

The Second Law of Thermodynamics allows to determine which processes are spontenouis.

$$\Delta S_{univ} = \Delta S_{sys} + \Delta S_{surr} \geq 0$$

The problem with applying such criteria is that you must account for the whole Universe.

Surrounding stays in equilibrium at temperature T and pressure p.



There are two natural tendencies behind spontaneous processes:

- the tendency for a *lower energy state*

- the tendency toward a *more energy dispersed state*

Spontaneity involves

ΛS

Heat that flows into or out of the system changes the entropy of the surroundings.

The entropy of the surroundings.
$$\Delta S_{surr}\Big|_{T=const} = \frac{-q_{surr}}{T}$$

$$q_{surr}\Big|_{p=const} = -q_{p} = \Delta H_{sys}$$

$$\Delta S_{\rm surr} = \frac{-q_{\rm surr}}{T} = \frac{-\Delta H_{\rm sys}}{T}$$

$$\Delta S_{\rm univ} = \Delta S_{\rm sys} + \Delta S_{\rm surr}$$

$$\Delta S_{\rm univ} = \Delta S_{\rm sys} + \frac{-\Delta H_{\rm sys}}{T}$$

$$-T\Delta S_{\rm univ} = -T\Delta S_{\rm sys} + \Delta H_{\rm sys}$$



Definiton of a new function - *Gibbs free energy - it allows to focus on the system.*

$$\Delta G_{sys} = -T\Delta S_{univ}$$
$$\Delta G_{sys} = -T\Delta S_{sys} + \Delta H_{sys}$$

At constant p and T, ΔG_{sys} can be used in the place of $\Delta S_{universe}$ to predict the direction of a process.

The surrounding enters only in a generic, anonymous way, through one number, its temperature T.



Gibbs 1839-1903 Professor of Mathematical Physics, Yale University

The enthalpic part of Gibbs free energy is a measure of the average potential energy of interaction between molecules, and the entropic part is a measure of the order or intermolecular correlations.



Extensive property State function

 ΔG allows us to predict whether a process is spontaneous or not (under constant T and p):

 $\Delta G < 0 \Rightarrow$ spontaneous in forward direction $\Delta G > 0 \Rightarrow$ non-spontaneous in forward direction $\Delta G = 0 \Rightarrow$ at equilibrium

 ΔG is the energy which is "free" to do work.

Second law restated for a system

For any spontaneous process or reaction: $\Delta G_{sys} < 0$

$$Gibbs \ Free \ Energy \ of \ reaction$$

$$aA + bB + \dots \rightleftharpoons gG + hH \dots$$

$$\Delta G = G(\text{final state}) - G(\text{initial state})$$

$$\Delta G = g \ \mu_{\text{G}} + h \ \mu_{\text{H}} + \dots - a \ \mu_{\text{A}} - b \ \mu_{\text{B}} - \dots$$

$$from \ \mu_{\text{i}} = \mu_{\text{i}}^{0} + RT \ln C_{\text{i}} \text{ it follows:}$$

$$\Delta G = g \ \mu_{\text{G}}^{0} + h \ \mu_{\text{H}}^{0} + \dots - a \ \mu_{\text{A}}^{0} - b \ \mu_{\text{B}}^{0} - \dots + RT \ln \frac{[G]^{\text{g}}[\text{H}]^{\text{h}} \dots}{[\text{A}]^{\text{a}}[\text{B}]^{\text{b}} \dots}$$

$$\Delta G = \frac{\Delta G^{0}}{\Delta G^{0}} + RT \ln \frac{[G]^{g}[H]^{h} \dots}{[A]^{a}[B]^{b} \dots}$$

The direction of the reaction depends upon

- Energy content of participants
- Reactant-to-product ratio

 ΔG does <u>not</u> indicate the rate of the process.

A reaction in an equilibrium

$$0 = \Delta G^{0} + RT \ln \left(\frac{[G]^{g}[H]^{h} \dots}{[A]^{a}[B]^{b} \dots} \right)_{Eq} \Delta G^{0} = -RT \ln \left(\frac{[G]^{g}[H]^{h} \dots}{[A]^{a}[B]^{b} \dots} \right)_{Eq} = -RT \ln K$$

The *standard free-energy of reaction* (ΔG^{0}) is the free-energy change for a reaction when it occurs under standard-state conditions.

 ΔG° , the standard free energy change for a reaction determines *the relative concentrations of reactants and products that will be found at thermodynamic equilibrium*.



 $\Delta G^0 < 0 \qquad \Delta G^0 = -RT \ln K \qquad \Delta G^0 > 0$



K and ΔG^0 reveal nothing about the rates of process nor information about the height of the barrier between these states.

The rate of a chemical reaction is dependent on the free energy of activation (transition state).

$$rate = A \cdot e^{-E_a/RT}$$

A catalyst (enzyme) lowers the Energy of activation but does not alter reactant or product free energies.





Protein denaturation

$$\Delta G = \Delta H - T \Delta S$$

T is the temperature (in Kelvin) where the reaction is at equilibrium. $\Delta G = 0$

 ΔH^{o} – includes the components of the binding energy and intra- and inter-molecular interactions

 ΔS^{o} – includes effect of molecular rearrangement.



Protein denaturates at $T \ge 60^{\circ}C$ where the T ΔS term carries more weight and offsets the large positive ΔH .