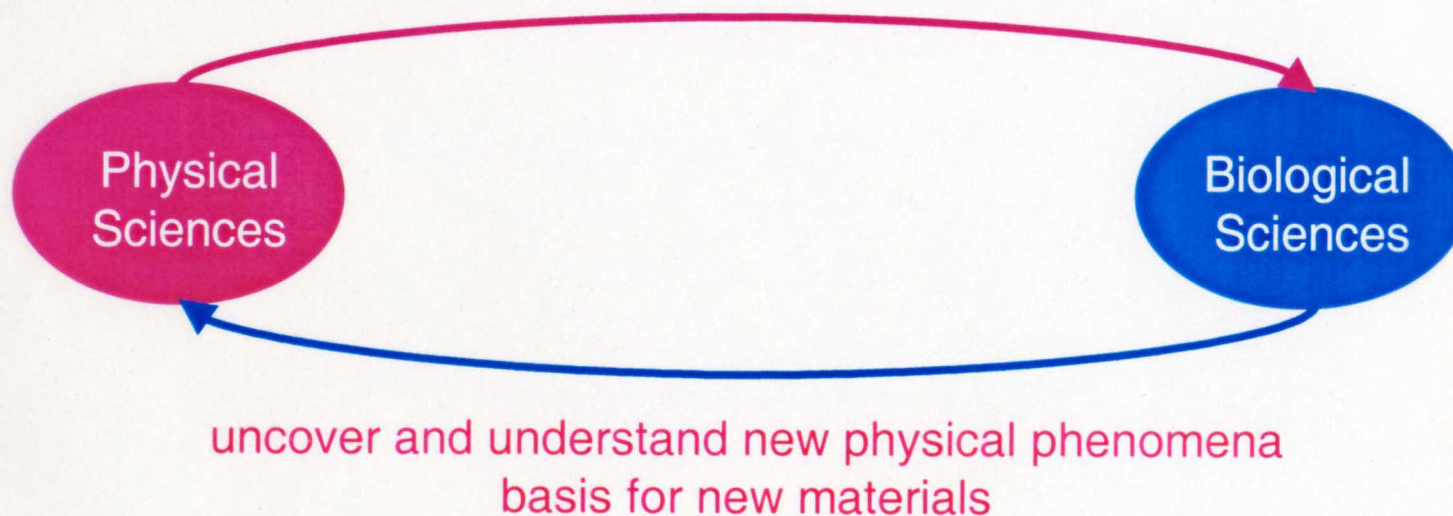


LIPID-PROTEIN INTERACTIONS AT INTERFACES

- Many functions crucial to life are carried out by membrane proteins bound to or embedded in lipid bilayers
- A wide variety of diseases results from deficient or abnormal lipid-protein interactions

Interface between **biological** and **physical** sciences

elucidate the normal functions of these proteins
mechanism by which toxicity is introduced in disease



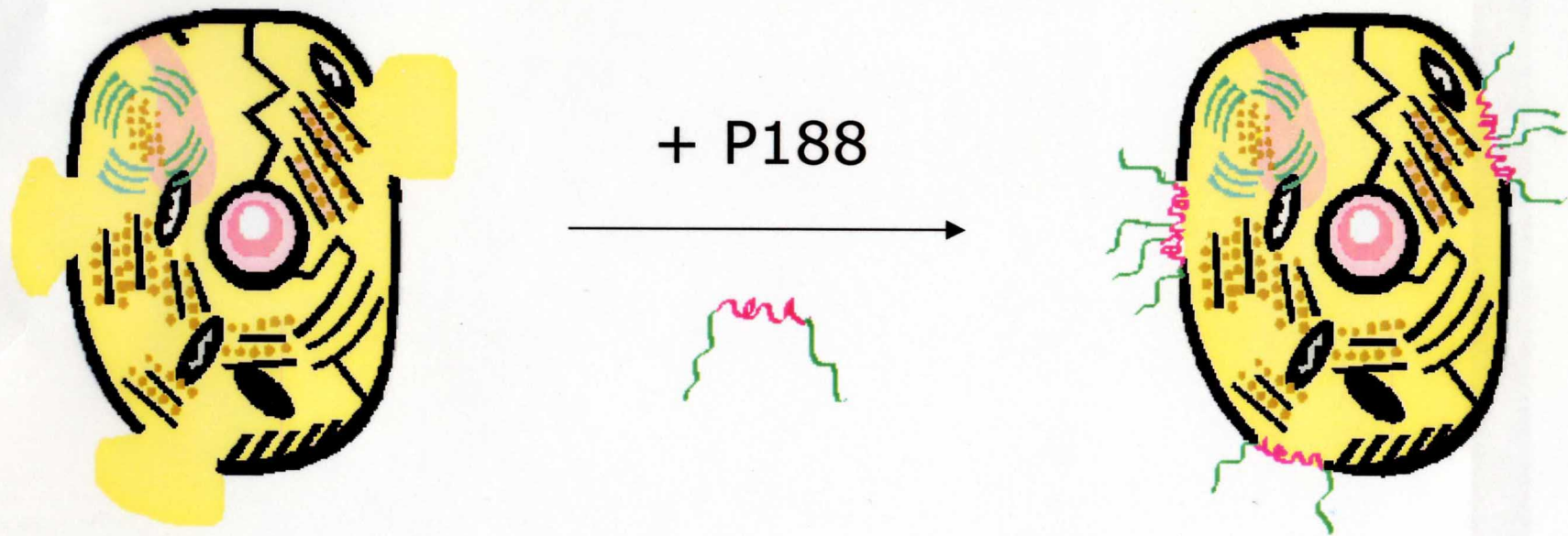
LOSS OF MEMBRANE STRUCTURAL INTEGRITY

Electrically shocked cells suffer a loss of structural integrity.



Electrically shocked cell

COPOLYMER SURFACTANT AS MEMBRANE SEALANT

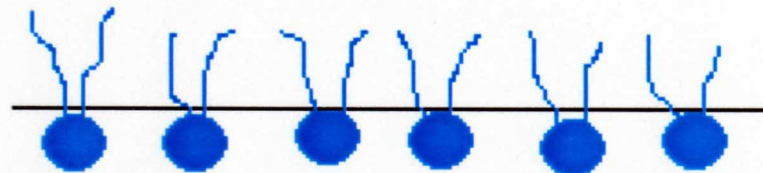


Lee, R.C.; River, L.P.; Pan, F.; Ji, L.; Wollmann, R. L. Proc. Natl. Acad. Sci. USA May 1992, 89, 4524-4528.

QUESTIONS TO BE ADDRESSED

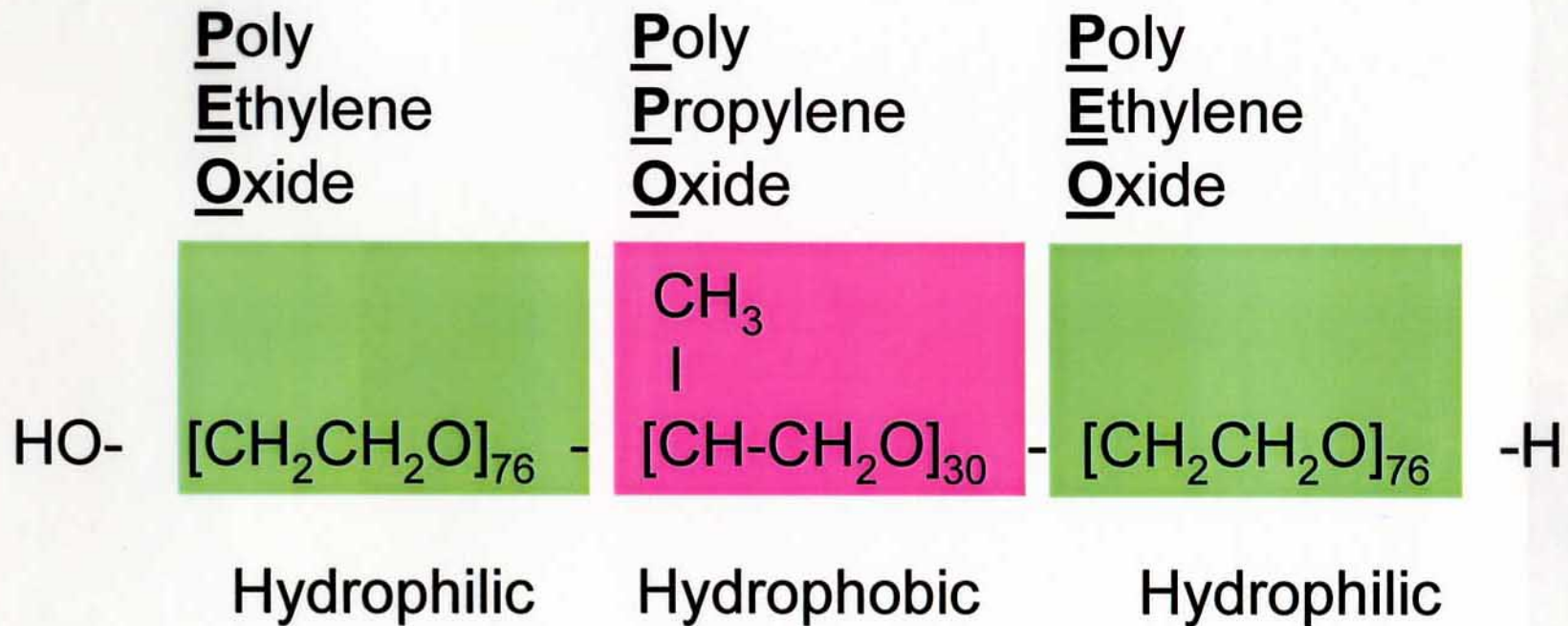
- How does P188 interact with the cell membrane?
 - Does P188 insertion affect membrane morphology?
 - Is the sealing effect localized?
 - What happens to the polymer as the cell heals?
 - Effective treatment strategies? Designer polymers?
-
- How membrane packing regulates P188 insertion
 - How lipid electrostatics influence P188 insertion
 - Morphological changes using FM
 - High resolution imaging using AFM

Lipid monolayers as models for outer leaflet of bilayer

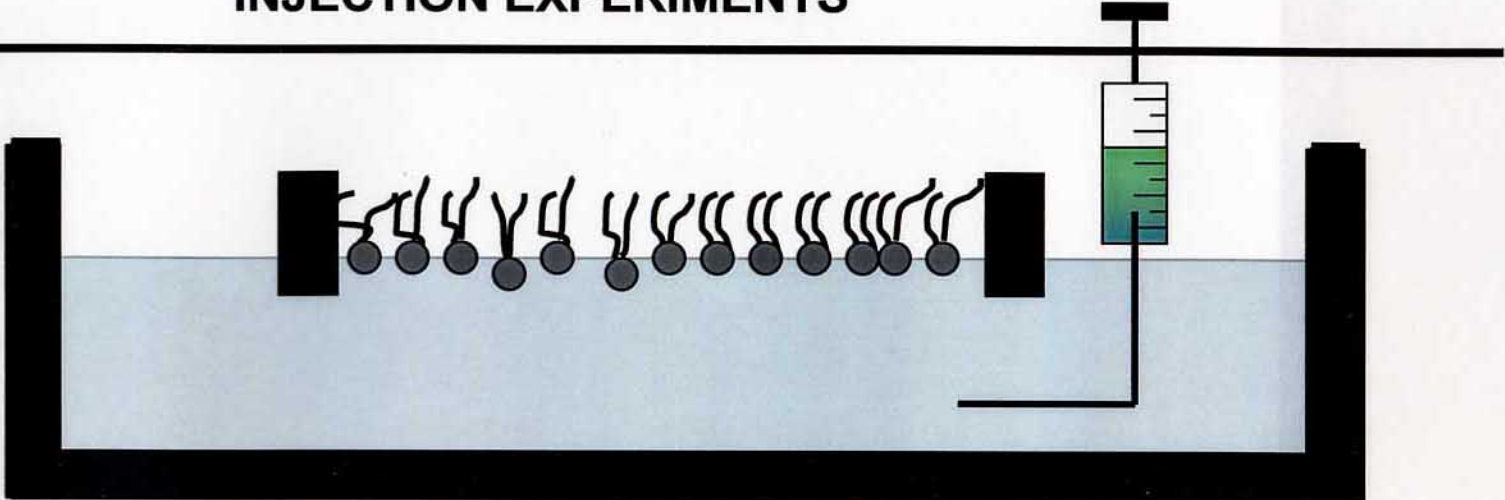


Lipid monolayer
at air-water interface

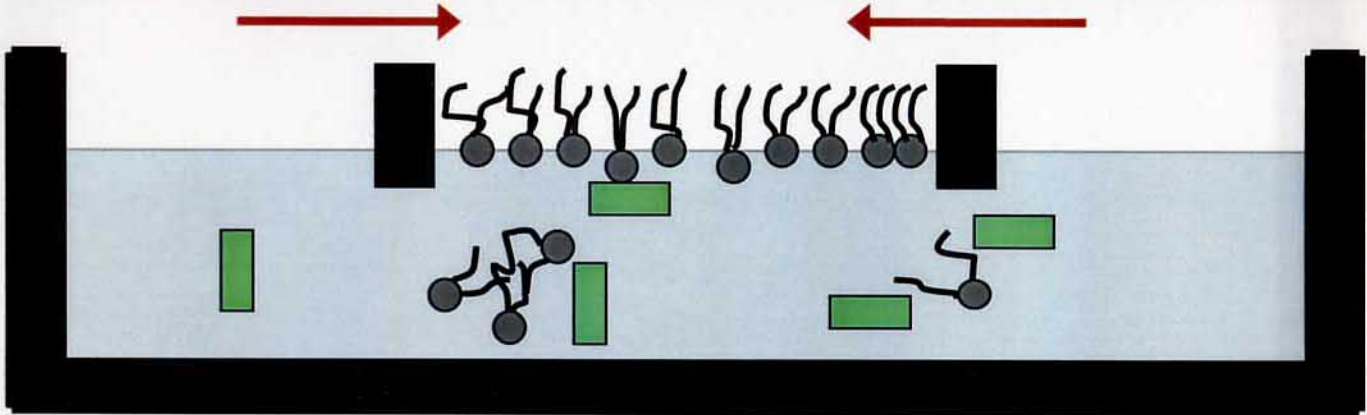
TRIBLOCK COPOLYMER P188



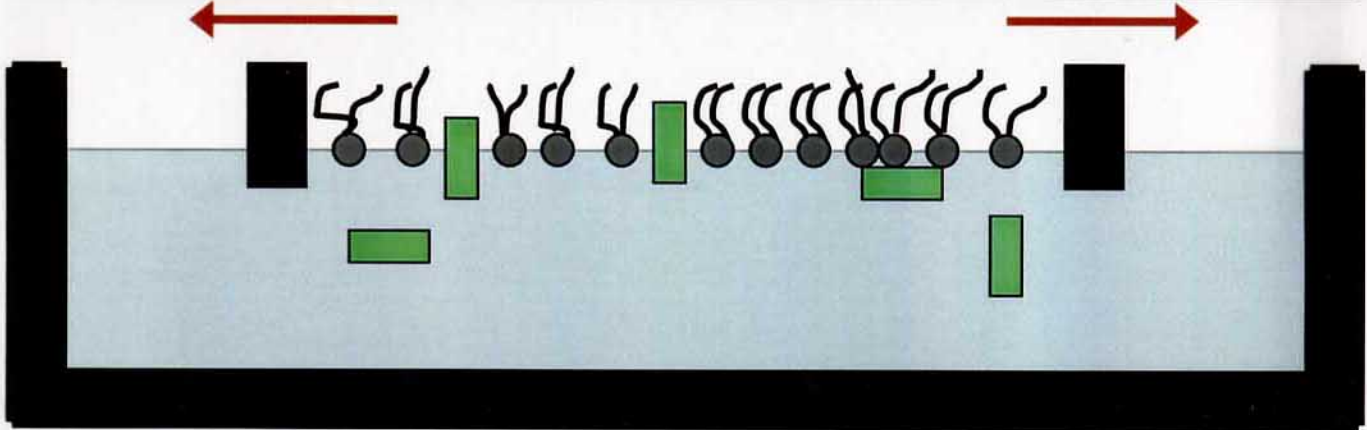
INJECTION EXPERIMENTS



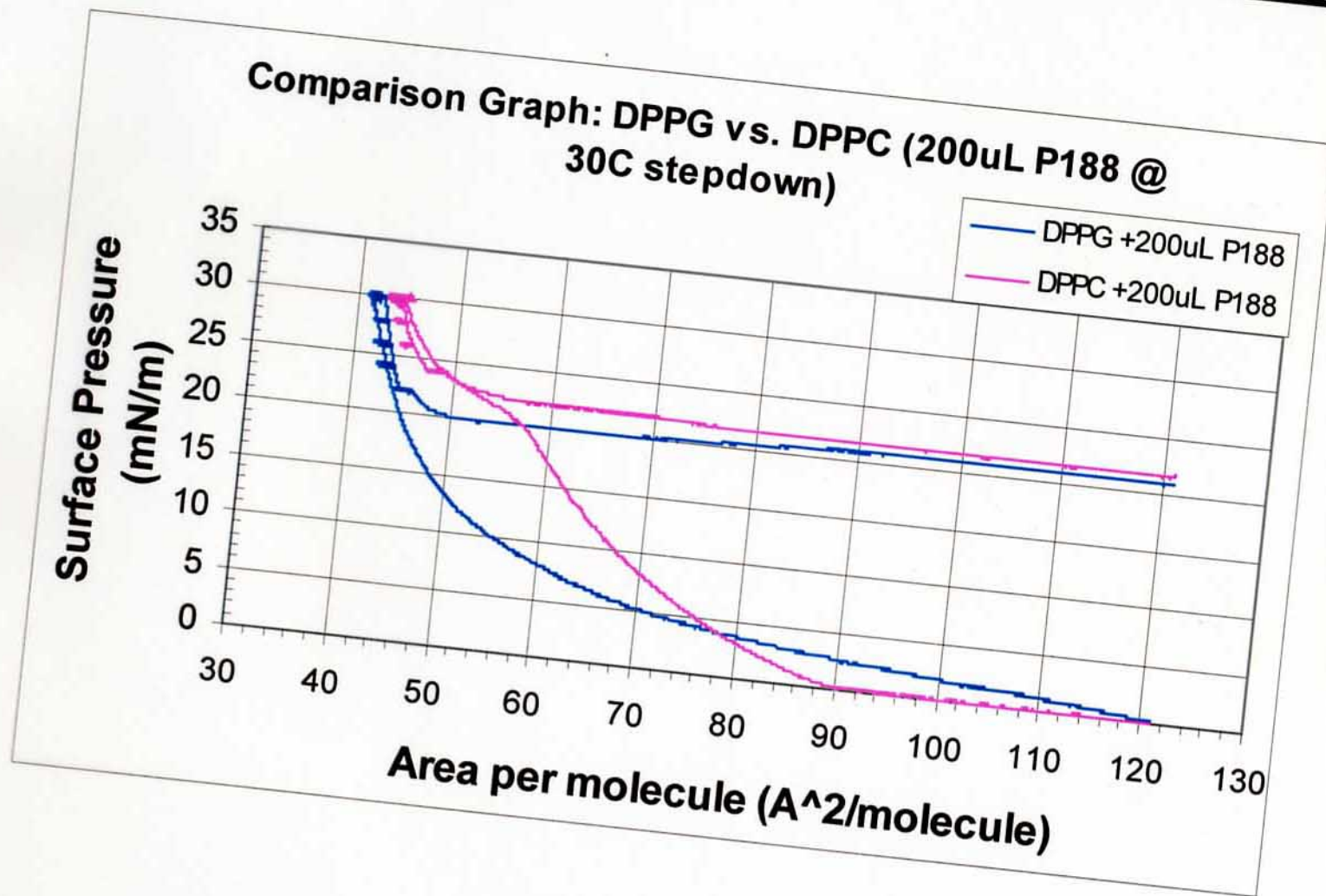
Desorption



Insertion

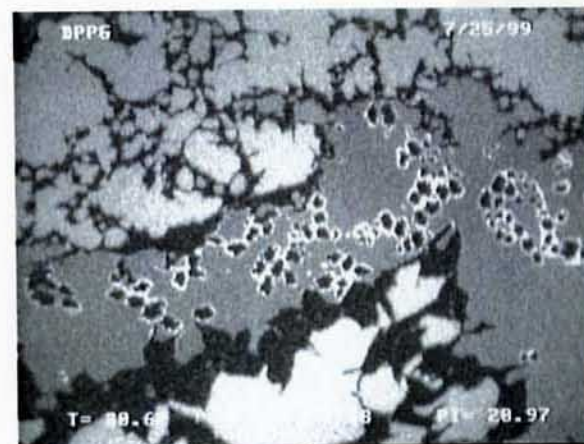
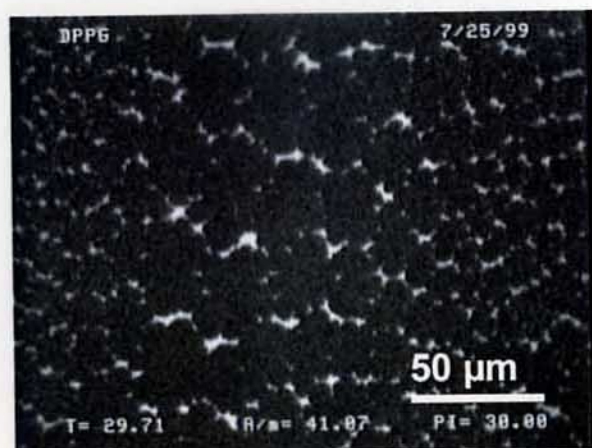


P188 INSERTION REGULATED BY LIPID PACKING DENSITY



DPPC + P188
DPPG + P188

P188 INSERTION INTO DPPG MONOLAYERS



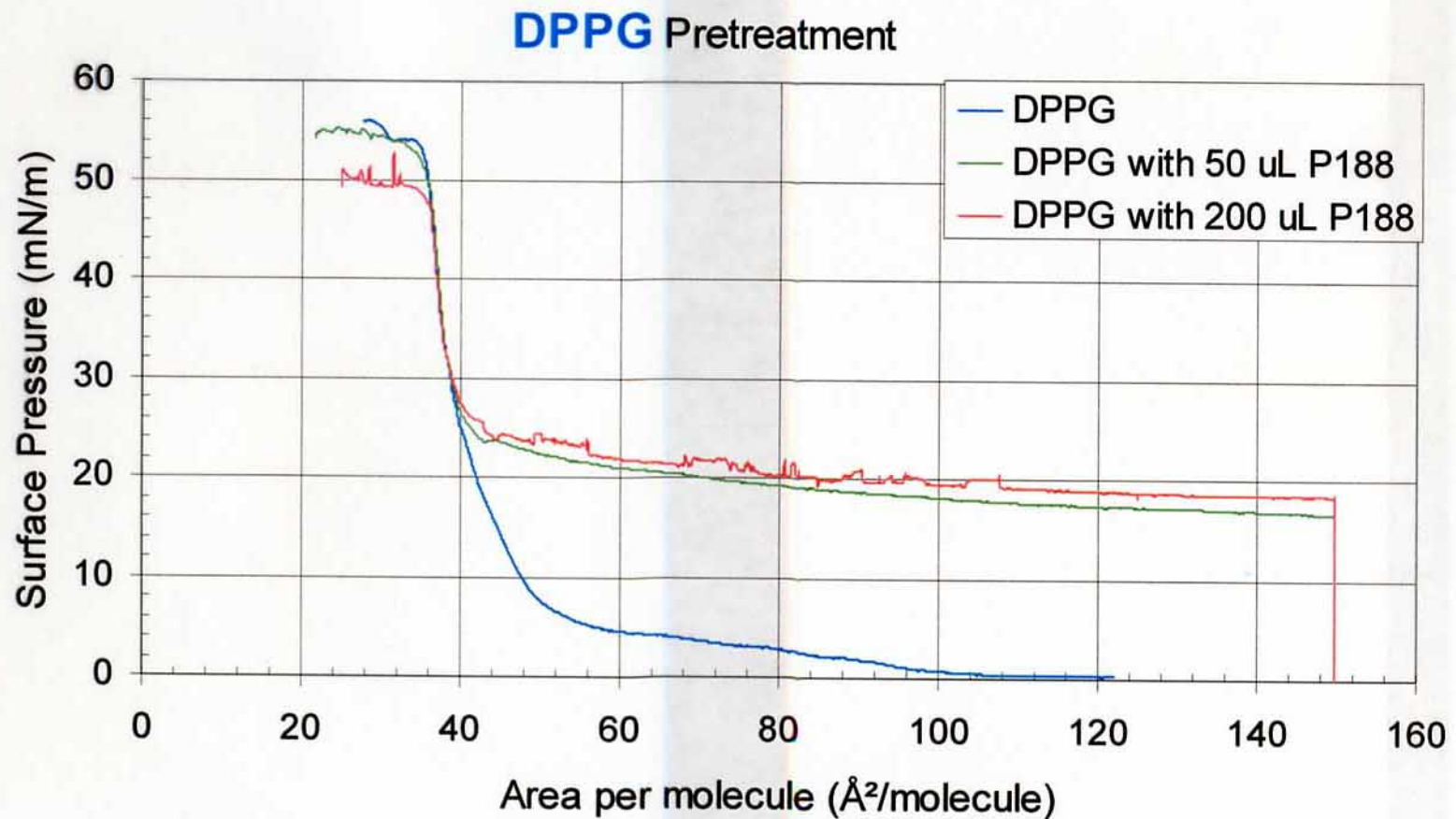
Pure **DPPG** at 30 mN/m
before P188 injection

DPPG + P188 at 20 mN/m
Area/ PG molecule = 78 \AA^2

DPPG + P188 at 20 mN/m
Area/ PG molecule = 120 \AA^2

PRETREATMENT EXPERIMENTS WITH DPPG MONOLAYERS

P188 is added before the lipid monolayer is compressed



EFFECT OF INCREASED MOLECULAR WEIGHT

Poloxamer 238 (P238)

Same Hydrophobic (20% PPO): Hydrophilic (80% PEO) weight ratio as P188

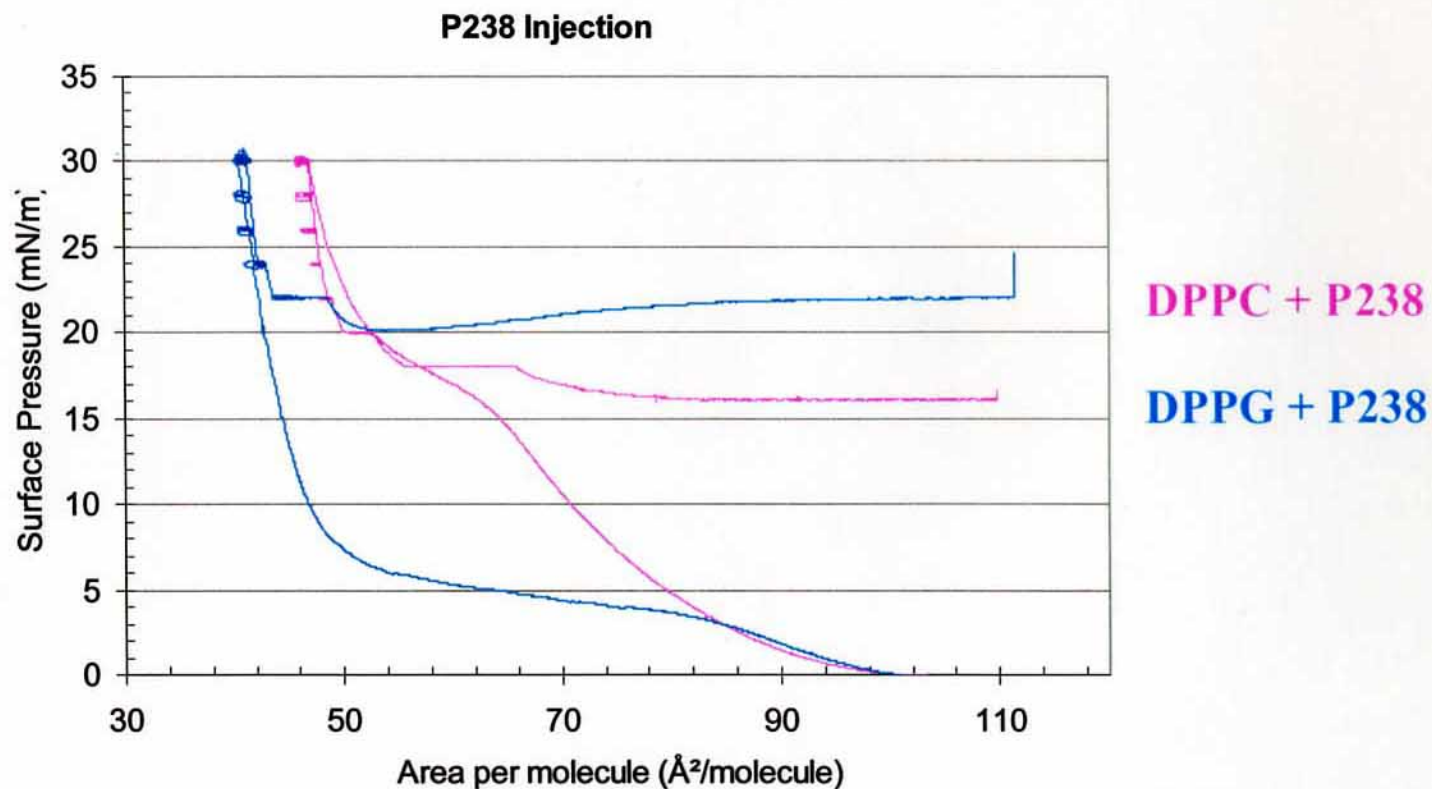


MW (P238) = 10,600 g/mol



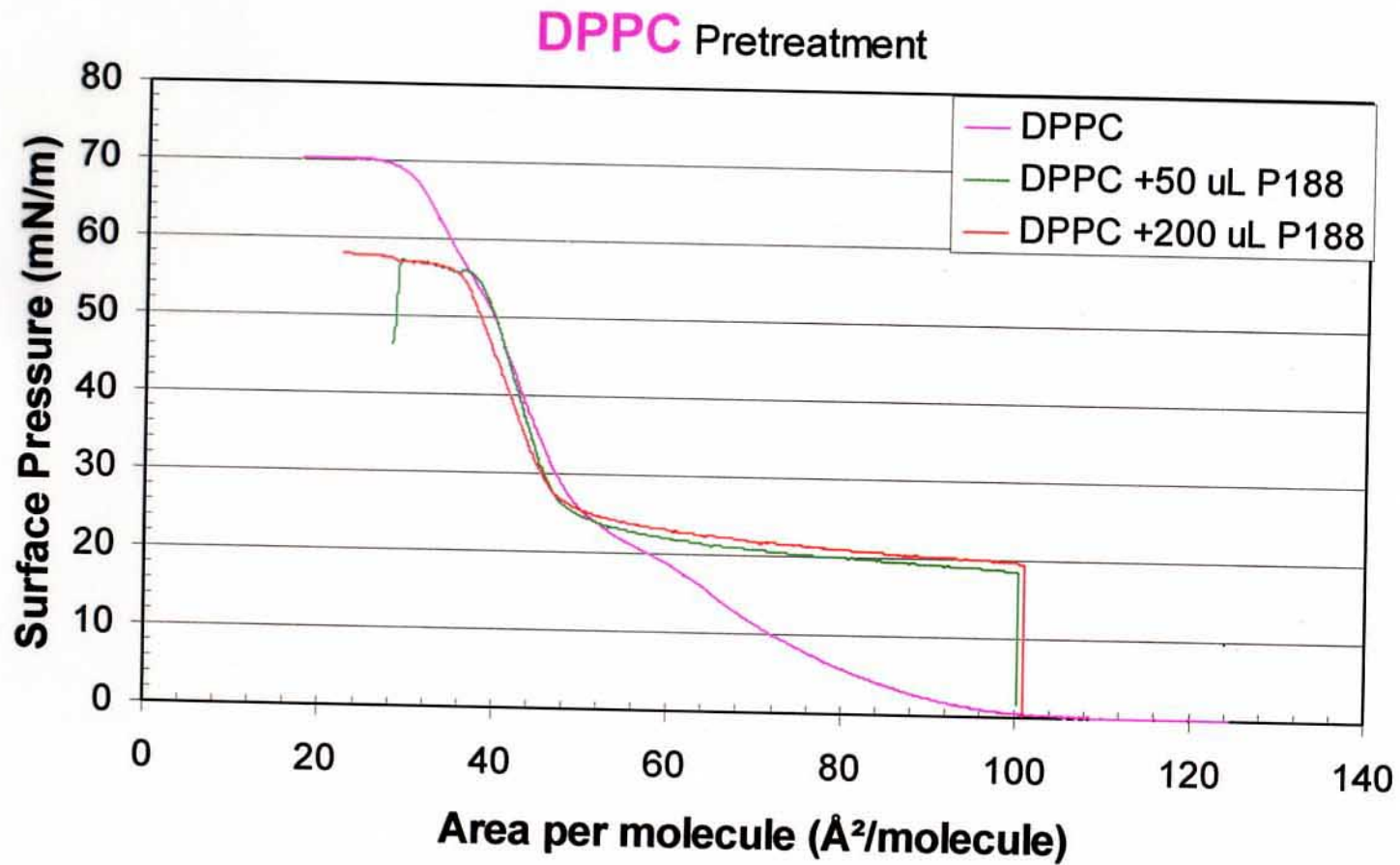
MW (P188) = 8400 g/mol

P238 INJECTED INTO DPPC AND DPPG MONOLAYERS AT 30 mN/m



PRETREATMENT EXPERIMENTS WITH DPPC MONOLAYERS

P188 is added before the lipid monolayer is compressed



Poloxamer 338 (P338)

Same Hydrophobic (20% PPO): Hydrophilic (80% PEO) weight ratio as P188

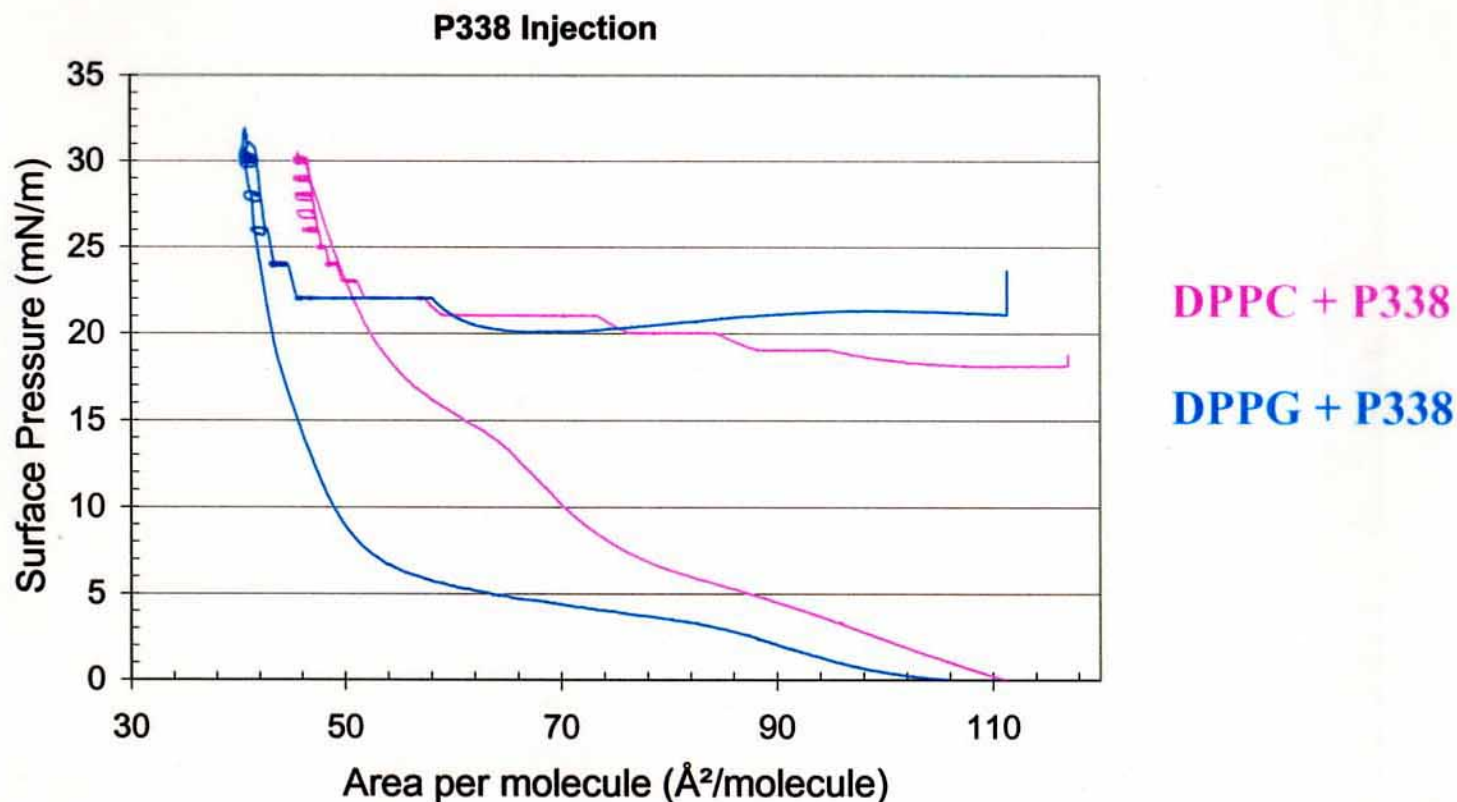


MW (P338) = 14,100 g/mol



MW (P188) = 8400 g/mol

P338 INJECTED INTO DPPC AND DPPG MONOLAYERS AT 30 mN/m



EFFECT OF INCREASED HYDROPHOBICITY

Poloxamer 217 (P217)

P217 weight percentages

30 % PPO (hydrophobic)

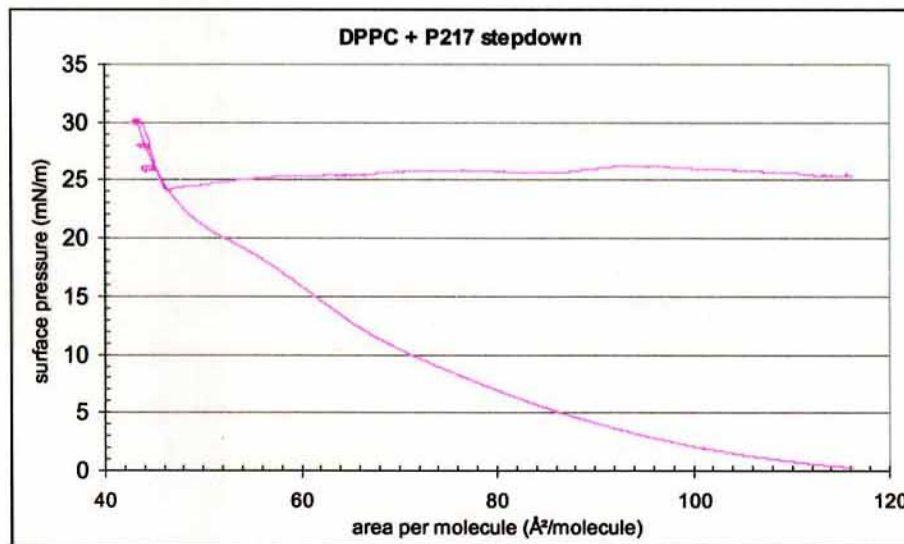
70 % PEO (hydrophilic)

versus

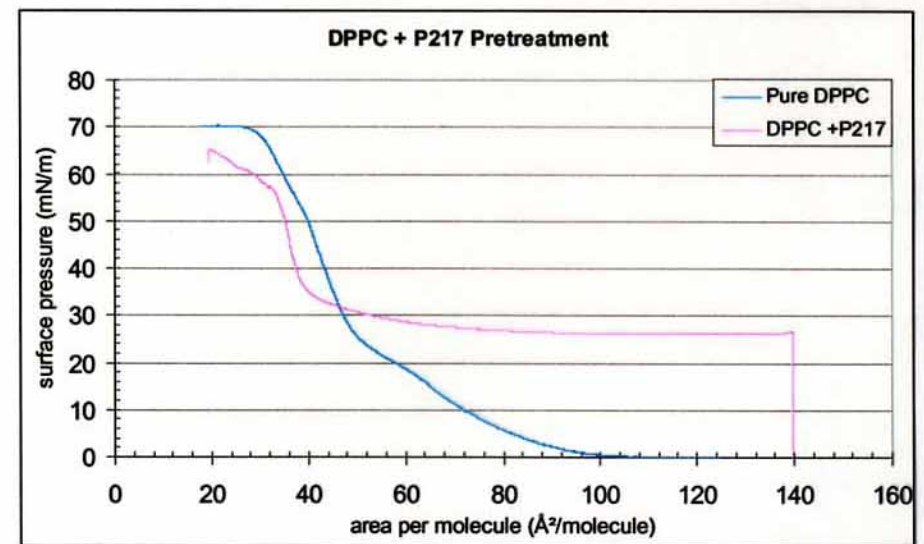
P188 weight percentages

20 % PPO (hydrophobic)

80 % PEO (hydrophilic)

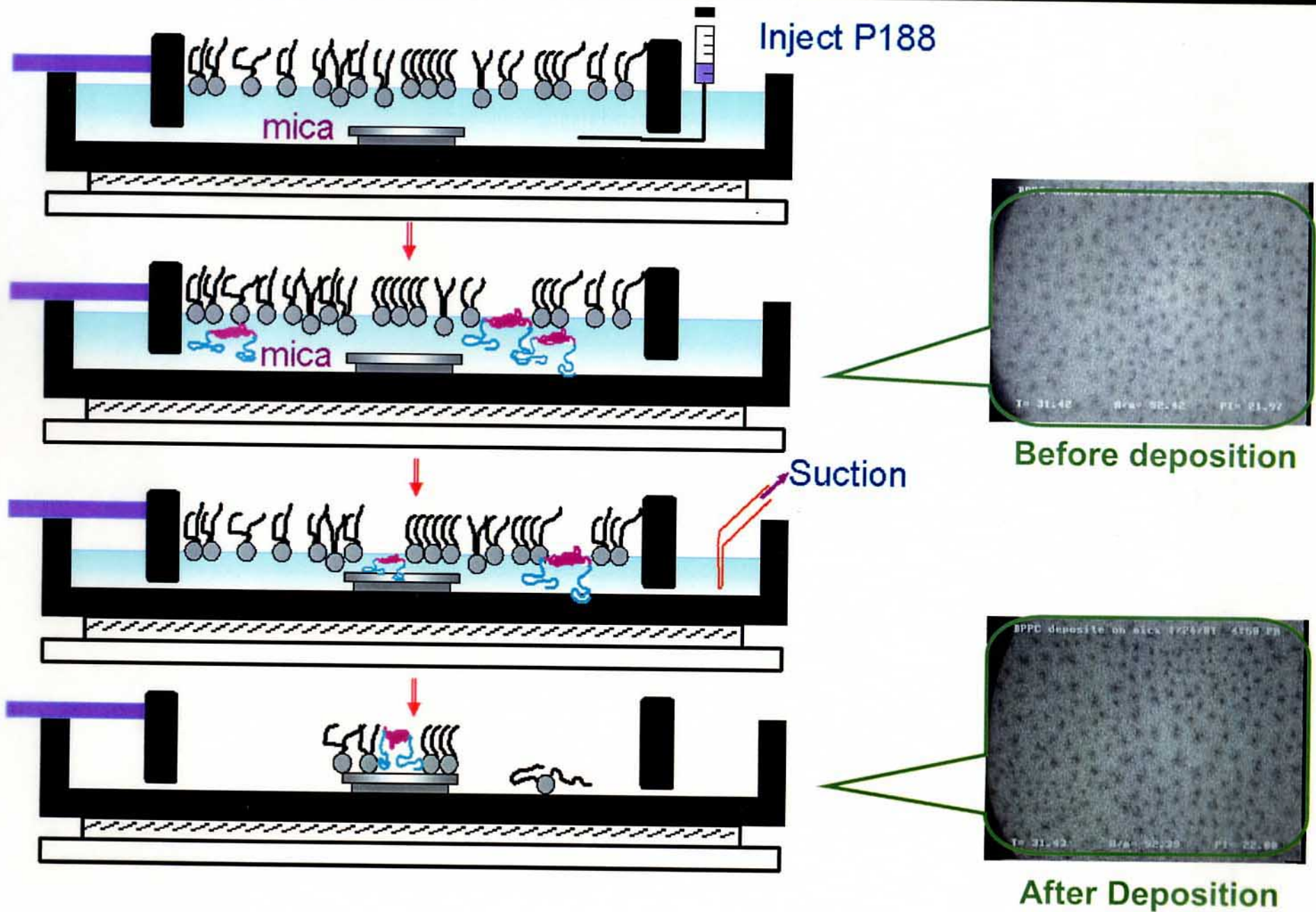


DPPC + P217 Stepdown at 30°C



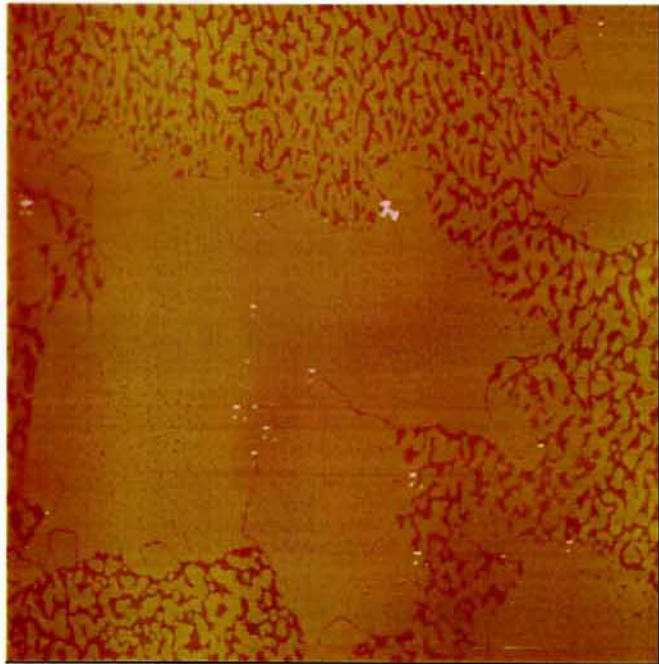
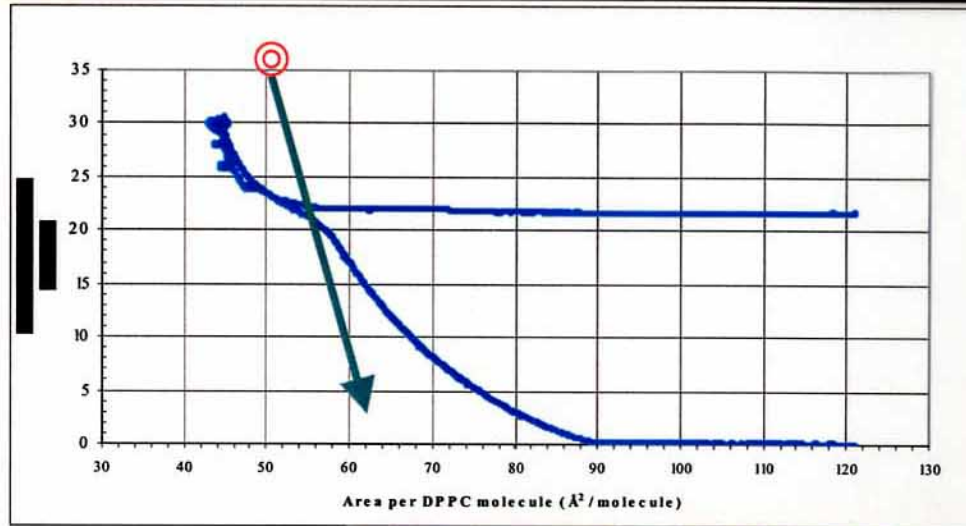
DPPC + P217 Pretreatment at 30°C

DEPOSITION OF MONOLAYERS ONTO SOLID SUBSTRATES

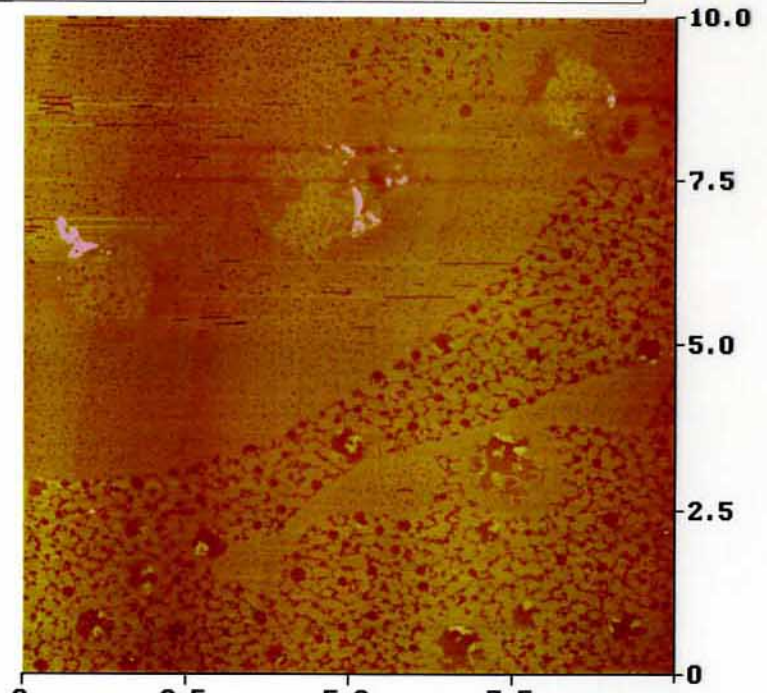


AFM HEIGHT IMAGES - SUPPORTED DPPC FILM AT 24 mN/m

$\pi = 24 \text{ mN/m}$

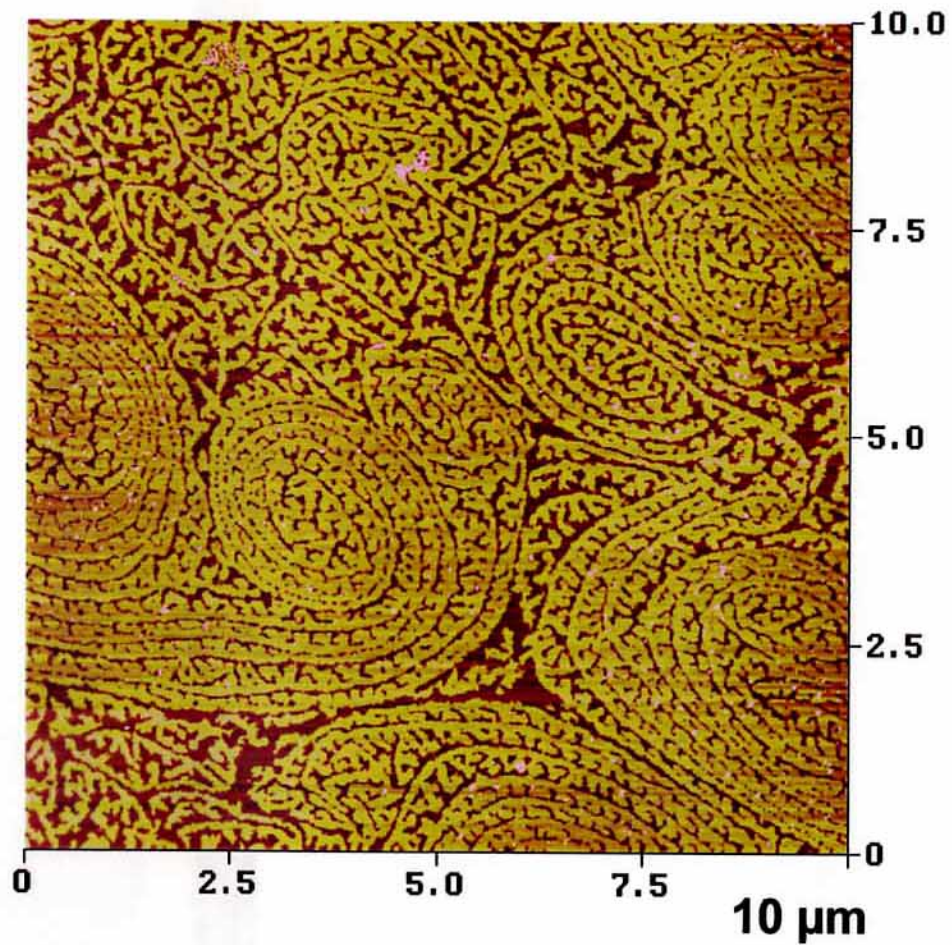


DPPC 15 μm

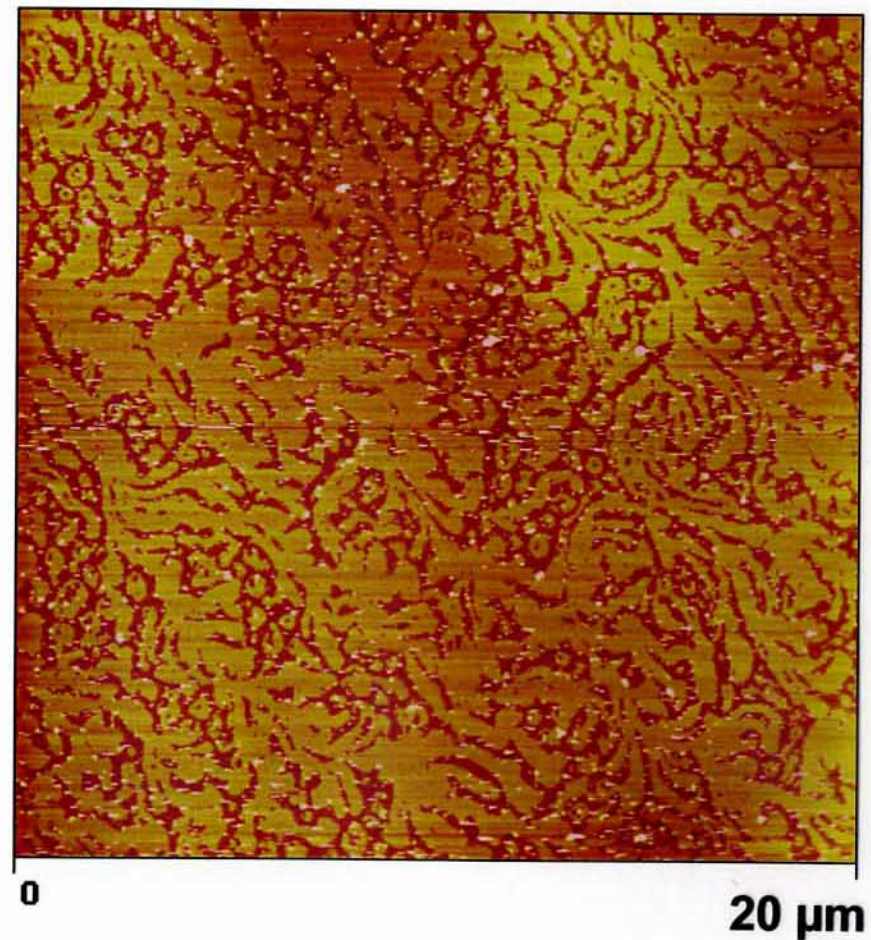


DPPC/P188 10 μm

AFM HEIGHT IMAGES - SUPPORTED DPPC FILM AT 22 and 20 mN/m

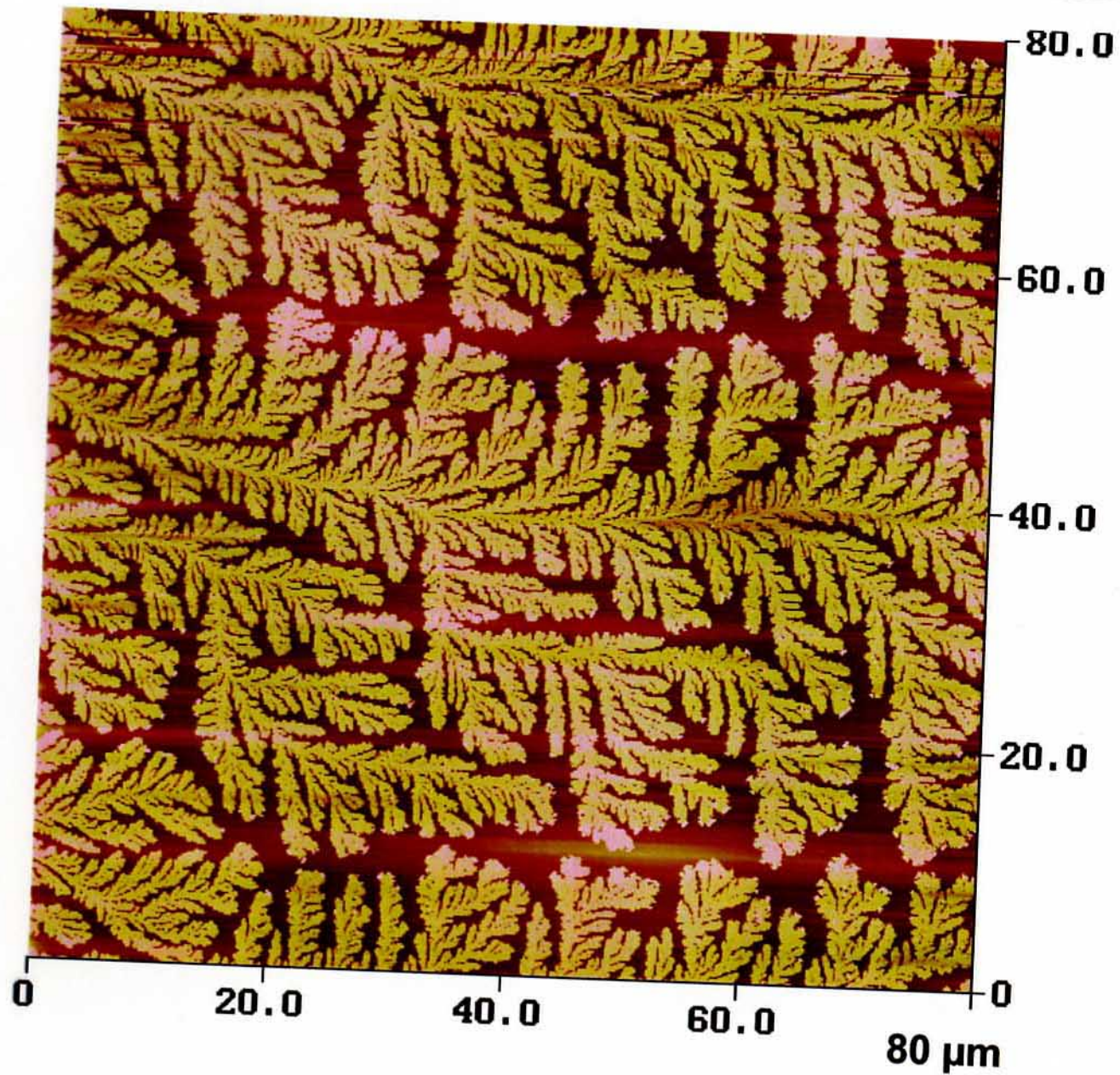


DPPC/P188
 $\pi = 22 \text{ mN/m}$

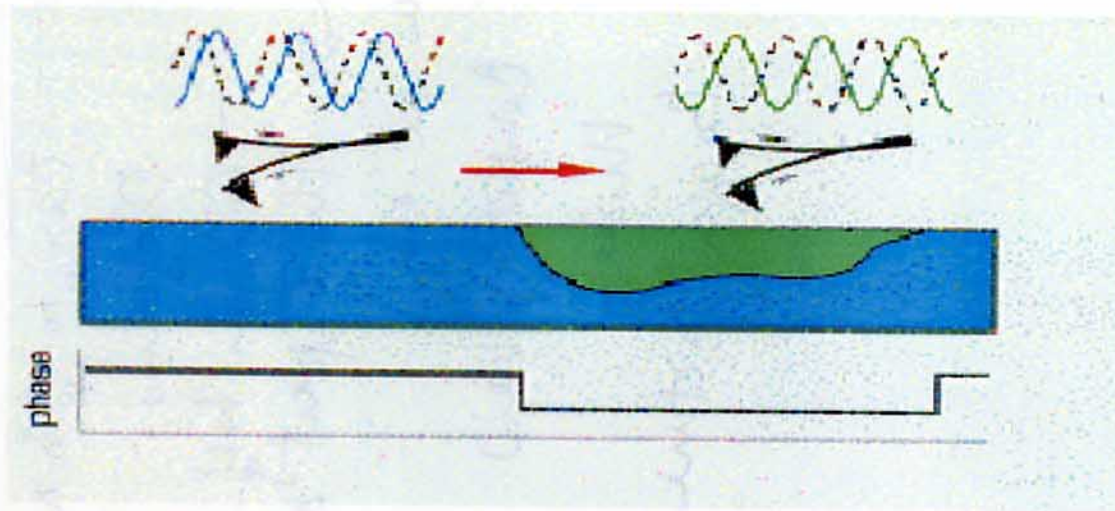


DPPC/P188
 $\pi = 20 \text{ mN/m}$

AFM HEIGHT IMAGES OF SUPPORTED P188

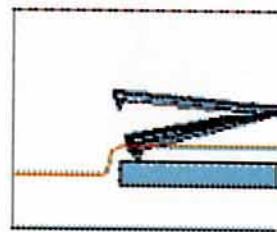


PHASE MODE OF AFM

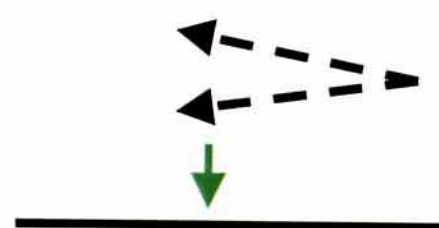
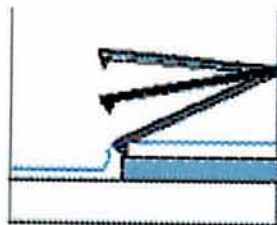


A_0 : the amplitude of freely oscillating cantilever
 A_{sp} : the set amplitude used for feedback control

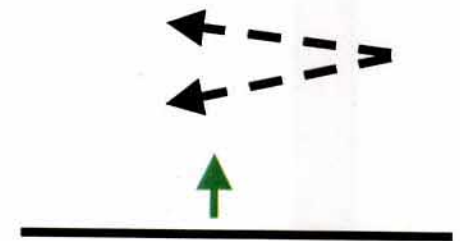
Light tapping
(A_0 small)



Hard tapping
(A_0 large)



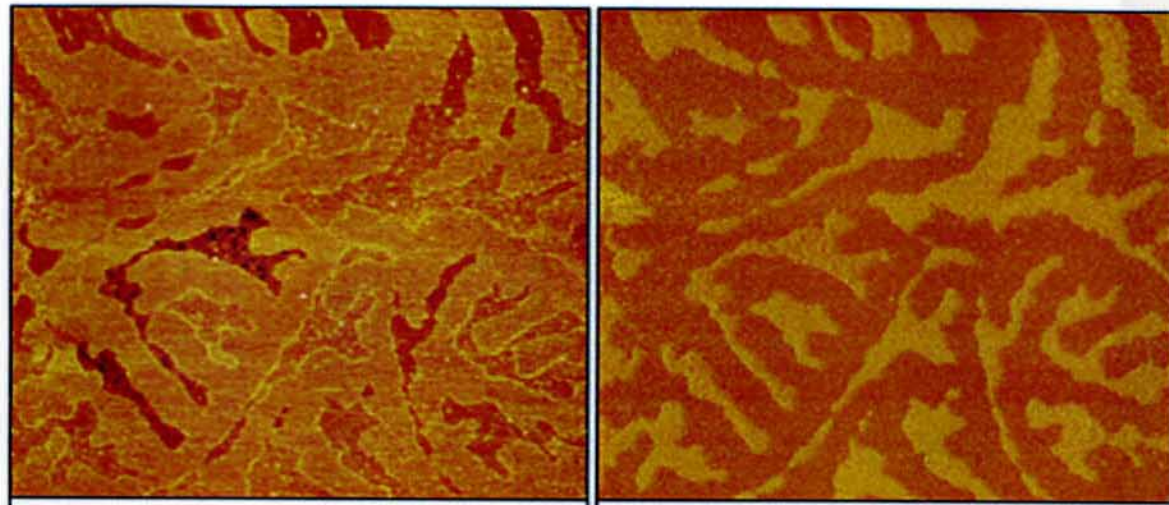
Attractive
Tip-sample interaction



Repulsive
Tip-sample interaction

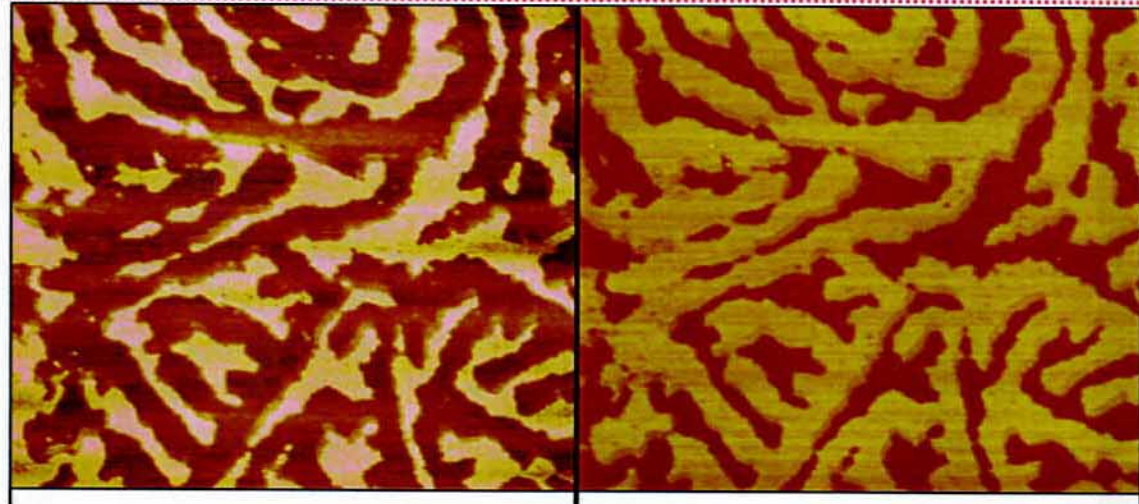
AFM DEMONSTRATES TWO SPECIES PRESENT ON SUPPORT

Light tapping
(Asp = 14 nm
Ao = 18 nm)



0 2.00 μm 0 2.00 μm
Data type Height Phase
Z range 5.000 nm 100.00 $^{\circ}$

Hard tapping
(Asp = 40 nm
Ao = 53 nm)



0 2.00 μm 0 2.00 μm
Data type Height Phase
Z range 5.000 nm 180.0 $^{\circ}$

DPPC+P188
@ 20 mN/m, 30°C

CONCLUSIONS

- P188 will only adsorb into damaged regions of cells
- P188 insertion insensitive to headgroup charge
- P188 is "squeezed out" when cell heals
- AFM can identify to location of the polymer
- Insertion of P188 results in interesting morphology

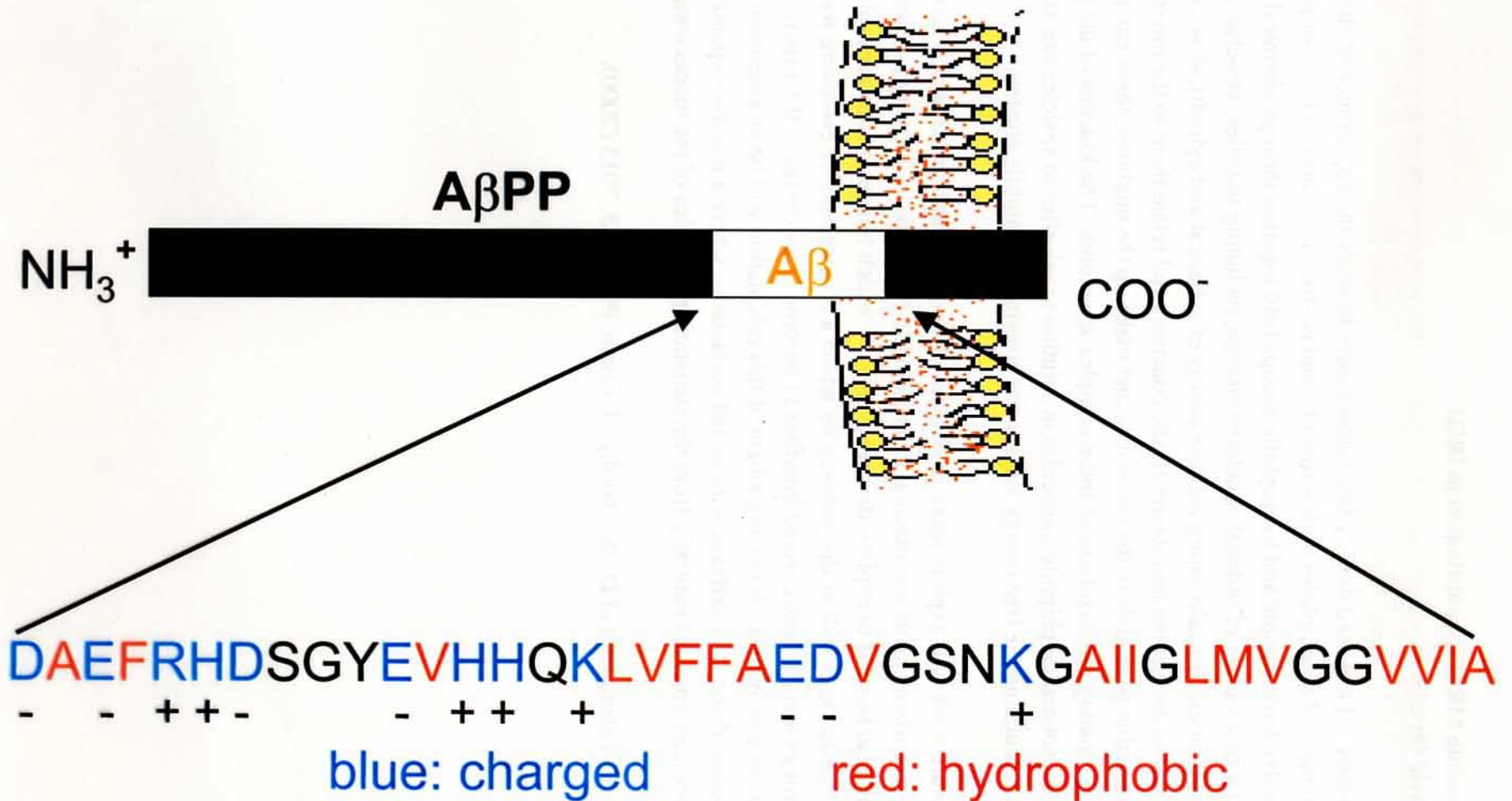
ALZHEIMER'S DISEASE (AD)

- Clinical dementia consistent with AD has long been known
- Bavarian psychiatrist Alois Alzheimer (1906) called attention to
 - > neurofibrillary tanglements
(altered form of the neuronal microtubule protein, tau)
 - > senile plaques
(spherical deposits of extracellular amyloid β -protein)
 - > vascular deposits
($A\beta$)
- Sequence

Glenner & Wong, *Biochem. Biophys. Res. Comm.* (1984) **120**: 885-890
Masters et al., *Proc. Nat'l. Acad. Sci. USA* (1985) **82**: 4245-4249
- Gene encoding the amyloid precursor protein (β APP)

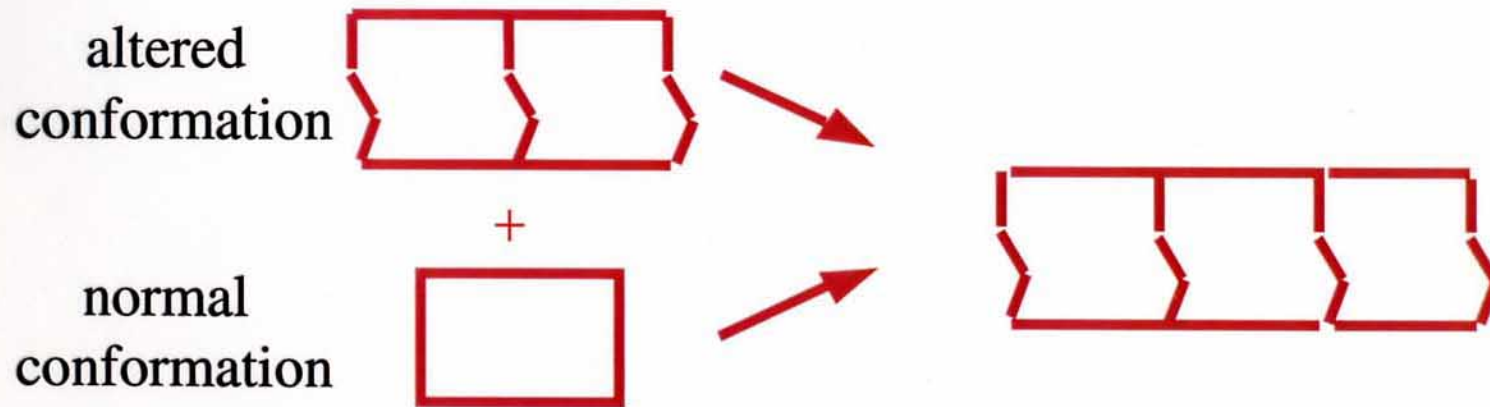
Kang et al., *Nature* (1987) **325**: 733-736

AMYLOID-BETA (A β) PROTEIN



Self-Perpetuating Change in Protein Conformation

- A mechanism of disease
Alzheimer's Disease $A\beta$,
Huntington's Disease, Parkinson's Disease,
Systemic Amyloidoses, Prion Diseases (Mad Cows)?
- A mechanism of inheritance
Yeast Ψ Element, [URE3], Het-S, etc.



AMYLOID-BETA (A β) PROTEIN

- Formed by proteolytic processing
 β APP \rightarrow A β
- 39 – 43 amino acid residues
 A β 1-42 \rightarrow major component in senile plaques
 A β 1-40 \rightarrow major component in vascular deposits
- Neurotoxicity studies in vitro
 A β \rightarrow causal agent for cell death
 Yankner *et al.*, *Science* (1989) **245**: 417-420
- For neurotoxic response
 A β specimens need to be aged or aggregated
- Aggregation structures and kinetics depend on
 - \rightarrow pH
 - \rightarrow ionic strength
 - \rightarrow peptide concentration
 - \rightarrow temperature
 - \rightarrow mixing protocol Synder *et al.*, *Biophys. J.* (1994) **67**: 1216-1228

LIPID-INDUCED CONFORMATIONAL CHANGE

- Absence of lipid

low peptide concentration -> random coil

high peptide concentration -> β -sheet

- Presence of negatively charged lipid vesicles

low peptide concentration

random coil -> β -sheet

- Presence of neutral lipid vesicles

-> no change in conformation detected at low peptide concentration

- Presence of 0.1 M NaCl and negatively charged lipid vesicles

-> no change in conformation detected at low peptide concentration

Terzi *et al.*, *Biochemistry* **33** (1994) 7434-7441

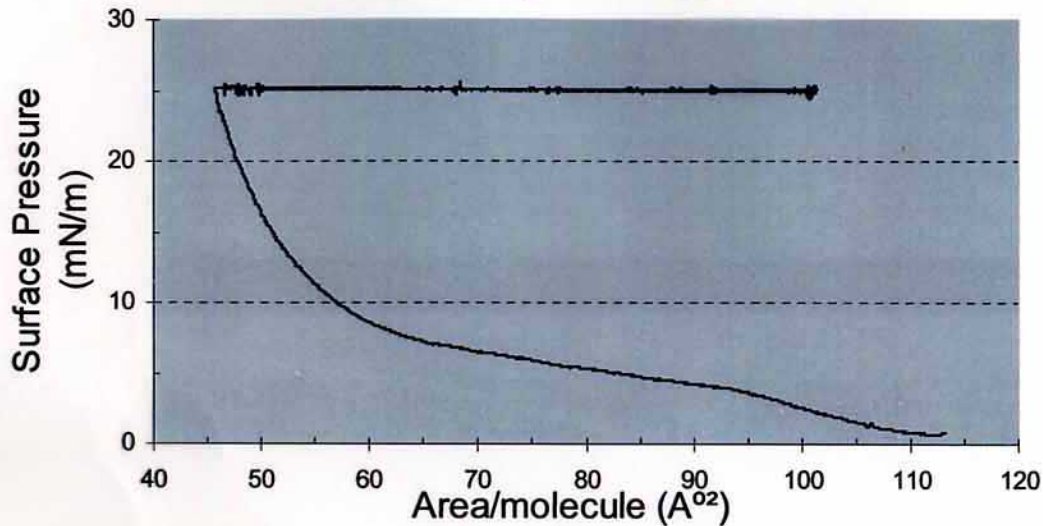
Terzi *et al.*, *J. Mol. Biol.* **252** (1995) 633-642

Yanagisawa *et al.*, *Nature Medicine* **1** (1995) 1062-1066

Choo-Smith & Surewicz, *Febs Letters* **402** (1997) 95-98

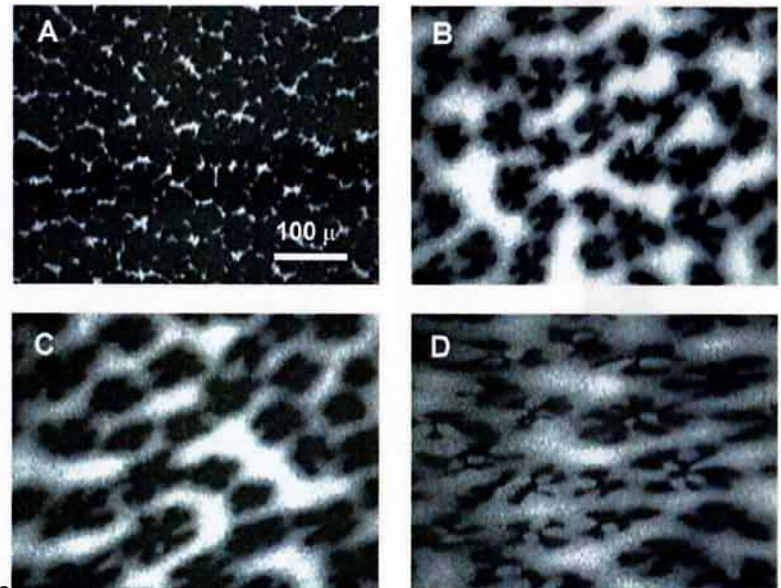
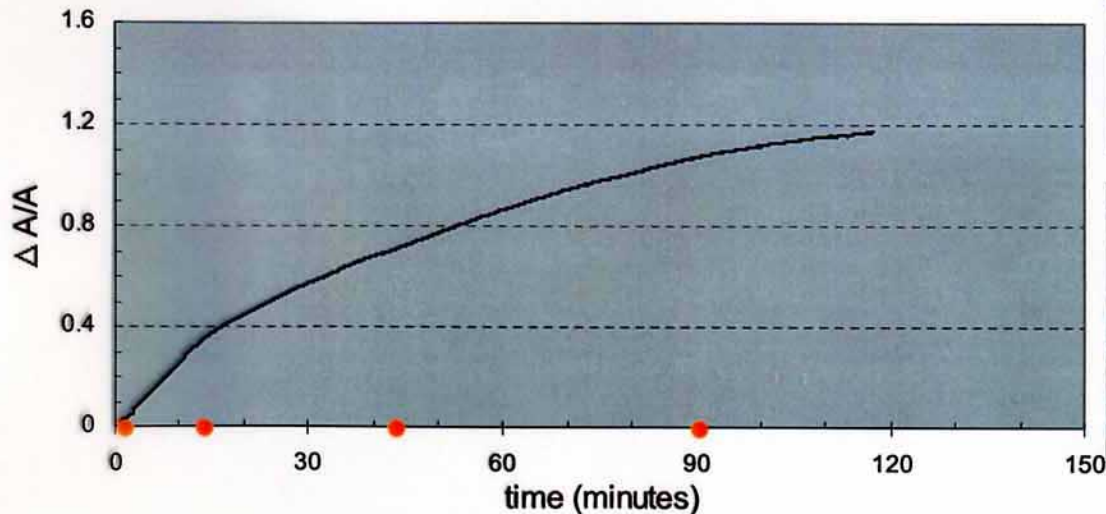
INSERTION OF A β DISSOLVED IN PURE WATER INTO DPPG FILMS

AB1-40 injection into 25 mN/m DPPG on a pure water subphase T=30°



The fluorescence micrographs below shows the change in domain morphology of a DPPG monolayer at 25 mN/m after AB40 is injected into the subphase. The images were taken just before injection, and 15, 45 and 90 minutes after injection of the peptide (in alphabetical order). The corresponding locations are marked on the lower graph.

Relative increase in the Area/molcule after AB40 injection



ISOTHERM RESULTS

		Pure Water	Buffer + Salt
D	20 mN/m	200+	52
P	23 mN/m		23
P	25 mN/m	115	10
G	30 mN/m	30	
D	20 mN/m	25	82
P	23 mN/m	<3	29
P	25 mN/m	0	14
C	30 mN/m		2
D	20 mN/m	88	
T	25 mN/m	80	35
A			
P	30 mN/m	54	10

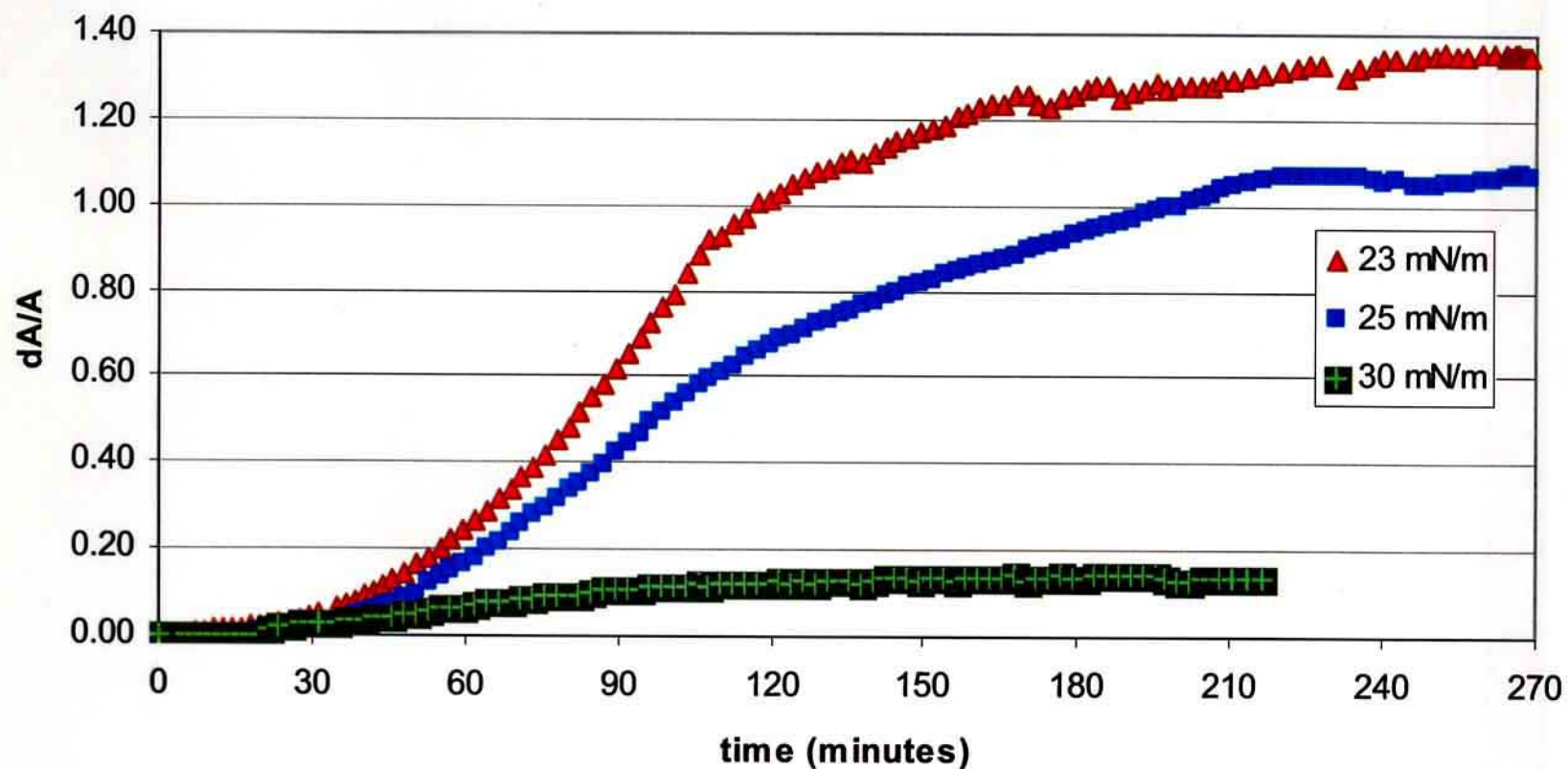
		Pure Water	Buffer+Salt
D P P C	15 mN/m	30	
	20 mN/m	25	82
	23 mN/m	<3	29
	25 mN/m	0	14
	30 mN/m		<2
D M P E	15 mN/m	130+	50
	20 mN/m	50+	20
	23 mN/m		12
	25 mN/m	9	

The numbers here are the total percentage change in the area per phospholipid molecule after A β 40 was injected and the system equilibrated. A β 40 was dissolved in H₂O prior to injection. The temperature

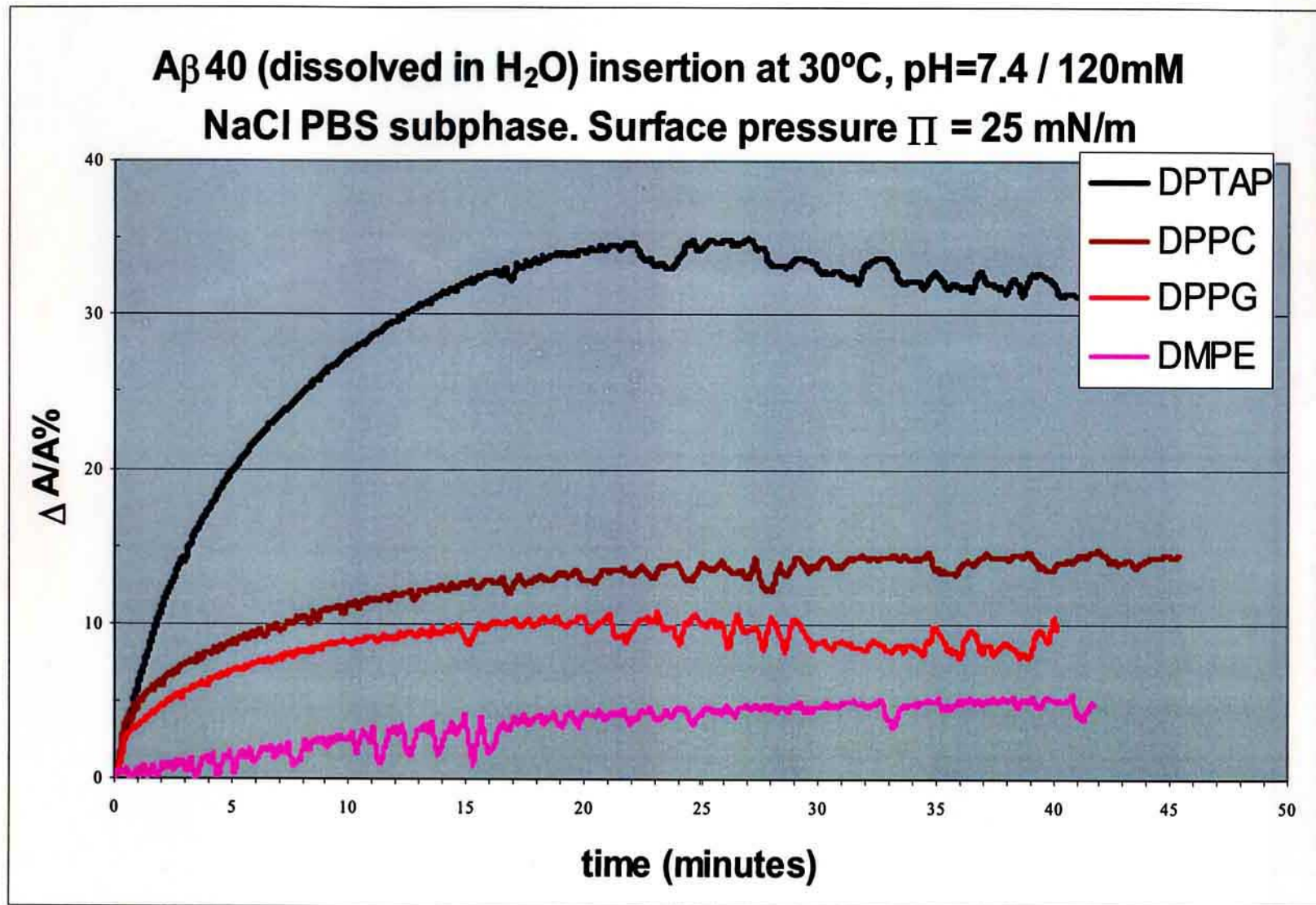
in all the experiments shown here are 30 °C. The subphase is pH=7.4 phosphate buffer with 120mM NaCl where indicated. The peptide concentration in the total subphase volume is 250 nM.

A β DISSOLVED IN DMSO: INSERTION RESULTS

DPPG, pure water subphase, T=30°C

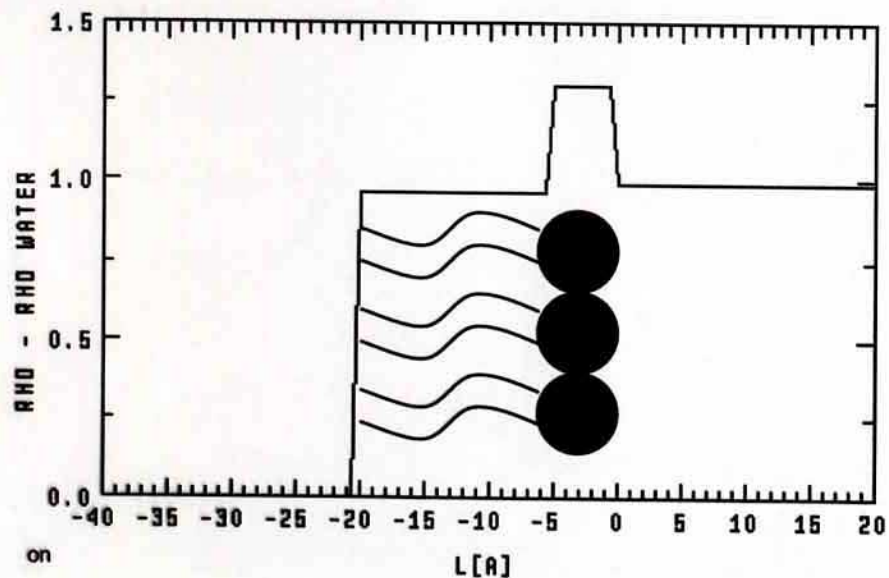
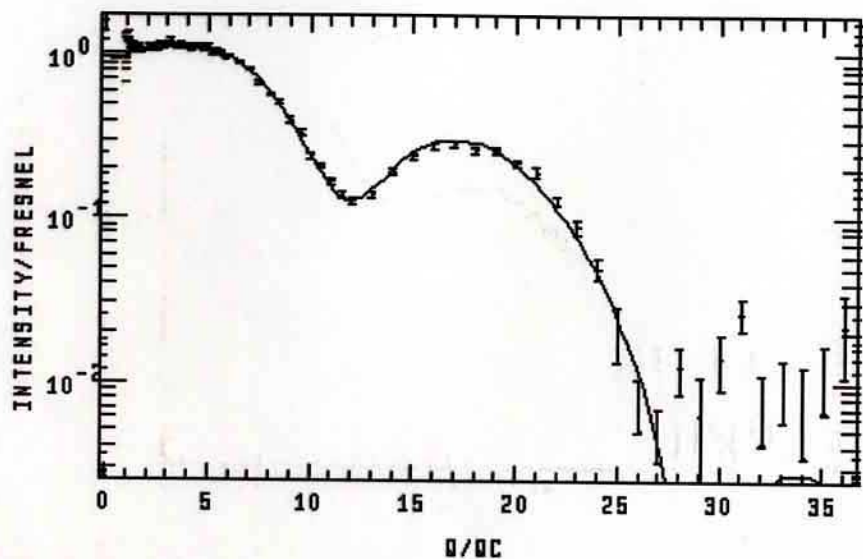


COMPARISON OF DIFFERENT LIPIDS

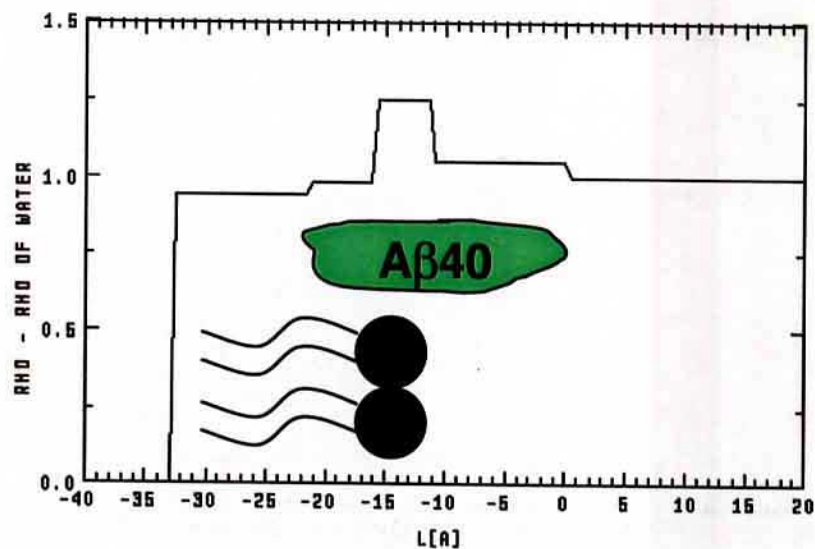
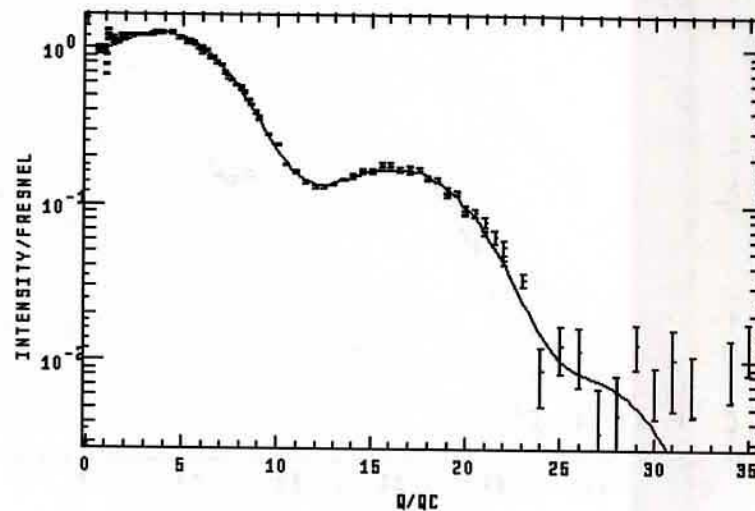


DPTAP-A β 40 interaction in WATER

DPTAP, $T=30$ C, $\Pi=30$ mN/m



DPTAP + A β 40, $T=30$ C, $\Pi=30$ mN/m

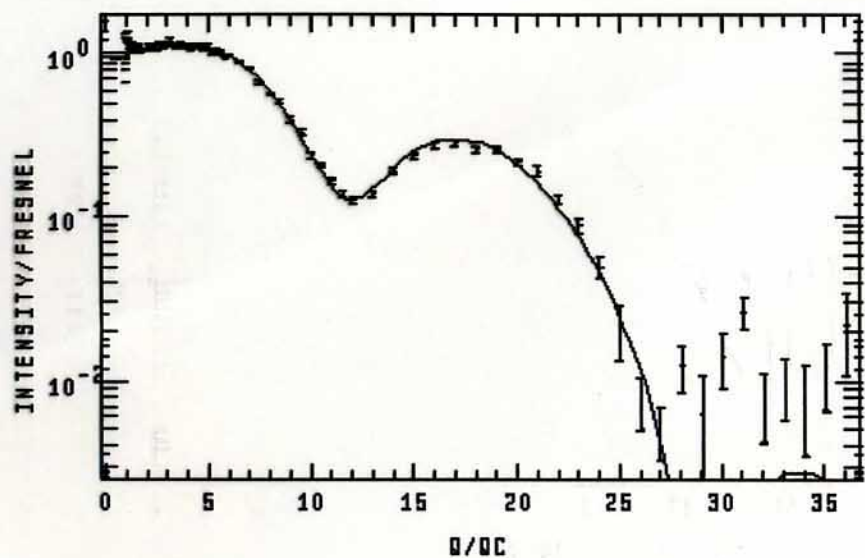


SURFACTANT-LIKE BEHAVIOR

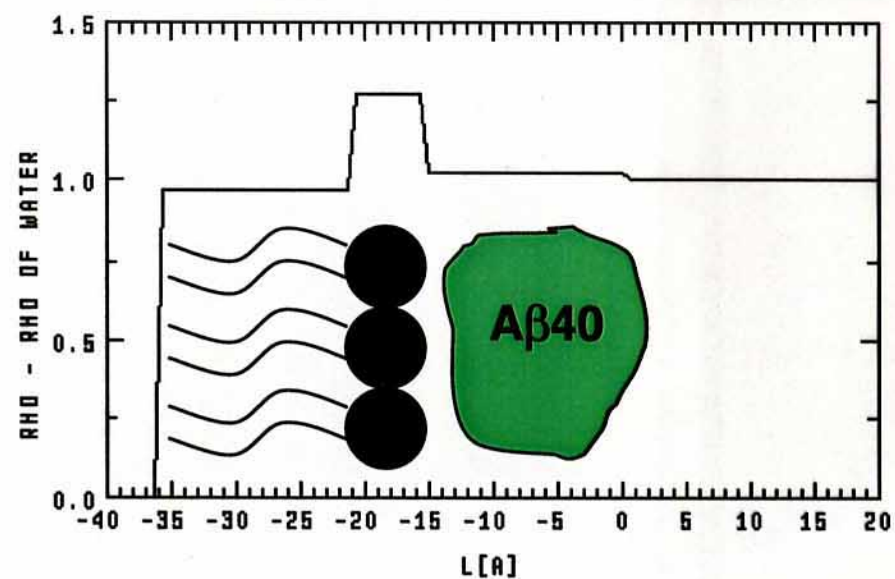
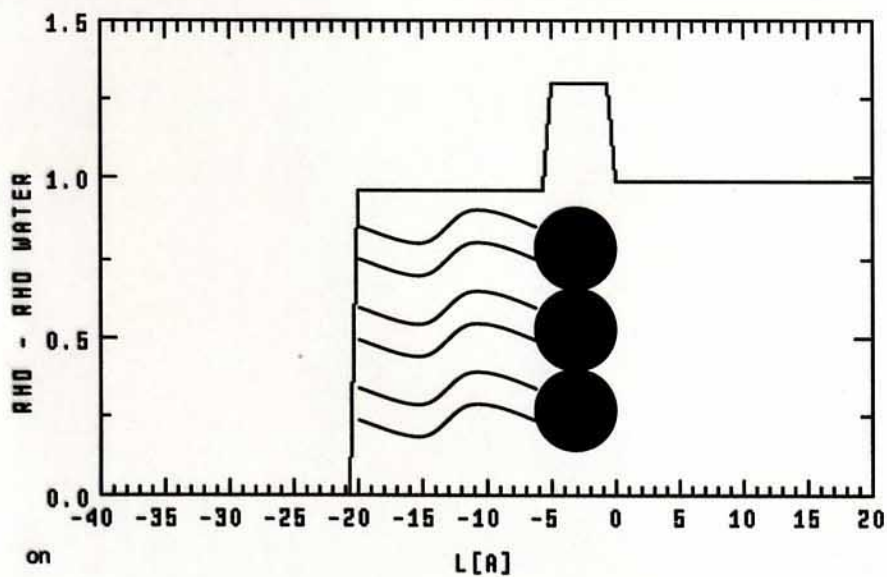
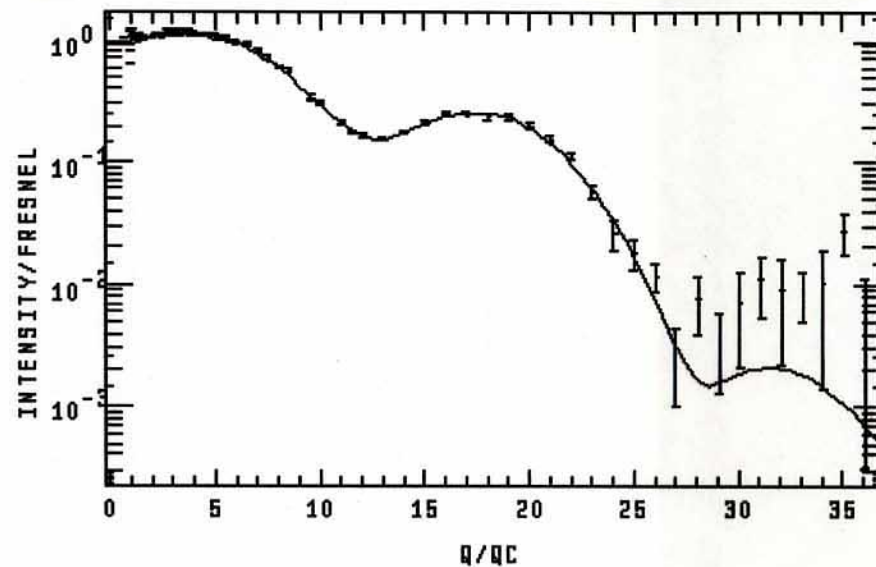
- N-terminus 1-28
 - > polar extracellular domain
 - C-terminus 29-(39 to 42)
 - > hydrophobic transmembrane region
 - Amphipathic in nature
 - Behaves like a surfactant
 - > lowers surface tension
 - > "critical micelle concentration" exists
- Soreghan *et al.*, *J. Biol. Chem.* (1994) **269**: 28551-28554
- 33 residues or shorter
 - > no surface response
 - > transmembrane portion is necessary

DPTAP-A β 40 interaction in PBS

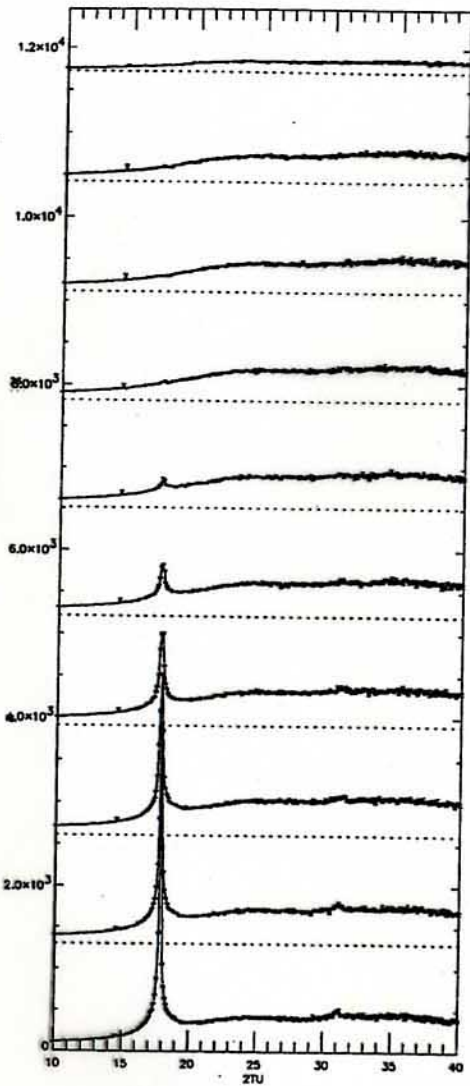
DPTAP, $T=30$ C, $\Pi=30$ mN/m



DPTAP + A β 40, $T=30$ C, $\Pi=30$ mN/m



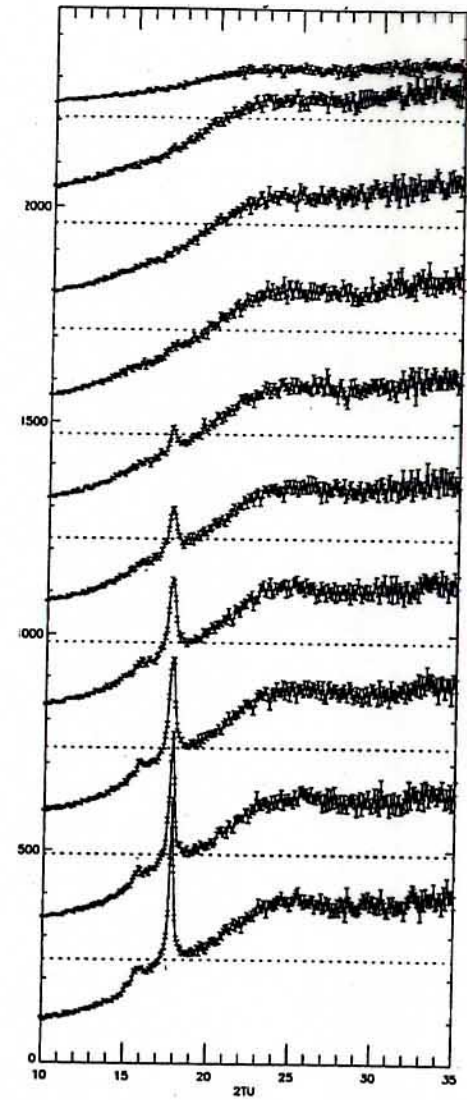
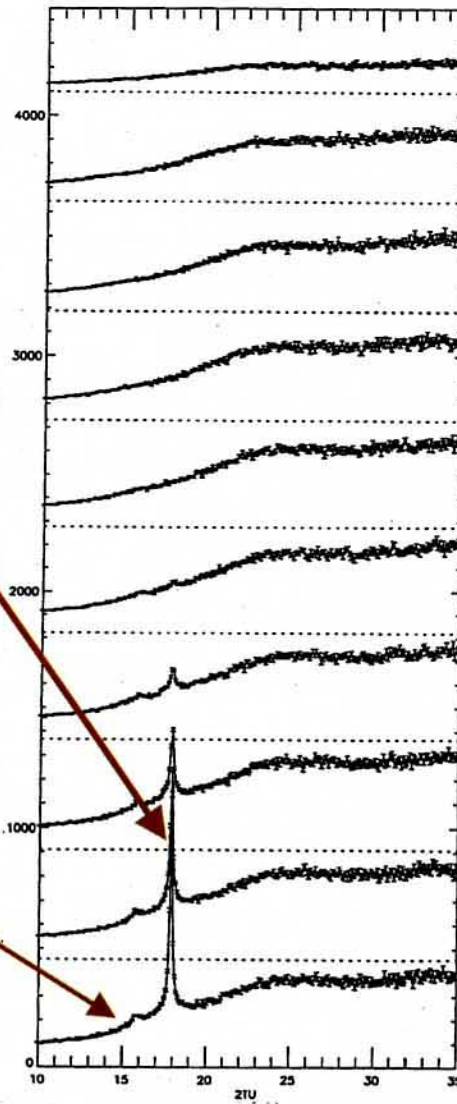
GIXD RESULTS



$T = 30^\circ\text{C}$
 $\Pi = 30\text{mN/m}$

intensity has
decreased

notice
2nd peak

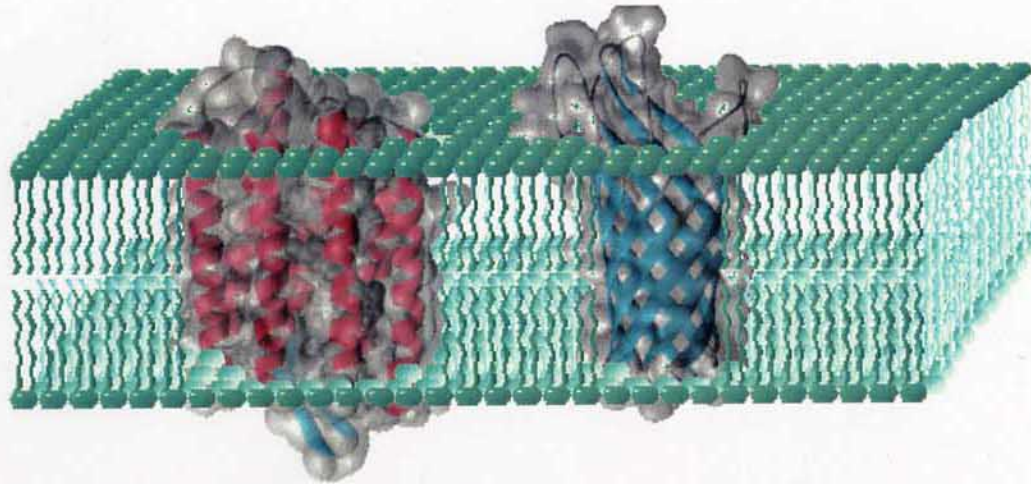


no Aβ40, DPPG only

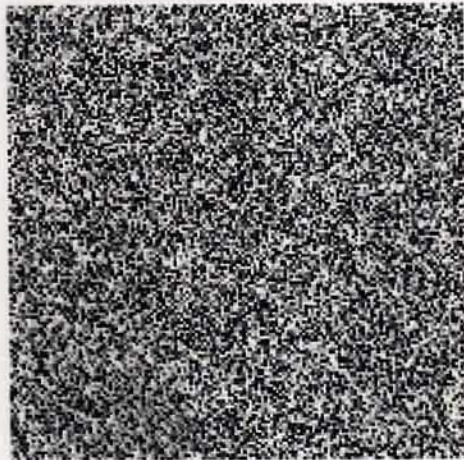
DPPG with Aβ40

DPPG with Aβ40, 2nd scan

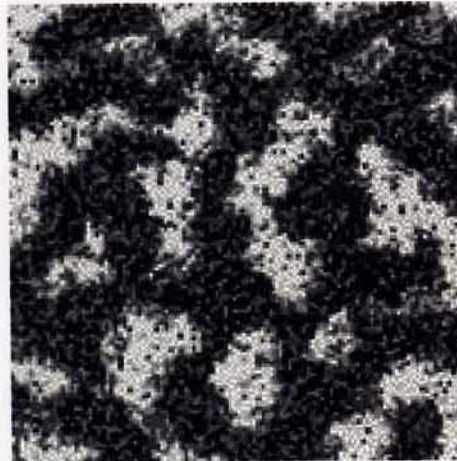
Is the lipid bilayer structured or not?



Fluid mosaic model of Nicolson and Singer
Science 1972, 175, 720; the bilayer is homogenous



homogenous bilayer



bilayer with lateral heterogeneity

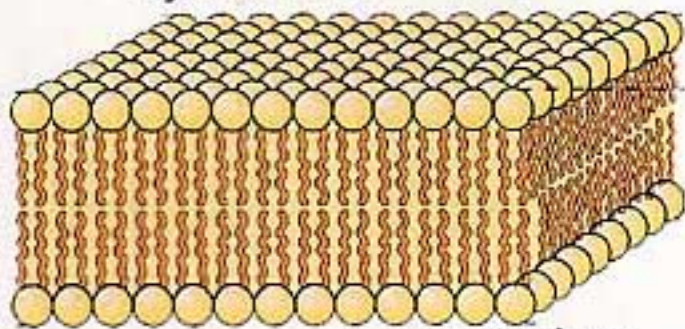
Lateral heterogeneity can be:

- Density fluctuations in a one component bilayer
- Compositional fluctuations in a two component bilayer
- Gel-fluid phase separation pattern

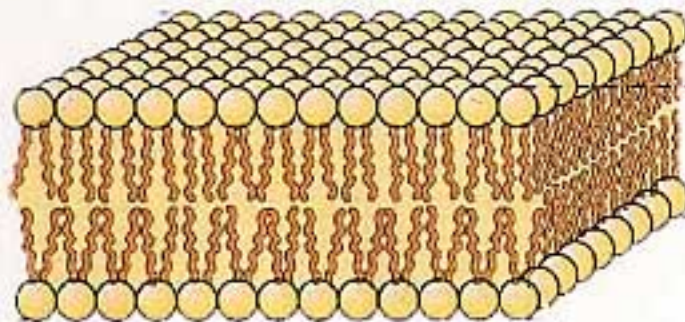
Model membrane composed of a binary mixture

gel phase--low temperatures

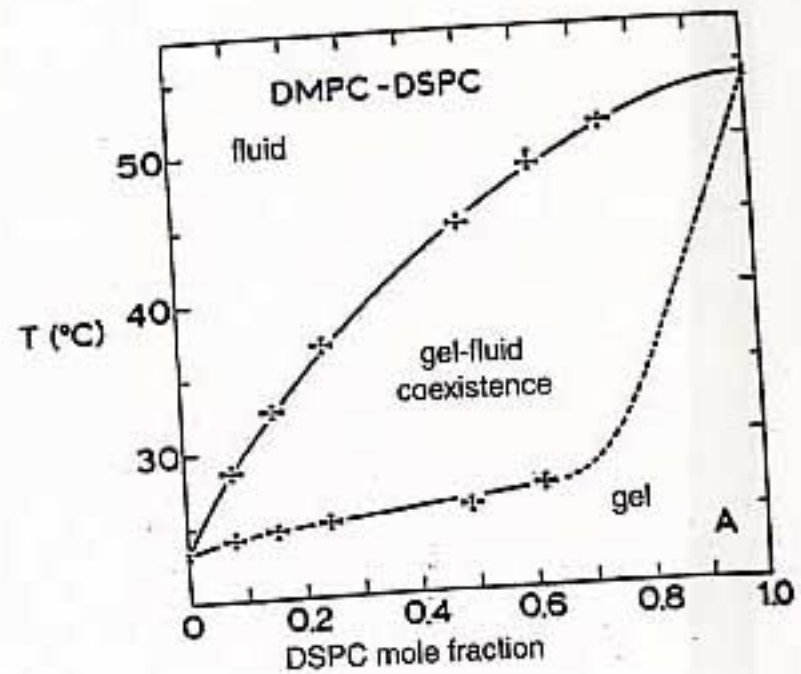
hydrocarbons are tightly packed



at higher temperatures—moves to fluid phase



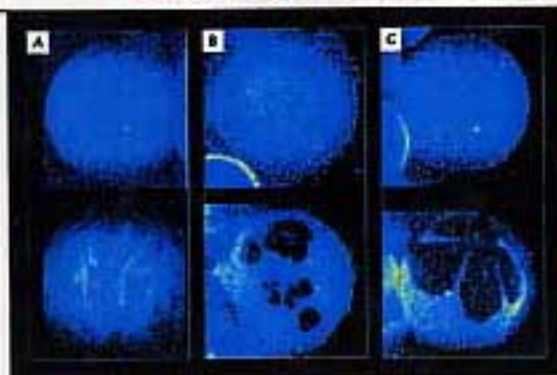
bilayer "melts", movement is allowed



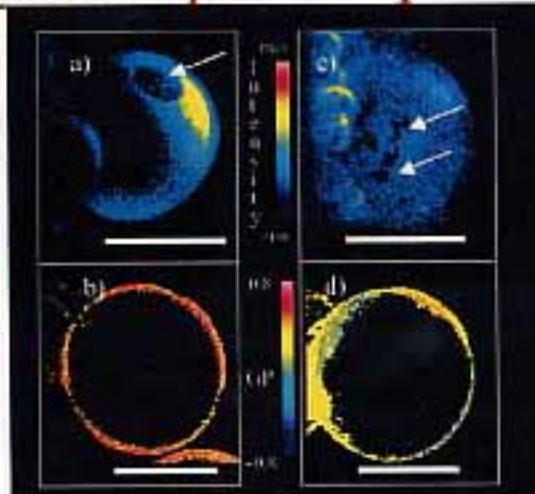
Shimshick, E. J.; McConnell, H. M.
Biochemistry 1973, 12, 2351-2360.

Phase diagram of a DSPC/DMPC mixed bilayer

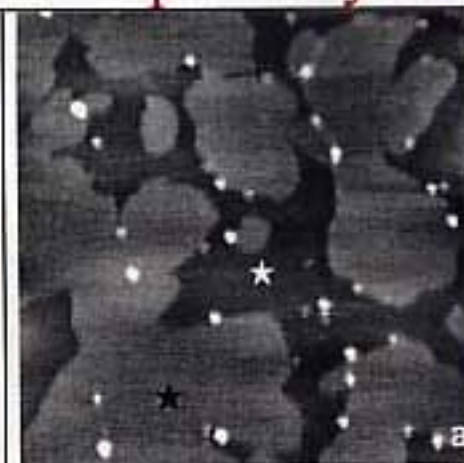
Previous work on lateral phase separation in lipid bilayers



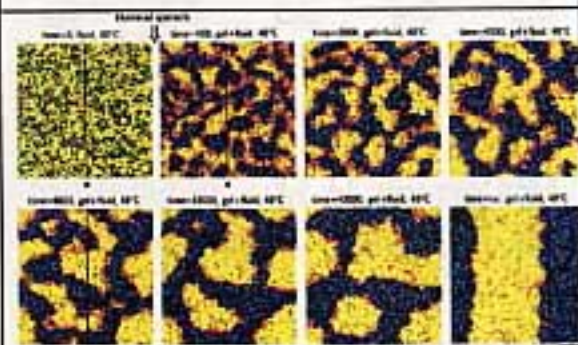
BPJ cover july 2000 DLPC/DPPC (A), DLPC/DSPC (B), DLPC/DAPC (C). Fluid phase top, fluid-gel coexistence bottom. **Fluorescence, Micron scale**



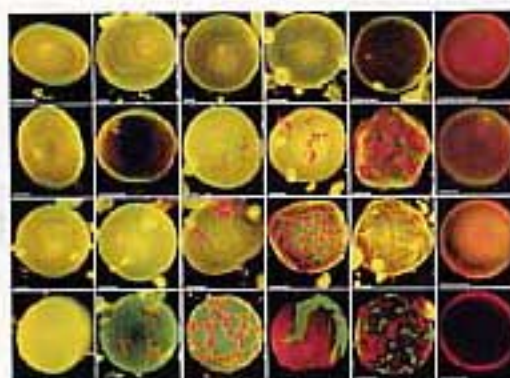
BPJ cover march 2001 Brush border membrane with (a&b) and w/o (c&d) cholesterol. **Fluorescence, Micron scale**



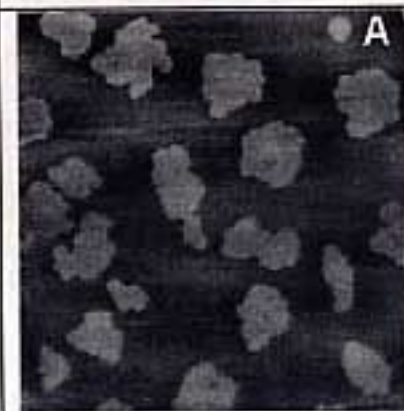
Langmuir 2000, 16, 1473 DPPC/DOPS deposited at room temperature, scan size 2.8 μm . **Atomic force microscopy (AFM)** Micron/Nanometer scale



JPC cover dec 14: 2000: Monte Carlo simulations of a DSPC/DMPC bilayer time evolution after quenching. Image size 55nm. **Simulations, Nanometer scale**



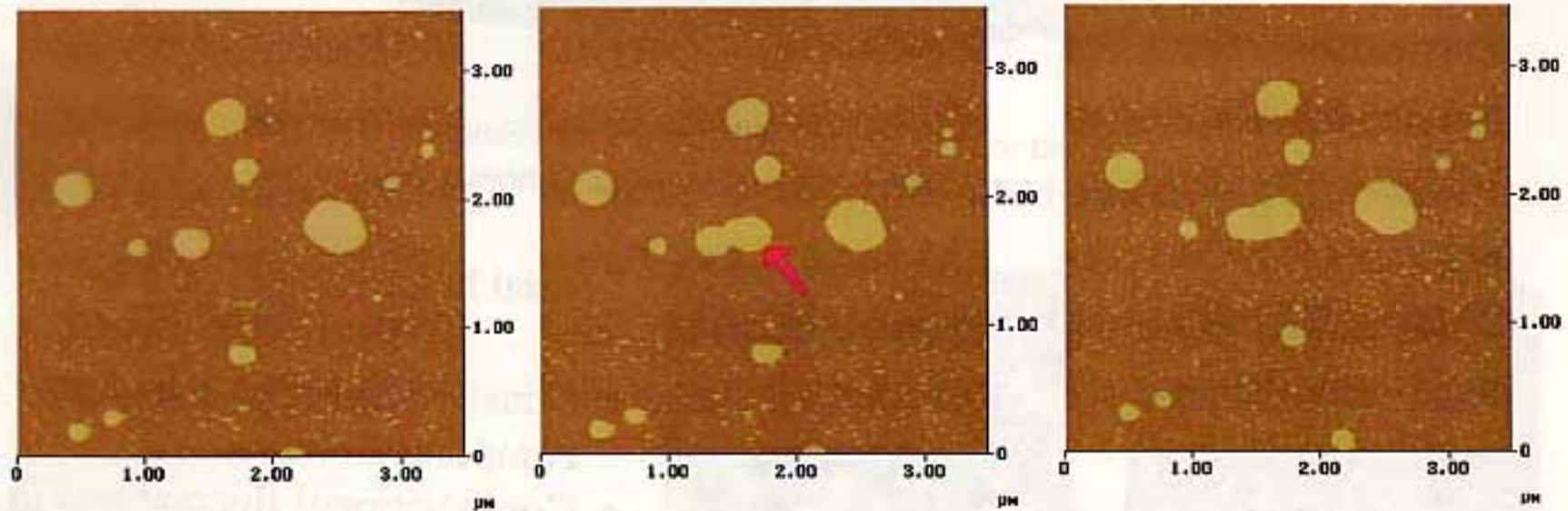
BPJ cover july 2001: DPPC/DLPC/Chol mixture in GUV. Horizontal: DPPC increases. Vertical: cholesterol increases. **Fluorescence, Micron scale**



Langmuir, 2001,17, 1653 DPPC/DOPC in PBS quenched to room temperature. Scan size 10 μm **AFM** Micron scale

Observation of Individual Vesicle Fusion

Egg-PC bilayer domains imaged in fluid using Tapping Mode. The concentration of the vesicle suspension is $.3\mu\text{g}/\text{ml}$ diluted in 30mM MgCl_2 .



time: 0 minutes

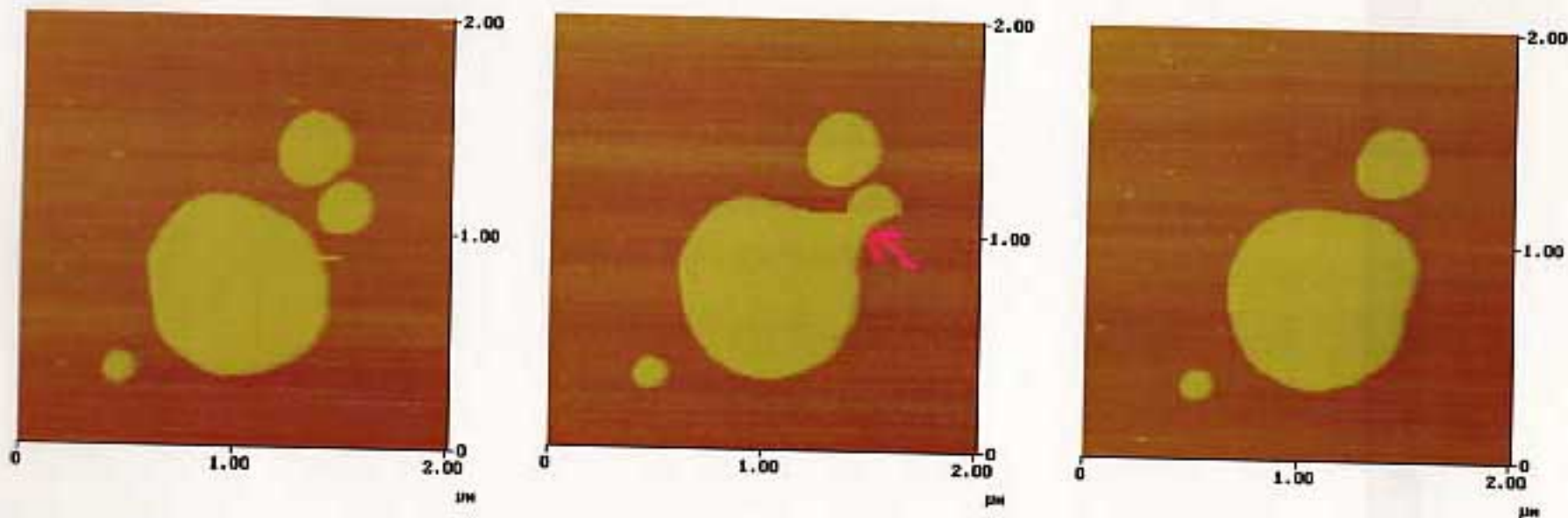
7 minutes

13 minutes

Individual vesicle fuses overlapping with an already adsorbed vesicle (red arrow). The resulting domain decreases perimeter.

Intermediated Fusion Between Bilayer Domains.

Egg-PC bilayer domains imaged in fluid using Tapping Mode. The concentration of the vesicle suspension is $.3\mu\text{g/ml}$ diluted in 30mM MgCl_2 .



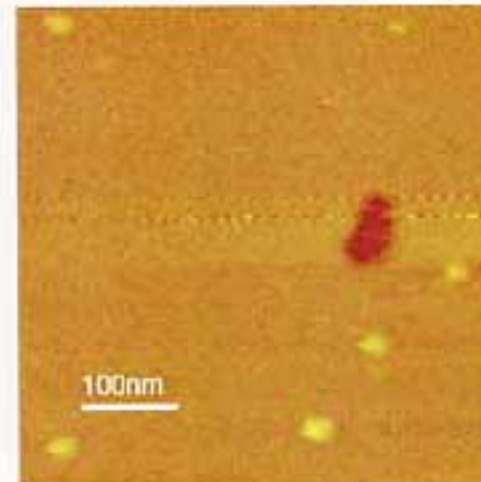
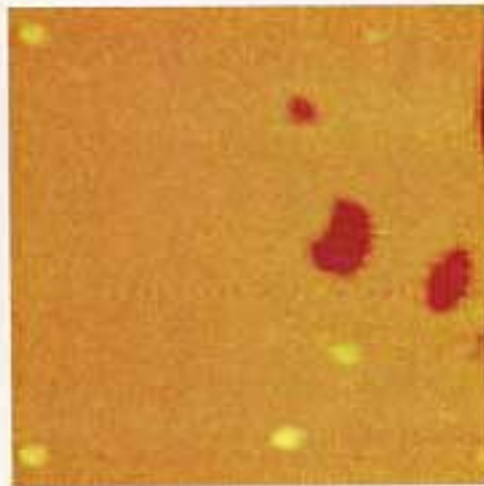
time: 0 minutes

4 minutes

9 minutes

Example of two initially separated bilayer domains that are fused together by a vesicle adsorbing in between (red arrow). The line tension transforms the final bilayer domain shape into a quasi-circular one.

... TO COMPLETE BILAYER COVERAGE

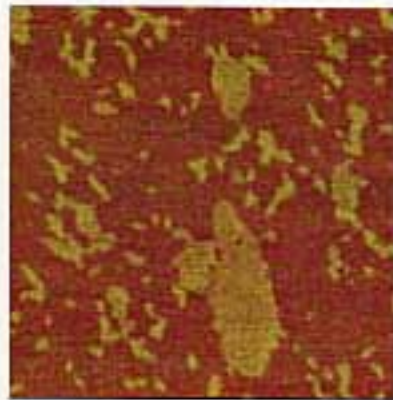


Healing of holes observed at the final stage
of the formation of a continuous lipid bilayer

Muresan and Lee, J. Phys. Chem. B 105 (2001) 852-855

Nanoscale Phase Separation in Supported Bilayers

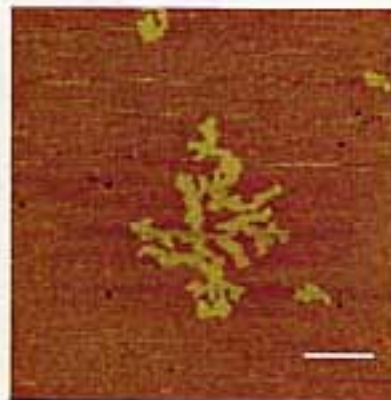
3:7 DSEPC:DMPC
results in
compact domains



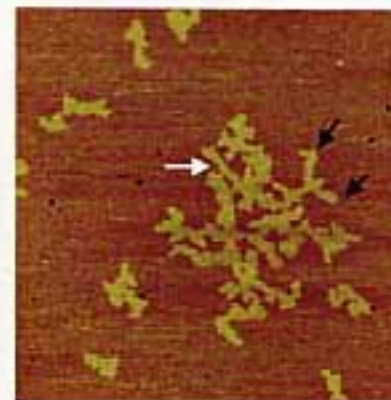
1:1 DSEPC:DMPC
results in
branched domains



Growth of aggregates
in 1:1 mixture



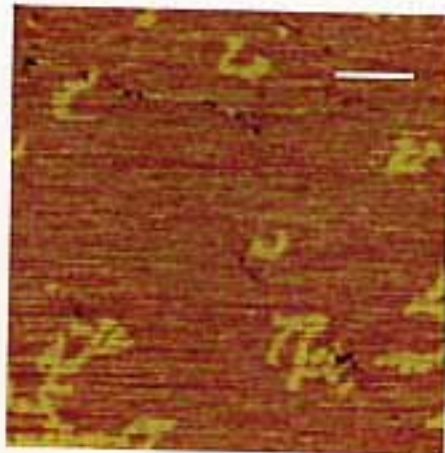
$t = 0$



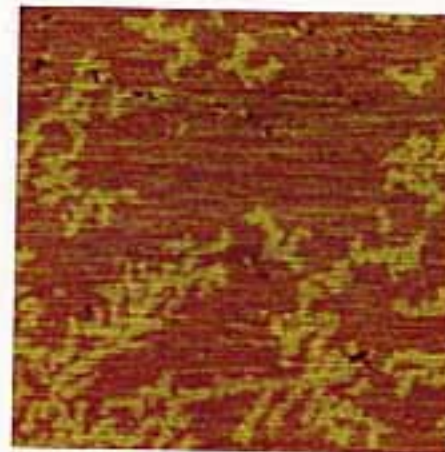
$t = 1$ hour

Formation of Fluid Nanoscale Compartments

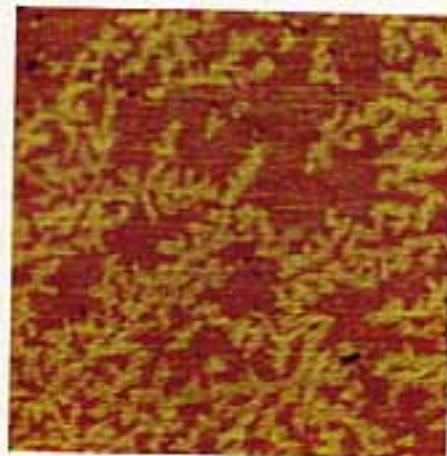
3:7 DSEPC:DMPC



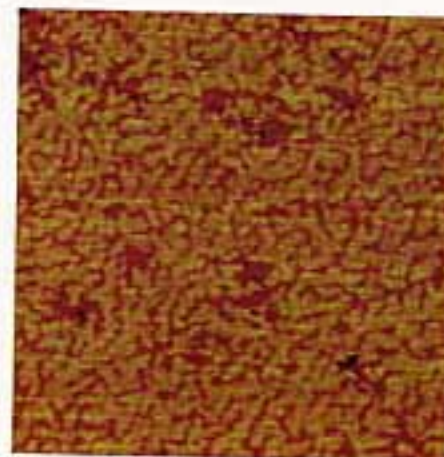
$t = 0$



$t = 33$ minutes



$t = 46$ minutes



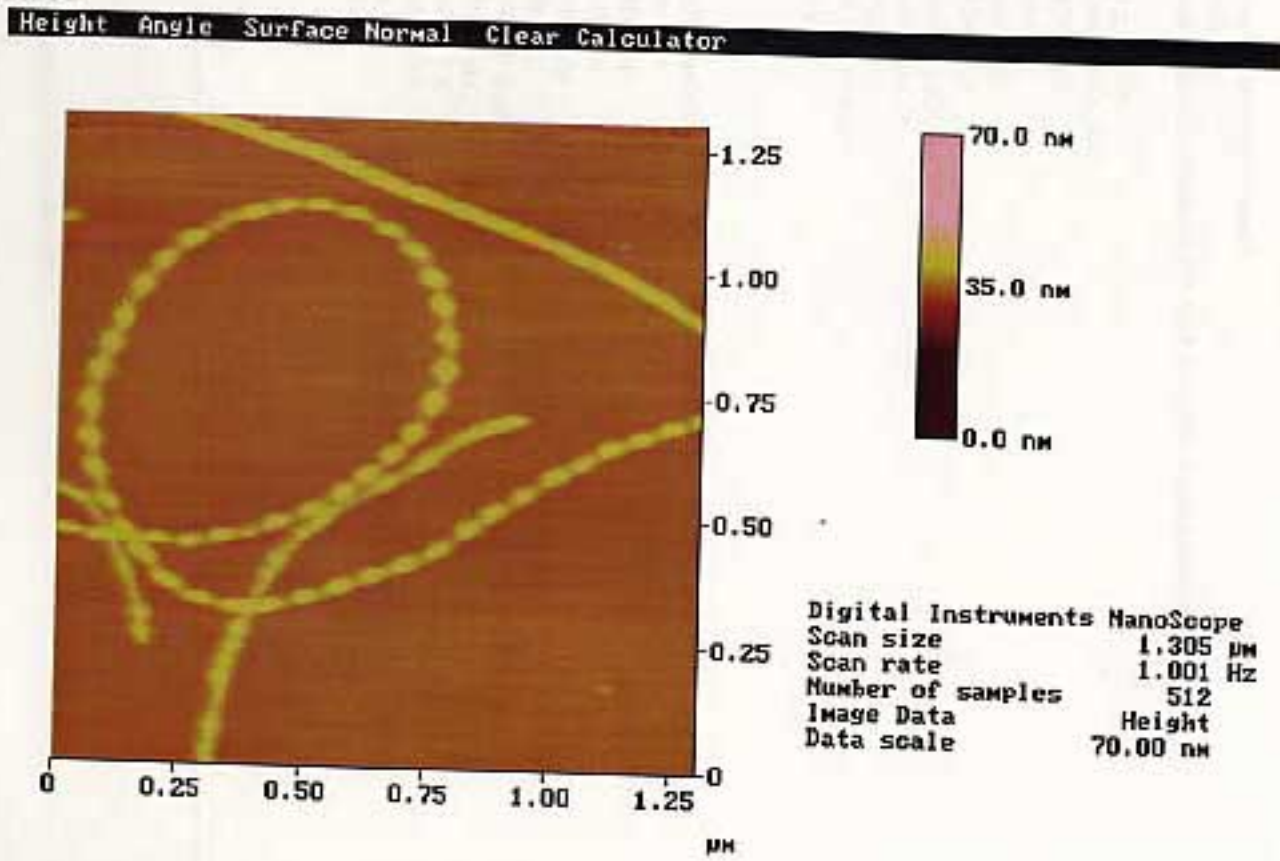
$t = 74$ minutes

A.S. Muresan, H. Diamant and K.Y.C. Lee, *JACS* 122 (2001) 6951-6952

Scale bar: 500nm

Results

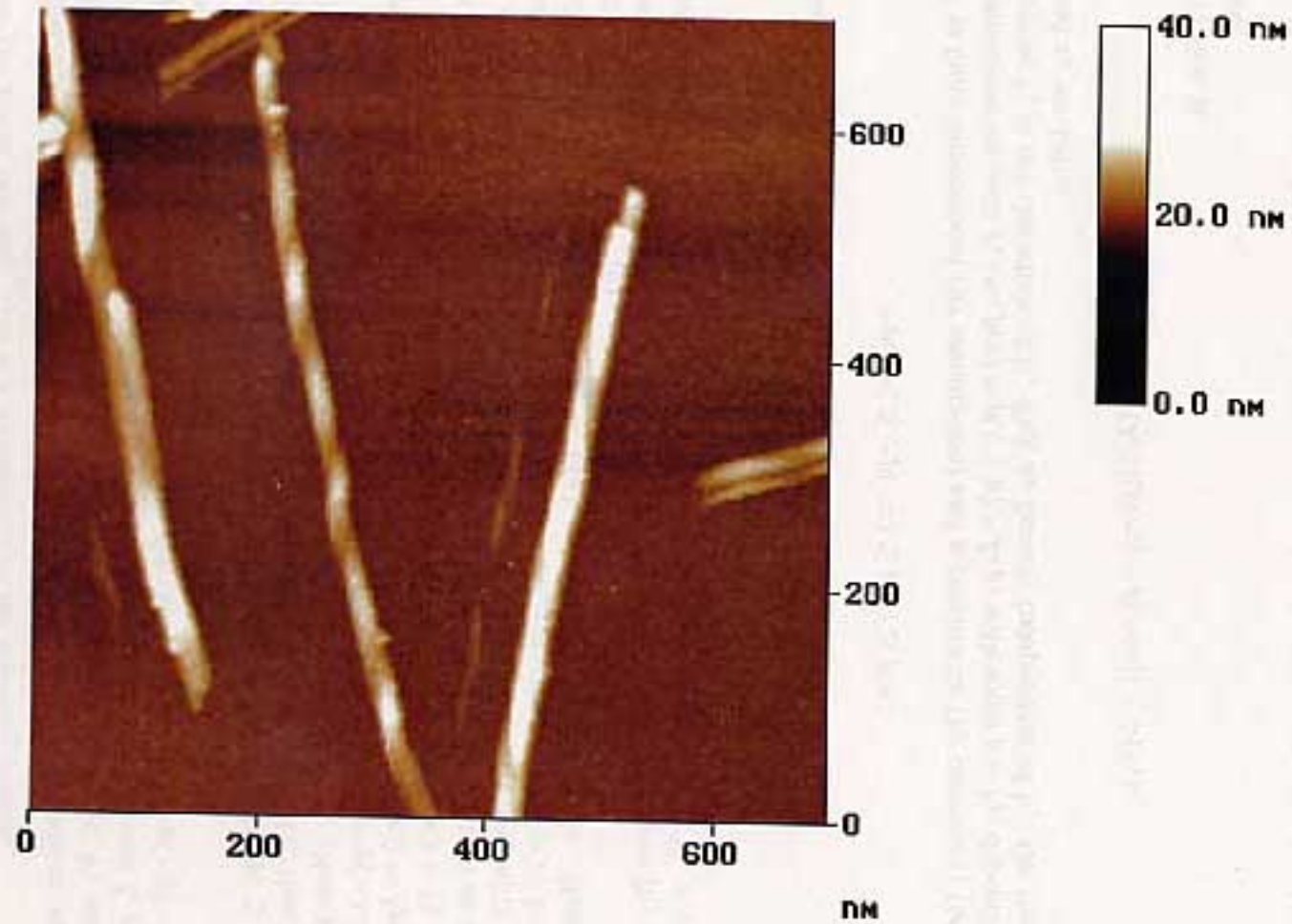
Tapping Mode in air shows bead-like structure for the A β fibrils while Tapping Mode in air resolves the helical structure of the fibrils.



07262158.003

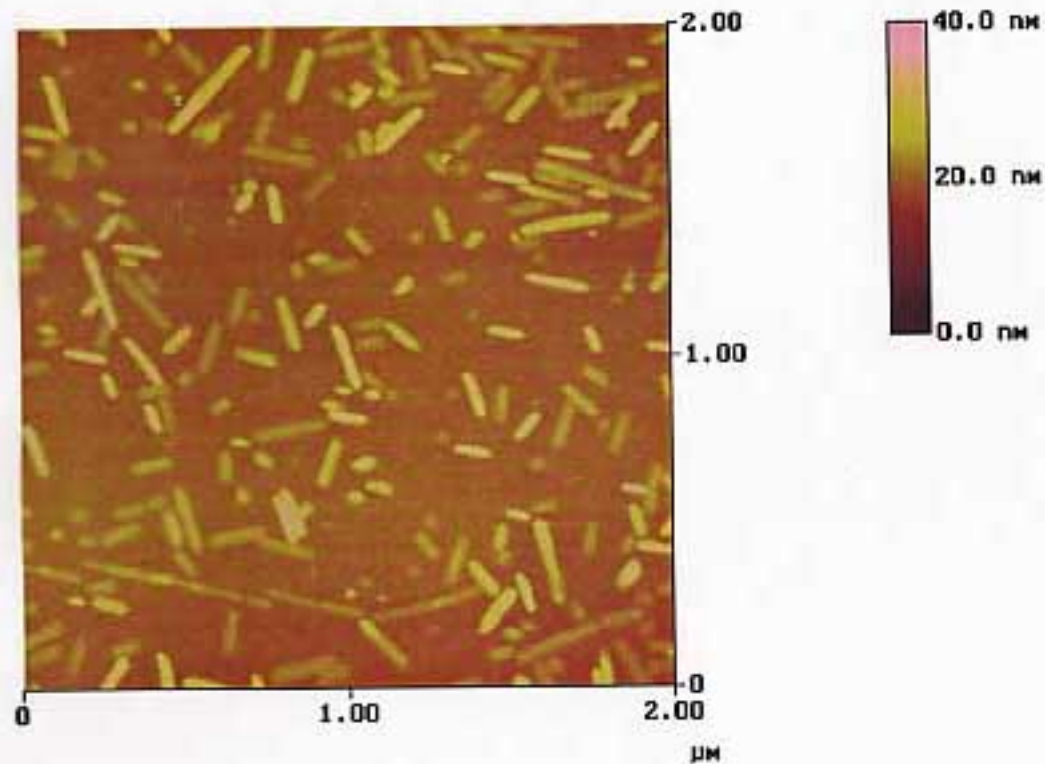
Height

Aged A β 1-40 imaged in air in TM.



Aβ10-35 fibrils imaged using Tapping Mode in fluid.

PEG does not inhibit fibril formation
but inhibits fibril-fibril association



A β -PEG fibrils imaged in solution on a graphite surface

Unlike A β 10-35 fibrils, these fibrils neither overlap nor entangle with one another

No axial periodicity is observed

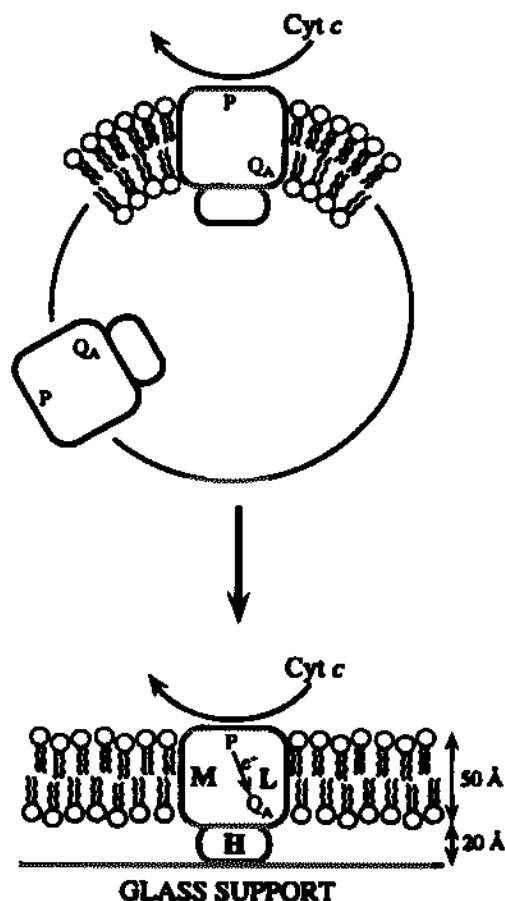


FIGURE 1: Schematic diagram of the bacterial photosynthetic reaction center in a bilayer membrane. The three protein subunits (L, M, and H) and the relevant functional components, the special pair primary electron donor, P, and the primary quinone acceptor, Q_A , are indicated. The RC is depicted in an orientation that is consistent with the results presented and a mechanism in which the vesicles fuse to the glass support by opening out.

membranes (Sui et al., 1988; Plant et al., 1995; Sackmann, 1996), though most of this work does not involve membranes which contain protein.

The RC is the smallest isolatable unit which performs the initial photoinduced charge separation steps in photosynthesis. The three-dimensional structures of RCs from two species of photosynthetic bacteria are known to atomic resolution (Deisenhofer et al., 1995). It is comprised of three subunits, termed L, M, and H, whose total molecular mass is approximately 100 kDa (see Figure 1). Its physiological role is to transduce light energy into a charge separation across the native membrane, which in turn leads to a transmembrane electrochemical potential used by the organism to store energy. The RC accomplishes this charge separation by transferring an electron from the excited state

EXPERIMENTAL PROCEDURE

Materials. The dye-lipid conjugate, 1,2-dihexadecanoyl-*sn*-glycerol-3-ethylammonium salt (Texas R-1, 1,3-diazol-4-yl)-1,2-dihexadecanoylphoethanolamine, triethylamine, and dithiothreitol were purchased from Molecular Probes. Dithiothreitol was also purchased from Molecular Probes. Biotin, 4-hydroxyazobenzene, and avidin (Immunopure avidin) were purchased from Molecular Probes. Chemical Co. Horse heart cytochrome c was purchased from Molecular Probes Chemical Co.

Reaction Centers. RCs were isolated from *Rhodospirillum rubrum* wild-type or the mutant *R. rubrum* (M189) by the methods that are described in Taguchi et al. (1986). The RCs isolated by this method contain only the primary quinone, Q_A , and five cysteine residues at positions H156, H234, and H234. From an examination of the cysteine residue at H156 and the other four are buried from the surface). *Rhodospirillum rubrum* RCs isolated as described in Taguchi et al. (1986) have cysteine residues at positions H156 and L247, all of which are buried from the surface. The same three-dimensional structure is shared by analogous *Rb. sphaeroides* RCs (Krogh et al., 1992). The surface cysteine residue was isolated from *Rb. capsulatus* RC by replacing the cysteine at position M189 to create the mutant *R. rubrum* (M189) in detail in Boxer et al. (1992).

RC-Biotin and RC-Rhodamine Conjugates. The surface cysteine of the *R. rubrum* RCs was modified using reagents and procedures described in Boxer et al. (1992). This was used to modify the RCs with rhodamine conjugates at the cysteine H156 (Debus et al., 1986). The RCs were dialyzed into 10 mM Tris, pH 8.0, and 0.025% BSA to about 75 μ M using a Centricon 30 (Amicon Co.). The RCs were then dialyzed into 10 mM Tris, pH 9.0, and 0.025% LDAO buffer. The changes of buffer containing 0.025% LDAO and a 10-fold molar excess of the dye-lipid conjugate were used to modify the RCs.

Unilamellar Vesicles

- Freeze-thaw extrusion
- lipid (dissolved in CHCl_3) dried under N_2 stream forming a thin film
- dried overnight under vacuum
- rehydrate & vortex
- 5-10 freeze thaw cycles
- extrusion using desired pore size
or
probe sonicate for ~ 50 nm diam. vesicles.

Proteoliposomes

- protein added to vesicles
at certain desired ratio
- run down Sepharose column
previously equilibrated with lipid.

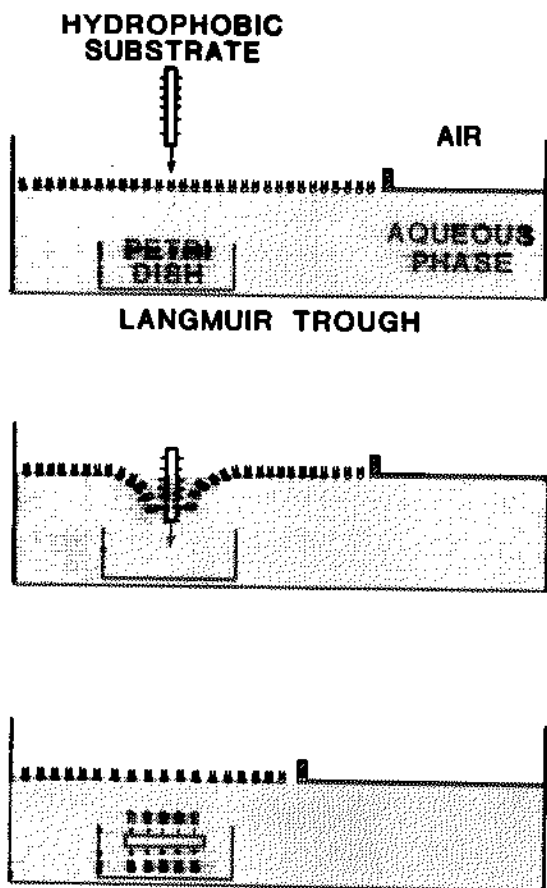


Fig. 1. Transfer of a phospholipid monolayer from the air-water interface to a solid support. The support, which has been alkylated to render it hydrophobic, is lowered vertically through the phospholipid monolayer at the air-water interface in a Langmuir trough. Supported monolayers can also be prepared by lowering the substrate horizontally through the air-water interface (for example, see Ref. 24).

hydrophilic solid supports; again, these can be glass, quartz or single crystal silicon. Fig. 1 shows the transfer of phospholipid monolayers onto an alkylated support. The alkylated support is lowered through the monolayer at the air-water interface in a Langmuir-Blodgett trough. Sometimes the support is kept in a vertical orientation during immersion (as in Fig. 1) and sometimes a 'face-down' (horizontal) immersion is used.

There are at least two methods of preparing supported bilayers. In Fig. 2, a supported bilayer

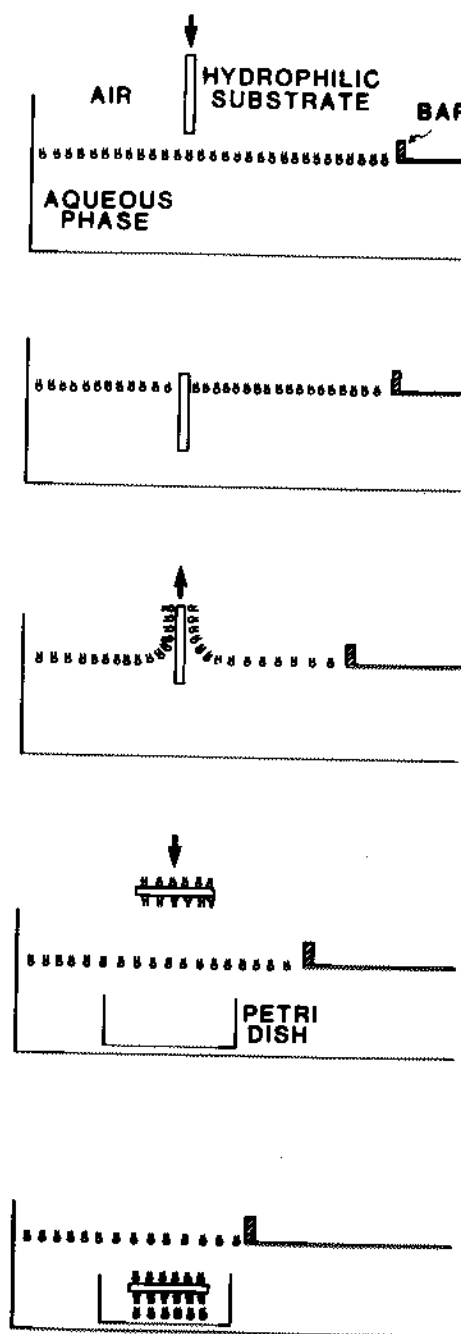


Fig. 2. Preparation of supported bilayers. Two monolayers are spread at the air-water interface in a Langmuir-Blodgett trough. The clean hydrophilic silicon substrate is immersed vertically through the monolayer. The significant change in surface pressure is observed. The substrate is then pulled out to permit water to drain from the surface. The surface pressure observed is then constant over the area of the trough. The loss in

Formation of Supported Bilayers

① Monolayer transfer by Langmuir-Blodgett techniques

- automatic dep. process
- process can be followed by ellipsometry or microfluorescence
- both layers can be manipulated

② Vesicle Rupture

- reconstitute with protein in native orientation

③ Monolayer + Vesicle Rupture

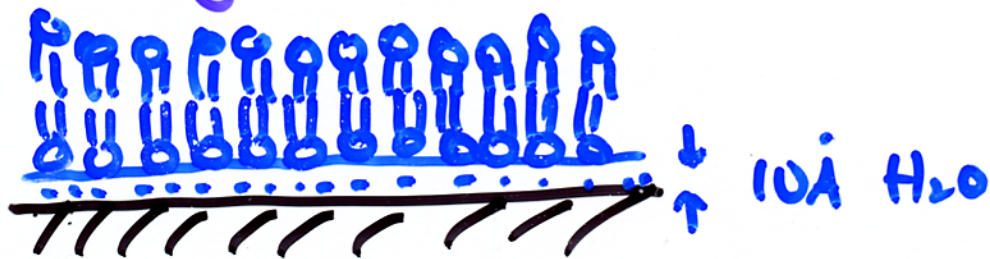
- Allow the presentation of correct side as an outer leaflet
- maintain mobility of lower leaflet

Self-Assembly of Supported bilayers

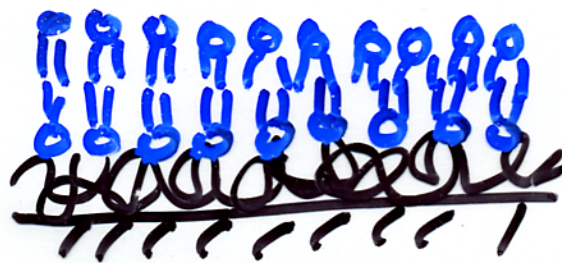
- ① Integrated bilayers w/ lower leaflet fixed to substrate either covalently or by ion bridges



- ② Supported bilayer with 2 free leaflets



- ③ Bilayer resting on thin, soft, hydrated polymer film



To immobilize monopolar proteins

formed with ultrathin polymer films hydrophobized by coupling of long alkyl chains to hydrophilic polymer backbone

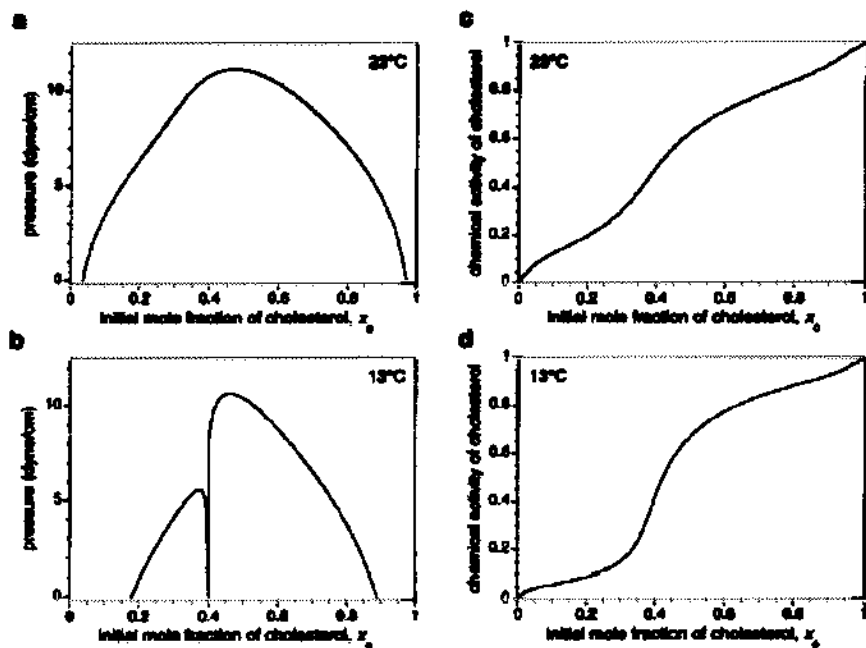


Figure 4. Theoretical phase diagrams and chemical activities. Panels a and b show calculated pressure (x_0)-composition (x_0) phase diagrams for reactive mixtures containing cholesterol and phospholipid at 23 and 13 °C, respectively. Compositions of coexisting phases were found by the method of double tangent construction using the free energy of eq 2 and the parameters listed in the text. Panels c and d show calculated chemical activity-composition curves for reactive mixtures of cholesterol and phospholipid at a pressure of 20 MPa and at temperatures of 23 and 13 °C, respectively for the same parameters. The normalized equilibrium constant K_{eq}^0 is 0.3, and the cooperativity n is 2.

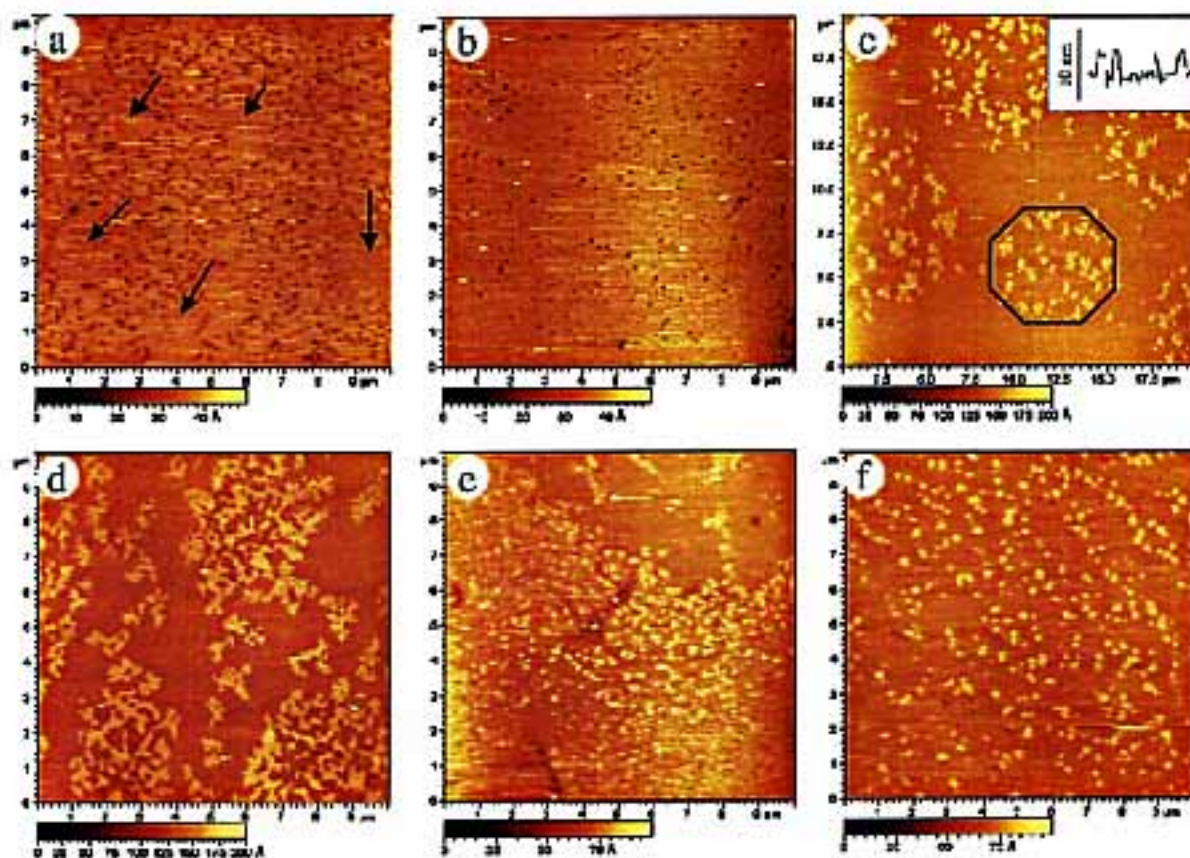


FIGURE 5 AFM images for hybrid bilayers of SPM/DOPC/cholesterol (1:1:1) with and without 5% GM1. (a and b) SPM/DOPC/cholesterol (1:1:1) bilayers deposited at 10 mN/m (a) and 35 mN/m (b). The arrows indicate more tightly packed areas of the bilayer. (c) SPM/DOPC/cholesterol (1:1:1) bilayer with 1% GM1 deposited at 10 mN/m with one of the large domains containing small GM1-rich microdomains outlined in black; the x scale for the action analysis shown in the inset is the same as the image. (d-f) SPM/DOPC/cholesterol (1:1:1) bilayers with 5% GM1 deposited at 10 mN/m (d), 30 mN/m (e), and 35 mN/m (f).

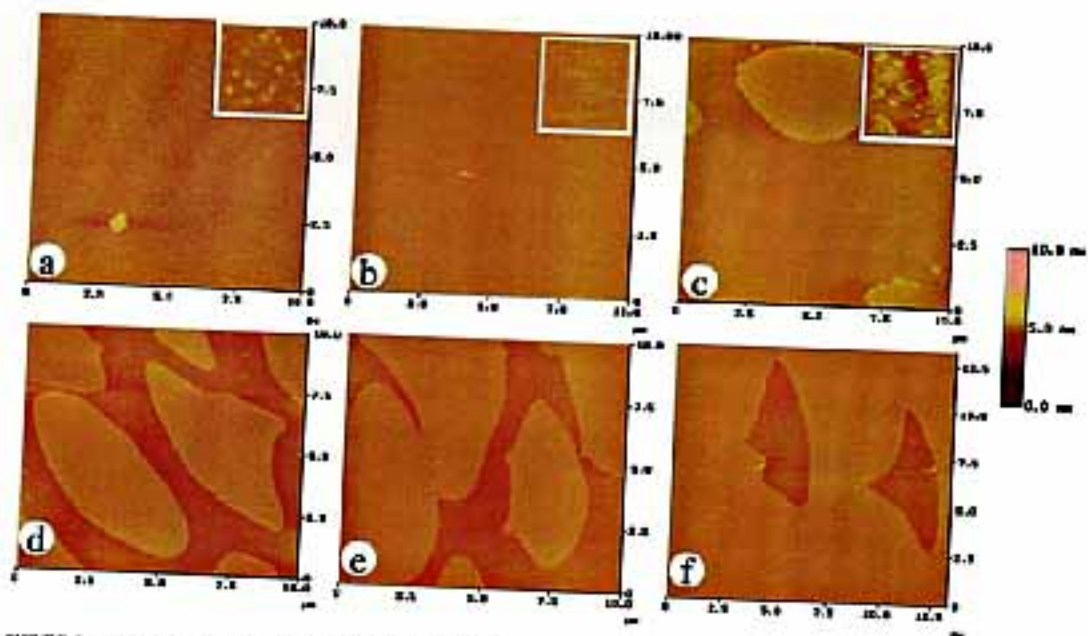


FIGURE 3 AFM images for SPM/DOPC and SPM/DOPC/cholesterol monolayers. (a) SPM/DOPC (1:1) monolayer deposited at 7 mN/m. The inset is a small-scale image ($500 \times 500 \text{ nm}^2$). (b) SPM/DOPC (1:1) monolayer with 10% cholesterol transferred at 7 mN/m. The inset is a small-scale image ($300 \times 500 \text{ nm}^2$). (c) SPM/DOPC (1:1) monolayer with 20% cholesterol at 7 mN/m. The inset is a large-scale image ($40 \times 40 \mu\text{m}^2$). (d) SPM/DOPC (1:1) monolayer with 33% cholesterol at 7 mN/m. (e) SPM/DOPC (1:1) monolayer with 33% cholesterol at 10 mN/m. (f) SPM/DOPC (1:1) monolayer with 33% cholesterol at 30 mN/m. The z scale is 10 nm for all images (see color bar) except image a, which is 5 nm.

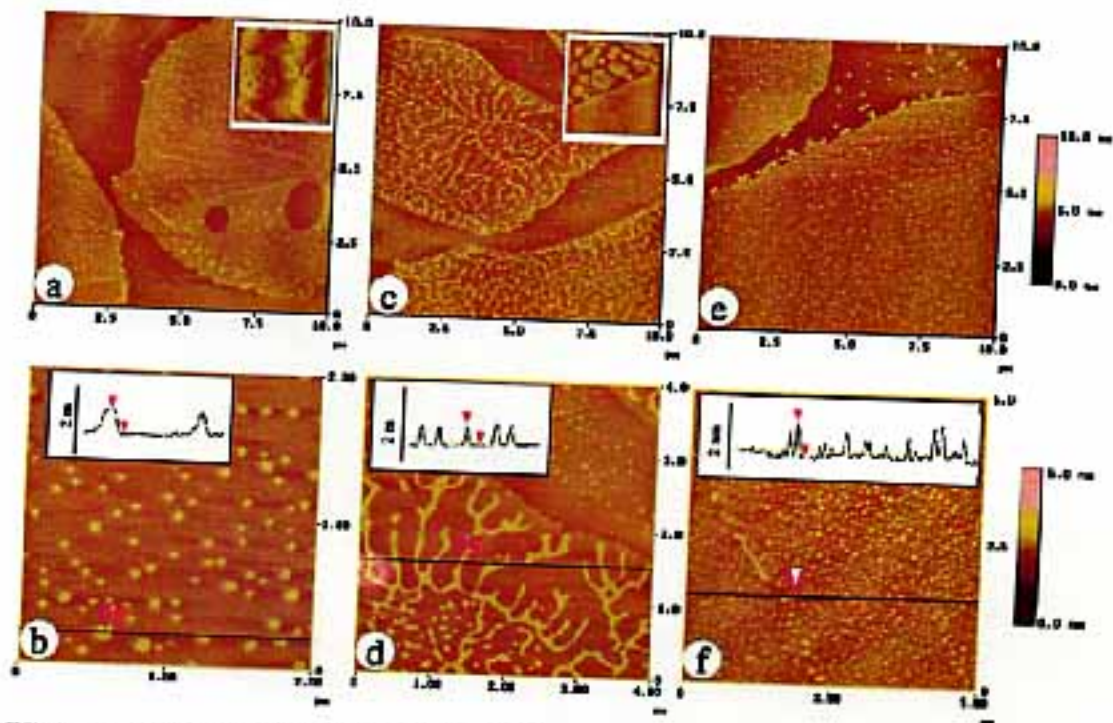


FIGURE 4 AFM images for monolayers of SPM/DOPC/cholesterol (1:1:1) with 1% GMI (a, b at 10 mN/m, and c at 30 mN/m) and 5% GMI (d, e, f at 40 mN/m). The insets in a and c are large-scale images ($40 \times 40 \mu\text{m}^2$) for the same sample. The z scale (see color bar) is 5 nm for b, d, and f and 10 nm for a, c, and e. Section analyses (for the solid line on the images) are shown as insets in b, d, and f; the x scale for the inset is the same as that for the image.

The Size of Lipid Rafts: An Atomic Force Microscopy Study of Ganglioside GM1 Domains in Sphingomyelin/DOPC/Cholesterol Membranes

Chunbo Yuan, Jennifer Furlong, Pierre Burgos, and Linda J. Johnston

Seascale Institute for Molecular Sciences, National Research Council of Canada, Ottawa, Ontario K1A 0R6, Canada

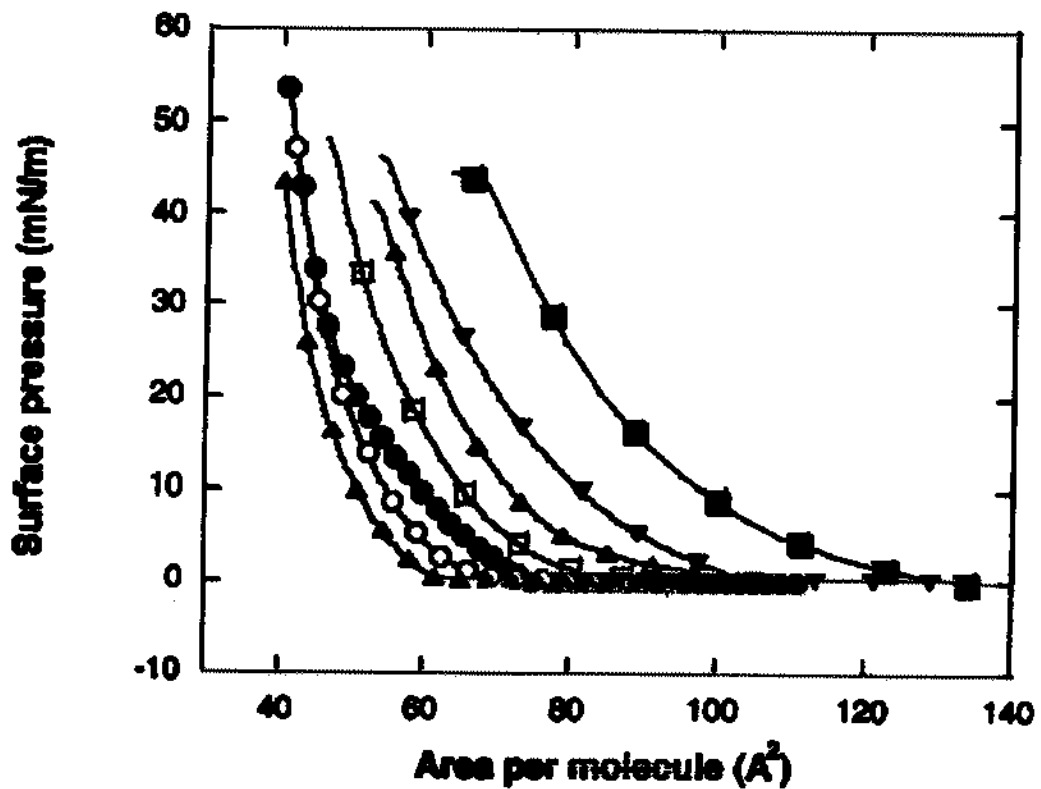


FIGURE 1 Surface pressure-area isotherms for SPM (●), DOPC (■), SPM/DOPC (1:1) (▼), SPM/DOPC (1:1) mixtures with 10% (▲), 20% (□), and 33% (○) cholesterol, and a SPM/DOPC/cholesterol (1:1:1) mixture with 5% GM1 (▲).

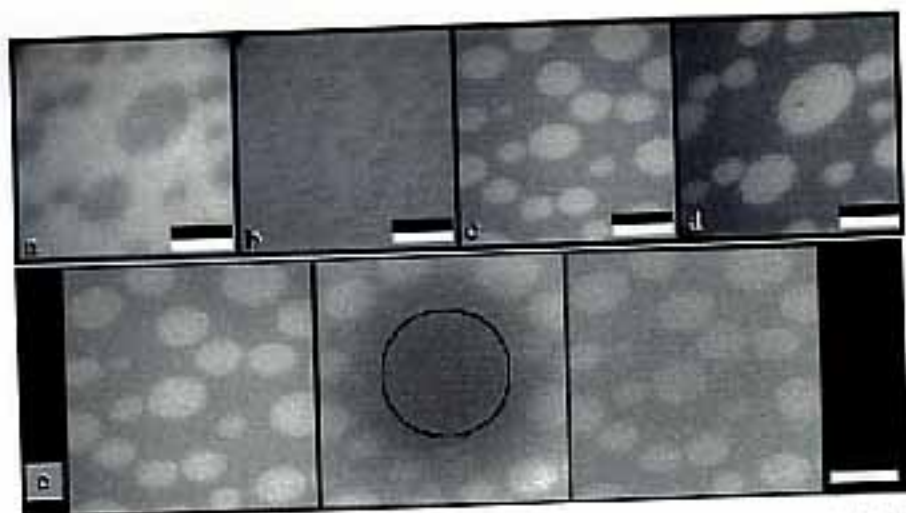


FIGURE 4 Fluorescence micrographs of planar supported phospholipid bilayers made from POPC/cholesterol/sphingomyelin (2:1:1) with 1 mol % GM1 (distal to glass support) and egg PC (proximal to glass support). Only distal lipid layers contain fluorescent lipid dyes: (a) 0.1 mol % TR-DPPe; (b) 0.5 mol % FL-DPPe; (c) 0.5 mol % NBD-DPPe; (d) FL-CTB. Note that images a and d were obtained from the same bilayer preparation, which was double labeled with TR-DPPe and FL-CTB (for 10 min at 1 μ g/ml in PBS), then, 10 μ m. (e) Fluorescence from NBD-DPPe equilibrates between liquid-ordered and liquid-disordered regions (left) before photobleaching; (middle) after 4 min of continuous photobleaching; (right) after 4 min of recovery. The circle in the central panel indicates the size of the field stop during the bleaching period. To achieve a significant bleaching rate, neutral density filters in the illumination path of the HBO100 were removed during bleaching. Size, 10 μ m. All images were recorded at room temperature (24°C).

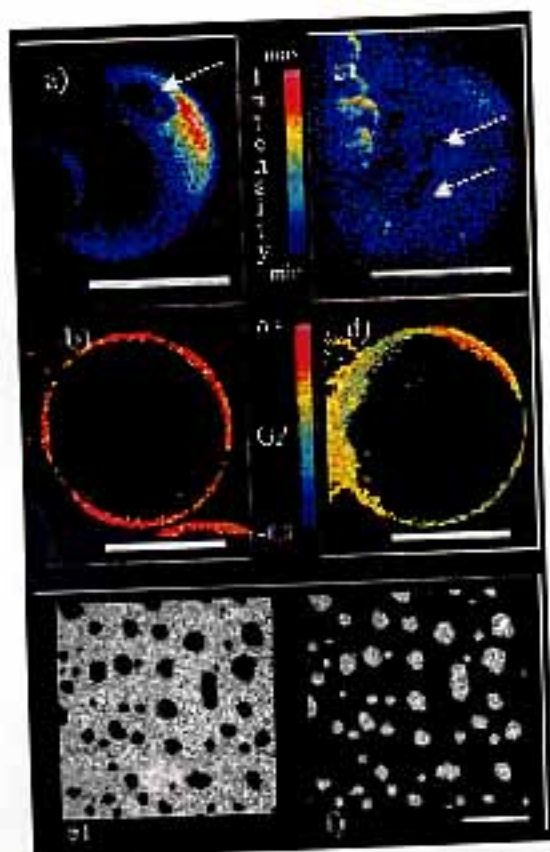


FIGURE 7 Two-photon fluorescence images of polar sections (a and c) and GP images of septal sections (b and d) of LAURDAN-labeled GUV formed from extracted brain border membrane lipids with cholesterol present (a and b) at 28°C or with cholesterol extracted (c and d) at 15°C [see Materials and Methods]. Size, 50 μ m. (e and f) Planar supported BHM monolayer with 0.2 mol % TR-DPPe and 1 mol % GM1 after incubation with FL-CTB (1 μ g/ml in PBS for 15 min) as observed in fluorescence channel (e) and fluorescence channel (f). Bar for planar supported lipid layer is 10 μ m.

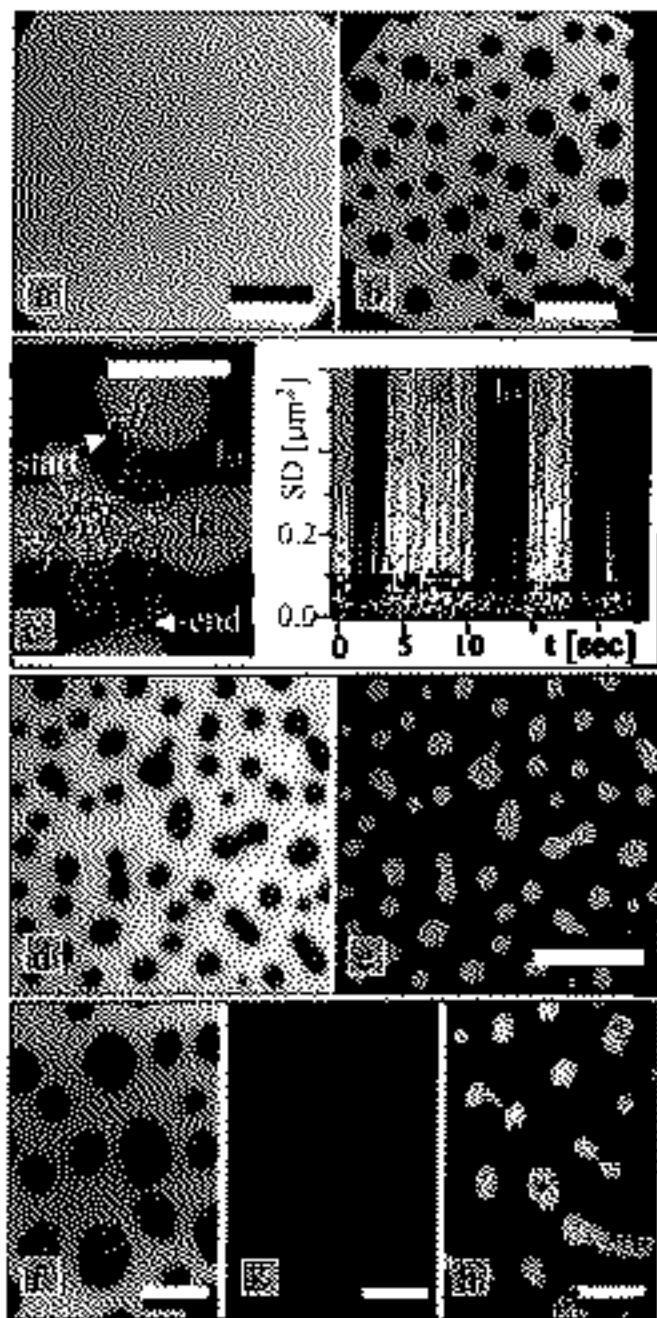


FIGURE 3 Fluorescence micrographs of phospholipid membranes supported by silanized glass substrates at room temperature (see Materials and Methods). (a) Non-raft domain (POPC/cholesterol 2:1); (b) Raft domain (DOPC/cholesterol/phosphatidylcholine 2:1:1). Both contain 0.5 mol % PL-DPPC. (c, left panel) DOPC/cholesterol/phosphatidylcholine 1:1:1) containing 0.5 mol % PL-DPPC and 1 mol % GM1 with ovalbium trajectory of gold particle (50–60 nm) as visualized by SPT (Gustav and Lindman, 1997). GM1 was functionalized to bind specifically to PL-DPPC lipid rafts that are enriched in the liquid-disordered (ld) lipid phase and appear bright in the fluorescence image while being depleted from the liquid-ordered (lo) phase (see text). (Right panel) Square jump distance (SD) between successive video frames ($\Delta t = 33$ ms) for a trajectory. The fluorescence image presented correlation between parts of the particle trajectory pertaining to the two different lipid phases. Trajectory analysis (see Materials and Methods) for this particle yielded $D_{ld} = 1.1 \times 10^{-7}$ cm²/s and $D_{lo} = 0.33 \times 10^{-7}$ cm²/s. (d and e) Double-stained POPC/cholesterol/phosphatidylcholine (2:1:1) phospholipid bilayer containing 1 mol % GM1 and 0.1 mol % TR-DPPC after a 10-min incubation with PL-CTB (1 μ g/ml in PBS) as observed in fluorescence channel (d) and fluorescence channel (e). (f–h) DOPC/cholesterol/phosphatidylcholine (1:1:1) containing 1 mol % GM1 with 0.5 mol % PL-DPPC as observed in fluorescence channel before (f) and after (g) a 30-min incubation with streptococcal cytolysin (Stx) (Toxin X-100 in PBS). It shows rafts after a post-attachment incubation with PL-CTB (1 μ g/ml in PBS for 10 min). With the assumption of $c = 1$ μ m, all scale bars are 10 μ m. All experiments were conducted at room temperature (20°C).

Lipid Rafts Reconstituted in Model Membranes

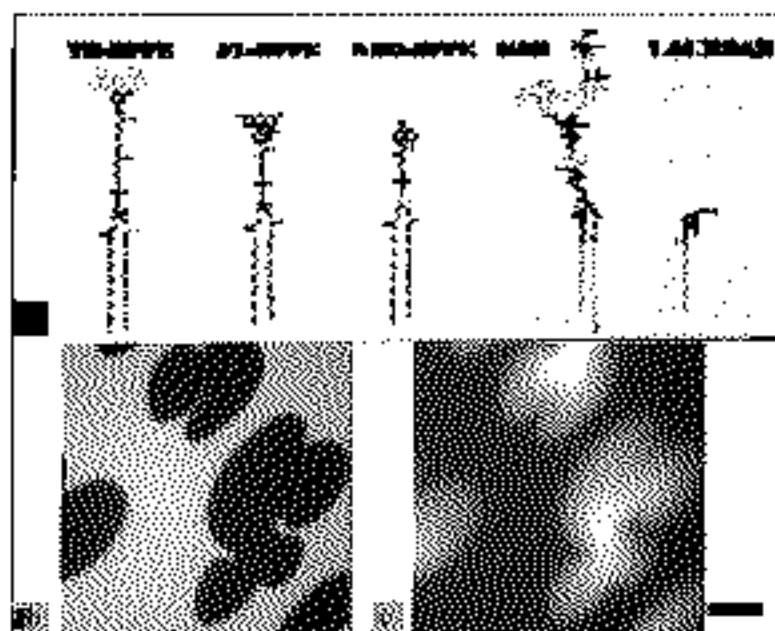
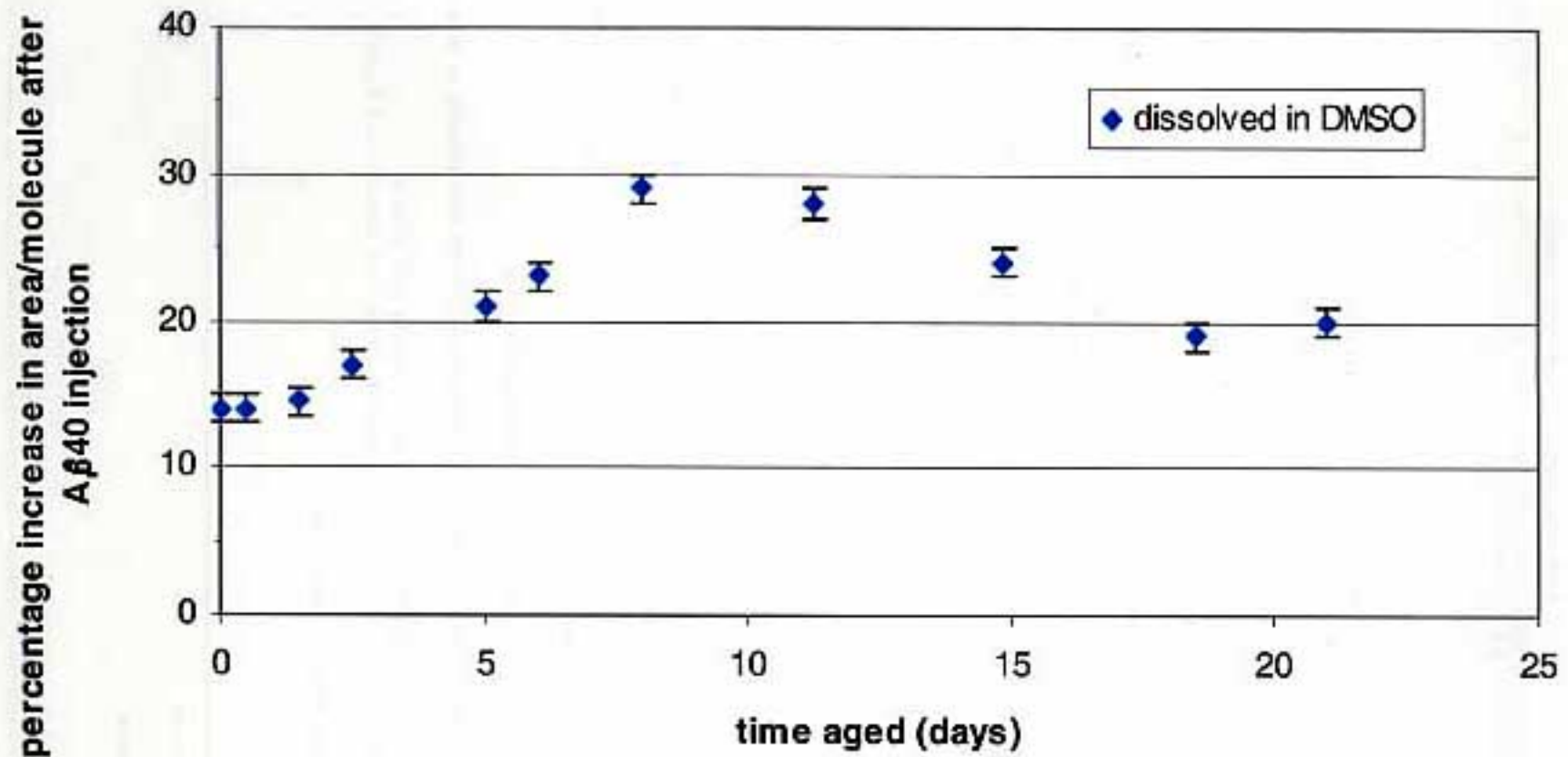
G. Datta,^{1*} L. A. Baginski,¹ Z. M. Wang,² H. L. Trossman,² M. Lee,³ K. Jacobson,^{4,5} and E. Sezgin¹¹Department of Cell Biology and Anatomy, ²Department of Physics, and ³Levinthal Computational Center, University of South Florida, Tampa, Florida 33620, ⁴Center for Biomedical Research, Department of Physics, University of Illinois at Urbana-Champaign, Urbana, Illinois 61801, and ⁵Department of Medicine, University of Texas Southwestern Medical Center at Dallas and Veterans Affairs Medical Center, Dallas, Texas 75216

FIGURE 2 Fluorescence microscopy of the vesicles. Fluorescent lipid analogs were used and the fluorescence images were obtained during the preparation of lipid DPPC vesicles with 0, 25, 50, 100, and 1 mol % cholesterol in addition to the total liposomes classed as 1. The control DPPC membrane was prepared in the vesicles buffer for lipid and put them at 3 degrees and 20°C from the outside interface, to a premixed DPPC membrane that was prepared previously to a phase transition at 5 degrees. To make stable, the vesicles were incubated with Biotinylated antibodies (Ab) for at least days 10–20.

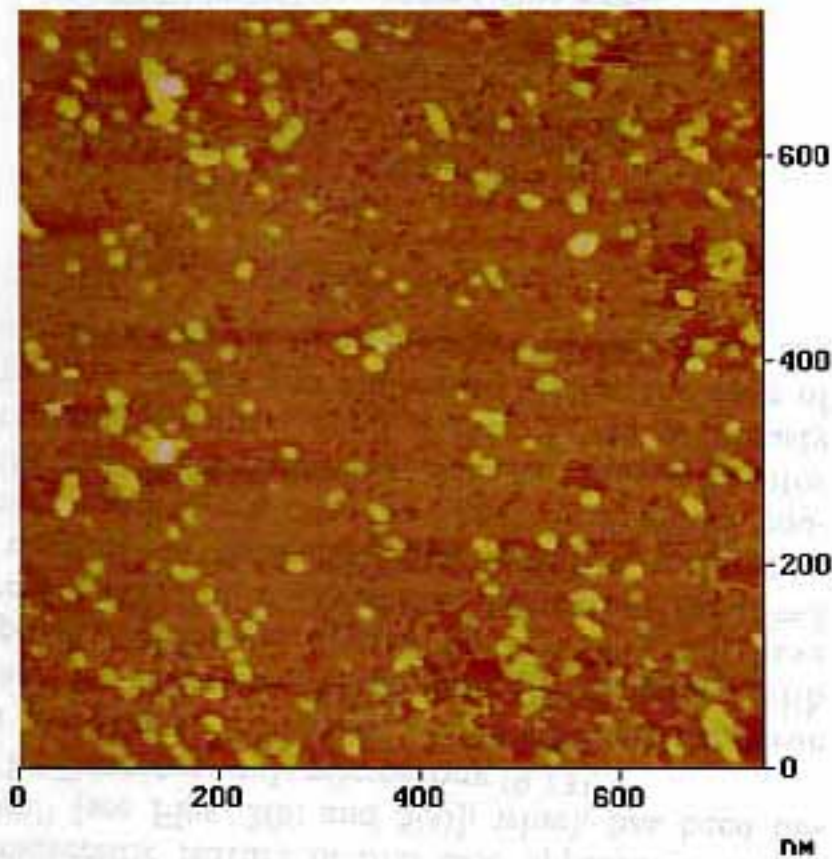
CONCLUSIONS

- A β 40 inserts into lipid membranes under physiologically relevant conditions, and the amount of insertion decreases with increasing surface pressure.
- Monolayer morphology changes drastically after A β 40 injection even under conditions with no observable trough expansion (apparent insertion).
- Lipid membranes need not be acidic for A β insertion: positively charged DPTAP and negatively charged DPPG show similar insertion profiles. Therefore, electrostatics may not be the driving force for peptide insertion.
- Aging the peptide promotes insertion.
- Dissolving A β 40 in the solvent DMSO causes up to a 40 minute delay in Insertion in pure water at high pressures (≥ 25 mN/m) and changes the insertion profile.
- Supported bilayers retains their fluidity and can be imaged by AFM.
- Lateral phase separation in bilayers can be detected.
- A β fibril shows helicity; modified with PEG do not form cross-mesh.
- Incubation with lipid vesicles leads to formation of protofilament on membrane surface.
- GIXD results confirms the disruption of condensed phase in lipid layer.

INSERTION PROFILE OF AGED A β 40 INTO A DPPG MONOLAYER



T=30 °C. Peptide concentration while aging was 12 μ M (just below the CMC). Peptide concentration in the subphase *after* injection is 250 nM. Subphase is pure water.



A β incubated in a solution of DOPC/GM1/A β 9/1/1 molar ratio for one day. A β concentration = 60 μ M. Vesicle concentration Pores on the membrane can be seen together with structures that resemble A β protofibrils.

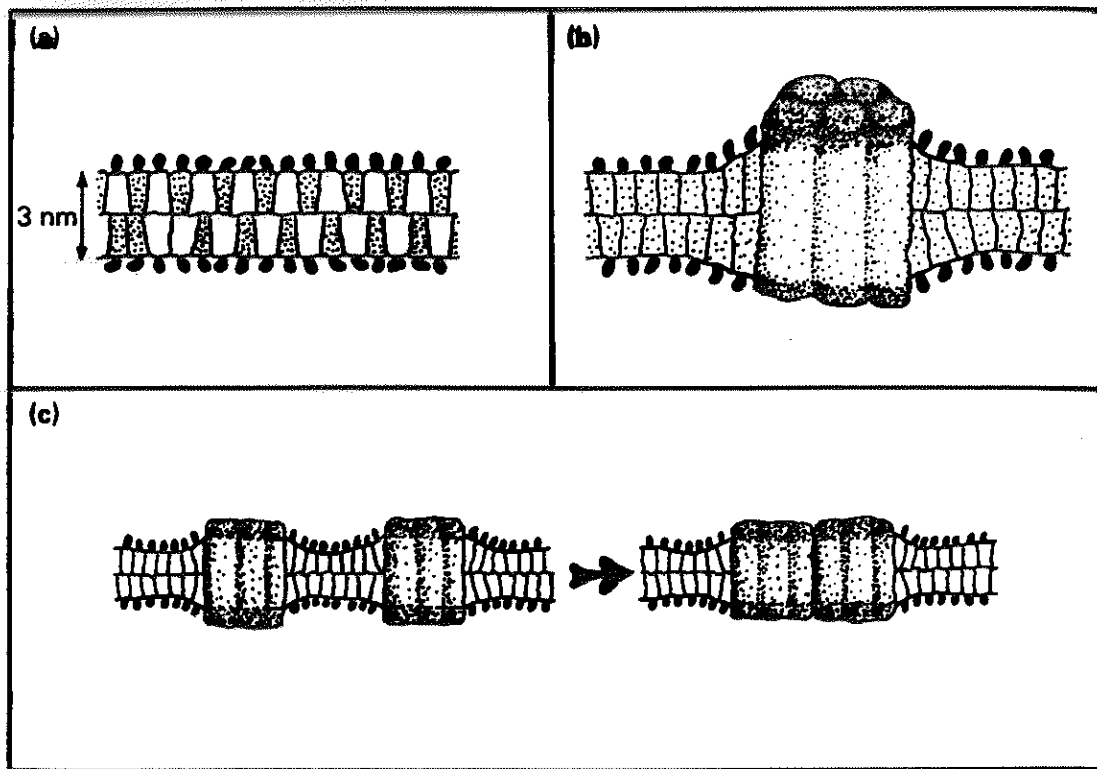


Fig. 17.7. (a) Mixture of two different lipids packing together within a planar membrane. The shaded lipids are cone-shaped ($v/a_0l_c < 1$); the white lipids are wedge-shaped ($v/a_0l_c > 1$). (b) Packing constraints induced in the hydrocarbon chain regions of lipids around a protein molecule, which may be relaxed when proteins aggregate, as shown in (c).

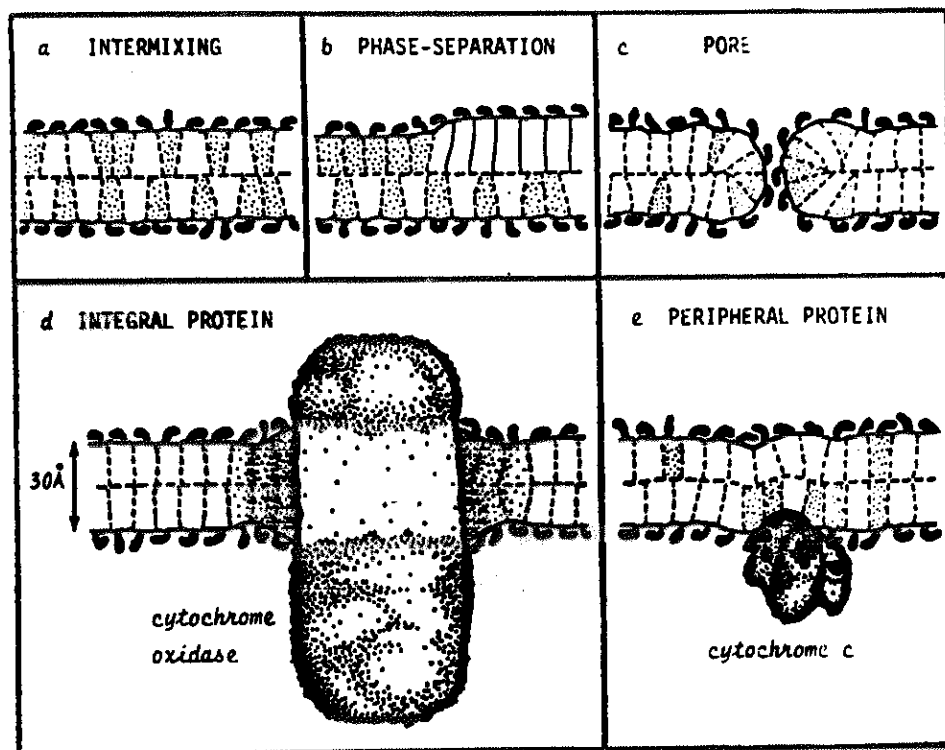


Fig. 17.9. Mean packing conformations of mixed lipid and lipid-protein membranes, showing how local packing stresses may cause clustering of specific lipids and/or non-bilayer shapes (shaded lipid regions). All the figures have been drawn to scale. Note the relatively large size of the cytochrome oxidase protein molecule which protrudes greatly from the bilayer. (From Israelachvili *et al.*, 1980a.)

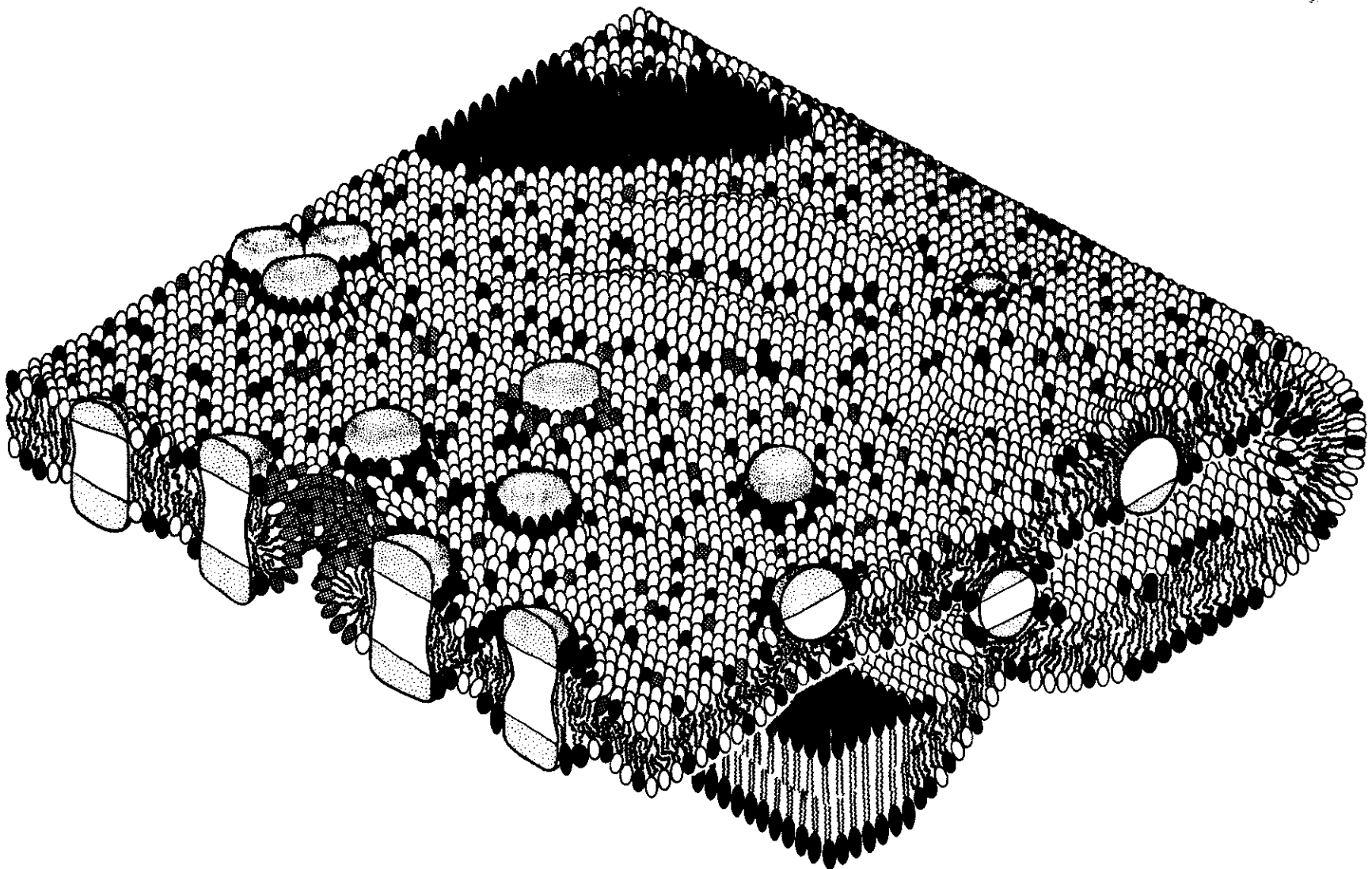


Fig. 17.8. Schematic figure of a biological membrane. The lipid contents of different membranes vary from as little as 25% up to 80% by weight. At highly curved regions the outer convex face contains mainly cone-shaped lipids while the inner concave face has more wedge-shaped lipids. Note how stressed regions may adopt a locally non-bilayer structure, some of which are shown in more detail in Fig. 17.9.

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Israelach

Supported Bilayers as model membranes

Dyn. + structural properties studied by

- FTIR
- NMR
- TIRF
- X-ray scattering
- neutron scattering
- lateral diff. measurements.
- imaging ellipsometry
- reflection interference contact microscopy
- SFA

Bilayer on beads → NMR

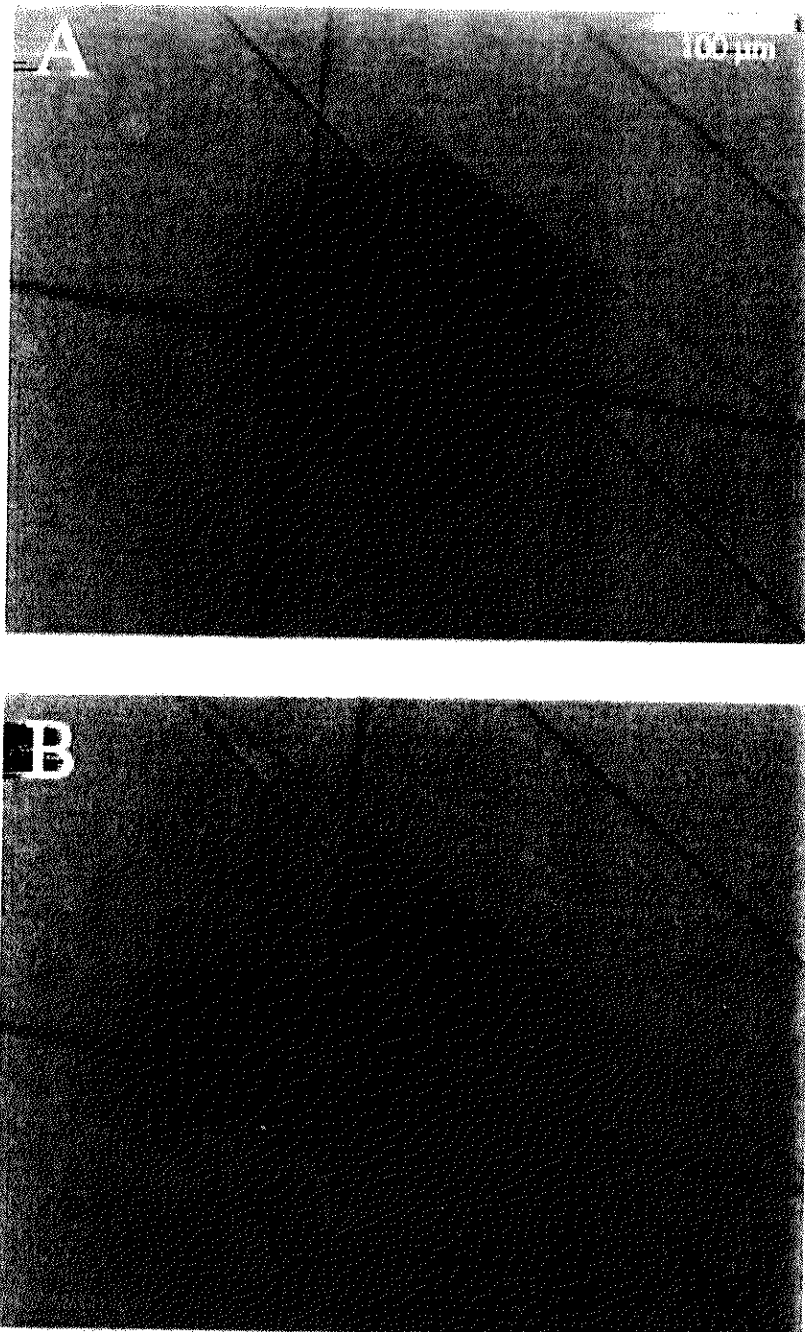


FIGURE 6: (A) Photograph of fluorescence from supported bilayers containing RCs on glass. The mole ratio of RC:lipid in the sample shown is about 1:700. One mole percent of Texas Red DHPE was used as the fluorescent lipid. The sharp dark lines are scratches in the lipid bilayer surface that create physical boundaries to lateral diffusion of the lipids. Prior to being scratched, the surface appeared homogeneous. A circular region of the lipid was shadowed out by the microscope aperture and photobleached with the excitation light. (B) The same field of view 20 min after photobleaching. The photobleached area now fills the central bounded triangular region; other unbounded regions outside the triangle have recovered a substantial amount of fluorescence, as long-range lateral diffusion of fluorescent and photobleached lipid leads to mixing. These results demonstrate that the lipid component is freely diffusing in two

cysteine-reactive fluorescent rhodamine for imaging the protein component. These supported bilayers were composed of PE instead of the Texas red lipid. The fluorescence spectrum of the supported bilayers is uniform and exactly matches the scratch-dye-lipid probe for a given protein. The fluorescence intensity were measured following bleaching of a region. Unlike the lipids described above, the protein components in a native membrane diffuse on this time scale. Diffusion coefficients between 1 and 5 $\mu\text{m}^2/\text{s}$ (Gennis) in the supported bilayers are observed.

An identical experiment was performed with the *Rb. capsulatus* RC mutant (M)189 and the fluorescent dye R-492. This experiment was designed to investigate whether the lack of mobility in the *Rb. sphaeroides* RCs might be due to the fluorescent label and the protein. The engineered cysteine is a surface cysteine (H-156 cysteine) in the *Rb. capsulatus* RC) from a construct that is a combination of the *Rb. sphaeroides* RC and the *Rb. capsulatus* RCs. This was verified experimentally, as the mobility of these RCs was observed to be high. The engineered cysteine is situated on the opposite side of the membrane from the mobility of these RCs was observed.

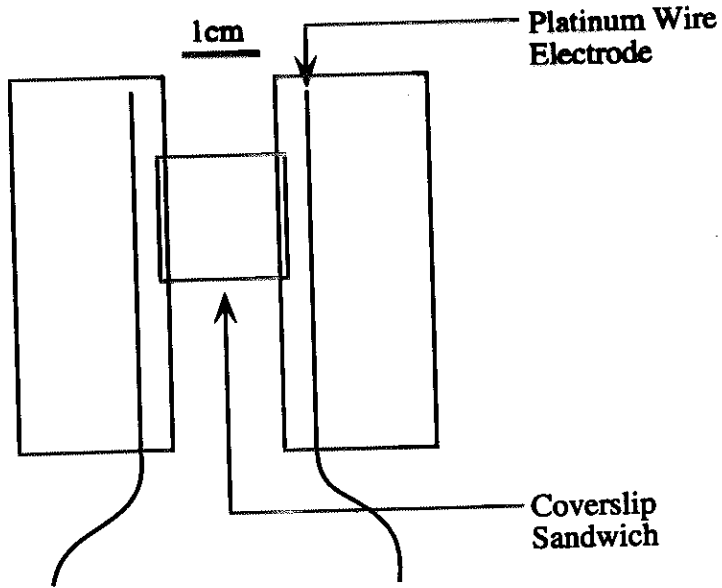
DISCUSSION

Our goal in this work was to demonstrate that oriented RCs on a solid support can be prepared that RC-containing proteoliposomes preformed SUVs in a highly oriented cytochrome *c* binding site facing the surface with the result reported by Sakaguchi, which was substantiated by the biotin labeling of the cytochrome *c* binding site with the cyt *c* binding experiment. This result is genuinely oriented in a transmembrane orientation.

The RCs remain oriented in the membrane with the cytochrome *c* binding site facing the surface.

FIGURE 1

A.



B.

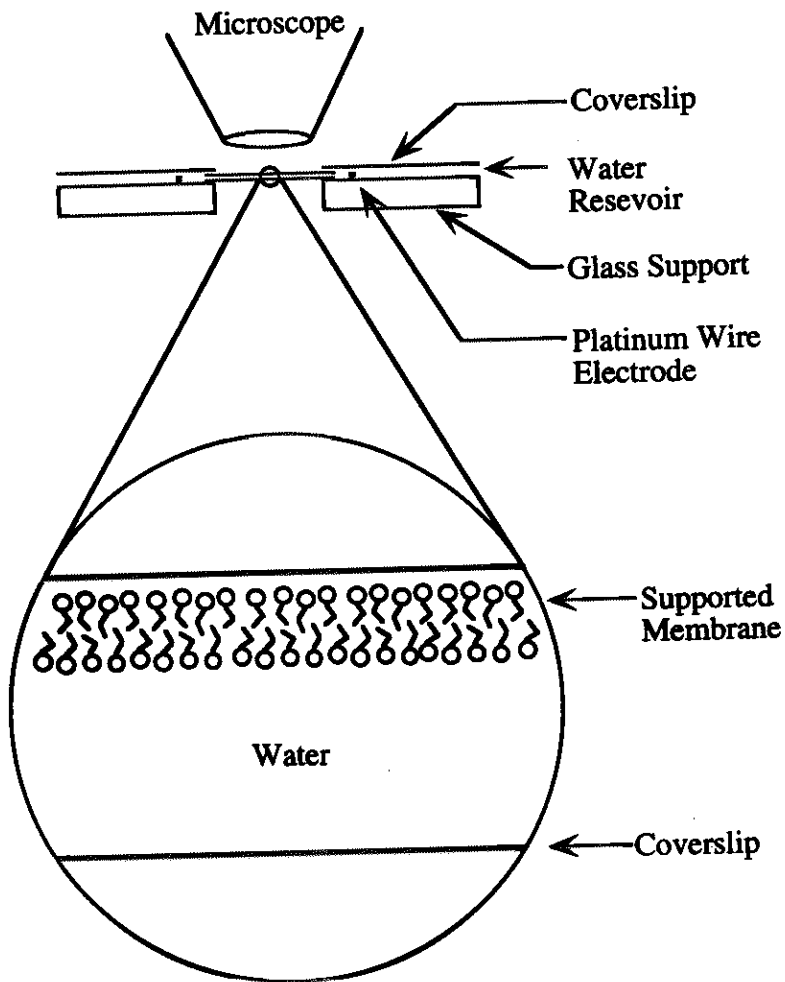


FIGURE 1
Groves et al.

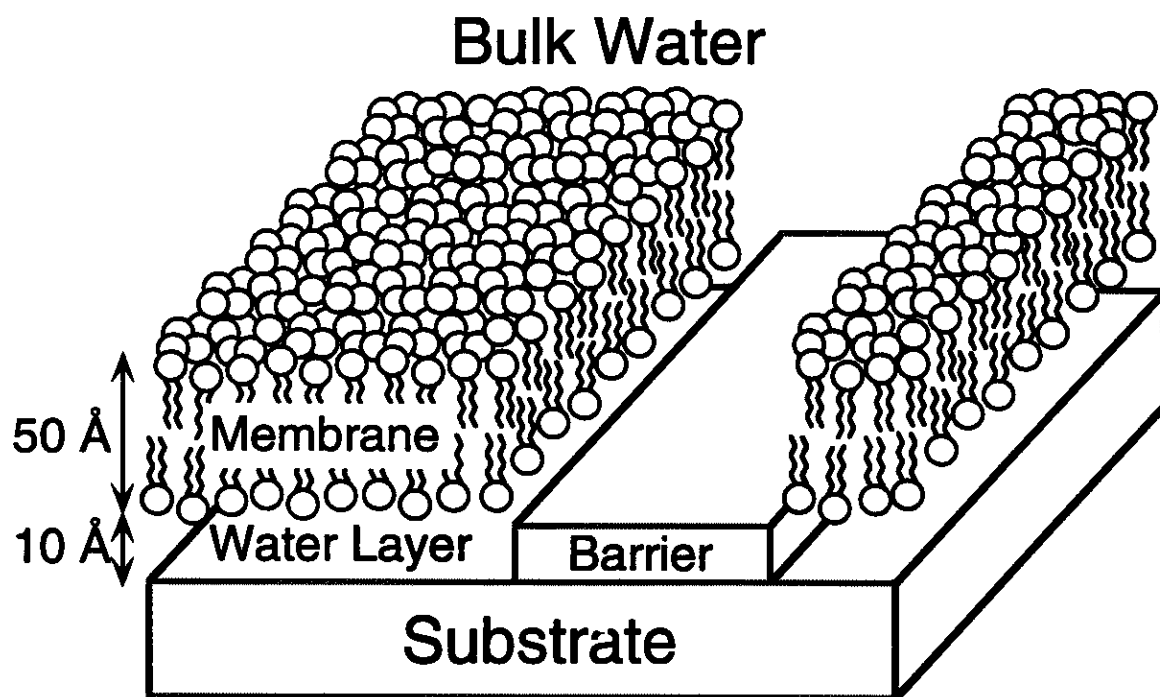
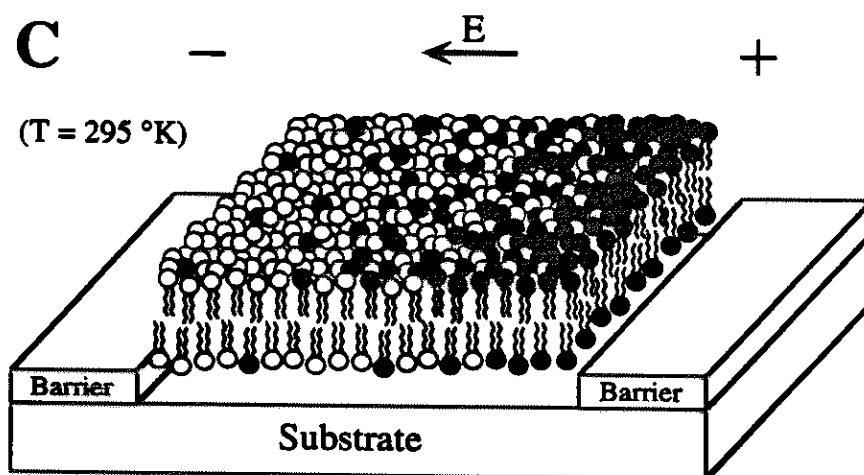
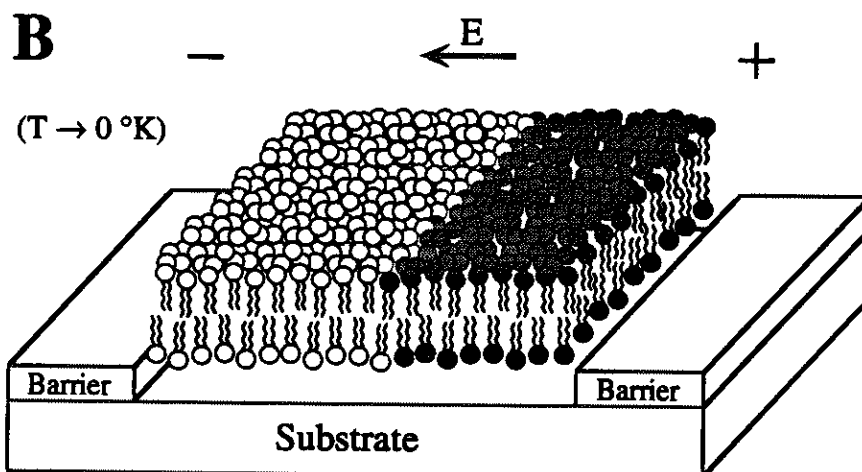
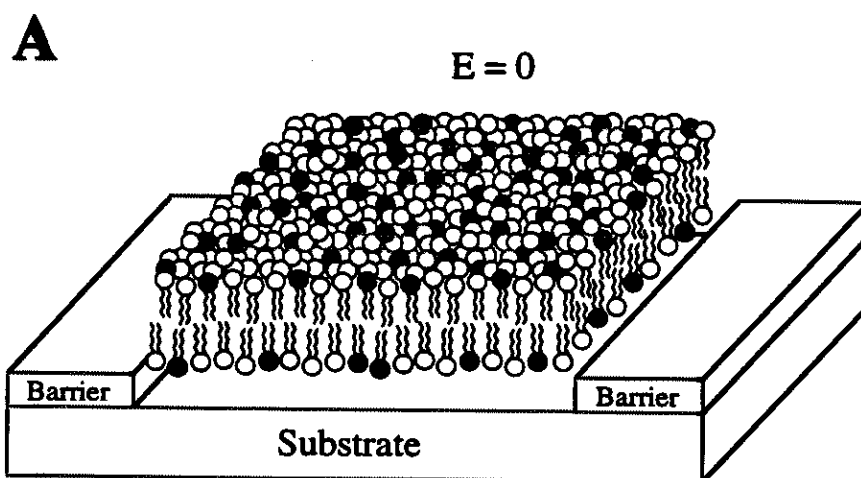
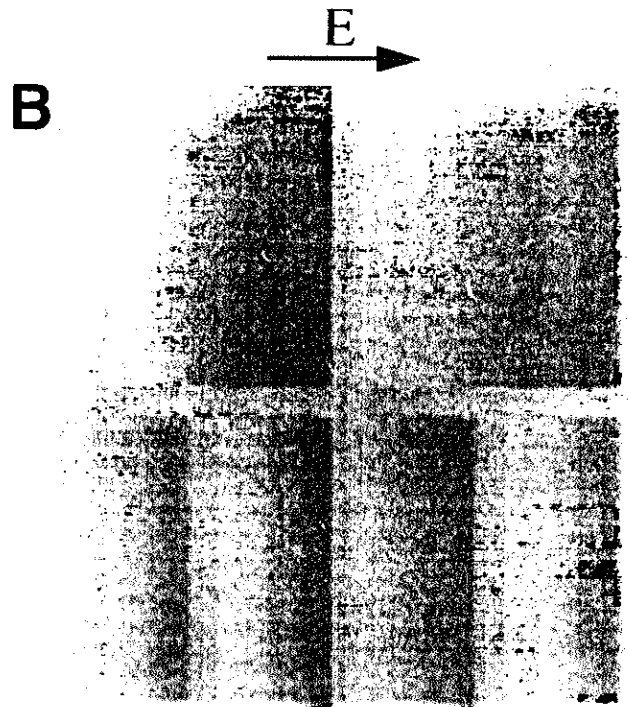
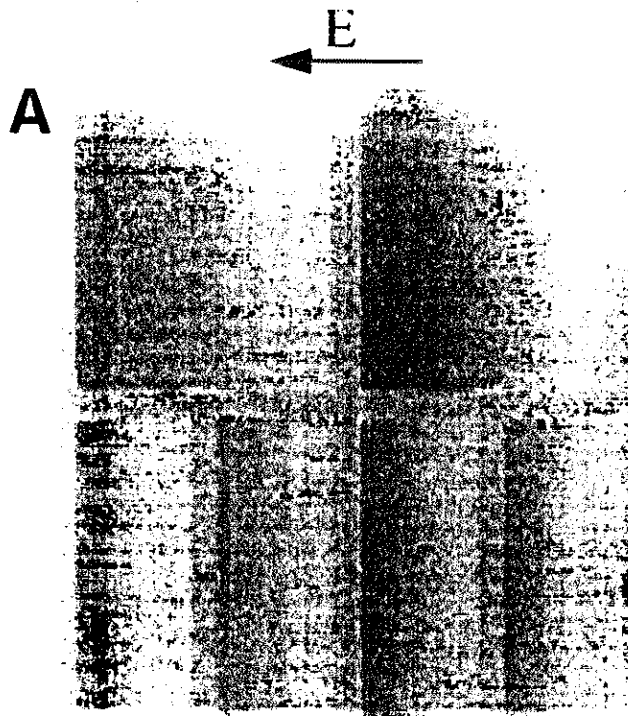


FIGURE 1
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functionalization of solid surface

for anchoring polymer

mixed monolayer of long chain surfactant
exhibiting flat head group - binding
& alcohol - non binding

Immobilization of protein

Streptavidin - biotin

functionalizing deposited membrane

incorp. nanoparticle receptor

Reversible coupling

Chelating lipid

w/ divalent ions: His tag binds

w/out " " : detach^t occurs

(EDTA)

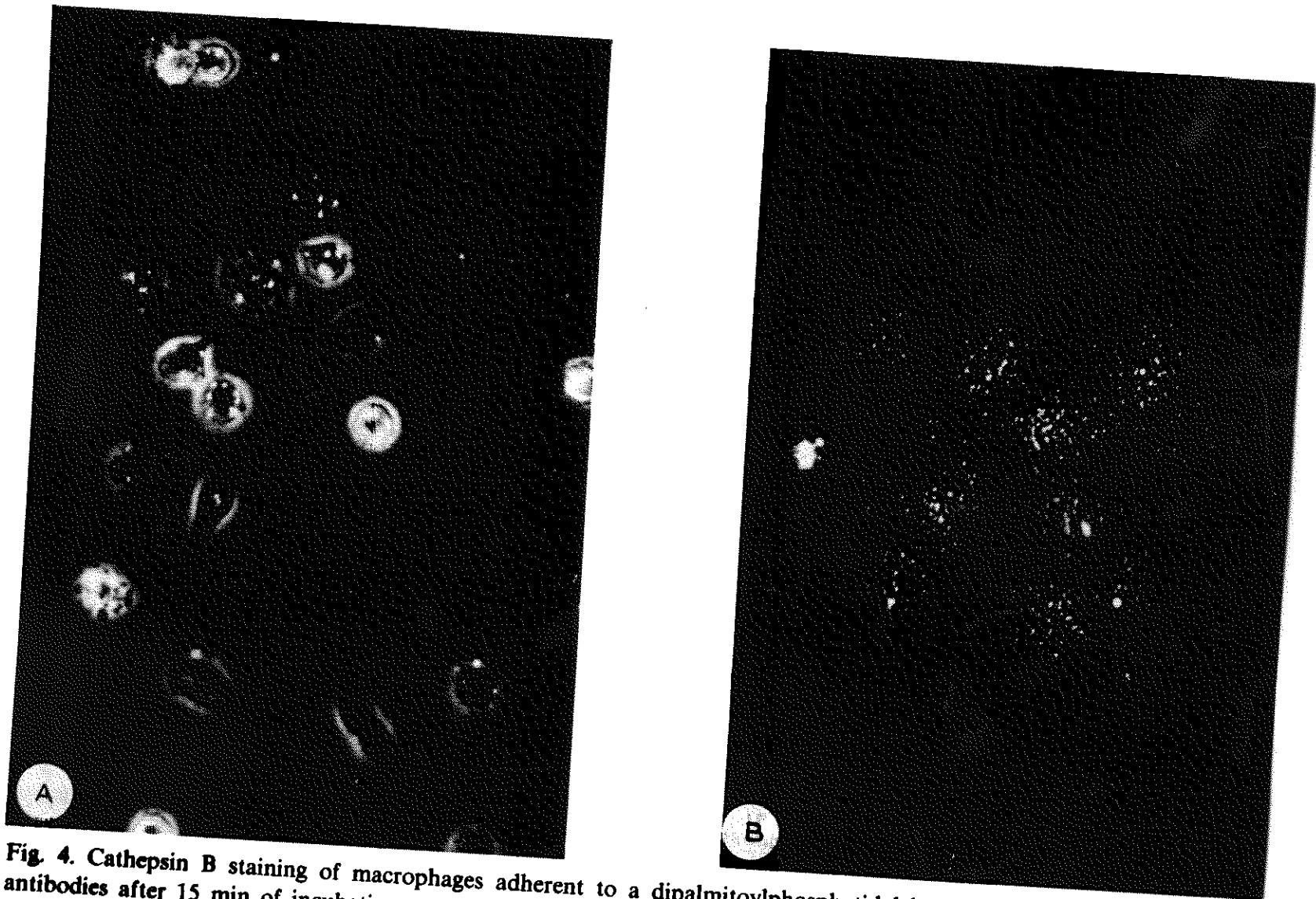


Fig. 4. Cathepsin B staining of macrophages adherent to a dipalmitoylphosphatidylcholine monolayer with lipid-hapten-bound antibodies after 15 min of incubation at 37°C. The yellow-orange granules of fluorescent precipitate formed by the enzyme are localized both inside the cells and outside the cells (associated with the monolayer). (A) Combined phase contrast and epifluorescence photomicrograph taken in the median focal plane of the cells. (B) Epifluorescence photomicrograph taken in a focal plane between the monolayer-coated cover glass and adherent macrophages, showing accumulation of fluorescent precipitate. From Hafeman et al. [29].

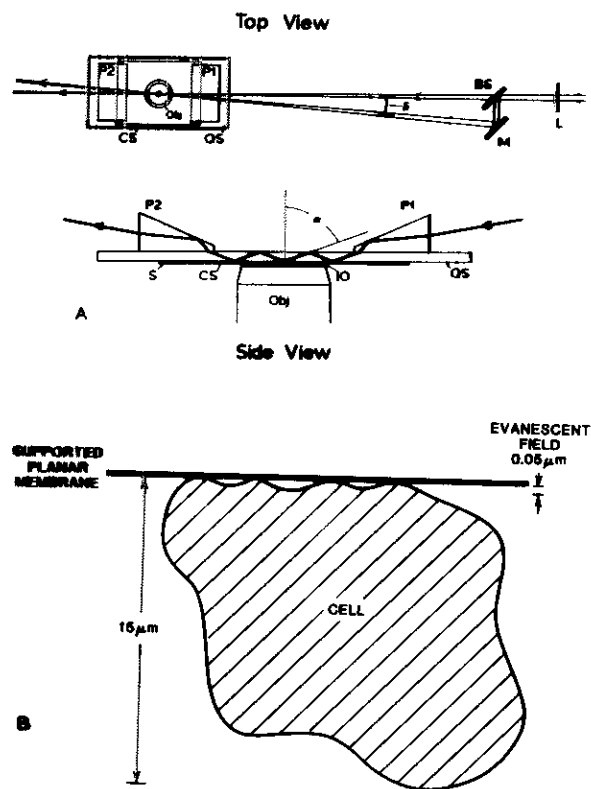


Fig. 5. Evanescent illumination of supported planar membranes. (5A) Apparatus for producing an evanescent radiation field so as to stimulate fluorescence of molecules that are within 100 nm of the lipid monolayer coating the lower surface of the quartz microscope slide (QS). The prisms P1 and P2 are used to introduce and remove the vertically polarized laser beam. Cells are bound to the monolayer on the lower surface of the quartz slide, and are bathed in cell buffer localized between the monolayer and a coverslip (CS). A spacer (S) maintains a fixed distance, 24 microns, between the quartz slide and the coverslip. An objective lens with immersion oil (IO) is used to view cells bound to the monolayer. The top view shows the setup employed to produce an interference pattern in the evanescent wave field; the beam splitter (BS) divides the beam into two beams of equal intensity, and the mirror (M) directs one beam so as to produce an interference pattern over the objective lens of the microscope. Taken from

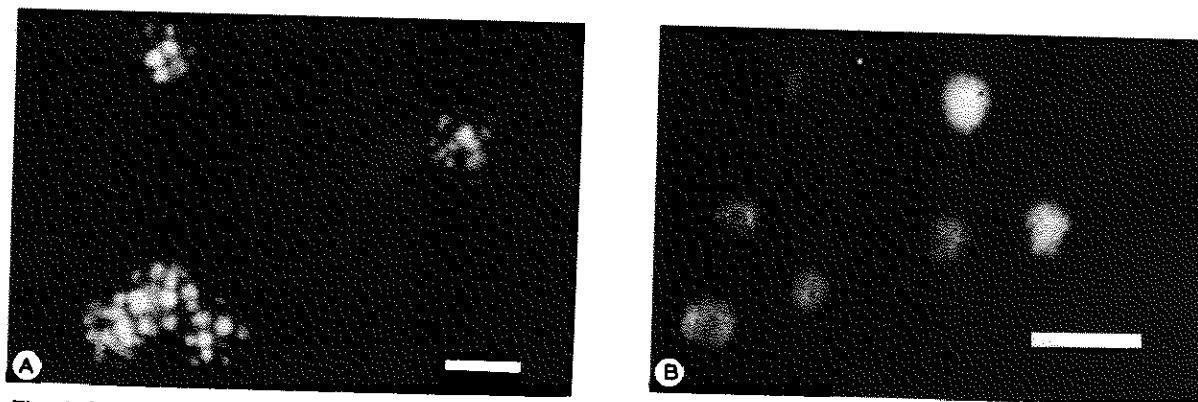


Fig. 6. Rat basophil leukemia cells bound to dipalmitoylphosphatidylcholine monolayers containing two mol% dinitrophenyl lipid-hapten. The scale bars are 25 μm . Before binding to the monolayer membranes, cells were incubated with either fluorescein-labelled monoclonal anti-dinitrophenyl-IgE (6A) or 1:1 anti-dinitrophenyl-IgE: fluorescein-labelled rat myeloma IgE (6B). Cells were illuminated by total internal reflection as described in Fig. 5. Monoclonal anti-dinitrophenyl-IgE (clone 26.82, [41]) was obtained from F.T. Liu and D.H. Katz, Scripps Clinic and Research Foundation, La Jolla, CA. Rat myeloma IgE was obtained from Henry Metzger, National Institutes of Health, Bethesda, MD. The rat basophil leukemia cells were incubated with the monolayers in complete cell buffer: 135 mM NaCl/5 mM KCl/1 mM MgCl_2 /1.8 mM CaCl_2 /5.6 mM glucose/0.5 mg per ml gelatin/10 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid at pH 7.2.

would have been very difficult to reach using conventional cell culture techniques.

Supported planar membranes containing H-2K^k, prepared by the method shown in Fig. 3, were more effective in inducing secondary cytotoxic responses in vitro, than the small unilamellar vesicles from which the supported planar membranes were prepared [18]. In addition, through the use of supported planar membranes, in this case made from vesicles containing H-2k^k spread at an air-water interface, it has been possible to show that cloned cytotoxic T cells bind specifically to target membranes containing only purified allo-antigen [37]. The T cell clone used in these studies was derived from conventional mixed lymphocyte culture, and selected on the basis of its cytotoxicity for H-2K^k-bearing target cells.

The phenomenon of recognition of allo-antigen

the generality of this model for allo-recognition follows from the fact that T cells, which were selected only on the basis of their cytotoxicity for target cells bearing the usual assortment of cell-surface proteins, bind planar membranes containing only the allo-antigen.

In general, it has been difficult to stimulate helper T cell responses with small unilamellar vesicles. Curman et al. [16] were able to stimulate a secondary mixed lymphocyte reaction using partially purified I-A in phospholipid vesicles. The only report of an antigen-induced class II restricted T cell response to purified major histocompatibility complex molecules in liposomes is by Walden et al. [17]. In these experiments, antigen was covalently linked at high concentrations to the target liposomes. We have used affinity purified I-A^d and peptide antigen to stimulate

Polymer-induced forces

- rep. if anchored to the interface.
- If exchange is possible
 - repulsive @ large dist.
 - attractive @ short "

Dynamic membrane roughness

- determined by out-of-plane motions of lipids

2-3 Å

- thermally excited membrane undulations.
 - create pronounced bending undulations with amplitudes $\sim 10 \mu\text{m}$

Undulation free control cell adhesion,

flickering - shape fluctuation

- prevents sticking
- cell aply

- strong undulation $\sim 100 \text{ nm}$ for surface couples them' specific short-range forces repulsive long range disjoining pressure.

Hydrated Membranes as Electrooptical Biosensors

bilayer used as

- 1) a very thin electrical insulator
- 2) a matrix for incorporatⁿ of receptor
- 3) suppression of nonspecific ligand binding

Support: gold or $\text{SnO}_2/\text{ZnO}_2$ or metal oxide.
semiconductor (MOSFET)

ligand binding: capacitance measurement
current

$$C_m = \frac{\epsilon}{d_m} \text{ surface charge } ds$$

Impedance Spectroscopy

- SPR (1.3 - 1.8 μm) water & glycerol disrupt
- surface acoustic waves
- lateral distribution of membrane-bound ligands may be organized by 2D electrophoresis
→ separate bound & free receptors

Supported Membranes as Plasmalemma Cells

interplay of specific & universal forces

polymer-induced forces (up)

dyn. repulsive forces

- areal density of receptor

→ control lock & key interaction

- Cell adhesion in immunological response
MHC replace antigen presenting cells
to stimulate T helper cells

Polymer-Lipid Composite Films

- Reconstitution of membrane. spanning proteins & the reorg. of receptors by 2D electrophoresis

- Study stability of membrane proteins coupled to macromolecular network

- Mimic cytoskeleton & polymer cushions (highly cross-linked)

ex. like spectrin-actin network.

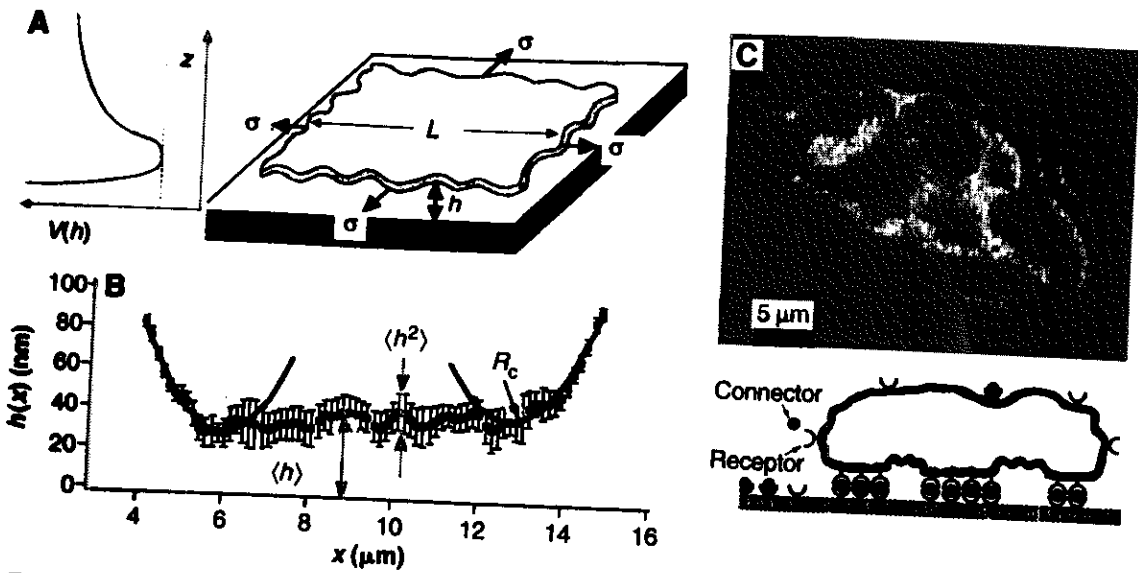


Fig. 7. Analysis of undulation forces between lipid vesicle and rigid wall. (A) Schematic view of a dynamically rough piece of membrane of dimensions L by L at distance h above the solid surface with lateral tension σ . The excitation may be decomposed into plane-wave Fourier modes. The effect of the wall is represented by an interfacial potential $V(h)$. (B) Representation of parameters characterizing weakly adhering vesicles simultaneously measurable by RICM. These include (i) mean distance $\langle h \rangle$ (~ 400 Å), (ii) mean square amplitude $\langle h^2 \rangle$ (~ 20 nm²), (iii) lateral tension ($\sigma \sim 0.001$ mN/m), (iv) force constant of potential $\partial^2 V / \partial z^2$, (v) contract curvature R_c , and (vi) spatial correlation function of undulation amplitude $u(x)$ (27). (C) RICM visualization of a dimyristoyl phosphatidylcholine (DMPC) vesicle containing 1 mol % biotin-labeled lipid as the receptor interacting with a surface covered by streptavidin. The formation of adhesion plaques is caused by interplay of the strong attraction between receptors and the repulsive undulation forces.

by increasing the tension, a transition from a weakly to a strongly bound state may occur and one may therefore speak of tension-induced adhesion (33). Transitions between free and bound states may arise if swelling is induced in vesicles or cells, leading to an increase of the lateral tension.

The remarkable strength of undulation forces is demonstrated in Fig. 7C. Vesicles containing ~ 1 mol % biotinylated lipid (as model "receptors") adhere to substrates covered with streptavidin to form tight binding sites. Although the specific attraction is very strong, only adhesive plaques

by 2D electrophoresis (isoelectric focusing), (ii) fundamental studies of the mobility of membrane proteins coupled to macromolecular networks, and (iii) the deposition of self-healing (defect free) membranes to increase the electric resistivity and suppress nonspecific binding (Fig. 5). Composite monolayer-polymer films are also versatile model systems for studies of fluid-fluid wetting by viscous fingering or solitary waves (14) as well as dewetting processes. To mimic the physical properties of cytoskeletons beneath cell plasma membranes (such as the spectrin-actin network of erythro-

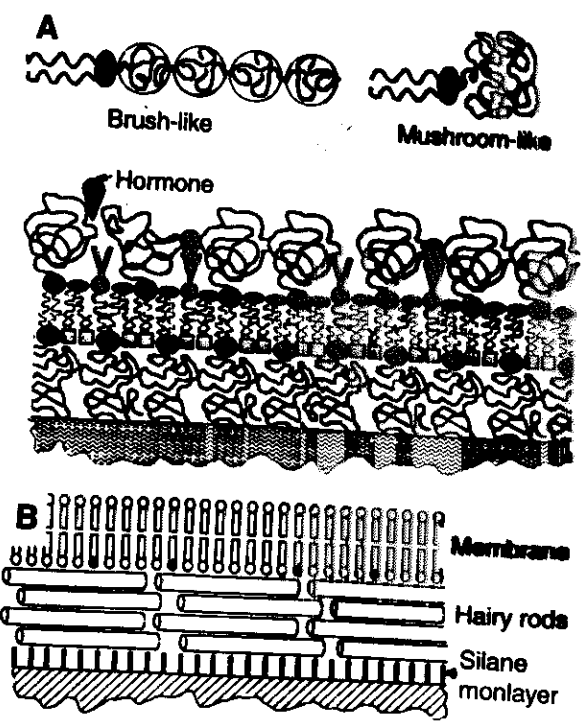


Fig. 8. (A) Separation of a bilayer from substrate by stealths formed by (clusters of) polymer brushes (or mushrooms) of hydrophilic polymers (for example, polyethyleneoxide grafted to long-chain lipids). Also shown is the formation of artificial glycocalyx on the outer monolayer by the mixing of phospholipids, lipid-coupled receptors, and lipopolymers. (B) Membrane deposition on ultrathin layers of hairy rods prepared by the LB technique from hydrophobic hairy rods (cellulose-derivatives), which are hydrophilized by shaving of hydrophobic hairs in HCl vapor.

varied by means of the lateral pressure because the polymer can be switched between the stretched (brush-like) and more expanded (mushroom-like) conformations (36). Lipopolymers in the outer monolayer allow one to synthesize artificial glycocalices to study the influence of steric repulsion on cell-substrate interactions or the