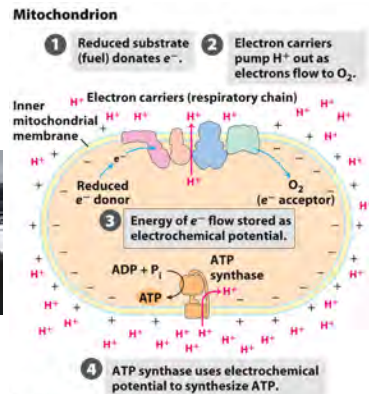


Phosphorylation

Hypothesis: The Respiratory Chain and ATP Synthase Produce ATP by a Chemiosmotic Mechanism



Peter Mitchell, 1920-1992



Hypothesis: The Respiratory Chain Produce ATP by a substrate-level-phosphorylation mechanism

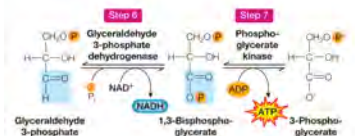


Efraim Racker, 1913-1991

The search was for a compound, like 1,3 bisphosphoglycerate, that would be used by the ATP Synthase to make ATP.

Substrate-Level Phosphorylation

Glycolysis



Phosphorylation

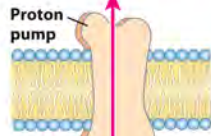
Chemiosmotic Theory

- $ADP + P_i \rightarrow ATP$ is **highly thermodynamically unfavorable**.
- How do we make it possible?
- Phosphorylation of ADP is **not** a result of a direct reaction between ADP and some high-energy phosphate carrier.
- Energy needed to phosphorylate ADP is provided by the **flow of protons down the electrochemical gradient**. This can be calculated.
- The energy released by electron transport is used to transport protons against the electrochemical gradient. Primary and secondary transport principles are at work.
- **If all that was needed was a proton gradient, could one be established without the ET chain and still drive ATP biosynthesis?**

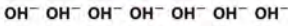
Phosphorylation

Chemiosmotic Theory

p side
 $[H^+]_p = C_2$



n side
 $[H^+]_n = C_1$



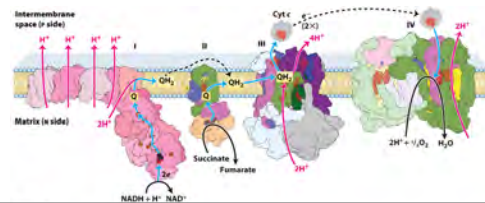
$$\Delta G = RT \ln (C_2/C_1) + ZF\Delta\psi$$

$$= 2.3RT \Delta pH + F\Delta\psi$$

$$\Delta G' = 1.0 + 1.4 = 2.4 \text{ kcal/mol}$$

As a consequence, it will take
 ~3 protons per ATP. But, how
 many precisely?

- The proteins in the electron-transport chain created the **electrochemical proton gradient** (proton-motive force) by one of three means:
 - actively transporting protons across the membrane
 - Complex I and Complex IV
 - releasing protons into the intermembrane space
 - oxidation of QH_2 at Complex III
 - chemically removing protons from the matrix
 - reduction of CoQ (Complex I, II, & III)
 - reduction of oxygen (Complex IV)



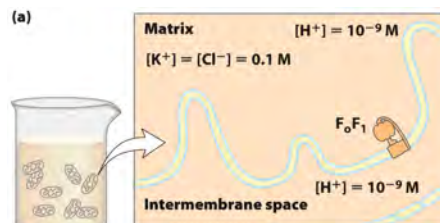
Phosphorylation

Chemiosmotic Theory



Nobel Prize in Chemistry
 1978

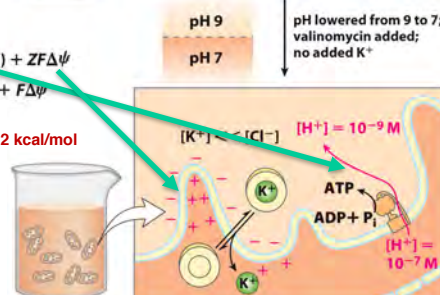
Mitchell's experiment:



$$\Delta G = RT \ln (C_2/C_1) + ZF\Delta\psi$$

$$= 2.3RT \Delta pH + F\Delta\psi$$

For 3 protons:
 $\Delta G' = 3.0 + 4.2 = 7.2 \text{ kcal/mol}$

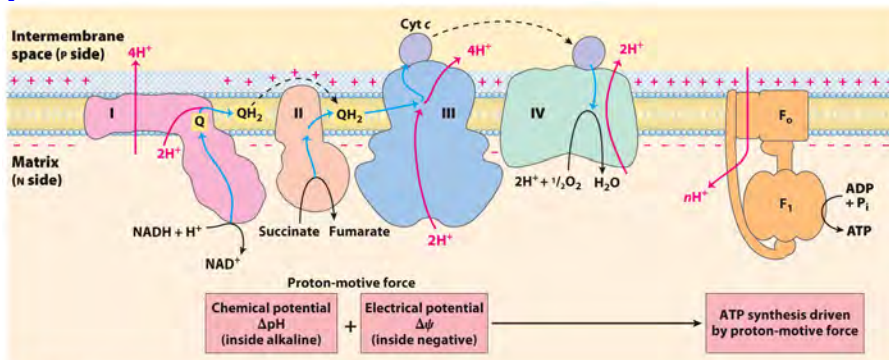
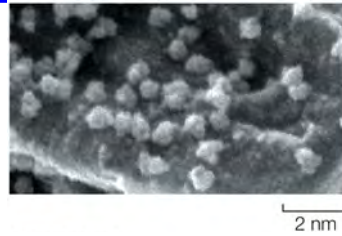


In the absence of electron transport, an artificial H^+ gradient is sufficient for ATP synthesis by mitochondria.

Phosphorylation

Chemiosmotic Theory for ATP Synthesis:

- **Electron transport** sets up a proton-motive force.
- Energy of proton-motive force **drives synthesis of ATP**.



Phosphorylation

Chemiosmotic Theory

Chemiosmotic Energy Coupling Requires Membranes

- The proton gradient needed for ATP synthesis can be stably established across a membrane that is impermeable to ions.
 - plasma membrane in bacteria
 - inner membrane in mitochondria
 - thylakoid membrane in chloroplasts
- The membrane must contain proteins that **couple** the “downhill” flow of **electrons** in the **electron-transfer chain** with the “uphill” flow of **protons** across the membrane.
- The membrane must contain a protein that **couple**s the “downhill” flow of **protons** to the **phosphorylation of ADP**.

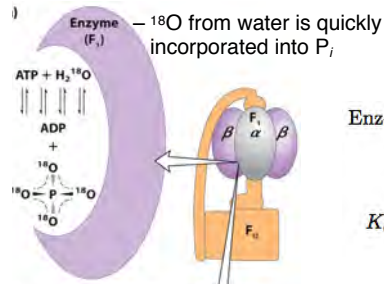
Phosphorylation

Mechanism: BINDING

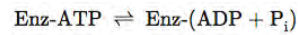
– Isotope studies

– Structural studies

The F_1 catalyzes $ADP + P_i \rightleftharpoons ATP + H_2O$

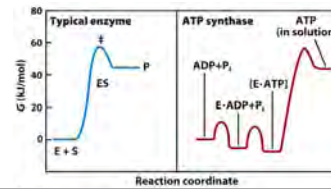


– Kinetic studies



$$K'_{eq} = \frac{k_{-1}}{k_1} = \frac{24 \text{ s}^{-1}}{10 \text{ s}^{-1}} = 2.4$$

– Compare to $>10^5$ for in solution



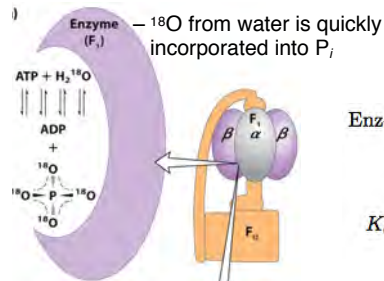
Phosphorylation

Mechanism: BINDING

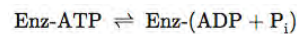
– Isotope studies

– Structural studies

The F_1 catalyzes $ADP + P_i \rightleftharpoons ATP + H_2O$

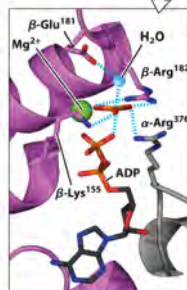
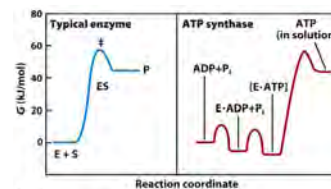


– Kinetic studies



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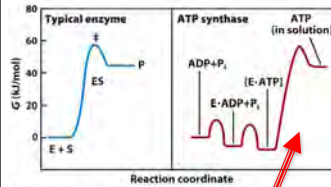
– Compare to $>10^5$ for in solution



Phosphorylation

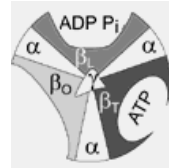
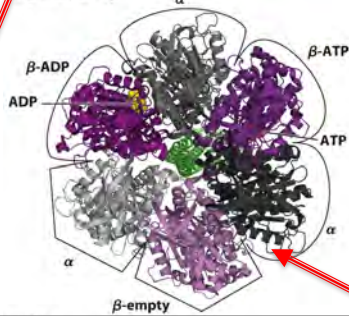
Mechanism: RELEASE

–MORE Structural studies



- Hexamer arranged in three $\alpha\beta$ dimers
- Dimers can exist in three different conformations:
 - open: empty
 - loose: binding ADP and P_i
 - tight: catalyzes ATP formation and binds product

Top view of F_1



WHERE does the energy come from for the release?

WHAT drives the γ -subunits motion?

– This is where ALL the energy of PMF is used to release ATP!

Phosphorylation

Mechanism

Binding-Change Model

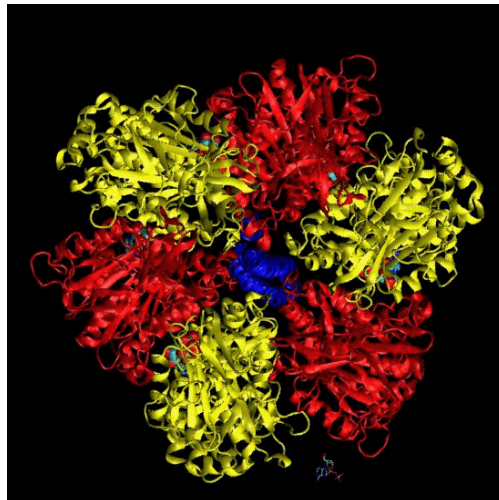


Paul Boyer



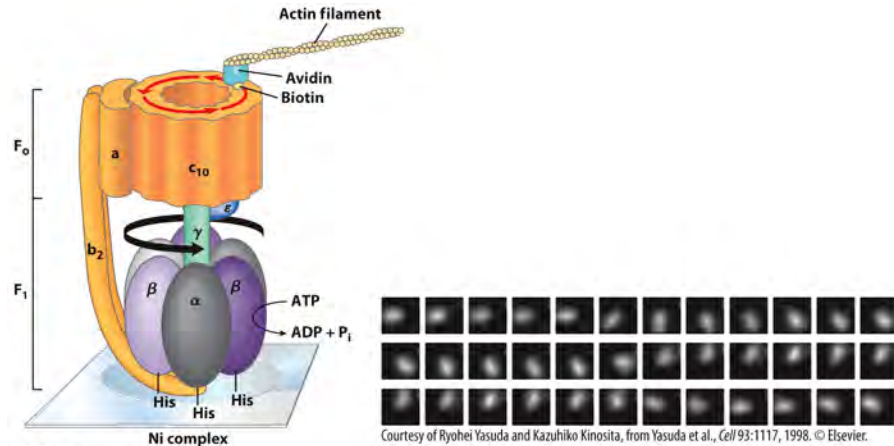
John E. Walker

Nobel Prize in Chemistry 1997



Phosphorylation

Experimental evidence for the rotary/binding-change mechanism

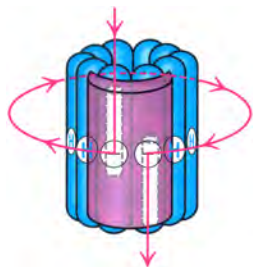


ATP Synthase moving a bead

(<https://www.youtube.com/watch?v=oFgMTdVRi6I>)

Phosphorylation

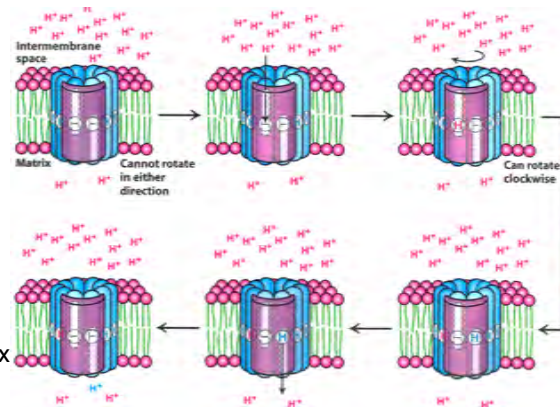
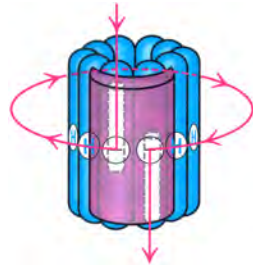
Mechanism



- Protons in excess outside the matrix forces them on a specific Asp residue in the C-subunit.
- But, it must enter through the **a-subunit**.
- Once bound, it causes a conformational change to rotate the c-ring and make available another proton binding site.
- Protons can't get access to the matrix until a complete revolution and contact with the **a-subunit** again, which has a tunnel to the matrix.
- Proton translocation causes a rotation of the F₀ subunit and the **central shaft γ**.

Phosphorylation

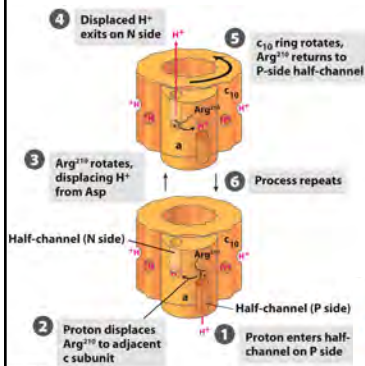
Mechanism



- Protons in excess outside the matrix forces them on a specific Asp residue in the C-subunit.
- But, it must enter through the **a-subunit**.
- Once bound, it causes a conformational change to rotate the c-ring and make available another proton binding site.
- Protons can't get access to the matrix until a complete revolution and contact with the **a-subunit** again, which has a tunnel to the matrix.
- Proton translocation causes a rotation of the F_0 subunit and the **central shaft γ** .

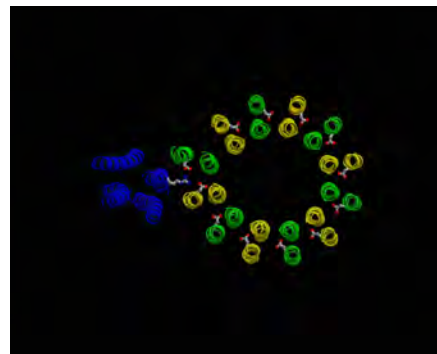
Phosphorylation

Mechanism



H. sapiens

S. cerevisiae



E. coli

The number of c-subunits in the F_0 ring varies from 8-17.

Phosphorylation

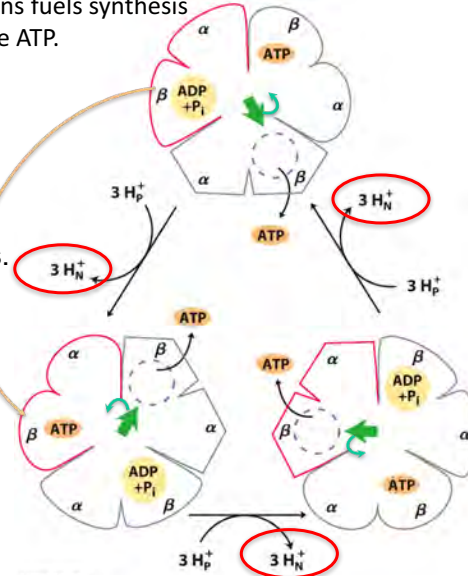
Mechanism

Coupling Proton Translocation to ATP Synthesis

(Let's assume there are 9 c-subunits in F_0)

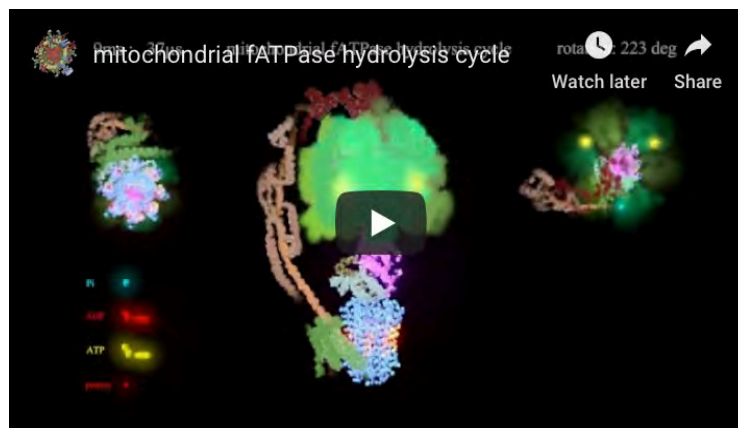
- Proton translocation causes a rotation of the F_0 subunit and the central shaft γ .
- This causes a conformational change within all the three $\alpha\beta$ pairs.
- The conformational change in one of the three pairs promotes condensation of ADP and P_i into ATP.
- The conformational change in another drives the release of ATP.
- This conformational change opens the binding site for ADP and P_i .

Translocation of three protons fuels synthesis of one ATP.



Phosphorylation

The Respiratory Chain and ATP Synthase Produce ATP by a Chemiosmotic Mechanism

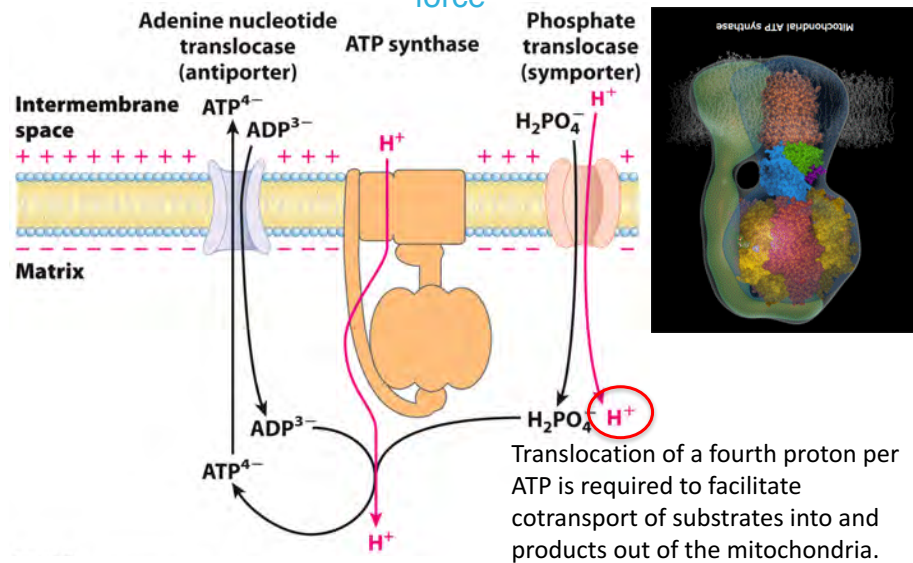


Animation of ATP Synthase

(<https://www.youtube.com/watch?v=E1qTSwe1o3U>)

Phosphorylation

Transport of ADP and P_i into the Matrix also uses proton-motive force



Oxidative Phosphorylation

Net Production of ATP by Oxidation of NADH

- 10 protons pumped to P-side.
- For an F_0 with 9 c-subunits, 9 protons are allowed down their concentration gradient and one 360° rotation of the F_1 subunit produces 3 ATP molecules; or 3 protons per ATP.
- But, one proton is needed to get substrate P_i into matrix, so the total is 4 protons per ATP.
- So, for each $2e^-$ from NADH, the 10 protons will yield 2.5 ATP molecules
- For $2e^-$ from $FADH_2$, the 6 protons will yield 1.5 ATP molecules.
- Different organisms have different numbers of c-subunits, so this number varies species to species. Humans have 8.

Oxidative Phosphorylation

Net Production of ATP by Oxidation of Glucose

- In prokaryotic systems, organelles do not segregate machinery, so all electron carriers can easily feed directly into the electron-transport chain.
- In eukaryotic systems, organellar segregation prevents NADH from the cytosol from directly entering the electron-transport chain at Complex I.
 - NAD⁺ pools are kept segregated and cannot directly cross the mitochondrial inner membrane.
 - Two methods are used to feed the electrons from NADH from the cytosol into the mitochondria:
 - glycerol-3-phosphate shuttle
 - malate-aspartate shuttle

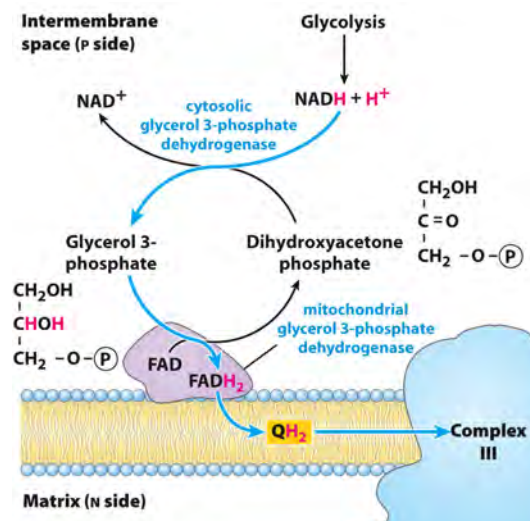
Oxidative Phosphorylation

Converting Cytosolic Electron Carriers (NADH) to the Mitochondria

Glycerol-3-Phosphate Shuttle

Malate-Aspartate Shuttle

- This more complicated shuttle is mostly present in liver, heart, and kidney.
- Will be discussed when we do amino-acid degradation
- It moves NADH equivalents from cytosol to NADH equivalents to the mitochondria.



Oxidative Phosphorylation

Net Production of ATP via Catabolic Pathways

TABLE 19-5 ATP Yield from Complete Oxidation of Glucose

Process	Direct product	Final ATP
Glycolysis	2 NADH (cytosolic) 2 ATP	3 or 5 ^a 2
Pyruvate oxidation (two per glucose)	2 NADH (mitochondrial matrix)	5
Acetyl-CoA oxidation in citric acid cycle (two per glucose)	6 NADH (mitochondrial matrix) 2 FADH ₂ 2 GTP	15 3 2
Total yield per glucose		30 or 32

^aIf the malate/aspartate shuttle is used to transfer reducing equivalents into the mitochondrion, yield is 5 ATP. If the glycerol 3-phosphate shuttle is used, the yield is 3 ATP.

Oxidative Phosphorylation

- Primarily regulated by substrate availability
 - NAD⁺ and ADP/P_i
- Inhibition of OxPhos leads to accumulation of NADH.
 - causes feedback inhibition cascade up to PFK-1 in Glycolysis
- Inhibitor of F₁ (IF₁)
 - prevents hydrolysis of ATP during low oxygen
 - IF₁ only active at lower pH in matrix, encountered when electron transport is stalled (i.e., low oxygen)

Regulation

