Physical Principles in Biology Biology 3550 Spring 2024

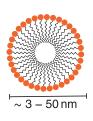
Lecture 32

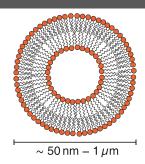
Membrane Permeability and Introduction to Protein Folding

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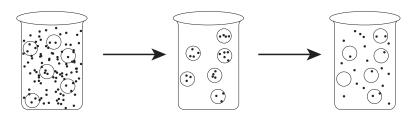
#### Micelles and Vesicles





- Micelles are formed by detergents and soaps; vesicles are formed by phospholipids.
- Micelles are made up of a single shell of detergent or soap molecules; vesicles are made up of lipid bilayers.
- Micelles are generally much smaller than vesicles.
- Different shapes and sizes of the micelles and vesicles reflect the different shapes of detergents and soaps (~conical) and phospholipids (~cylindrical).

## Using Vesicles to Measure Permeability of Bilayers



- Form vesicles in presence of molecules of interest.
- Separate vesicles from external molecules.
- Allow molecules to diffuse across bilayers for a period of time.
- Separate vesicles from external molecules and measure concentrations.
- What determines rate of escape? Fick's first law!

# Diffusion Across Vesicle Bilayer

Fick's first law:  $J = -D \frac{dC}{dx}$ 

Concentration gradient: 
$$\frac{dC}{dx} \approx \frac{C_{\text{in}} - C_{\text{out}}}{\text{Bilayer thickness}}$$

A parameter commonly used in this and other contexts: permeability coefficient, P (not to be confused with pressure!).

Represents combination of diffusion coefficient and membrane thickness:

$$P = \frac{D}{\Delta x}$$

Units:

$$m^2/s \div m = m/s$$

#### Fick's First Law Expressed Using the Permeation Coefficient

$$J = -D\frac{dC}{dx} \approx -D\frac{\Delta C}{\Delta x} = -P\Delta C$$

■ In practice: Measure flux and calculate P

$$P = -\frac{J}{\Delta C}$$

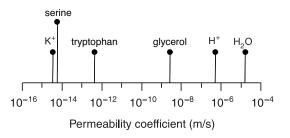
- What do we need to know?
  - Concentration of molecules inside vesicle and out.
  - Rate of molecules diffusing (moles/time).
  - Surface area of vesicles (J has units of mol  $\cdot$  m<sup>-2</sup>s<sup>-1</sup>).
  - Don't need to know membrane thickness!
- *P* reflects both molecule and bilayer (or other kind of membrane)

#### Clicker Question #1

What kind of ions or molecules would you expect to have the **smallest** permeability coefficients for phospholipid bilayers?

- A) Sugars
- B) Amino acids
- C) Water
- D) Nucleotides
- E) Small ions like Na<sup>+</sup>, K<sup>+</sup> or Cl<sup>-</sup>
  All answers count for now!

# Measured Permeability Coefficients



- Range of permeabilities is extremely wide: 9 orders of magnitude.
- Charged ions have very low permeability.
- Polar small molecules have low to medium permeabilities.
- Permeabilities of water and H<sup>+</sup> are actually quite high!

# Comparing Permeability Across Bilayers with Diffusion Coefficients in Water

$$P = D/\Delta x$$

■ Can calculate an "effective diffusion coefficient" by assuming a value for  $\Delta x$ .

$$D = P\Delta x$$

Assume  $\Delta x = 4 \text{ nm}$ 

For ions (other than H<sup>+</sup>):

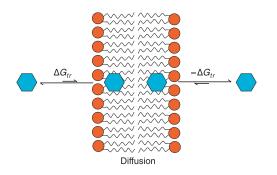
$$D = P\Delta x = 10^{-14} \text{ m/s} \times 4 \times 10^{-9} \text{ m}$$
  
=  $4 \times 10^{-23} \text{ m}^2/\text{s}$ 

For small polar molecules:

$$D = P\Delta x = 10^{-10} \text{ m/s} \times 4 \times 10^{-9} \text{ m}$$
  
=  $4 \times 10^{-19} \text{ m}^2/\text{s}$ 

Compare to  $D = 10^{-10} \,\mathrm{m^2/s}$  for small molecules in water.

### Quantitative Model for Permeability



- Molecule rapidly equilibrates between aqueous and lipid phases.
- Molecule diffuses across lipid phase.
- Diffusion is rate limiting, but overall rate depends on fraction of molecules in the lipid phase:

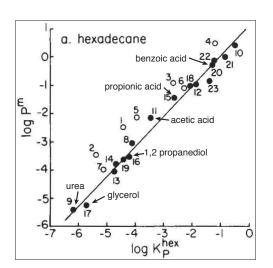
$$P = K_{\rm tr} D_{\rm l}/\Delta x$$

 $K_{tr}$  is equilibrium constant between phases.

 $D_1$  is diffusion coefficient in lipid.

■ Model predicts a correlation between measured  $\Delta G_{\rm tr}$  and P for molecules of similar size.

# Correlation Between Permeability and Solubility in Non-polar Liquids



- Values are for small polar molecules.
- P<sup>m</sup>: Bilayer permeability coefficient (cm/s).
- K<sub>p</sub><sup>hex</sup>: Partition coefficient from water to hexadecane.
- Correlation supports "solubility diffusion" model.

Walter, A. & Gutknecht, J. (1986). Permeability of small nonelectrolytes through lipid bilayer membranes. *J. Membrane Biol.*, 90, 207–217.

http://dx.doi.org/10.1007/BF01870127

#### An Alternative Model

- Bilayers randomly form holes that quickly reseal.
- What would this model predict?
- Some sort of transient pore model likely explains high permeabilities of water and H.

### Some Questions about the Origins of Life on Earth

What are some of the fundamental things that living organisms have to do?

- Collect nutrients from environment.
- Convert nutrients into other compounds and useful forms of energy.
- Build complicated macromolecules, including enzymes and genetic material.
- Create compartments bounded by membranes.
- Reproduce themselves.

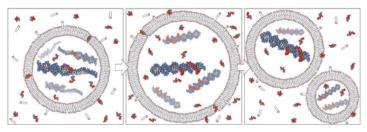
Which came first, proteins or nucleic acids?

- Genetic information is encoded by nucleic acids.
- Proteins are needed to synthesize nucleic acids.
- Now know that some RNA molecules have catalytic activities, suggesting that something like RNA may have come before proteins and DNA.

#### When Did Membranes Enter the Picture?

- Compartmentalization would favor local reactions and prevent competing reactions from stealing reagents.
- Without pores, bilayers prevent escape of molecules, but also their entry!
- Some fatty acids can form bilayers that small polar and charged molecules can cross, but that larger molecules can't.

#### A Model for Primitive Proto-cells



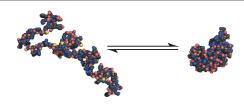
- Primitive reactions lead to formation of polymers, possibly RNA-like.
- Polymers become trapped in semi-permeable vesicles.
- Precursors to polymers diffuse into vesicles and add to the polymers, effectively trapping them.
- Polymers grow, forcing vesicles to grow, and eventually divide.

# Warning!



**Direction Change** 

# Another Thermodynamic Process in Biology: Protein Folding and Unfolding



- Unfolded proteins:
  - Broad ensembles of rapidly interconverting conformations.
- Folded (native) proteins:
  - Compact, well-defined conformations.
  - Usually the functional state.
  - Motions are restricted, but can be essential for function.
- Three-dimensional structure forms after (or during) synthesis.
- Many proteins can unfold reversibly, and the process has been extensively studied in vitro.

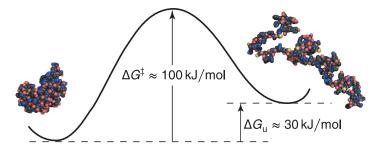
# Protein Unfolding: A Simplified Summary

For small, single-domain proteins ( $\approx$  100 amino acids), unfolding is well described as a "two-state" process:

$$N \rightleftharpoons U$$

Partially folded molecules are rarely detected at equilibrium.

■ Free energy profile for unfolding and refolding:



## **Equilibrium Constant for Unfolding**

■ Calculate  $K_u$  from  $\Delta G_u^{\circ}$  at 298 K

$$\begin{split} &\Delta G_{\rm u}^\circ = -RT \ln K_{\rm u} \\ &K_{\rm u} = e^{-\Delta G_{\rm u}^\circ/(RT)} = e^{-(30\times 10^3\,{\rm J/mol})/(8.314\,{\rm J/(mol\cdot K)}\times 298\,{\rm K})} \\ &K_{\rm u} = \frac{[{\rm U}]_{\rm eq}}{[{\rm N}]_{\rm eq}} \approx 6\times 10^{-6} \end{split}$$

- Only a very small fraction of molecules are unfolded at any instant.
- But, unfolding can be relatively fast, and individual molecules will sample the unfolded state at some point.
- What determines the overall equilibrium between native and unfolded states?
- What determines which three-dimensional structure a particular sequence will form?