

Virtual Biomedical and STEM/STEAM Education

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PÉCSI TUDOMÁNYEGYETEM UNIVERSITY OF PÉCS

U. PORTO







The structure of eukaryotic cells

György Sétáló, MD, PhD Gergely Berta, MD, PhD Oktávia Tarjányi, MD, PhD Krisztián Tanics









>>>> The cell membrane and biological membranes of intracellular organelles

- Phospholipid bilayer with embedded proteins (ratios approx.50-50%)
- fluid mosaic model
- glycoproteins and glycolipids contain carbohydrate chains
- membranes allow selective transport, separate but also connect to ' environment (via receptors involved in signaling)





- The largest organelle
- Its shape can be round, flat, rod or a string
- Size: 5-10 µm
- It stores most of the cell's DNA, i.e. the genetic material
- It is the cell's control center: gene expression \rightarrow cellular structure and function
- It is the site of DNA replication, transcription (= RNA synthesis) and RNA processing



- Consists of an outer and inner membrane + perinuclear space
- Abounds in nuclear pore complexes: intensive transport (e.g. protein import, RNA export)
- Nuclear lamina: a protein layer lining the inner surface of the inner nuclear membrane



- Chromatin: DNA + proteins + RNA + ions
- Histone and nonhistone proteins bind to DNA
 - > Euchromatin: looser structure, transcription takes place in it
 - Heterochromatin: more condensed structure, inactive regarding transcription
 - Nucleolus: manufactures ribosomal subunits
 - Nucleoplasm: the inner substance of the nucleus



• An extensive membrane system connected to the outer nuclear membrane





 Rough endoplasmic reticulum (RER)
Smooth endoplasmic reticulum (SER)







There are bound ribosomes on its surface Functions:

- > Protein synthesis, for various organelles:
 - o RER
 - Golgi apparatus
 - o cell membrane
 - o lysosomes
 - o secretory proteins
- > Protein processing (maturation):
 - Folding: formation and stabilization of conformation with the help of chaperone proteins
 - Disulfide bond formation
 - Chemical modifications e.g. glycosylation (addition of carbohydrates)
- > Quality control of proteins



>>>> Smooth ER

- There are no ribosomes on its surface
- Its structure is not as well-ordered as that of the RER
- Functions:
 - > Synthesis of lipids (e.g. phospholipids, steroids)
 - ➤ Ca²⁺ storage
 - Detoxification: biotransformation of foreign molecules
- In muscle: sarcoplasmic reticulum (a special SER): Ca²⁺ release from it \rightarrow muscle contraction









Golgi apparatus

- is the next station of intracellular protein transport (along the "secretory pathway")
- It is composed of flat compartments (cisternae)
 - cis-Golgi
 - median-Golgi
 - trans-Golgi

Functions:

- protein maturation (e.g. glycosylation)
- sorting, and packaging of proteins into transport vesicles (by the trans-Golgi)

- The transport of materials (proteins, lipids, liquids, etc.) between the RER, Golgi apparatus, and SER is performed via transport vesicles (membrane-enclosed sacs)
- Along 2 major pathways:

endocytic pathway

endocytosis: materials are taken up from the outside of the cell by a membrane invagination process

- o phagocytosis (e.g. leukocytes engulf bacteria)
- pinocytosis (liquid uptake, "cell drinking")
- o receptor-mediated endocytosis (e.g. uptake of LDL particles)

Endocytosis \rightarrow endosome \rightarrow merging with primary lysosomes \rightarrow secondary lysosome (or alternatively storage, exocytosis)

Secretory pathway → exocytosis (secretion): transport of materials into the extracellular space: RER → transport vesicle → Golgi-apparatus → transport (secretory-) vesicle → exocytosis







- Organelles of degradation and digestion. A single biomembrane surrounds a special content:
 - primary lysosomes: low pH (approx. 5), degradative enzymes (acid hydrolases). They originate form the trans-Golgi by budding.
 - secondary lysosomes: formed when primary lysosomes are fused to other vesicles. They contain materials which are being degraded (and are usually bigger in size).
- Degradation of molecules from different sources:
 - taken up from the cell's surroundings (by endocytosis, phagocytosis): heterophagy
 - > intracellular molecules: e.g. worn out/damaged organelles (autophagy)







Lysosome





- Secondary bonds are formed between protein monomers during a polymerization process → filaments
- Roles:
 - ➤ determines cell shape
 - provides mechanical support
 - cell movement, shape alterations
 - chromosome movement during cell division
 - intracellular transport (movement of vesicles)
 - roles in intracellular signaling
 - 3 types: microfilaments,
 - intermediate filaments and
 - microtubules



- The organizing center of microtubules
- composed of 2 centrioles
- in a centriole, 9 microtubule triplets form a cylinder
- microtubules are small tubes, composed of 13 paralel protofilaments of tubulin proteins



microtubule



https://people.maths.bris.ac.uk/~matbl/images/mousefibroblasts.j



Mitochondrion

- Size ~ 0,5-1 µm
- Wide range in numbers: 1 thousands of mitochondria /cell
- Role: "power plant of the cell", has a major role in ATP synthesis
- Cristae can show a laminar or tubular structure







>>> The structure of mitochondria

- Outer membrane: highly permeable
- Therefore, the Intermembrane space has a composition similar to that of the cytosol
- Inner membrane: highly impermeable, it contains important proteins:
 - transport proteins (e.g. H⁺/pyruvate symporter, ADP/ATP antiporter) necessary due to the high degree of impermeability
 - Respiratory chain (electron transport chain) proteins
 - > ATP synthase

cristae are to increase the surface area

• **Matrix**: contains a lot of enzymes and also the mitochondrial DNA



>>>> The function of mitochondria

• ATP synthesis through aerobic cellular respiration, in the breakdown process of glucose:

$1 C_6 H_{12}O_6 = 6 CO_2 + 6 H_2O$ and 36 ATP

- 3 phases:
 - Glycolysis in the cytoplasm
 - > Citric acid cycle in the mitochondrial matrix
 - Terminal oxidation along the inner mitochondrial membrane

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Glycolysis

- Glucose (with 6 carbons) \rightarrow 2 pyruvates (with 3 carbons each)
- In the absence of oxygen: e.g. lactic acid (lactate) is produced (fermentation, anaerobic cellular respiration) Glucos
- A small amount of ATP molecules, and reduced coenzymes (NADH) are also produced
- In the mitochondrial matrix, pyruvate is turned into acetyl groups (2 carbons), which are bound to Coenzyme A (CoA)



- Also called Szentgyörgyi-Krebs or citrate cycle
- Oxaloacetate (4 Catoms) + acetyl group (2 C-atoms) → citric acid (6 C-atoms)
- products:
 - $> 2 CO_2$
 - ≻GTP
 - Reduced coenzymes: NADH, FADH₂ (they contain high energy electrons)



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Terminal oxidation

- NADH, FADH₂→electrons → respiratory chain (electron transport chain) proteins
- Respiratory chain members $I. \rightarrow II. \rightarrow III. \rightarrow IV.$
- Cytochromes participate: they contain iron (Fe²⁺ or Fe³⁺)
- Respiratory chain proteins transport protons from the matrix into the intermembrane space
- This maintains a H⁺ concentration difference: electrochemical proton gradient
- The energy of this gradient is harnessed by the ATP synthase protein in the inner membrane: protons are streaming back into the matrix, whilst ADP + P \rightarrow ATP
- Peter D. Mitchell: chemiosmosis theory









Summary: breakdown of glucose and other nutrients with the help of mitochondria



Mitochondrial DNA

• Small, circular

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- Codes for: mRNA, tRNA, rRNA molecules of mitochondria
- But most mitochondrial proteins are imported from the cytosol (synthesized on free ribosomes)
- The presence of DNA is supportive of endosymbiosis:
- Mitochondria derived from ancient prokaryotes



Endosymbiosis theory



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