Biochemistry

Metabolic pathways

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Energy metabolism

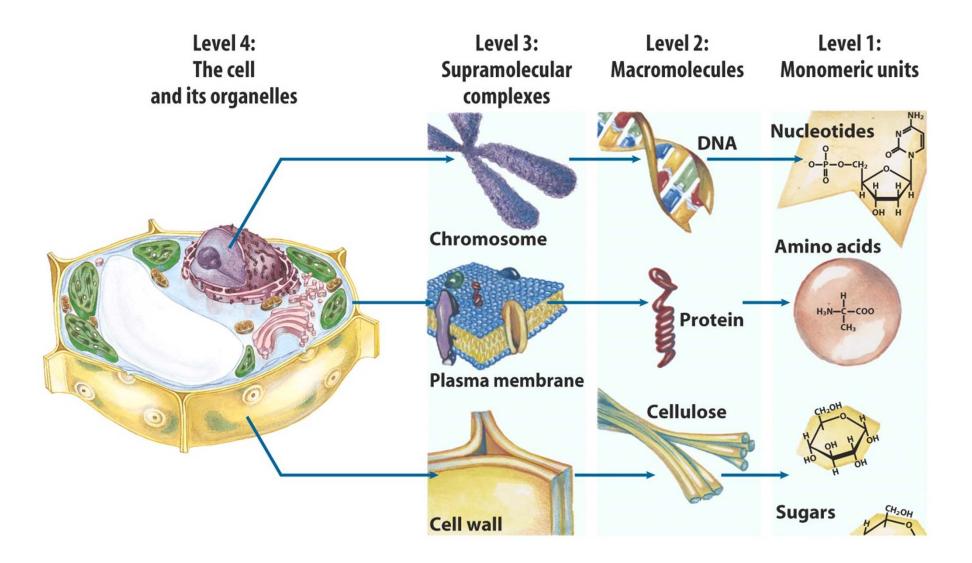
high-energy donors coupled reactions co-factors: NAD*/NADH; FAD/FADH₂; TPP; PLP; CoA; Biotin glycolysis/gluconeogenesis citric acid cycle: regulation (tissue-dependent) *GABA-shunt*; glyoxylate cycle respiratory chain and oxidative phosphorylation glycogen metabolism (activated glucose) pentose phosphate pathway (tissue specificity) photosynthesis: RUBISCO *C*4-plants Nitrogen metabolism

N₂ assimilation via reduction to NH₃ (nitrogenase complex) NH₃ metabolism: glutamate-dehydrogenase glutamate synthase glutamine synthetase glutamine amidotransferase urea cycle C₁ metabolism (PLP, THF, SAM, homocystein) nucleotide metabolism: biosynthesis of purines and pyrimidines from RNA to DNA (NDP reductase) salvage pathway, HGPRT deficiency cytostatic drugs catabolism (ADA-deficiency, urate)

Signal transduction

GPCR - glucagon signalling RTK - insulin signalling G-proteins

Structural hierarchy in the molecular organization of cells



Lehninger Principles of Biochemistry. 5th Ed. Nelson & Cox

Essential Questions?

- What are the properties of **regulatory** enzymes?
- How do regulatory enzymes sense the momentary needs of cells?
- What molecular mechanisms are used to regulate enzyme activity?
- Factors that influence enzymatic activity
- General features of **allosteric regulation**
- The kind of **<u>covalent modification</u>** that regulates the activity of enzymes
- Is the activity of some enzymes controlled by both allosteric regulation and covalent modification?
- Special focus: is there an example in nature that exemplifies the relationship between quaternary structure and the emergence of allosteric properties? hemoglobin and myoglobin paradigms of protein structure and function

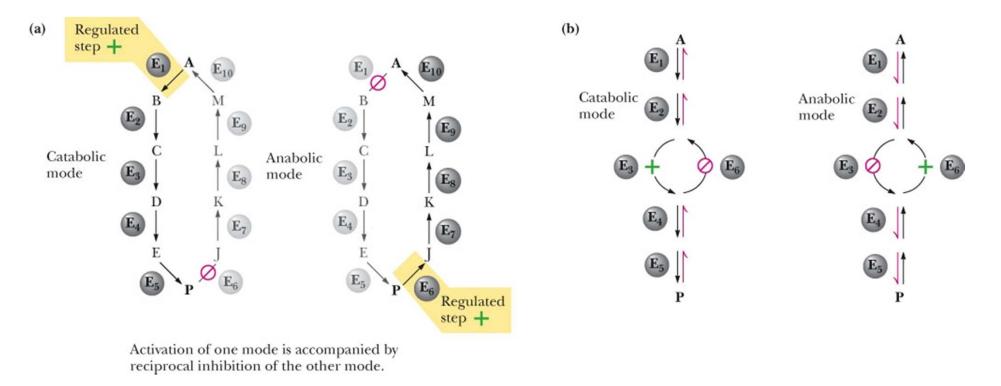
What Factors Influence Enzymatic Activity?

- The availability of substrates and cofactors usually determines how fast the reaction goes.
- Km values ~ the prevailing *in vivo* concentration of the substrates
- As product accumulates, the apparent rate of the enzymatic reaction will decrease as equilibrium is approached
- Genetic regulation of enzyme synthesis and decay determines the amount of enzyme present at any moment
 - Induction = activation of enzyme synthesis
 - **Repression** = shutdown of enzyme synthesis
 - By controlling the [E], cells can activate or terminate metabolic routes.
 - Other factors: Zymogens, isozymes, and modulator proteins may play a role

Principal Characteristics of Metabolic Pathways

- 1. Metabolic pathways are irreversible.
- 2. Catabolic and anabolic pathways must differ.
- 3. Every metabolic pathway has a first committed step.
- 4. All metabolic pathways are regulated.
- 5. M.p. in eukaryotic cells occur in specific subcellular compartments.

Metabolic Regulation Requires Different Pathways for Oppositely Directed Metabolic Sequences



Parallel pathways of catabolism and anabolism must differ in at least one metabolic step in order that they can be regulated independently. Shown here are two possible arrangements of opposing catabolic and anabolic sequences between **A** and **P**. (a) Parallel sequences proceed by independent routes. (b) Only one reaction has two different enzymes.

Metabolic Pathways are Compartmentalized within Cells

- Eukaryotic cells are compartmentalized by an endomembrane system advantageous for metabolism
- Each organelle (compartment) dedicated to specialized metabolic functions and contains appropriate enzymes confined together.

• Advantages:

- Allow analyses of respective functions because they can be separated
- Enzymes are isolated from competing pathways
- Temporal compartmentalization
- Intermediates are spatially and chemically segregated
- Genes of metabolism show a circadian pattern of regulated expression
- Example: glucose-6 phosphatase that converts glucose-6-phosphate to glucose is localized in the ER.

Where metabolic processes occur at the organ level

Liver

- Liver is the center of metabolism maintains blood glucose levels and regulates the concentration of metabolites in the blood.
- Stores glycogen that can be made into glucose-6-phosphate, then glucose.
- Makes glucose by gluconeogenesis (from pyruvate, *de novo*).
- Synthesizes FA, cholesterol and bile salts.
- Produces ketone bodies but cannot use them (no CoA transferase in the liver)
- Only the liver and kidneys contain glucose-6-phosphatase

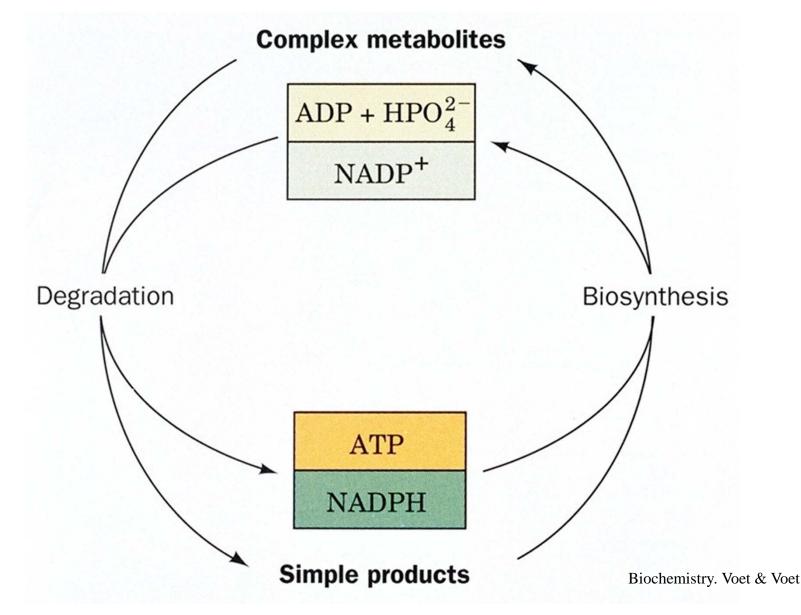
Glucose regulation in the liver

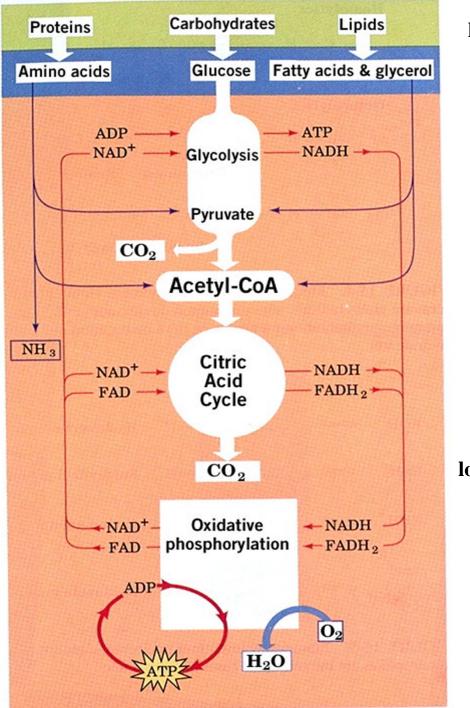
- When blood glucose is high (<u>fed state</u>), FA are synthesized by the liver, converted into triacylglycerols and packaged into VLDL that are secreted into the blood.
- When blood glucose is low (<u>fasting state</u>), the liver produces ketone bodies to fuel the heart and muscle to preserve glucose for the brain. Eventually brain is fed by ketone bodies.

Other organs in metabolism

- Brain Glucose is the primary fuel; only after prolonged fasting (not eating) does the brain use ketone bodies for fuel (last resort).
 - Brain has no capacity to store fuel needs a constant supply
 - Consumes a lot of energy 120 g of glucose per day.
 - Glucose is transported into the brain by GLU3 glucose transporter (crosses the membrane).
 - [glucose] in brain maintained around 5 mM so glucose is saturated under normal conditions. If drops to 2.2 mM the brain is in trouble.
- Muscle uses glucose, FA and ketone bodies for fuel; have stores of glycogen that is converted to glucose when needed for bursts of activity.
- Intestines
- Kidney
- Adipose tissue
- Heart

<u>Metabolic principle:</u> Degradation is coupled to the formation of **ATP** (energy store) and **NADPH** (reduction equivalents), that represent sources of free energy for biosynthetic reactions.



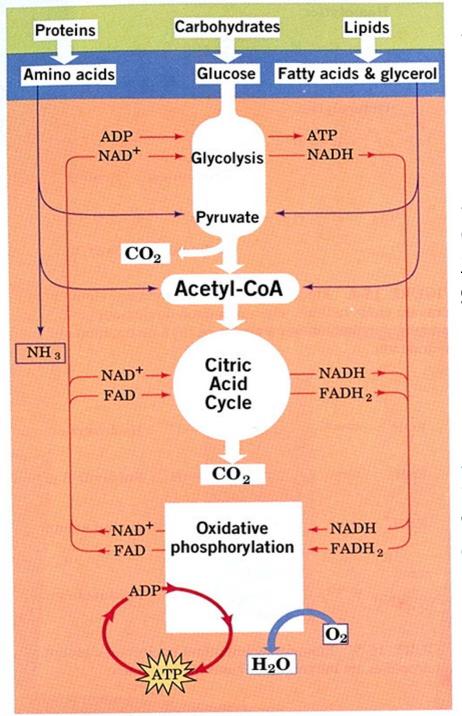


high-enthalpy, low-entropy

Overview of catabolism

low-enthalpy, high-entropy

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Stage 1: Proteins, polysaccharides and lipids are broken down into their component building blocks.

Stage 2: The building blocks are degraded into the <u>common</u> <u>product</u>, generally the acetyl groups of acetyl-CoA.

Stage 3: Catabolism converges to three principal end products: water, carbon dioxide, and ammonia.

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The central role of ATP for energy exchange in biological systems was discovered in 1941 by Fritz Lipmann and Herman Kalckar.

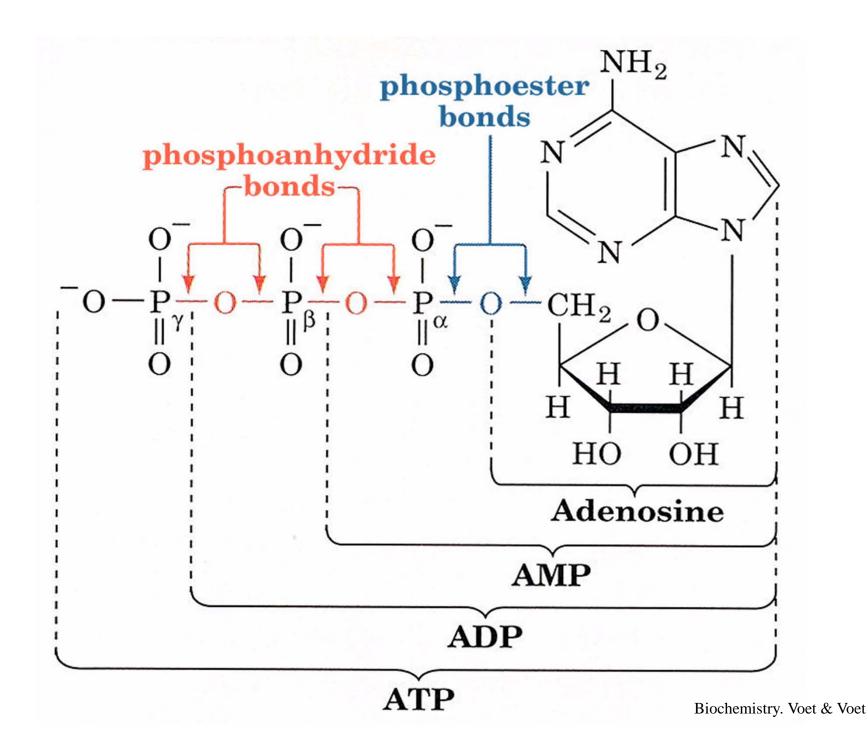
ATP-turnover: 40 kg/day

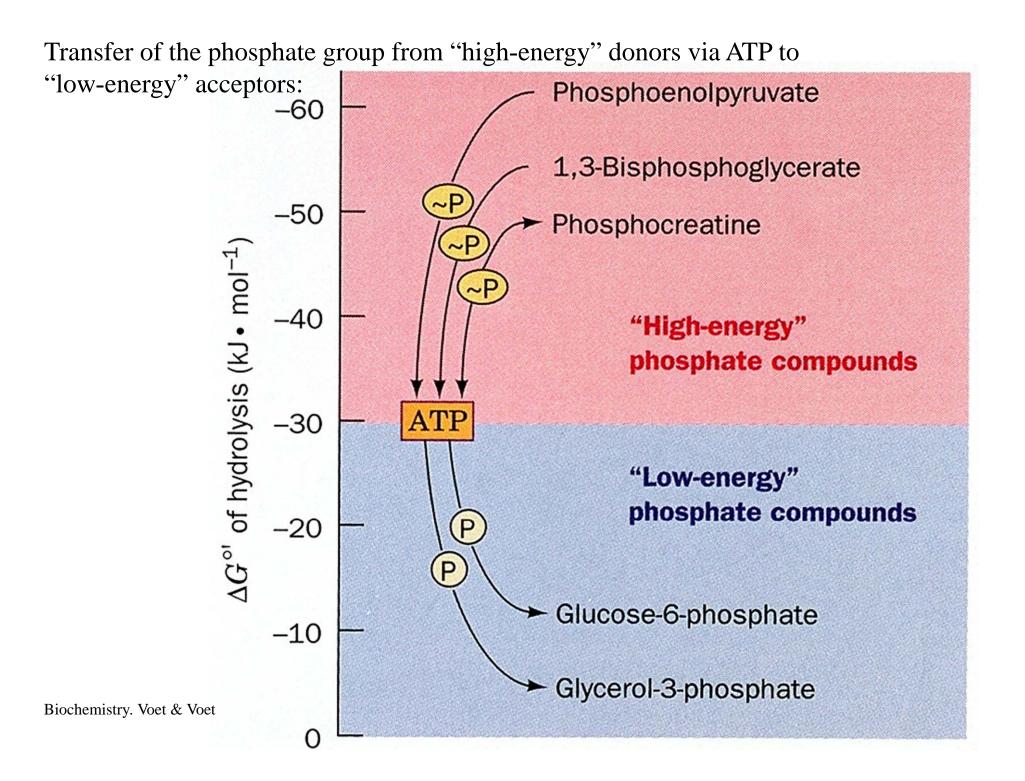
Energy rich compounds: the *triphosphate subunit*, contains 2 *phosphoric acid anhydride linkages*, which upon hydrolysis release high amounts of energy:

 $\begin{array}{ll} \text{ATP} + \text{H}_2\text{O} & \rightleftharpoons \text{ADP} + \text{Pi} + \text{H}^+ & \Delta \text{G}^\circ` = -30,5 \text{ kJ/mol} \\ \text{ATP} + \text{H}_2\text{O} & \rightleftharpoons \text{AMP} + \text{PPi} + \text{H}^+ & \Delta \text{G}^\circ` = -30,5 \text{ kJ/mol} \end{array}$

ATP, ADP, AMP, interconversionable: *adenylate-kinase*: ATP + AMP \Rightarrow ADP + ADP

ATP is **continuously** formed and hydrolysed.





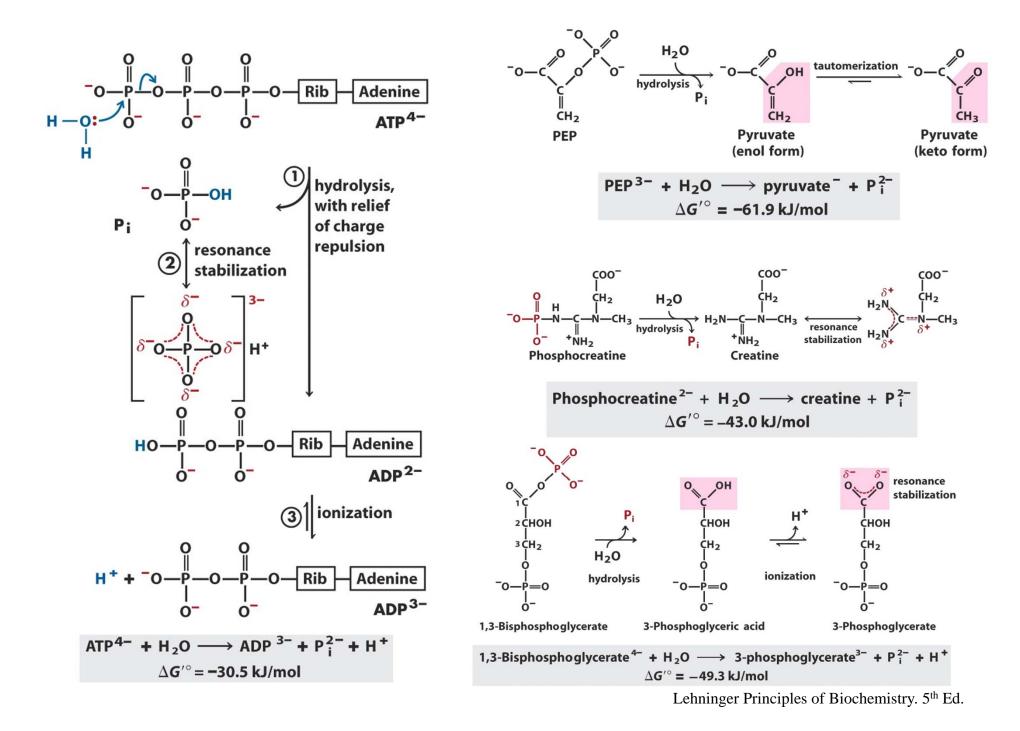


TABLE 13–5Adenine Nucleotide, Inorganic Phosphate, andPhosphocreatine Concentrations in Some Cells

	ATP	ADP [†]	AMP	Ρ _i	PCr		
Rat hepatocyte	3.38	1.32	0.29	4.8	0		
Rat myocyte	8.05	0.93	0.04	8.05	28		
Rat neuron	2.59	0.73	0.06	2.72	4.7		
Human erythrocyte	2.25	0.25	0.02	1.65	0		
E. coli cell	7.90	1.04	0.82	7.9	0		

Concentration (mu)*

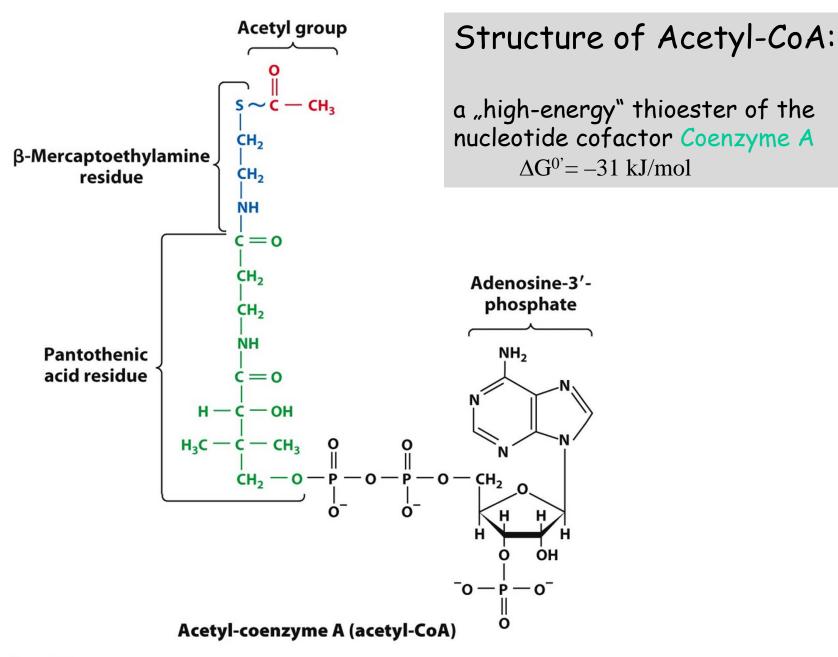
*For erythrocytes the concentrations are those of the cytosol (human erythrocytes lack a nucleus and mitochondria). In the other types of cells the data are for the entire cell contents, although the cytosol and the mitochondria have very different concentrations of ADP. PCr is phosphocreatine, discussed on p. 505.

[†]This value reflects total concentration; the true value for free ADP may be much lower (see Box 13-1).

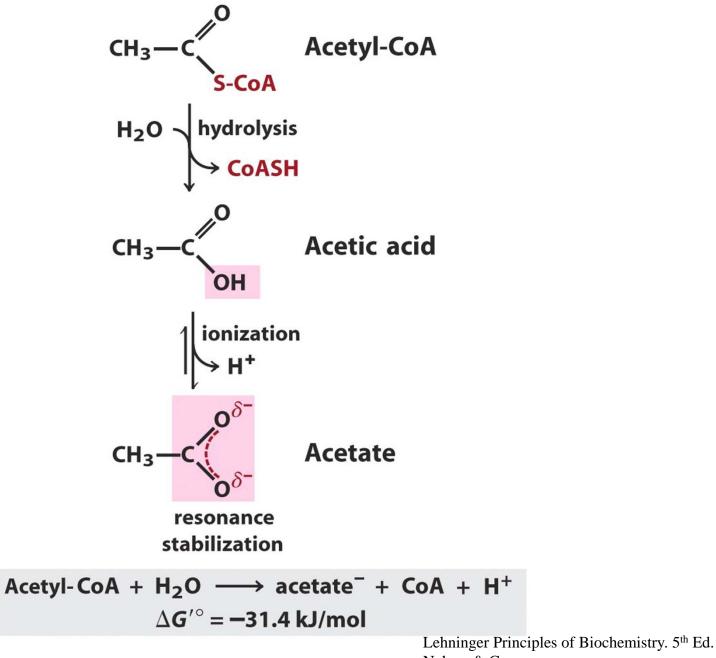
When a sudden demand of energy depletes ATP, the PCr reservoir is used to replenish ATP at a rate considerably faster than possible via catabolism.

Vice versa, when demand of energy is reduced, ATP produced by catabolism is used to replenish the PCr reservoir.

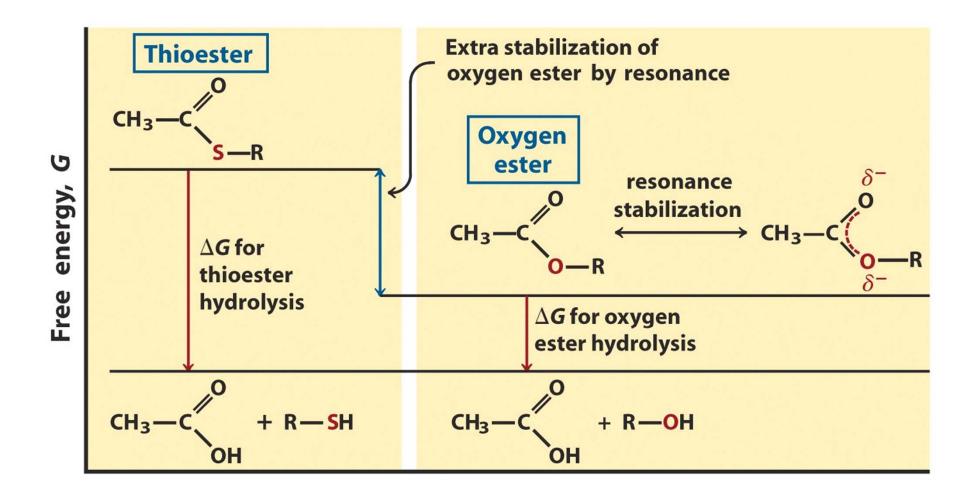
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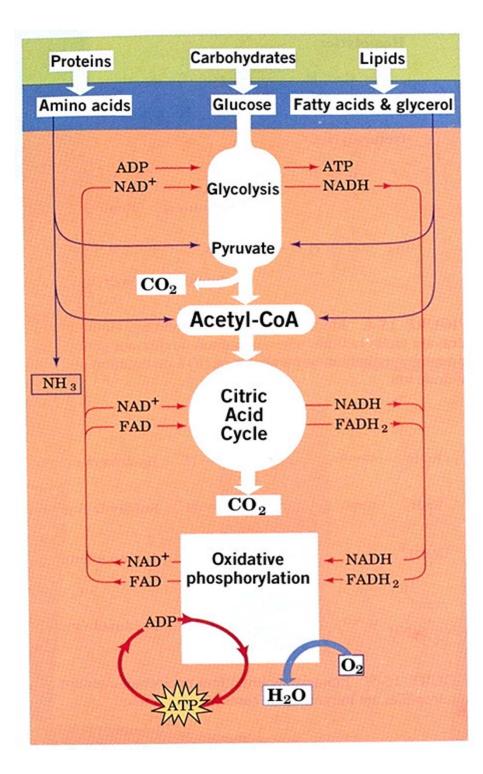




Nelson & Cox



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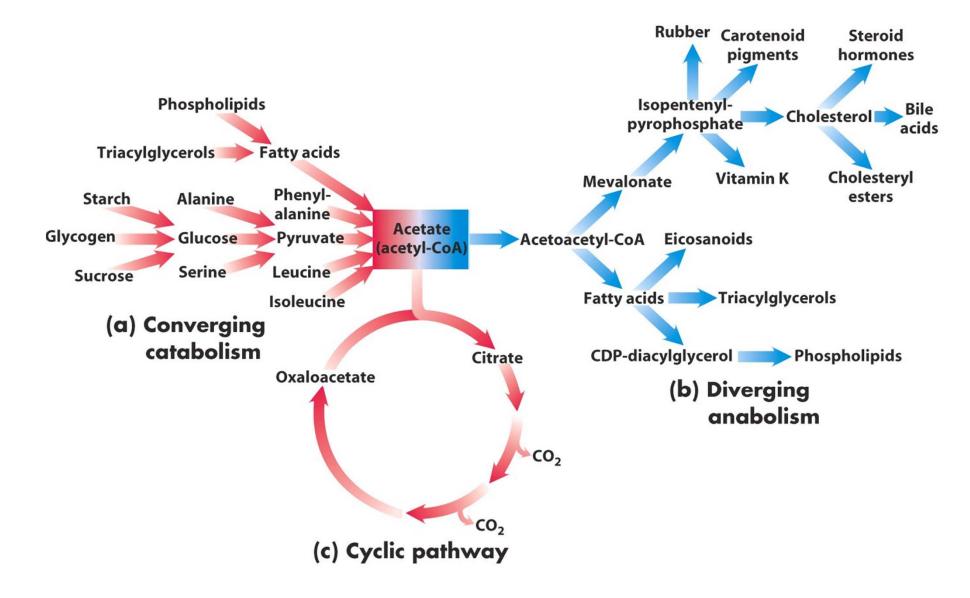
The central role of Acetyl-CoA in

metabolism:

Citric acid cycle Metabolic intermediate: Fatty acid metabolism Carbohydrate metabolism Amino acid metabolism Precursor of cholesterol and steroid hormones Acetyl-group donor: Choline: Acetylcholine (neurotransmitter) Lys of histones

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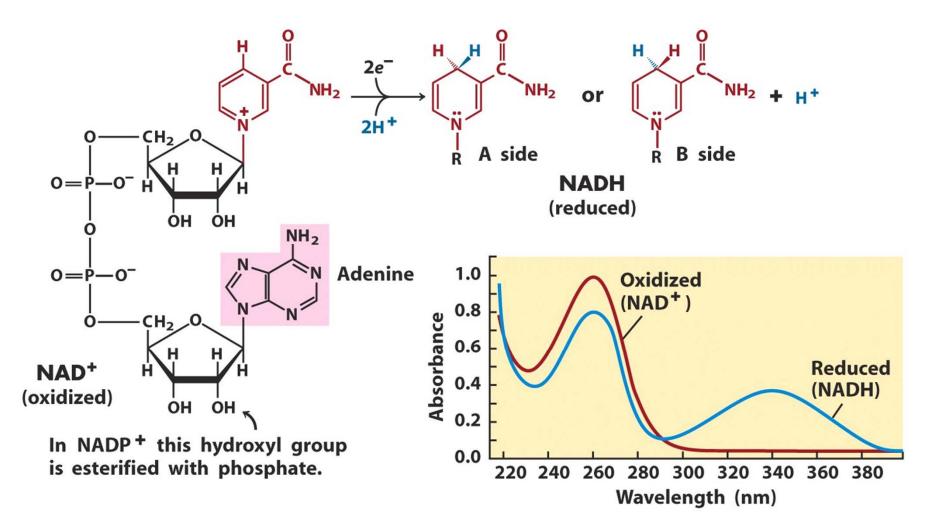
Three types of nonlinear metabolic pathways



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NAD⁺ and NADP⁺ accept e⁻ only pairwise (,,parking" an hydride ion)

Niacin (nicotinic acid, Vit. B3)



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Enzyme	Coenzyme	Stereochemical specificity for nicotinamide ring (A or B)	Text page(s)
Isocitrate dehydrogenase	NAD ⁺	А	610
lpha-Ketoglutarate dehydrogenase	NAD^+	В	610
Glucose 6-phosphate dehydrogenase	NADP ⁺	В	540
Malate dehydrogenase	NAD ⁺	А	612
Glutamate dehydrogenase	NAD ⁺ or NADP ⁺	В	665
Glyceraldehyde 3-phosphate dehydrogenase	NAD ⁺	В	530
Lactate dehydrogenase	NAD ⁺	А	538
Alcohol dehydrogenase	NAD ⁺	А	540

TABLE 13-8Stereospecificity of Dehydrogenases That Employ NAD+ or NADP+ as Coenzymes

Lehninger Principles of Biochemistry. 5th Ed. Nelson & Cox NAD⁺ and NADP⁺, NADH and NADPH are cofactors in more than **200** reactions (type electron transfer):

 $AH_2 + NAD(P)^+ \implies A + NAD(P)H + H^+$

Enzymes: Oxidoreductases/Dehydrogenases

Alcohol-dedydrogenase

 $CH_3CH_2OH + NAD^+ \iff CH_3CHO + NADH + H^+$

 $[NAD^+] \gg [NADH]$

Enables catabolic oxidations

 $[NADP^+] \ll [NADPH]$

Total concentration $\cong [10^{-5} \text{ M}]$

Total concentration $\cong [10^{-6} \text{ M}]$

Enables anabolic reductions

Niacin deficiency causes Pellagra (3 Ds: Dermatitis Diarrhoea Dementia).



An inability to absorb niacin (vitamin B3) or the amino acid tryptophan may cause pellagra, a disease characterized by scaly sores, mucosal changes and mental symptoms

ADAM.



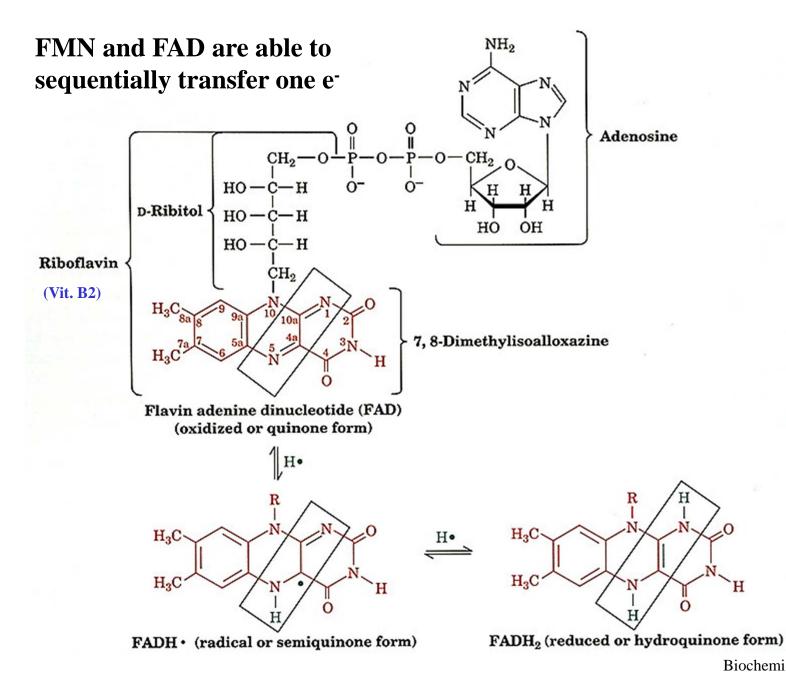


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Fig. 6.8 Clinical findings of macin deficiency before (A) and after (B) therapy in an alcoholic patient.

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TABLE 13–9Some Enzymes (Flavoproteins)That Employ Flavin Nucleotide Coenzymes

	Flavin	Text
Enzyme	nucleotide	page(s)
Acyl-CoA dehydrogenase	FAD	638
Dihydrolipoyl dehydrogenase	FAD	605
Succinate dehydrogenase	FAD	612
Glycerol 3-phosphate dehydrogenase	FAD	714–715
Thioredoxin reductase	FAD	869
NADH dehydrogenase (Complex I)	FMN	696-697
Glycolate oxidase	FMN	767

