

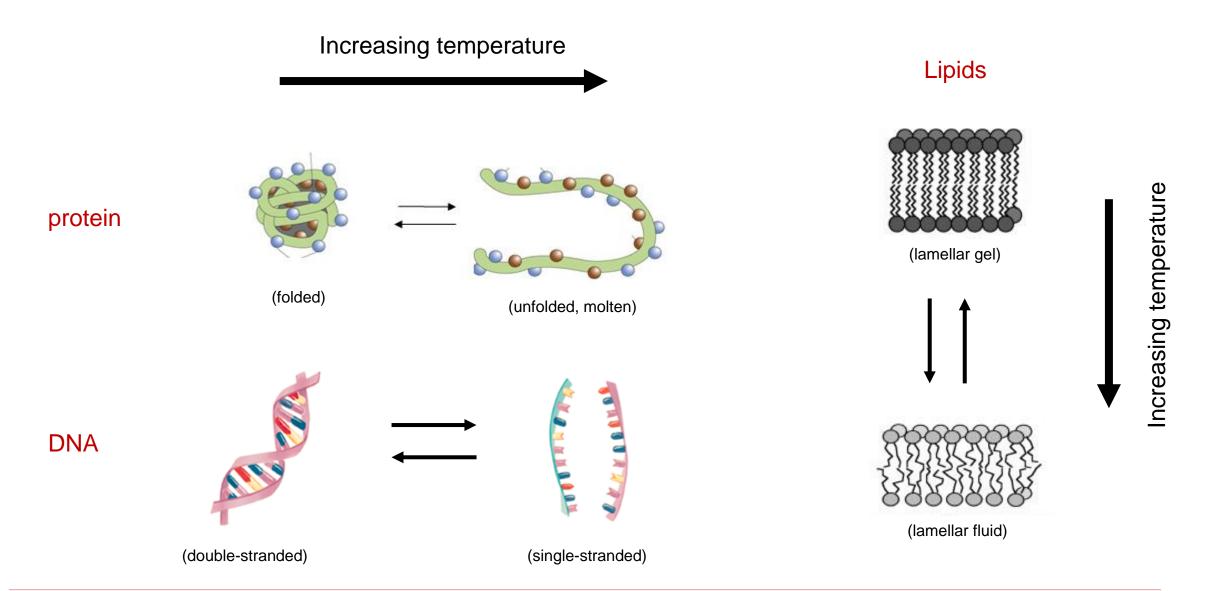


Lecture 6

Soft Matter PhysicsThermal & phase transitions

- Thermal denaturation of proteins
- Cooperativity in thermal melting
- Phase transitions and classifications
- Unmixing of liquid phases

Thermal denaturation of proteins, DNA and lipids



Recap: Chemical potential & Mass action law

 $\mu_0 \rightarrow$ Standard state chemical potential

(at standard conditions: p = 101.3 kPa, T = 298.15 K, $c_0 = 1 \text{ M}$)

For mixture(s):
$$\mu_i = \mu_{i,0} + k_B T \ln \left(\frac{c_i}{c_{i,0}} \right)$$

Relation chemical potential (μ) – Gibbs free energy (G):

$$\mu(T, p, N) = \left(\frac{\partial G}{\partial N}\right)_{T, p}$$

Chemical reaction:

$$-v_A A + -v_B B + ... \rightarrow ... + v_Y Y + v_Z Z$$
(stoichiometry factors v_i)

Free energy change:

$$\Delta_r G = \underbrace{\sum_{i} \nu_i \mu_i}_{A G^0} + k_B T \sum_{i} \ln \left(\frac{c_i}{c_{i0}} \right)^{\nu_i} = \Delta_r G^0 + k_B T \sum_{i} \ln \left(\frac{c_i}{c_{i,0}} \right)^{\nu_i}$$

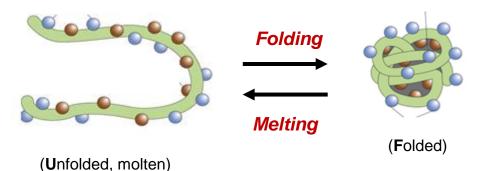
Recap: Mass action law

Chemical reaction:
$$-v_A A + -v_B B + ... \rightarrow ... + v_Y Y + v_Z Z$$
 $(v_A \text{ and } v_B \text{ negative})$

Free energy change:
$$\Delta_r G = \Delta_r G^0 + k_B T \sum_i \ln \left(\frac{c_i}{c_{i,0}} \right)^{v_i} = \Delta_r G^0 + k_B T \ln \left(\prod \left(\frac{c_i}{c_{i,0}} \right)^{v_i} \right)$$

$$\Delta_r G = \Delta_r G^0 + k_B T \ln \left(\underbrace{\prod_{i} c_i^{\nu_i} \prod_{i} c_{i,0}^{-\nu_i}}_{K_{eq}} \right) \longrightarrow \Delta_r G = \Delta_r G^0 + k_B T \ln \left(c_0 K_{eq} \right)$$

Example: protein



$$U \rightleftharpoons F$$

$$[P]_{total} = [U] + [F]$$

Free energy of folding:

$$\Delta_r G_{fold} = \Delta_r G_{fold}^0(p,T) + k_B T \ln \left(\frac{[F]}{[U]}\right)$$
 "substrate"

What happens at melting temperature (T_m) ?

50% fraction Folded + 50% fraction Unfolded

$$[U] = [F] = [P]_{total}/2$$

At melting temperature (T_m) :

$$[U] = [F] = [P]_{total}/2$$

$$\Delta_r G_{fold} = \Delta_r G_{fold}^0(p, T) + \underbrace{k_B T \ln \left(\frac{[F]}{[U]}\right)}_{0}$$

$$\Delta_r G_{fold} = \Delta_r G_{fold}^0 = \Delta_r H_{fold}^0 - T \Delta_r S_{fold}^0$$

During folding reaction:

$$\Delta_r H_{fold}^0 < 0; \qquad \Delta_r S_{fold}^0 < 0$$

(heat release: e.g. H-bond formation) (lower configurational entropy)

What happens when we change T?

T small (folding):

 $\Delta_r G_{fold} < 0 \dots \Delta_r H_{fold}^0$ dominates

T high (melting):

 $\Delta_r G_{fold} > 0 \dots \Delta_r S_{fold}^0$ dominates

$$T = T_m$$
:

$$\Delta_r G_{fold} = 0$$
 ... both states eq. populated

When
$$T = T_m$$
:

When
$$T = T_m$$
: $\Delta_r G_{fold} = \Delta_r H_{fold}^0 - T_m \Delta_r S_{fold}^0 = 0$

("Equilibrium")

$$T_m = rac{\Delta_r H_{fold}^0}{\Delta_r S_{fold}^0}$$
 "competition" between $\Delta_r H$ and $\Delta_r S$

Use mass action law to study changes in state fraction as function of T

$$K_{eq} = K_{fold} = e^{-\Delta_r G_{fold}^0 / k_B T} = \frac{[F]}{[U]} = \frac{[P] - [U]}{[U]}$$

$$[P] - [U] = [U] K_{fold} \Leftrightarrow [P] = [U] K_{fold} + [U]$$

$$[P] = [U](K_{fold} + 1)$$

Fraction of unfolded protein

$$\rho_{unfold} = \frac{[U]}{[P]} = \frac{1}{K_{fold} + 1}$$

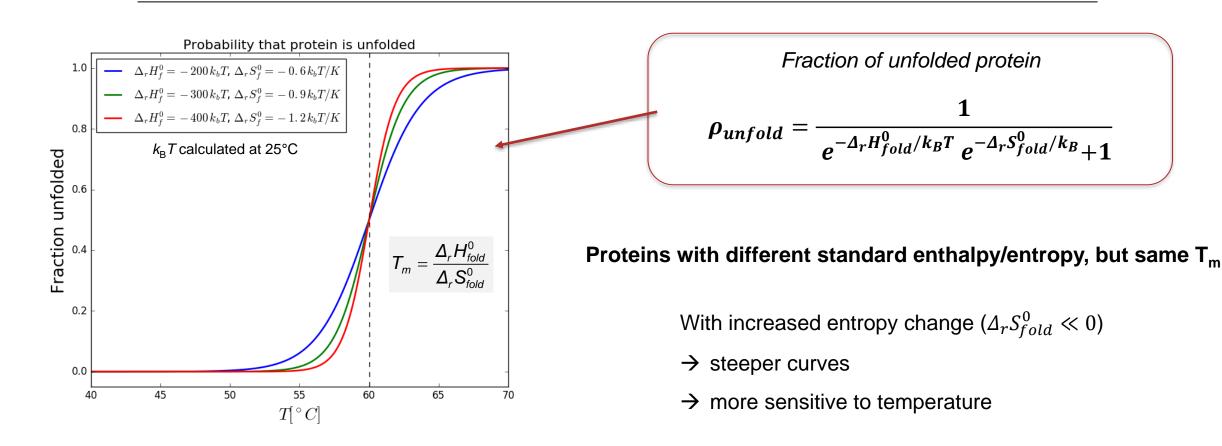
$$\rho_{unfold} = \frac{1}{e^{-\Delta_r G_{fold}^0 / k_B T} + 1}$$

$$\rho_{unfold} = \frac{1}{e^{-\Delta_r H_{fold}^0 / k_B T} e^{-\Delta_r S_{fold}^0 / k_B} + 1}$$

When
$$T = T_m$$
:

When
$$T = T_m$$
: $\Delta_r G_{fold} = \Delta_r H_{fold}^0 - T_m \Delta_r S_{fold}^0 = 0$

$$T_m = rac{\Delta_r H_{fold}^0}{\Delta_r S_{fold}^0} \qquad \qquad \qquad \begin{tabular}{ll} "competition" \\ between \\ \Delta_r H \ and \ \Delta_r S \ \end{tabular}$$



Heat capacity during denaturation of a 2-state system

Enthalpy changes $\Delta_r H_{fold}^0$ can be directly measured using calorimetry \rightarrow Measure of heat taken up in the reaction

$$H_{total} = \rho_{fold}H_{fold} + \rho_{unfold}H_{unfold} = \rho_{fold}H_{fold} + \rho_{unfold}(H_{fold} - \Delta_r H_{fold}^0)$$

$$H_{total} = \rho_{fold}H_{fold} + \rho_{unfold}H_{fold} - \rho_{unfold}\Delta_r H_{fold}^0 = H_{fold} - \rho_{unfold}\Delta_r H_{fold}^0$$

 $H_{fold}(T) = constant$

Heat capacity at constant pressure

$$c_p = \left(\frac{\partial H_{tot}}{\partial T}\right)_p$$

$$c_p(T) = -\Delta_r H_{fold}^0 \left(\frac{\partial \rho_{unfold}}{\partial T}\right)_p = \frac{\left(\Delta_r H_{fold}^0\right)^2}{k_B T^2} \frac{e^{-\Delta_r H_{fold}^0/k_B T} e^{\Delta_r S_{fold}^0/k_B}}{\left(e^{-\Delta_r H_{fold}^0/k_B T} e^{\Delta_r S_{fold}^0/k_B} + 1\right)^2}$$

Heat capacity during denaturation of a 2-state system

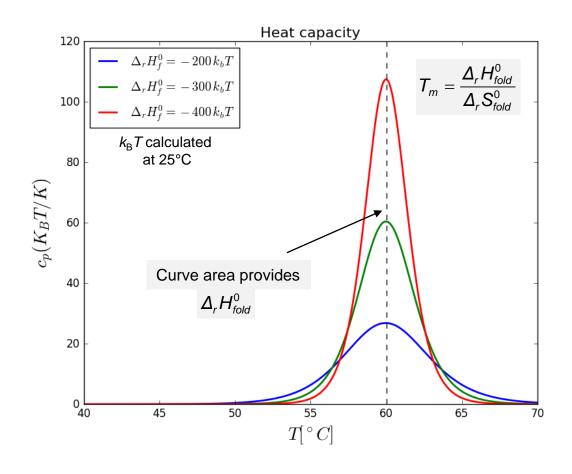
Heat capacity as function of *T*

$$c_p(T) = -\Delta_r H_{fold}^0 \left(\frac{\partial \rho_{unfold}}{\partial T} \right)_p$$

$$c_{p}(T) = \frac{\left(\Delta_{r} H_{fold}^{0}\right)^{2}}{k_{B} T^{2}} \frac{e^{-\Delta_{r} H_{fold}^{0}/k_{B} T} e^{\Delta_{r} S_{fold}^{0}/k_{B}}}{\left(e^{-\Delta_{r} H_{fold}^{0}/k_{B} T} e^{\Delta_{r} S_{fold}^{0}/k_{B}} + 1\right)^{2}}$$

Characteristic bell-like curve near T_m

- \rightarrow T range with strongest change in ρ_{unfold}
- \rightarrow Curve width ... indicates sharpness of transition (extract $\Delta_r H_{fold}^0$ and $\Delta_r S_{fold}^0$)
- $\rightarrow \int \rho_{unfold} dT = 1$, curve area provides $\Delta_r H_{fold}^0$



Proteins with different standard enthalpy/entropy, but same T_m

Differential scanning calorimetry (DSC)



Differential scanning calorimetry (DSC)

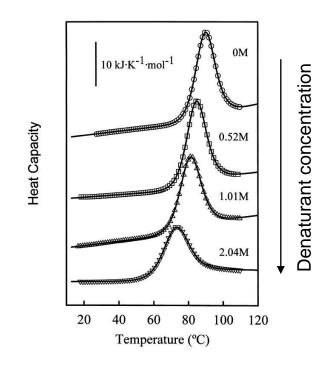
Calorimeter scans heat flux for given ΔT with respect to reference volume

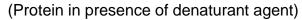
Heat capacity as function of
$$T$$

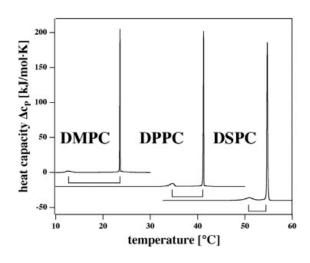
$$c_p = \left(\frac{\partial H_{tot}}{\partial T}\right)$$

DSC is widely applied folding/melting and phase transitions

(e.g., biomolecules, polymers, liquid crystals, ...)

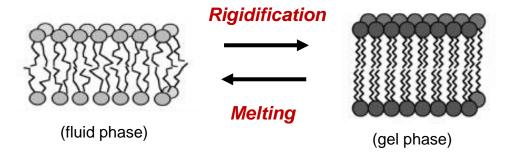


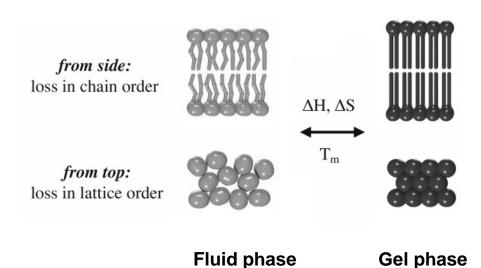




(Lipid bilayer membranes undergoing phase transition)

Cooperativity during thermal melting





disordered lattice

Lipid bilayer membrane

$$Fluid \rightleftharpoons Gel$$

 $[L]_{total} = [F] + [G]$

Heat capacity as function of T

$$\rho_{melted} = \frac{[fluid\ lipid]}{[total\ lipid]} = \frac{1}{K_{eq}+1}$$

$$c_p(T) = -\Delta_r H_{rig}^0 \left(\frac{\partial \rho_{melted}}{\partial T} \right)_p$$

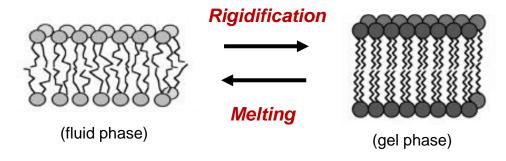
$$c_p(T) = \frac{\left(\Delta_r H_{rig}^0\right)^2}{k_B T^2} \frac{K_{eq}}{\left(K_{eq} + 1\right)^2}$$

These terms don't include cooperativity yet!

ordered lattice

Cooperativity during thermal melting

Cooperativity factor *n* (like Hill's coefficient)





$$\Delta_r H_{rig}^0 \to \mathbf{n} \ \Delta_r H_{rig}^0$$

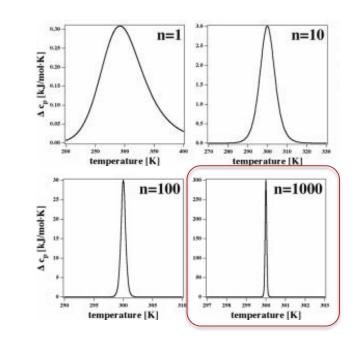
$$\Delta_r S_{rig}^0 \to \mathbf{n} \ \Delta_r S_{rig}^0$$

$$\Delta_r G_{rig}^0 \to \mathbf{n} \ \Delta_r G_{rig}^0$$

Heat capacity as function of *T*

$$c_{p}(T) = \frac{\left(\mathbf{n}\Delta_{r}H_{rig}^{0}\right)^{2}}{k_{B}T^{2}} \frac{e^{-\mathbf{n}\Delta_{r}H_{rig}^{0}/k_{B}T}e^{\mathbf{n}\Delta_{r}S_{rig}^{0}/k_{B}}}{\left(e^{-\mathbf{n}\Delta_{r}H_{rig}^{0}/k_{B}T}e^{\mathbf{n}\Delta_{r}S_{rig}^{0}/k_{B}} + 1\right)^{2}}$$

Melting not independent \rightarrow in cluster of n lipids



Cooperativity:

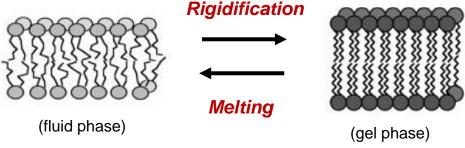
Larger *n*



- → Sharper peak
- → Narrow transition

Cooperativity during thermal melting

Cooperativity factor *n* (like Hill's coefficient)





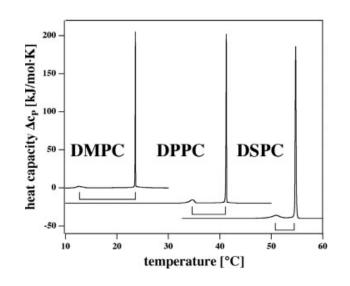
$$\Delta_r H_{rig}^0 \to \mathbf{n} \ \Delta_r H_{rig}^0$$

$$\Delta_r S_{rig}^0 \to \mathbf{n} \ \Delta_r S_{rig}^0$$

$$\Delta_r G_{rig}^0 \to \mathbf{n} \ \Delta_r G_{rig}^0$$

Heat capacity as function of *T*

$$c_{p}(T) = \frac{\left(\mathbf{n}\Delta_{r}H_{rig}^{0}\right)^{2}}{k_{B}T^{2}} \frac{e^{-\mathbf{n}\Delta_{r}H_{rig}^{0}/k_{B}T}e^{\mathbf{n}\Delta_{r}S_{rig}^{0}/k_{B}}}{\left(e^{-\mathbf{n}\Delta_{r}H_{rig}^{0}/k_{B}T}e^{\mathbf{n}\Delta_{r}S_{rig}^{0}/k_{B}} + 1\right)^{2}}$$



Cooperativity:

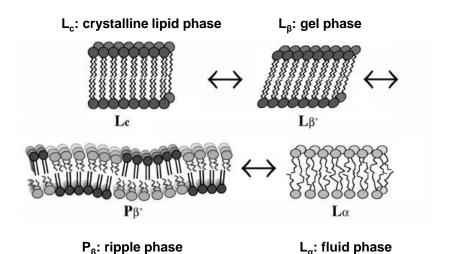
Larger *n*

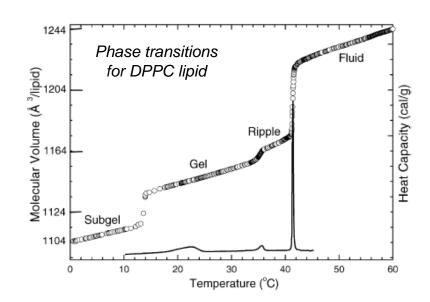


- → Sharper peak
- → Narrow transition

Melting not independent \rightarrow in cluster of n lipids

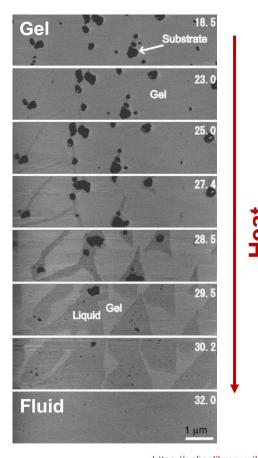
From thermal melting to phase transitions (e.g., in lipids)



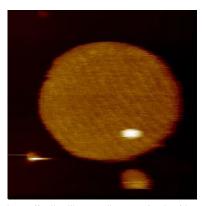


AFM images/movies of DMPC supported lipid bilayers

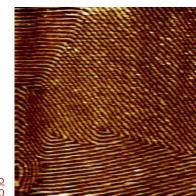
Main transition



Pre-transitionRipple-phase <-> Fluid phase

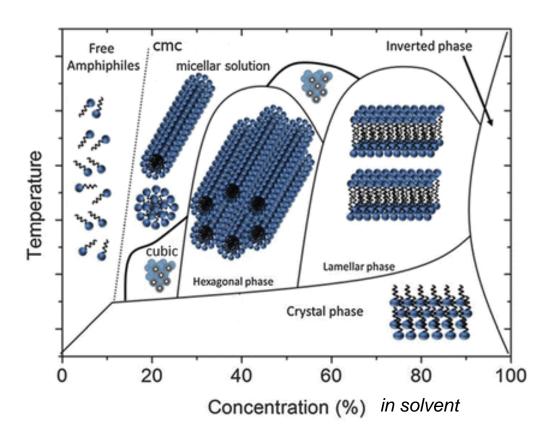


https://onlinelibrary.wiley.com/action/download Supplement?doi=10.1002%2Fsmll.201601549 &file=smll201601549-sup-0002-S2.mov

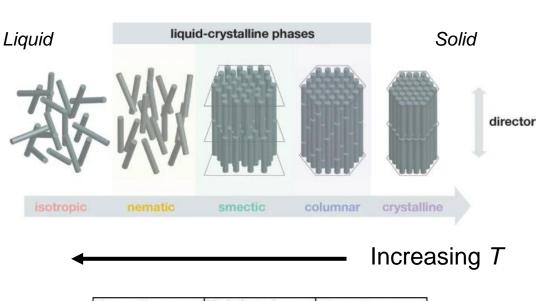


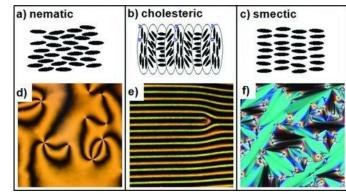
Phase transitions can be complex (e.g., in liquid crystals)

Amphiphiles, such as lipids, are <u>lyotropic</u> liquid crystals (depend on \underline{T} and \underline{C} in solvent)



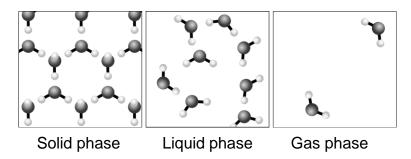
Thermotropic liquid crystals \rightarrow mesophase depends on \underline{T}

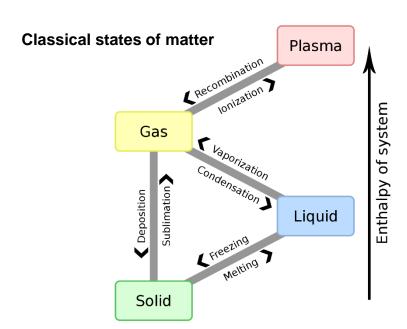




Phase transitions of a substance

Example: Water



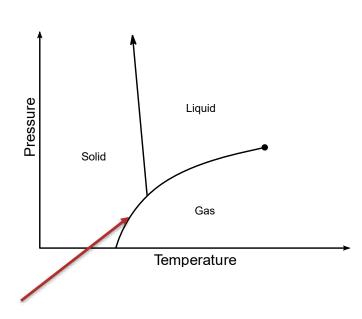


Phase transition of substance: spontaneous/abrupt transition between <u>different states</u> of qualitative uniform properties

Phase transitions occur at specific transition temperatures (e.g. $T_{\rm m}$) at given pressure

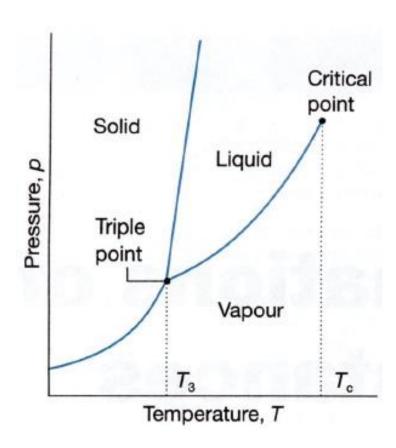
Phase diagrams show *T*, *p* regions at which phases are thermodynamically stable

Phase diagram



Phase boundaries show *T*, *p* regions at which sets of two phases coexist in equilibrium

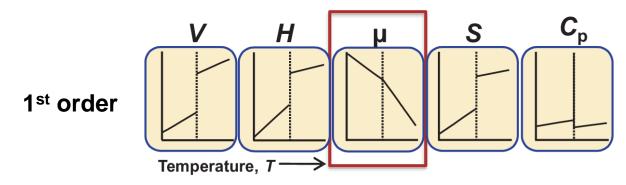
Phase transitions of a substance



Triple point: *T, p* point with *s-l-g* coexistence

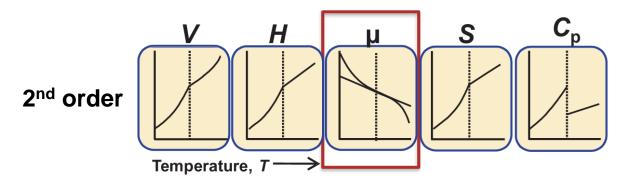
Critical point: T, p so high \rightarrow super-critical fluid

Ehrenfest classification of phase transitions



e.g., melting/freezing, evaporation/condensation

First derivate of μ discontinuous



e.g., (super-)conducting, certain solid-solid transitions

Second derivate of μ discontinuous

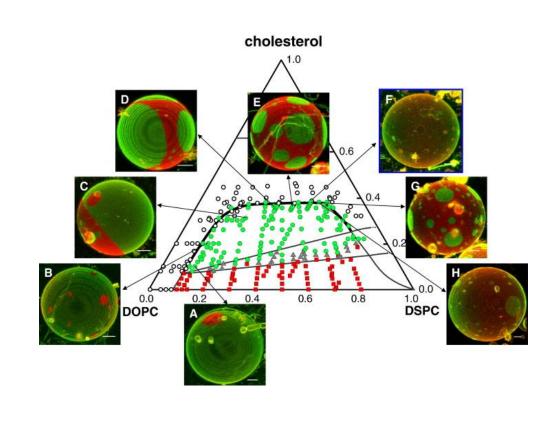
Phase transitions of a mixture of substances

Phase transitions for mixture of substances (e.g., n (im-)miscible compounds)

Phase separation with binary mixtures of lipids

$T_{\rm m}$ = -2 $^{\rm o}$ C (a) fluid Temperature [°C] 30 **POPC** coexistence T_m = 41°C ge -10 20 40 60 80 100 DPPC [%] **DPPC**

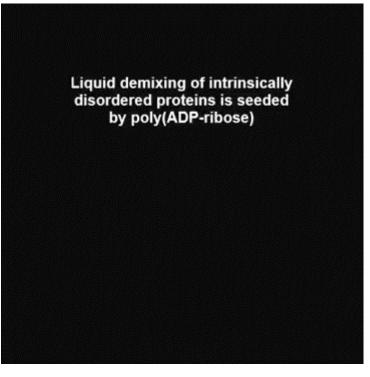
Phase separation with ternary mixtures of lipids



Phase transitions of a mixture of substances

Liquid phase-sep. compartments in cells:





https://www.youtube.com/watch?v=QQFWPRMXzI0



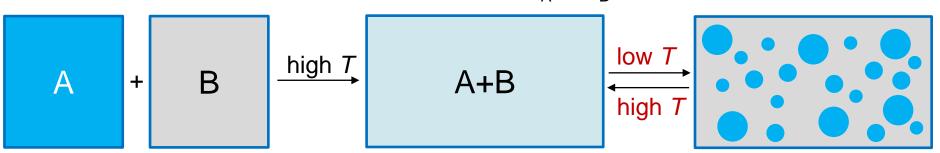
https://www.youtube.com/watch?v=t5jbxh0C0UU

All examples shown here describe the unmixing of two liquid phases (either in 2D or 3D)

Liquid-liquid unmixing

Regular solution model:

Volume fractions: $\Phi_A + \Phi_B = 1$



Two liquids at *T, V* constant



Goal to calculate ΔF_{mix}

$$\Delta F_{mix} = \underbrace{F_{A+B}}_{mix} - \underbrace{(F_A + F_B)}_{unmix}$$

Using Boltzmann formula $S = -k_{\rm B} \sum_i p_i \ln p_i$ (Shannon Entropy), calculate $S_{mix} = -k_{\rm B} (\Phi_A \ln \Phi_A + \Phi_B \ln \Phi_B)$

Assumptions:

- 1) Molecules arranged into lattice with z nearest neighbors
- 2) Interaction energy terms are pairwise additive (ε_{AA} , ε_{BB} , ε_{AB})
- 3) Given site (A & B molecule) has $z\Phi_A$ A neighbors and $z\Phi_B$ B neighbors

Internal energy for mixed vs. unmixed states:

$$U_{mix} = \frac{1}{2} \left(\underbrace{z\Phi_A^2 \varepsilon_{AA} + z\Phi_B \Phi_A \varepsilon_{AB}}_{\text{A on site}} + \underbrace{z\Phi_A \Phi_B \varepsilon_{AB} + z\Phi_B^2 \varepsilon_{BB}}_{\text{B on site}} \right)$$

$$U_{unmix} = \frac{1}{2} \left(\underbrace{z \Phi_A \varepsilon_{AA}}_{\text{site A}} + \underbrace{z \Phi_B \varepsilon_{BB}}_{\text{site B}} \right)$$

Liquid-liquid unmixing

$$\Phi_A + \Phi_B = 1$$

A + B
$$\frac{\text{high } T}{\text{high } T}$$

$$U_{mix} = \frac{1}{2} \left(\underbrace{z\Phi_A^2 \varepsilon_{AA} + z\Phi_B \Phi_A \varepsilon_{AB}}_{\text{A on site}} + \underbrace{z\Phi_A \Phi_B \varepsilon_{AB} + z\Phi_B^2 \varepsilon_{BB}}_{\text{B on site}} \right)$$

$$U_{unmix} = \frac{1}{2} \left(\underbrace{z\Phi_A \varepsilon_{AA}}_{\text{site A}} + \underbrace{z\Phi_B \varepsilon_{BB}}_{\text{site B}} \right)$$

$$\Delta U_{mix} = U_{mix} - U_{unmix} = \frac{z}{2} (\Phi_A^2 \varepsilon_{AA} + \Phi_B^2 \varepsilon_{BB} + 2\Phi_B \Phi_A \varepsilon_{AB} - \Phi_A \varepsilon_{AA} - \Phi_B \varepsilon_{BB})$$

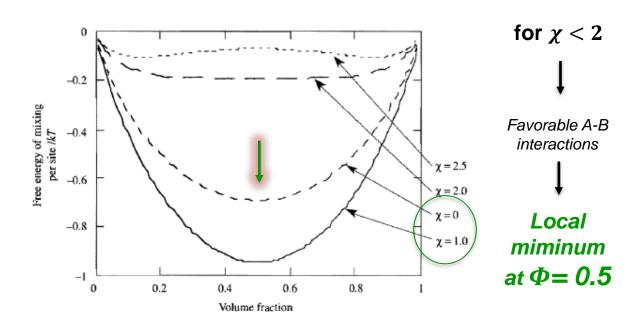
$$\Delta U_{mix} = \frac{Z}{2} \left(\underbrace{(\Phi_A^2 - \Phi_A)}_{-\Phi_A \Phi_B} \varepsilon_{AA} + \underbrace{(\Phi_B^2 - \Phi_B)}_{-\Phi_A \Phi_B} \varepsilon_{BB} + 2\Phi_B \Phi_A \varepsilon_{AB} \right)$$

$$\chi = \frac{\Delta U_{mix}}{k_B T \Phi_A \Phi_B} = \frac{z}{2} (2\varepsilon_{AB} - \varepsilon_{AA} - \varepsilon_{AA})$$

Defined dimensionless parameter

$$\frac{\Delta F_{mix}}{k_B T} = \frac{\Delta U_{mix}}{k_B T} - \frac{S_{mix}}{k_B} = \chi \Phi_A \Phi_B + \Phi_A \ln \Phi_A + \Phi_B \ln \Phi_B$$

$\chi(T)$ measure of the strength of A-B interactions

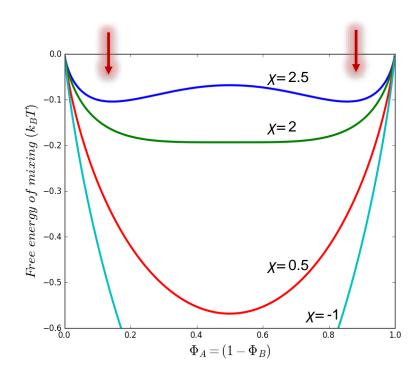


Free energy of mixing two substances

Unfavorable A-B interactions

for $\chi < 2$

Two local mimina at $\Phi \neq 0.5$



These mimima define the **binodal** separating unstable-stable states

Hypothetical generation of two compartments from mixed liquid with ϕ_0 :

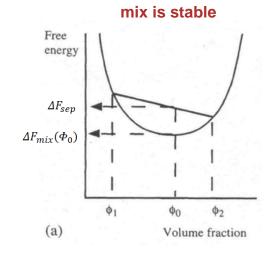
Since $\Phi_A + \Phi_B = 1$, shorten to $\Phi_A = \Phi$ and $\Phi_B = 1 - \Phi$

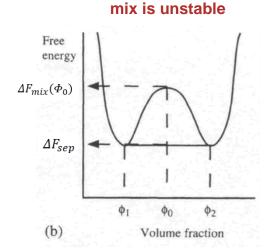
For a desired Φ_0 , we can then mix:

$$\phi_1 \leq \phi_0 \leq \phi_2$$

$$\alpha_1 \Phi_1 + \alpha_2 \Phi_2 = \Phi_0$$

Partial separated system





Free energy change of partially separated system:

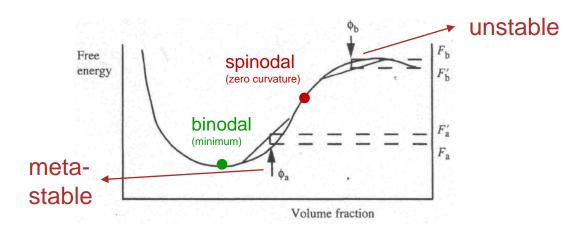
$$\Delta F_{sep} = \Delta F_{mix}(\Phi_1) + \Delta F_{mix}(\Phi_2)$$

Unstable if

$$\Delta F_{sep} < \Delta F_{mix}(\Phi_0)$$

Stability during unmixing

Stable vs. metastable vs. unstable

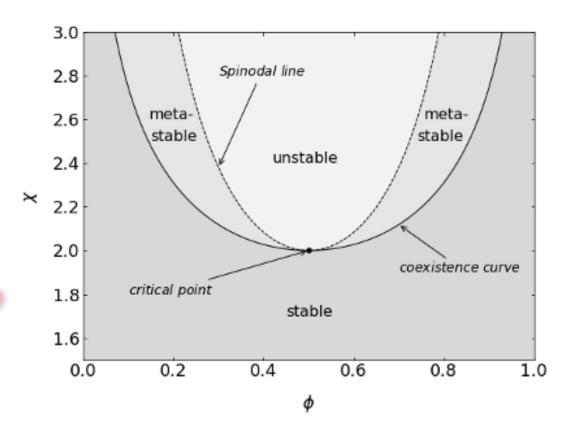


Sign of the curvature of the free energy curve:

$$rac{\mathrm{d}^2 F_{
m mix}}{\mathrm{d}\phi^2} < 0 \ldots ext{unstable} \ rac{\mathrm{d}^2 F_{
m mix}}{\mathrm{d}\phi^2} = 0 \ldots ext{spinodal line} \ rac{\mathrm{d}^3 F_{
m mix}}{\mathrm{d}\phi^3} = 0 \ldots ext{critical point} \ ...$$

 $> 0 \dots stable$

Phase diagram



 χ is proportional to 1/T

Additional literature for material covered in lectures 5 & 6

