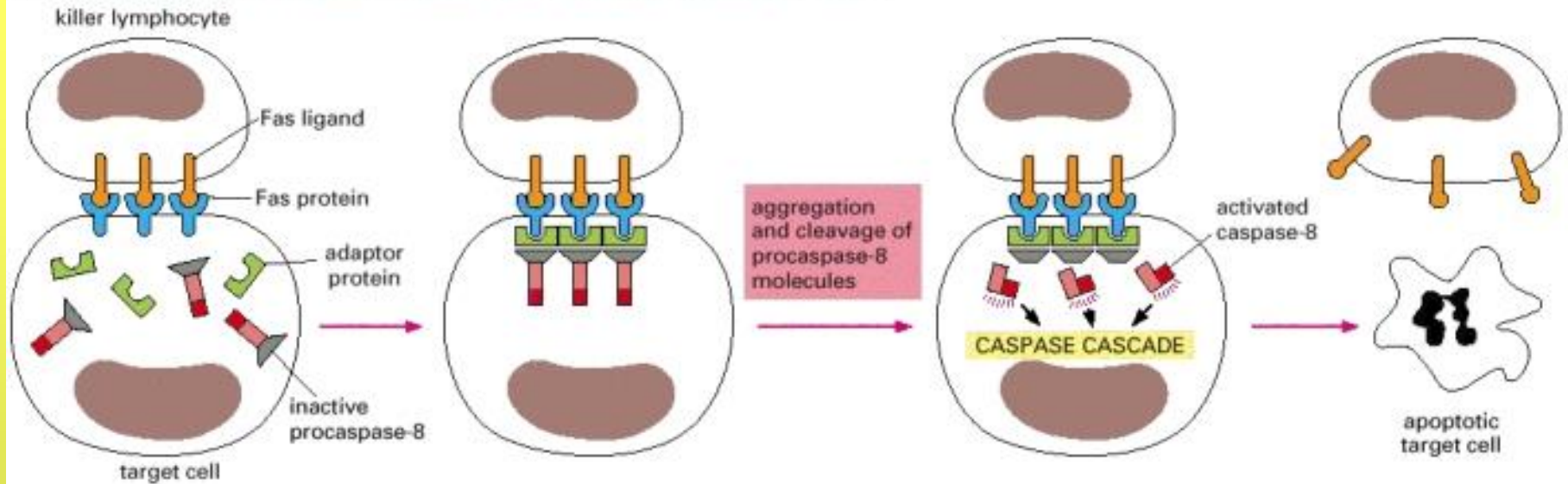
disruption of plasma membrane and organelles, release of the cell content into extracellular space → **inflammation**



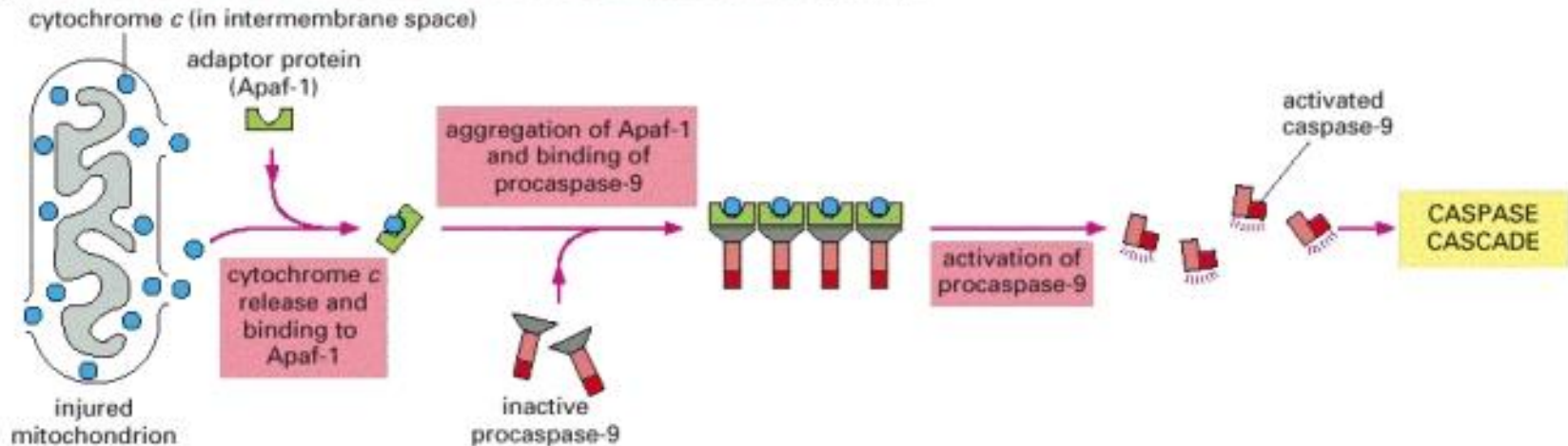
Caspases

are a family of cysteine proteases that play essential roles in apoptosis (programmed cell death), necrosis, and inflammation

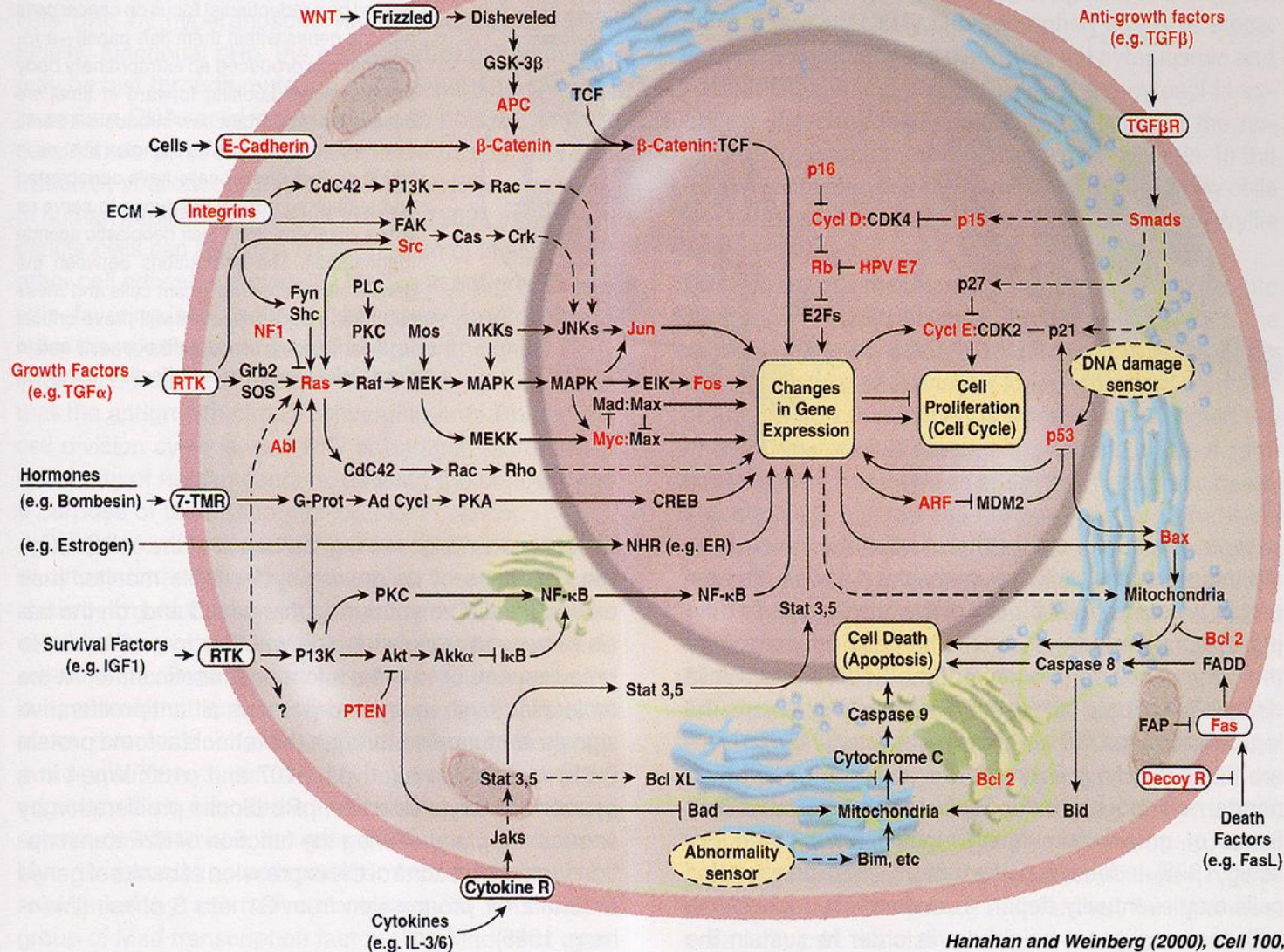
(A) **ACTIVATION OF APOPTOSIS FROM OUTSIDE THE CELL (EXTRINSIC PATHWAY)**



(B) **ACTIVATION OF APOPTOSIS FROM INSIDE THE CELL (INTRINSIC PATHWAY)**



Signal transduction pathways – Proliferation, apoptosis



**Thank you
for your attention**

Campbell, Neil A., Reece, Jane B., Cain Michael L., Jackson, Robert B., Minorsky, Peter V., **Biology**, Benjamin-Cummings Publishing Company, 1996 – 2010.



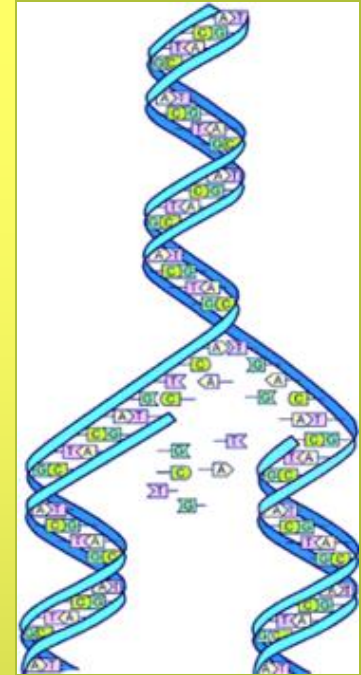
DNA replication

- **Semiconservative** = synthesis of the new strand to the old one

- **Replication begins simultaneously at multiple sites (in Eukaryotes)**

at the beginning of S phase – single fragments
at the end of S phase – accumulation of double fragments

- **Asynchronous replication:** some parts are replicated earlier: constitutional genes (permanently active), euchromatin, other later: heterochromatin



- **Antiparalel strand** is replicated in the form of fragments (**Okazaki fragments**)
- **Initiation** of replication: **RNA primer** – DNA polymerase can not start synthesis from free nucleotides; it can only add nucleotides to the 3' end of a preexisting polynucleotide

Enzymes required for replication :

- **DNA polymerase** (DNA-dependent DNA-polymerase) - polymerization
- **primase** (DNA-dependent RNA-polymerase) – primer synthesis
- **helicase** – unwinding of the DNA double helix
- **gyrase** – unfolding and folding of high order spiralizations of chromatine fibre
- **ligase** – connecting fragments