

The Respiratory Chain

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BIO TEAM 429

بسم الله الرحمن الرحيم

RESPIRATORY BLOCK

The Respiratory Chain

إعداد الطالبات

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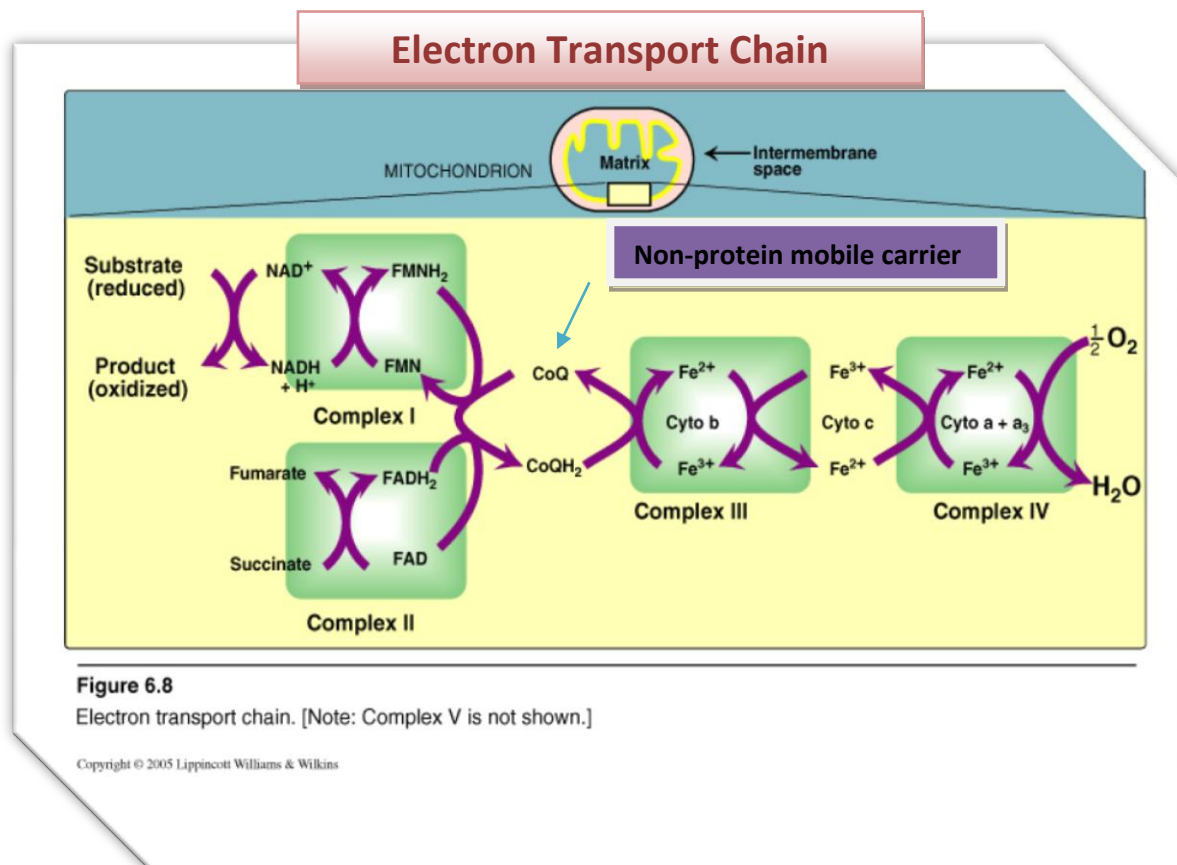
رهام الحناكي

دعواتكم لنا بالتوفيق في الدارين

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- **Energy is produced by** metabolizing energy – rich molecules
 - Ex. Glucose is metabolized by a series of oxidation reaction \longrightarrow $\text{CO}_2 + \text{H}_2\text{O} + \text{electrons}$
- The electrons are donated to specific co enzymes \longrightarrow $\text{NAD}^+ + \text{FAD}$
- The coenzymes are reduced to energy rich forms \longrightarrow $\text{NADH} + \text{FADH}_2$
- NADH and FADH_2 donate a pair of electrons to a specialized set of electron carriers called:
Electron transport chain
- The electrons lose their free energy which is stored with $\text{ADP} + \text{P}_i$ (**inorganic phosphate**) to form **ATP. HOW?**
- Like this: $\text{ADP} + \text{P}_i + \text{energy} \longrightarrow \text{ATP}$ (**is oxidative phosphorylation**)
- The remainder of the energy not used in ATP formation**is used to generate heat and Ca^{+2} transport into mitochondria (drive ancillary reactions)**



Electron transport and ATP synthesis are tightly coupled processes

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❖ Mitochondria :

1) **Outer membrane:** contains special pores
Freely permeable to most ions and small molecules

2) **Inner membrane:** highly convoluted to increase surface area → cristae

- Unusually rich in proteins → half are directly used in electron transport and oxidative phosphorylation

❖ Need specialized carriers or transport systems for impermeable:

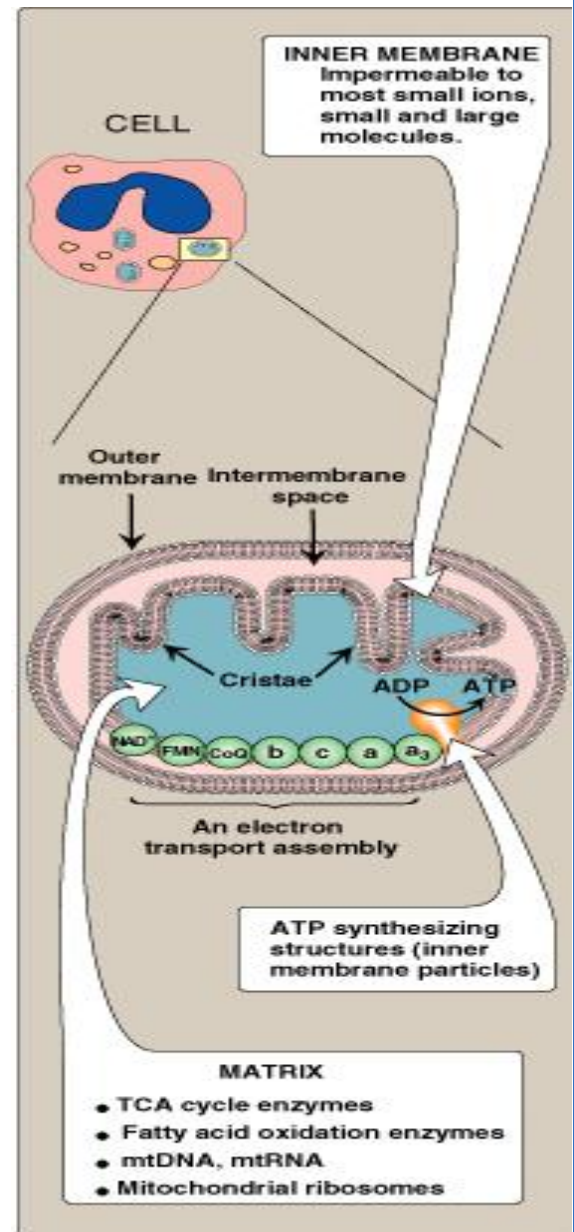
- Ions: H^+ , Na^+ , K^+
- Small molecules: ATP, ADP, pyruvate, metabolites for mitochondrial function.

❖ **ATP synthase complexes:** protein complexes containing domains

- protrude into the matrix and span entire membrane

3) **matrix:** get-like solution → 50% is protein

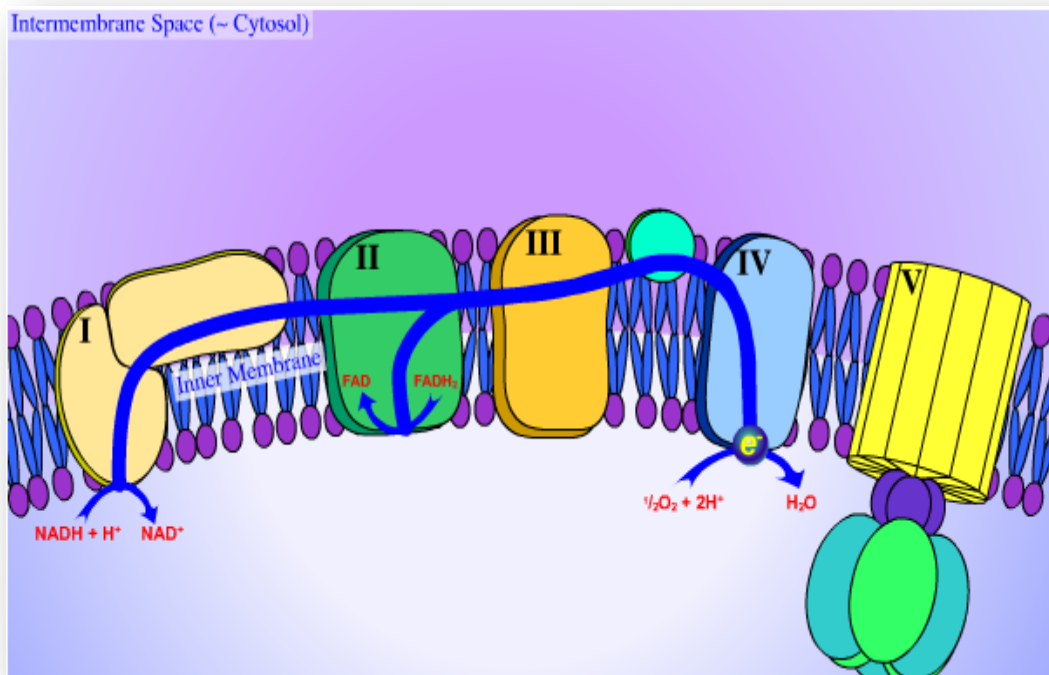
1. Enzymes: oxidation of pyruvate, amino acids, fatty acids responsible for TCA cycle
2. Oxidized coenzymes → hydrogen receptors (NAD^+ and FAD)
3. ADP and P_i
4. Mitochondrial DNA and RNA and ribosome synthesis of glucose, urea and heme



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Electron Transport Chain



- made of enzyme complexes : I, II, III, IV on the inner mitochondria membrane
- each complex donates electrons to **mobile carriers**
- carrier **receives** electrons from an electron donor and **donates them to** the next complex producing energy slowly
- **also called the** respiratory chain **because it** requires O_2
- electrons combine with O_2 and protein to form **water**
- all members of the chain are protein **except** coenzyme Q

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❖ reactions:

1. **formation of NADH** : NAD^+ is reduce to NADH
 - **Enzyme**: dehydrogenases
 - **reaction**: 2 electrons + 1 proton (from substrate) + NAD^+ \longrightarrow $\text{NADH} + \text{H}^+$
2. **Complex I**: receives NADH, most reduce complex
 - **Enzyme**: NADH dehydrogenase \longrightarrow has a tightly bound coenzyme contains several iron – sulfur centers \longrightarrow necessary for transfer of H to coenzyme Q
 - **Coenzyme**: FMN \longrightarrow flavin mononucleotide
 $\text{FMNH}_2 \longrightarrow$ when it accepts ($2 \text{ e}^- + 2 \text{ H}^+$)
3. **Complex II**: receives FADH_2 , dose nit produce H^+
 - **Enzyme**: succinate dehydrogenase
4. **Coenzyme Q**: a quinine derivative with hydrophobic tail accepts H atom from FMNH_2 and FADH_2 links both flavoproteins to the cytochromes
5. **Complex III**: cytochromes b and a, receives electrons from coenzyme Q
 - **Cytochromes**: is a heme group + iron atom
 - Iron atom is reversibly converted from Fe^{3+} to Fe^{2+} reversible electron carrier
6. **Cytochromes c**
7. **Complex IV**: cytochromes a + a_3 or cytochromes oxidase
 - The only electron carrier which molecular oxygen can react directly with the heme iron
 - Contain bound copper for the following reaction to occur :
 $2 \text{ electron} + \text{molecular O}_2 + \text{free portions} \longrightarrow \text{H}_2\text{O}$
 - **Enzymes**: the free energy is released as electrons
 - Electrons are transferred as **hydride ions** (H^-) to NAD^+ or as hydrogen atom to FMN, coenzyme Q and FAD or as electrons to cytochromes
 - **Free energy**: is used for phosphorylation of ADP to ATP
 - The excess is used for the reaction or produced as heat
 - $\text{NADH} \longrightarrow 3 \text{ ATP}$ (pump 3 H^+ from complex I, III, IV)
 - $\text{FADH}_2 \longrightarrow 3 \text{ ATP}$ (pump 2 H^+ from complex III, IV)

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Phosphorylation:

- 1) **Proton pump** : electron transport pumps protons (H^+) across the inner mitochondrial membrane to the intermembrane space
 - Causes an electrical gradient (outside \rightarrow more positive)
 - Causes a PH gradient (outside \rightarrow lower PH)
 - The proton gradient generates energy to drive ATP synthesis
 - A common intermediate that couples oxidation to phosphorylation.
- 2) **ATP synthase**: complex V Has two domains
 - ❖ **F₀**:
 - membrane spanning Domain
 - The protons reenter the matrix Through F₀ and rotate it.
 - ❖ **F₁** :
 - Extra membranous domain

✓ **Note** : electron movements :

Electron transport chain \rightarrow proton pump \rightarrow crosses inner mitochondria membrane

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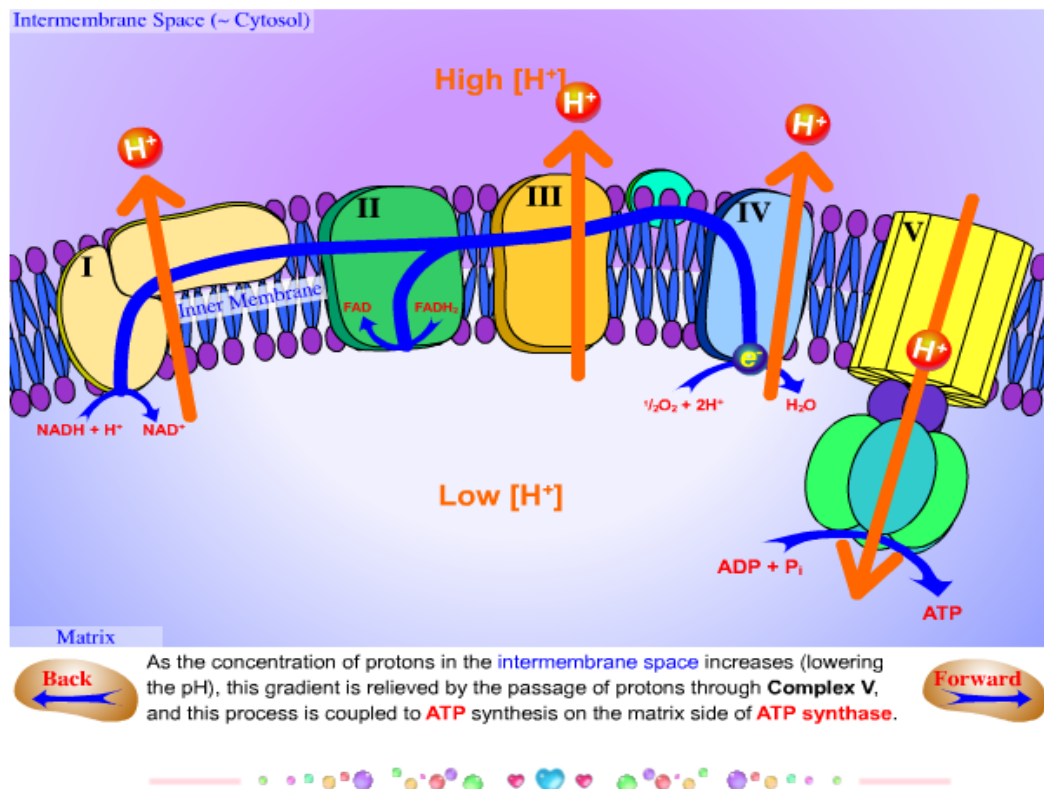
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Matrix \leftarrow F₁ \leftarrow F_n of complex V \leftarrow Electrical gradient and PH \leftarrow inter membrane space

- Undergoes conformational Changes after F₀ is rotated
 - Catalytic activity is then activated
 - ATP synthesis from ADP + Pi
 - dissipating the PH and electrical gradients

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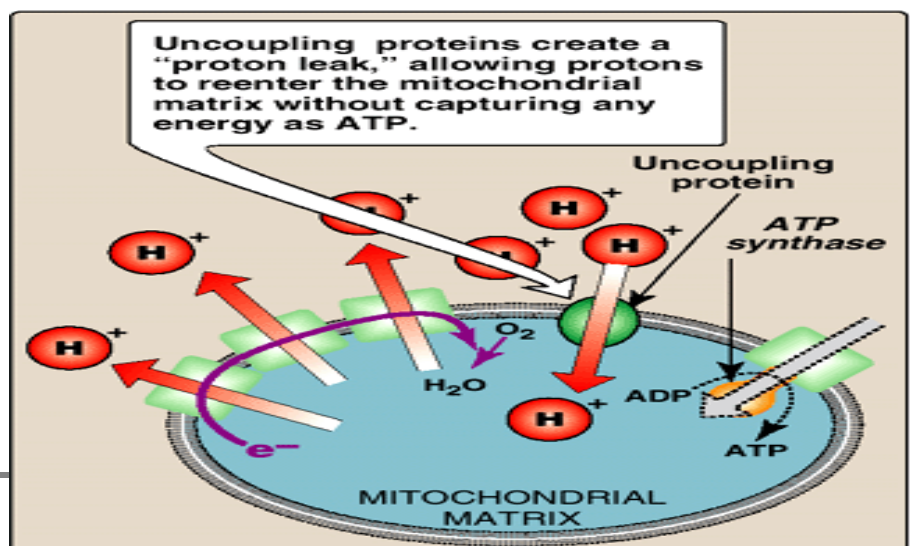
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Factors affecting phosphorylation and oxidation:

a) Respiratory control:

- Oligomycin:** drug that binds to F0 domain
 - Closes H^+ demands and prevents reentry into the matrix
 - Prevents reentry of protons into the matrix and phosphorylation
 - PH and electrical gradient cannot be dissipated
 - Difficulty pumping protons against the steep gradient prevent oxidation
- Decreased availability of ADP or P_i**



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b) Uncoupling proteins (UCP) :

- **Location:** inner mitochondrial membrane
- **Function:** the proteins create a “proton leak”
Allow protons to reenter the matrix without energy being captured as ATP.
- **UCP1:** found in brown adipocytes of mammals (**humans and animals**)
 - Function: activation FA oxidation and heat production
 - 90% of respiratory energy for thermogenesis (**unlike white fat**)
 - Responds to: cold, at birth, arousal in hibernating animals
 - In humans: little brown fat except infants
 - Doesn't play a major role in energy balance
- ✓ UCP2 and UCP3 are found but unknown significance

c) Synthetic uncouplers :

- Increase the permeability of inner mitochondrial membrane to protons
- Uncouple oxidative phosphorylation by readily diffusing
- Causes electron transport to proceed rapidly without a proton gradient
- Energy is released as heat rather than being used to synthesize ATP (**ATP is not synthesized**)

Ex:

- 1) 2,4 – dinitrophenol : a lipophilic protein carrier .
- 2) **Toxic dose of aspirin** and **salicylates** : accompanied by fever .

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MEMBRANE TRANSPORT SYSTEM :

A) **ADP**: cytosol → adenine nucleotide carrier → inner mitochondrial membrane

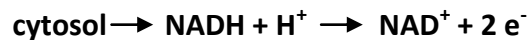
ATP: mitochondria → adenine nucleotide carrier → cytosol

P_i: cytosol → phosphate carrier → mitochondria

b) **Reducing equivalents**:

NADH: inner mitochondrial membrane lacks an NADH transport protein passes indirectly by **two different shuttles**.

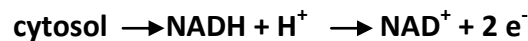
1) Glycerophosphate shuttle :



inner mitochondrial membrane → 2e^- + Glycerophosphate degeneration

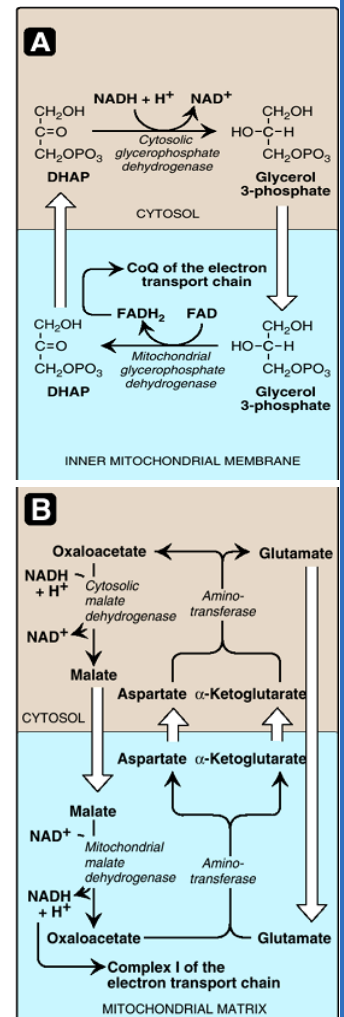
produces : $\text{FADH}_2 \rightarrow \text{COQ} \rightarrow 2\text{ATP}$ / cytosolic NADH oxidized .

2) Malate – Aspartate shuttle :



matrix → 2e^- + Malate degeneration

produces : $\text{NADH} \rightarrow 3 \text{ATP}$ / cytosolic NADH oxidized



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Apoptosis :programmed cell death.

Pores are formed on the **outer** membrane of mitochondria

Cytochrome c leave through the pores to the intermembrane space then cytosol.

In the cytosol: Cytochrome c + proapoptotic factors → activates a family of proteolytic enzyme (the caspases) → cleavage of key protein → morphological and biochemical change → Apoptotic cell death.

- **-Inherited defect in oxidative phosphorylation :**

Mt DNA : codes polypeptides required for oxidative phosphorylation

Polypeptides are synthesized in the mitochondria .

Mutation rate is **10 times** higher than nuclear DNA

Mutation:- defects in oxidative phosphorylation affects tissues with high ATP requirement

Ex: CNS , skeletal and heart muscle , kidney ,liver

Disease :- **mitochondrial myopathy** – **leber hereditary optic neuropathy** : neuroretinal degeneration

Damage to the optic nerve – bilateral loss of central vision .

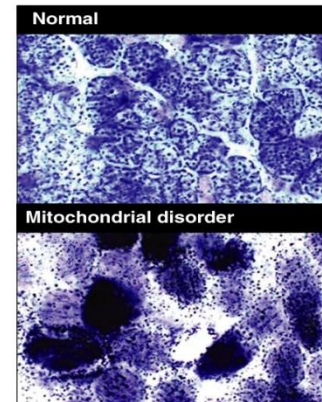


Figure 6.16
Muscle fibers from a patient with a mitochondrial myopathy show abnormal mitochondrial proliferation when stained for succinic dehydrogenase.

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