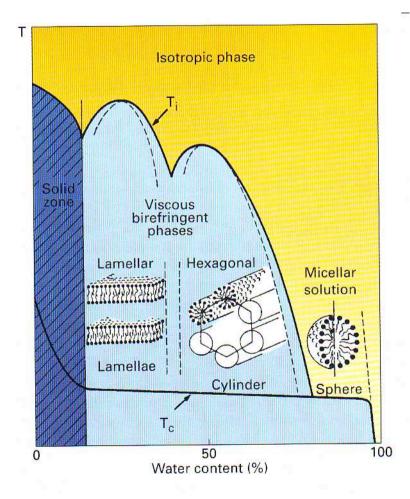
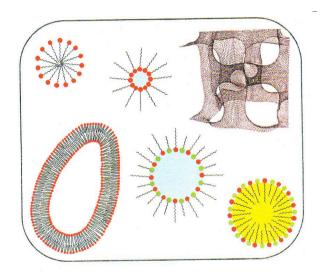
Self assembly & soft matter in nature





Literature

- Jones chp. 9 Self Assembly
- Jones chp. 10 Soft Matter in nature
- C. A. Angell, Formation of Glasses from Liquids and Biopolymers, Science 267, 1924 (31 March 1995)
- Materials Today: March and September issues (2006) Templating materials with self assembly

Self-assembly

Characteristics of self-assembly

Self assembly with surfactants & amphiphiles

- what/types
- energies
- aggregation geometries & phase behaviour

Self assembly with polymers

- phase separation
- structures

4 Characteristics of Soft Matter

- i) Length scales:
 - Structures of ≈10-1000 nm determine the properties
- ii) Time scales: processes from 10⁻¹² 10³ s
 - Dynamics processes over a wide time scales 10⁻¹² 10³ s
 - Very slow processes in non-equilibrium configurations

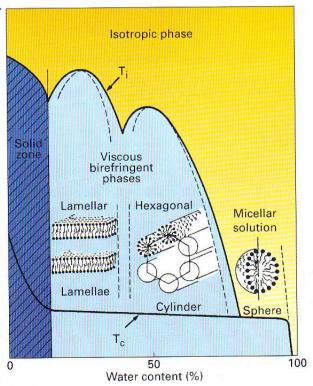
iii) Weak interactions

- Interactions ≈ kT
- iv) Self assembly
 - Hierarchical arrangement of structures
 - Competition between interaction energy and entropy

S

Self-assembly

- i) Hierarchical arrangement of structures
- ii) Competition between interaction energy and entropy
- iii) Structures of ≈10 to >1000 nm formed and determine the properties of the material



Amphiphilic molecule: two parts with very different affinities ⇒ surfactants, bolck co-polymers

Surfactant: an amphiphilic molecule with a tendency to adsorb onto free surfaces and interfaces (Swedish - tensid)

Amphiphilic molecule: two parts with very different affinities

Ex. hydrophilic head and hydrophobic tail



a single molecule is never really comfortable in a solvent
 ⇒ aggregation to minimize the interfacial energy
 ⇒ micro-phase separation, thermodynamically stable aggregates - micelles

Amphiphilic molecule:

two parts with very different affinities

Ex. hydrophilic head and hydrophobic tail

Hydrophobic tail: CH₃(CH₂)_n-

Hydrophilic head: *Anionic (-) Cationic (+) non-ionic block-copolymers*

Small molecules $M_W \approx 100-1000$

Amphiphilic molecule: two parts with very different affinities

Ex. hydrophilic head and hydrophobic tail

Example	Hydrophobic group	Hydrophilic group	Category	Where found
Sodium stearate	C ₁₈ H ₃₇	-COO-Na ⁺	Anionic	Soap
Sodium dodecyl sulphonate (SDS)	C ₁₂ H ₂₅	-OSO3Na+	Anionic	Detergents
Hexadecyl trimethyl- ammonium bromide (CTAB)	C ₁₆ H ₃₃	-N ⁺ (CH ₃) ₃ Br ⁻	Cationic	Mild disinfectants
C ₁₂ E ₅	C ₁₂ H ₂₅	$-(OCH_2CH_2)_5$	Non-ionic	Cosmetics
Pluronic P105	(OCH ₂ C ₂ H ₅) ₅₈	Two chains, each -(OCH ₂ CH ₂) ₃₇	Non-ionic, triblock copolymer	Cosmetics, pharmaceu- ticals
Lecithin	Two chains, each C ₁₂ H ₂₅	Phosphatadyl choline	Zwitterionic, phospholipid	Animal cell membranes, food

Amphiphilic molecule: two parts with very different affinities

Ex. hydrophilic head and hydrophobic tail

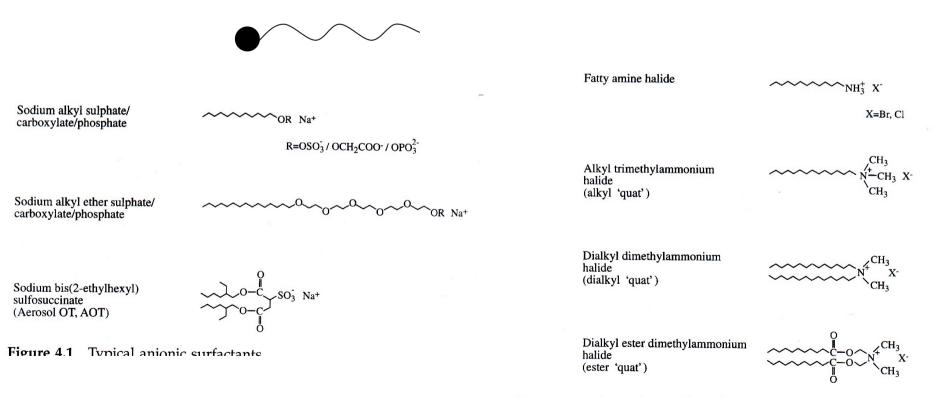
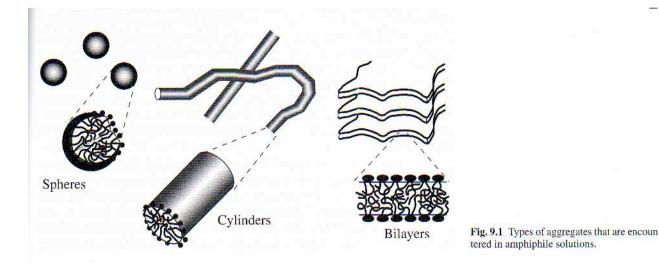


Figure 4.2 Typical cationic surfactants

Ex. hydrophilic head and hydrophobic tail



Self-assembly of molecules in order to minimise free energy ⇒ minimising the contact between water (solvent) and hydrophobic tail while keeping the hydrophilic head in contact with water



Macro vs. micro phase separation

Simple liquids (water & oil):

 $\varepsilon_N = \varepsilon_{\infty} + \frac{\alpha k_B T}{N^{1/3}}$ free energy when a molecule joins an aggregate $\alpha k_B T = 4\pi \gamma (3\nu/4\pi)^{2/3}$ free energy when a molecule joins an aggregate N - number of molecules in the aggregate γ - interfacial energy, ν - molecular volume

 ε_N - monotonic decreasing function of N \Rightarrow infinite aggregates - macroscopic phase separation

Amphiphiles:

 \Rightarrow finite size of aggregates - microscopic phase separation

Why?

Macro vs. micro phase separation

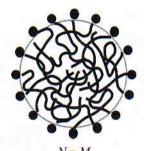
Amphiphiles:

 \Rightarrow micro-phase separation

 \Rightarrow finite size of aggregates entropic considerations more important

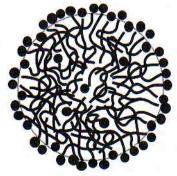






<u>Why?</u> Balance between protecting the hydrophobic tail and keeping the head in contact with the water

associating tails vs. repulsion of head groups



N > M

Fig. 9.3 The optimum aggregation number, M for a spherical micelle. Micelles containing a smaller number of molecules have too large an area for each head-group, causing unfavourable water/hydrocarbon interactions, while in micelles larger than the optimum number some head-groups are forced inside the hydrophobic core.

Shape of aggregates

micelles

Basic shapes:

- spheres –
- cylinders -
- bilayers
- vesicles

Superstructures:

- micellar crystals
- lamellar phases
- bicontinuous networks

Spheres Cylinders Bilayers remperature (°C) 40 BCC Micellar Hexagonal Lamellar solution 35 30 25 20 20 0 40 60 80 100

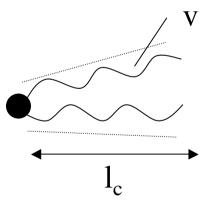
Amphiphile concentration (wt %)

Shape of aggregates

Factors determining the shape

- i) Optimum head group are, a_0
- ii) Critical chain length, I_c
- iii) Hydrocarbon volume, v
- I_c length of fully extended tail
- v volume of the tail# of tails, side groups, ...



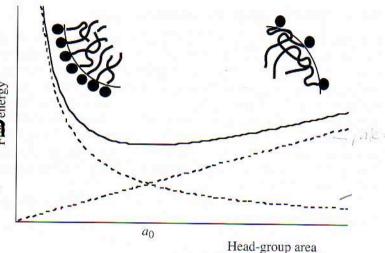


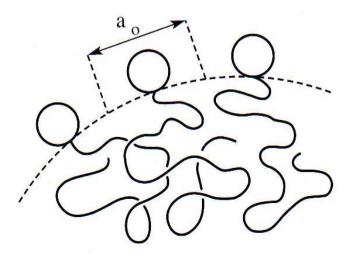
Optimum head group area, a_0

a_0 controlled by

repulsive forces: electrostatic or steric repulsion (can be altered by e.g salt concentration)

attractive forces: protecting the tail





Shape of aggregates

Spheres - smallest area/volume

Volume: $4\pi r^3/3 = Nv$ (N - # of molecules in aggregate)

Surface area: $4\pi r^2 = Na_0$

 \Rightarrow *r*=3*v*/*a*⁰ but r_{max}≤*l*_c

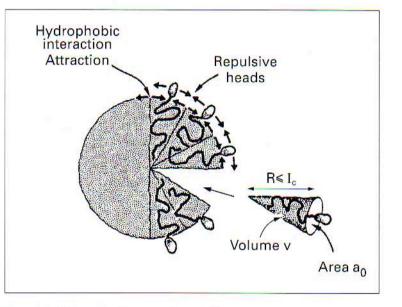


Fig. 4.4. Schematic diagram of a micelle

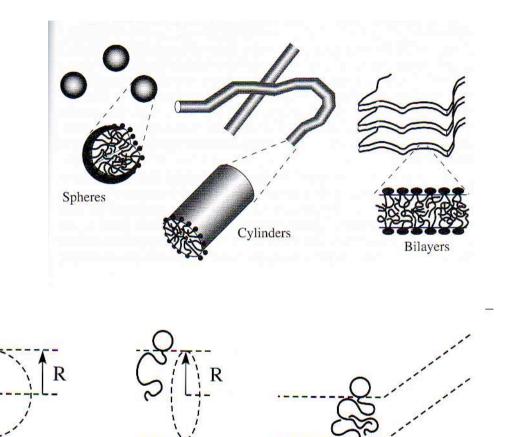
 $v/l_c a_0 \le 1/3$ for spherical micells

Shape of aggregates

а

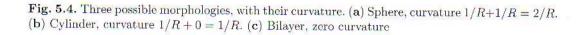
Spheres: $v/l_c a_0 \le 1/3$

Cylinders: $1/3 \le v/l_c a_0 \le 1/2$



С

Bilayer: $v/l_c a_0 > 1/2$



b

Spherical micelles

Spheres formed when: $v/l_c a_0 \le 1/3$

Size of micelles finite \Rightarrow ε_N has a minimum value at an optimum number M

$$\varepsilon_N = \varepsilon_M + \Lambda (N - M)^2$$
$$X_N = N \left[\frac{X_M}{M} \exp \left(\frac{-M \Lambda (M - N)^2}{k_B T} \right) \right]^{N/M}$$

 X_N - volume fraction of micelles with N molecules

 $\left< \left| N - M \right|^2 \right> = \frac{k_B T}{2M\Lambda}$

size distribution

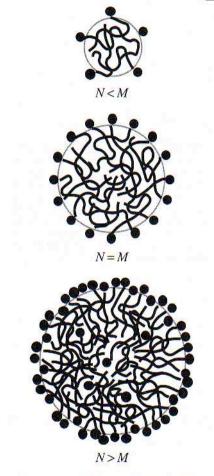


Fig. 9.3 The optimum aggregation number, M for a spherical micelle. Micelles containing a smaller number of molecules have too large an area for each head-group, causing unfavourable water/hydrocarbon interactions, while in micelles larger than the optimum number some head-groups are forced inside the hydrophobic core.

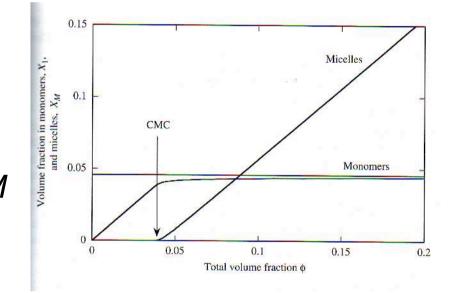
Critical Micelle Concentration

$$X_N = N \left[\frac{X_M}{M} \exp\left(\frac{-M\Lambda (M-N)^2}{k_B T}\right) \right]^{N/M}$$

Monomers vs. micelles

 ϕ - volume fraction of amphiphile $\phi < \phi_c$ - monomers $\phi > \phi_c$ - micelles of average size M

$$X_M = M \left[X_1 \exp\left(\frac{\Delta \varepsilon}{k_B T}\right) \right]^{-1} \qquad \Delta \varepsilon = \varepsilon_1 - \varepsilon_M$$



 $\phi_c = \exp(-\Delta \varepsilon / k_B T) - CMC$

Cylindrical micelles

Cylinders formed when: $1/3 < v/l_c a_0 \le 1/2$

Size of micelles finite but ε_N has a no minimum energy of a molecule in the cylinder is independent of the size \Rightarrow only $2\Delta E_{end}$

$$\varepsilon_N = \varepsilon_\infty + \frac{\alpha k_B T}{N}$$
 , $\frac{\alpha k_B T}{N} = \frac{2\Delta E_{end}}{N}$

- moderate local curvature C = 1 / RRRRC = 2/R
- ⇒ forming larger micelles leads to energy gain, but a loss in entropy

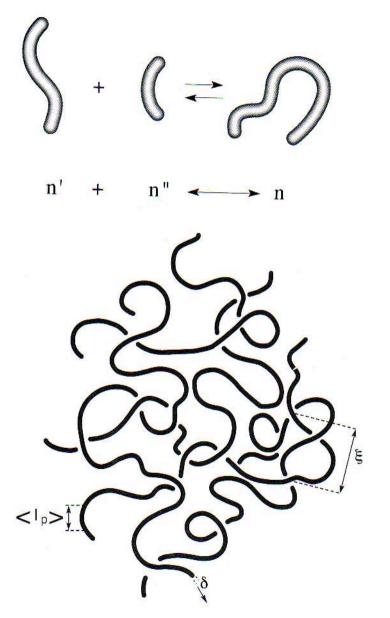
Cylindrical micelles

If ΔE_{end} is large \Rightarrow very long cylinder > μm

$$\langle L \rangle \propto \sqrt{\phi \exp \frac{2\Delta E_{end}}{k_B T}}$$

Polymer-like system but:

- i) Polymer length fixed
- ii) Micelles form and break
- iii) Micellar size varies with external conditions
- iv) Change in dynamic properties

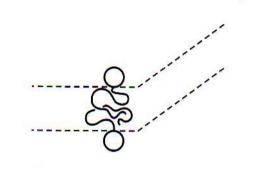


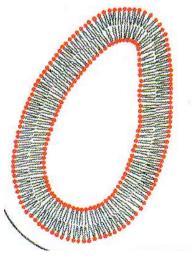
Bilayers

Bilayers formed when: $v/l_c a_0 > 1/2$

 $\Delta E_{edge}(A) \propto A^{1/2}$ (A - area of a layer) dependent on the size of the bilayer \Rightarrow formation of infinite layers

Vesicles - closed surface by a bilayer
⇒ model for cell membranes
⇒ drug delivery systems
⇒ colloidal behaviour





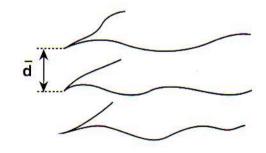
Bilayers

Bilayers are not flat - highly crumpled

Persistence length:

 $\xi_k = a \exp\left(\frac{4\pi k}{\alpha k_B T}\right)$

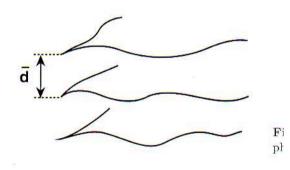
k - bending modulus of membrane For $r > \xi$ direction in membrane is lost



Lamellar phases can anyway be obtained with very large $d \Rightarrow$ confinement between two adjacent planes leads to loss in entropy

Bilayers

Lamellar phase can be distorted by fluctuations forming complex geometries



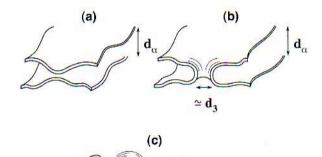


Fig. 5.21. Structure of the sponge phase.
(a) Two bilayers in the lamellar phase.
(b) Formation of an elementary passage.
(c) Proliferation of passages, leading to the sponge phase L₃

Phase behaviour

Self assembled structures are rich in phases

- ⇒ response by changing size and shape and arrangement of micelles
- ⇒ dramatic change in properties

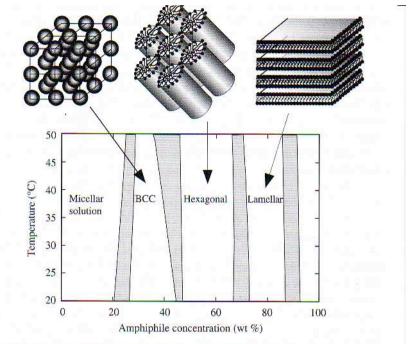
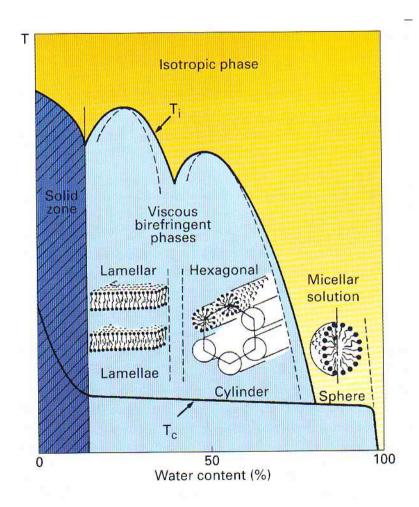
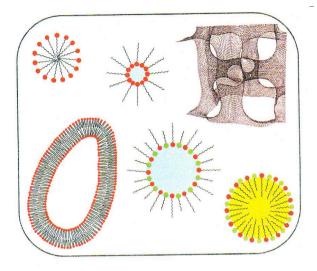


Fig. 9.7 The phase diagram of an amphiphilic copolymer in water, showing schematically the structures of the ordered phase. The material is a short triblock copolymer of ethylene oxide (EO) and propylene oxide (PO), with the structure $(EO)_{37}(PO)_{58}(EO)_{37}$. Data from P. Alexandris, D. Zhou, and A. Khan, *Langmuir*, **12**, 2690 (1996).

Self-assembly





More complexity from:

- several types of amphiphiles
- mixed solvents
- changing pH, temperature, ...

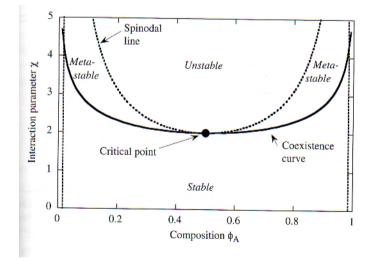
Self assembly in polymers

Mixing polymers

Generally polymers don't mix - why?

Free energy of mixing: $\frac{F}{k_B T} = \phi_A \ln \phi_A + \phi_B \ln \phi_B + \chi \phi_A \phi_B$ (two molecular liquids)

entropic interaction energy



regular solution model

Interaction parameter χ : χ < 2 always single phase χ > 2 phase separation

Mixing polymers

Generally polymers don't mix - why?

For two polymer with N monomers

$$\frac{F^{pol}}{k_B T} = \phi_A \ln \phi_A + \phi_B \ln \phi_B + N \chi \phi_A \phi_B$$

$$\frac{F^{mon}}{k_B T} = \frac{\phi_A}{N} \ln \phi_A + \frac{\phi_B}{N} \ln \phi_B + \chi \phi_A \phi_B$$

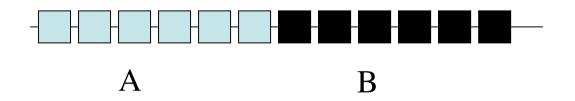
(Flory-Huggins free energy)

Interaction parameter χ : $\chi < 2/N$ always single phase $\chi > 2/N$ phase separation

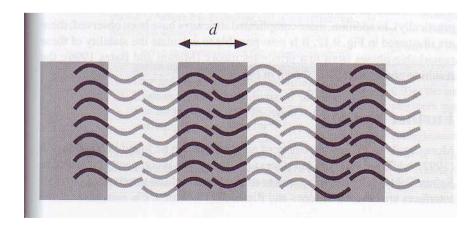
N is usually $\approx 10^3 - 10^6$ \Rightarrow even for small χ we get macro-phase separation

Block copolymers

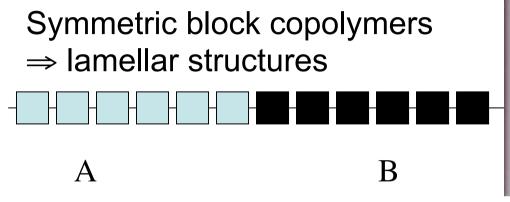
Blocks of different polymers covalently linked together

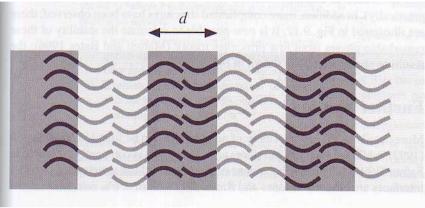


Even if $\chi < 2/N$ the system cannot macro-phase separate (similar to amphiphiles)



Block copolymers





Asymmetric structures ⇒ more complex morphology



Fig. 9.11 A lamellar morphology (top) is less favoured for an asymmetric diblock copolymer, as it requires the two blocks to be stretched to different degrees. Instead, a morphology with curved interfaces (bottom) is favoured.

Templating with block copolymers

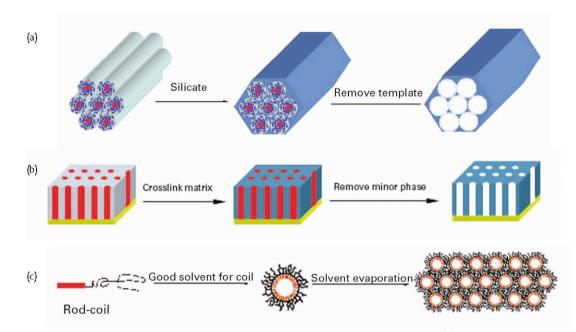


Fig. 6 Schematic of three approaches to generate nanoporous and mesoporous materials with block copolymers. (a) Block copolymer micelle templating for mesoporous inorganic materials. Block copolymer micelles form a hexagonal array, silicate species (in blue) then occupy the spaces between the cylinders. The removal of the micelle template leaves hollow cylinders. (b) Block copolymer matrix for nanoporous materials. Block copolymers form a hexagonal cylinder phase in a bulk or thin film state. Subsequent crosslinking fixes the matrix. Hollow channels are generated by removing the minor phase. (c) Rod-coil block copolymer for microporous materials. Solution-cast micellar films consist of multilayers of hexagonally ordered arrays of spherical holes. (Adapted from⁸⁵ and reprinted with permission. © 1999 American Association for the Advancement of Science.)

Templating with block copolymers

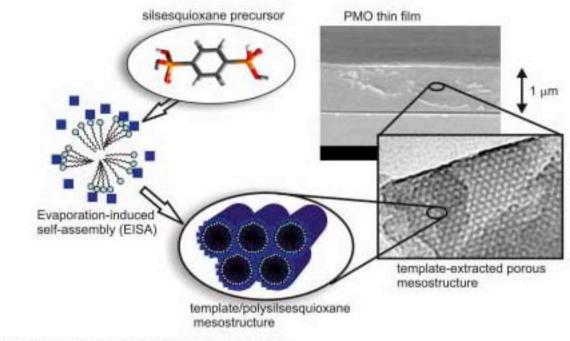
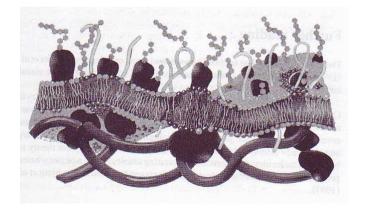


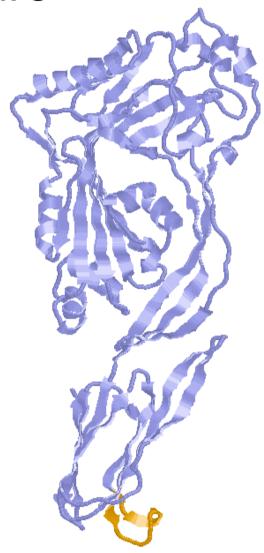
Fig. 6 Schematic Nustration of the synthesis and structure of PMO thinf line.

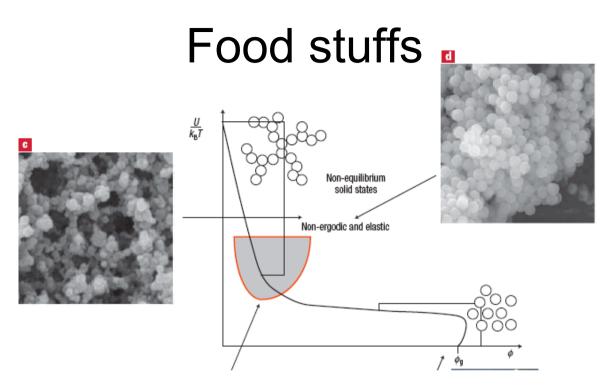
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Soft matter in nature

- Biopolymers
 - DNA
 - Proteins, protein folding
 - Polysaccharides
- Biomembranes







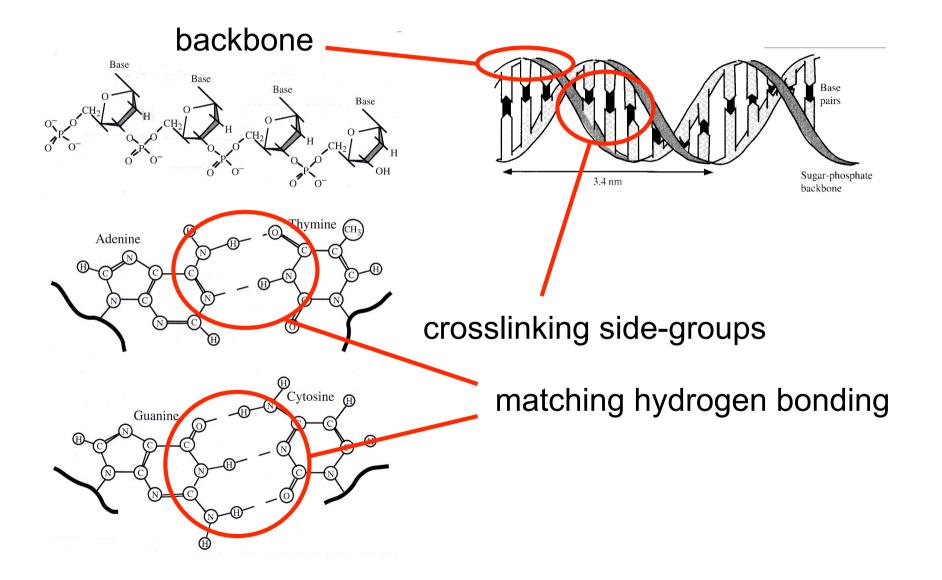
- Colloids (micelles, emulsions, foams...)
- Bio-polymers (proteins, sugars,..)
- Gels (gelatin)
- Amphiphiles (lipids)
- Glass phases
- Liquid crystal phases

Biopolymers

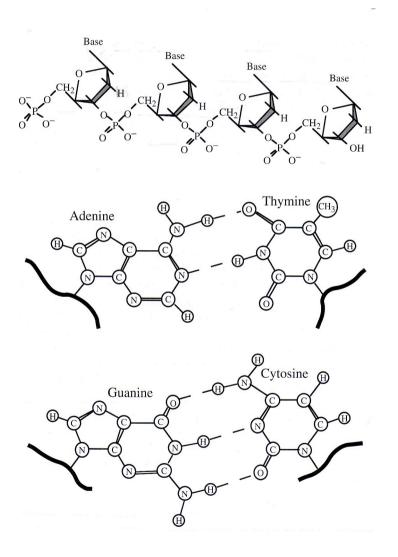
Biopolymers - sequenced copolymers

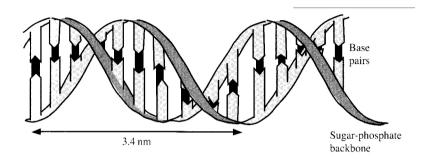
- Nucleic acids sequenced copolymers ex. DNA, RNA
- Proteins ordered sequence of amino acids ex. enzymes, structural elements
- Polysaccharides random copolymers ex. cellulose, starch

DNA



DNA





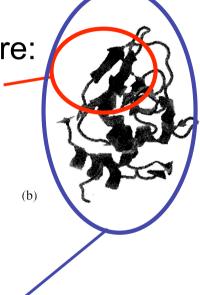
very stiff: *I_p≈200-300 units* long polymer: > 10⁶ monomers contour length µm - mm end-to-end 200nm - 75µm

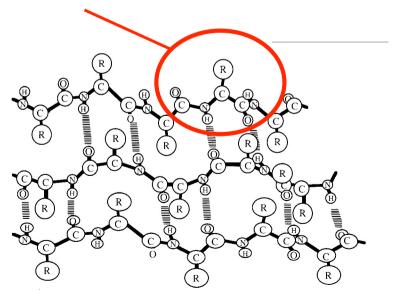
Proteins

Proteins self-assemble and form hierarchical structures

Primary structure: Backbone sequence of amino acids Secondary structure:

- α -helix
- β-sheets

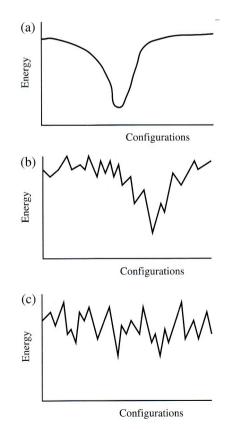


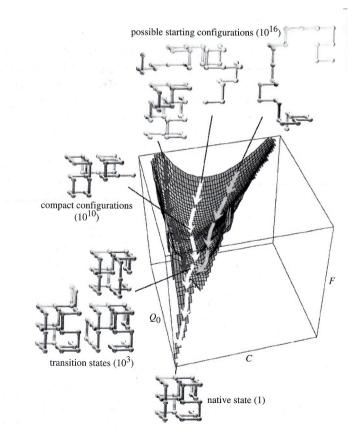


Tertiary structures: Full 3-D arrangement Only one configuration provides the function

Protein folding

Not a random exploration of the energy landscape in the folding process (ms)



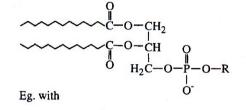


Membranes

-P-O-CH2CH2NH3

Phospholipids

Sphingolipid

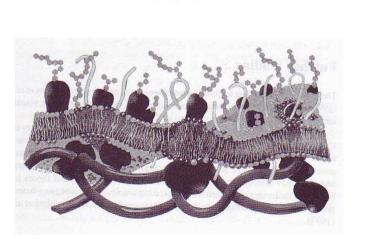


 $R = CH_2CH_2NH_3^+$: Phosphatidylethanolamine

 $R = CH_2CH_2N(CH_3)_3^+$: Phosphatidylcholine (lecithin)

✓ HC = CH - CHOH

-NH-CH



(A sphingomyelin)

Phospholipids from bilayers: 2 chains gives large v $\Rightarrow v/l_c a_0 > 1/2$

Real cell membrane:

- several amphiphiles
- phase separated regions
- membrane proteins
- modifying molecules
- in-plane motions