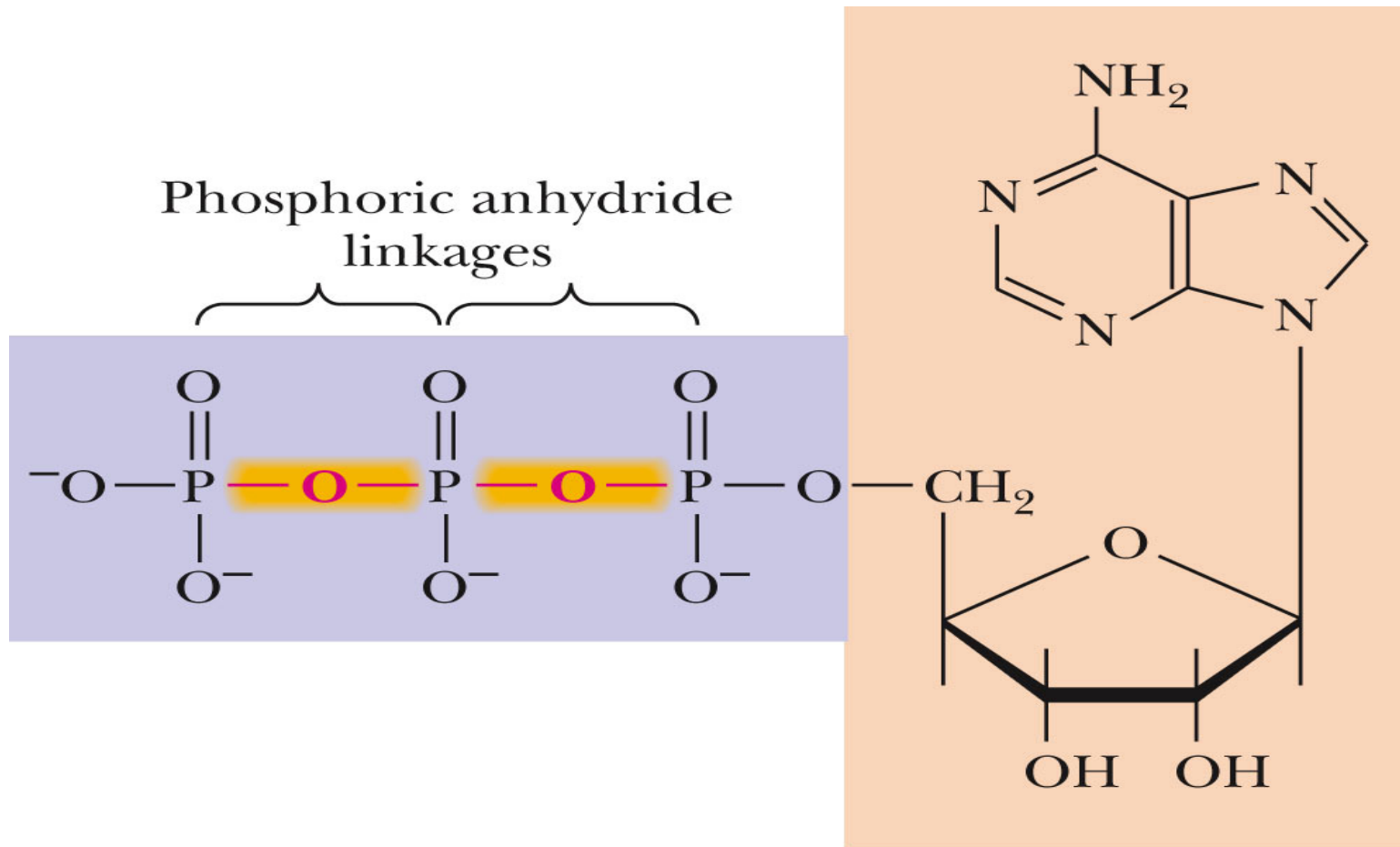


Know the structure of ATP



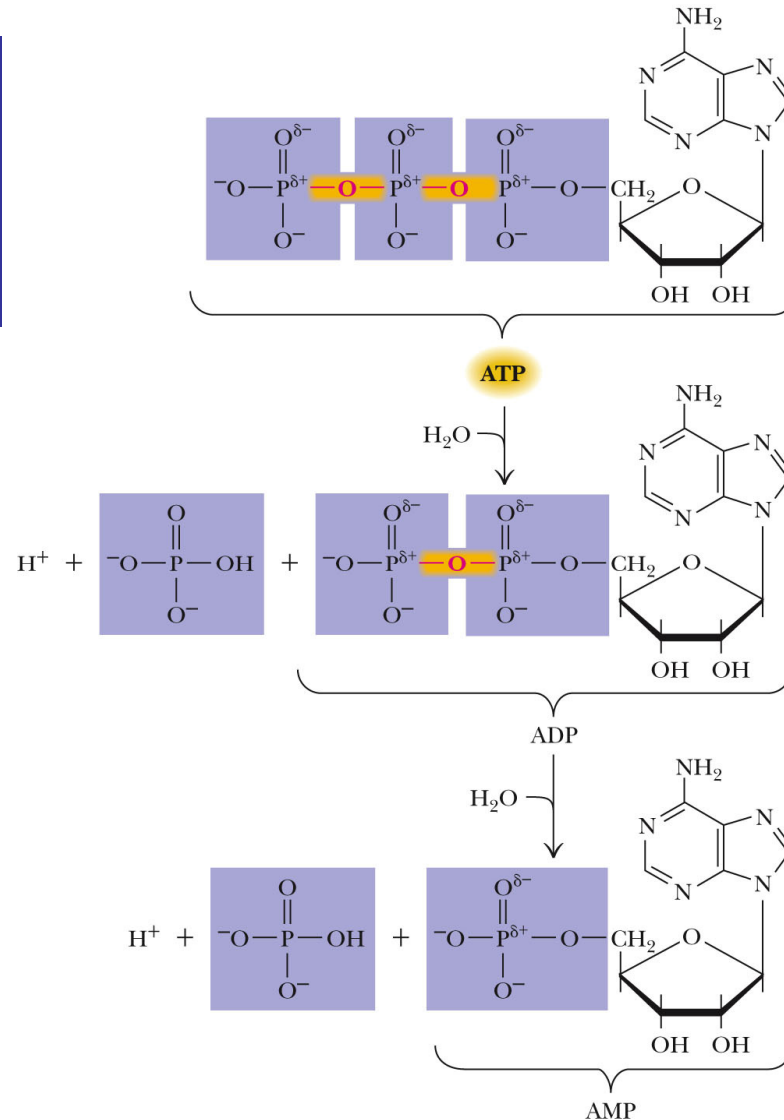
ATP
(adenosine-5'-triphosphate)

Hydrolysis of ATP is favourable

Bond strain in the anhydride due to electrostatic repulsion

Hydrolysis products resonance stabilised

Hydrolysis causes increased entropy

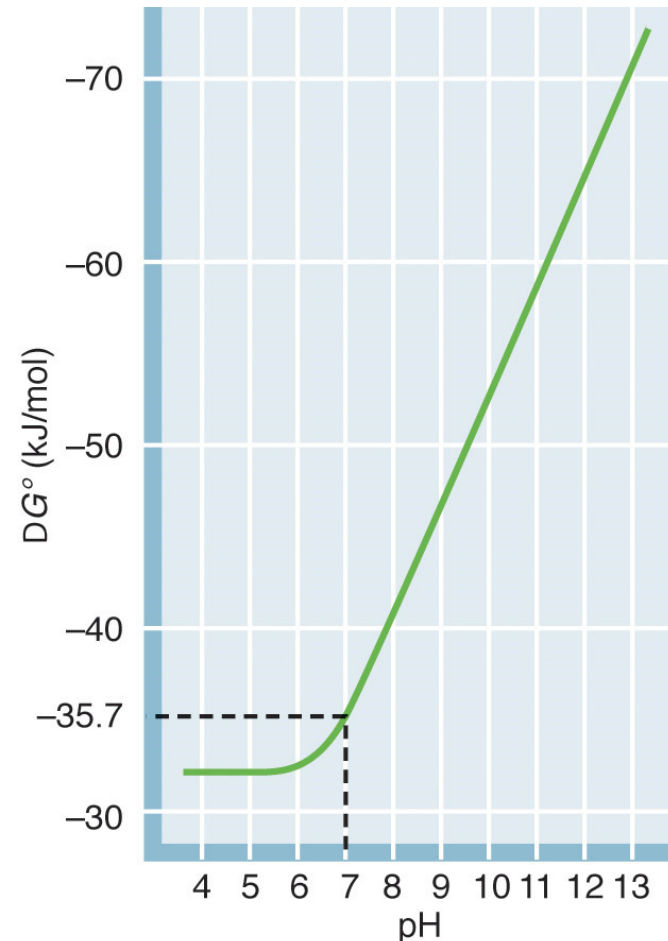


Complex equilibria involved in ATP hydrolysis

- pH dependent

Highest pKa of one of the protons is ~7

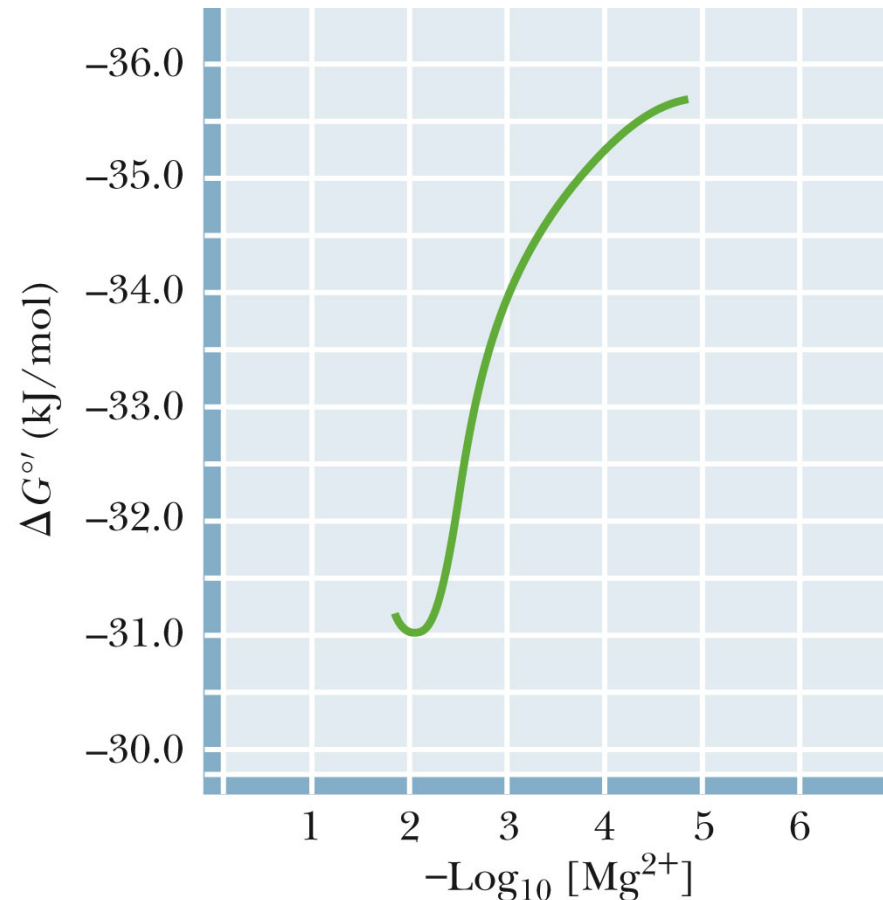
Multiple ionisations of ATP, ADP and phosphate affect the K_{eq} and hence the ΔG° of hydrolysis



Complex equilibria involved in ATP hydrolysis

- Binds metal cations

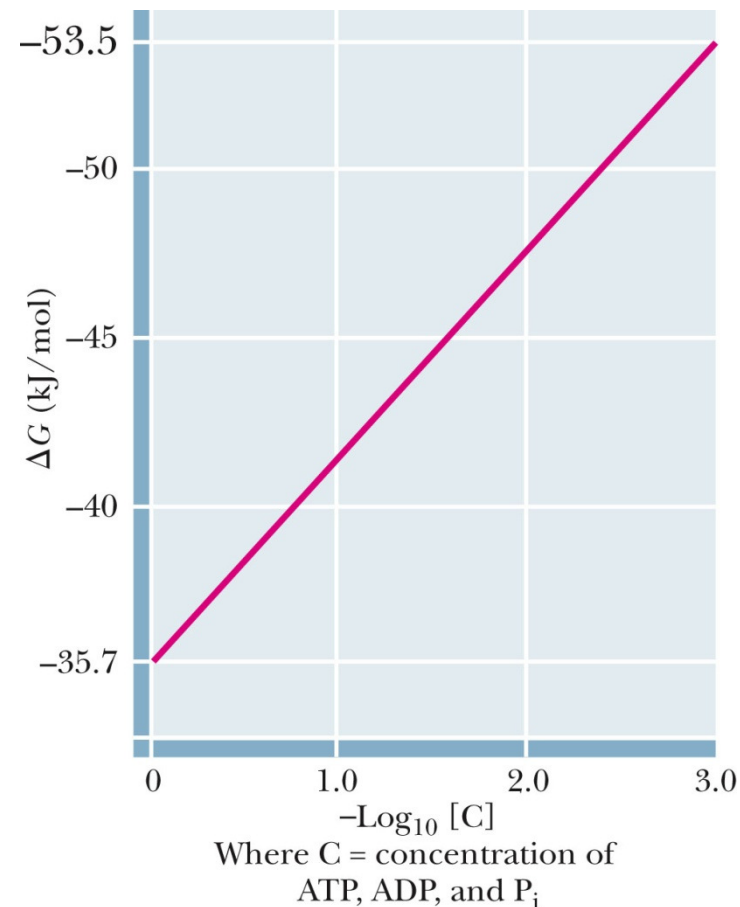
Less negative ΔG° of hydrolysis of ATP when in the presence of physiological concentrations of cations



Complex equilibria involved in ATP hydrolysis

- Far lower concentrations than standard conditions

At physiological concentrations, free energy of hydrolysis of ATP becomes more negative compared to under standard conditions



Daily human requirement for ATP

- Average human consumes:
 - 50% efficiency
 - Moles of ATP used each day:
 - Mr ATP:
 - Mass of ATP hydrolysed per day:
-
- Recycled:
 - Commercial cost:

How ATP changes the Keq

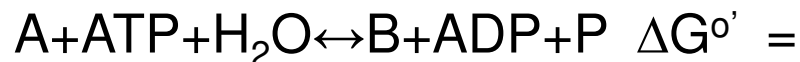
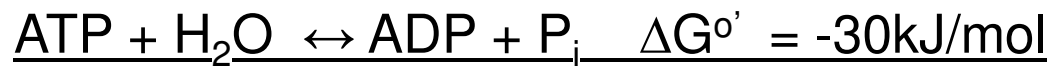
Consider $A \leftrightarrow B$ with $\Delta G^{\circ'} = + 20 \text{ kJ/mol}$

$$\Delta G^{\circ'} = -RT \ln(\text{Keq}) \quad (R=8.31 \text{ J/K.mol}; T=298\text{K})$$

$$\text{Keq} = \frac{[B]}{[A]}$$

1 molecule of product formed for every _____ molecules of reactant

Suppose now, that $A \leftrightarrow B$ is coupled to ATP hydrolysis



$$\text{Keq} =$$

Given that in a typical cell, $[\text{ADP}] = 8 \times 10^{-3}$, $[\text{P}] = 10^{-3}$, $[\text{ATP}] = 8 \times 10^{-3}$

$$[B]/[A] =$$

Metabolism: An Overview

Chapter 17



Overview

- Metabolism:
- Sum of the chemical changes that convert **nutrients (substrate)** into **energy** into **products** via specific chemical **intermediates**
- Hundreds of enzymatic reactions organised into discrete pathways

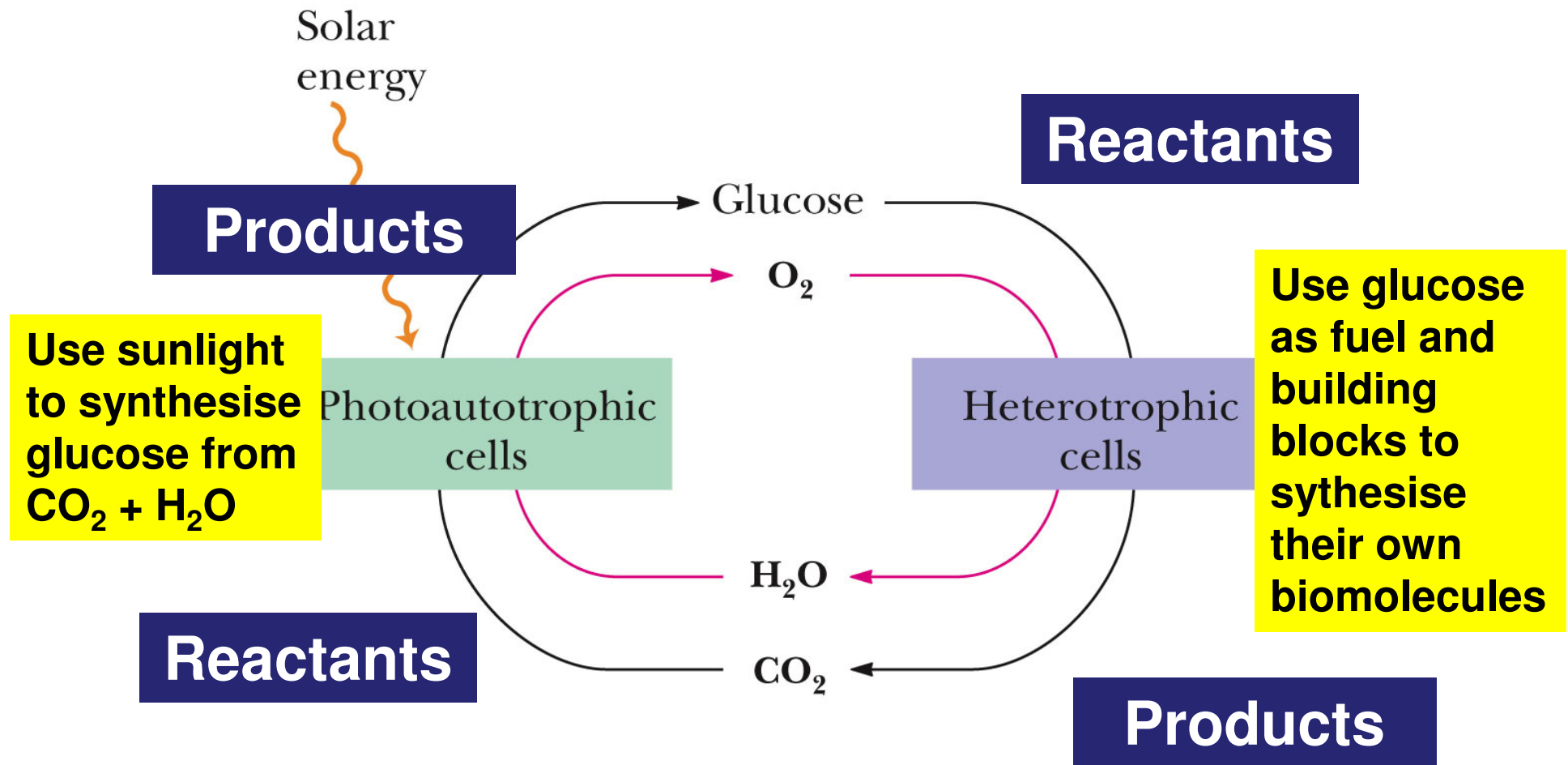
Metabolic Diversity

- Basic set of metabolic pathways
- Different cells and organisms may express alternate pathways

Diversity to metabolic processes

- Organisms are classified by how they obtain **carbon** and **energy**

Flow of energy in the Biosphere



Metabolic maps

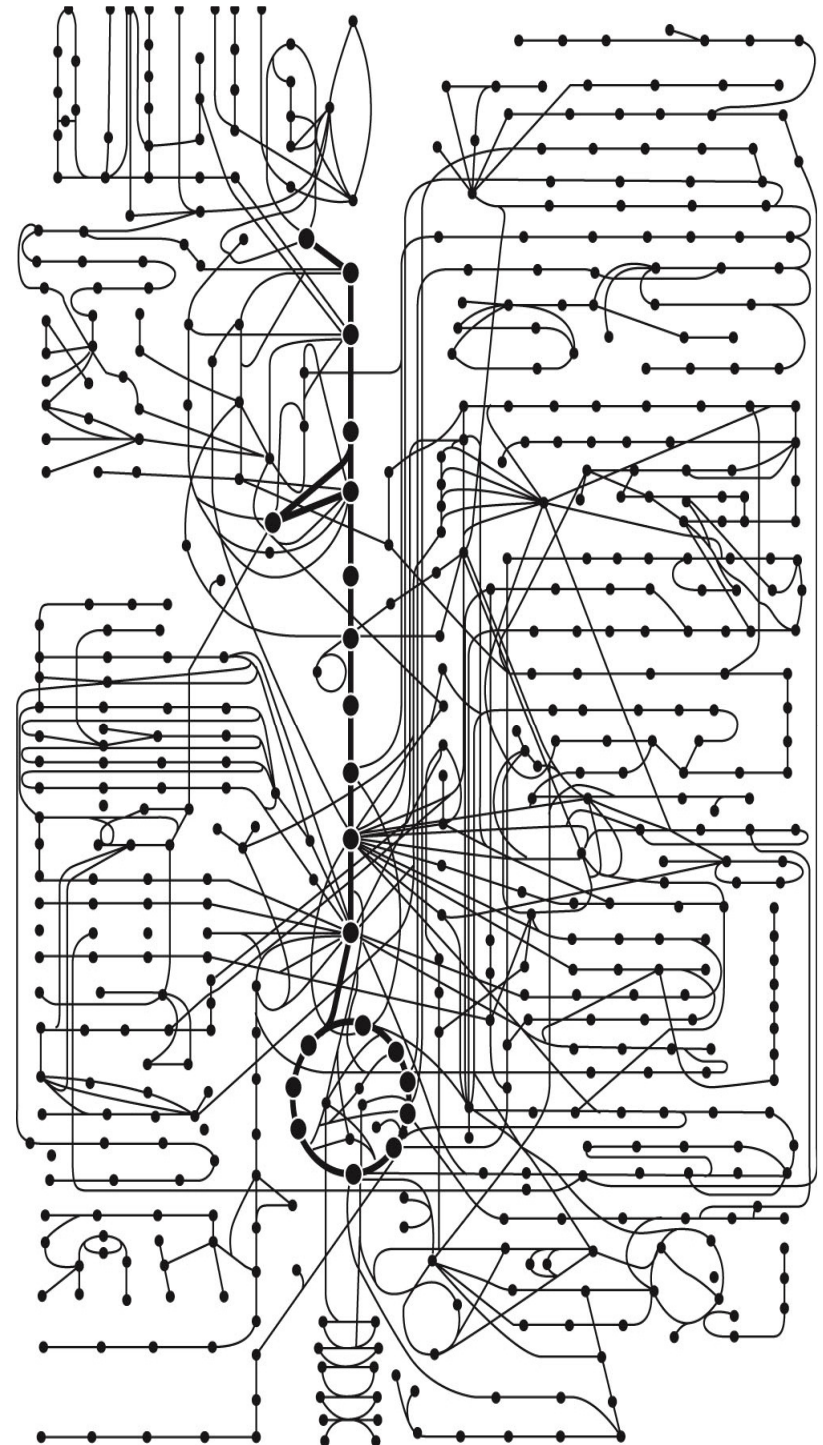
- Indicate the **principal reactions** of metabolism of
- Very complex
- Become easy to follow once the major metabolic routes are known and their functions understood

Carbohydrates
Lipids
Amino acids
Nucleotides

- Simplification: just 2 symbols
- **Dot** =
- **Line** =

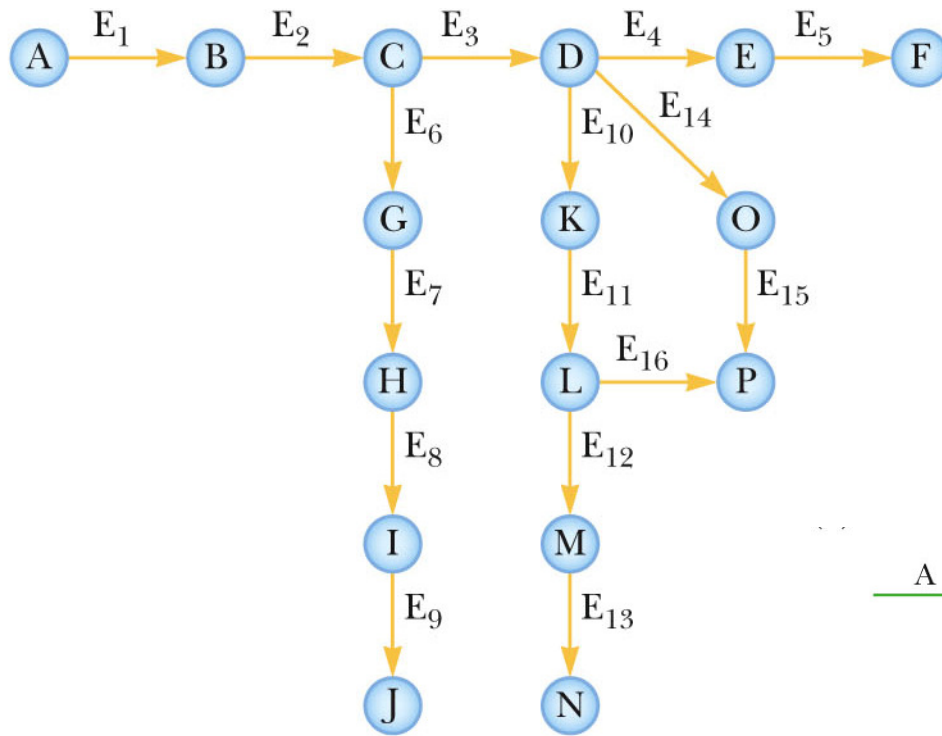
TABLE 17.2 Number of Dots (Intermediates) in the Metabolic Map of Figure 17.2, and the Number of Lines Associated with Them

Lines	Dots
1 or 2	410
3	71
4	20
5	11
6 or more	8



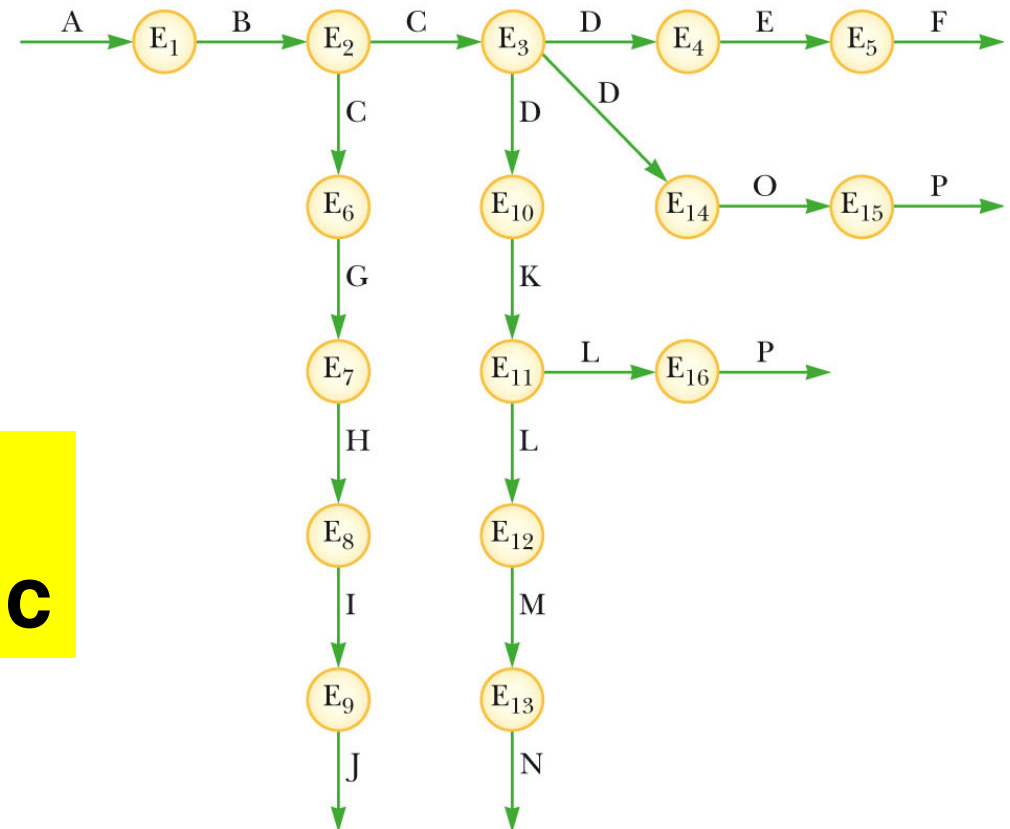
Alternative mapping of metabolic reactions

- Simpler portrayal of chemistry occurring in the organism
- **Spatial** and **temporal** portrayal of metabolites and enzymes
- Recast the metabolic maps in a **protein-centric** rather than metabolite-centric manner



Traditional view:
metabolite-centric

Recent view:
protein-centric



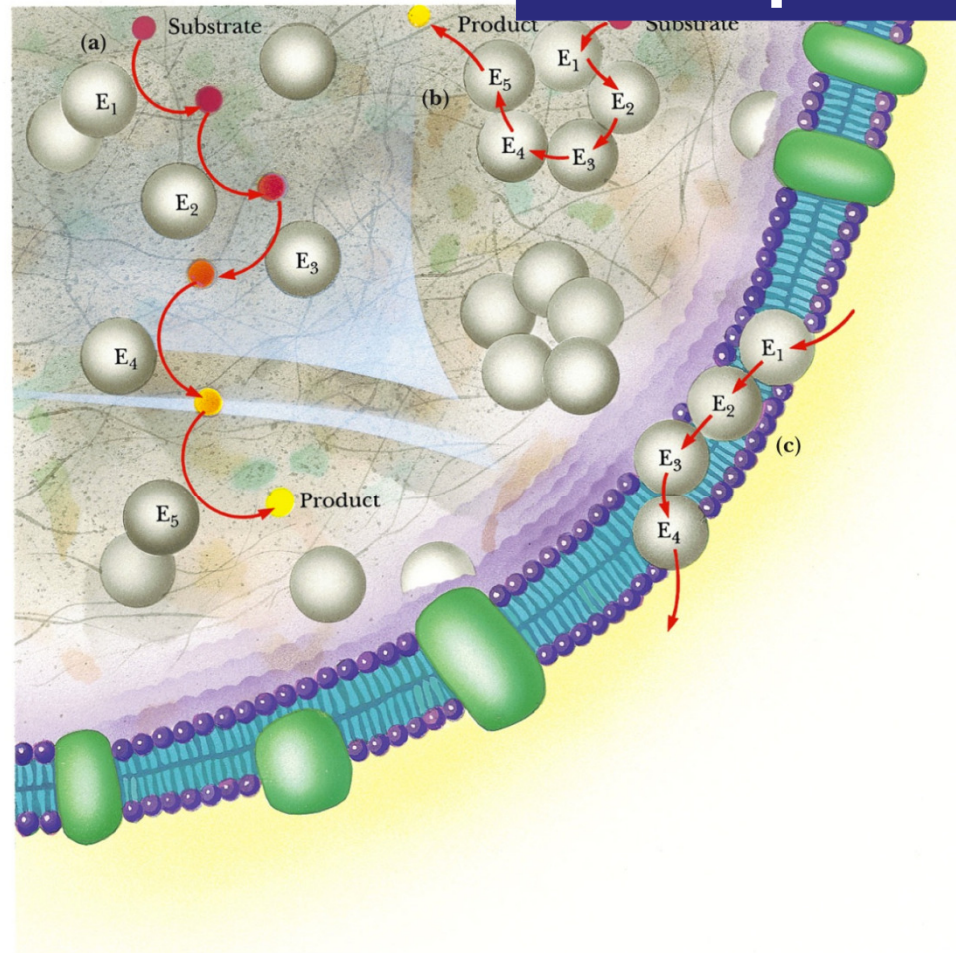
```

graph LR
    A --> E1((E1))
    E1 -- B --> E2((E2))
    E2 -- C --> E3((E3))
    E2 -- C --> E6((E6))
    E3 -- D --> E4((E4))
    E3 -- D --> E10((E10))
    E3 -- D --> E14((E14))
    E4 -- E --> E5((E5))
    E5 -- F --> F_out[ ]
    E6 -- G --> E7((E7))
    E7 -- H --> E8((E8))
    E8 -- I --> E9((E9))
    E9 -- J --> J_out[ ]
    E10 -- K --> E11((E11))
    E11 -- L --> E16((E16))
    E11 -- L --> E12((E12))
    E12 -- M --> E13((E13))
    E13 -- N --> N_out[ ]
    E14 -- O --> E15((E15))
    E15 -- P --> P_out[ ]
    E16 -- P --> P_out
  
```

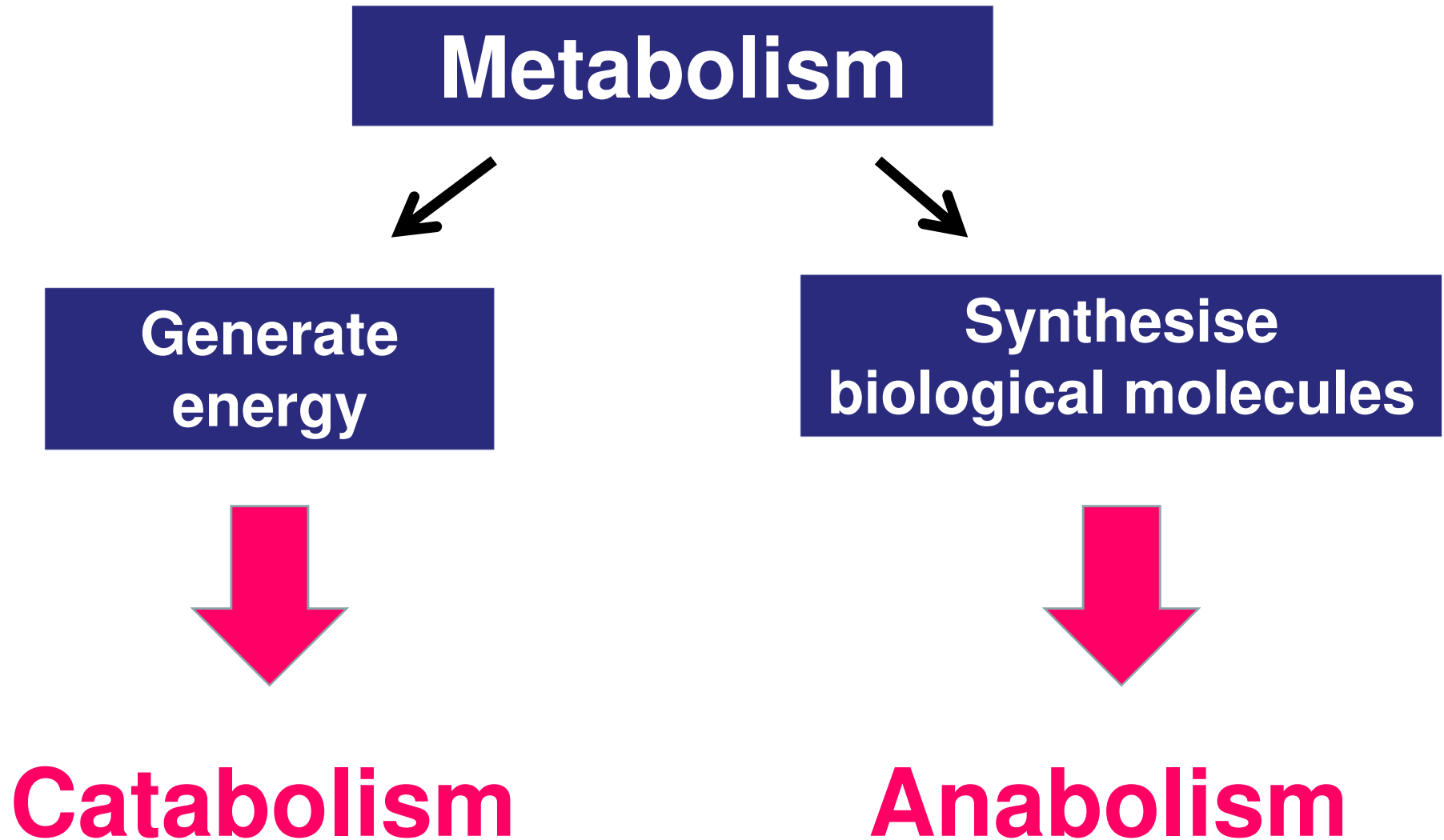
The diagram illustrates the glycolysis pathway, showing the sequence of enzymes and their associated cofactors. The pathway starts with substrate A, which enters the 'Enzymes 1 and 2' box. From this box, a green arrow labeled 'C' leads to a yellow circle labeled E_3 . From E_3 , a green arrow labeled 'D' leads to the 'Enzymes 4 and 5' box. From this box, a green arrow labeled 'F' leads to the right. From the 'Enzymes 1 and 2' box, a green arrow labeled 'C' leads down to the 'Enzymes 6 to 9' box. From E_3 , a green arrow labeled 'D' leads down to the 'Enzymes 10 and 11' box. From E_3 , another green arrow labeled 'D' leads down to the 'Enzymes 14 and 15' box. From the 'Enzymes 14 and 15' box, a green arrow labeled 'P' leads to the right. From the 'Enzymes 10 and 11' box, a green arrow labeled 'L' leads down to the 'Enzymes 12 and 13' box. From the 'Enzymes 10 and 11' box, another green arrow labeled 'L' leads down to a yellow circle labeled E_{16} . From E_{16} , a green arrow labeled 'P' leads to the right. From the 'Enzymes 12 and 13' box, a green arrow labeled 'N' leads down to the bottom. All boxes are yellow with black text, and all circles are yellow with black text. All arrows are green.

Multienzyme systems

Multienzyme
complex

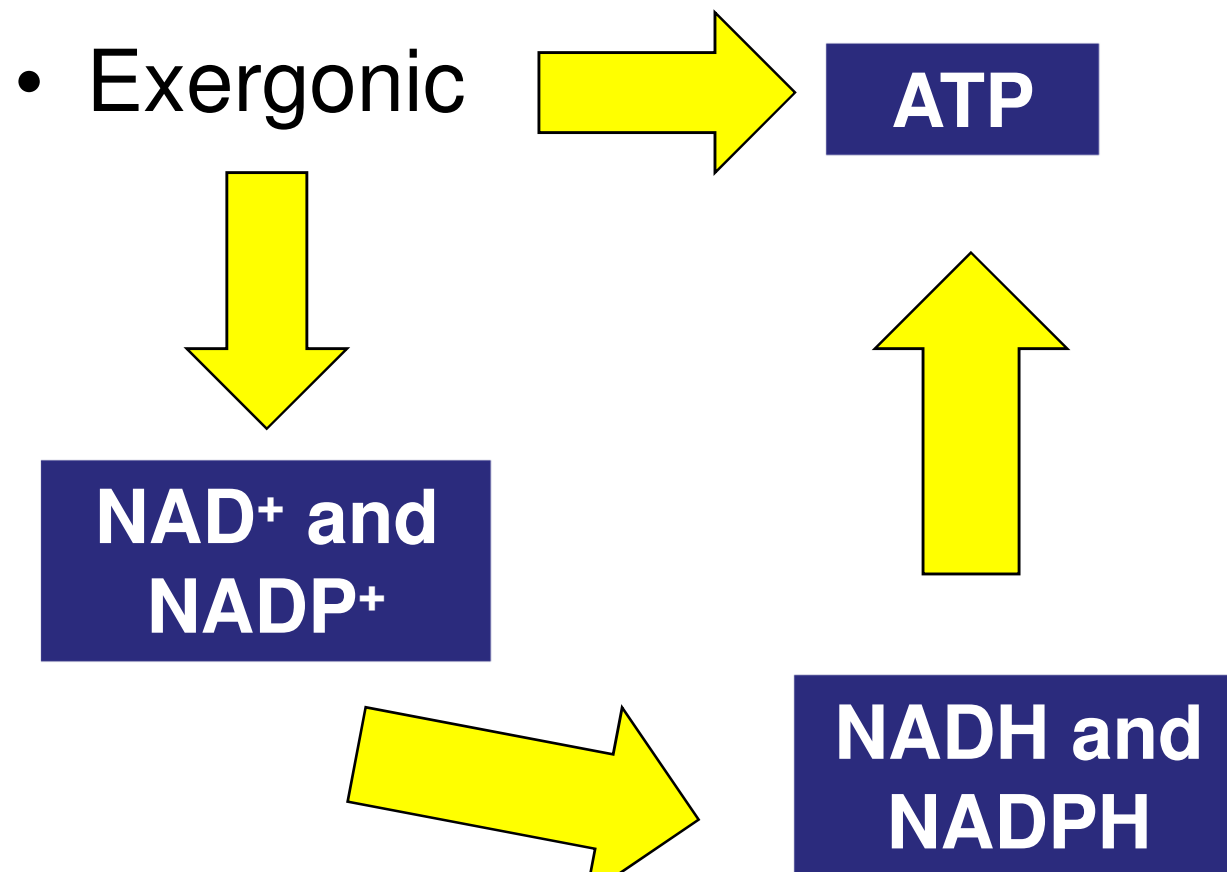


Anabolism and Catabolism



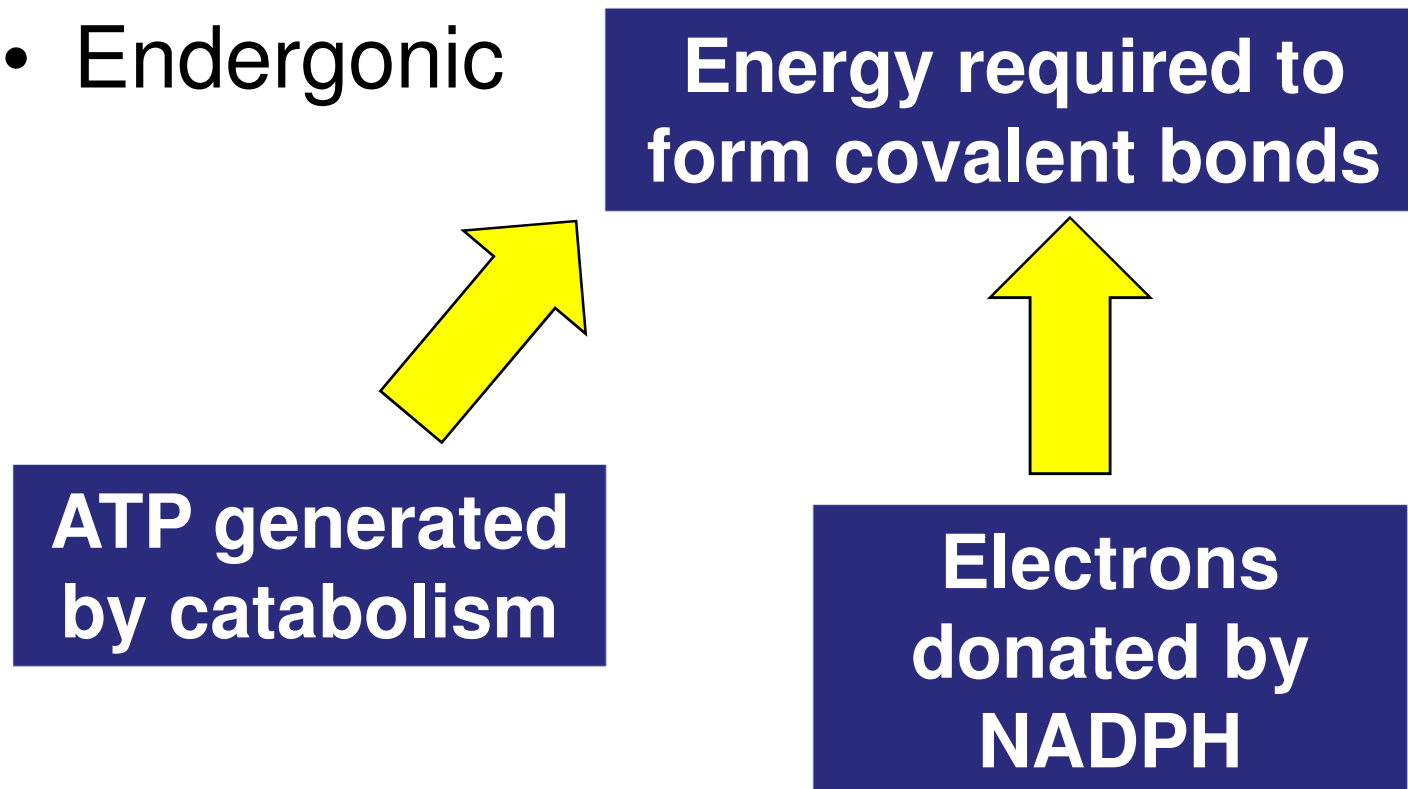
Catabolism

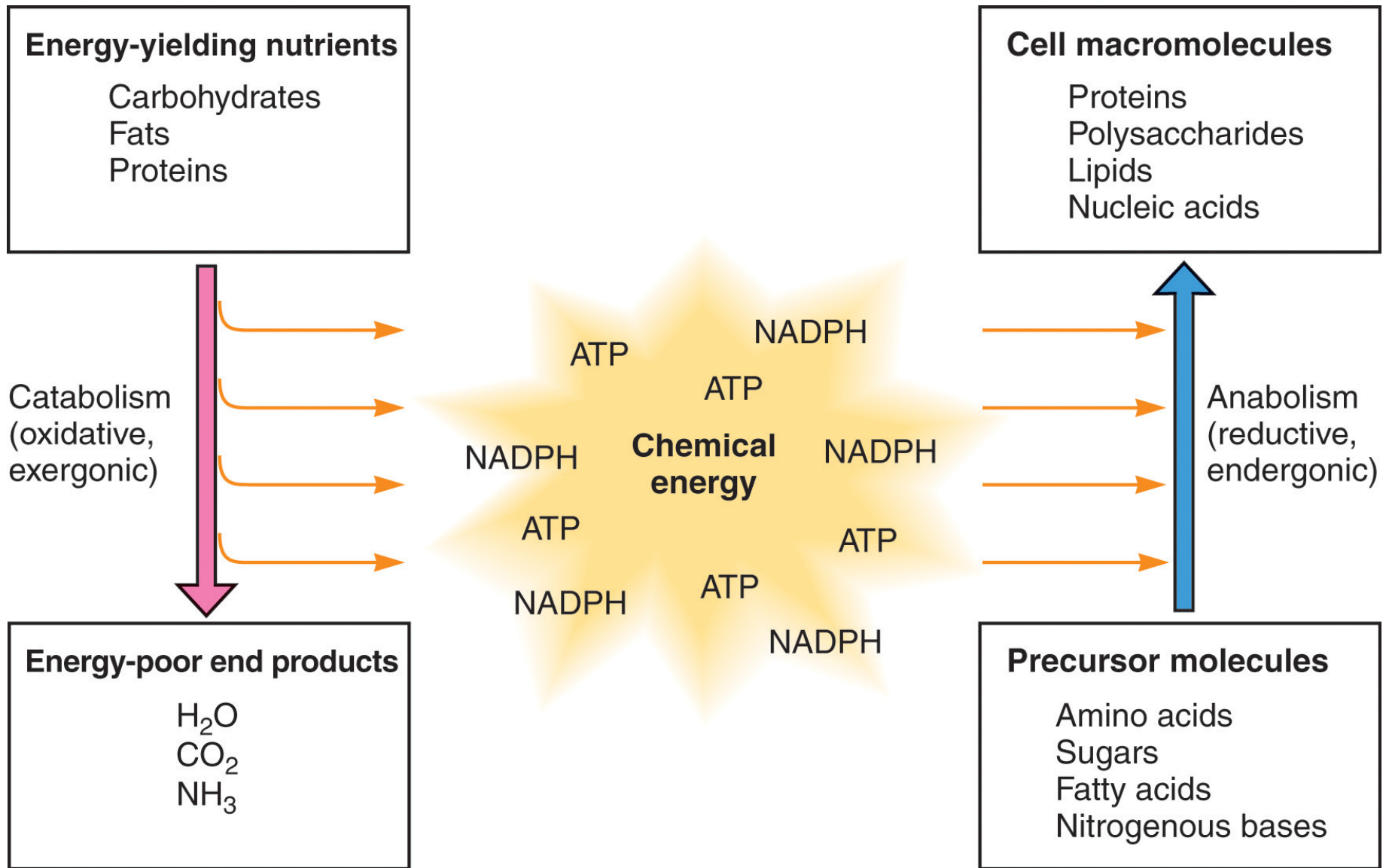
- Breakdown of complex nutrients



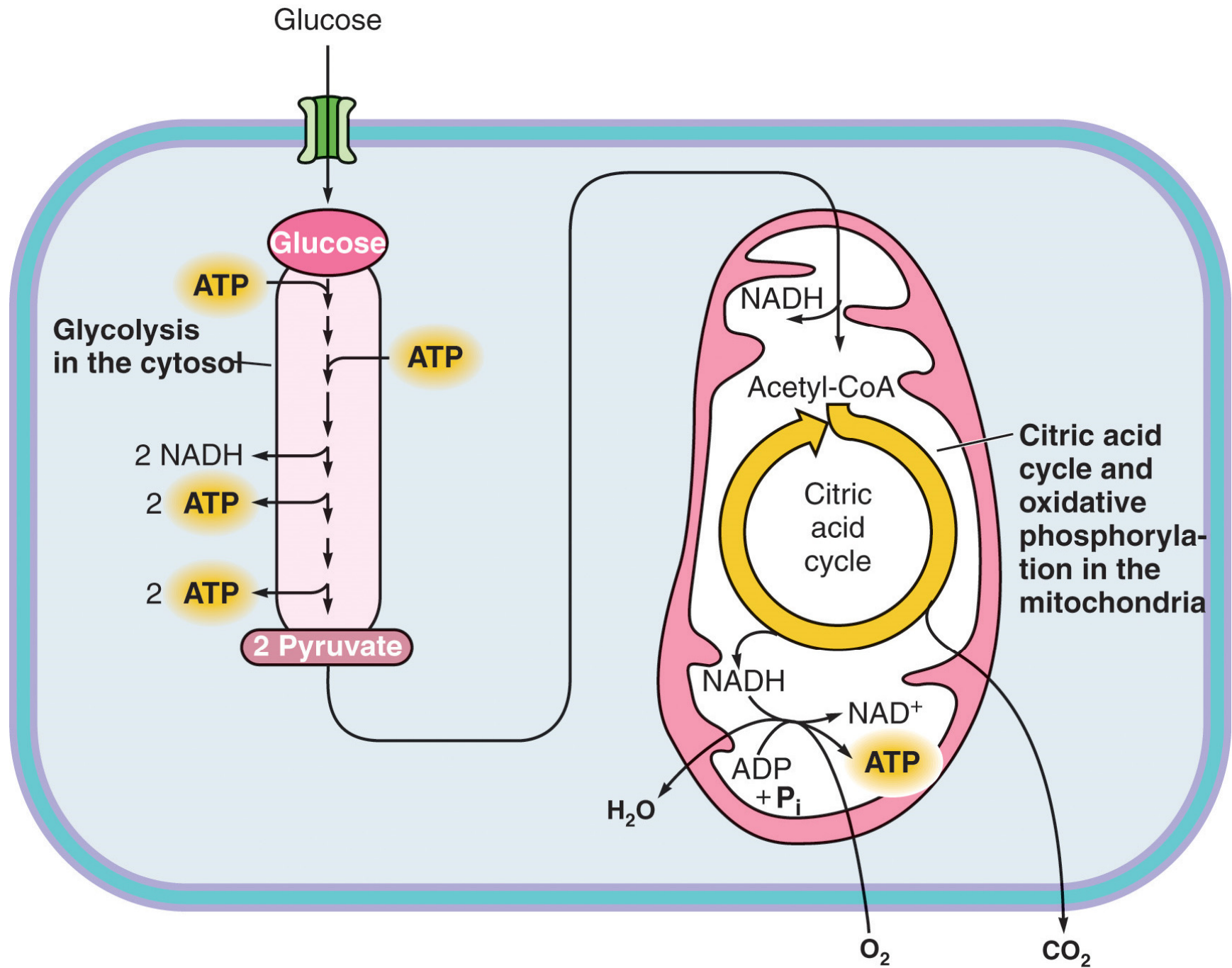
Anabolism

- Synthesis of complex biomolecules from simple precursors
- Endergonic





- Tight control: metabolic needs served in an immediate and orderly fashion
- Competing metabolic pathways localised in **different cellular compartments** to avoid any interference
- E.g. enzymes involved in **catabolism of fatty acids** localised in **mitochondria** & enzymes for **fatty acid biosynthesis** located in **cytosol**



3 Stages of Aerobic Catabolism

STAGE 1

Nutrients degraded into building blocks

STAGE 2

Building blocks degraded further into limited set of metabolic intermediates

STAGE 3

Acetyl-CoA converted to waste via citric acid cycle and oxidative phosphorylation

Stage 1:

The various kinds of proteins, polysaccharides, and fats are broken down into their component building blocks, which are relatively few in number.

Large biomolecules

Proteins

Polysaccharides

Lipids

Building block molecules

Amino acids

Pentoses,
hexoses

Glucose

Glycerol, fatty acids

Stage 2:

The various building blocks are degraded into a common product, the acetyl groups of acetyl-CoA.

Glycolysis

Glyceraldehyde-3-phosphate

Pyruvate

Acetyl-CoA

Common
degradation product**Stage 3:**

Catabolism converges via the citric acid cycle to three principal end products: water, carbon dioxide, and ammonia.

Citric acid
cycle

Oxidative phosphorylation

End
productsSimple, small
end products of
catabolism NH_3 H_2O CO_2

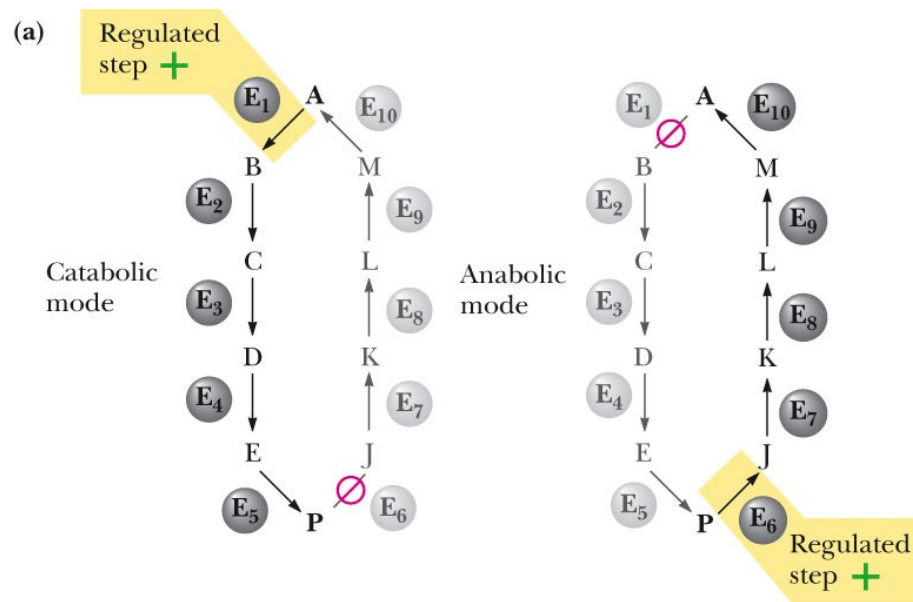
Catabolism, Anabolism and Amphibolism

- Catabolism **converges** to a common intermediate
- Anabolism **diverges** from a small group of simple intermediates to a wide variety of biomolecules
- Some pathways have roles in both catabolism and anabolism and are said to be **amphibolic**

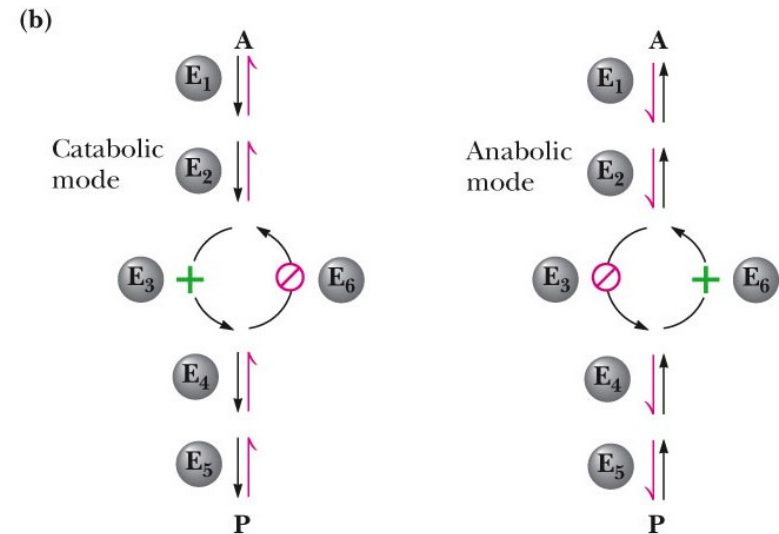
Catabolism and Anabolism

- Anabolic synthesis of a given end product does **not** correspond to the catabolic pathway of the same substance
- Eg catabolism of glucose to pyruvate vs. biosynthesis of glucose from pyruvate

Parallel pathways of catabolism and anabolism must differ by at least one step so that they can be regulated independently



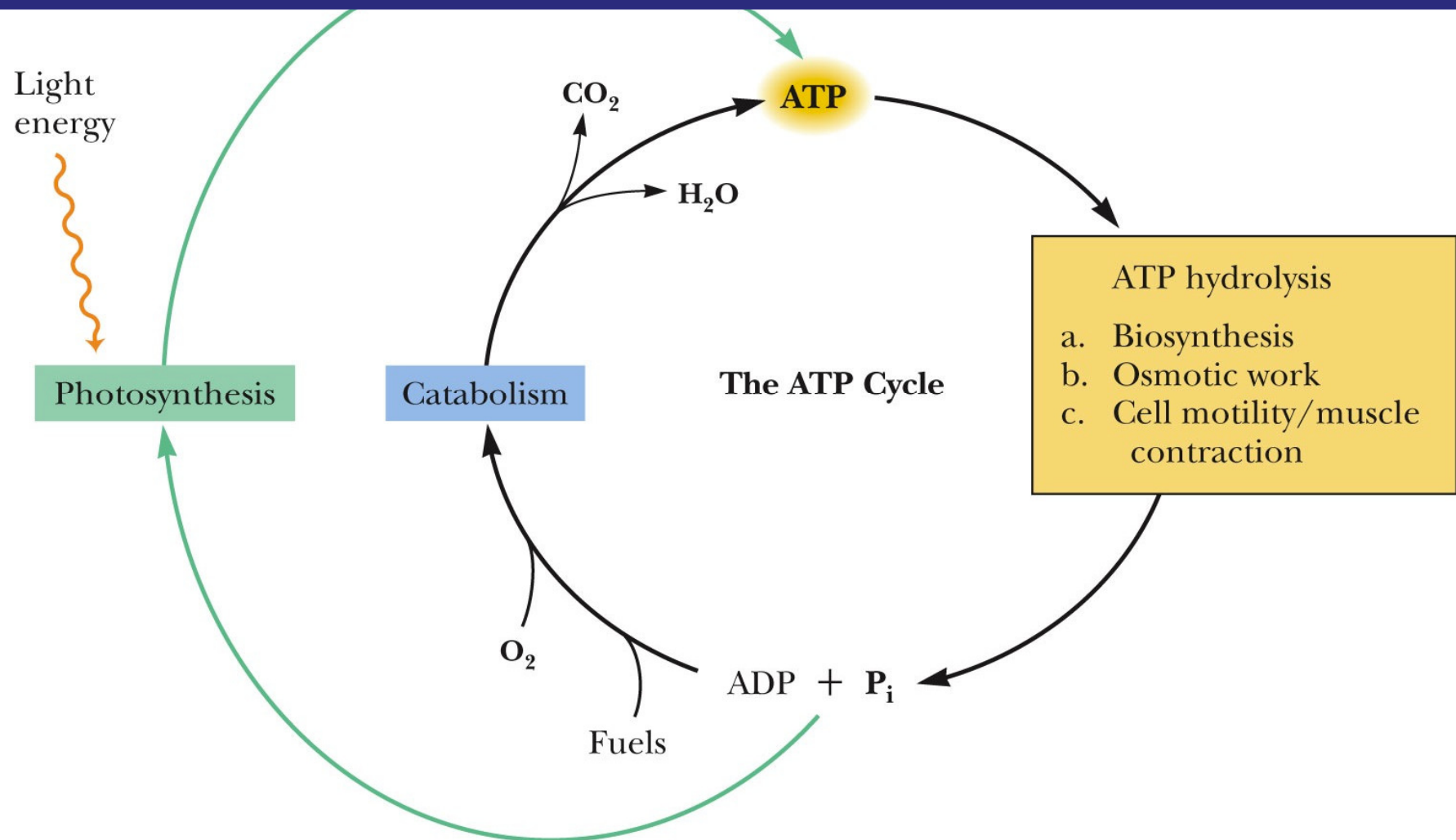
Independent routes between A and P



Only one reaction has 2 different enzymes

The ATP Cycle

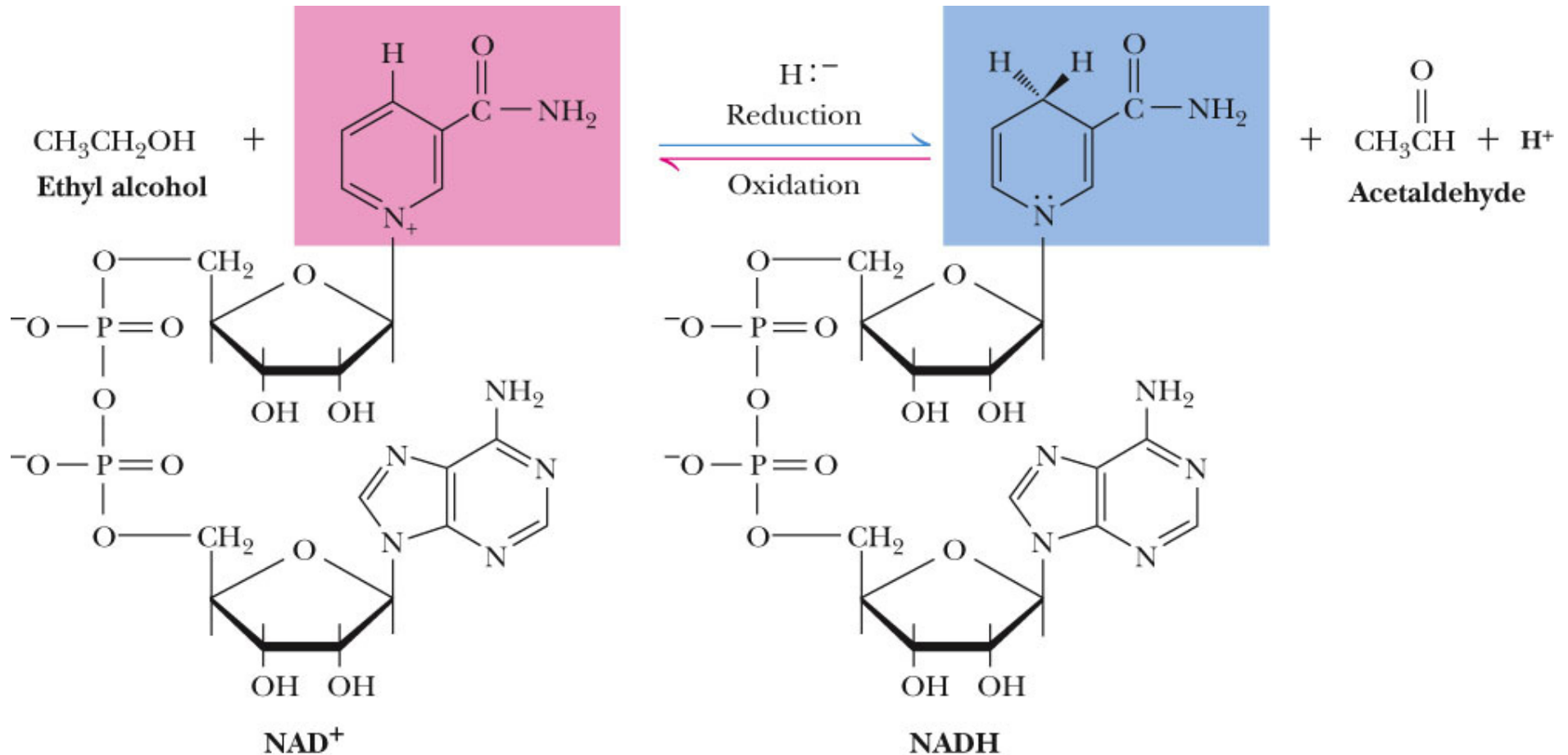
ATP produced by phototrophs and heterotrophs



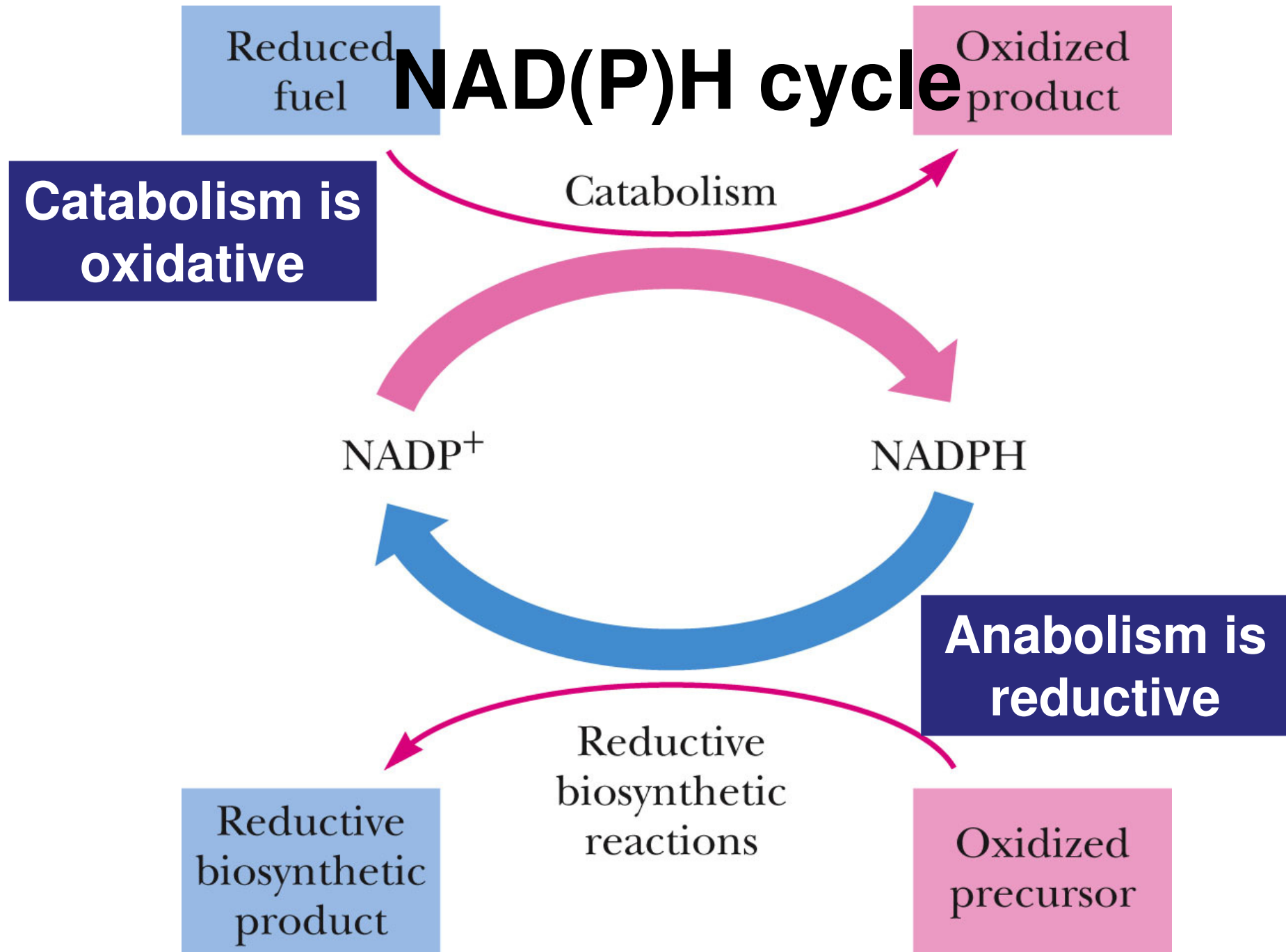
The NAD⁺/NADP⁺ Cycle

- Oxidative reactions of catabolism → **hydride ions** (H :⁻) released
- Hydride ions transferred to NAD⁺
- NAD⁺ **reduced** to NADH
- NADH is **oxidised** to NAD⁺ during anabolism
- Final electron acceptor is O₂ which becomes reduced to water

Hydride transferred to NAD⁺ to form NADH + H⁺



Nicotinamide Adenine Dinucleotide



Coenzymes and Vitamins

E.g. NAD^+
and NADP^+

“Vital
amines”

- Vitamins cannot be synthesised by the organism
- Different organisms require different vitamins
- Low Mr substances that bring **unique chemical functionality** to enzymatic reactions