

*Reconstruction of a transmembrane  
protein tetraspanin (CD9) into lipid  
bilayer by interaction of ganglioside  
GM3 and tetraspanin*

Glycobiology World Congress

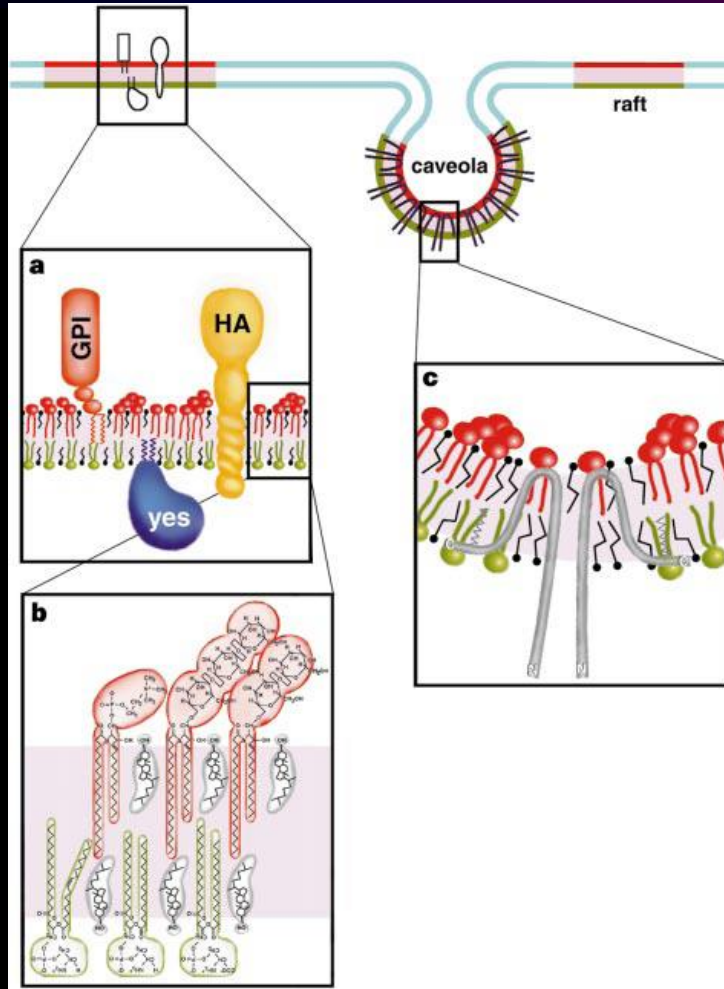
August 12, 2015

Tokyo University of Science, Japan

Shigeomi Horito

# Model for the organization of rafts

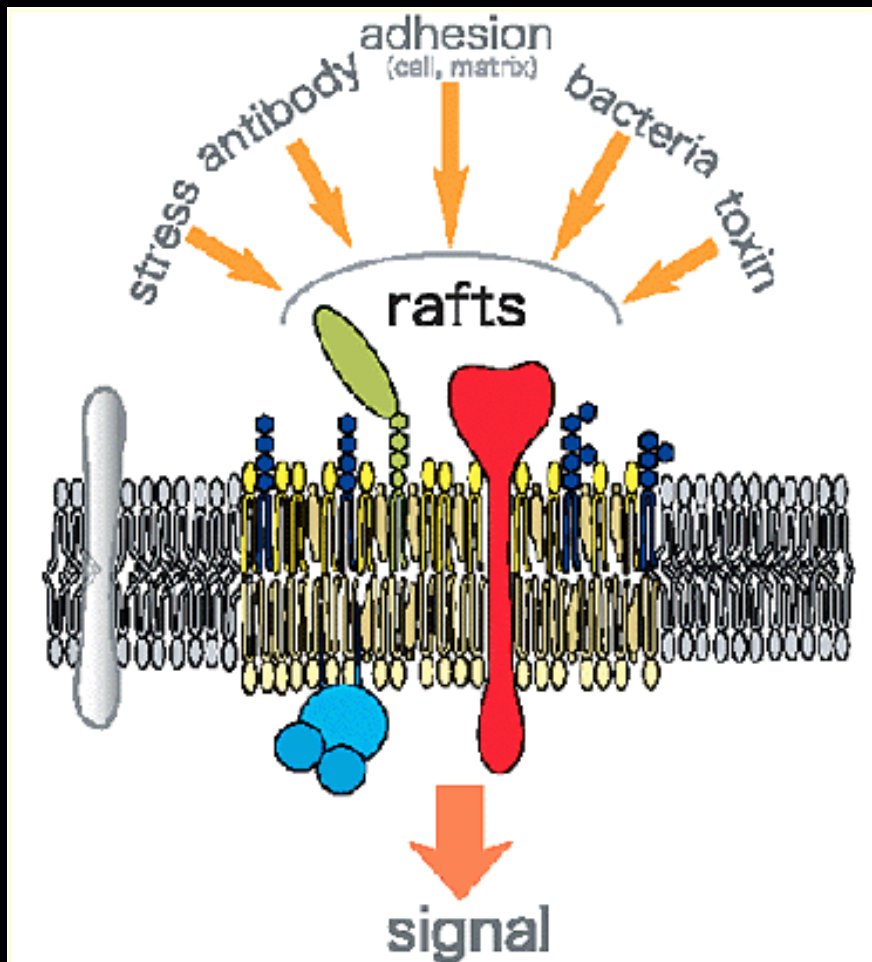
Kai Simons & Elina Ikonen, NATURE | VOL 387 | 5 JUNE 1997



The rafts (red) segregate from the other regions (blue) of the bilayer.

Rafts contain proteins attached to the exoplasmic leaflet of the bilayer by their GPI anchors, proteins binding to the cytoplasmic leaflet by acyl tails (the Src-family kinase Yes is shown), or proteins associating through their transmembrane domains, like the influenza virus proteins neuraminidase and haemagglutinin (HA).

# *Function of ganglioside rafts*



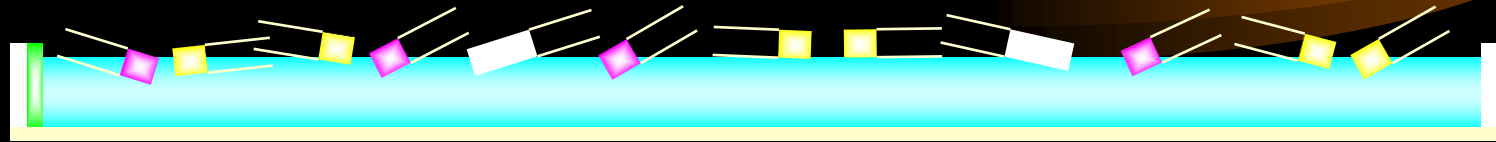
On outer membrane of raft, it is very clear that many ligands link to the corresponding receptors.

How is the signal transduced on inner membrane of raft?

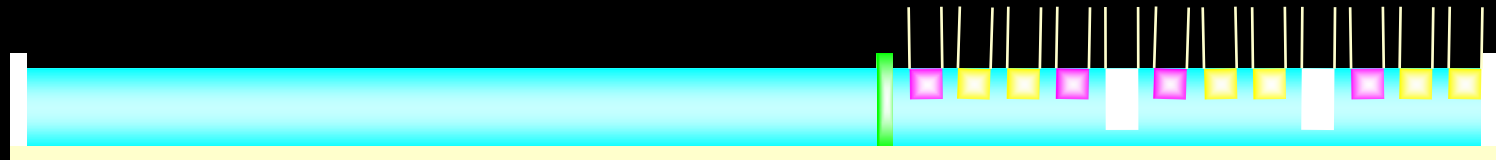
For example, autophosphorylation enzyme or cutting enzyme of signal peptides.

How is the enzyme recruited?

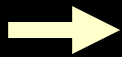
# *Preparation monolayers by Langmuir-Blodgett technique*



Lipids solution were spread on the subphase using microsyringe.



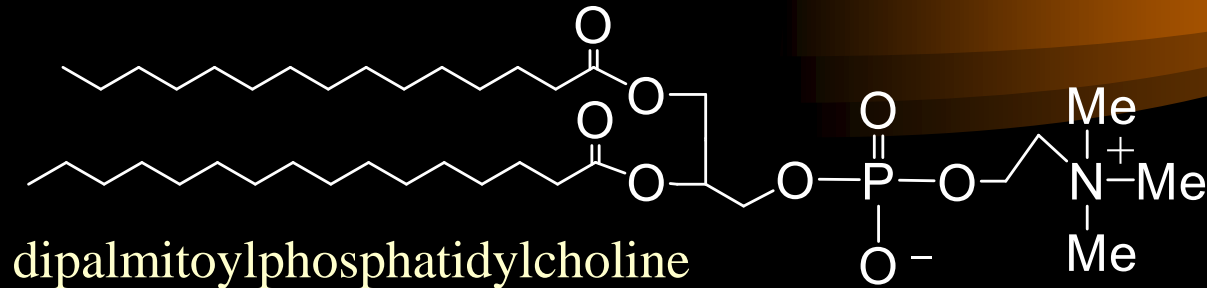
After evaporation of the solvent, the monolayer was compressed at  $50 \text{ cm}^2 / \text{min}$ .



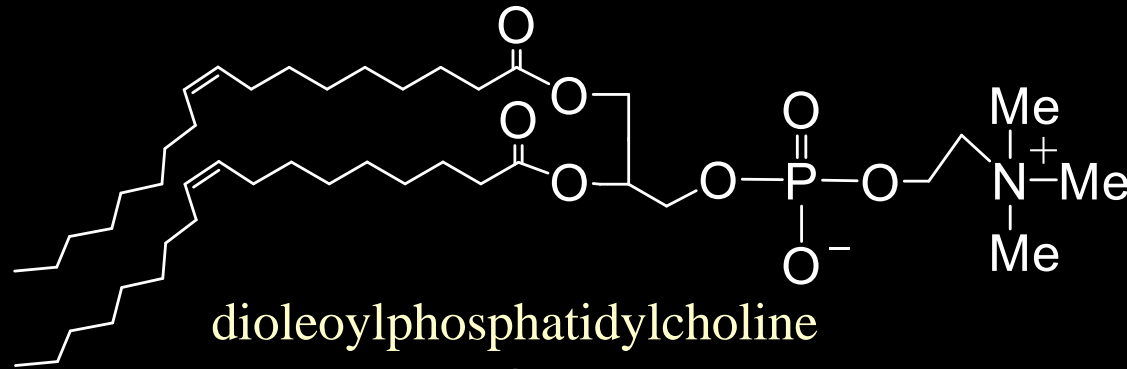
**The half side of cell membrane can be obtained.**

# *Molecular structures of lipids for L-B film preparation*

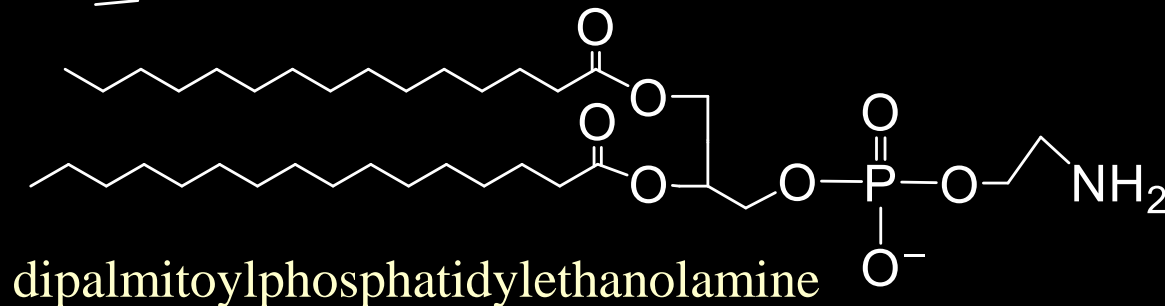
**DPPC**



**DOPC**

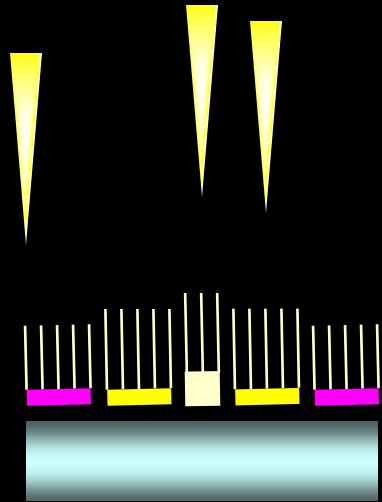


**DPPE**

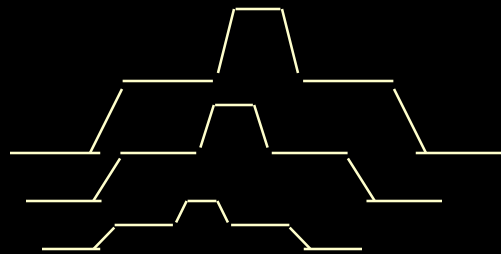


# *About atomic force microscopy (AFM)*

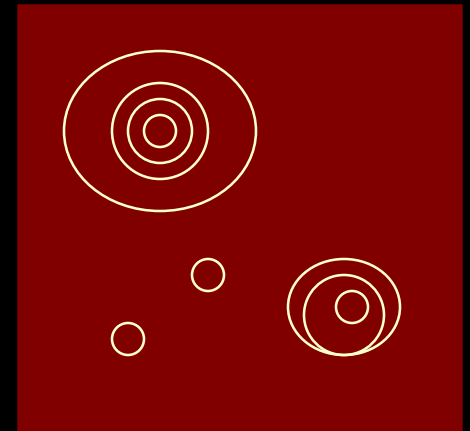
**Atomic Force between probe and sample is able to measure.  
Same value of atomic force means same distance.**



**sample**



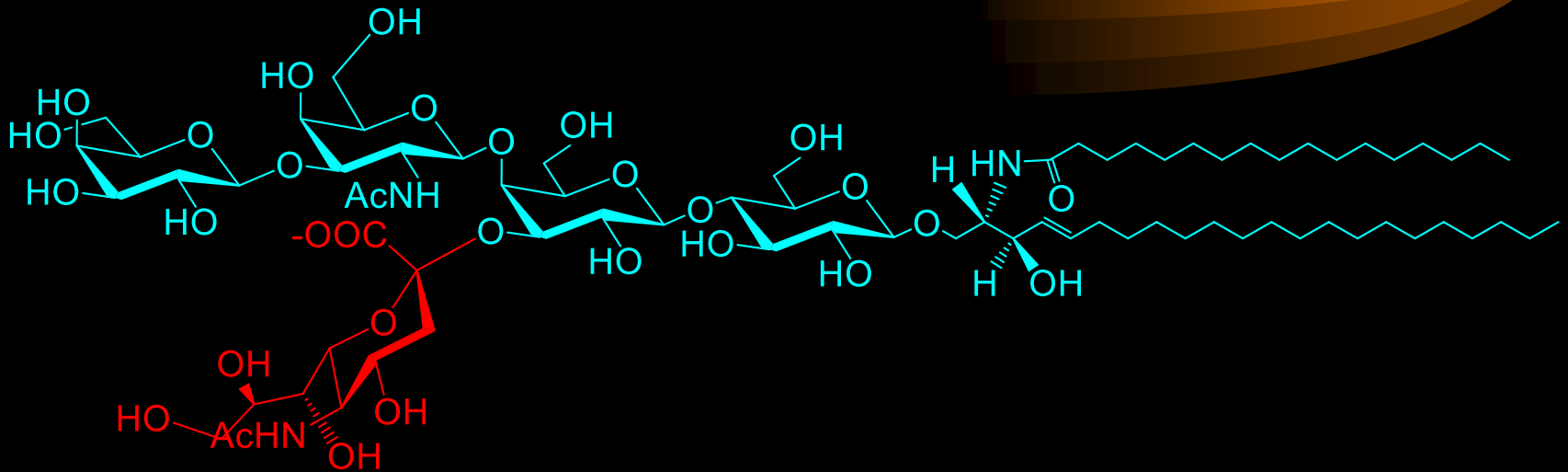
**Hight of sample surface  
was obsurbed from  
position of probe.**



**hight map**

**Resolution: about 0.2 nm**

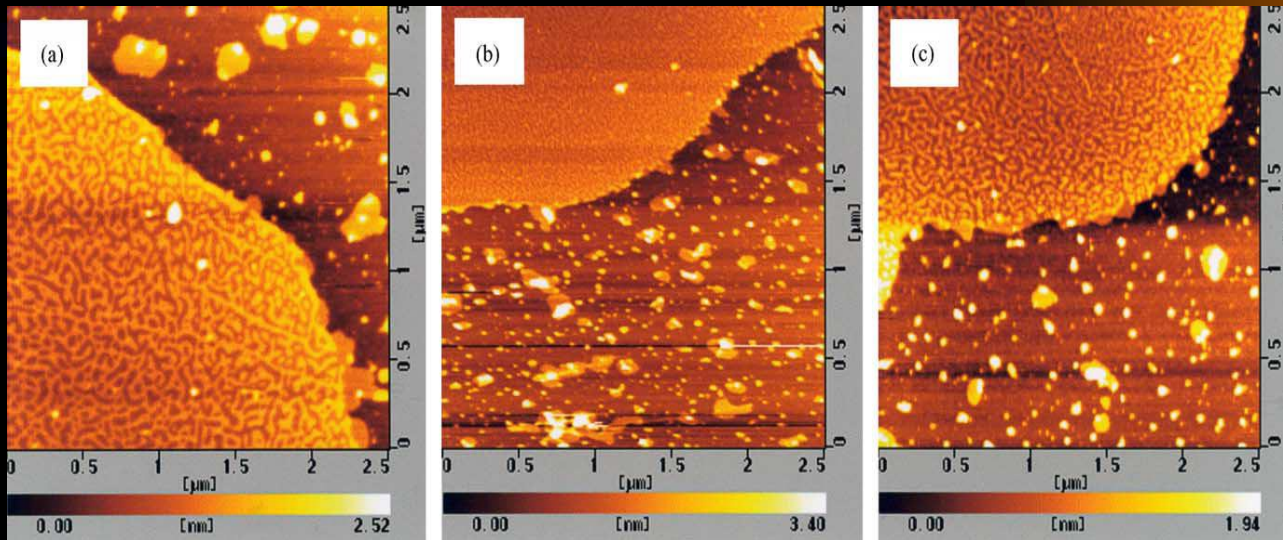
# *Chemical structure of ganglioside GM1*



Ganglioside was composed of carbohydrate part and lipid part (Ceramide).

Ganglioside GM1 has a sialyl  $\alpha 2 \rightarrow 3$  galactose residue.

# *Membrane properties of binary and ternary systems of ganglioside GM1/DPPC/DOPC*



Effect of surface pressure on the AFM images for the GM1/DPPC/DOPC (2:9:9) monolayer. Surface pressure: (a) 30 mN/m; (b) 35 mN/m; (c) 40 mN/m.

Conclusion: The percolation pattern in the GM1/DPPC/DOPC monolayer changed as the surface pressure was varied.

Yumiko Ohta, Shoko Yokoyama, Hideki Sakai, Masahiko Abe  
Colloids and Surfaces B: Biointerfaces 34 (2004) 147–153.

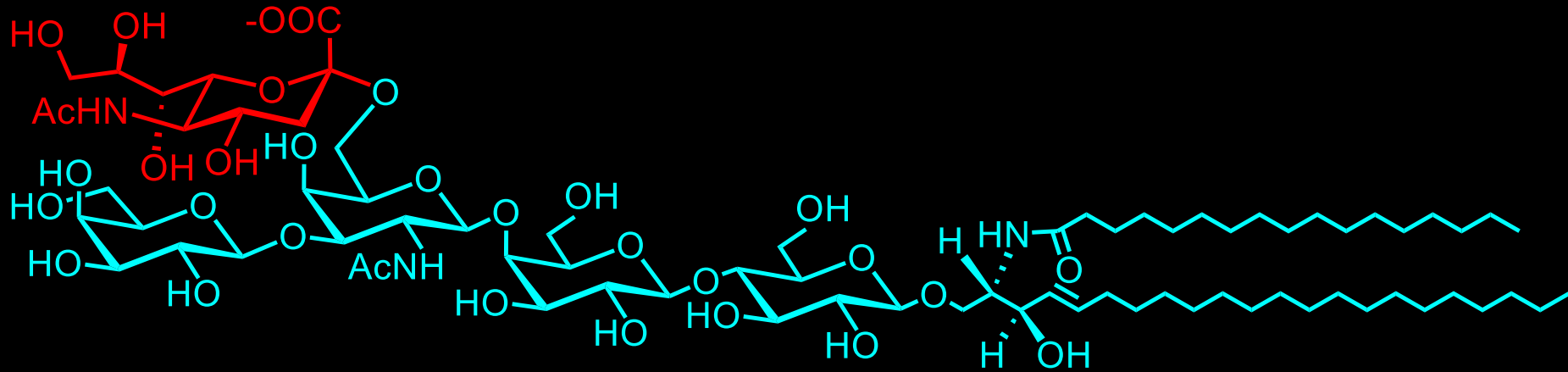


# *1. Research on ganglioside GM1a raft*



- GM1 $\alpha$  is not commercially available, We must synthesize it.
- GM1 $\alpha$  in DOPC/DPPC monolayers and hybrid bilayers are prepared.
- Distribution of GM1 $\alpha$  is investigated by AFM.

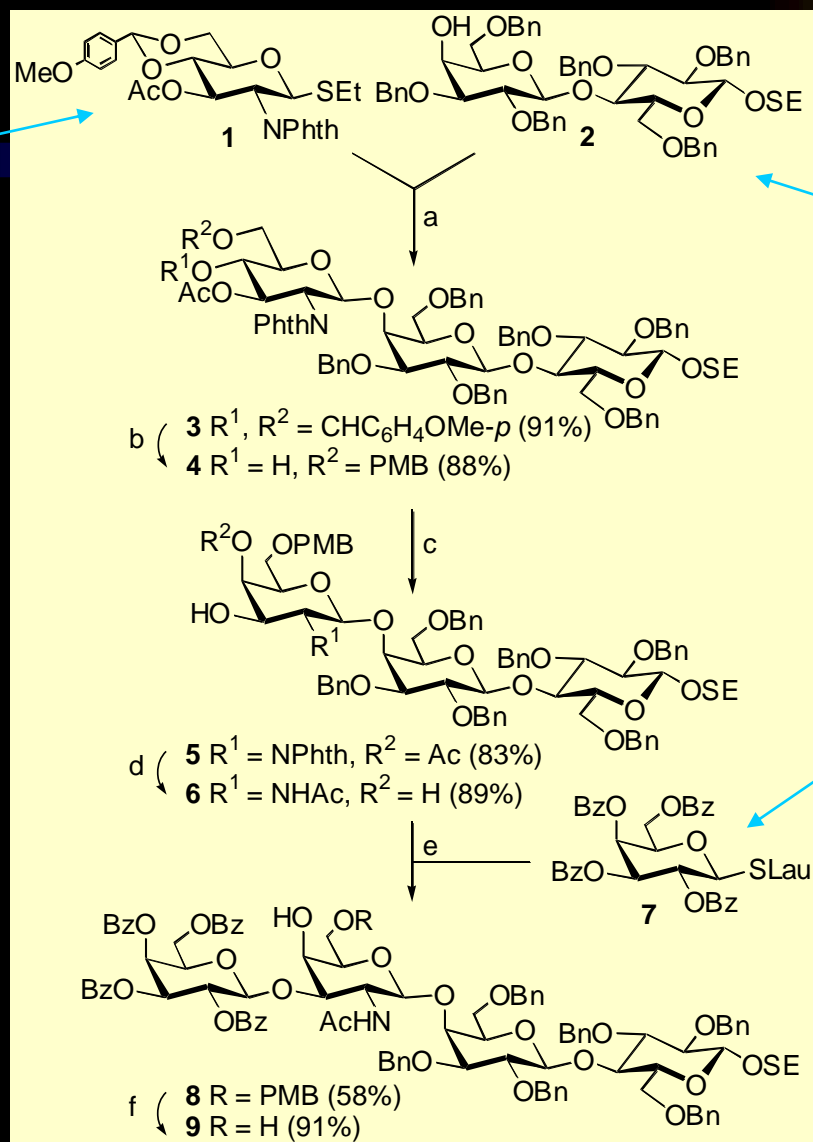
# *Chemical structure of GM1 $\alpha$*



$\alpha$ -Series ganglioside containing a sialyl  $\alpha$ -2 $\rightarrow$ 6-*N*-acetylglucosamine residue are exclusively localized on cholinergic neurons.

# Chemical synthesis of GM1 $\alpha$ (1)

6 steps from  
glucosamine

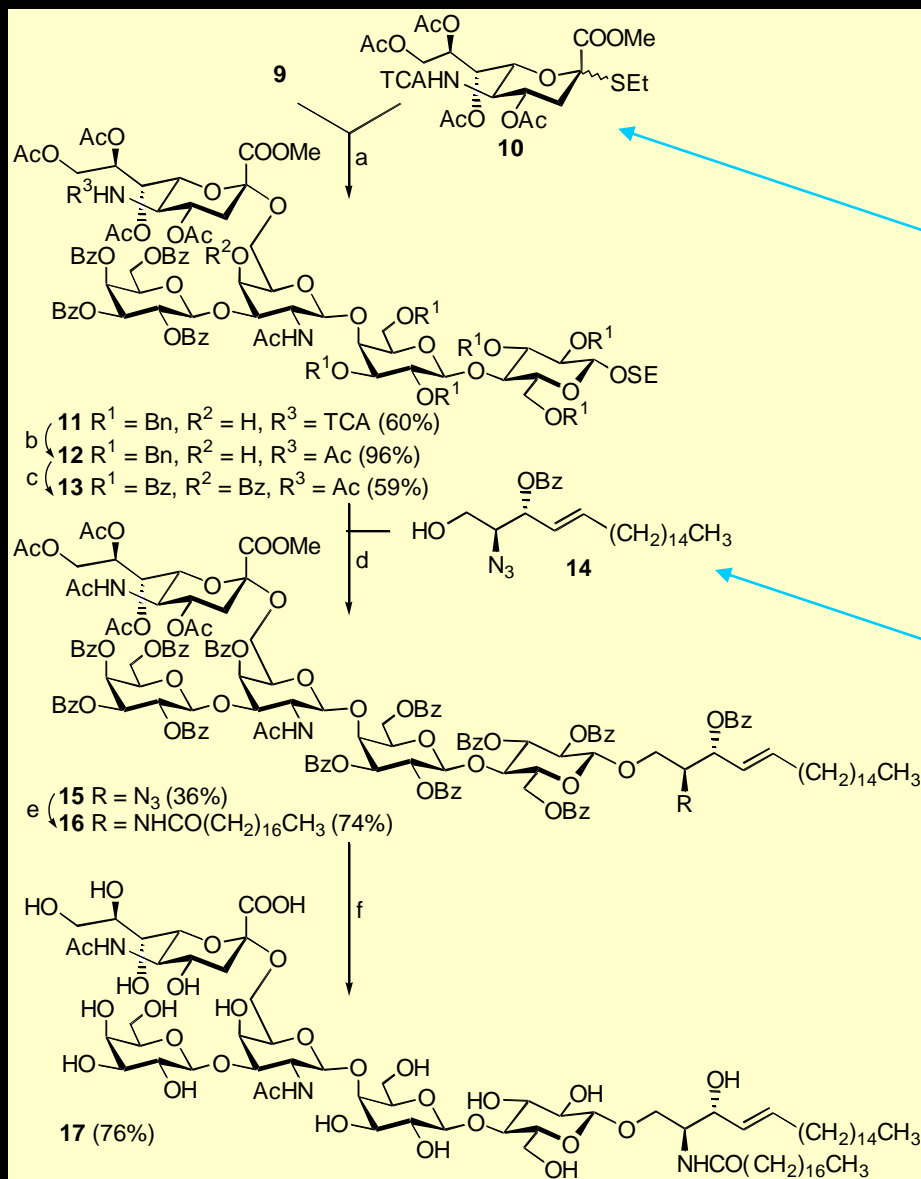


7 steps from lactose

4 steps from galactose

23 steps

# Chemical synthesis of GM1a (2)



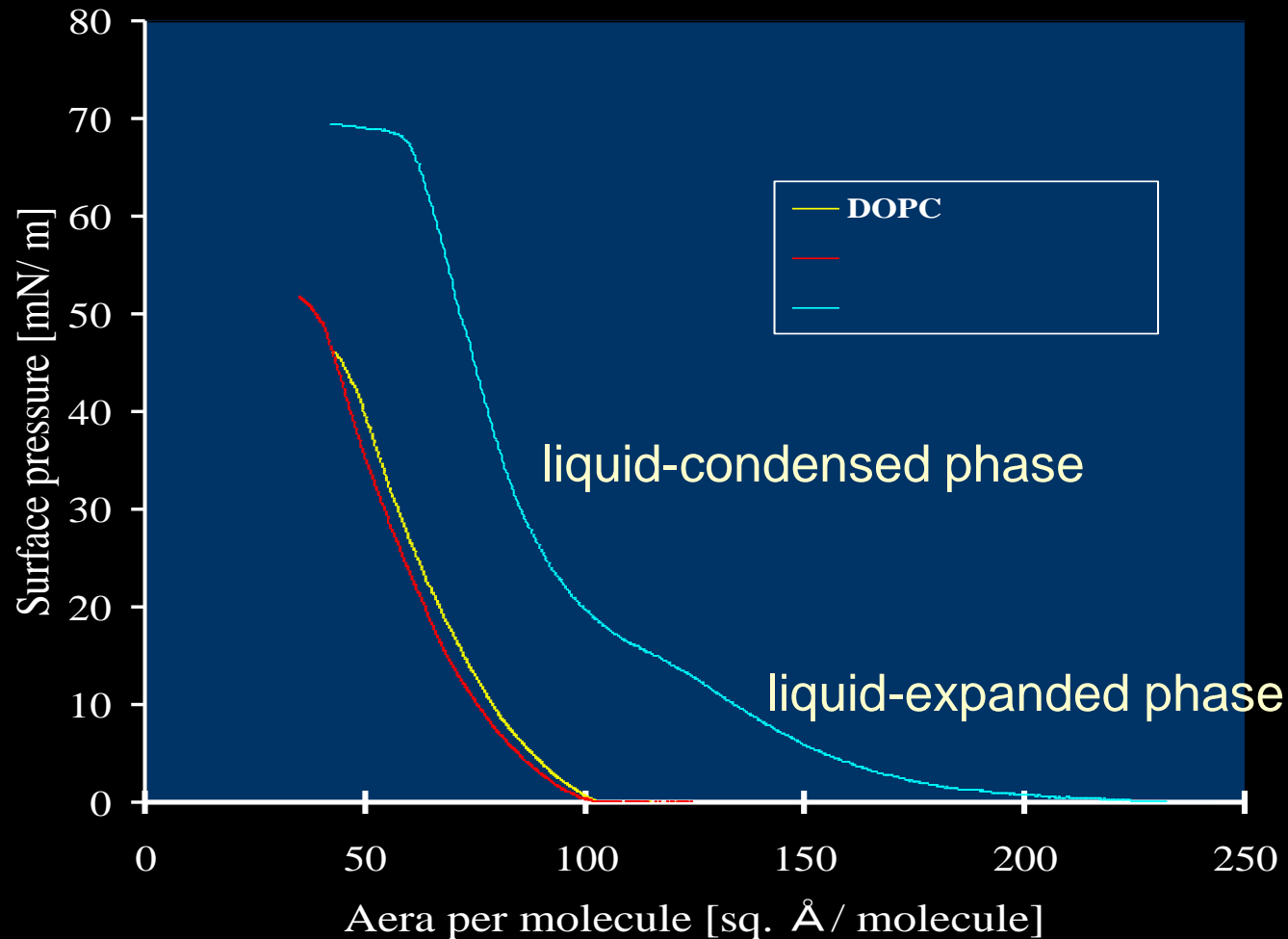
9 steps from glucosamine

7 steps from galactose

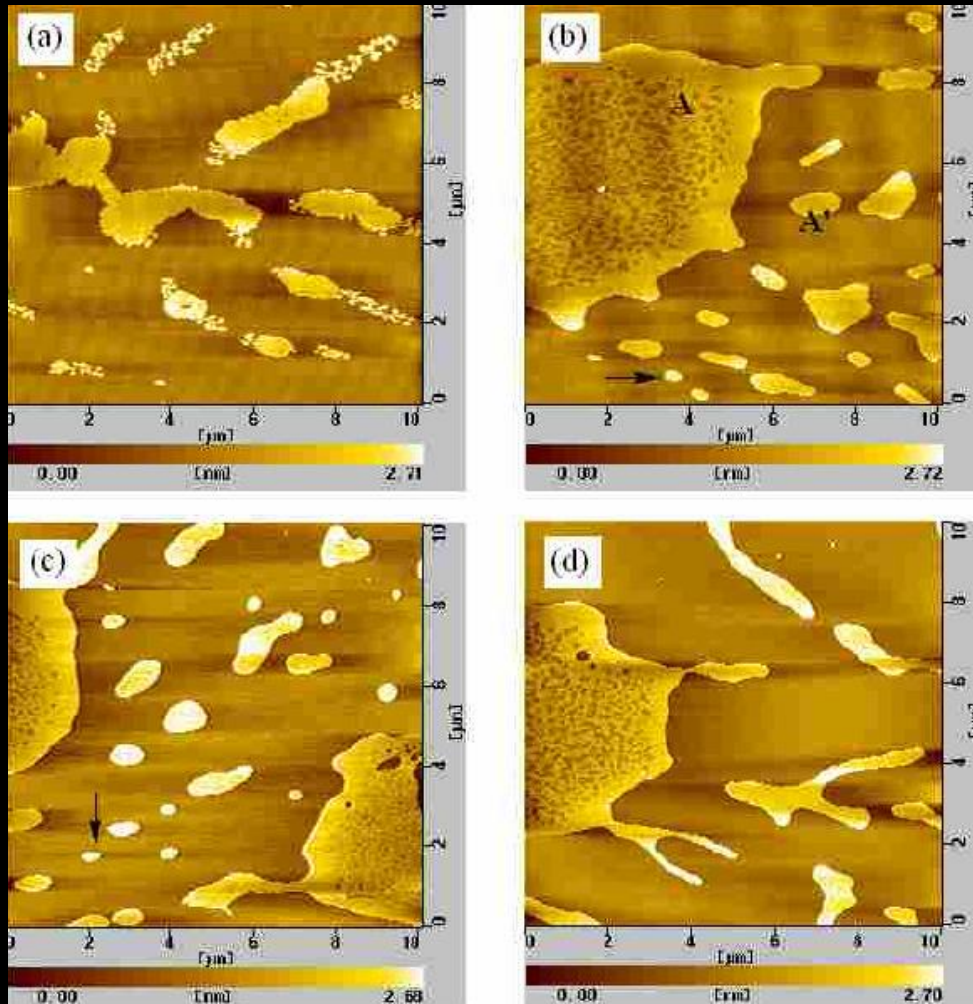
Total steps: 45

(33 steps were known in publication.)

# Surface pressure vs. area per molecule isotherms for GM1a/DOPC/DPPC(1:9:9)



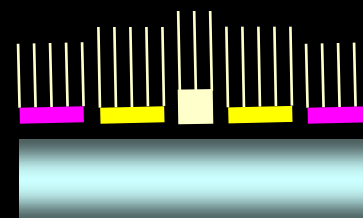
# AFM images for GM1 $\alpha$ /DOPC/DPPC (1:9:9) monolayers



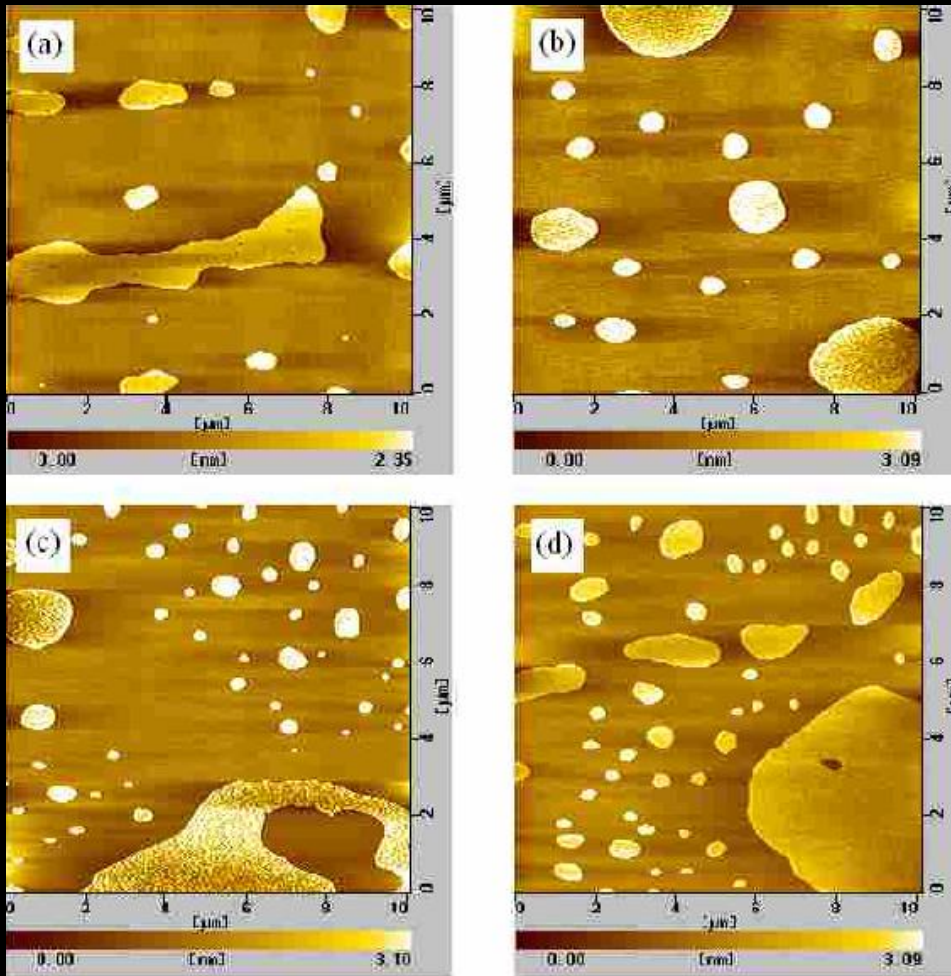
- a) 15 mN/m
- b) 25 mN/m
- c) 30 mN/m
- d) 40 mN/m

A, A': DPPC-rich domains  
→: GM1 $\alpha$ -raft

Subphase: 2.0mM NaCl

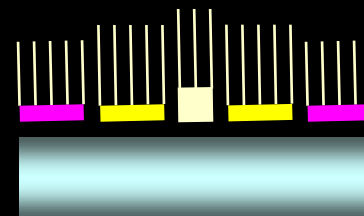


# *AFM images for GM1 $\alpha$ /DOPC/DPPC (1:9:9) monolayers on physiological saline*

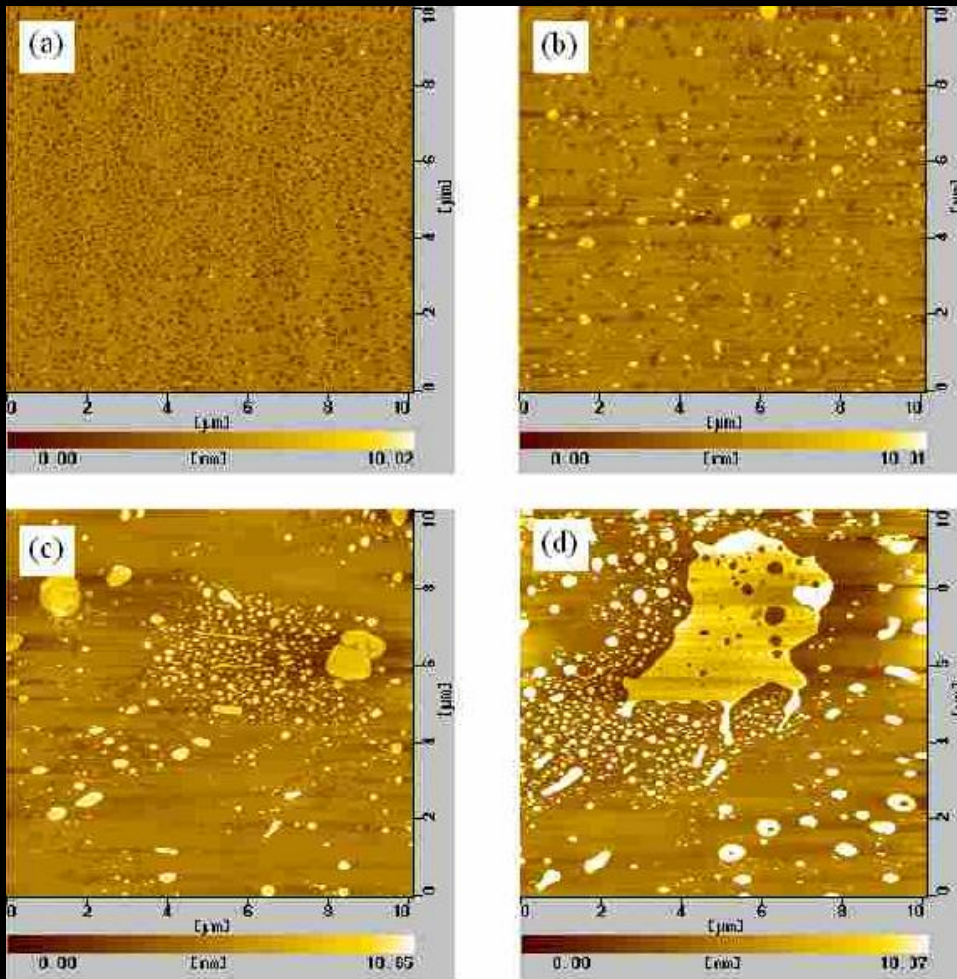


- a) 15 mN/m
- b) 25 mN/m
- c) 30 mN/m
- d) 40 mN/m

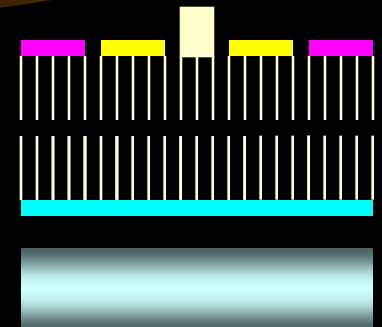
Subphase: 157 mM NaCl



# AFM images for hybrid bilayers of GM1 $\alpha$ /DOPC/DPPC (1:9:9)



- a) 15 mN/m
- b) 20 mN/m
- c) 25 mN/m
- d) 30 mN/m



First layer: DPPE  
Subphase: 157 mM NaCl

Second layer:  
GM1 $\alpha$ :DOPC:DPPC=1:9:9  
Subphase: 2.0 mM NaCl



# *Self-assembly by surface pressure*



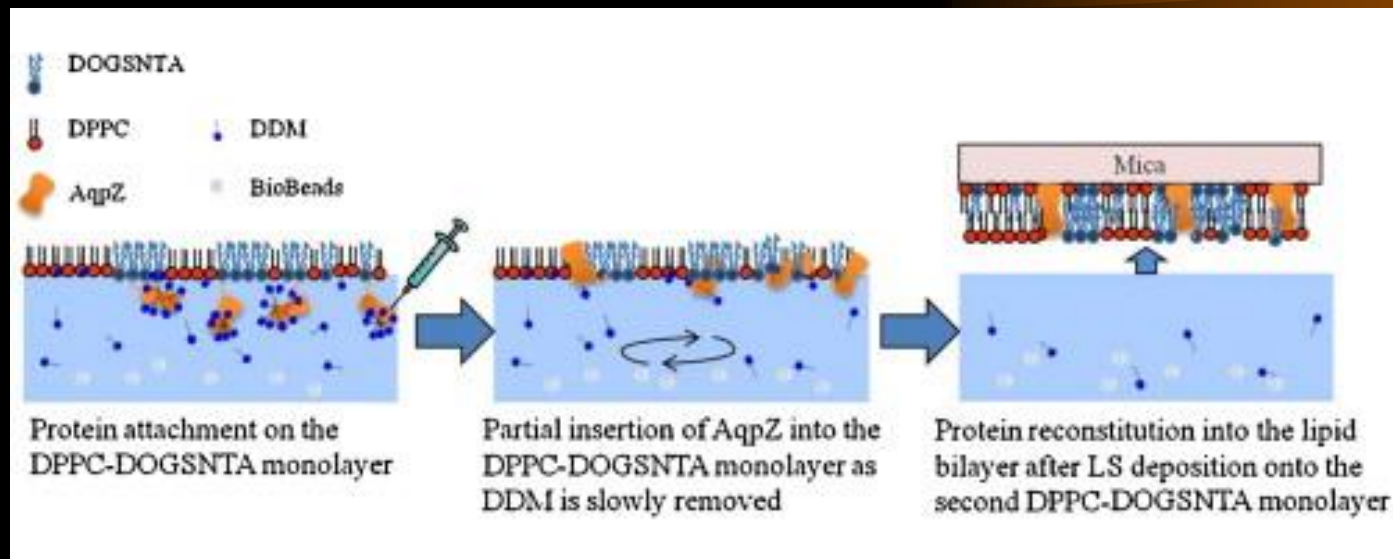
- GM1 $\alpha$ /DOPC/DPPC ternary monolayer and hybrid bilayer obviously shows GM1 $\alpha$ -rafts.
- The raft was associated, separated and associated again, according to increase of surface pressure.
- The association of rafts causes enzyme-localization on signal transduction.



*2. Reconstruction of  
transmembrane protein*

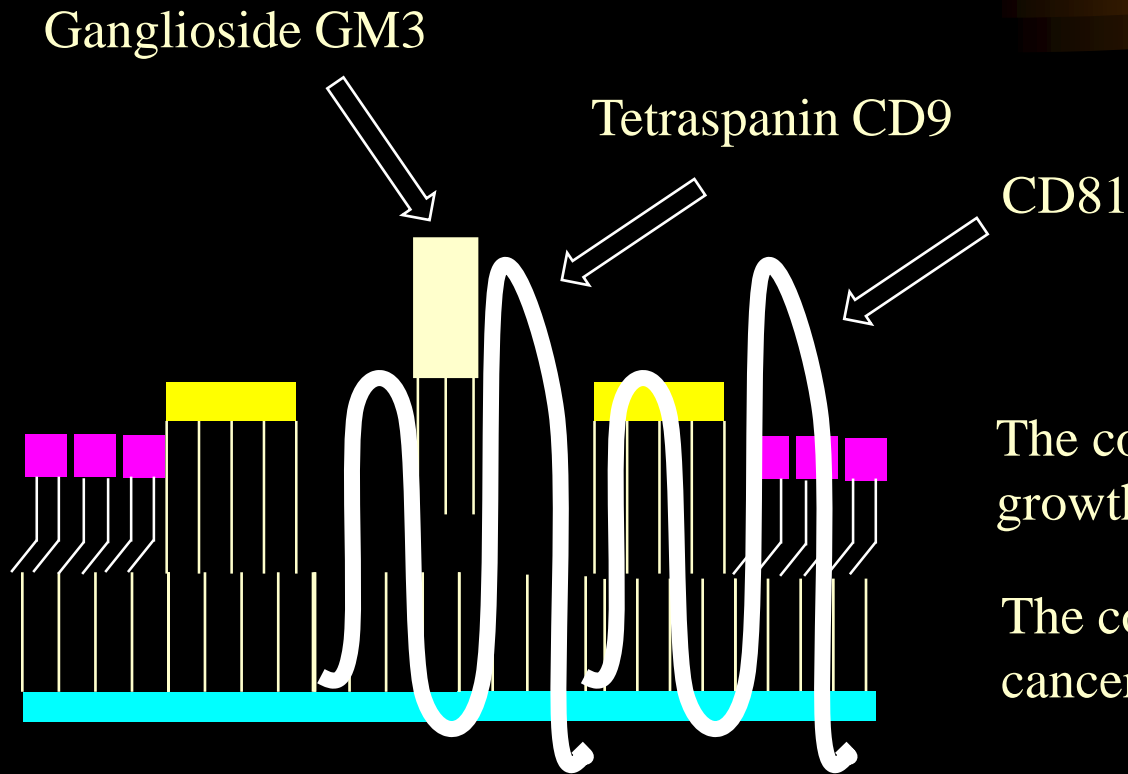
# *A novel method of AquaporinZ incorporation via binary-lipid Langmuir monolayers*

Guofei Suna, Hu Zhou<sup>b</sup>, Yi Li<sup>a</sup>, Kandiah Jeyaseelanc, Arunmozhiarasi Armugam<sup>c</sup>, Tai-Shung Chung, *Colloids and Surfaces B: Biointerfaces* **89** (2012) 283–288.



The incorporation of Histidine-tagged AquaporinZ by Nickel chelating lipids is investigated with Langmuir–Blodgett technology for the first time. Detergent removal by BioBeads in the Langmuir–Blodgett system is studied. AquaporinZ and detergent interaction during protein insertion is elaborated and a protein incorporation mechanism is proposed.

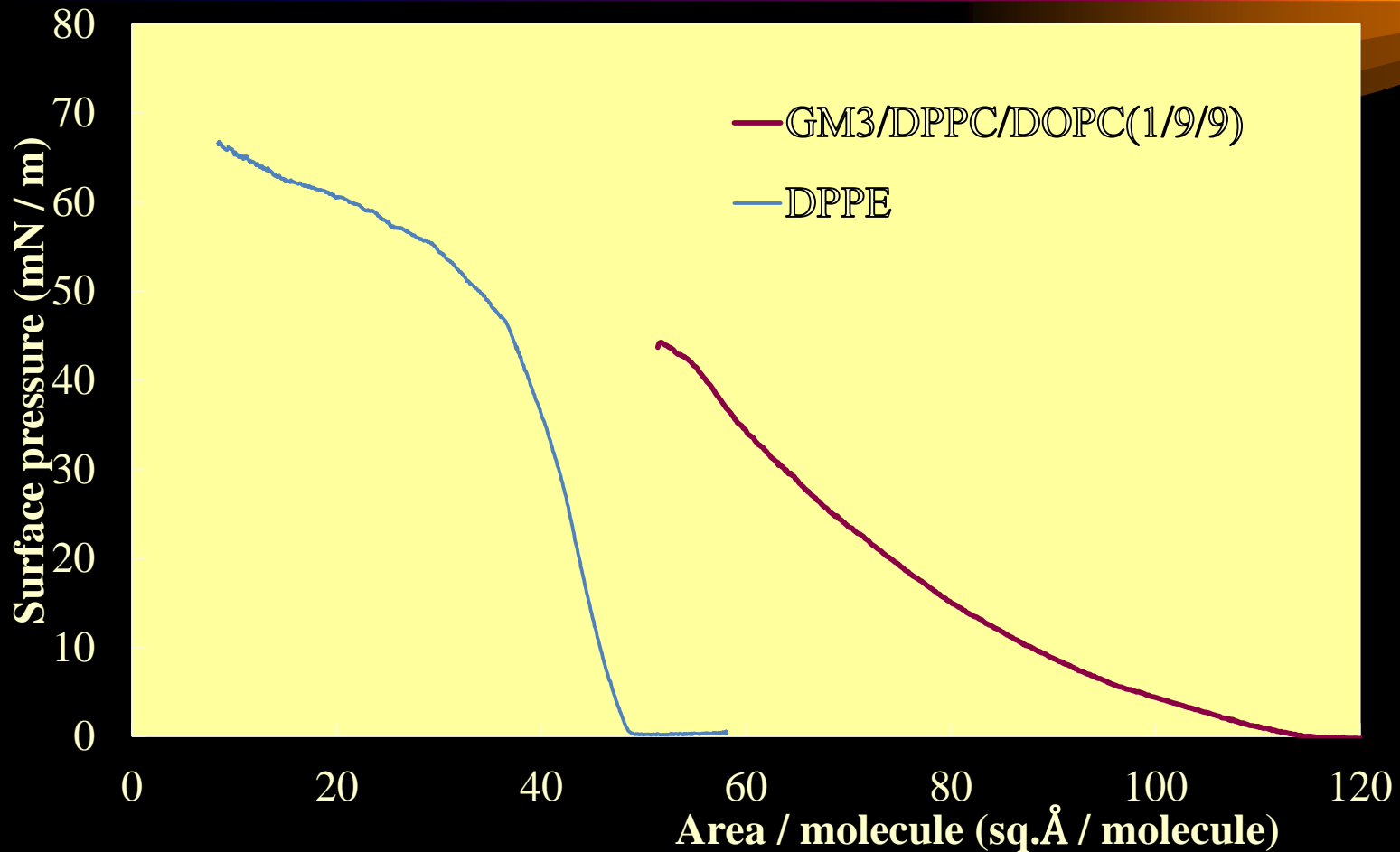
# *GM3/CD9/CD81 complex*



The complex regulates cell growth, proliferation.

The complex regulates also cancer cell proliferation.

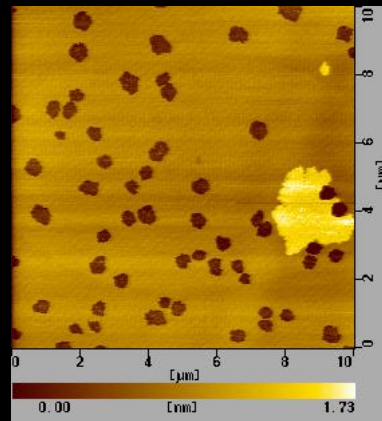
# *Surface pressure vs. area per molecule isotherms for GM3/DPPC/DOPC(1:9:9) ternary monolayer*



# AFM images for GM3/DOPC/DPPC (1:9:9) monolayers

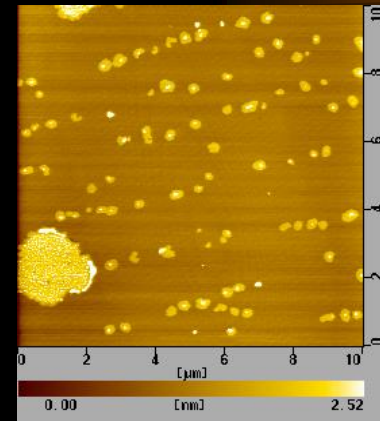
Surface pressure:  
10 mN/m

9 images / 9 points



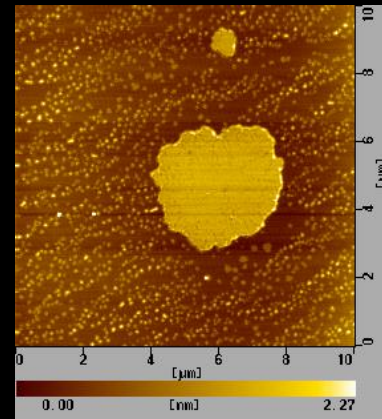
Surface pressure:  
20 mN/m

9 images / 9 points



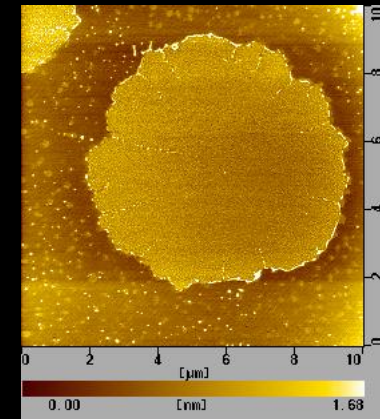
Surface pressure:  
30 mN/m

8 images / 9 points

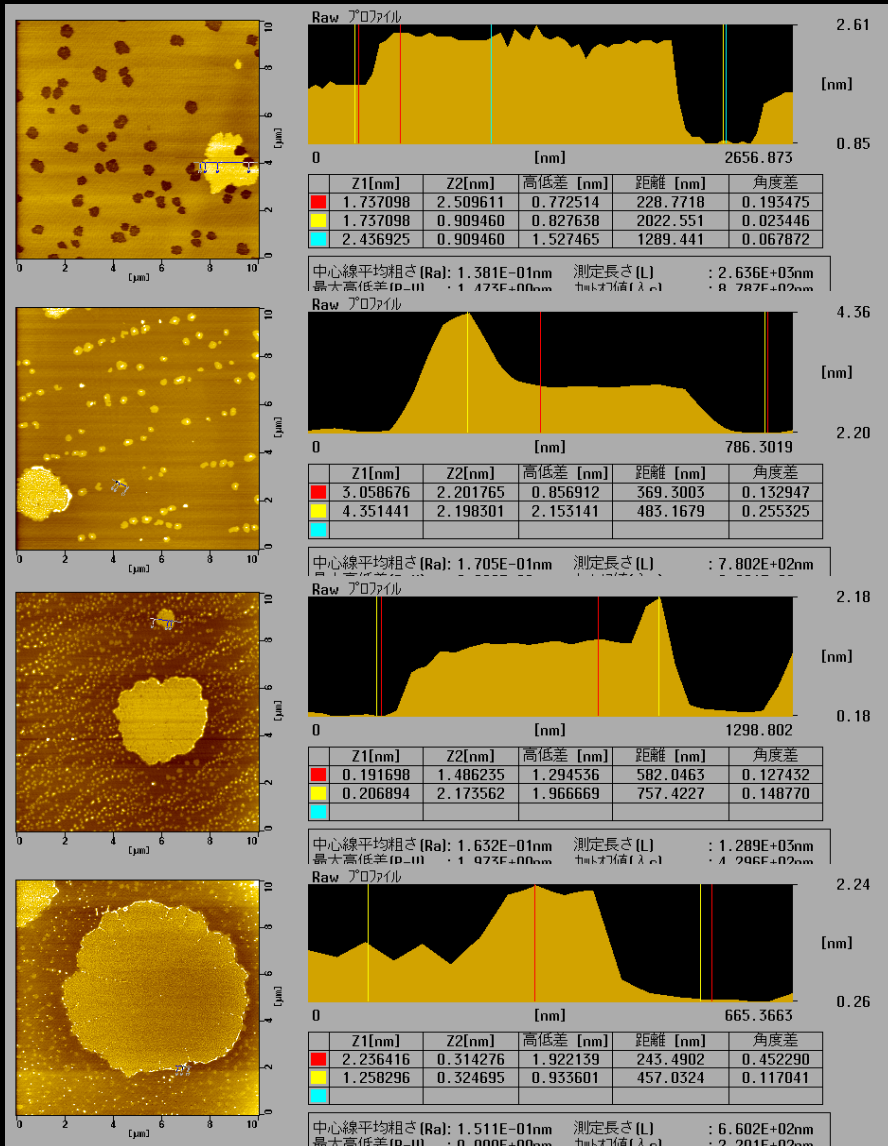


Surface pressure:  
40 mN/m

9 images / 9 points



# Cross-sectional study of rafts



Surface pressure: 10 mN/m

Hight:  $0.9 \pm 0.1$  nm

Surface pressure: 20 mN/m

Hight:  $0.8 \pm 0.1$  nm,  $2.1 \pm 0.2$  nm

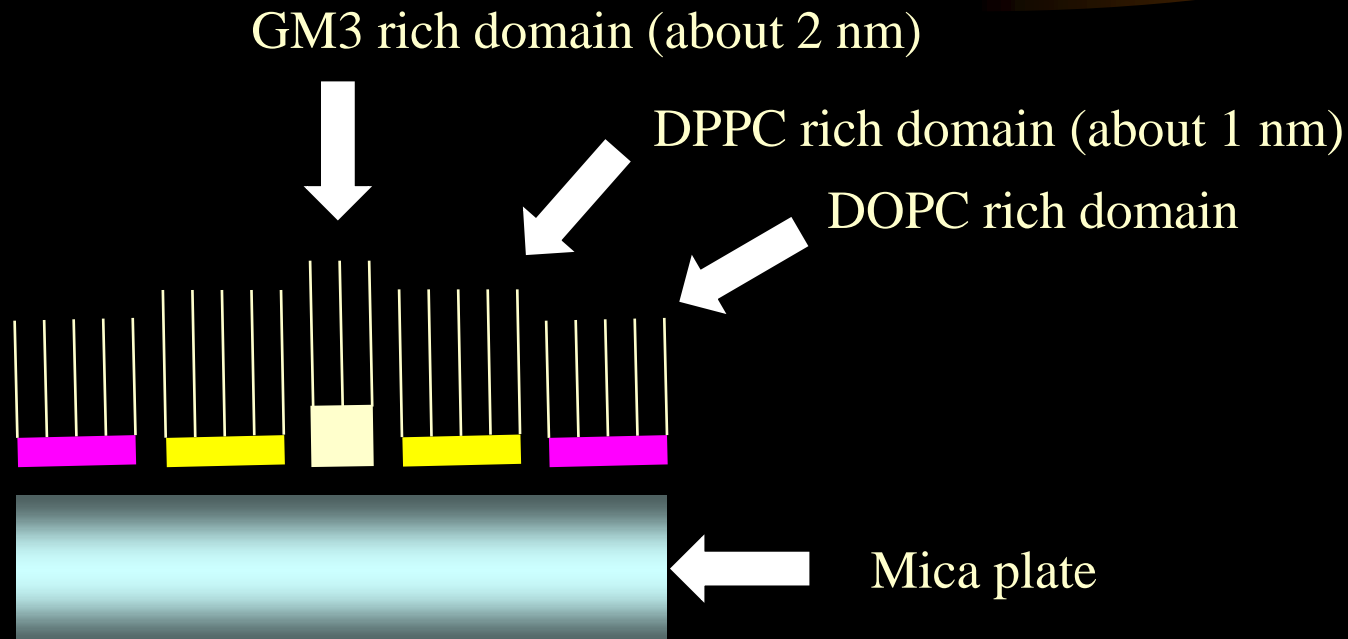
Surface pressure: 30 mN/m

Hight:  $0.5 \pm 0.2$  nm,  $1.7 \pm 0.5$  nm

Surface pressure: 40 mN/m

Hight:  $0.5 \pm 0.2$  nm,  $1.6 \pm 0.3$  nm

# *Proposed structure of ganglioside raft*

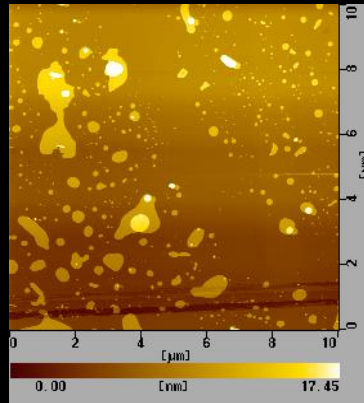




# *AFM images for hybrid bilayers of GM3/DOPC/DPPC (1:9:9)*

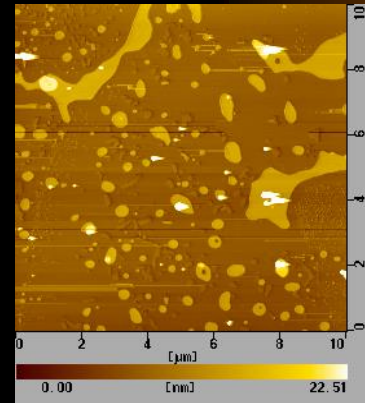
Surface pressure:  
10 mN/m

6 images / 9 points



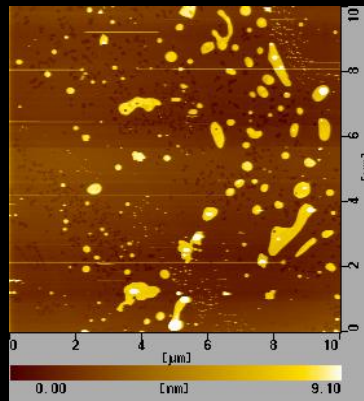
Surface pressure:  
20 mN/m

7 images / 9 points



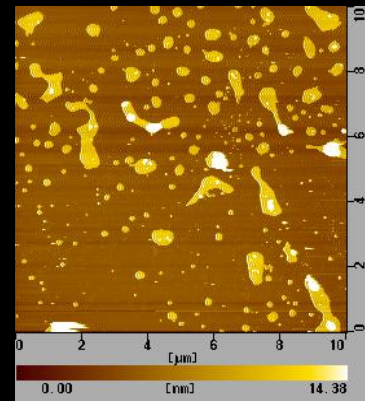
Surface pressure:  
30 mN/m

6 images / 9 points

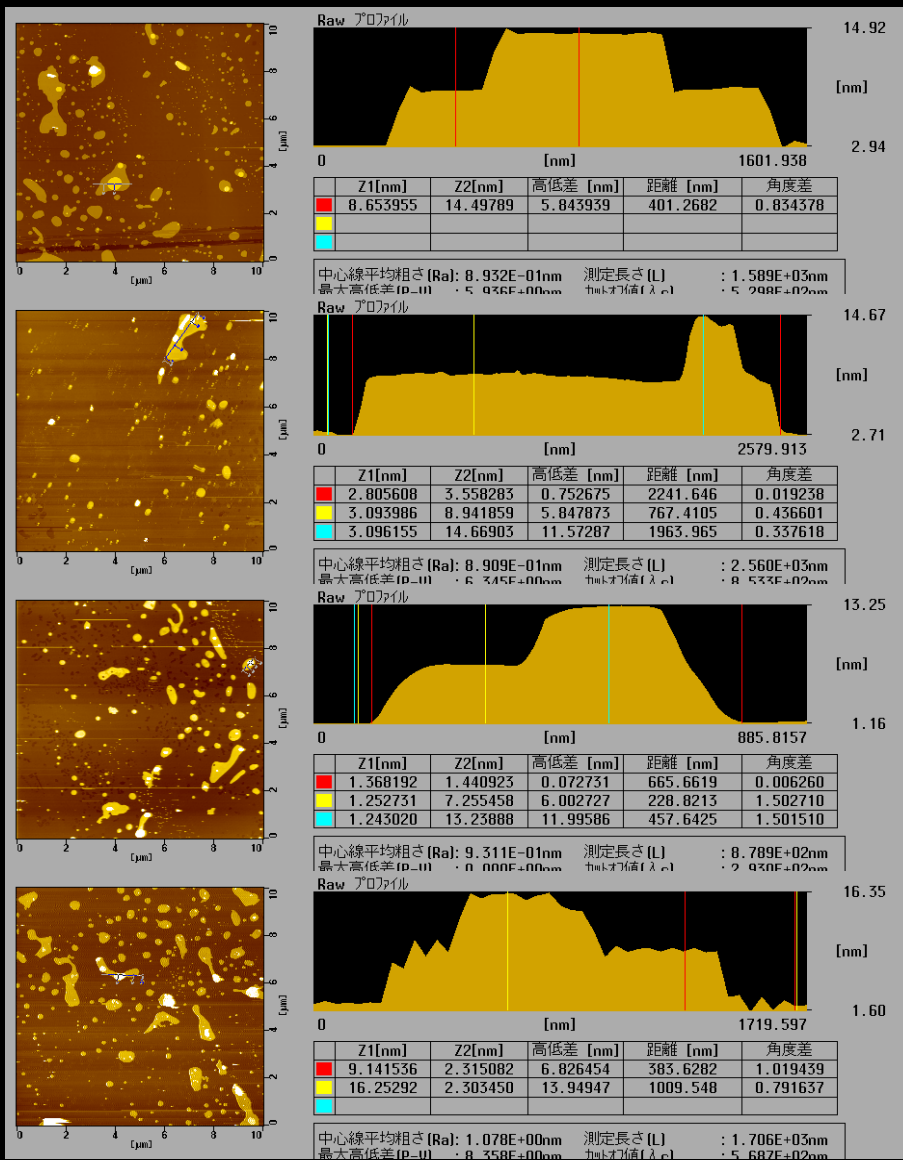


Surface pressure:  
40 mN/m

5 images / 9 points



# Cross-sectional study of rafts



Surface pressure: 10 mN/m

Hight:  $6.3 \pm 0.7$  nm,  $11.5 \pm 0.4$  nm

Surface pressure: 20 mN/m

Hight:  $6.7 \pm 0.6$  nm,  $12.5 \pm 0.2$  nm

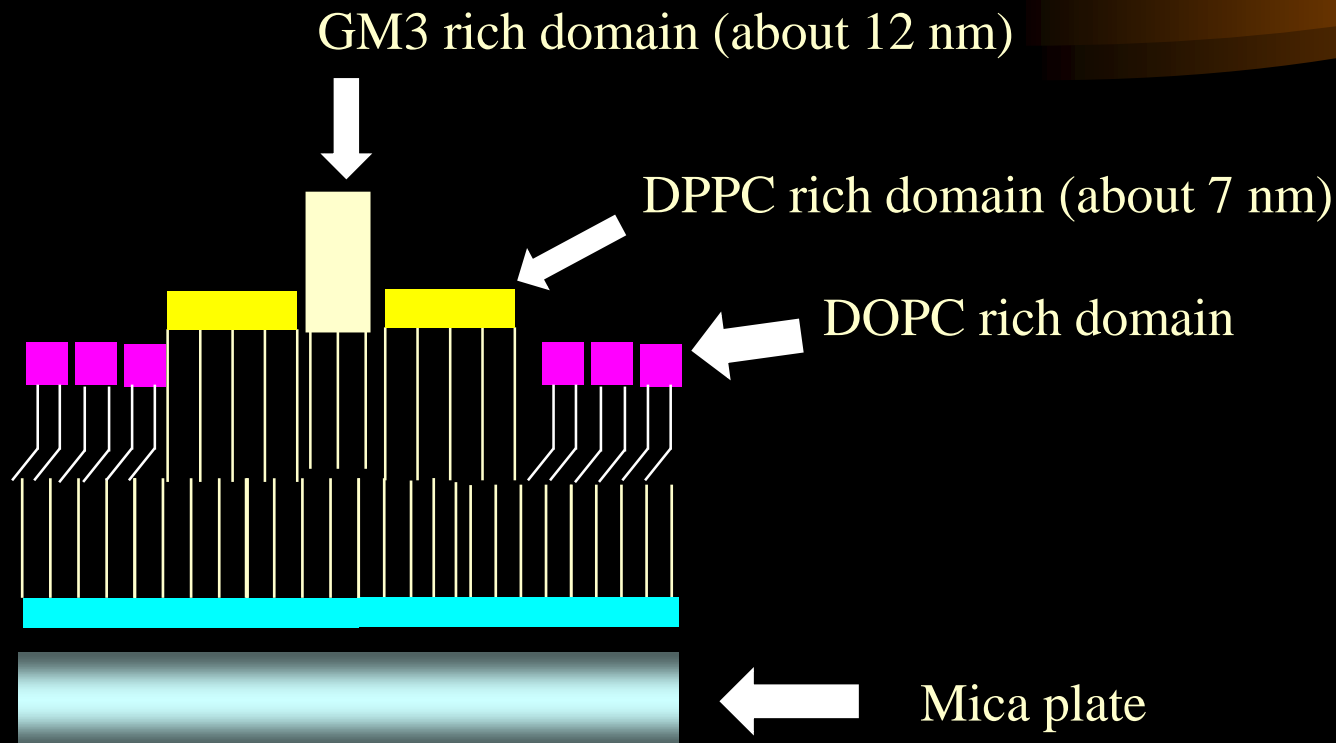
Surface pressure: 30 mN/m

Hight:  $6.2 \pm 0.4$  nm,  $11.7 \pm 0.5$  nm

Surface pressure: 40 mN/m

Hight:  $6.5 \pm 0.5$  nm,  $11.2 \pm 0.3$  nm

# *Proposed structure of ganglioside raft*

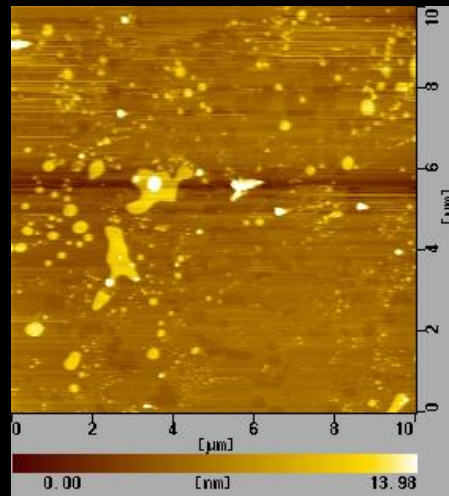


# *Reconstruction of tetraspanin (CD9) into lipid bilayer*

The hybrid bilayer was kept in CD9 solution (2  $\mu\text{g}/\text{ml}$ ) for 5 min.

Surface pressure:  
30 mN/m

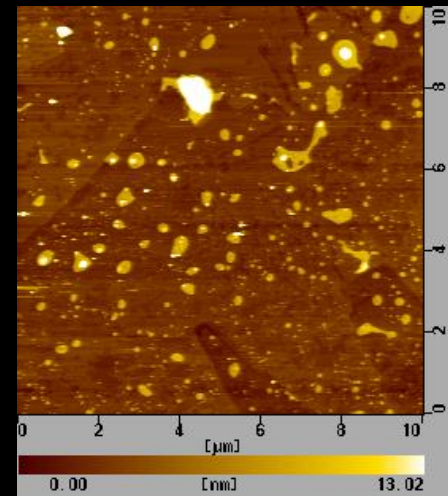
5 images / 9 points  
were changed.



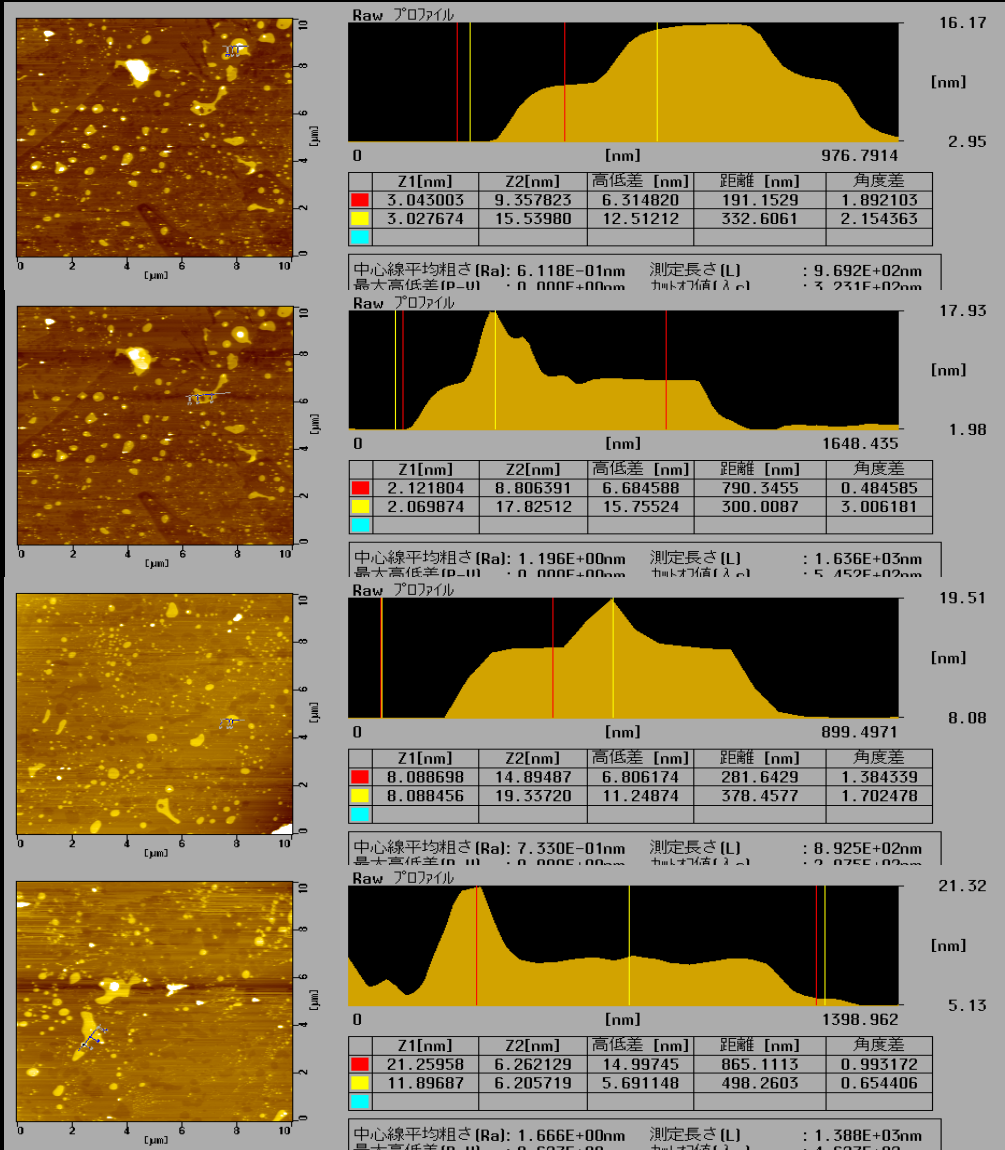
Surface pressure:  
40 mN/m

2 images / 9 points  
were no change.

7 images / 9 points  
were changed.



# Cross-sectional study of rafts reconstructed by tetraspanin



Surface pressure: 30 mN/m

Height:  $6.1 \pm 0.5$  nm,  $11.7 \pm 1.1$  nm

Surface pressure: 30 mN/m

Height:  $6.1 \pm 0.5$  nm,  $16.4 \pm 0.9$  nm

Surface pressure: 40 mN/m, 2 / 9

Height:  $6.4 \pm 0.5$  nm,  $11.3 \pm 0.8$  nm

Surface pressure: 40 mN/m, 7 / 9

Height:  $6.4 \pm 0.5$  nm,  $15.6 \pm 1.7$  nm

# *Proposed structure of ganglioside raft with tetraspanin*

GM3 rich domain (about 12 nm) + tetraspanin (about 17 nm)

