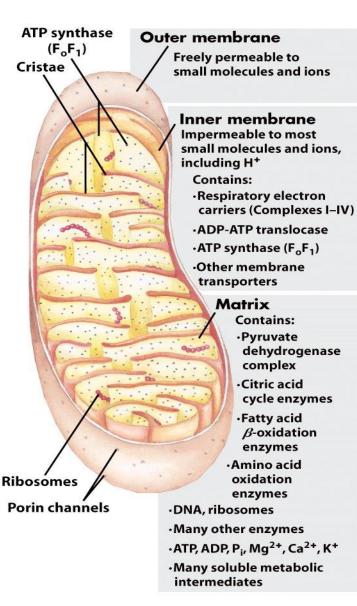
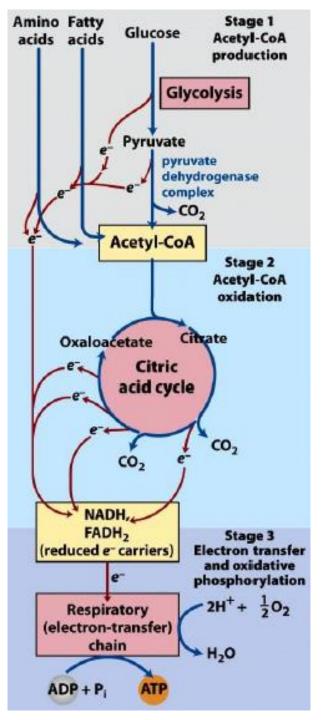
Mitochondrial transport processes, respiratory chain and ATP synthesis

From the Chemistry Exam to the Final Exam in Biochemistry Dr. Lengyel Anna

Biochemical anatomy of a mitochondrion





What phase of cellular respiration has the highest ATP yield?

- a) Oxidative phosphorylation
- b) Gluconeogenesis
- c) Krebs cycle
- d) Glycolysis
- e) Fermentation

Given a healthy individual with a normal metabolic rate, which of the following compounds is the most energy rich?

- A. GTP
- B. ATP
- C. FADH₂
- D. NADH

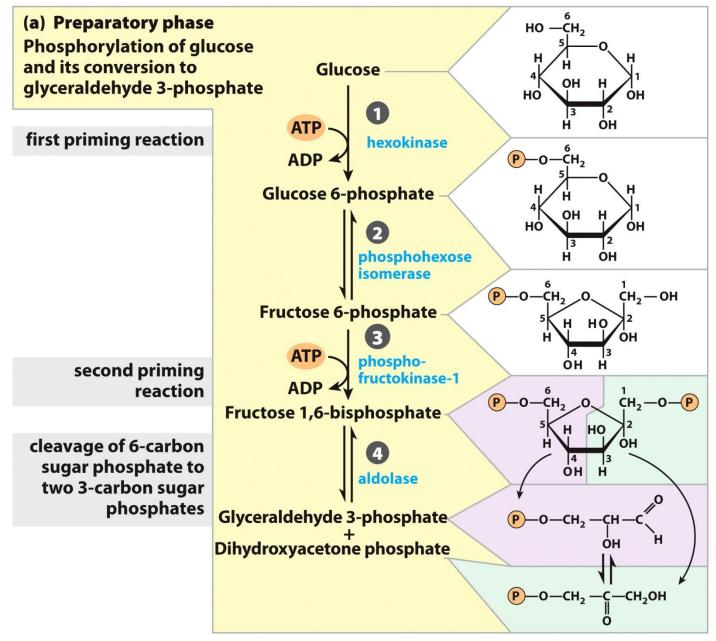


Figure 14-2 part 1

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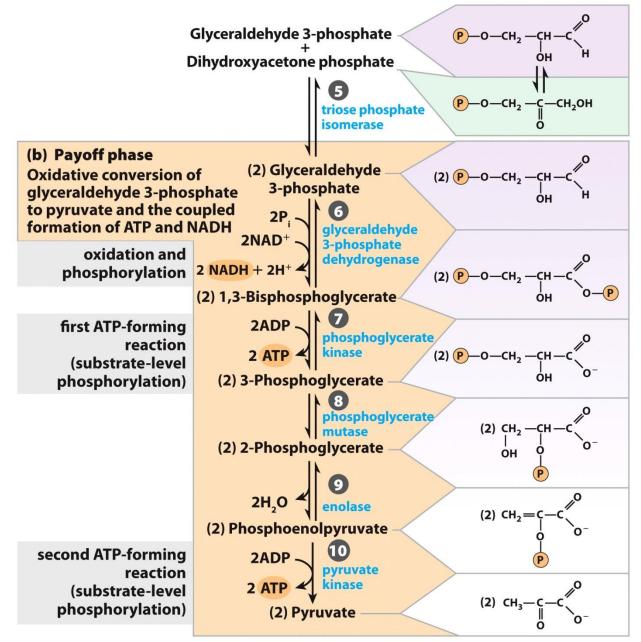


Figure 14-2 part 2

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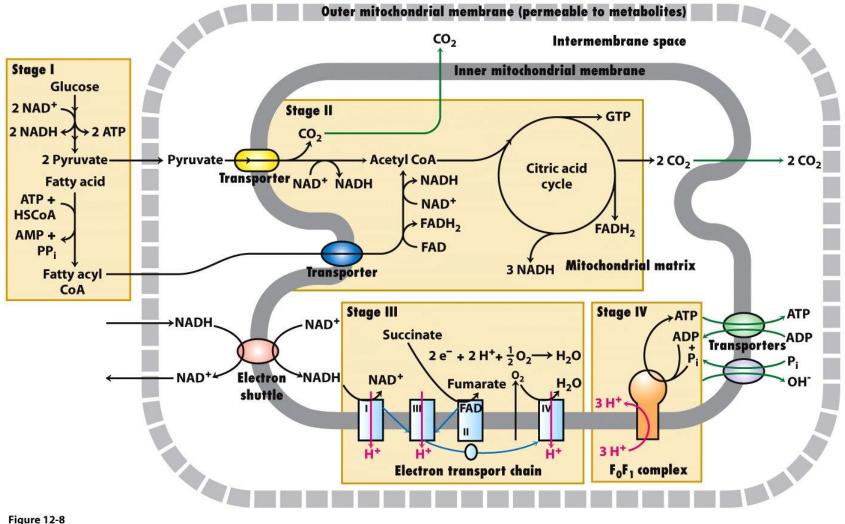
In substrate level phosphorylation...

- A. ATP synthesis is linked to dissipation of proton gradient.
- B. High energy intermediate compounds cannot be isolated.
- C. Oxidation of one molecule of substrate is linked to synthesis of more than one ATP molecule.
- D. Only mitochondrial reactions participate in ATP formation.
- E. The cleavage of the high-energy bond in the substrate provides the energy required for ATP synthesis.

The energy of oxidation is initially trapped as a high-energy phosphate compound and then used to form ATP. Which of the following intermediates of glycolysis is a high energy compound?

- A. Fructose-6-P
- B. Glyceraldehyde-3-P
- C. Fructose-1,6 bisphosphate
- D. Glucose-6-P
- E. Phosphoenol pyruvate

Mitochondrial transports



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Malat – aspartate shuttle

P/O = 2.5

From cytosolic NADH+H⁺ mitochondrial NADH+H⁺

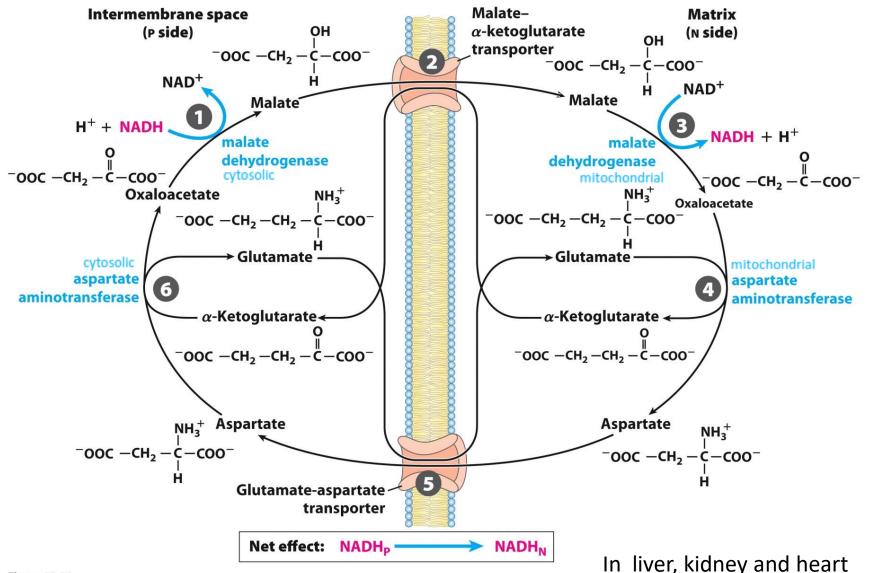
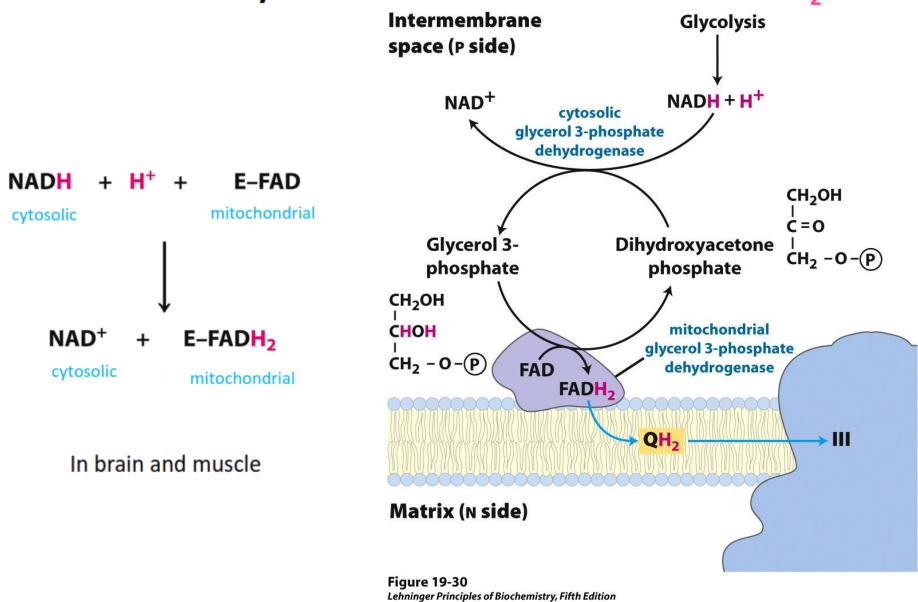


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Glycerol 3-phosphate shuttle

-phosphate shuttle P/O = 1.5 From cytosolic NADH+H⁺ mitochondrial FADH₂



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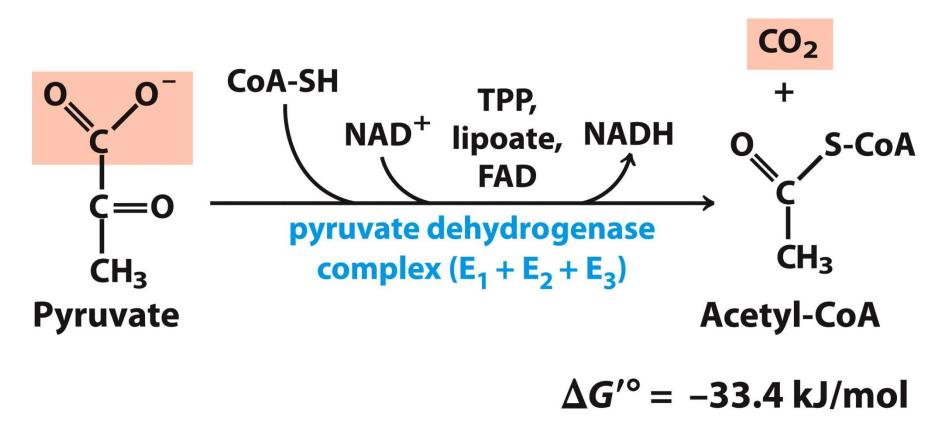
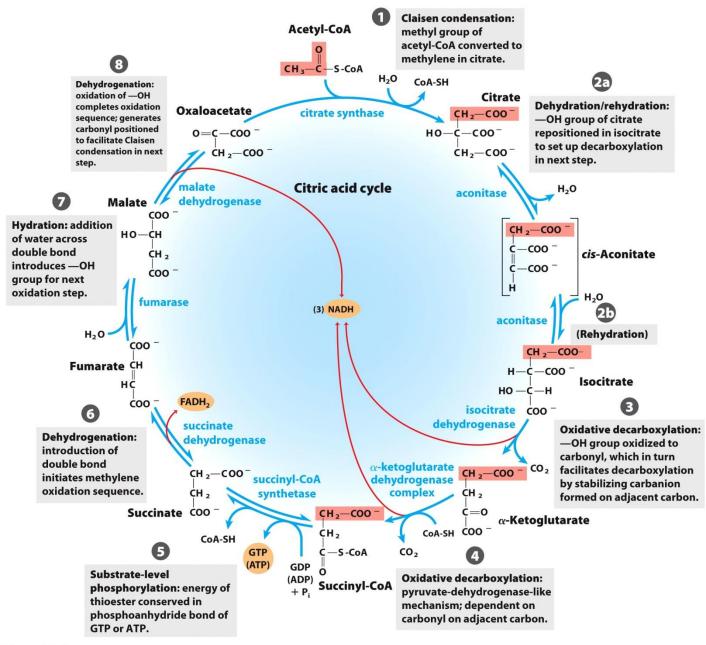


Figure 16-2 *Lehninger Principles of Biochemistry*, Sixth Edition © 2013 W. H. Freeman and Company

Which one of the following enzymes catalyzes substrate level phosphorylation in TCA cycle?

- A. Malate dehydrogenase
- B. Succinyl-CoA synthetase
- C. α-ketoglutarate dehydrogenase complex
- D. Isocitrate dehydrogenase
- E. Succinate dehydrogenase

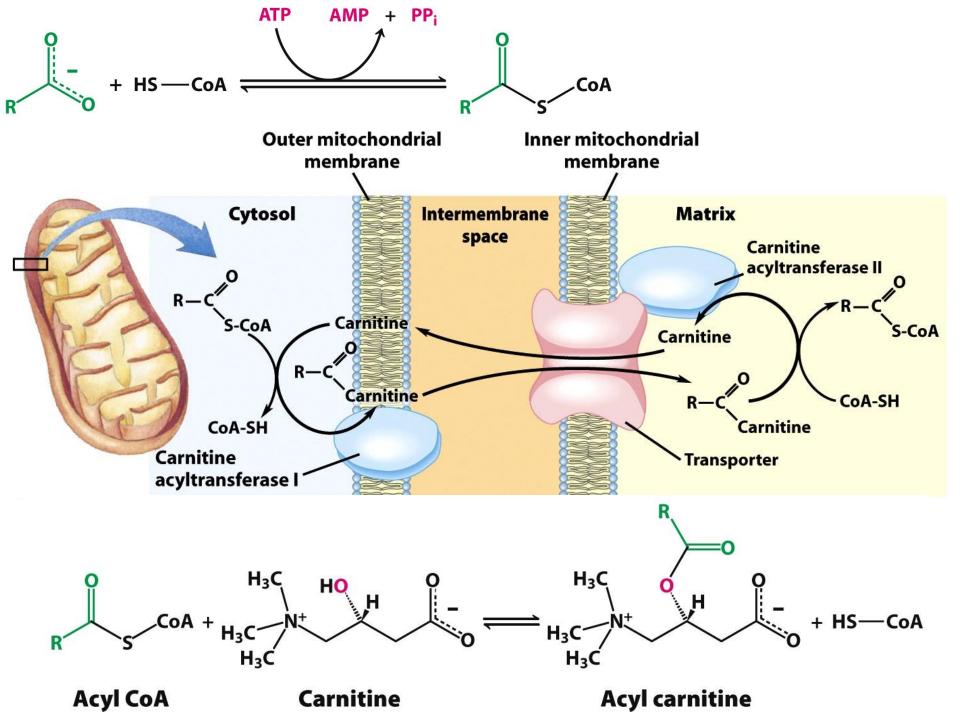


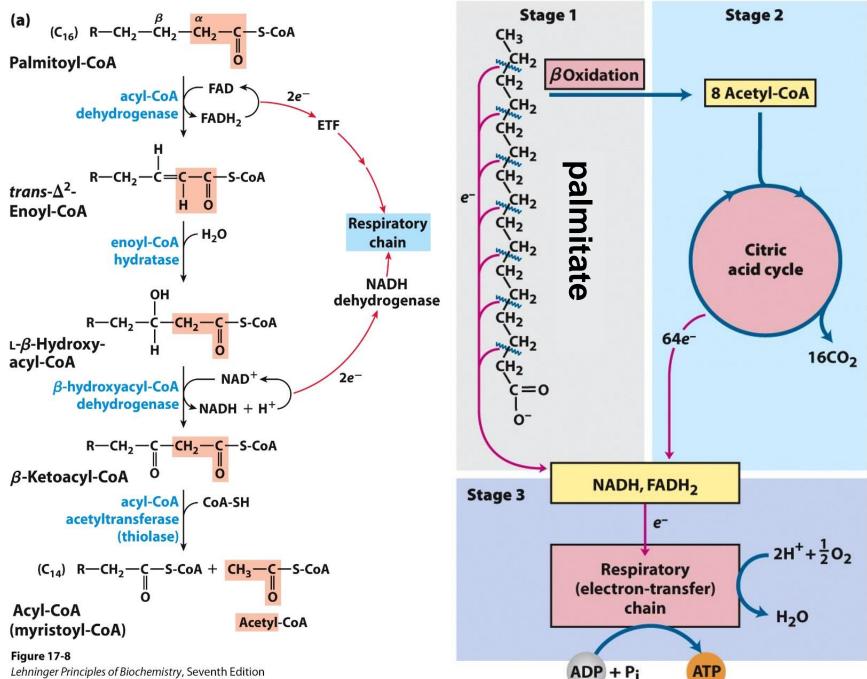


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During aerobic respiration, which of the following pathways correctly orders the process of cellular metabolism after glycolysis in eukaryotic cells?

- A. Citric acid cycle \rightarrow Pyruvate decarboxylation \rightarrow Oxidative phosphorylation
- B. Pyruvate decarboxylation \rightarrow Oxidative phosphorylation \rightarrow Citric acid cycle
- C. Citric acid cycle \rightarrow Oxidative phosphorylation \rightarrow Pyruvate decarboxylation
- D. Pyruvate decarboxylation \rightarrow Citric acid cycle \rightarrow Oxidative phosphorylation





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All of the following except one are NAD⁺ requiring enzymes:

- A. Acyl-CoA dehydrogenase
- B. Glyceraldehyde-3-P dehydrogenase
- C. Pyruvate dehydrogenase complex
- D. Malate dehydrogenase
- E. Lactate dehydrogenase

The respiratory chain

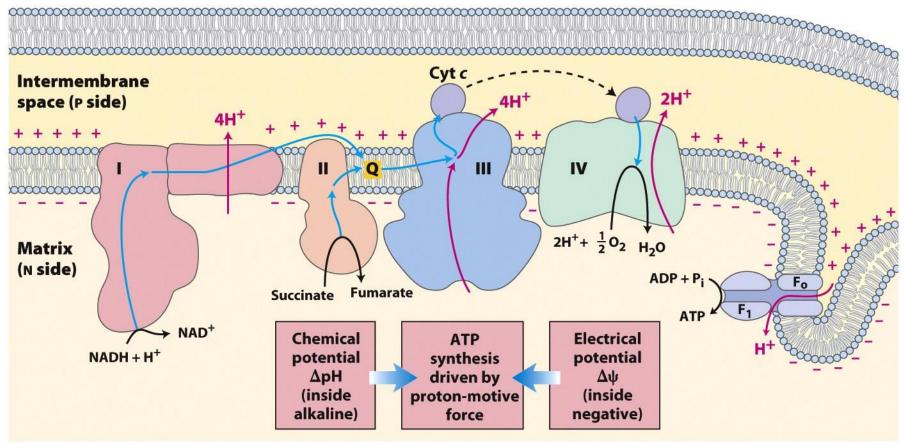


Figure 19-19

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The e^- transport is an exergonic process (ΔG <0), it covers the energy requirement of the p^+ transport which is an endergonic process (ΔG >0).

The primary purpose of the electron transport chain of mitochondria is _____.

- a) to directly phosphorylate ADP
- b) to synthesize ATP synthase
- c) to directly phosphorylate AMP
- d) to carry ADP into the mitochondrial matrix
- e) the generation of energy to sequester protons in the intermembrane space

Which of the following areas of the mitochondria has the lowest pH?

- A. The mitochondrial matrix
- B. The intermembrane space
- C. The cytosol
- D. The mitochondrial cristae

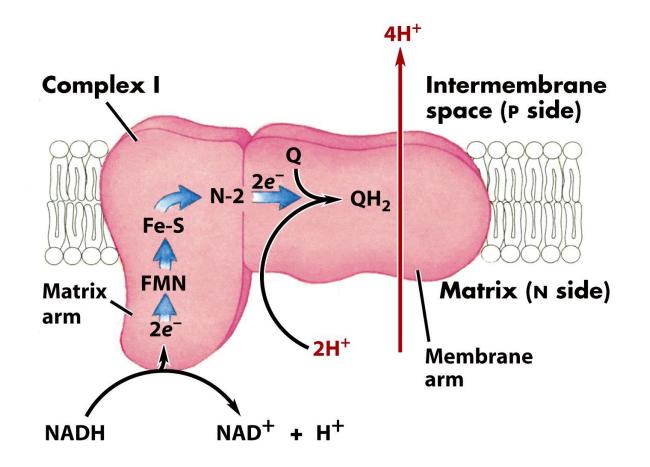
Why is oxygen necessary in aerobic cellular respiration?

- A. It provides the hydrogen nuclei needed to create a proton gradient in the intermembrane space.
- B. It is the final electron acceptor in the electron transport chain.
- C. It is needed for glycolysis, which begins the process of respiration in cells.
- D. It is important in creating oxaloacetate in the Kreb's cycle.

redox process (ox + $n e^- \rightarrow red$)	n	ε°' (V)
$^{1\!\!/_2}\mathrm{O}_2\left(g\right)+2\:\mathrm{H}^++2\:\mathrm{e}^-\to\mathrm{H}_2\mathrm{O}\left(l\right)$	2	0.81
cytochrome- a_3 (Fe ³⁺) + e ⁻ \rightarrow cytochrome- a_3 (Fe ²⁺)	1	0.55
cytochrome-a (Fe ³⁺) + e ⁻ \rightarrow cytochrome-a (Fe ²⁺)	1	0.29
cytochrome-c (Fe ³⁺) \rightarrow cytochrome-c (Fe ²⁺)	1	0.25
cytochrome-c ₁ (Fe ³⁺) \rightarrow cytochrome-c ₁ (Fe ²⁺)	1	0.22
cytochrome-b (Fe ³⁺) \rightarrow cytochrome-b (Fe ²⁺)	1	0.07
ubiquinone + 2H+ + 2e- \rightarrow ubiquinol	2	0.04
NADH dehydrogenase (FMN) + 2H ⁺ + 2e ⁻ \rightarrow NADH dehydrogenase (FMNH ₂)	2	-0.03
NADP ⁺ + H ⁺ + 2 e ⁻ \rightarrow NADPH	2	- 0.32
$NAD^+ + H^+ + 2 e^- \rightarrow NADH$	2	- 0.32
$2 H^+ + 2 e^- \rightarrow H_2(g) (pH = 7)$	2	- 0.41

Complex I. NADH-dehydrogenase (NADH: ubiquinone oxidoreductase)

NADH + H⁺ + Q \longrightarrow NAD⁺ + QH₂ inhibitors: amytal, rotenone, piericidin A



Complex II. Succinate-dehydrogenase

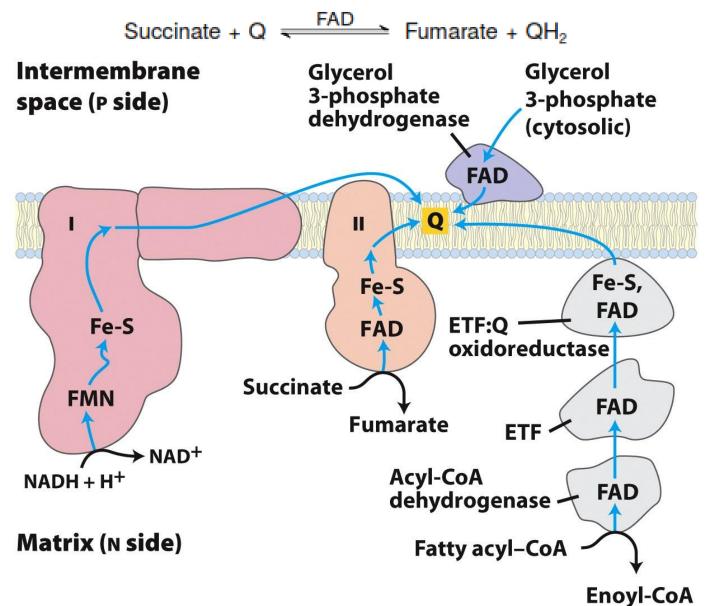


Figure 19-8 *Lehninger Principles of Biochemistry, Fifth Edition* © 2008 W.H. Freeman and Company

Complex III. Ubiquinone-cytochrome c-oxidoreductase

 $QH_2 (red) + 2 cyt c (ox) + 2 H_N^+ \longrightarrow Q (ox) + 2 cyt c (red) + 4 H_P^+$ Inhibitor: antimycin A, myxothiazol

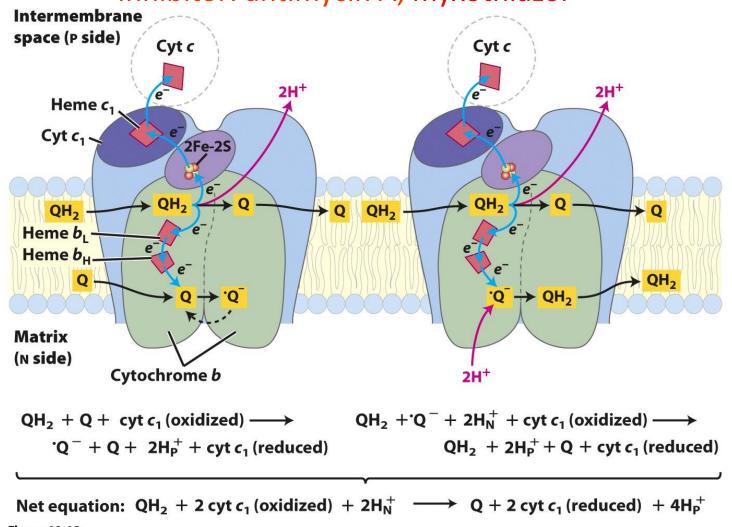
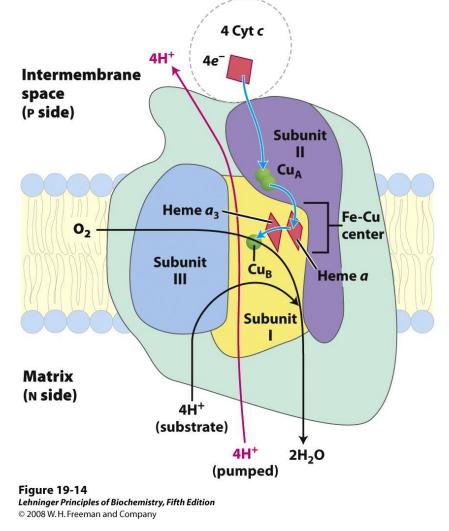


Figure 19-12 Lehninger Principles of Biochemistry, Fifth Edition © 2008 W.H. Freeman and Company

Complex IV. Cytochrome c-oxidase

4 Cyt-c (red.) + 8H⁺(n)+ $O_2 \longrightarrow$ 4 Cyt-c (ox.) + 4H⁺(p) + 2 H₂O Inhibitors: cyanide, CO, hydrogen sulfide, azides



$$2 \operatorname{cyt} c (\operatorname{red}) + 4 \operatorname{H}_{N}^{+} + \frac{1}{2} \operatorname{O}_{2} \longrightarrow 2 \operatorname{cyt} c (\operatorname{ox}) + 2 \operatorname{H}_{P}^{+} + \operatorname{H}_{2} \operatorname{C}_{2}$$

Which of the following components of electron transport chain does not contain iron sulfur center?

- A. NADH dehydrogenase complex
- B. Ubiquinone-cytochrome c-oxidoreductase
- C. Succinate dehydrogenase
- D. Cytochrome c-oxidase

The ATP synthase

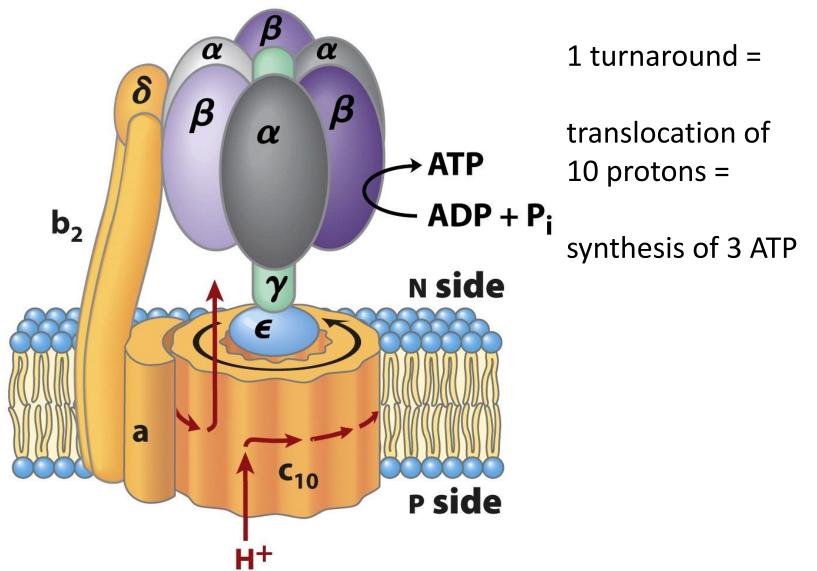
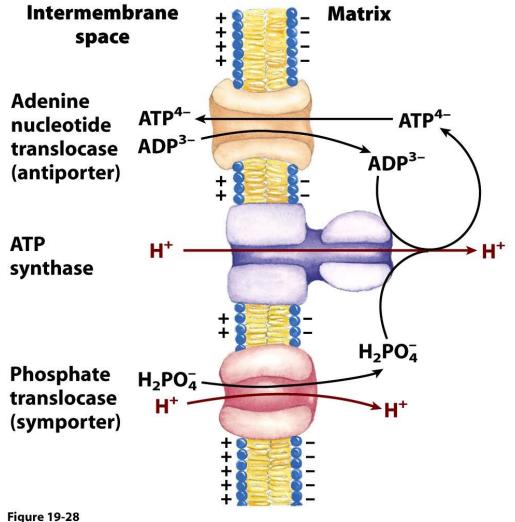


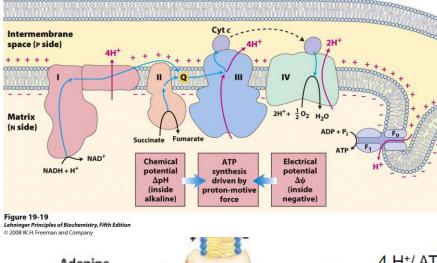
Figure 19-25f *Lehninger Principles of Biochemistry, Fifth Edition* © 2008 W.H. Freeman and Company

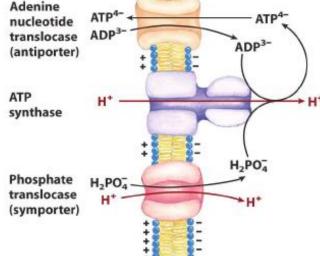
Adenine nucletide and phosphate translocases



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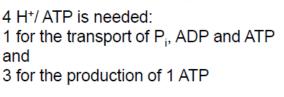
The P/O Ratio: How many ATP is synthetized from the energy released by the reduction of an oxygen atom?

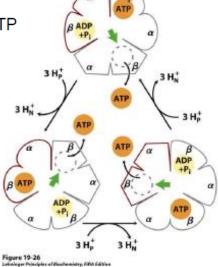




1 oxygen atom is reduced by 2 electrons
Both NADH and FADH₂ provide two electrons
During their oxidation in the respiratory chain: NADH "pumps" 10 protons into the IMS
FADH₂ "pumps" 6 protons into the IMS
Translocation of 10 protons into the matrix through ATP synthase results in the synthesis of 3 ATP molecules

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The P/O Ratio: How many ATP is synthetised from the energy released by the reduction of an oxygen atom?

- The P/O ratio was expected to be 3 ATP / NADH and 2 ATP / FADH₂ (outdated!!!).
- Today we have experimentally determined results, which show ~2.5 ATP / NADH and ~1.5 ATP / FADH₂.

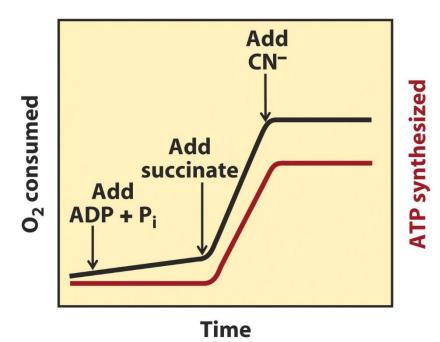
Consider: the pumping of protons into the intermembrane space is NOT a stoichiometric process.

ATP synthesis is coupled with the redox reactions of the respiratory chain (electrontransfer). In an uncoupled mitochondrium, oxidation of NADH or succinate (without oxidative phophorylation [ATP synthesis]) leads to heat production.

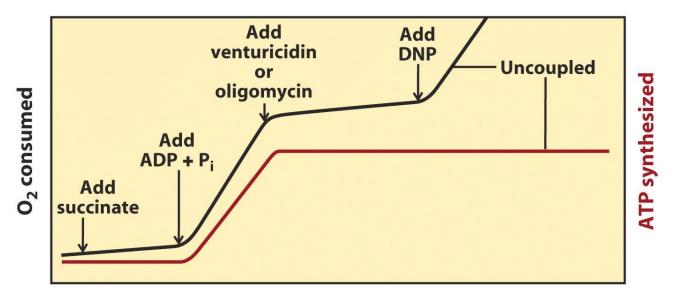
e.g.: thermogenin (UCP1) (physiologic uncoupling protein).

Isolated mitochondria + ADP + Pi + substrate (succinate) + buffer

- Substrate (succinate) is oxidized
- O₂ is consumed
- ATP is synthesized
- O₂ consumption in coupled mitochondria can be inhibited by CIII and CIV inhibitors: Antimycin A, Cyanid, CO



Isolated mitochondria + ADP + Pi + substrate (succinate) + buffer + inhibitors



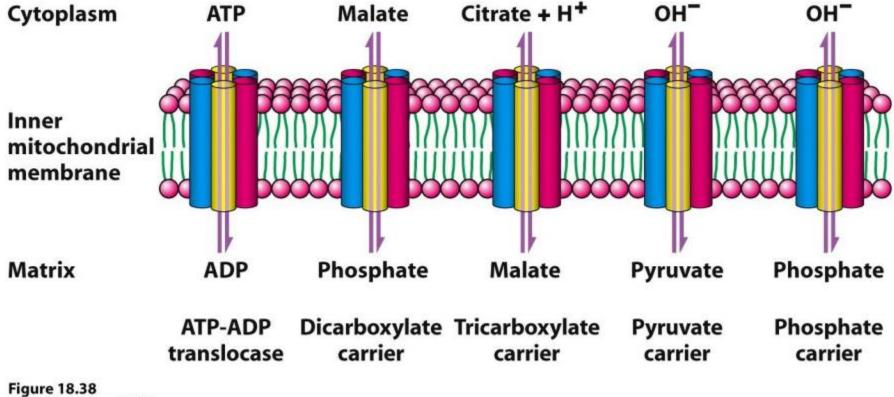
Time

- In intact (coupled) mitochondria the inhibition of ATP synthesis (Fo-F1) blocks electron transfer. Inhibitors: Venturicidin, Oligomycin, Aurovertin
- Uncoupling of oxidation and phosphorylation can also be demonstrated using chemical compounds. Respiration increases, but no ATP is produced.

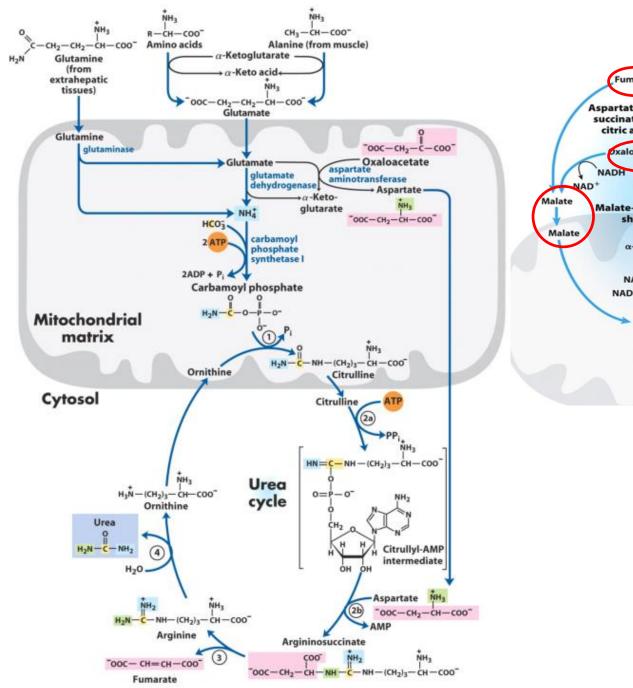
Uncoupling with chemicals: 2,4 Dinitrophenol (DNP), FCCP

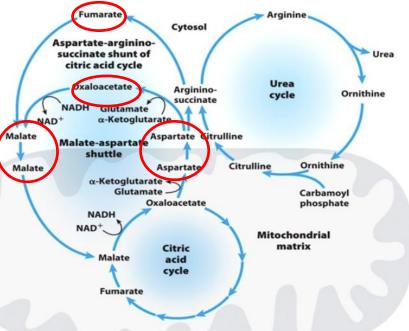
Summary

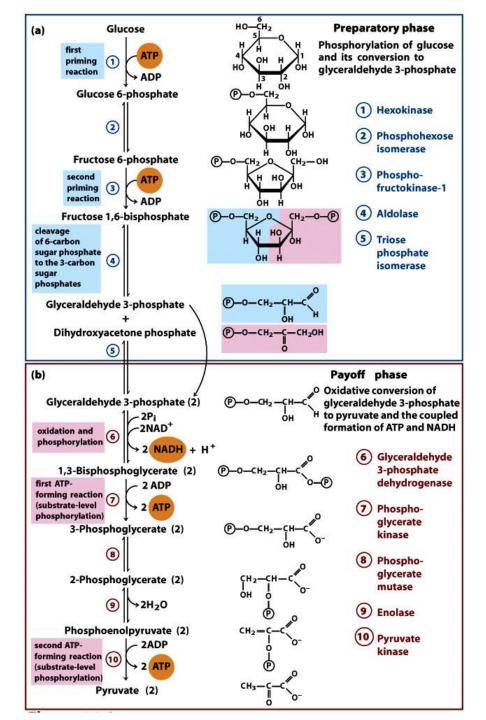
- Oxidative phosphorylation occurs in the inner membrane of eukaryotic mitochondrion. Proton pumps of the respiratory chain and the F₀F₁ ATP synthase work in this process together, they are coupled.
- 4 complexes of the respiratory chain (many subunits and redox centers) transfer electrons from reduced coenzymes to the terminal electron acceptor O₂, which will be reduced to water.
- In this process, a proton gradient arises between the outer and inner sides of the membrane. Protons reenter the matrix through the F₀F₁ ATP synthase and drive ATP synthesis.
- Proton gradient arose by the oxidation of NADH produces ~2,5 mol ATP, in the case of FADH₂ ~1,5 mol ATP. This is the P/O-ratio.
- Oxidation and phosphorylation can be uncoupled! The uncoupling protein, thermogenin induces heat production instead of ATP synthesis.
- Oxidative phosphorylation is regulated by the ADP level.
- Communication between cytosol and mitochondrium is fullfilled by several transporters located in the inner membrane.

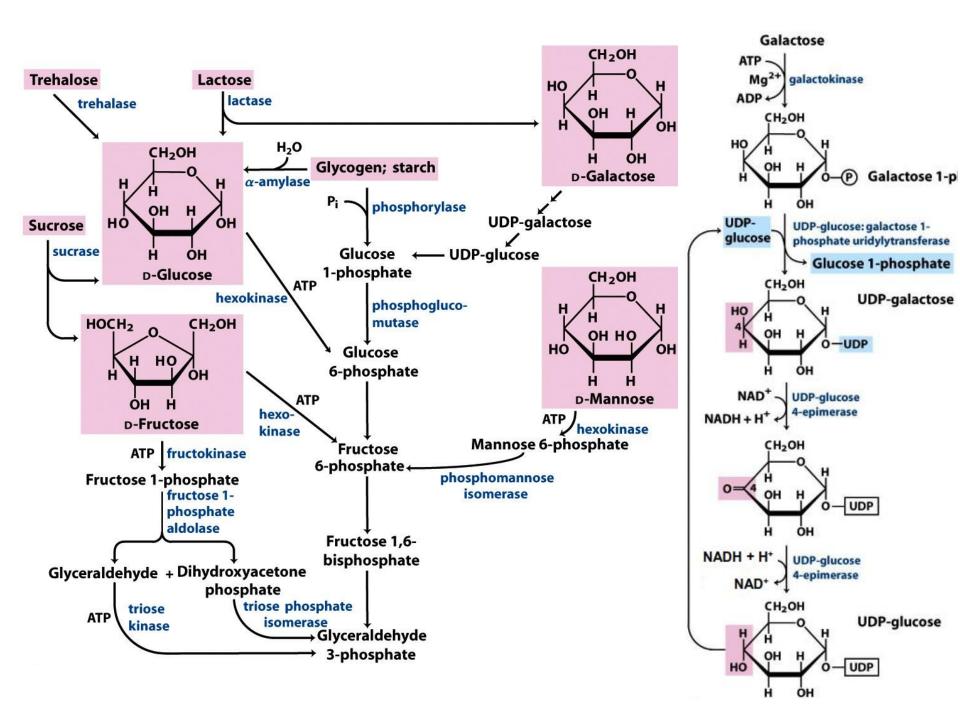


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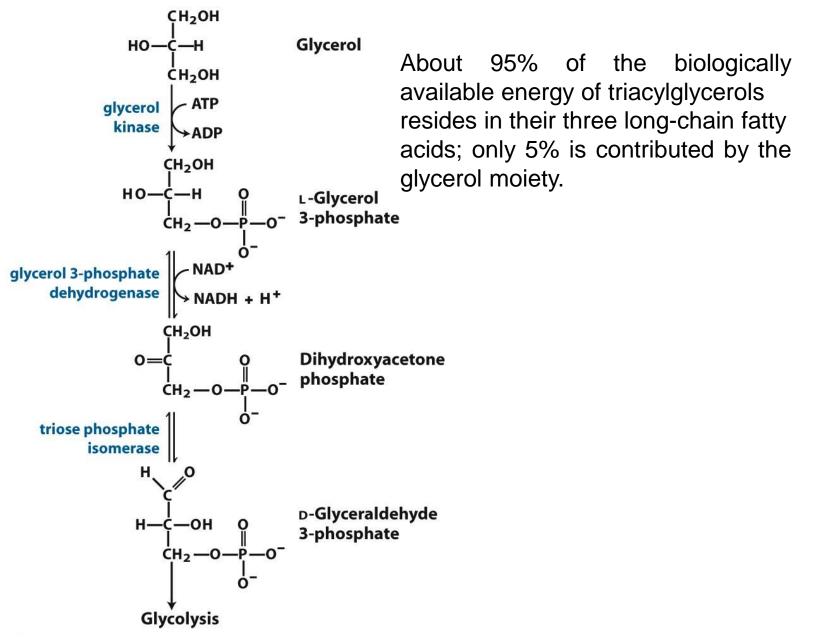


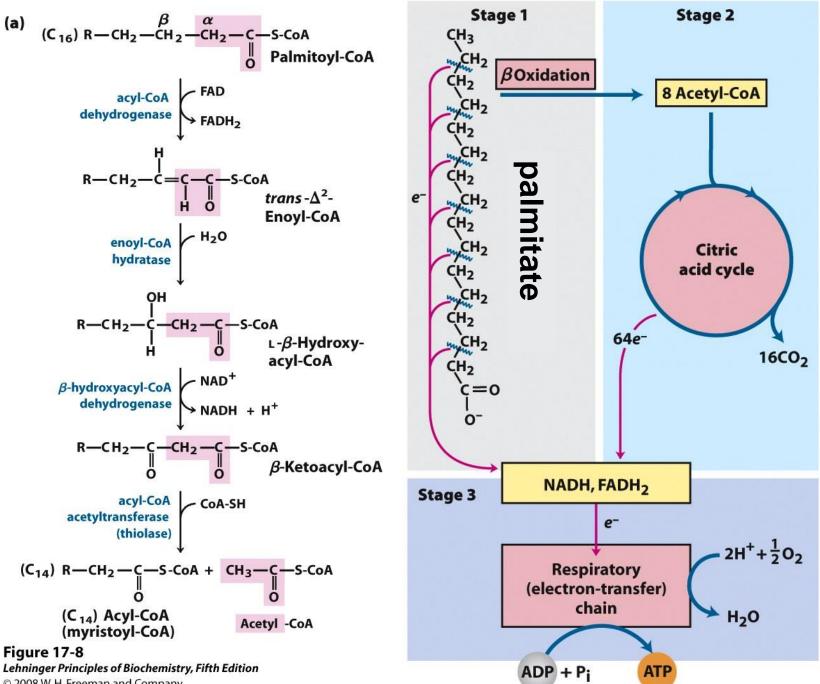




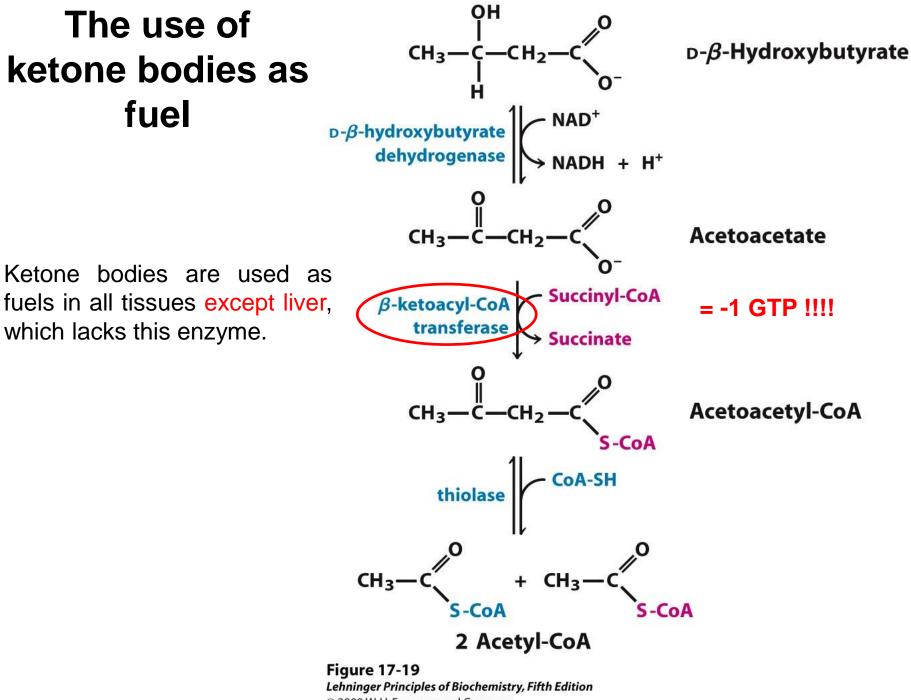


Entry of glycerol into glycolysis





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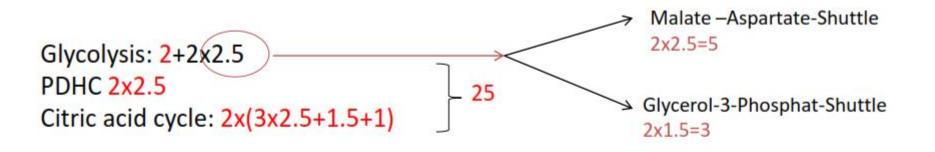


TABLE 16-1	Stoichiometry of Coenzyme Reduction and ATP Formation in the Aerobic Oxidation of Glucose via Glycoly the Pyruvate Dehydrogenase Complex Reaction, the Citric Acid Cycle, and Oxidative Phosphorylation				
Reaction		Number of ATP or reduced coenzyme directly formed	Number of ATP ultimately formed*		
Glucose \longrightarrow g	glucose 6-phosphate	-1 ATP	-1		
Fructose 6-pho	sphate —— fructose 1,6-bisphosphate	-1 ATP	-1		
2 Glyceraldehy	de 3-phosphate	2 NADH	3 or 5 [†]		
2 1,3-Bisphosp	hoglycerate	2 ATP	2		
2 Phosphoenol	pyruvate> 2 pyruvate	2 ATP	2		
2 Pyruvate —	→ 2 acetyl-CoA	2 NADH	5		
2 Isocitrate —	$\rightarrow 2 \alpha$ -ketoglutarate	2 NADH	5		
2 α-Ketoglutara	ate \longrightarrow 2 succinyl-CoA	2 NADH	5		
2 Succinyl-CoA \longrightarrow 2 succinate		2 ATP (or 2 GTP)	2		
2 Succinate —	→ 2 fumarate	2 FADH ₂	3		
2 Malate \longrightarrow	2 oxaloacetate	2 NADH	5		
Total			30-32		

*This is calculated as 2.5 ATP per NADH and 1.5 ATP per FADH₂. A negative value indicates consumption.

[†]This number is either 3 or 5, depending on the mechanism used to shuttle NADH equivalents from the cytosol to the mitochondrial matrix; see Figures 19–30 and 19–31.

Table 16-1 Lehninger Principles of Biochemistry, Fifth Edition © 2008 W. H. Freeman and Company

palmitoyl-CoA + 23 O₂ + 108 Pi + 108 ADP \rightarrow CoA + 108 ATP + 16 CO₂ + 23 H₂O

TABLE 17-1Yield of ATP during Oxidation of One Molecule of Palmitoyl-CoA to CO, and H,O

Enzyme catalyzing the oxidation step	Number of NADH or FADH ₂ formed	Number of ATP ultimately formed*
Acyl-CoA dehydrogenase	7 FADH ₂	10.5
eta-Hydroxyacyl-CoA dehydrogenase	7 NADH	17.5
Isocitrate dehydrogenase	8 NADH	20
lpha-Ketoglutarate dehydrogenase	8 NADH	20
Succinyl-CoA synthetase		8†
Succinate dehydrogenase	8 FADH ₂	12
Malate dehydrogenase	8 NADH	20
Total		108

*These calculations assume that mitochondrial oxidative phosphorylation produces 1.5 ATP per FADH₂ oxidized and 2.5 ATP per NADH oxidized.

[†]GTP produced directly in this step yields ATP in the reaction catalyzed by nucleoside diphosphate kinase (p. 510).

The energetic cost of activating a fatty acid is equivalent to **2** ATP, and the net gain per molecule of **palmitate** is 106 ATP.

