THERMODYNAMICS OF METABOLISM

- Overall process of catabolism RELEASES energy
- Overall process of anabolism REQUIRES energy input

How do we define amount of energy?

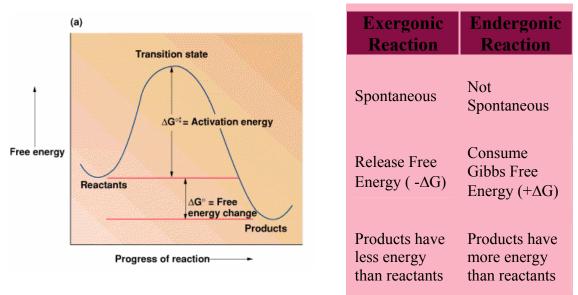
Let's look at a hypothetical spontaneous chemical reaction:

The quantity of usable energy (chemical potential) in a reaction is called the **Gibbs Free Energy** (ΔG). ΔG is the difference between the energy contained in the products of a reaction and the reactants:

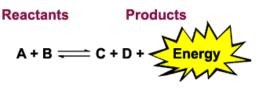
ΔG = (energy of products) - (energy of reactants)

Chemical reactions are classified as being either **exergonic** or **endergonic**. That just means that a reaction can either **release** energy useful for work (an **exergonic** reaction) or **requires** energy to proceed (an **endergonic** reaction).

The spontaneous reaction above is an **exergonic** reaction and ΔG will be negative. Thus, a negative ΔG value tells you that that reaction is possible



- Use ΔG Thermodynamic term
 - Measure of energy available to do useful work
 - ΔG is a constant for a reaction under a defined set of conditions
- We will use ΔG° The STANDARD free energy change
 - Energy change occurring when a reaction proceeds under standard biological conditions
 - 1 ATM, 25°C and all reactants at 1M
 - $\circ~$ Not really biologically relevant so we define ΔG° as ΔG° at pH 7.0, 1 ATM, 25°C and all reactants at 1M
 - Since standard conditions DO NOT occur in cells, absolute values of ΔG° are not realistic. Can be used to COMPARE energy requirements.



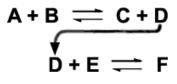
- ΔG° ' defines the difference between the energy content of products and reactants
 - $\circ \Delta G^{\circ}$ Energy of products minus reactants

$$\Delta \mathbf{G}^{\circ} = \mathbf{G}^{\circ}_{\text{products}} - \mathbf{G}^{\circ}_{\text{reactants}}$$

- Common in biological reactions to **COUPLE** unfavorable reactions with favorable ones to make the overall reaction favorable!
 - We have already discussed that some reactions are, while others are endergonic. To get the energy to those endergonic reactions, they are **paired up** with energy-releasing exergonic reactions. Like a locomotive that gets the train car over the hill, an exergonic reaction can "pull" an endergonic reaction along to its destination (products). The reactions can be hooked together, or coupled, via a common intermediate.



Thermodynamics lets us predict that the reaction will proceed if the overall ΔG of the two reactions is negative. When two reactions are coupled, the overall ΔG is the sum of the ΔG s of the component reactions.



ΔG° VALUES ARE ADDITIVE!

- Reactions that are not spontaneous can be coupled to spontaneous reactions
- Under these conditions, the overall reaction can be spontaneous (favorable)

٠	A + B	\rightarrow C + D	ΔG° reaction 1
•	D + E	$\rightarrow \underline{F}$	$\Delta \underline{G^{\circ}}$ reaction 2
•	A + B +	$E \rightarrow C + F$	ΔG° overall reaction

- ΔG° overall reaction = ΔG° reaction 1 + ΔG° reaction 2
- The coupling of metabolism, particularly to ATP hydrolysis, can drive thermodynamically unfavorable reactions forward.
- An example:
 - 1. glucose-6-phosphate \rightarrow fructose-6-phosphate $\Delta G^{\circ\circ} = +1.7 \text{ kJ mol-1}$
 - 2. fructose-6-phosphate + ATP \rightarrow fructose-1,6-bisphosphate + ADP $\Delta G^{\circ\circ} = -14.2 \text{ kJ mol-1}$

OVERALL REACTION:

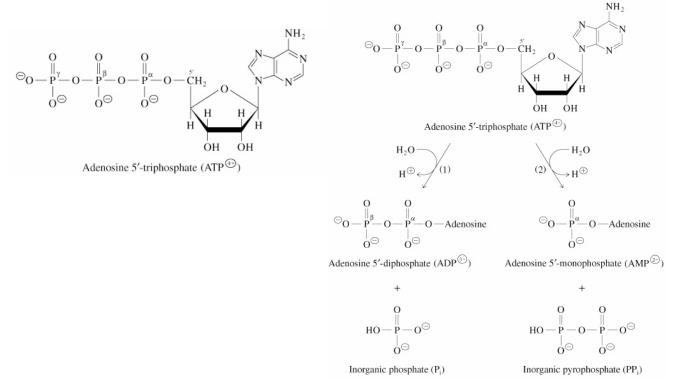
• glucose-6-phosphate + ATP \rightarrow fructose-1,6-bisphosphate + ADP $\Delta G^{\circ\circ} = -12.5 \text{ kJ mol}-1$

ATP AS AN ENERGY SOURCE

- ATP is a common form of "energy currency" within cells
- Commonly use ATP hydrolysis to drive unfavorable reactions

 $ATP + H_20 \rightarrow ADP + Pi + H+$

- ATP hydrolysis has a high ΔG° (-30.5 kJ mol-1)
- The equilibrium for this reaction lies heavily towards the products



o A large amount of energy is released in the hydrolysis of the phosphoanhydride bonds of ATP

Energy of phosphoanhydrides

- (1) <u>Electrostatic repulsion</u> among negatively charged oxygens of phosphoanhydrides of ATP
- (2) <u>Solvation of products</u> (ADP and P_i) or (AMP and PP_i) is better than solvation of reactant ATP
- (3) <u>Products are more stable than reactants</u> There are more delocalized electrons on ADP, P_i or AMP, PP_i than on ATP