

Krebs Cycle

TCA cycle

Citric acid cycle

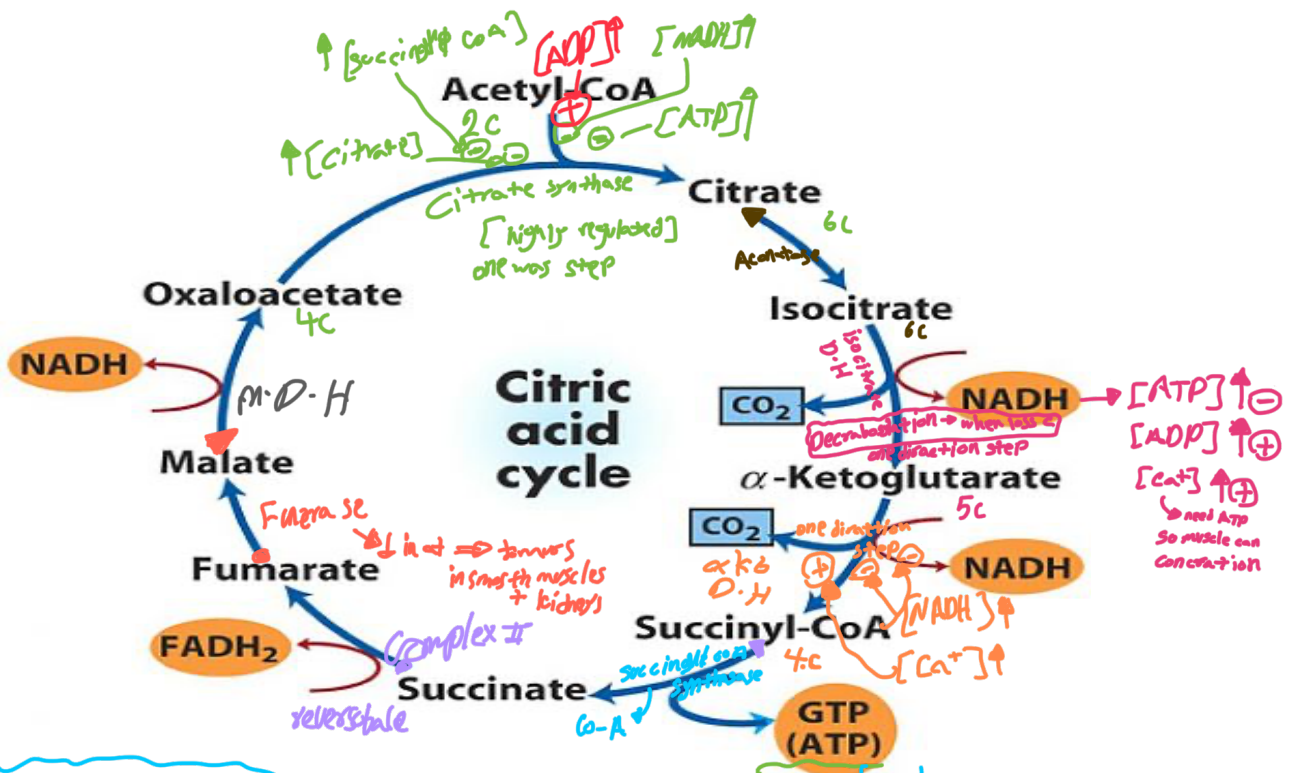
TCA Cycle

1. Acetyl-CoA donates its acetyl group to the four-carbon compound oxaloacetate to form the six-carbon citrate. Citrate is then transformed into isocitrate, also a six-carbon molecule
2. dehydrogenated with loss of CO₂ to yield the five-carbon compound Alpha-ketoglutarate
3. alpha-Ketoglutarate undergoes loss of a second molecule of CO₂ and ultimately yields the four-carbon compound succinate.
4. Succinate is then enzymatically converted in three steps into the four-carbon oxaloacetate—which is then ready to react with another molecule of acetyl-CoA.



The tricarboxylic acid (TCA) cycle

- Krebs cycle
- The citric acid cycle
- The enzymes of the TCA cycle are located in the **mitochondrial matrix**
- Fuel Molecules lose electrons, get oxidized and donate those electrons to energy carriers producing NADH and FADH₂
- **Main function of the TCA cycle is to release high-energy electrons that power the synthesis of ATP via oxidative phosphorylation**



(cofactor)
 α-k-G → histone deacetylase
 → going to remove ethyl groups
 → going to bind and not allow
 → to happen which uncontrolled growth ⇒ died

- Acetyl-CoA adds two carbons to oxaloacetate to start the cycle.
- Isomerization takes place by removing H₂O and then adding it back.
- A CO₂ is lost and a NADH is produced
- Another CO₂ is lost and another NADH is produced.
- A substrate-level phosphorylation. ATP
- FAD⁺ is reduced to form FADH₂
- another NADH is produced

Each turn of the cycle forms one turn

• 1 GTP

+ 2 CO₂

• 3 NADH molecules

• 1 FADH₂ molecule

• Note: oxygen is not required until later

• Oxygen is required to recycle

• NADH \square NAD⁺

• FADH₂ \square FAD

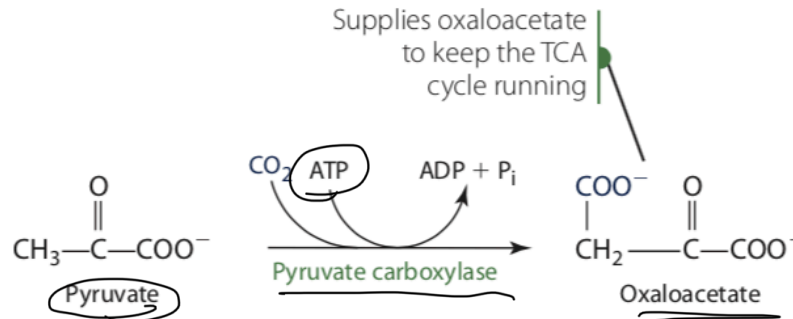
• (Thus generating ATP)

• Cofactors must be recycled to be reused in the Krebs Cycle

- The role of the citric acid cycle not confined to the oxidation of acetate only
- The citric acid cycle is a “hub of intermediary metabolism”
- Four- and five-carbon end products of many catabolic processes feed into the cycle and serve as fuels.
- ^{4c}Oxaloacetate and ^{5c}alpha ketoglutarate are produced from ^{A.A}aspartate and ^{D.A.A}glutamate
- Under some metabolic circumstances, drawn out of the cycle to serve as precursors of the amino acids aspartate and glutamate by simple transamination reactions
- Oxaloacetate is converted to glucose in gluconeogenesis → by a reaction converting it to pyruvate
- Succinyl- CoA is a central intermediate in the synthesis of the porphyrin ring of heme groups, which serve as oxygen carriers (in carriers' (in hemoglobin and myoglobin
- As intermediates of the citric acid cycle are removed to serve as biosynthetic precursors, they are replenished by anaplerotic reactions
- The most important anaplerotic reaction in mammalian liver and kidney is the reversible carboxylation of pyruvate by CO₂ to form oxaloacetate
- Pyruvate carboxylase is a regulatory enzyme and is virtually inactive in the absence of acetyl-CoA.

- To keep the TCA cycle functioning, oxaloacetate—and other TCA cycle intermediates that can lead to oxaloacetate— must be **replenished** in the cycle.
- Oxaloacetate, fumarate, succinyl-CoA, and a-ketoglutarate can all be formed from certain amino acids
- We need CO_2 (from amino acids)

High levels of acetyl CoA activate pyruvate Carboxylase



Requires Biotin

Anaplastic reaction

Figure 3.22 Formation of oxaloacetate from pyruvate and CO_2 .

Regulation of the Citric Acid Cycle

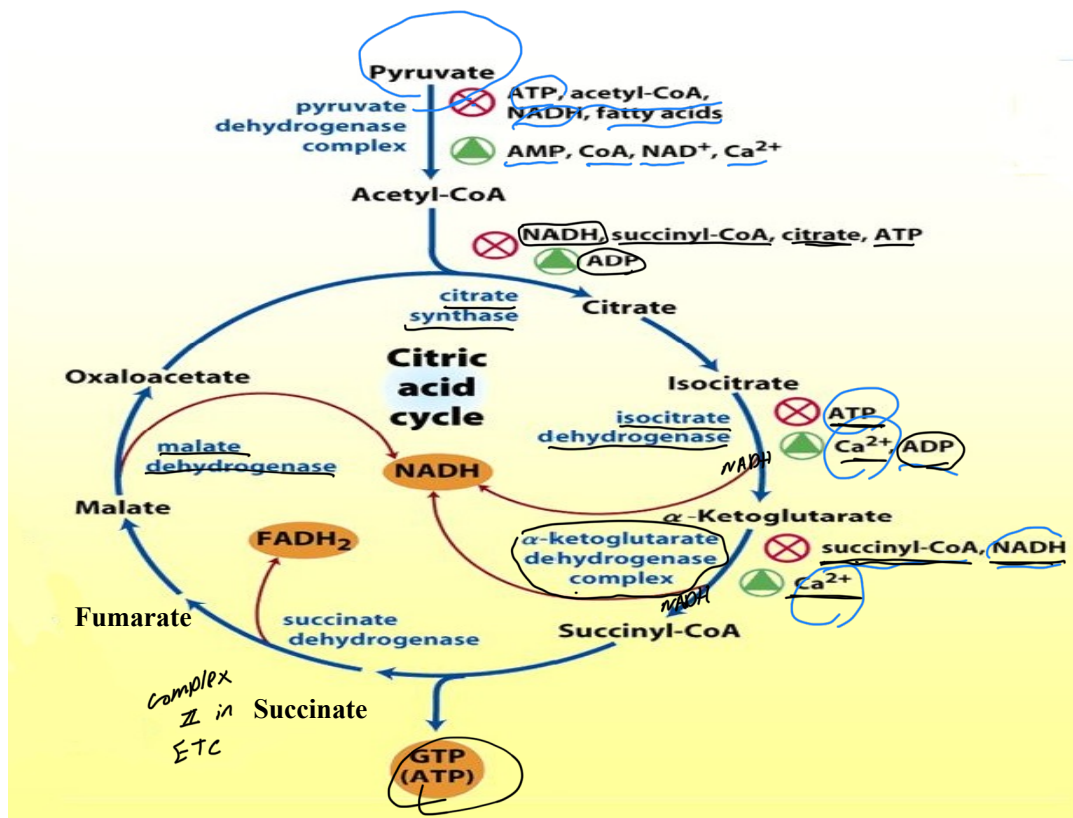
- No hormonal regulation
- Allosteric regulation only (molecules in cell bind to enzymes part of the TCA cycle, not on active site (on a different allosteric site), causing enzyme to undergo a **conformational change**, either making enzyme work better or be inhibited)
- Substrate availability

Example

- Low concentration of acetyl CoA will slow down citric acid cycle
- Citrate being shuttled out of citric acid cycle for FA synthesis \square Slow down cycle $\Rightarrow \downarrow \square \square$
- In starvation state, amino acids break down and enter citric acid cycle as alpha ketoglutarate or oxaloacetate \square more substrate \square speeding cycle

Allosteric regulation

- The PDH complex (Pyruvate → acetyl CoA)
 - Inhibited by fatty acids and acetyl-CoA and when the cell's [ATP]/[ADP] and [NADH]/[NAD⁺] ratios are high
 - Regulatory protein enzymes- The PDH complex is inhibited by reversible phosphorylation of E1 complex (protein kinase and protein phosphatase)
- **The Citric Acid Cycle Is Regulated at Its Three Exergonic Steps**
 - Citrate synthase
 - Isocitrate dehydrogenase
 - alpha-ketoglutarate dehydrogenase



Formation of ATP

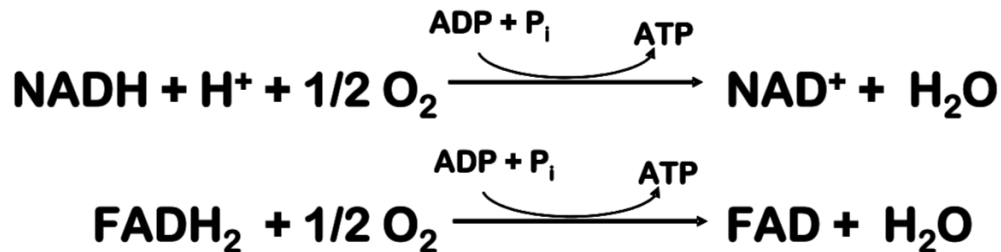
- Acts as the main energy currency and must be continually synthesized from the energy provided by macronutrients
 - Some ATP are synthesized by direct phosphorylation involving high-energy phosphate donors, referred to as **substrate-level phosphorylation**
 - Two reactions in glycolysis
 - one reaction in the TCA cycle produce GTP
- The production of ATP in mitochondria by **oxidative phosphorylation** (*Electron transport chain*)

Oxidative phosphorylation is the process of making ATP by using the proton gradient generated by the ETC.

- The production of ATP in mitochondria by oxidative phosphorylation begins with the oxidation of fuel molecules by the TCA cycle and the release of electrons and protons.
- Electrons obtained from nutrients and metabolic intermediates are transferred to NAD⁺ and FAD
- $AH_2 + NAD^+ \rightarrow A + NADH + H^+$ **TCA CYCLE AND**
- $BH_2 + FAD \rightarrow B + FADH_2$ **GLYCOLYSIS**
- The electrons and protons are captured by NADH and FADH₂ and delivered to the inner mitochondrial membrane.

NAD⁺ and FAD must be recycled

Recycling is accomplished by oxidation and transfer of electrons to oxygen.



NAD⁺ and FAD are then available for additional oxidative metabolism. The energy released during electron transport is coupled to ATP synthesis.

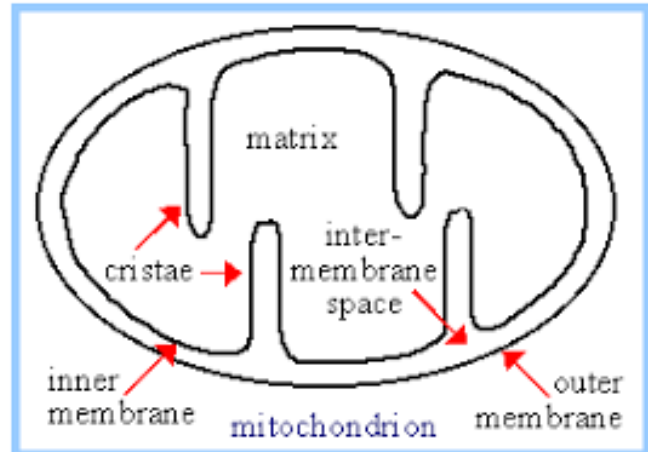
The electrons and protons are captured by NADH and FADH₂ and delivered to the inner mitochondrial membrane.

Location of mitochondrial complexes

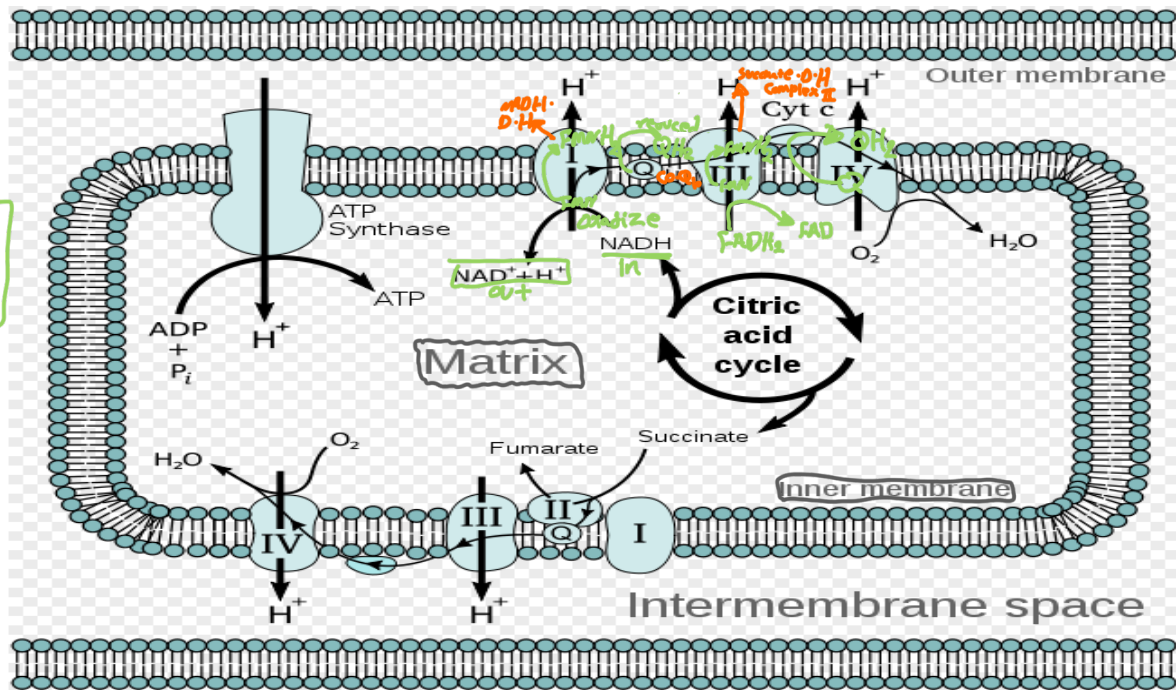
• Inner mitochondrial membrane:

Electron transport chain:
oxidizes reduced coenzymes

ATP synthase: machinery to synthesize ATP



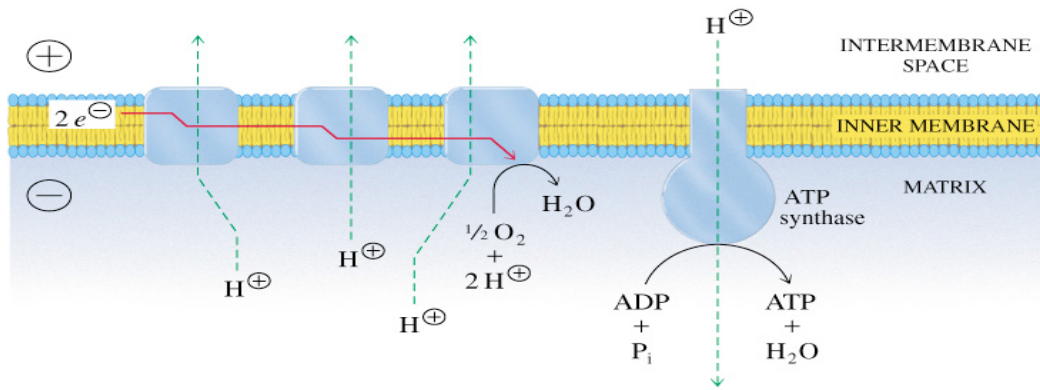
OF CAC
6 NADH
2 FADH₂



when complex I be in ↑ Energy levels going to pass the e⁻, and H⁺ going to intermembrane space.
 Complex I be in ↑ E levels due to 2 e⁻ + H⁺ don't have a pump to pump H⁺ going to pass e⁻ to Q.

Complex II → Fe²⁺ → Fe³⁺ → Fe³⁺ → Fe²⁺ going to activate the e⁻ → Fe³⁺ → Fe²⁺ → activate a pump [H⁺]
 Cyt-C → Fe²⁺ → Fe³⁺
 Complex III → Fe²⁺ → Fe³⁺ → Fe²⁺ → Fe³⁺ → there going to be O₂ → e⁻ → 2H⁺ → H₂O
 ↳ final e⁻ acceptor

- The flow of electrons from NADH or FADH₂ to O₂ through protein complexes located in the mitochondrial inner membrane leads to the pumping of protons out of the mitochondrial matrix. The resulting uneven distribution of protons generates a pH gradient and a transmembrane electrical potential that creates a proton-motive force
- ATP is synthesized when protons flow back to the mitochondrial matrix through an enzyme complex (ATP synthase). Thus, the oxidation of fuels and the phosphorylation of ADP are coupled by a proton gradient across the inner mitochondrial membrane



- The steps are coupled. Electrons do not flow to oxygen unless ATP is needed
- Each NADH produces 2.5 ATP
- Each FADH₂ produces 1.5 ATP

- **Four protein complexes in the Inner membrane make up the ETC**

- Complexes I, II, III, IV

- Work together in succession to catalyze **redox reactions**

- **Integral membrane proteins with prosthetic groups to move electrons**

- Electrons move through the complexes in order

- Electrons from **NADH** enter at **Complex I**

- Electrons from **FADH₂** enter at **Complex II**

- In each reaction, an **electron donor** is **oxidized** and an **electron acceptor** is **reduced**

- **Compounds differ from one another in how readily they will**
 Compounds differ from one another in how readily they will be oxidized or reduced

- Electrons flow downhill – spontaneously moving from molecules that are strong electron **DONORS** to strong electron **ACCEPTORS** = move from high energy state to low energy state
- Flow of electrons is spontaneous and **thermodynamically** favorable because the next carrier has greater affinity for electrons than the previous
- NADH is a strong electron donor: because its electrons are held in a high-energy linkage, the free-energy change for passing its electrons to many other molecules is favorable
- NADH is a good molecule for donating electrons to the respiratory chain, while O₂ is a good electron *Acceptor*

Co-factors in Electron Transport

- Protein components use **metal-containing prosthetic groups** or flavins to carry electrons
- Metal-containing groups such as iron-sulfur clusters, copper ions, hemes

Co-factors in Electron Transport

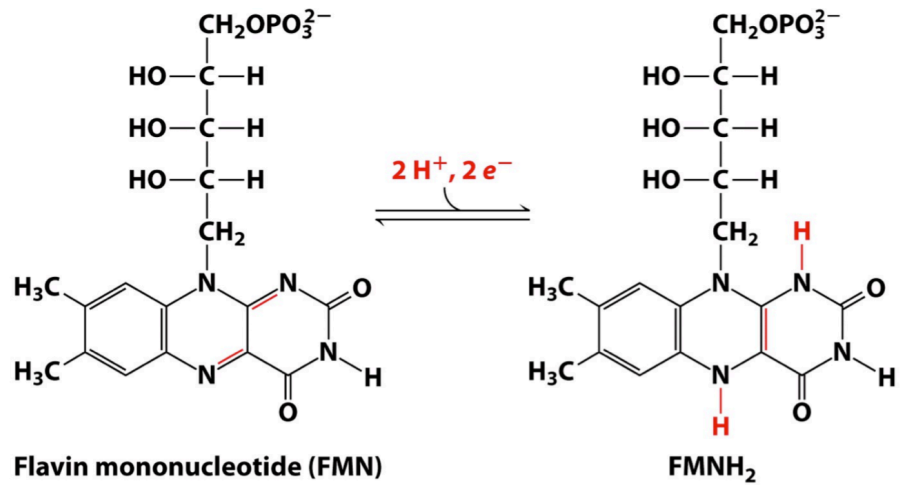
① Flavins:

(Complex I) FMN

- FMNH₂

(Complex II) FAD

- FADH₂

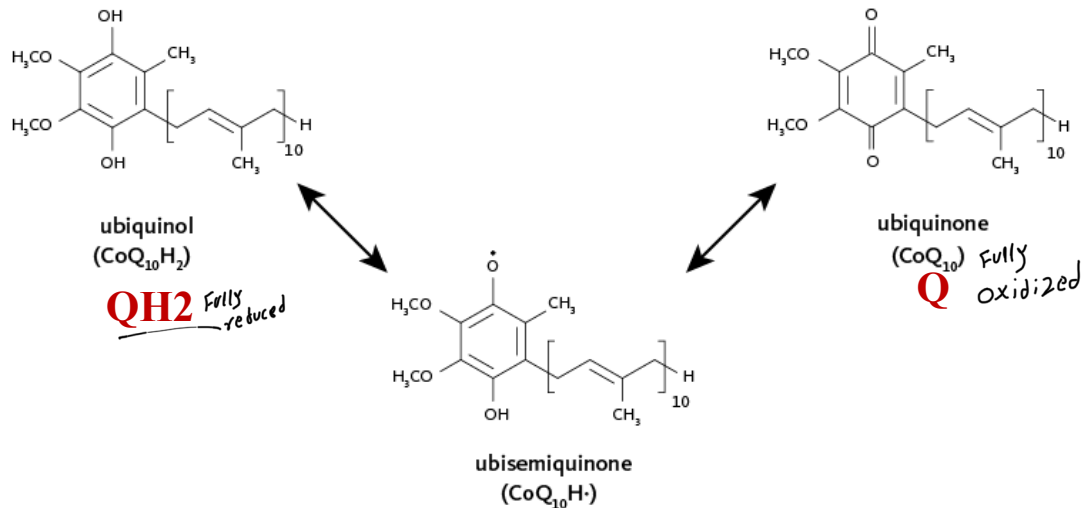


Co-factors in Electron Transport

② Ubiquinone (Q) (Mobile electron carrier)

- Also called coenzyme Q
- A membrane-soluble compound
- Long hydrophobic tail keeps Q anchored in the mitochondrial inner membrane
- lipid soluble molecule that diffuses within the lipid bilayer, and shuttles electrons from Complexes I and II and pass them to III
- Not a part of any complex

Figure 1. The Different Redox Forms of Coenzyme Q₁₀



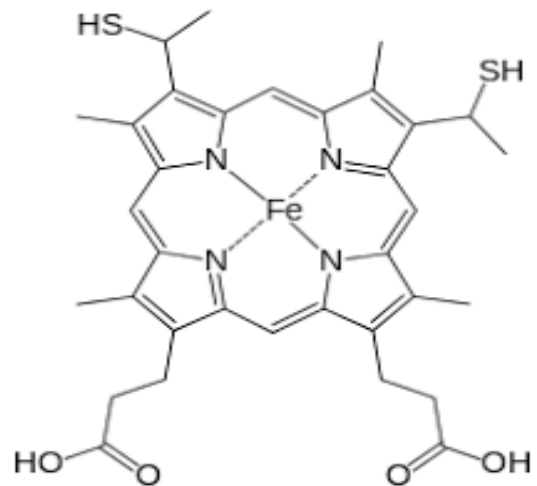
Coenzyme Q₁₀ exists in three oxidation states: the fully reduced ubiquinol form (CoQ₁₀H₂), the radical semiquinone intermediate (CoQ₁₀H•), and the fully oxidized ubiquinone form (CoQ₁₀).

<https://lpi.oregonstate.edu/book/export/html/352>

Co-factors in Electron Transport

③ Cytochrome c

- A peripheral membrane protein associated with the outer face of the membrane, transports electrons from III to IV
- Not a part of any complex
- Shuttles electrons and protons from Complex III to Complex IV

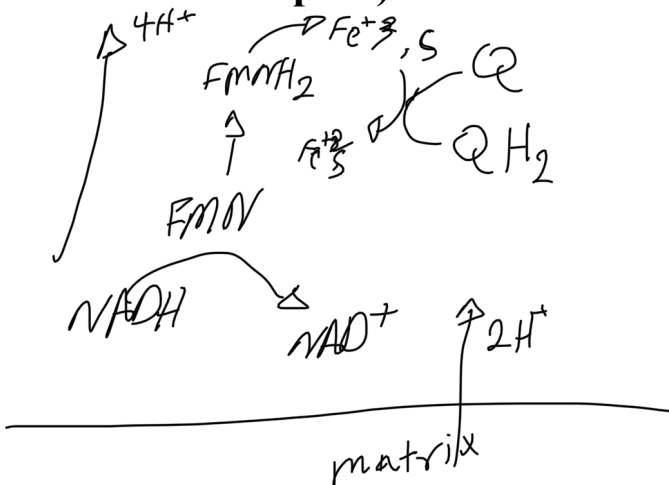


MIDTERM

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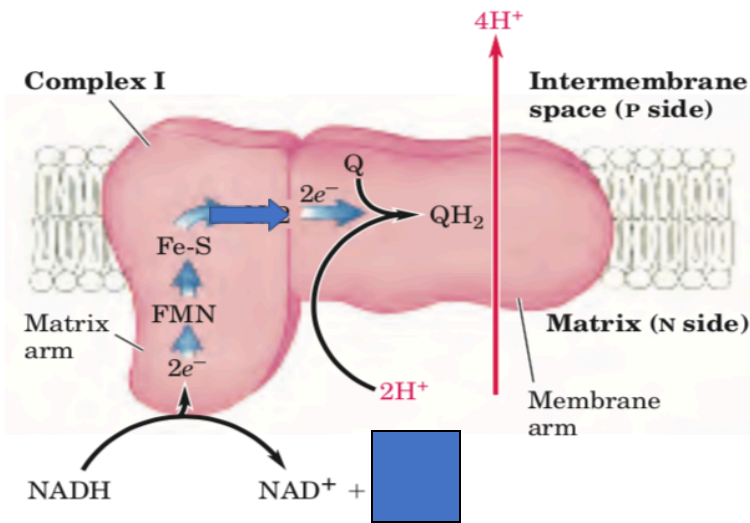
Complex I

- Complex I includes a **flavoprotein** (contains FMN – related to FAD) and proteins with **Fe-S centers (iron-sulfur clusters)**
- Electrons flow from NADH to Flavin mononucleotide (FMN)
- Electrons then flow to a prosthetic group on an iron sulfur cluster
- Iron cycles between 3+ and 2+ states
- Reduction of Q to QH₂ requires 2 e⁻
- **About 4 H⁺ translocated per 2 e⁻ transferred (pumped from matrix to intermembrane space)**

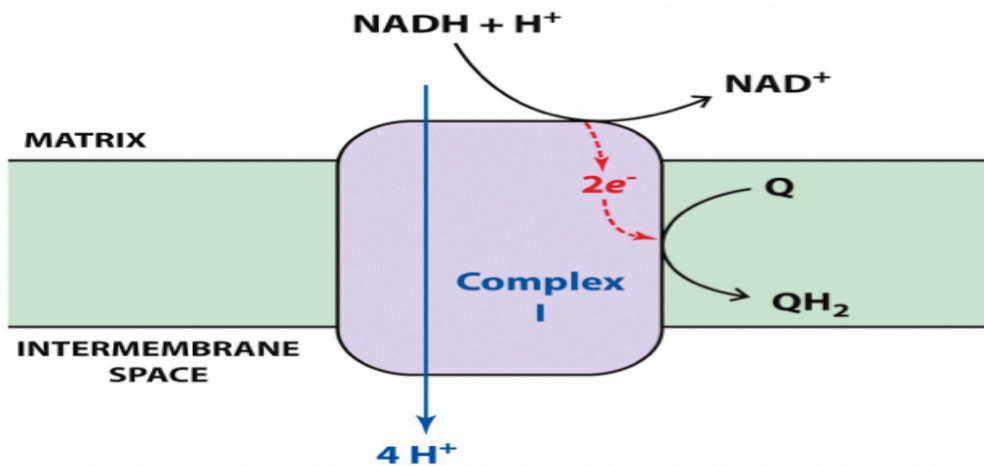


Complex 1

NADH dehydrogenase, is a very large, L-shaped structure that functions to accept high energy electrons from NADH molecules.



NADH binds to the vertical component of complex (within the matrix)
NADH gives off two electrons onto an acceptor molecule called flavin mononucleotide (FMN)
the FMN also uptakes two hydrogen ions (one from the matrix and one from the NADH molecule) to form the fully reduced FMNH₂.

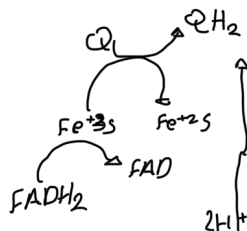


The electrons then move through a series of iron-sulfur clusters (Iron cycles between +2 and +3 end up being transferred onto ubiquinone.(Q) Q uptakes two protons from the matrix to form ubiquinol (QH₂)

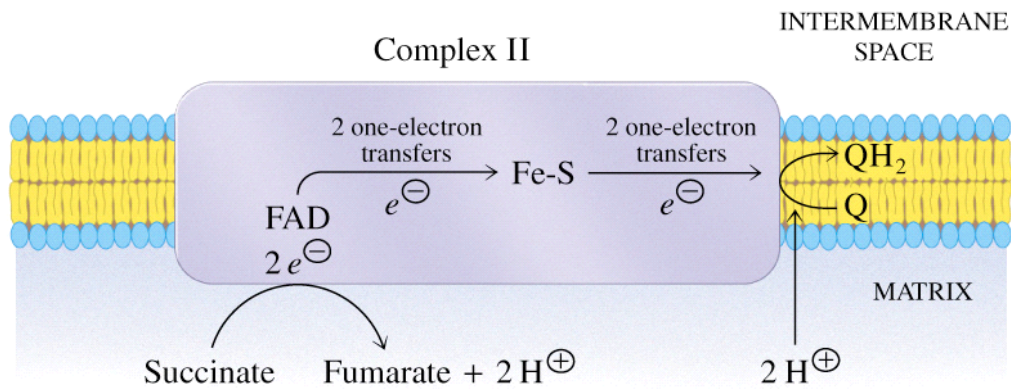
- The transfer of electrons to CoQ occurs one electron at a time.
- The Q molecule is lipid soluble and freely moves through the hydrophobic core of the membrane. Once it is reduced, (QH₂), ubiquinone delivers its electrons to the next complex in the electron transport chain.

Complex II

- **Contains succinate dehydrogenase**, a component of the TCA cycle
- Complex II contains a FAD protein and Fe-S centers
- succinate is converted to fumarate in the TCA cycle, FAD is reduced to FADH₂.
- FADH₂ (produced in kreb cycle) **remains bound to complex**
- Complex II can oxidize the FADH₂ back into FAD and move the free electrons through a series of iron-sulfur clusters and onto ubiquinone, thereby forming ubiquinol (The FADH₂ is oxidized by electron transfer through the Fe-S centers to reduce Q to QH₂)



no H⁺ pumps.



Complex II does **NOT** contribute to **proton gradient**, but supplies electrons from succinate. Less ATP generated from FADH₂ compared to NADH

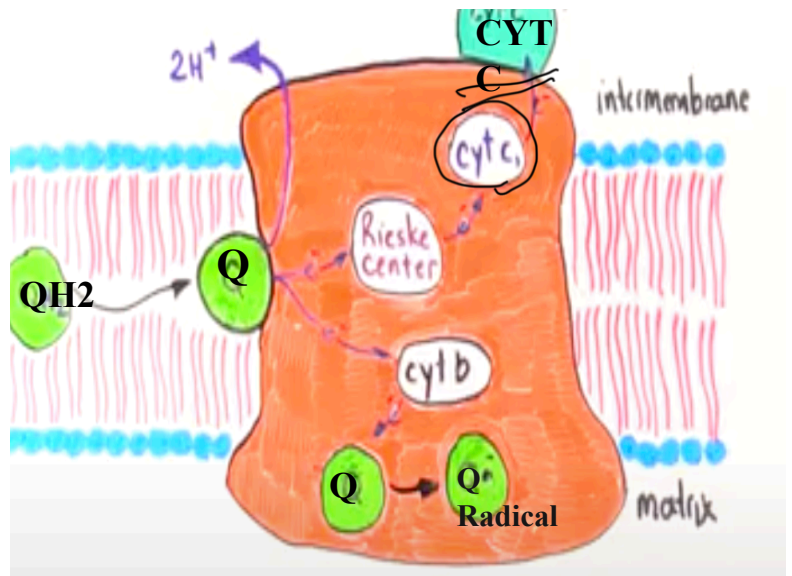
Electron transfer and proton flow in Complex III

- Electron transfer from Ubiquinol (QH₂) to cytochrome *c*
- The electrons derived from both NADH and FADH₂ are passed from QH₂ to cytochrome *c* through the reactions of Complex III
- Complex III contains two different cytochromes and a Rieske center that contains the 2Fe-2S center group
 - **Cytochrome b** (two heme groups)
 - **Cytochrome c** (single heme group, carries one electron)
 - **Rieske center** that contains the 2Fe-2S center group

Complex III

* $\text{cyt } c_1$ and $\text{cyt } c$ is different

- The process by which the electrons are transferred from the ubiquinol to cytochrome c is known as the **Q cycle** \square **Two half cycles**
- In the first half of the cycle, one electron flows to cytochrome c . A second electron flows to cytochrome b where it transfers to a Q molecule, creating an unstable semiquinone ion.



- One of these electrons moves onto the Rieske center, then onto cytochrome c_1 and finally onto cytochrome c . Note that cytochrome c , unlike ubiquinone, can only carry a single electron at any given time.
- The other electron that comes from QH₂ follows a different pathway and moves through the heme groups of cytochrome b and onto ubiquinone to form a partially reduced species called a semiquinone radical ion. The two protons that were originally attached to ubiquinol are transferred into the intermembrane space.
- In the second half-cycle of the Q cycle, another ubiquinol (QH₂) attaches onto complex III. Upon binding, the two protons are moved into the intermembrane space and the two electrons follow the same pathways as before. The fourth electron flows to cytochrome b where it joins the semiquinone (Q radical) to form QH₂.

Q cycle half one:

QH₂ attach to complex 3 \rightarrow

can be \leftarrow bind one e^- \rightarrow $\text{cyt } c_1$ \rightarrow $\text{cyt } c$

$2 \rightarrow \text{Fe-S}$

$2 \rightarrow \text{cyt } b$

$Q \rightarrow Q^{\bullet}$
Radical

Complex III

- In the second half-cycle of the Q cycle, another ubiquinol attaches onto complex III. Upon binding, the two protons are moved into the intermembrane space and the two electrons follow the same pathways as before

Summary

A single Q cycle

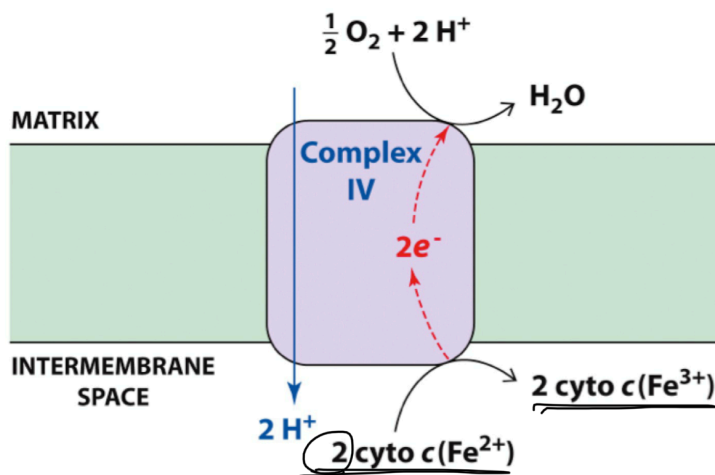
- oxidation of 2 QH₂
- Reduces two cytochrome c molecules
- forms a single ubiquinol molecule
- umps four protons into the intermembrane space

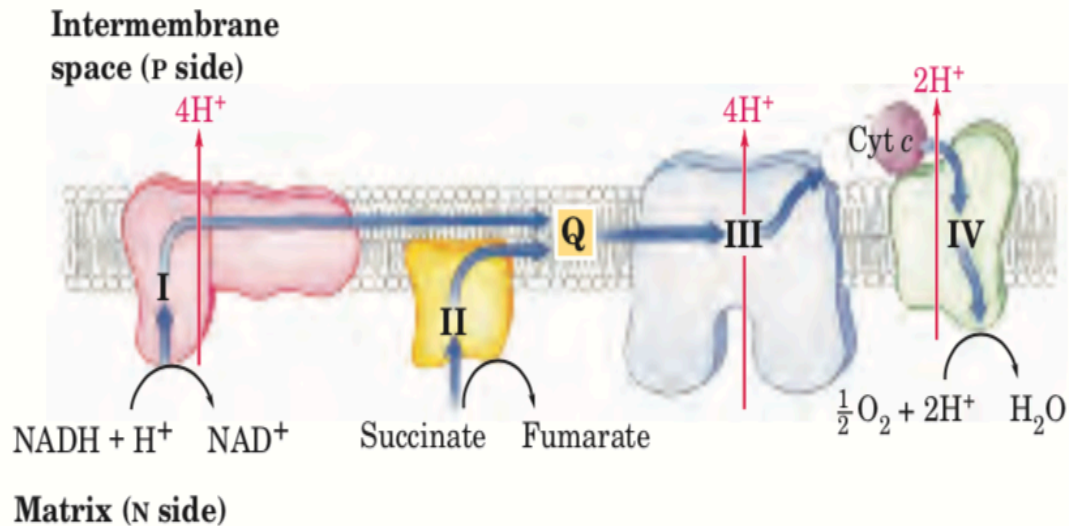
Complex IV

- Accepts electrons from cytochrome c
- Contains **2 cytochromes** (a and a₃) and proteins with copper centers that provide multiple centers for oxidation-reduction
- Cytochromes a and a₃ are the only species capable of direct transfer of electrons to oxygen
- two heme groups (one in each of the cytochromes a and a₃)

- oxygen molecule held very tightly between the iron and copper ions until the oxygen is completely reduced
- Catalyzes reduction of oxygen to form water.
- The metal ions cycle between their oxidized (Fe^{+3} , Cu^{+2}) and reduced (Fe^{+2} , Cu^{+})
- Moves H^{+} into the intermembrane space and contributes to the proton gradient

Complex IV





Oxidative phosphorylation. Electrons from NADH and FADH₂ travel along the electron transport chain a total of **10 protons** are translocated from the matrix to the intermembrane space.

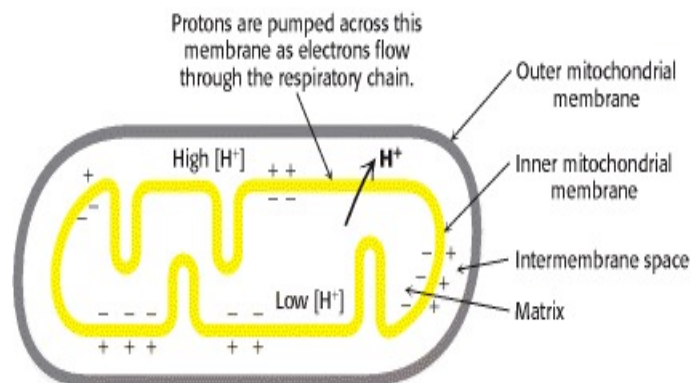
ATP Synthesis

- How is a concentration gradient of protons transformed into ATP?
- **Complex I, Complex III and Complex IV pump protons** across the inner mitochondrial membrane
- Pumping uses the energy liberated from the oxidation of NADH and FADH₂
- Pumping generates a membrane potential because it generates an electrochemical gradient
 - Negative inside, positive outside
 - Alkaline inside, acidic outside

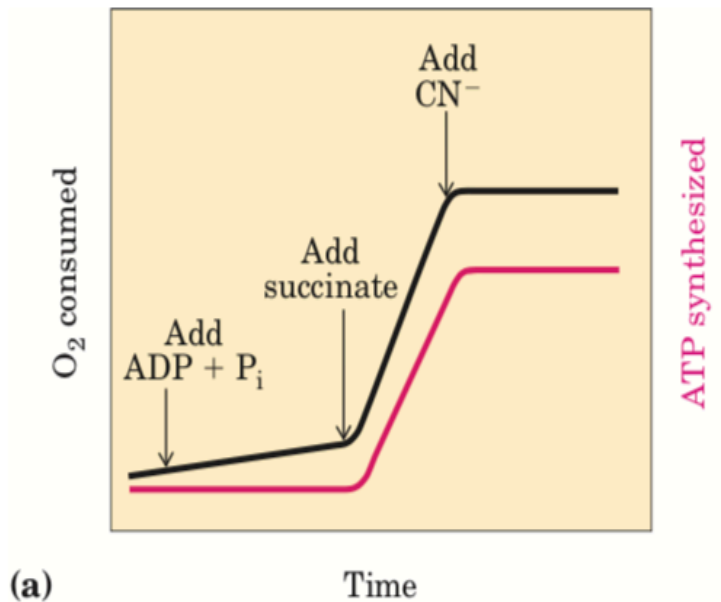
ATP Synthesis

The Chemiosmotic Hypothesis

- A proton concentration gradient serves as the energy reservoir for driving ATP formation
 - Protonmotive force (Δp) is the energy of the proton concentration gradient
 - Protons that are translocated into the intermembrane space by electron transport, flow back into the matrix via ATP synthase
 - H^+ flow forms a circuit (similar to an electrical circuit)
 - The transmembrane protein, **ATP synthase**, catalyzes the phosphorylation of ADP in a reaction driven by movement of H^+ across the inner membrane into the matrix
-
- In chemiosmotic theory transmembrane ATP synthases are very important. They convert energy of spontaneous flow of protons through them into chemical energy of ATP bonds.
 - The proton gradient is **COUPLED** with ATP synthesis

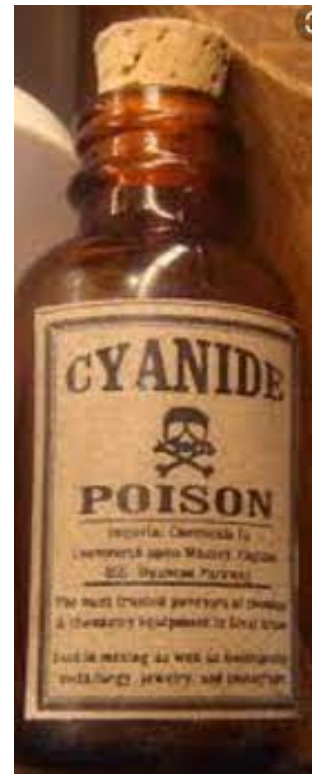


- In experiments to demonstrate coupling, mitochondria are suspended in a buffered medium and an O₂ electrode monitors O₂ consumption (Oxygen disappearance from closed vessel)
- At intervals, samples are removed and assayed for the presence of ATP.
- Addition of ADP and P_i alone results in little or no increase in either respiration (O₂ consumption; black) or ATP synthesis (red). When succinate is added, respiration begins



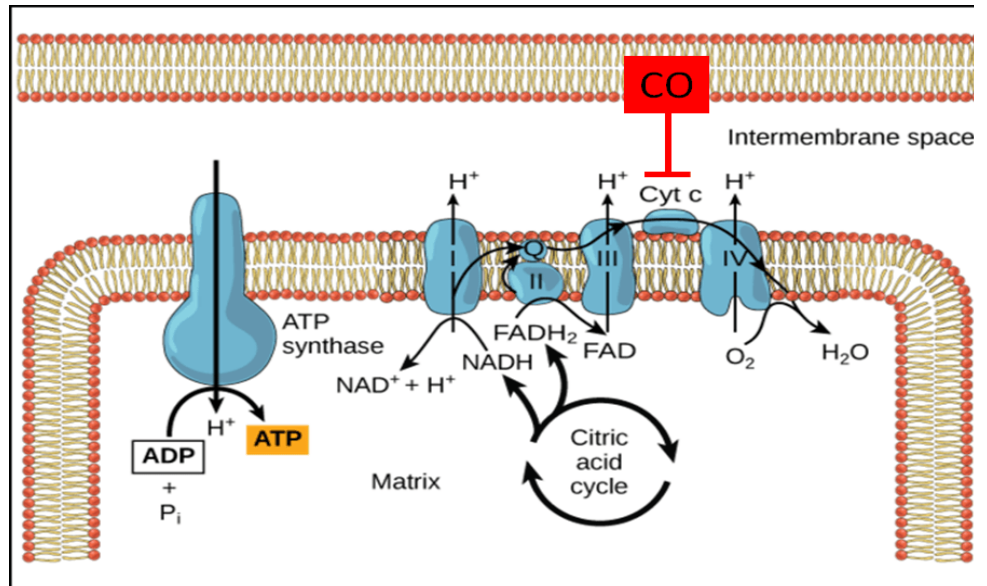
When succinate is added, respiration begins immediately and ATP is synthesized. Addition of cyanide (CN⁻), which blocks electron transfer between cytochrome oxidase and O₂, inhibits both respiration and ATP synthesis.

- Cyanide is a naturally occurring chemical, found in many plants, that has been used in conventional warfare
- Cyanide poisons the mitochondrial electron transport chain within cells and renders the body unable to derive energy (adenosine triphosphate—ATP) from oxygen.



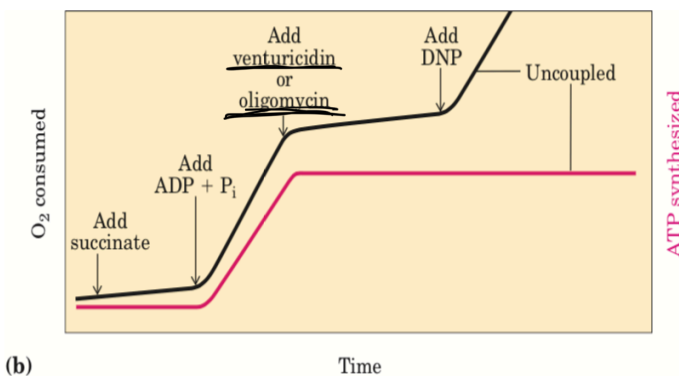
Cyanide poisons the mitochondrial electron transport chain within cells and renders the body unable to derive energy (adenosine triphosphate—ATP) from oxygen. Specifically, it binds to the a₃ portion (complex IV) of cytochrome oxidase and prevents cells from using oxygen, causing rapid death

Carbon Monoxide works similarly to **Cyanide**. **Carbon monoxide** binds to and inhibits **cytochrome c oxidase (complex IV)**.



<https://blogs.brown.edu/emergency-medicine-residency/winter-is-coming-think-co-toxicity/>

Experiments demonstrate that the reverse is true as well! inhibition of ATP synthesis blocks electron transfer in intact mitochondria (providing O₂ and oxidizable substrates, but no ADP) no ATP synthesis can occur and electron transfer to O₂ does not proceed



venturicidin or oligomycin, inhibitors of ATP synthase, blocks both ATP synthesis and respiration

Dinitrophenol (DNP) is an uncoupler, allowing respiration to continue without ATP synthesis.

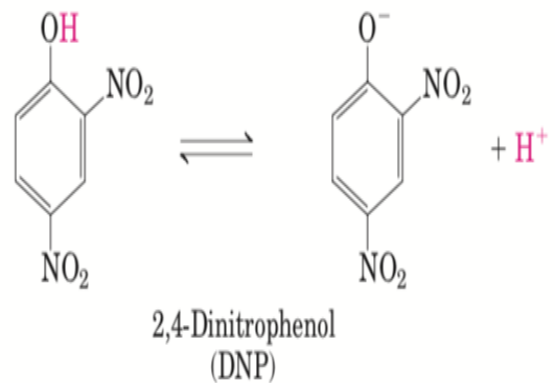
- What would happen to the energy stored in the proton gradient if it weren't used to synthesize ATP or do other cellular work?

- The proton-motive force builds up

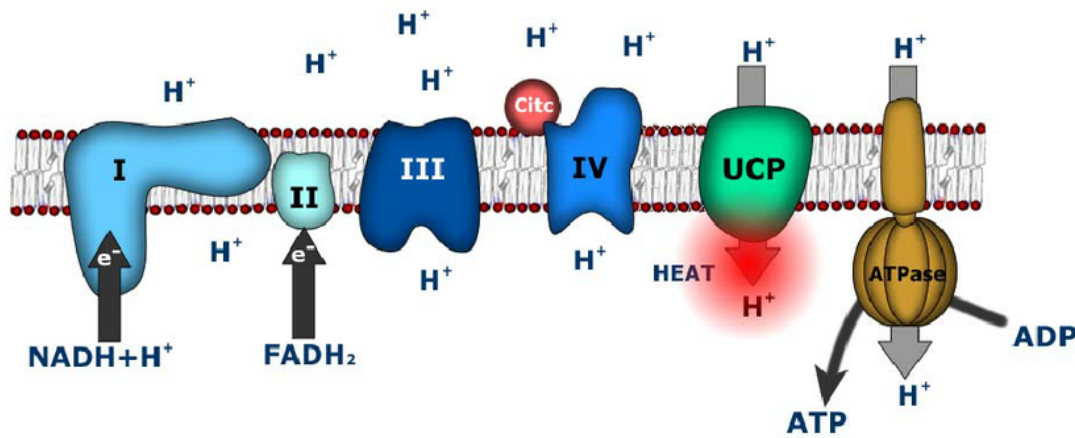
- **Uncoupling agents:** molecules that disrupts oxidative phosphorylation in by dissociating the reactions of ATP synthesis from the electron transport chain

- Mitochondria still capable of catalyzing electron transfer from succinate or NADH to O₂

- O₂ consumed **but no ATP generated**



DNP has dissociable proton/ hydrophobic molecule. Diffuse readily across mitochondrial membranes. After entering the matrix in the protonated form, they can release a proton



Mitochondrial transporters present in the inner membrane of mitochondria

Allow proton-leak : they act as proton channels to redirect protons back into the matrix and away from ATP synthase
dissipation of oxidation energy as heat

Implicated in thermogenesis

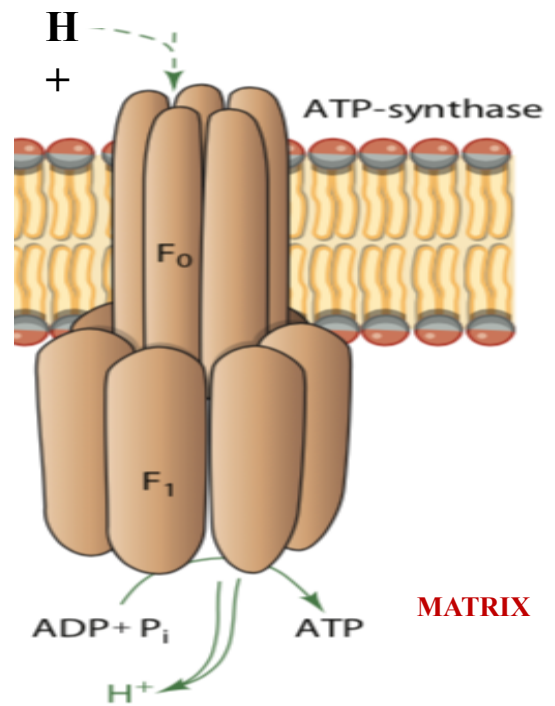
Valle, Adamo & Oliver, Jordi & Roca, Pilar. (2010). Role of Uncoupling Proteins in Cancer. C 567-591. 10.3390/cancers2020567.

Uncoupling Protein-1 (UCP1)

- Uncoupling proteins found in mammalian brown adipose tissue (BAT)
- “Brown fat” **dark color due to high levels of mitochondria which contain thermogenin (uncoupling protein)- produce heat- non shivering thermogenesis**
- In humans and other large mammals, BAT disappears after infancy

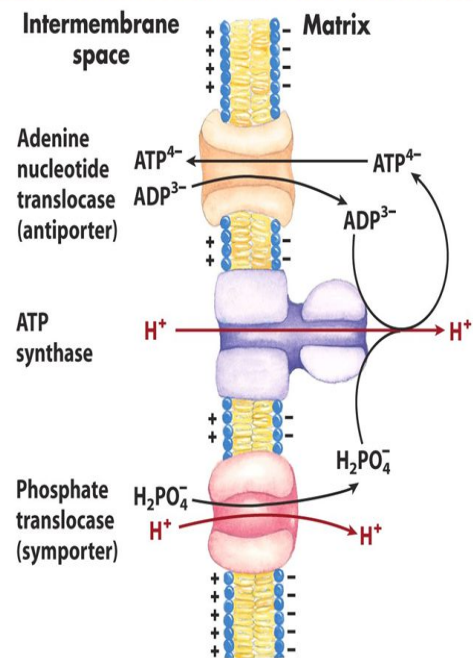
ATP Synthase

- Knob like Projections into the matrix side
- Two units
- F1 Contains the catalytic site for ATP synthesis
- F0 transmembrane channel for H⁺
- Proton flow back into the matrix through (F0) component of the ATP synthase while F1 catalyzes the synthesis of ATP

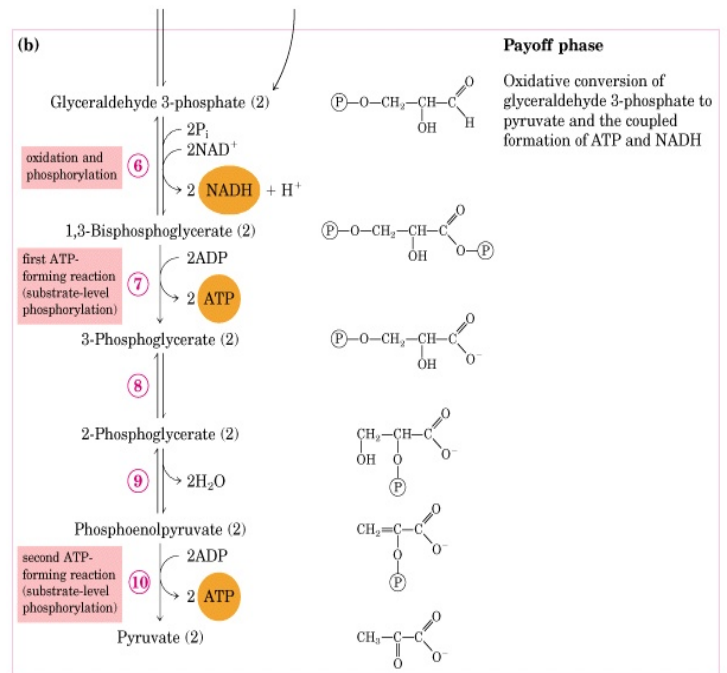


- ATP is synthesized in mitochondria but must be moved to the cytosol to supply energy for the cell. The enzyme *ATP-ADP translocase* shuttles ATP out of the mitochondria and ADP in.
- **ATP must be transported to the cytosol, and ADP and Pi must enter the matrix**
- ADP/ATP carrier exchanges mitochondrial ATP⁴⁻ for cytosolic ADP³⁻
- The exchange causes a net loss of -1 in the matrix (draws some energy from the H⁺ gradient)

Adenine nucleotide and phosphate translocase



- Recall that the glycolytic pathway generates **NADH** in the cytosol in the oxidation of glyceraldehyde 3-phosphate
- **NAD⁺** must be regenerated for glycolysis to continue.
- Under anaerobic conditions, NADH in the cytosol is used in the lactate dehydrogenase reduction of pyruvate to lactate, thereby becoming reoxidized to NAD⁺ without involving oxygen



- NADH cannot simply pass into mitochondria for oxidation by the respiratory chain, because the inner mitochondrial membrane is impermeable to NADH and NAD⁺
- **How is cytosolic NADH reoxidized under aerobic conditions?**

- **Electrons from NADH, rather than NADH itself**, are carried across the mitochondrial membrane
- **Glycerol 3-phosphate shuttle**
- specific to certain tissues. The glycerol-3-phosphate shuttle functions in the brain and skeletal muscle
- NADH in the cytosol transfers its electrons to dihydroxyacetone phosphate, forming glycerol-3-phosphate that freely diffuses across the outer mitochondrial membrane.
- Electrons of glycerol-3-phosphate are then transferred to FAD that is associated with a membrane-bound isoform of glycerol-3-phosphate dehydrogenase located in the inner mitochondrial membrane

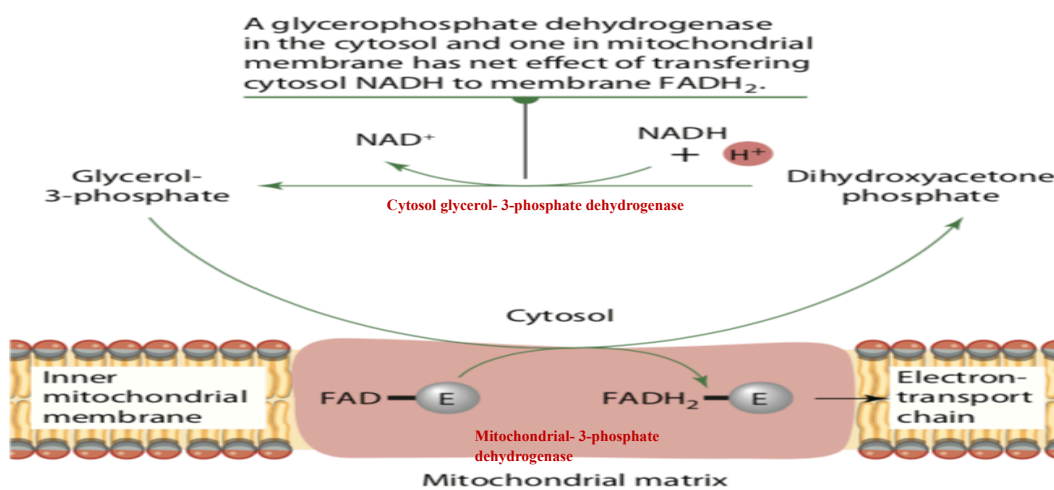
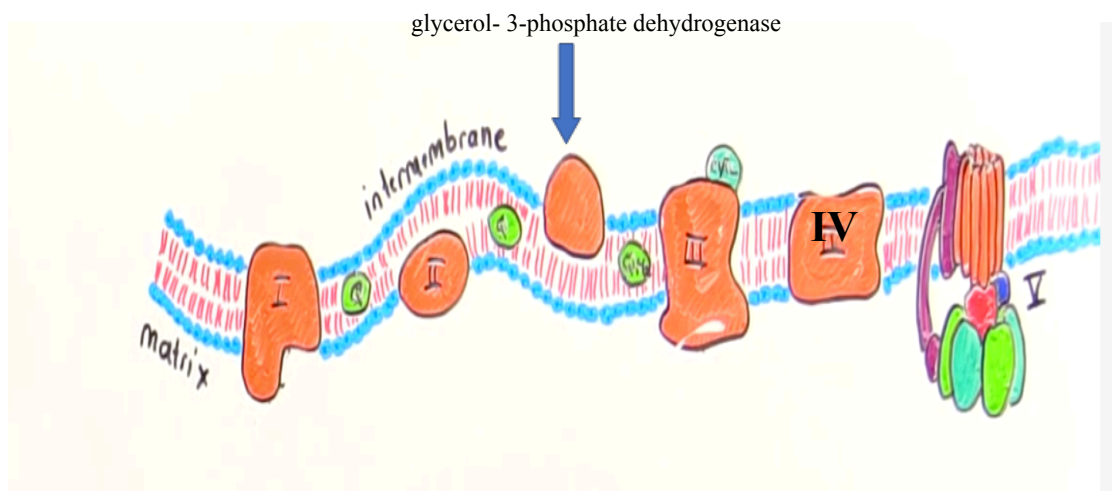


Figure 3.23 Glycerol-3-phosphate shuttle.

The first step in this shuttle is the transfer of a pair of electrons from NADH to dihydroxyacetone phosphate, a glycolytic intermediate, to form glycerol 3-phosphate. This process occurs in the cytoplasm.

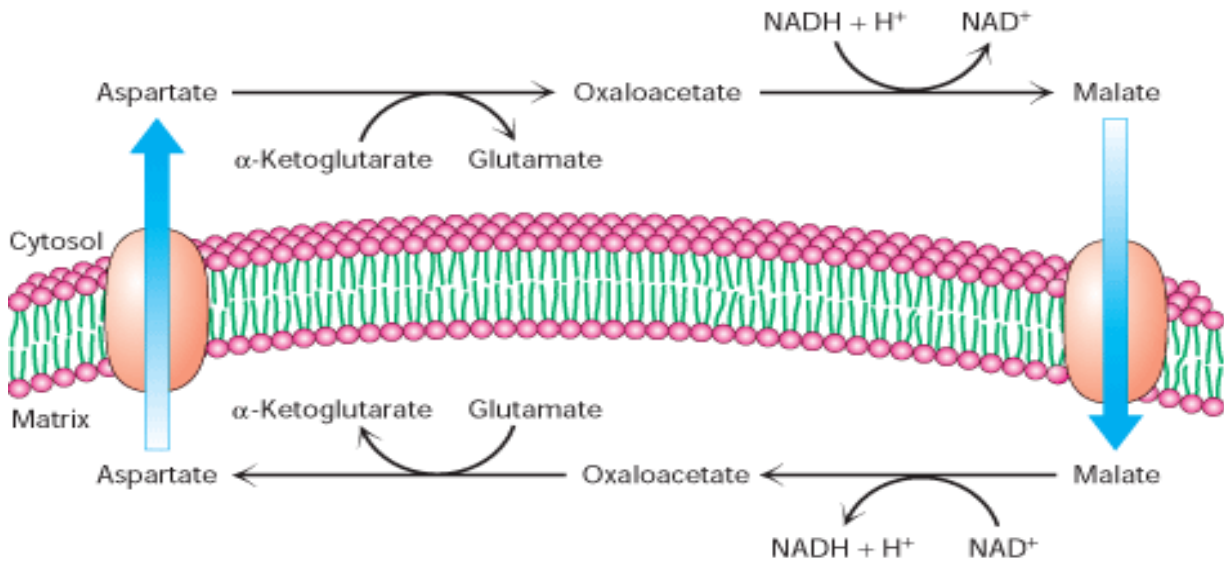
- Once G3P moves into the intermembrane space, it is then oxidized back into **DHAP** by the action of a membrane-bound enzyme called the mitochondrial **glycerol 3-phosphate dehydrogenase**
- An FAD molecule bound to this enzyme accepts the two high energy electrons and the two H⁺ ions to form FADH₂. The FADH₂ then gives up the two protons and electrons to a ubiquinone molecule to form ubiquinol. Ubiquinol then moves along the inner mitochondrial membrane to transfer the electrons onto complex III. Since this shuttle bypasses complex I, NADH molecules transported onto the ETC in this manner only produce a net result of 1.5 ATP molecules per NADH.



NADH molecules transported onto the ETC through the glycerol 3 phosphate shuttle produce 1.5 ATP molecules per NADH

Because bypassing complex 1

Malate-Aspartate Shuttle System



- The most active shuttle compound, malate, is freely permeable to the inner mitochondrial membrane. Oxaloacetate from the cytosol is reduced by the NADH to form malate and NAD⁺. The malate is oxidized by the enzyme malate dehydrogenase to oxaloacetate in the matrix of mitochondria, producing NADH that enters the electron transport chain and generates 2.5 moles of ATP per mole of NADH