Membranes

FIGURE 5-4 The faces of cellular

membranes. The plasma membrane, a single bilayer membrane, encloses the cell. In this highly schematic representation, internal cytosol (green stipple) and external environment (purple) define the cytosolic (red) and exoplasmic (black) faces of the bilayer. Vesicles and some organelles have a single membrane and their internal aqueous space (purple) is topologically equivalent to the outside of the cell. Three organelles-the nucleus, mitochondrion, and chloroplast (which is not shown)-are enclosed by two membranes separated by a small intermembrane space. The exoplasmic faces of the inner and outer membranes around these organelles border the intermembrane space between them. For simplicity, the hydrophobic membrane interior is not indicated in this diagram.



Cell membranes

Membranes are made of strongly anisotropic moleculesStrongly anisotropic molecules like to self-organizing.•a typical eukaryotic cell membrane contains 500–2000different lipid species



Cell membranes

What is so unique about membrane organization?



✓ Membranes are DYNAMIC with built-in Anisotropy

However, dynamics does not implicitly implies randomness and disorder! It is a many body problem with LOCAL (nm scale) order and structure

Milestones in membrane research



The Lipid Bilayer Is a Two-dimensional Fluid

Around 1970, researchers first recognized that individual lipid molecules are able to diffuse freely within the plane of a lipid bilayer. The initial demonstration came from studies of synthetic (artificial) lipid bilayers, which can be made in the form of spherical vesicles, called **liposomes** (**Figure 10–9**); or in the form of planar bilayers formed across a hole in a partition between two aqueous compartments or on a solid support.

Various techniques have been used to measure the motion of individual lipid molecules and their components. One can construct a lipid molecule, for example, with a fluorescent dye or a small gold particle attached to its polar head group and follow the diffusion of even individual molecules in a membrane. Alternatively, one can modify a lipid head group to carry a "spin label," such as a nitroxide



Figure 10–7 Packing arrangements of amphiphilic molecules in an aqueous environment. (A) These molecules spontaneously form micelles or bilayers in water, depending on their shape. Cone-shaped amphiphilic molecules (*above*) form micelles, whereas cylinder-shaped amphiphilic molecules such as phospholipids (*below*) form bilayers. (B) A micelle and a lipid bilayer seen in cross section. Note that micelles of amphiphilic molecules are thought to be much more irregular than drawn here (see Figure 10–26C).

Milestones in membrane research



Milestones in membrane research



Robert Huber





Johann Deisenhofer Prize share: 1/3

Hartmut Michel Prize share: 1/3 Prize share: 1/3

- 1985 Hartmut Michel
- Crystal structure of the first membrane protein
- Photosynthetic reaction ٠ center



Roderick MacKinnon Peter Agre

- · Crystal structure of the first ion channel
- KcsA, Aquaporin

The Nobel Prize in **Chemistry 2003**





Roderick MacKinnon Prize share: 1/2

Peter Agre Prize share: 1/2

Early models







Early models

Demonstration of Lateral Diffusion in Membranes



Frye and Edidin (1970) J. Cell Sci. 7: 319-335



Singer and Nicolson (1972) Science 175: 720-731

Lipids are in bilayer form

Lipids act as solvents for proteins and as permeability barrier and are in a fluid state

Proteins are like 'icebergs' in a viscous sea of lipids

Membrane proteins and lipids can freely diffuse laterally, but cannot rotate from one side of the membrane to the other side (flip-flop)

A small proportion of membrane lipids interact with specific membrane proteins and this could be essential for their function

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Singer and Nicolson (1972) Science 175: 720-731

Limitations of Fluid Mosaic Model

In some membranes, flip-flop of lipids is fast (ER, growing *E. coli*)

All membrane proteins are not free to move in the plane of the membrane

Non-bilayer structure of lipids is possible

There is evidence of lateral domains in membranes

Membranes can be crowded

Does not take into account : LOCAL ORDER, DOMAIN FORMATION

Source	Lipid	Protein	Cholesterol
Rat liver			
Plasma	30–50	50-70	20
Rough ER	15-30	60-80	6
Smooth ER	60	40	10
Inner mitochondria	20-25	70-80	<3
Outer mitochondria	30-40	60-70	<5
Nuclear	15-40	60-80	10
Golgi	60	40	8
Lysosomes	20-25	70-80	14
Rat brain			
Myelin	60-70	20-30	22
Synaptosome	50	50	20
Rat erythrocyte	40	60	24
Rat rod outer segment	50	40	<3
Escherichia coli	20-30	70	0
Bacillus subtilis	20-30	70	0
Chloroplast	35-50	50-65	0

are given.

ER, endoplasmic reticulum.

Source: Based on Jain, M. K., and R. C. Wagner, Introduction to Biological Membranes, 2nd ed. New York: Wiley, 1988, p. 34.



Engelman (2005) Nature 438: 578-580

Lateral distribution of molecules is heterogeneous, corresponding to An organization into DOMAINS

Current Model of Biological Membranes: Organization of Membranes into Domains



Current Model of Biological Membranes: Organization of Membranes into Domains



Mouritsen and Andersen (1998) *Biol. Skr. Dan. Vid. Selsk.* 49: 7-12 Life - As a Matter of Fat: Lipids in a Membrane Biophysics Perspective, Ole G. Mouritsen and Luis A. Bagatolli, 2nd Edn., 2016, Springer

Forces that hold membrane

The Hydrophobic Effect describes how an aqueous medium deals with non-polar substances

It forms the basis for the formation of a variety of organized molecular assemblies such as membranes, micelles, and folded proteins

It should not be confused with the force of interaction among two non-polar (hydrophobic) molecules which plays a very minor role in hydrophobic effect. The effect actually arises primarily from the strong attractive forces between water molecules and the entropic cost of incorporating a non-polar molecule among water molecules.

Tanford (1980) The Hydrophobic Effect John Wiley, New York

Hydrophobic forces

Hydrophobic forces are very relevant in biology. They are primarily driven by an energy cost of creating hydrocarbon-water contact. There is a reduction of entropy of water close of a hydrophobic surface: water becomes structured, even ice-like. It restricts the possible orientations close to the surface and decrease entropy.



Fig. 2.7 Water molecules adjacent to a hydrophobic molecule suffer restrictions in orientation as they form hydrogen bonds with other water molecules.

If one pictures a tetrahedral cage of four water molecules hydrogen bonding a central water molecule, the central water can donate its hydrogen atoms in any combination of two of its four neighbors. **This gives six ways to be fully hydrogen bonded**. Replacing one water of the cage by a hydrophobic, nonhydrogen-bonding neighbor reduces the number of ways this can happen by a factor of about two.

Hydrophobic forces

The restriction in orientation of vicinal water varies with temperature. It becomes harder and harder to order molecules as the temperature is raised. As a result, hydrocarbon–water contacts have a very high heat capacity. Raising the temperature gradually melts the ice-like vicinal water. Interestingly, and somewhat paradoxically, as the temperature rises and the entropy goes up, the hydrophobic effect does not get weaker, but instead gets slightly stronger. This is because the dispersion force becomes stronger with increasing temperature, and this compensates for the loss of entropic drive.

We must therefore view the hydrophobic force as entropy-driven at low temperatures (around room temperature) and enthalpy-driven at higher temperatures (near the boiling point of water). This in a limited range of temperatures.

Hydrophobic effect



Hydrophobic effect

Organized molecular assemblies of various types formed due to the Hydrophobic Effect



Phospholipid Supramolecular Assemblies

Bilayer







Multilamellar vesicle



(c)

Lipids

Water insoluble compounds (soluble in organic solvents)

Biological role:

- energy supply
- energy store
- components of cellular and organelle membranes



When in aqueous environment the heads have affinity for the water molecules, while the tails tend to avoid water by sticking together.







lipid bilayer

Fatty acids

Carboxylic acids with long hydrocarbon chains (12-24 -CH₂- units)



Some have one or more double bonds and are called unsaturated. The double bond is rigid and creates a kink in the chain; the rest of the chain is free to rotate

Stearic acid - saturated



hydrophobic tail



Oleic acid - unsaturated

Phospholipids

In phospholipids, two of the OH groups of glycerol are linked to fatty acids, while the third is linked to a phosphate group, which can be further linked to a polar group such as choline, serine, inositol, etc...



3

Sphingolipids



Sphingolipids are derivatives of sphingosine (E), an amino alcohol with a long hydrocarbon chain. Various fatty acyl chains are connected to sphingosine by an amide bond.

The sphingomyelins (SM), which contain a phosphocholine head group, are phospholipids.

Other sphingolipids are glycolipids in which a single sugar residue or branched oligosaccharide is attached to the sphingosine backbone.

Lipids nomenclature

- The nomenclature of fatty acids is rather complicated. There are at least five systems in use
- > The delta system numbers the double bonds from the carboxyl group (the α carbon)
- The omega system indicates where the first double bond is counting from the other end of the molecule (the ω carbon).

Trivial	Systematic	Colon	Delta	Omega
Stearic acid	Octadecanoic acid	18:0	Octadecanoic acid	-
Palmitic acid	Hexadecanoic acid	16:0	Hexadecanoic acid	-
Oleic acid	E-Octadec-9-enoic acid	18:1; n9	cis-∆9-octadecenoic acid	ω-9
Linoleic acid	9E, 12E-Octadeca-9, 12-dienoic acid	18:2; n9	<i>cis</i> , <i>cis</i> - $\Delta^{9, 12}$ -octadecadienoic acid	ω-6
Linolenic acid	6E, 9E, 12E-Octadeca-6, 9, 12-trienoic acid	18:3; n6	cis, cis, cis - $\Delta^{6,9,12}$ - octadecatrienoic acid	ω-3

Saturated vs Unsaturated Fatty Acids



The actual conformation of a molecule influences its size.

Temperature will lead to a rotation around the C-C bonds.

Only lipids with limited degree of disorder will fit into a bilayer structure.

Di-acyl PC lipids



Typical cross-sectional areas of the cylinders that describe average lipid conformation in the lipid bilayers= is about 0.63 nm², with average length from 1.0 to 1.5 nm (depending on number of C atoms, saturation).





Lipid polar head groups

Substituent	Chemical formula	Polar head group name	Ab ^{&}	
hydrogen	-Н	phosphatidic acid	PA	
choline	-CH ₂ CH ₂ N(CH ₃) ₃ ⁺	phosphatidylcholine	PC	
ethanolamine	- CH ₂ CH ₂ NH ₃ *	phosphatidylethanolamine	PE	
serine	- CH ₂ CH(NH ₃)COO	phosphatidylserine	PS	
glycerol	- CH ₂ CH(OH)CH ₂ OH	phosphatidylglycerol	PG	
<i>myo-</i> inositol	HO H HO H HO H HO H HO H	phosphatidylinositol	PI	

*Chemical formula for the substituent linked to the phosphate group at position 3 of the glycerol moiety. ⁸Abbreviation for the polar head group nomenclature.

Sphingosine based phospholipids



Carbon skeleton	Structure®	Systematic name ^c	Common name (derivation)	Melting point (°C)	Solubility at 30°C (mg/g solvent)	
					Water	Benzene
12:0	CH ₃ (CH ₂) ₁₀ COOH	n-Dodecanoic acid	Laurie acid (Latin <i>lauru</i> s, "laurel plant")	44.2	0.063	2600
14:0	CH ₂ (CH ₂) ₁₂ COOH	n-Tetradecanoic acid	Myristic acid (Latin myristica, nutmeg genus)	53.9	0.024	874
16,0	CH ₃ (CH ₂) ₁₄ COOH	n-Hexadecanoic acid	Palmitic acid (Latin palma, "palm tree")	63.1	0.0083	348
18:0	CH ₃ (CH ₂) ₁₆ COOH	n-Octadecanoic acid	Stearic acid (Greek stear, "hard fat")	69.6	0.0034	124
20:0	CH ₂ (CH ₂) ₁₄ COOH	n-Eicosanoic acid	Arachidic acid (Latin Arachis, legume genus)	76.5		
24:0	CH ₃ (CH ₂) ₂₂ COOH	n-Tetracosanoic acid	Lignoceric acid (Latin lignum, "wood" + cera, "wax")	86.0		
16:1 (A9)	CH ₃ (CH ₂) ₅ CH=CH(CH ₂) ₇ COOH	cis-9-Hexadecenoic acid	Palmitoleic acid	0.5		
18:1 (A9)	CH ₂ (CH ₂), CH=CH(CH ₂),COOH	cis-9-Octadecenoic acid	Oleic acid (Latin oleum, "oil")	13.4		
18:2(A9, 12)	CH ₃ (CH ₂)4 CH=CHCH ₂ CH=CH(CH ₂) ₇ COOH	cis-,cis-9,12- Octadecadienoic acid	Linoleic acid (Greek linon, "flax")	-5		
18:3(A9, 12, 15)	CH ₃ CH ₂ CH=CHCH ₂ CH=CHCH ₂ CH=CH(CH ₂) ₇ COOH	cis, cis, cis- 9,12,15- Octadecatrienoic acid	a-Linolenic acid	-11		
20:4(Δ5, 8, 11, 14)	CH ₃ (CH ₂) ₄ CH=CHCH ₂ CH=CHCH ₂ CH=CHCH ₂ CH=CHCH ₂ CH=CH(CH ₂) ₃ COOH	c/s,cis,cis,cis 5,8,11,14 Eicosatetraenoic acid	Arachidonic acid	-49.5		

More than 500 species of fatty acids !

⁶The prefix n indicates the normal unbranched structure. For instance, dodecanoic simply indicates 12 carbon atoms, which could be arranged in a variety of branched forms; n dodecanoic specifies the linear, unbranched form.

Source: Data from Nelson, D. L., and M. M. Cox, Lehninger Principles of Biochemistry, 4th ed. New York: W. H. Freeman, 2005.

Lipidomic survey of a budding yeast





All interfaces are covered with interfacially active molecules



- 1. Positive pressure resulting from headgroup repulsive forces
- 2. Negative pressure at the hydrophobic-hydrophilic interface the interfacial tension
- 3. Positive pressure resulting from entropic repulsion between acyl chains - chain pressure

Cholesterol and steroids

Steroids (such as cholesterol) have a rigid structure made up by 4 rings.





Cholesterol is an important component of the eukaryotic membranes and has a key role in controlling the membrane fluidity.



Effect of cholesterol

Lateral pressure profiles in DPPC/Cholesterol bilayer



Membrane physical properties



Membrane Physical Properties are Determined by its Lipid Composition

Nature (2014) 510: 48-57

Membrane physical properties



P= lipid volume/ (cross sectional are of the polar group x lipid length)

Lipid conformation



Conformation depends on temperature. It affects packing in the lipid bilayer. Indeed the shape itself is affected by the other molecules forming the aggregate.

Lipid shape is important for functioning. It is given by the compatibility between head and tail. We define`a packing parameter P:

$$P = v/al$$

P = 1 is a cilindrical shaped lipid molecules, fitting a lamellar structure with zero curvature.

Curvature although is important for many of the membrane processes

Lipids and membrane curvature



Lipids and membrane curvature



The more non-cylindrical are lipid shapes, the less stable the bilayer will be.

Each layer tend to elastically relax to a state of finite, spontaneous curvature, causing a curvature stress field.

If the bilayer cohesion does nor sustain the curvature stress, non lamellar structures form.

Lipid speak the language of curvature, in the many structures formed!

The inverted hexagonal structure (H_{II}), has long cilindrical rods of lipids, in a water filled tube, whose diameter can be varied with T, degree of hydration, pH (all change a/l ratio).

Lipids and membrane curvature



Cholesterol has an inverted conical shape (small OH, big steroid ring). Tends to promote the H_{II}. Stress field is mitigated by enzymes.

Hn

From research in microorganisms it appeared that curvature is a crucial parameter in regulating lipid synthesis/enzymatic activity of phospolipases-—lipid molecular shape/optimal packing is at the basis of curvature stress. Yet unknown which membrane-bound proteins are involved in curvature stress sensing-lipid synthesis.

NB: vesicles do not close because of curvature stress, but because of boundary conditions! (micron vs. nanometers)

Membrane physical properties



Lipids form soft interfaces

Membranes are soft interfaces. As polymers, exist in a condensed phase, but cannot be classified neither as solid, nor liquid. The physics of such interfaces is dominated by entropy.

Softness means high deformability but not necessarily high bulk compressibility! Soft matter is anisotropic, hierarchical, with structures spanning over different length scales, and is governed by self-assembling.

In liquid, the interfacial tension $\gamma = \left(\frac{\partial G^{S}}{\partial A}\right)_{V}$

with G^s being the Gibbs excess free energy, V, A volume and surface area acts to make the interface as small as possible, at the same time imparts a certain stiffness to the interface.

The introduction of interfacially active molecules (i.e. amphiphiles) lowers the interface tension.

If molecules are enough, the interface can be fully covered. Therefore the area is fixed and I.T. tends to zero.

Lipids form soft interfaces

Natural examples of soft interfaces: soap bubbles



Soap bubbles: two layers form, at the water-air interfaces, the outer and the inner surfactant layer.

Bubbles are stabilized for a particular size, a particular water layer thickness depending on:

-type of surfactant

- -quantity of surfactant
- -quantity of water

Self-organized monolayers (on liquid surfaces)



The term "molecular self-assembly" refers to spontaneous formation of an ordered molecular overlayer on the surface, often proceeding through several consecutive stages where 1D and 2D ordered structures can also exist.

Thermodynamically, molecular self-assembly proceeds toward the state of lower entropy, and must therefore be compensated by the establishment of intermolecular and molecule-surface interactions. 53

Self-organized monolayers (on solid surfaces)

