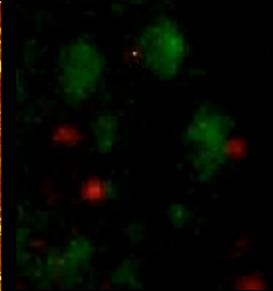
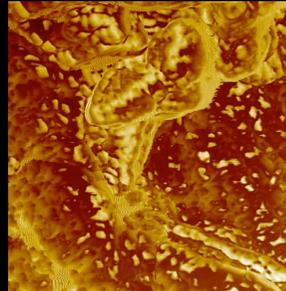
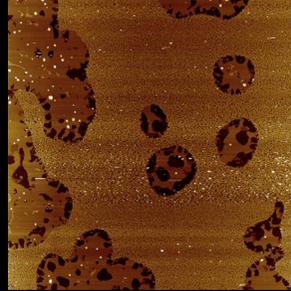
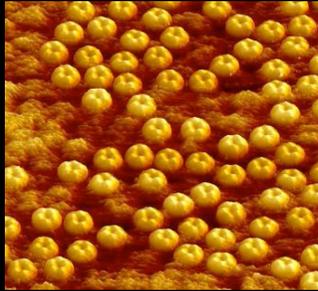


Centre de Biochimie Structurale
U1054 INSERM, UMR5048 CNRS, Université Montpellier



Summer School of AFM, Marcoule, 2011

Membrane Protein Reconstitution within Lipid Bilayer

Francesca Gubellini
Manuela Dezi
Daniel Levy



UMR168
CNRS 
institut **Curie**
Ensemble, prenons le cancer de vitesse.

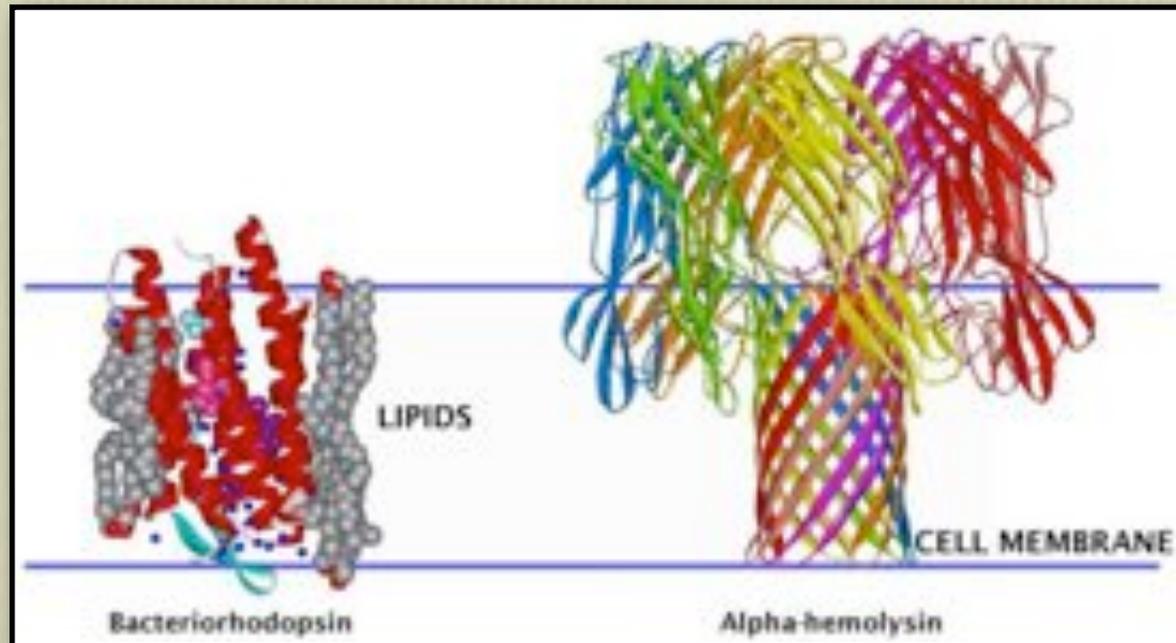


Outlines

- ▶ **Membrane proteins and cell functions**
- ▶ **Reconstitution of membrane proteins**
- ▶ **High resolution AFM imaging of transmembrane proteins reconstituted into artificial membranes**
 - ▶ **Direct incorporation**
 - ▶ **Lipid layer technique**
- ▶ **Other methods**

25% of the human genome encode transmembrane proteins

Target of 70% of the commercially available drugs



Alpha helix
(GPCR)

Beta barrels
(porin)



25% of the human genome encode transmembrane proteins
Target of 70% of the commercially available drugs

The Nobel Prize in Chemistry 1988 was awarded jointly to **Johann Deisenhofer, Robert Huber and Hartmut Michel** *"for the determination of the three-dimensional structure of a photosynthetic reaction centre"*.

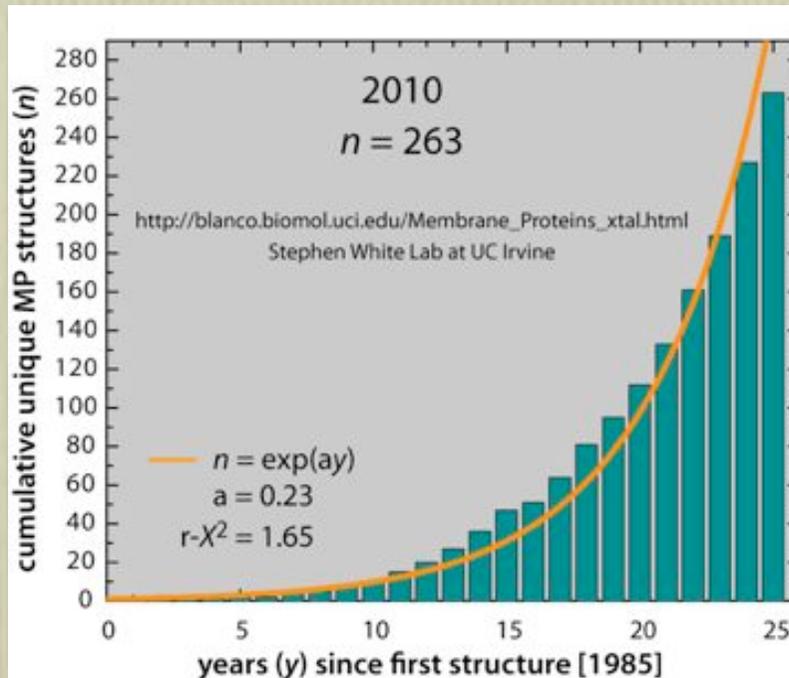
The Nobel Prize in Chemistry 1997 was divided, one half jointly to **Paul D. Boyer and John E. Walker** *"for their elucidation of the enzymatic mechanism underlying the synthesis of adenosine triphosphate (ATP)"* and the other half to Jens C. Skou *"for the first discovery of an ion-transporting enzyme, Na⁺, K⁺ -ATPase"*.

The Nobel Prize in Chemistry 2003 was awarded *"for discoveries concerning channels in cell membranes"* jointly with one half to **Peter Agre** *"for the discovery of water channels"* and with one half to **Roderick MacKinnon** *"for structural and mechanistic studies of ion channels"*.



Number of structures available in the pdb (protein data bank)

236 structures available in the PDB (~ 25 eukaryotic)



Monotopique

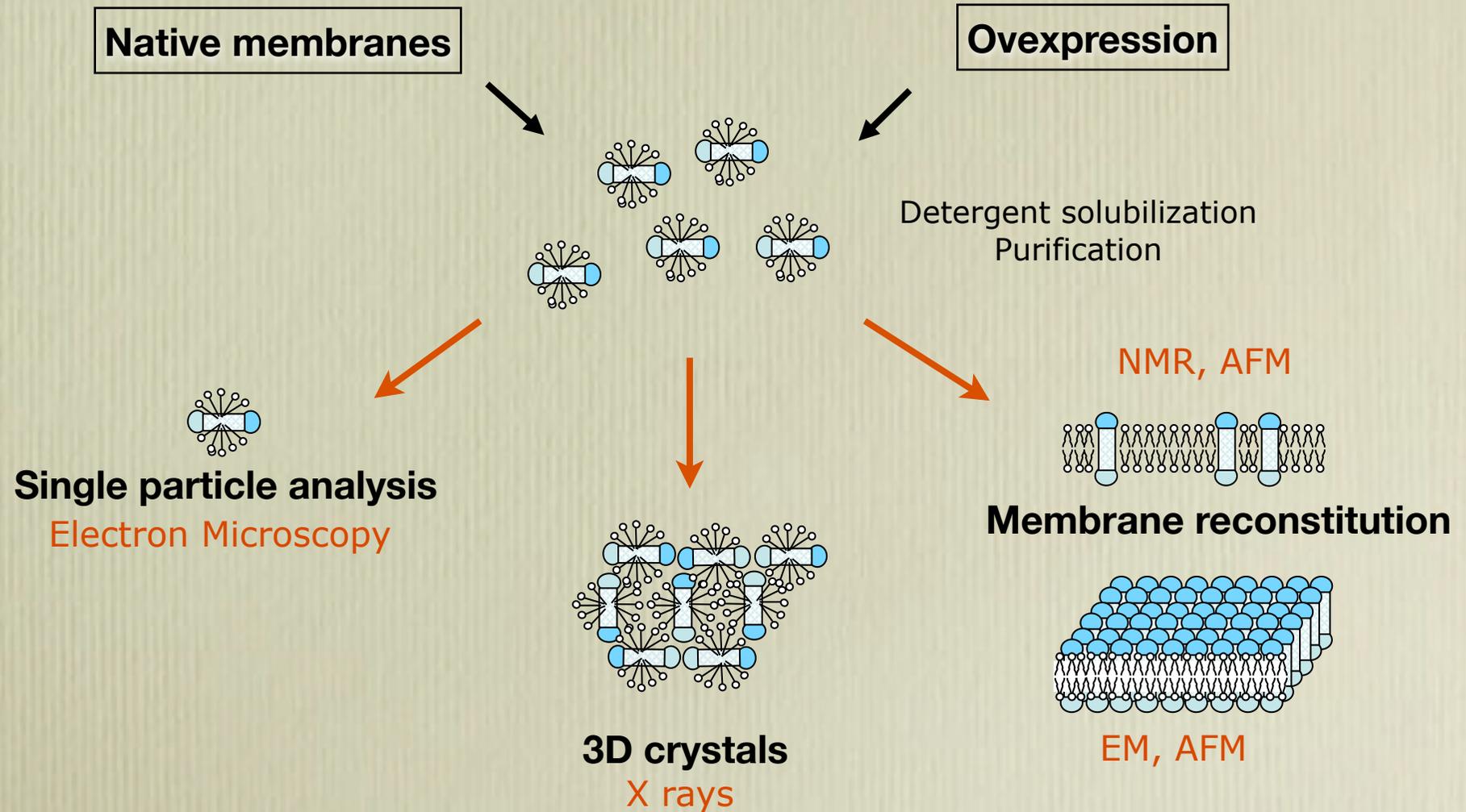
Prostaglandin H2 synthase-1
Cyclooxygenase-2
Monoamine Oxidase

Polytopique

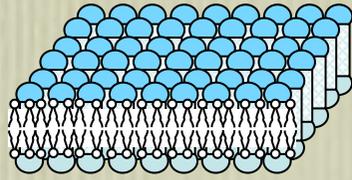
GPCR (Rhodopsin, $\beta 2$ adrenergic receptor)
Beta Barrel Membrane Proteins
Channels (ions, H_2O , NH_3)
Transporters (multidrug efflux, aa, ABC,...)
Proteins from photosynthetic complexes
ATPase
Respiratory proteins
Oxidases
Cytochrome

Mainly purified from biological membranes

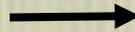
Classical strategies to solve the structure of transmembrane proteins



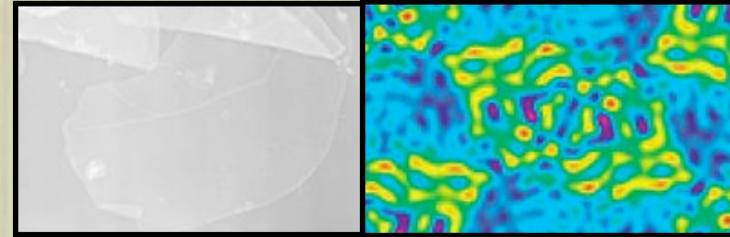
Structural analysis of reconstituted transmembrane proteins



2D crystals

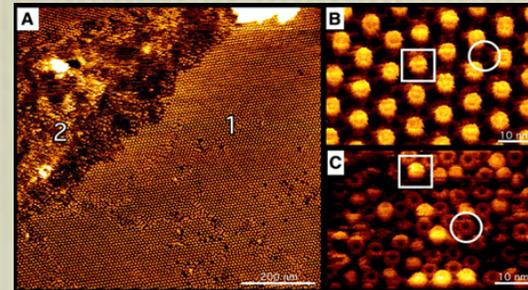


Electron Crystallography

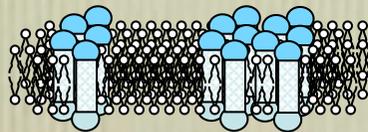


Chloride channel, J.A. Mindell et al, Nature (2001) 409, 219

Atomic Force Microscopy



Na⁺-ATP synthase S. Scheuring, Curie



High density

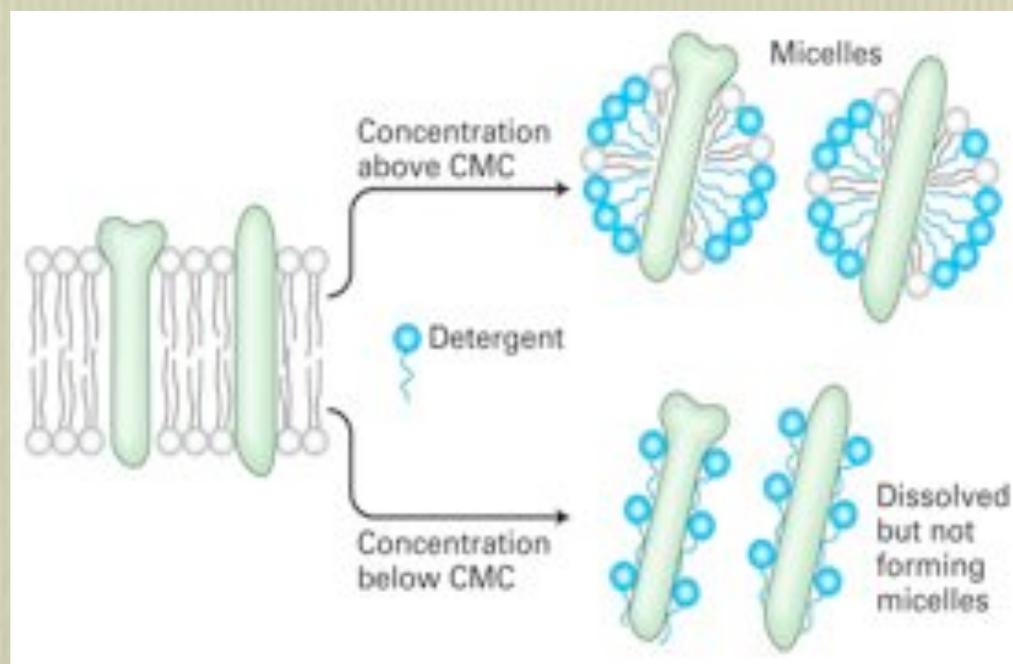


Atomic Force Microscopy ?



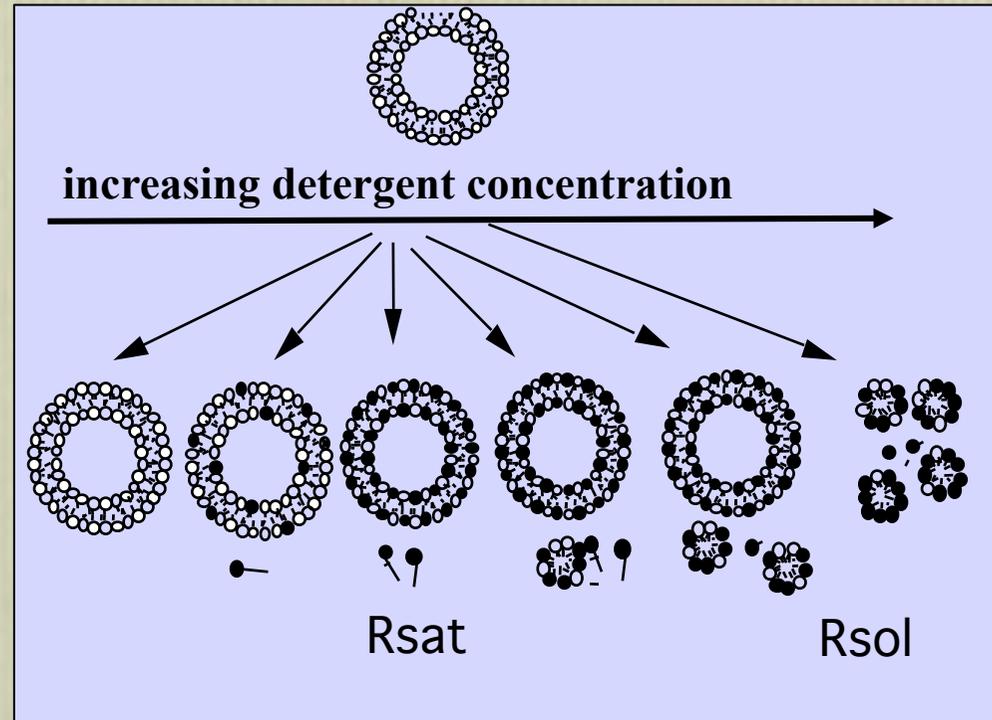
- Developing new strategies to escape the main critical steps
- Imaging under physiological conditions

Effect of detergents



Effect of detergents on lipid assemblies

(Lichtenberg, 1983; Almog, 1986; Ollivon, 1988; Paternostre, 1988;
Da Graça Miguel, 1989; Vinson, 1989...)



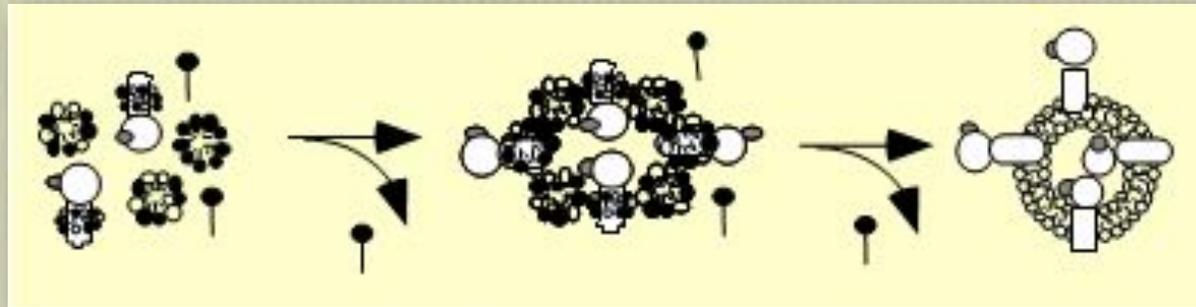
$$D_{total} = D_{water} + R_{eff}[Lip]$$

$R_{eff} = [det]/[Lip]$ in mixed aggregates

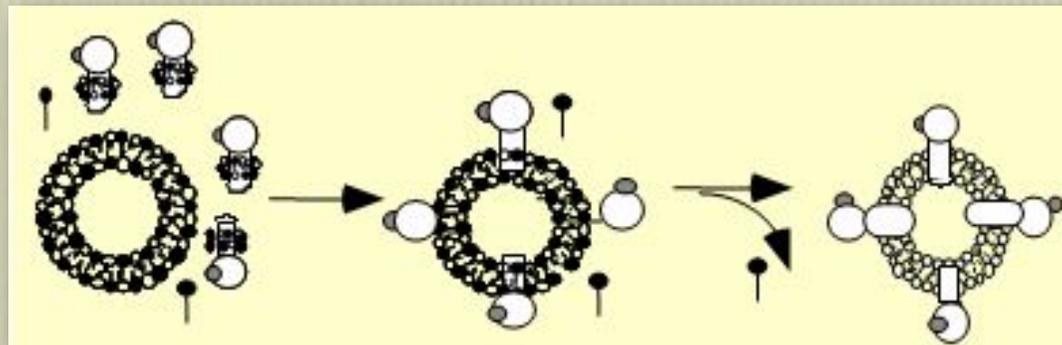
$$D_{sat} = cmc + R_{sat} \cdot [Lipide]$$

$$D_{sol} = cmc + R_{sol} \cdot [Lipide]$$

Direct incorporation of protein into detergent-saturated liposomes



Reconstitution from fully solubilized samples

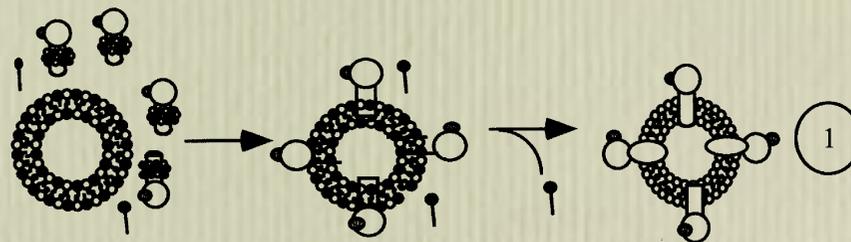


Direct incorporation in glycosylated detergent-destabilized liposomes

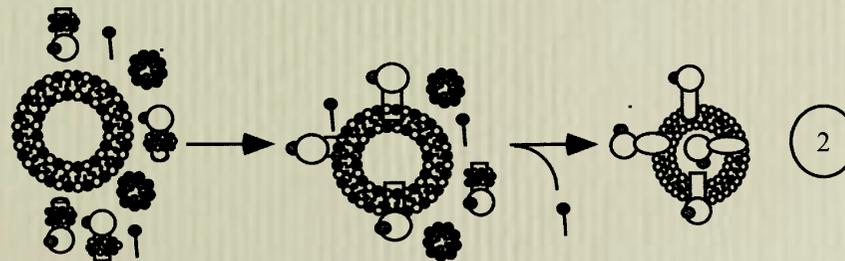
Unique orientation

Direct Incorporation of transmembrane proteins within supported lipid bilayers

The detergent induces the mechanism of incorporation



OG, DDM

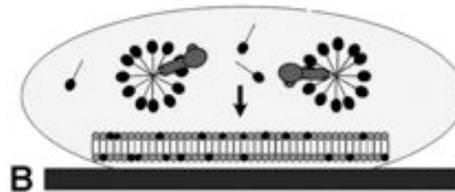
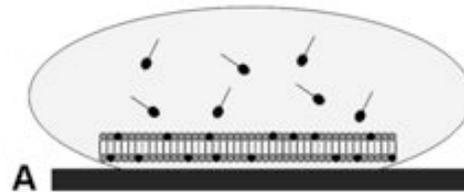


Triton X100

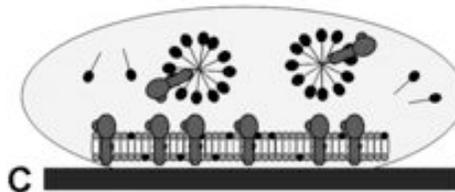


Direct Incorporation of transmembrane proteins within supported lipid bilayers

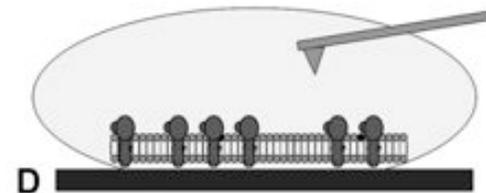
①



②



③

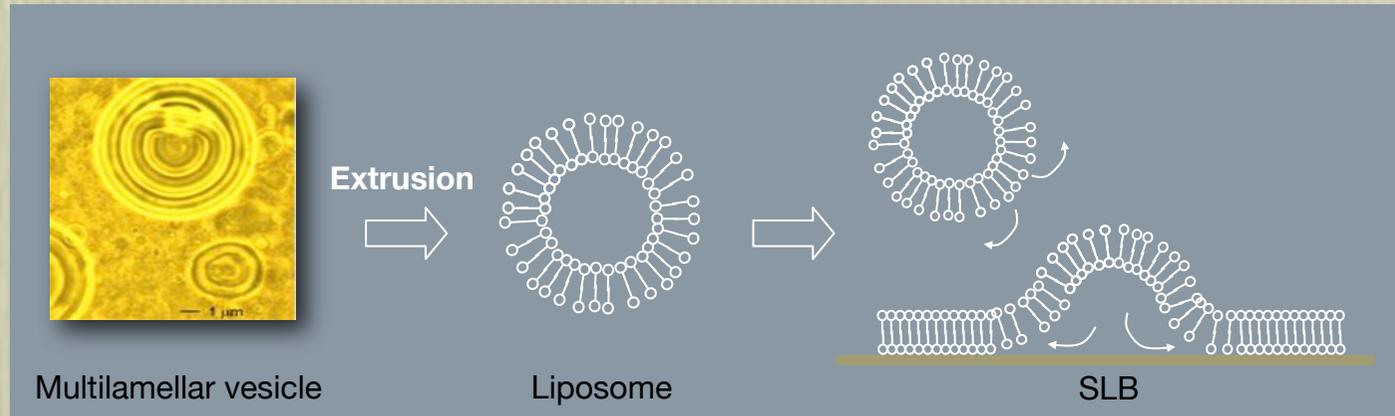


SLB destabilization
[detergent] \sim cmc

Incorporation

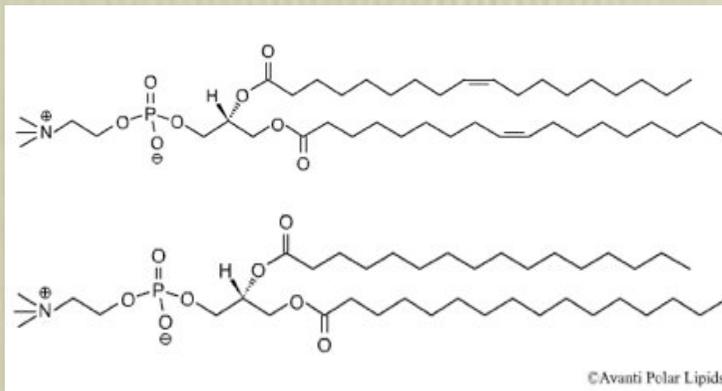
AFM Imaging
of non crystalline proteins

Direct Incorporation of transmembrane proteins within supported lipid bilayers



DOPC
 $T_m = -20^\circ\text{C}$

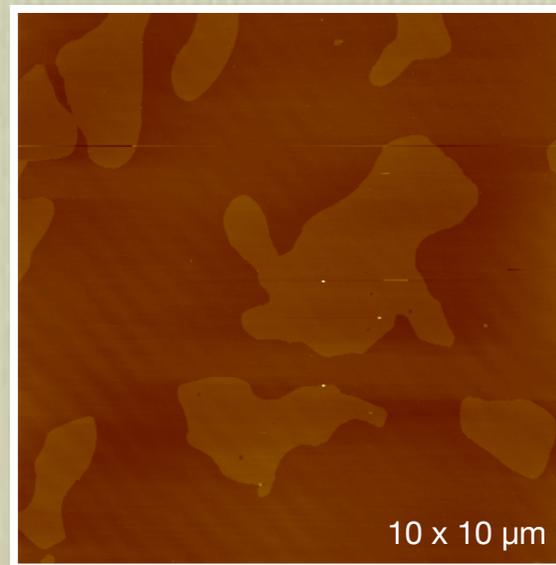
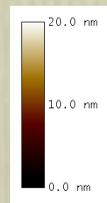
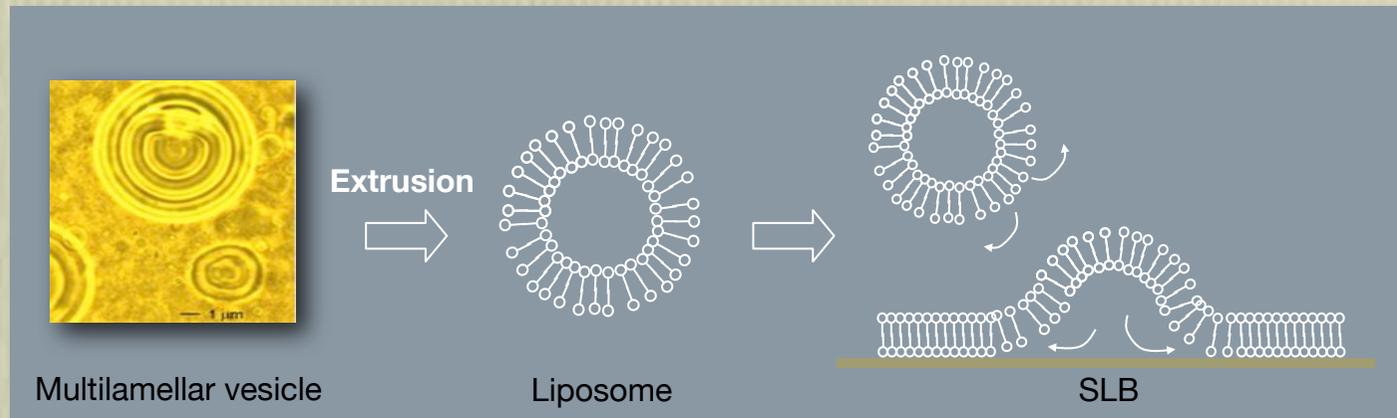
DPPC
 $T_m = 41^\circ\text{C}$



Fusion in PBS buffer
(20 mM phosphate, NaCl 150 mM, pH 7.4)
2 h at 65°C

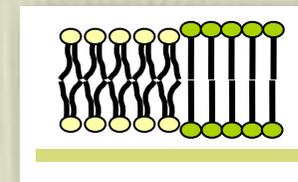


Direct Incorporation of transmembrane proteins within supported lipid bilayers



Contact mode

DOPC/DPPC

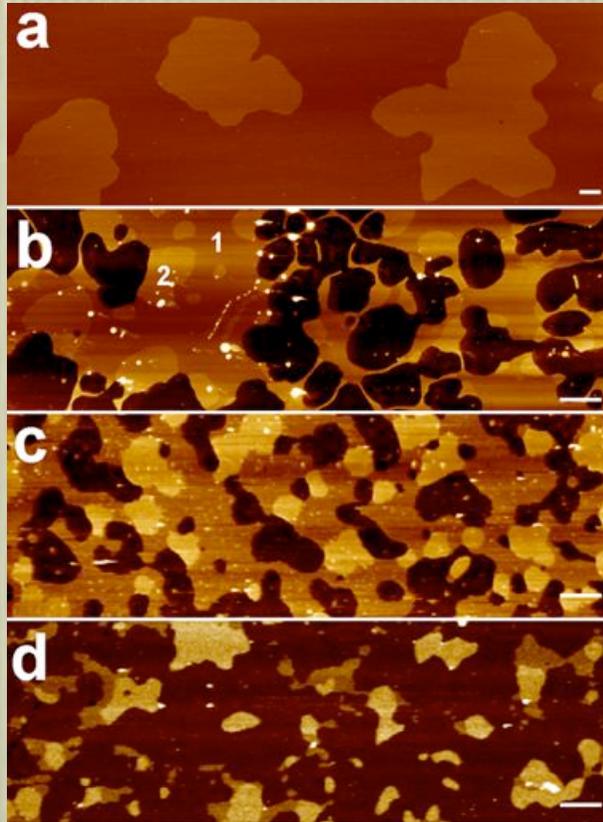


No leaflet decoupling
 $D = 2-5 \mu\text{m}^2/\text{s}$

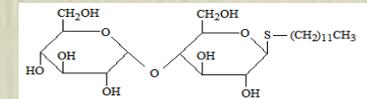
Imaging setup in liquid

Silicon nitride tip 100 mN/m
Applied force < 100 pN

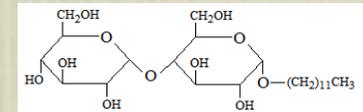
Membrane destabilization by detergents



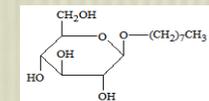
n-Dodecyl- β -D-Thiomaltopyranoside (DOTM)
cmc = 0.05 mM



n-Dodecyl- β -D-Maltopyranoside (DDM)
cmc = 0.17 mM



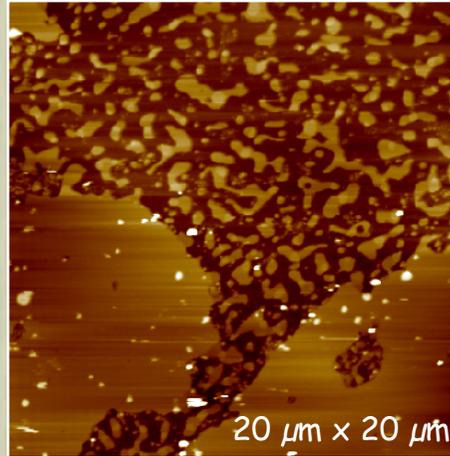
Octyl- β -D-glucopyranoside (OG)
cmc = 17 mM



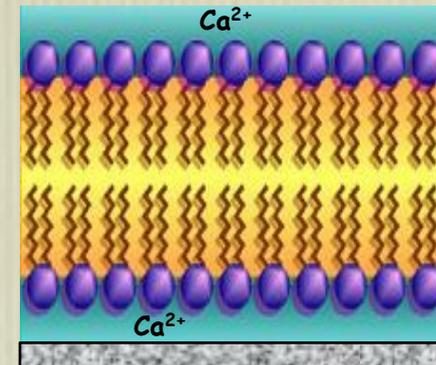
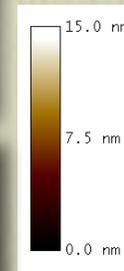
- SLB are stable above the cmc with low but not with high cmc detergent
- Planar lipid membrane are more resistant than liposomes
- Both gel and fluid phases are preserved

Membrane destabilization by detergents

0.2 mM DDM



5 mM CaCl₂



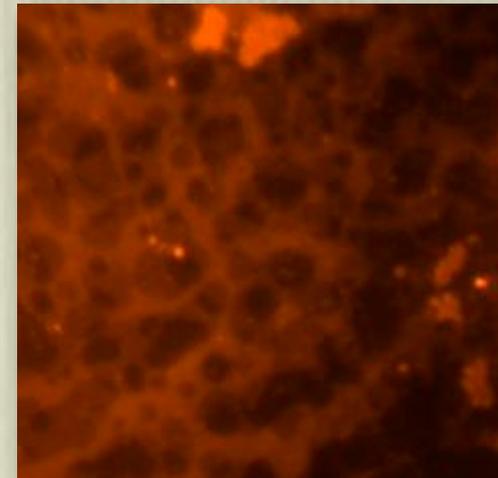
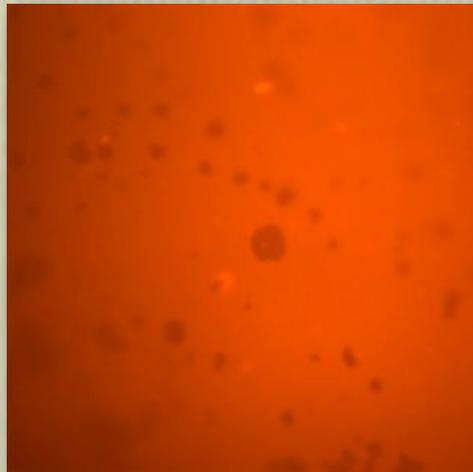
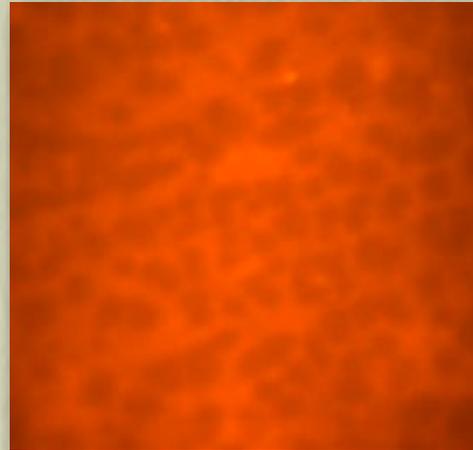
	- Ca ²⁺	+ Ca ²⁺
δh DOPC-mica	5.67 ± 0.56	4.36 ± 0.25
δh DPPC -mica	6.69 ± 0.35	5.37 ± 0.14

Membrane destabilization by detergents

Fluorescence



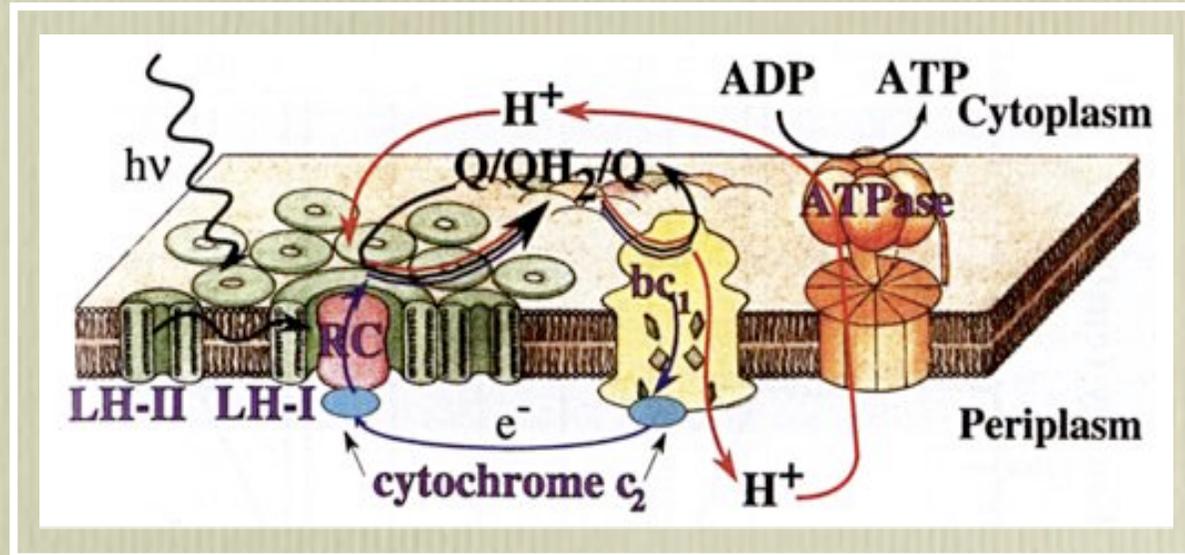
Upright microscope
LD objectives
HBO lamp



150 μ m

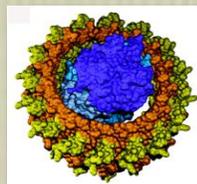


Direct Incorporation of transmembrane proteins within supported lipid bilayers

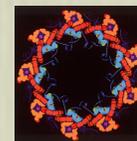


Photosynthetic apparatus of bacteria

Rhodobacter spheroides and *Rhodospseudomonas acidophila*



RC-LH1



LH2

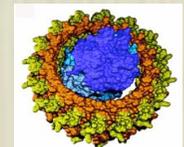
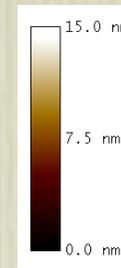
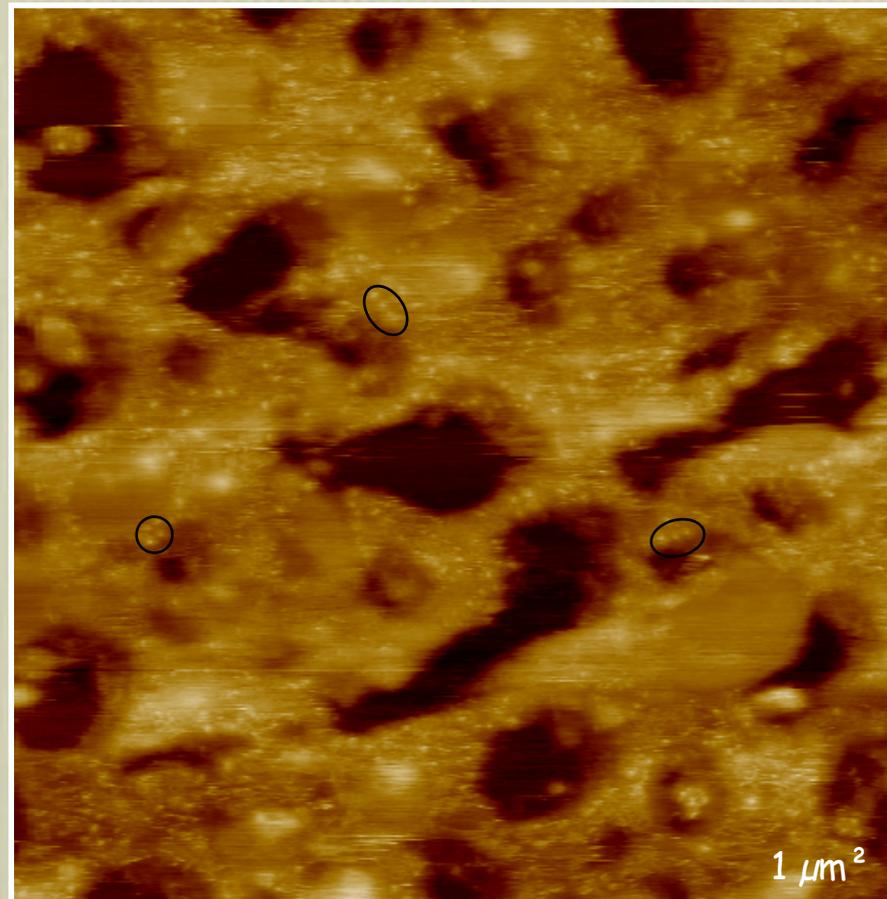
Incorporation of RC-LH1 from *Rhodobacter spheroides*

15 min incubation with RC-LH1

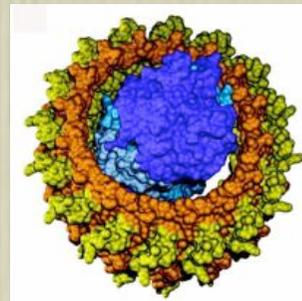
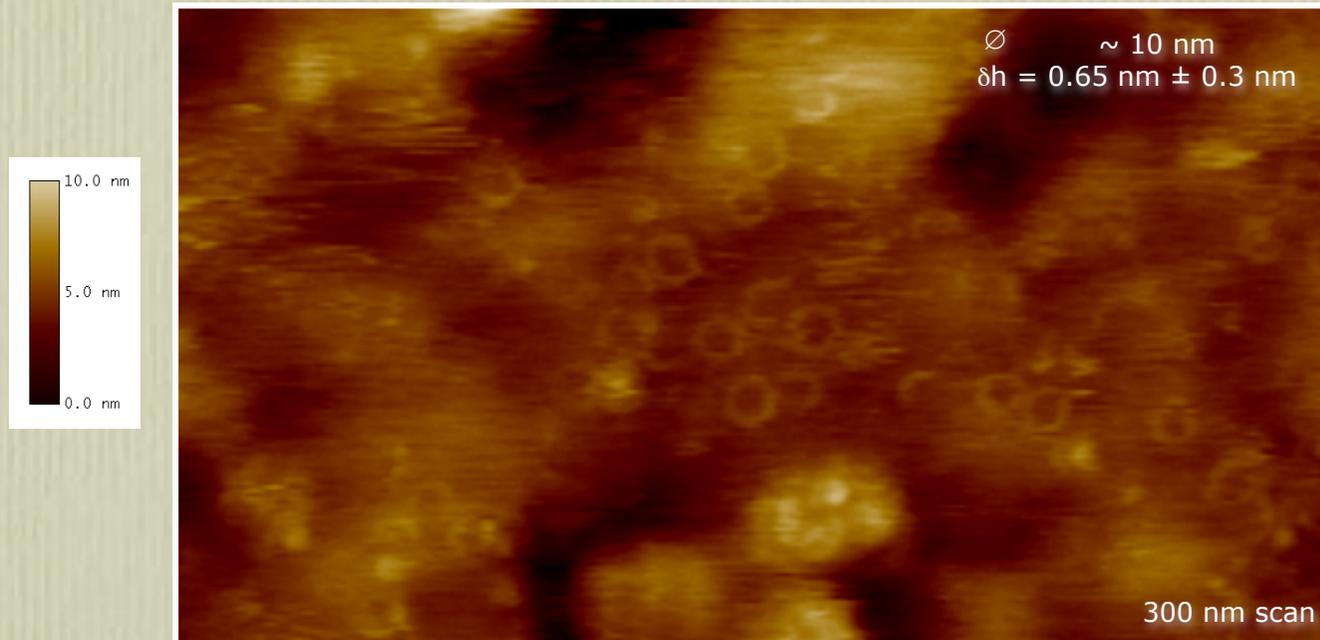
500 ng (1.5 picomole) in 0.075 mM DOTM, 150 mM KCl, 10 mM Tris pH 7.4

Contact mode

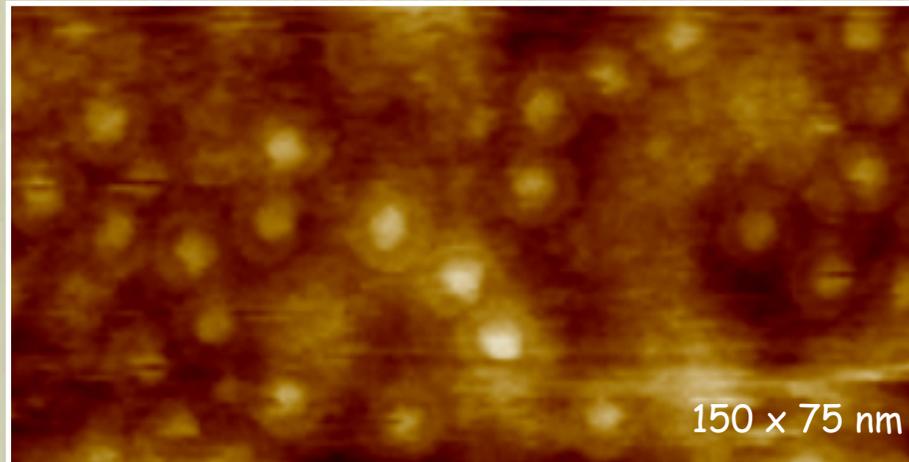
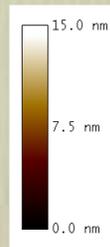
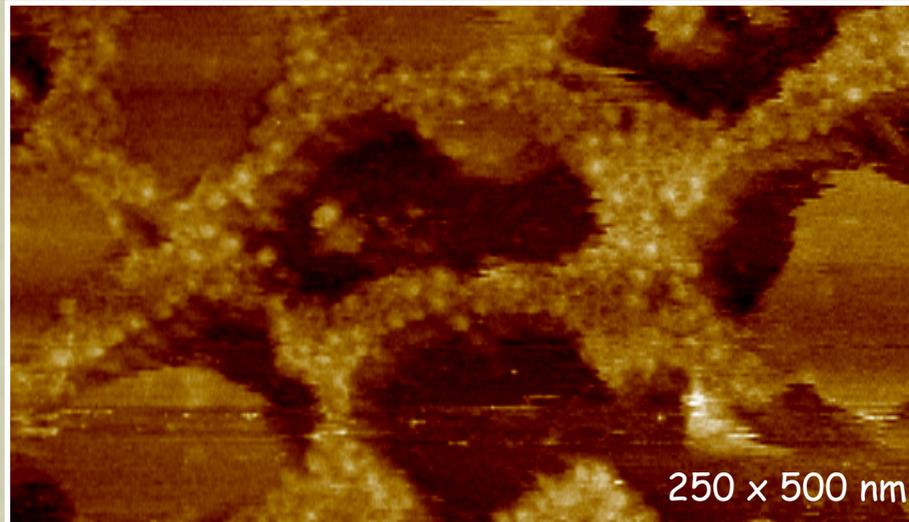
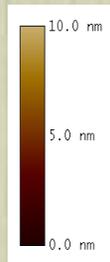
in 150 mM KCl, 10 mM Tris pH 7.4



Incorporation of RC-LH1 from *Rhodobacter spheroides*



Incorporation of RC-LH1 from *Rhodobacter spheroides*



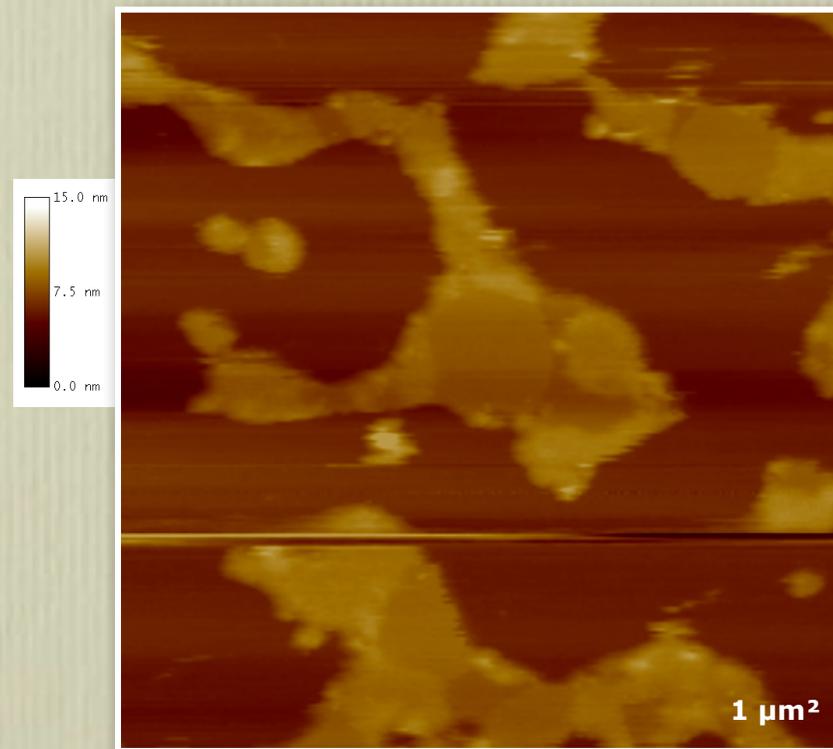
Unidirectional
incorporation

Proteins diffuse in the fluid phase and segregate in the lipid bilayer

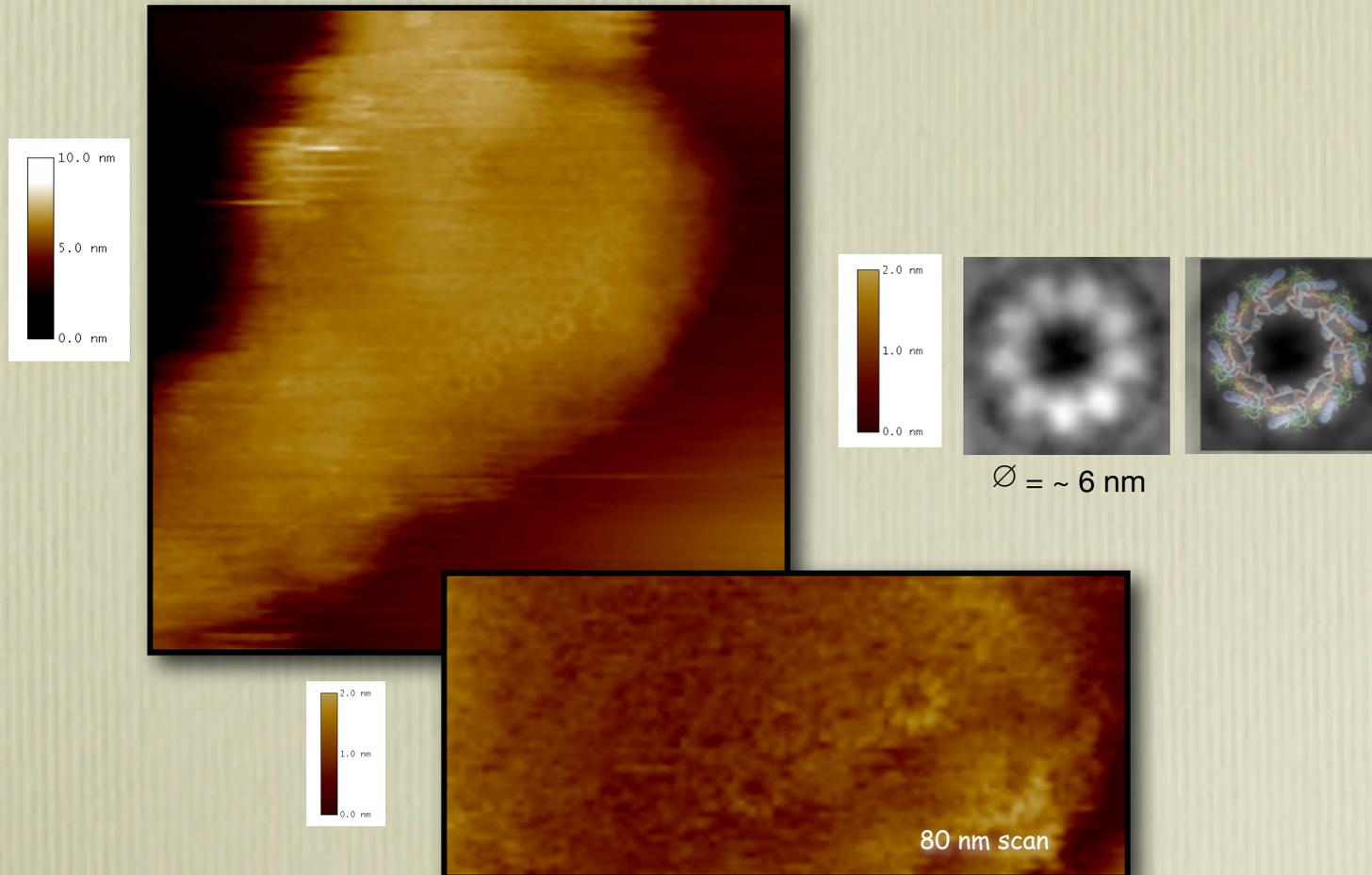
Incorporation of LH2 from *Rhodospseudomonas acidophila*

Experimental procedure

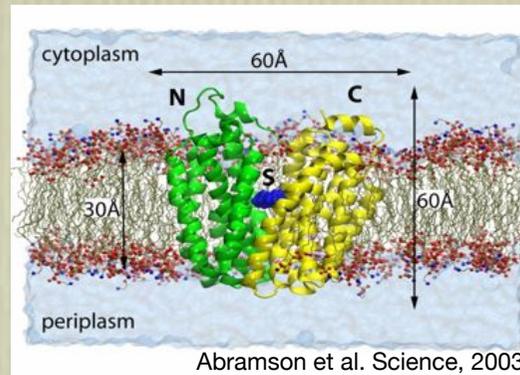
100 ng (1 picomole) in 0.075 mM DOTM, 150 mM KCl, 10 mM Tris pH 7.4
15 min incubation with LH2



Incorporation of LH2 from *Rhodospseudomonas acidophila*



Incorporation of the lactose permease Lac Y



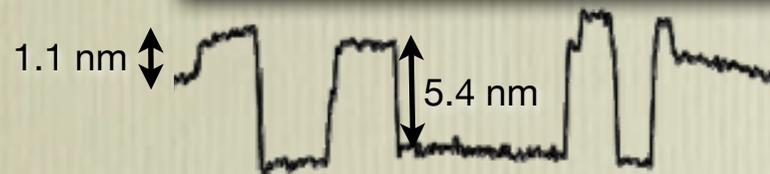
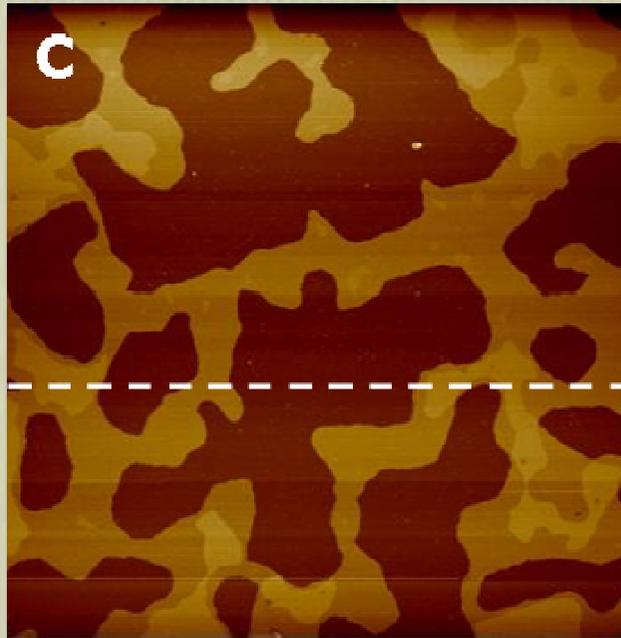
Laura Picas Escoffier
M. Teresa Montero
Jordi Hernández-Borrell



Incorporation of the lactose permease Lac Y

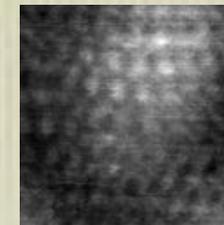
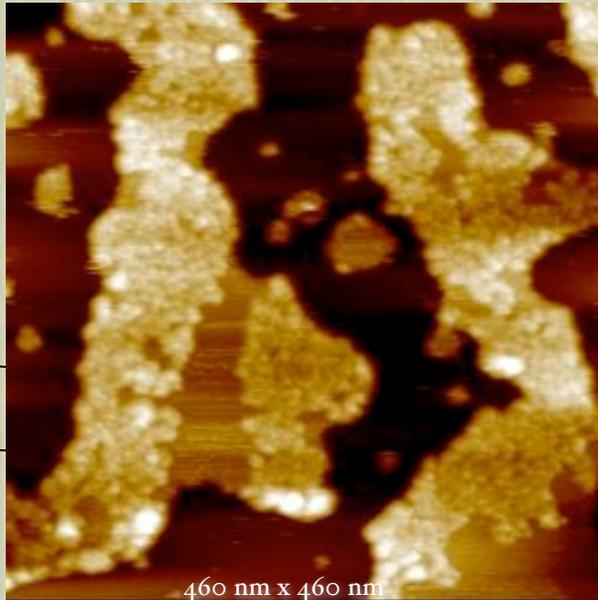
Detergent effect on POPE/POPG bilayer

n-Dodecyl- β -D-Maltopyranoside (DDM)(2 cmc)



Incorporation of the lactose permease Lac Y

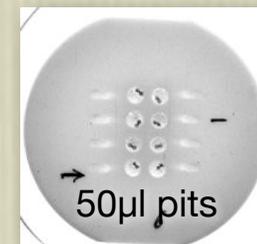
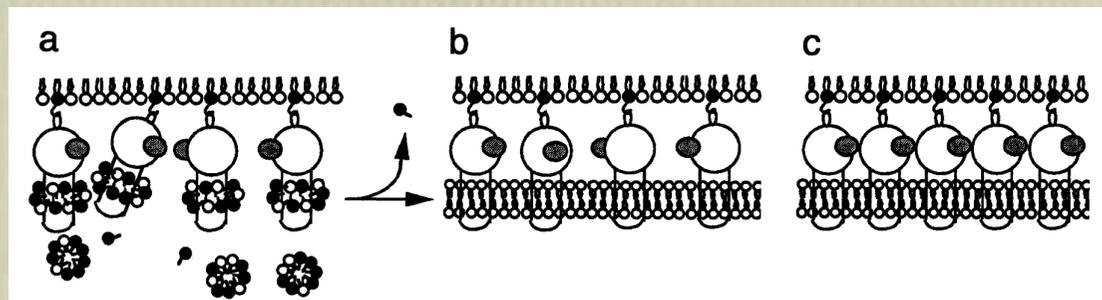
lipid bilayer
protein incorporation



Images acquired in TM-AFM

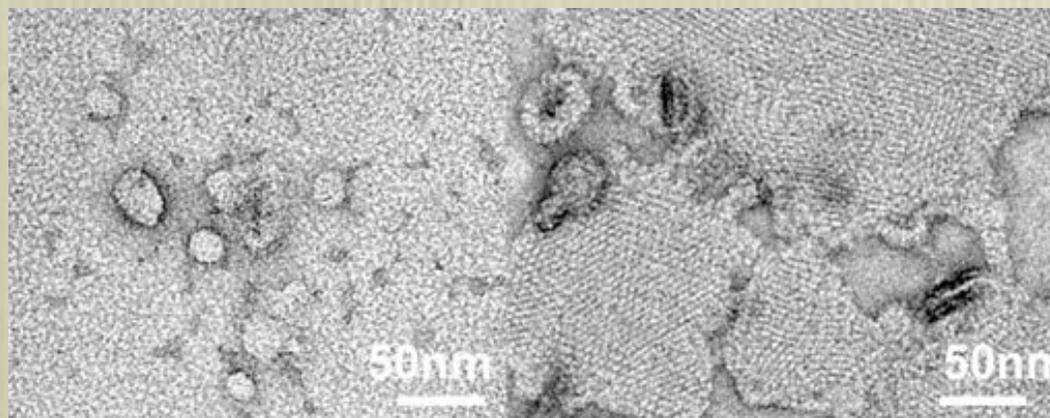


AFM imaging of proteins reconstituted using the lipid-layer 2D technique

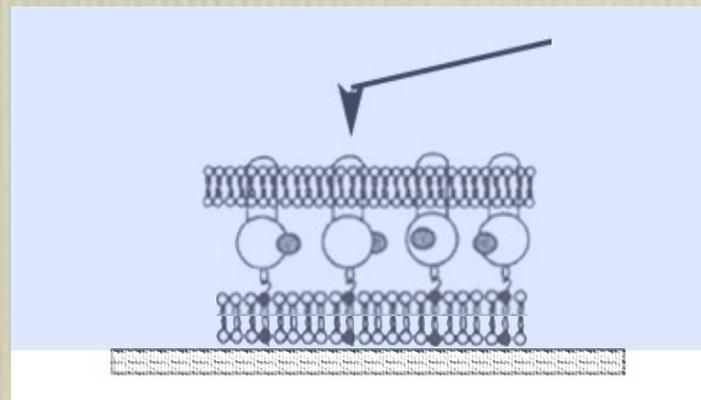
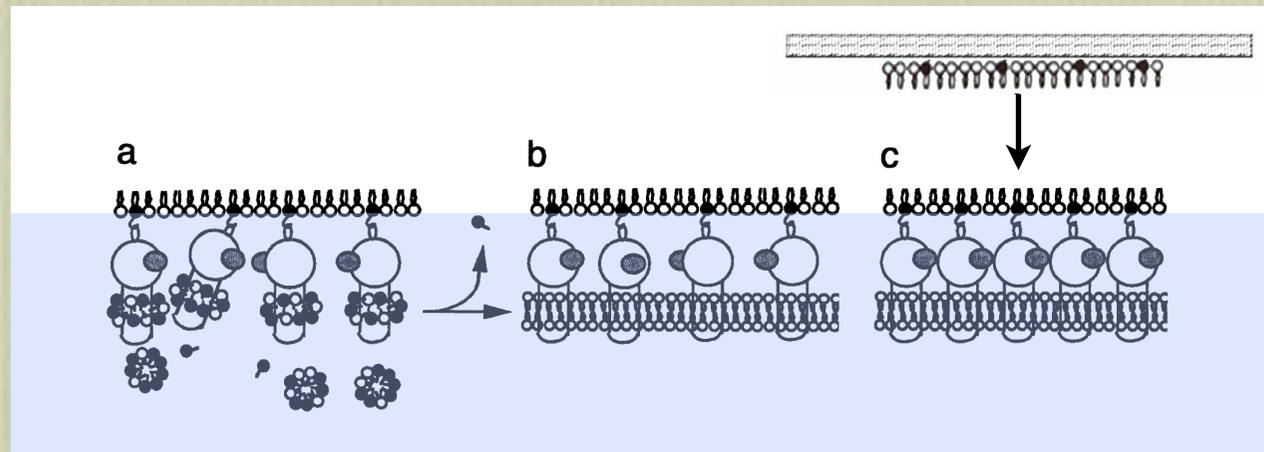


Levy et al, J Struct Biol (1999) 127:44-52

**Proteins in a native environment
1 µg per assay**

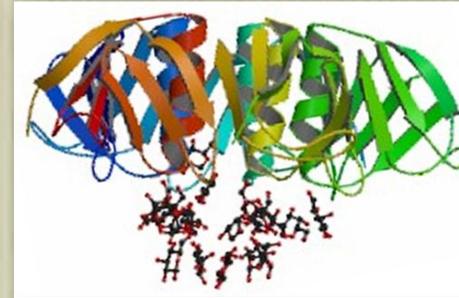
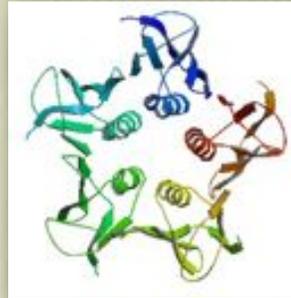


AFM imaging of proteins reconstituted using the lipid-layer 2D technique

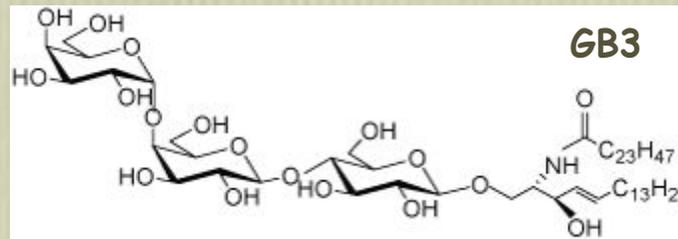


Transfer of shiga toxin B-subunit on lipid monolayer

$\emptyset = 6 \text{ nm}$

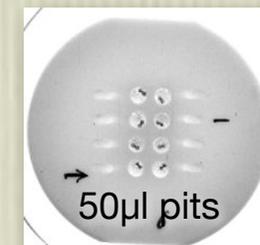
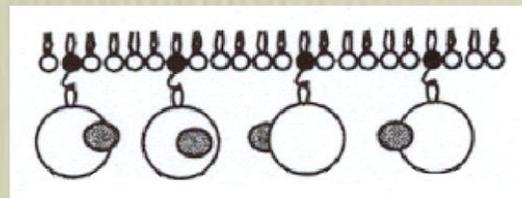


$\partial h = 2.5 \text{ nm}$

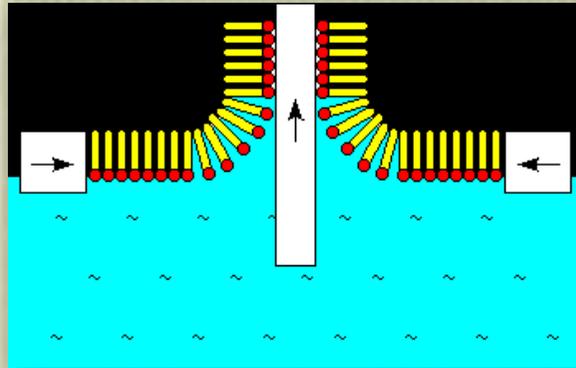


DOPC/ GB3

Shiga toxin
B-subunit

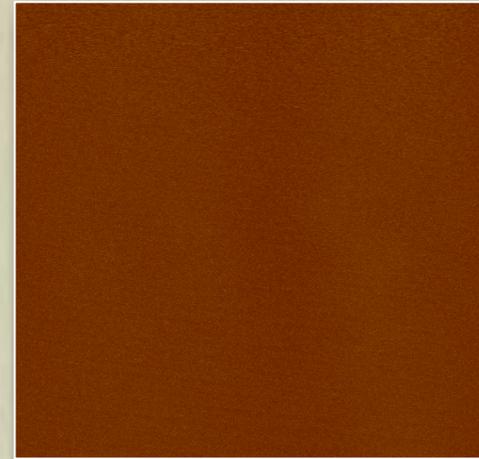


Transfer of shiga toxin B-subunit on lipid monolayer



Teflon barrier

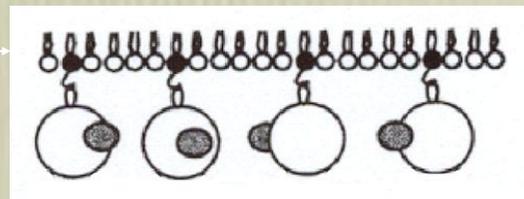
Langmuir-Blodgett transfer
30 mN/m



DOPC monolayer
Rms roughness = 0.8 Å



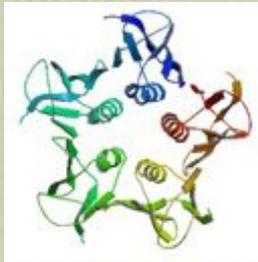
DOPC/ GB3
(Globotriaosylceramide)
(6:4)



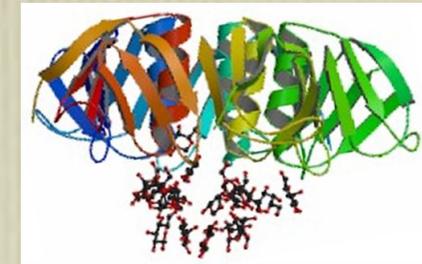
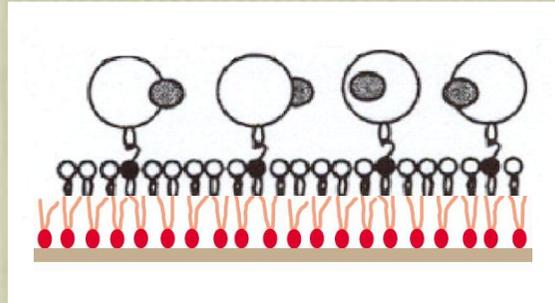
Shiga toxin
B-subunit



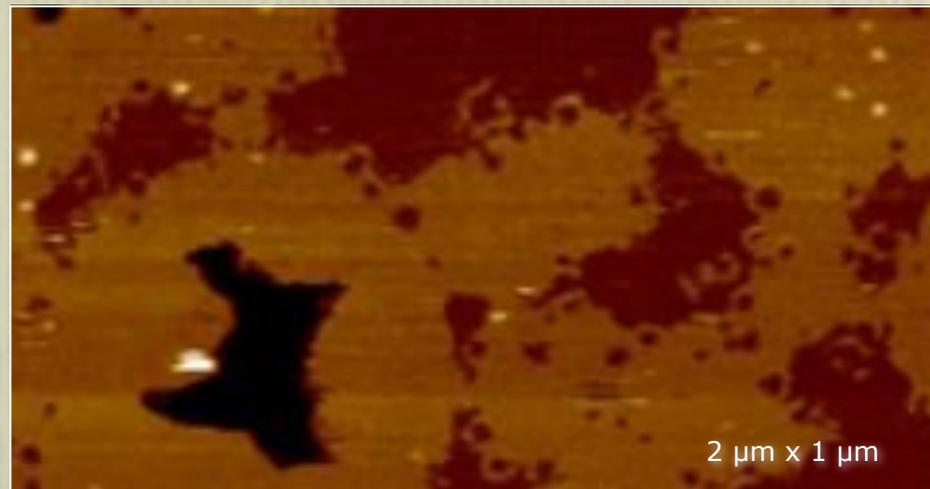
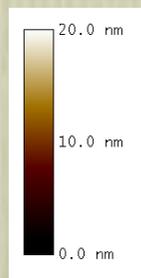
Transfer of shiga toxin B-subunit on lipid monolayer



$\varnothing = 6 \text{ nm}$

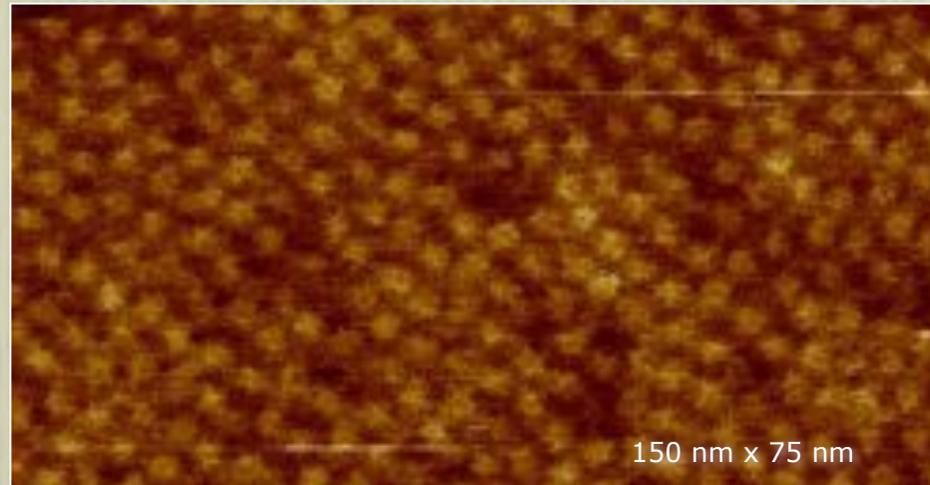
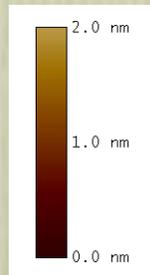
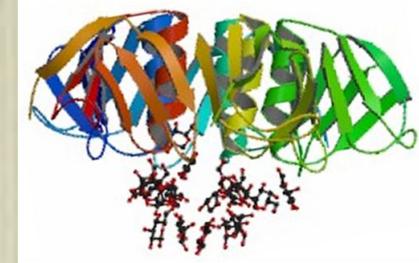
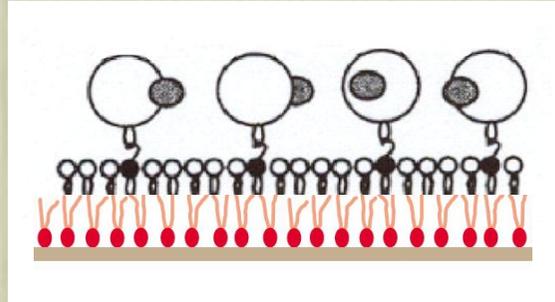
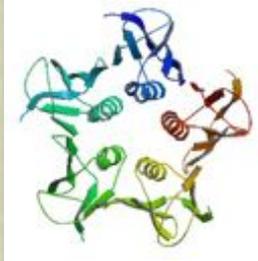


2.5 nm

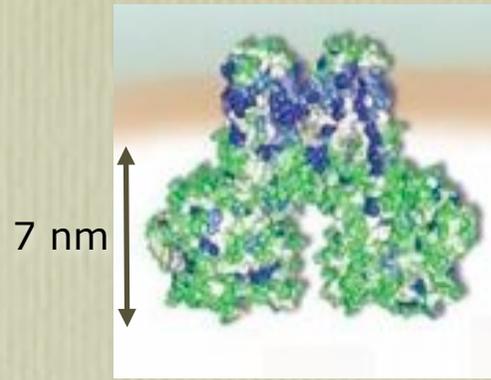


Contact mode

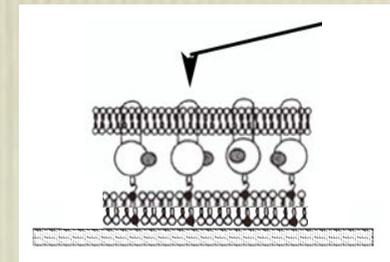
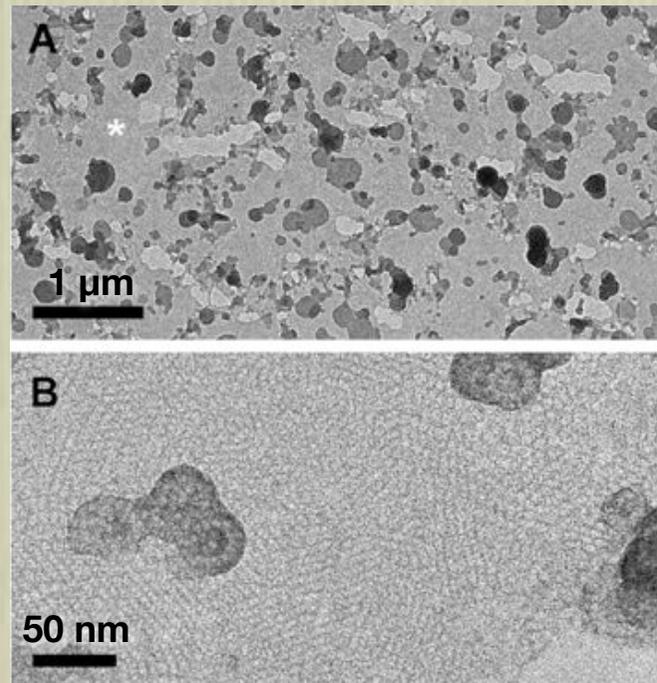
Transfer of shiga toxin B-subunit on a lipid monolayer



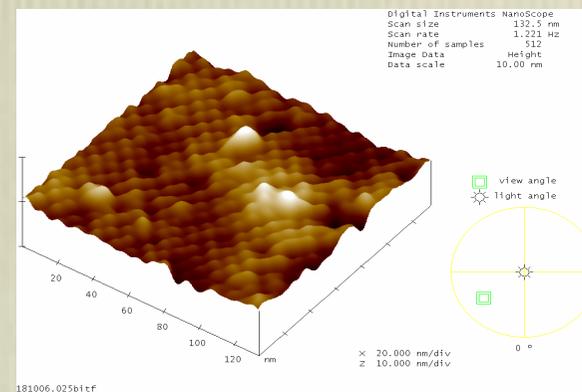
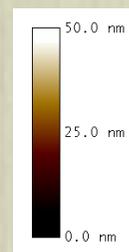
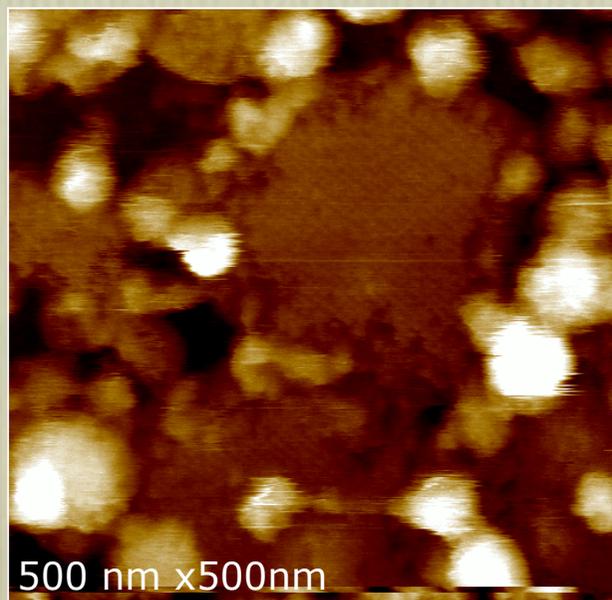
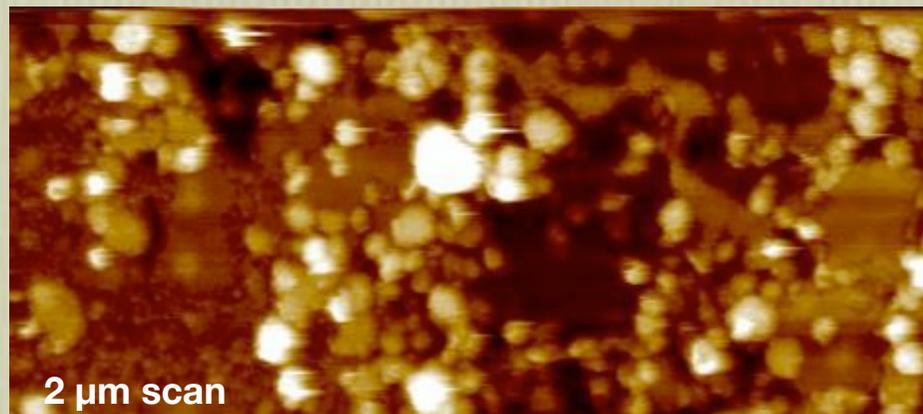
Transfer of reconstituted BmrA on a lipid monolayer



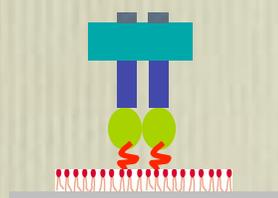
**ATP-binding cassette transporter
(ABC transporter)**



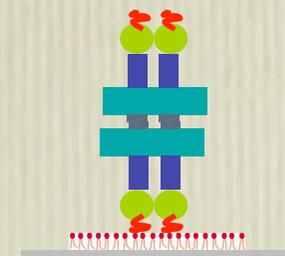
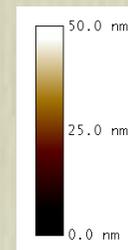
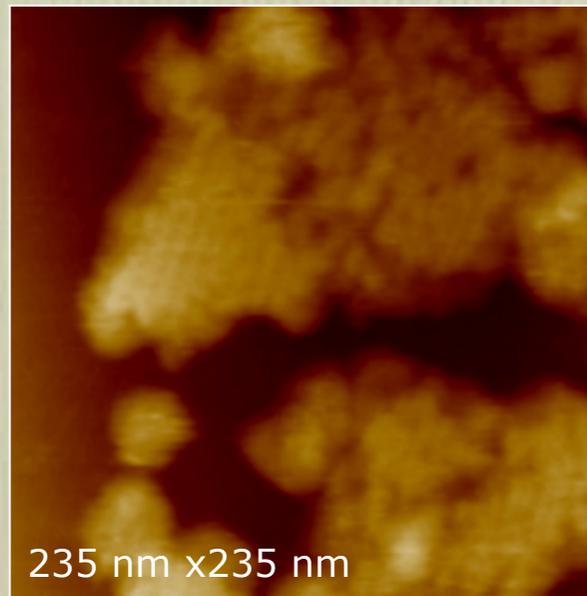
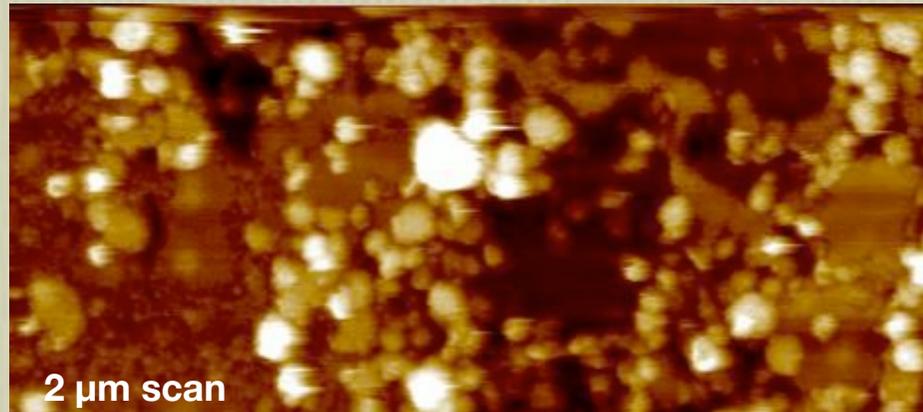
Transfer of reconstituted BmrA on HOPG



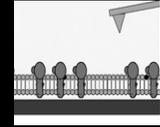
7 nm lattice, δh 1 nm



Transfer of reconstituted BmrA on HOPG



Summary

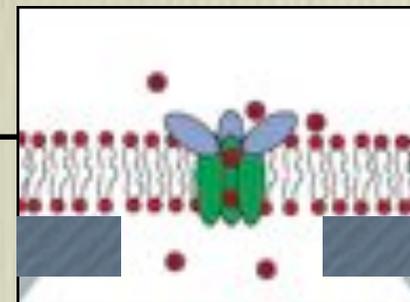
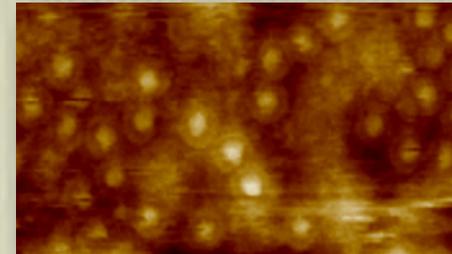


- * Low amount of protein
- * Single/control of the orientation
- * High resolution imaging

- * Proteins can segregate in the bilayer

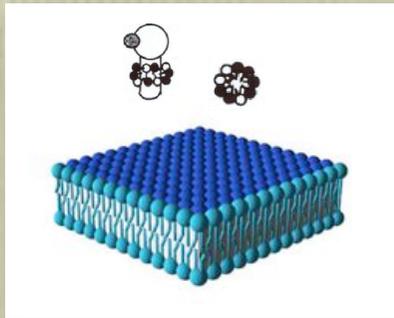
- * Suitable for functional and nano-biotechnological applications (transfer of large membrane patches)

- * Improvement of AFM resolution

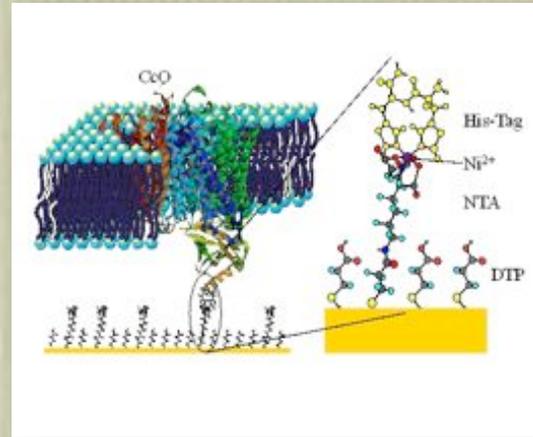


Membrane Biosensor

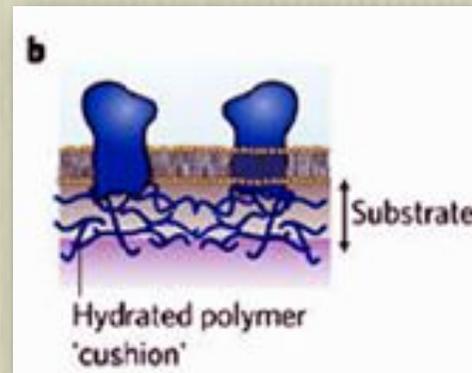
Comparison with other techniques



Incorporation



Tethered protein reconstitution



Polymer cushion



Tollin et al. 2003. *Trends Pharmacol Sci.* 24:655-9
Sinner et al. *Angew. Chem. Int. Ed.* 2006, 45, 1 – 5
Giess et al 2004. *Biophys J.* 87:3213-20.
Tanaka & Sackmann *Nature* 2005, 437, 656

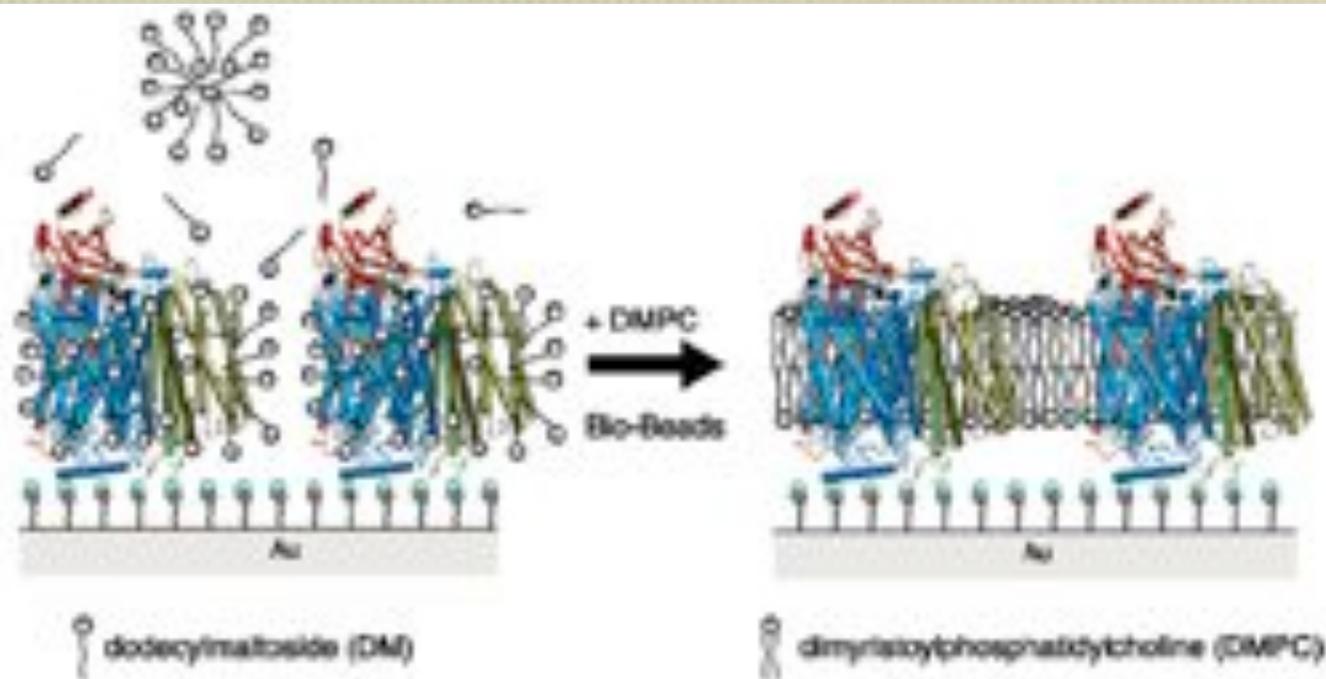
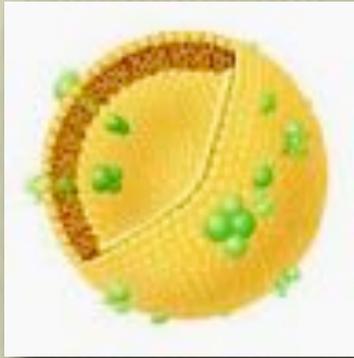


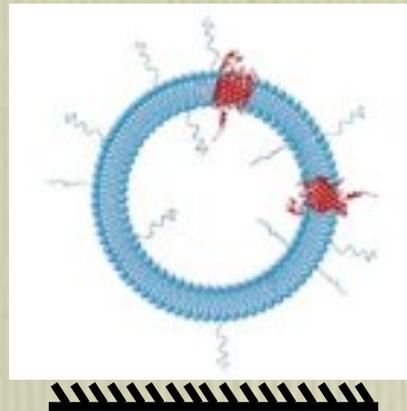
Figure 2. Schematic representation of the reconstitution of surface-adsorbed cytochrome *c* oxidase into a lipid bilayer. The surface-adsorbed cytochrome *c* oxidase via the affinity of the His-tag to the Ni-NTA surface (see Figure 1) is surrounded by the detergent dodecylmaltoside (left). After reaching maximum surface coverage, the protein is exposed to detergent-devalubilized lipid vesicles (DMPC), and microporous Bio-Beads are added. The retreat of the detergent molecules by the beads drives the lipids to form a layer around the membrane protein (right). Giess, *Biophys. J.* 2004, 87, 3213



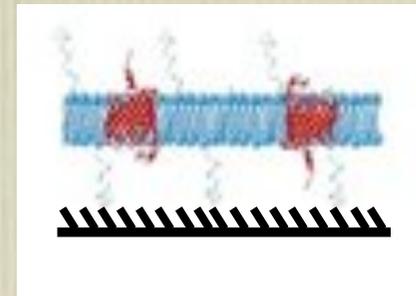
Proteoliposome fusion



Triton
removal
by
BioBeads



Vesicles
anchorage
and
disruption



DSPE-PEG-NHS



VDAC



POPC

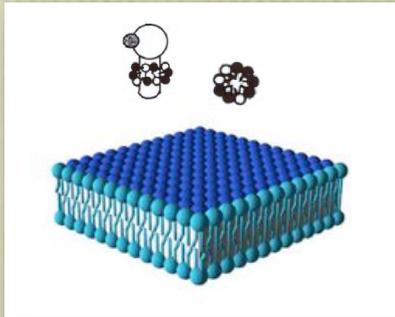


Amine-grafted surface
(gold-cysteamine or
glass-ADMS)



Triton-X100

Comparison with other techniques



Incorporation

Nanodisc?

THE NANODISC SOLUTION

- GPCRs: β_2 AR, RHO
BioTechniques (2006) 40, 601
JBC (2007) 282, 14875
- P450 +/- Reductase
ABB (2005) 430, 218
JBC (2007) 282, 7066
- Aromatase
BBRC (2008) 372, 379
- TAR Receptor
(1, 2, 3 - dimers)
PNAS (2006), 103, 11509
- Bacteriorhodopsin (mono/tri-mer)
ABB (2006) 450, 215
- SecYEG
EMBO J. (2007) 26, 1995
JBC (2009) 284, 7897
- Cholera Toxin
Anal. Chem. (2008), 80, 6245
- Coagulation Factors
JBC (2007) 282, 6556
- Methods Enzymol. (2009) 464, 211-231
FEBS Lett. (2010) 584, 1721-1727



Tollin et al. 2003. *Trends Pharmacol Sci.* 24:655-9
 Sinner et al. *Angew. Chem. Int. Ed.* 2006, 45, 1 – 5
 Giess et al 2004. *Biophys J.* 87:3213-20.
 Tanaka & Sackmann *Nature* 2005, 437, 656