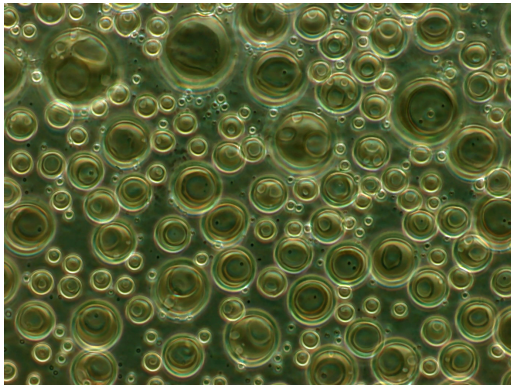
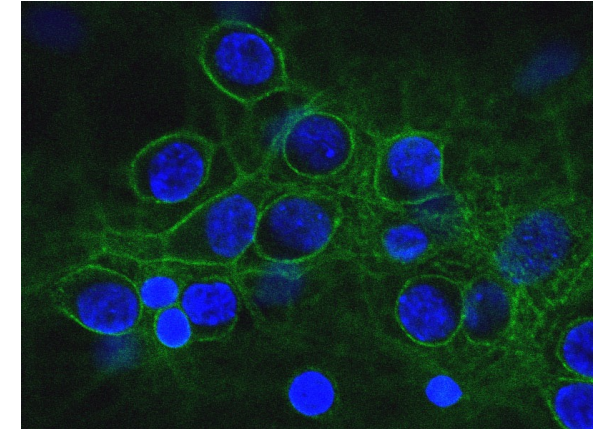
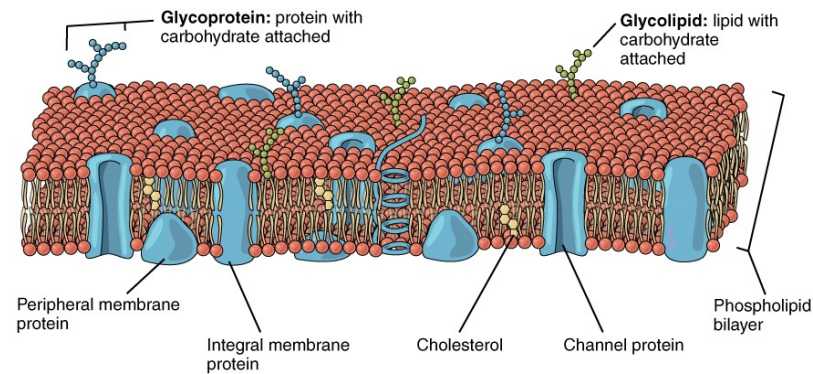
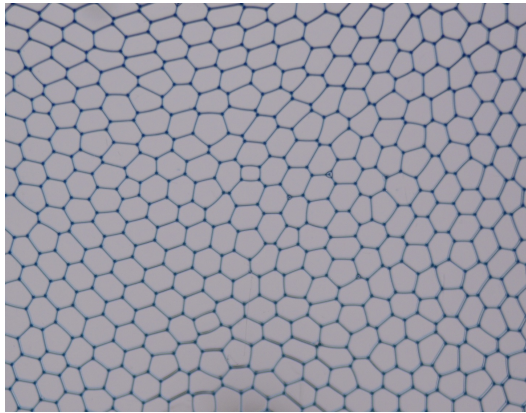


ME470: Mechanics of Soft and Biological Matter

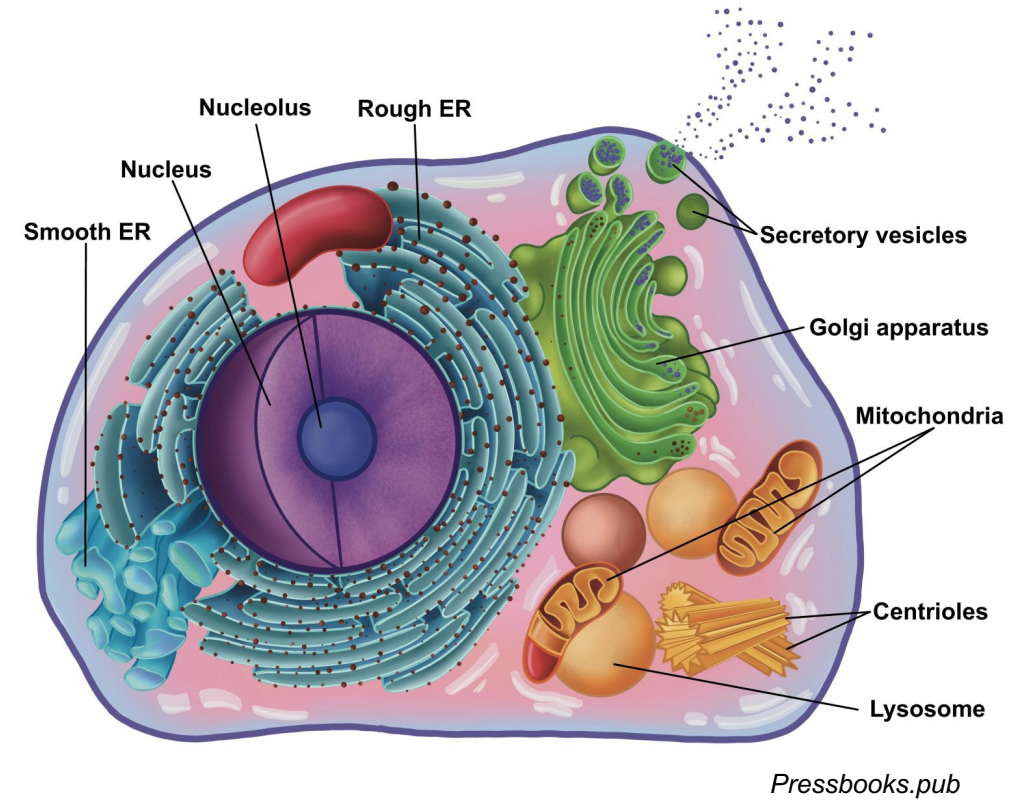
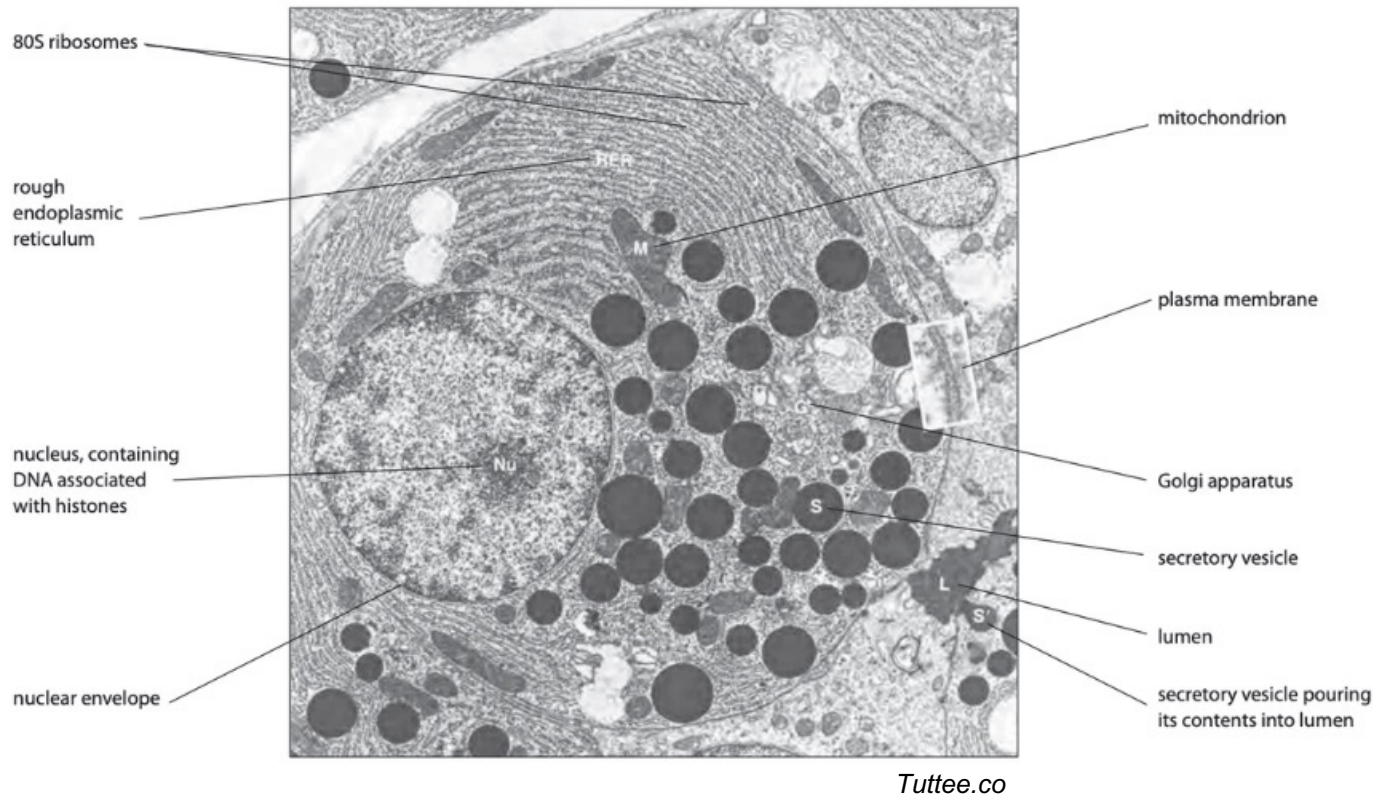
Lecture8: Membrane and Interfaces



Sangwoo Kim

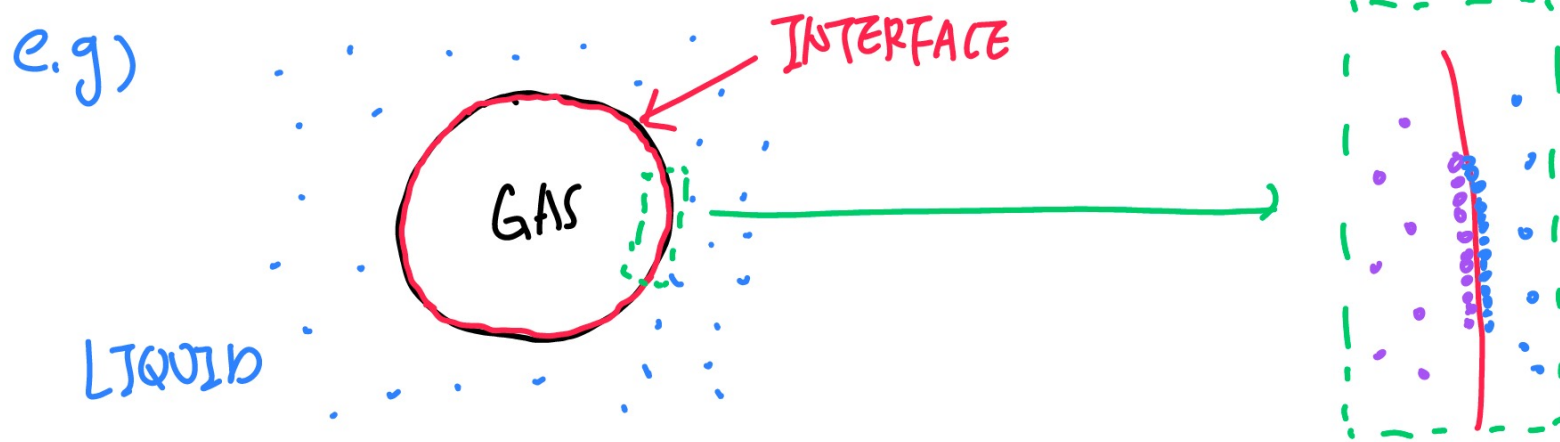
MESOBIO – IGM – STI – EPFL

Red Blood Cells



Membrane distinguishes inside and outside

➡ Mechanics of membrane!!



Energetically unfavorable
to be neighbors to unlike
molecules

⇒ Energy penalty $\propto A$

⇒ Change in free energy

$$dF = -SdT + \int \sigma_{ij} d\varepsilon_{ij} dV + \gamma dA$$

$-PdV$

Energy per area: interfacial tension

For isotropic stress (bubble)

→ Consider a system of bubble + liquid

$$dV_{gas} = -dV_{liq} = dV \quad \text{at constant } T$$

→ $dF = -P_g dV + P_l dV + \gamma dA = 0$



$$\Delta P = P_g - P_l = \gamma \frac{dA}{dV}$$

Young-Laplace law

For spherical bubble:

$$\Delta P = \frac{2\gamma}{R}$$

Note: looks like free enthalpy, $G = F_0 - \tau A$, where $\tau = \gamma$

Note: soap bubble in air, $\Delta P = 4\gamma/R$ (2 interfaces)

Note: such interface is NOT elastic.

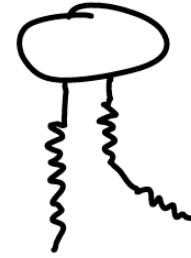
Detergent (e.g. SDS)



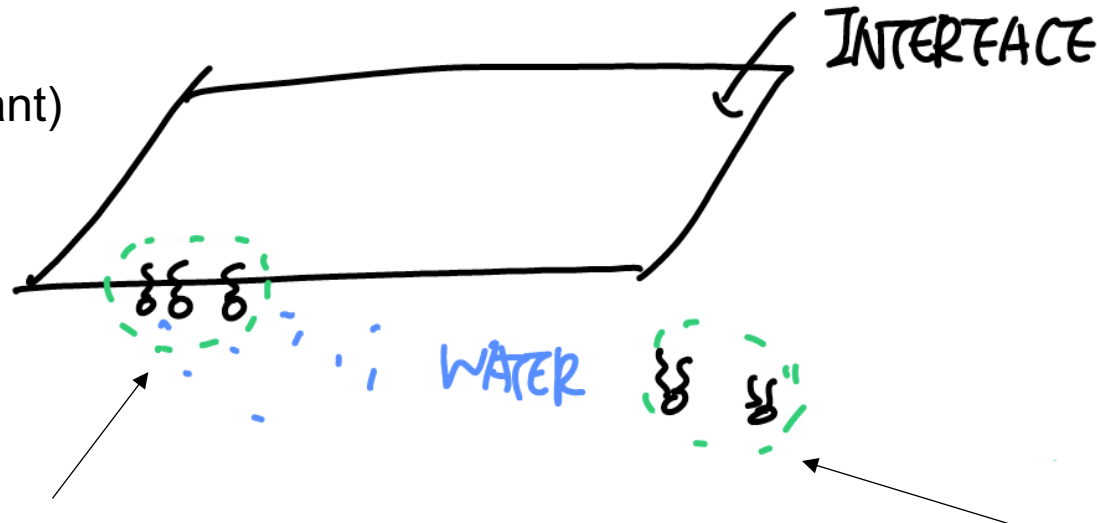
Hydrophilic head

Hydrophobic tail

Phospholipids (e.g. DOPC)

 $2 \sim 2.5 \text{ nm}$

(T=Constant)



Micelle

At interface state (2):
Lower energy per molecule

In bulk state (1):
Higher entropy

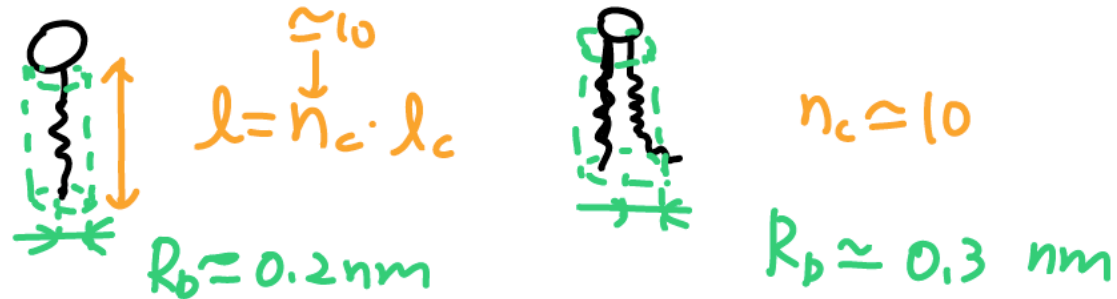
➡ Minimize free energy: $\Delta F = F_1 - F_2 = \Delta U - T\Delta S$

Estimate ΔU : classical approximation

Using surface tension of the hydrocarbon chain w.r.t water

$$\gamma_D \approx \gamma_P \approx 0.05 \text{ N/m}$$

Approximate geometry:
 $l_c = 0.126 \text{ nm}$



$$\Delta U \approx 2\pi\gamma R n_c l_c$$



$$\Delta U_D \approx 20 k_B T$$

$$\Delta U_P \approx 30 k_B T$$

➡ Entropy: $\Delta S = S_1 - S_2 \approx S_1$

Model as ideal gas (non-interacting molecules):

$$S_1 = k_B \left[\frac{5}{2} - \ln \left(\frac{nh^3}{(2\pi mk_B T)^{3/2}} \right) \right] \quad \text{Per molecules}$$

n = volume density

m = molecular mass

$h = 6.626 \times 10^{-34} \text{ J} \cdot \text{s}$ = Planck constant

Masses: $m_D \approx 400 \text{ Da}$, $m_P \approx 570 \text{ Da}$ ($1 \text{ Da} \approx 1.66 \times 10^{-27} \text{ kg}$)

ΔF can be either positive or negative depending on n

➡ Critical concentration n_{agg} , where interface gets preferentially populated!

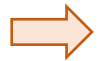
$$\Delta F = 0$$



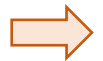
$$n_{agg} = \frac{(2\pi m k_B T)^{3/2}}{h^3} e^{5/2 - \Delta U/k_B T}$$

$$n_{agg}^{(D)} \approx 0.3M \quad (\text{large concentration})$$

$$n_{agg}^{(P)} \approx 2 \times 10^{-5}M \quad (\text{very small concentration})$$



Phospholipids aggregate rarely leave interface (insoluble surfactants) while detergents are very soluble

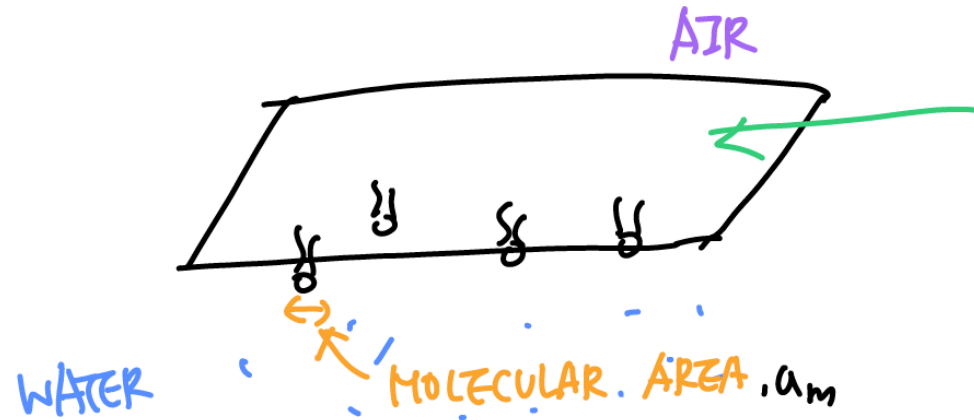


Tightly packed phospholipids at the surface : **MEMBRANE!**

Note: $n_{agg} = n_{micelle}$ (same energetic argument)

Note: some surfactants make **inverted** micelles in oil if the geometry is right





Area per molecule @ interface: a

$(a - a_m)$: surface exposed to air molecule

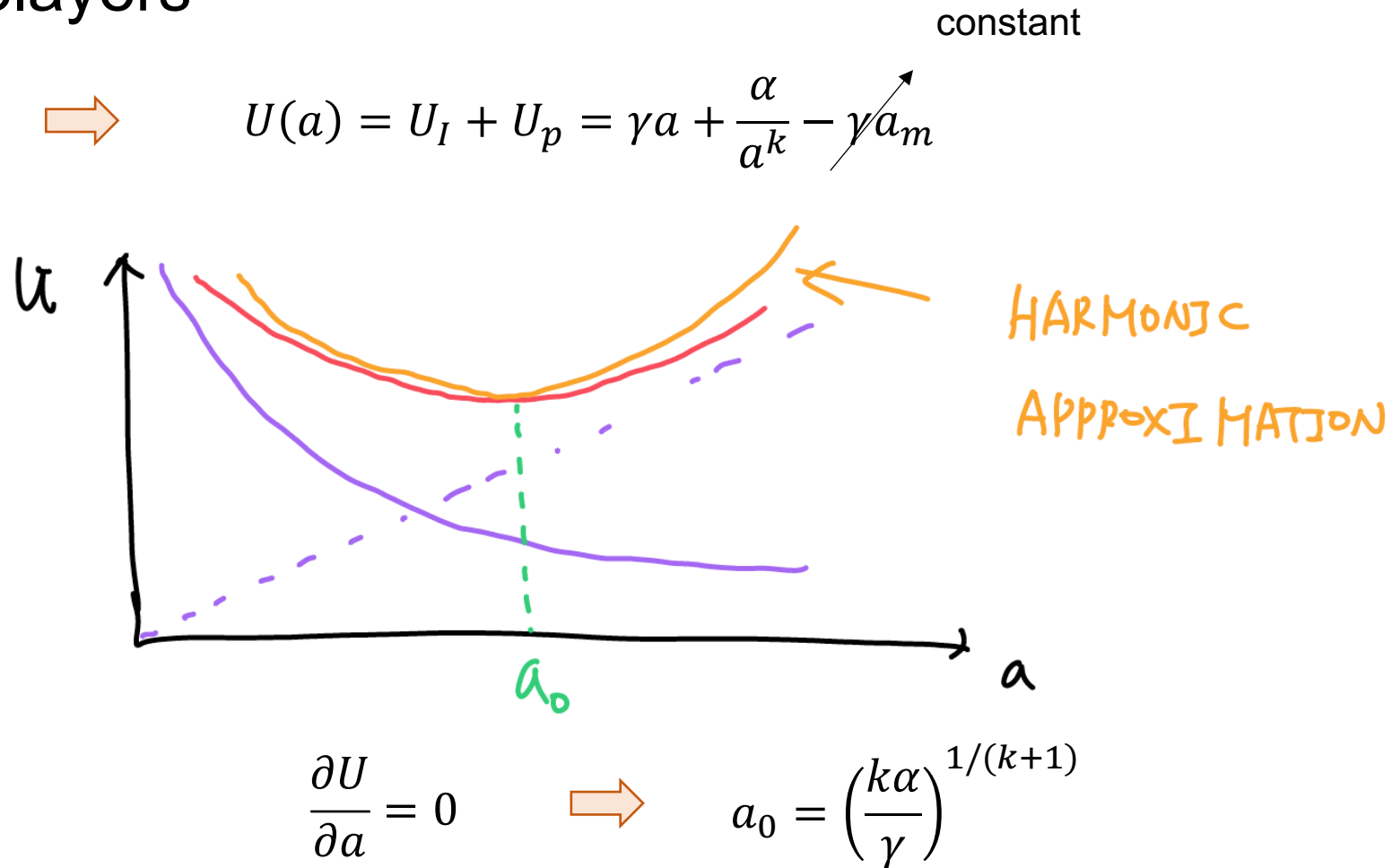
Surface energy cost: $U_I = \gamma(a - a_m)$

Repulsive interaction between molecules:

$$U_p = \alpha \frac{1}{a^k} \quad \alpha, k: \text{constant}$$

Theory: $k = 1$: Coulomb

$k = 2$: Steric repulsion



Taylor expansion (exercise):

$$U - U(a_0) = \frac{k+1}{2} \gamma \cdot \frac{(a - a_0)^2}{a_0} + H.O.T$$

Compare to 2D continuum material:

$$u_A = \frac{U - U(a_0)}{\partial a} = \frac{k+1}{2} \gamma \cdot \left(\frac{a - a_0}{a_0} \right)^2 = \frac{1}{2} K_A \varepsilon_{ll}^2$$



$$K_A = (k+1)\gamma$$

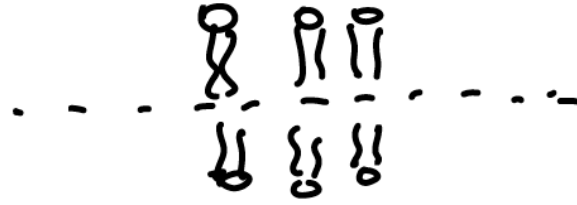
Langmuir trough measurements: often presented as effective surface pressure

$$\Pi_{eff} = -\frac{\partial U}{\partial a} = \frac{k\alpha}{a^{k+1}} - \gamma$$

Q: what will happen if we force PL molecules in bulk water?



MICELLE



LIPID BILAYER



LIPOSOME, VESICLES.

PL bilayers



Cell membrane

Goal: describe bilayer mechanics as 2D continuum

1. Extension/compression
2. Shear
3. Bending (not independent)