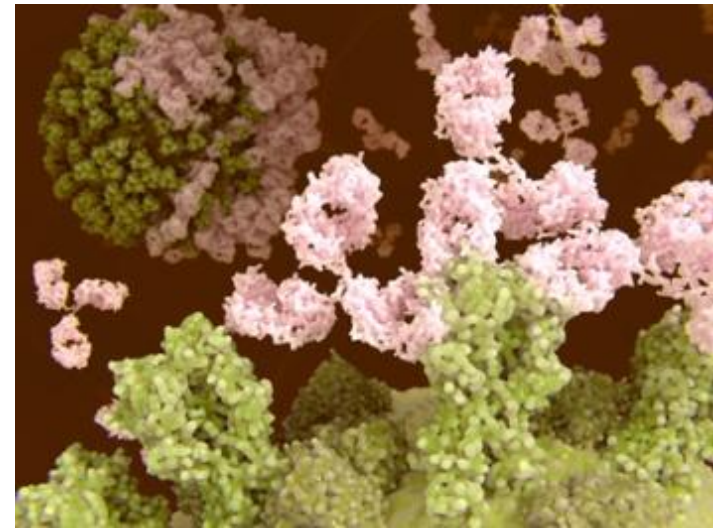


# Models of cell membrane domains for surface interactions structural investigation

Valeria Rondelli

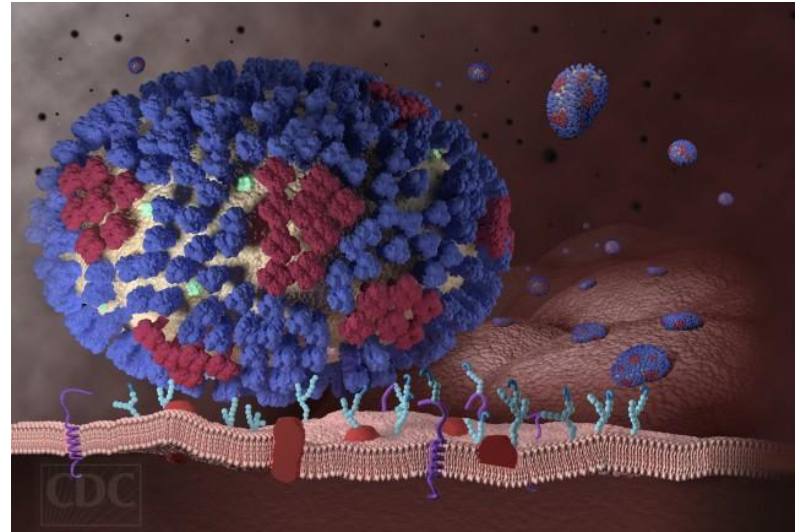
*Dept. of Medical Biotechnology and Translational Medicine*

*Università degli Studi di Milano*



# CELL MEMBRANE INTERACTION WITH APPROACHING BODIES

Biology question is far...  
what can we face?



→ *simplified* systems

keeping the *main 'bio'-features*

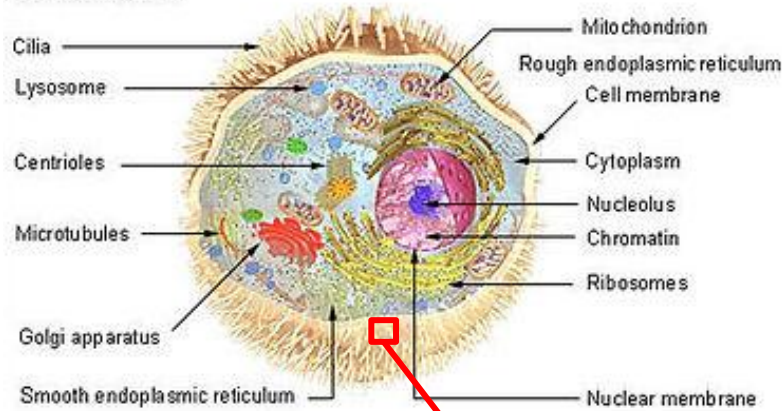
*Challenging aspects*    -> *The model*

-> *The possibility to investigate it (the technique)*

# THE SYSTEM

## Phospholipid membranes

### Cell Structure

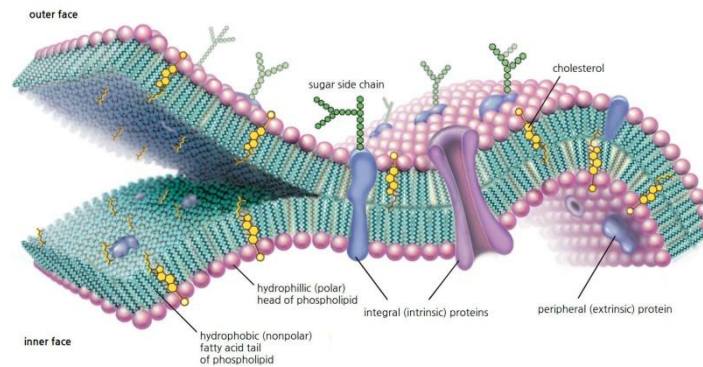
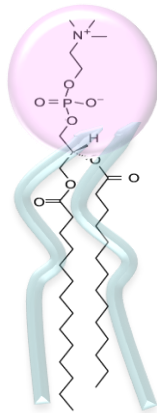


*Cell membrane*

*Structure*

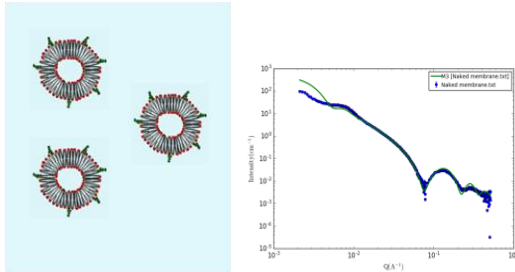
*Dynamics*

*Activity*



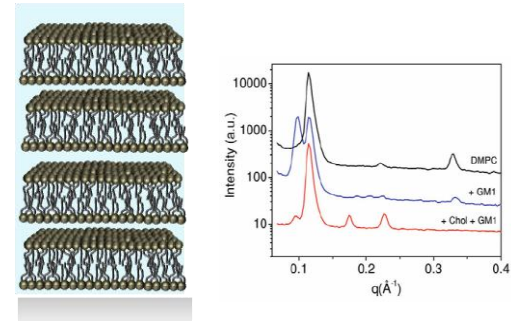
# COMPLEMENTARY TECHNIQUES FOR MEMBRANE STRUCTURAL INVESTIGATION

## *SANS/SAXS from monolamellar vesicles*



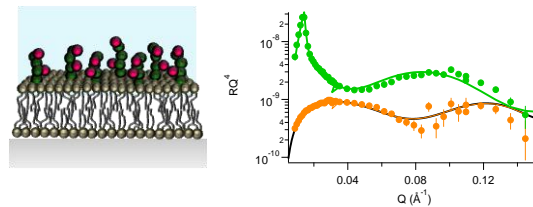
Membrane form factor  
Aggregate shape  
Structuring in solution

## *N/XD from membrane stacks*



Membrane  
thickness/thicknesses  
(lateral domains)

## *N/XR from single supported membranes*



Membrane transverse structure

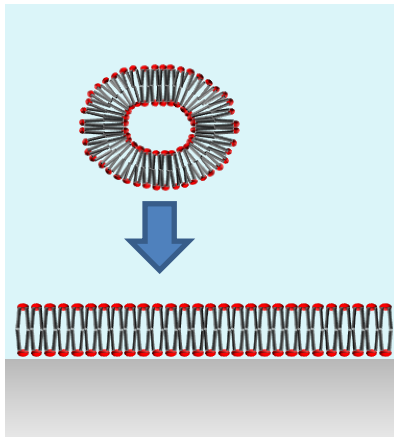
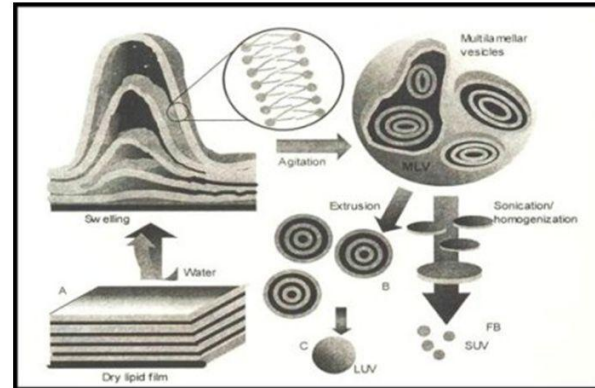


# DIFFERENT METHODS FOR MEMBRANE DEPOSITION

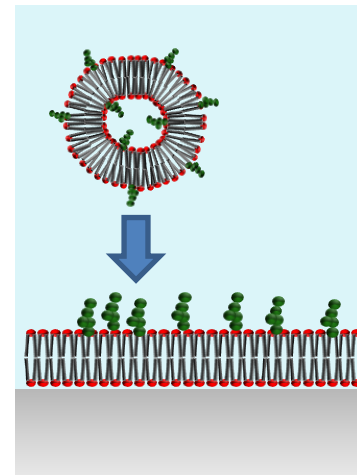
## Vesicle Fusion



### Liposomes Preparation



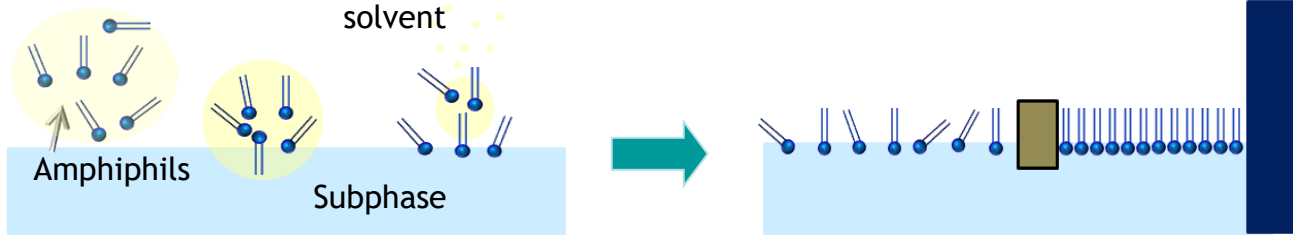
Fusion on solid supports



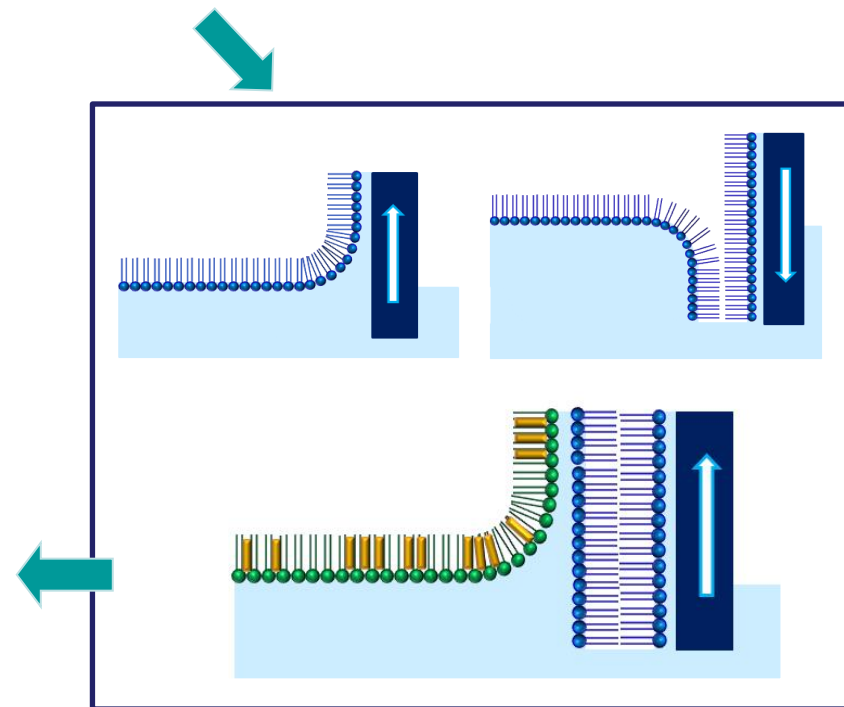
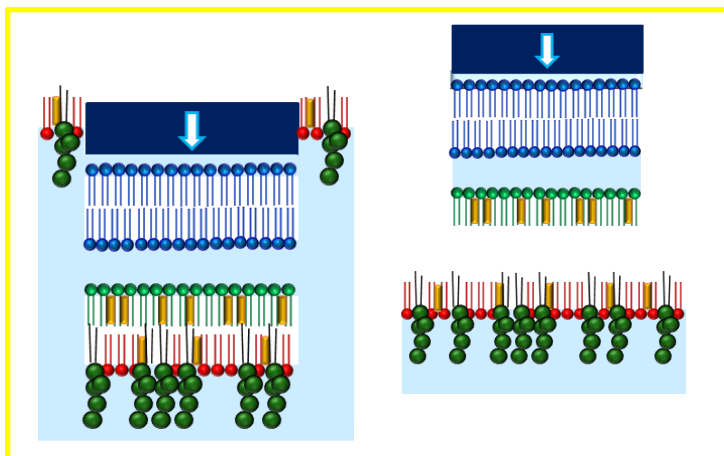
Multicomponent vesicle fusion

Spreading solution      Evaporation of solvent

Amphiphils      Subphase

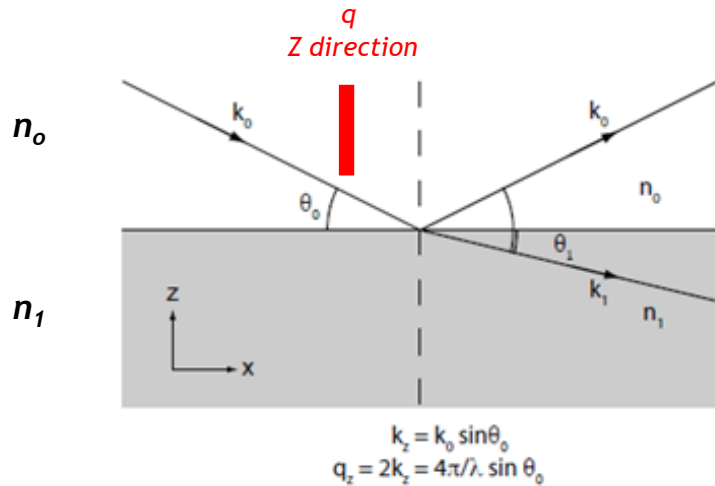


Langmuir Blodgett –  
Langmuir Schaefer  
technique



**HOMOGENEOUS DEPOSITION (*5 nm thick*) OVER LARGE AREAS (*5x8 cm<sup>2</sup>*)  
OF MONOLAYERS CONTROLLED IN **NUMBER** AND **COMPOSITION****

# SPECULAR REFLECTION FROM FLAT INTERFACES



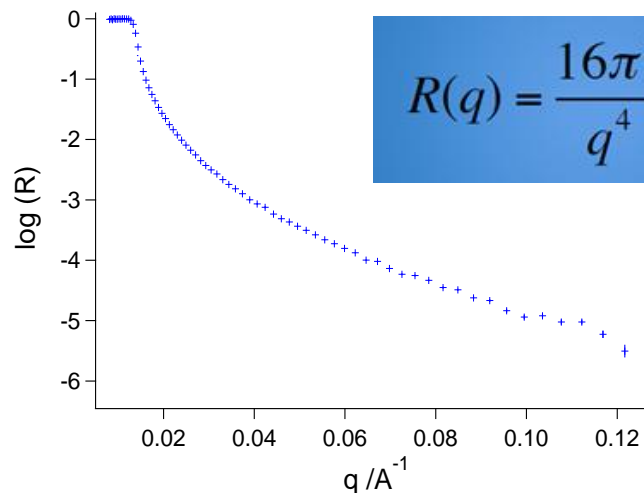
If  $\theta_i = \theta_r$



Specular reflection



The intensity of the reflected radiation depends on  $n_o$  and  $n_1$

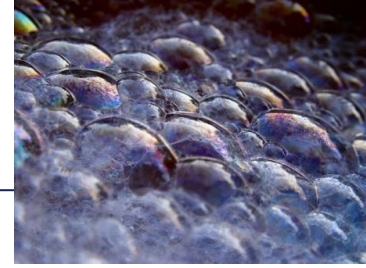


$$R(q) = \frac{16\pi^2}{q^4} |N'_b(q)|^2$$

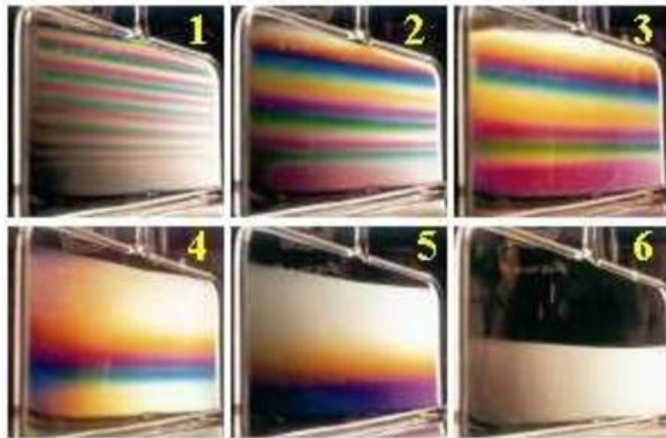
$$N_b = \frac{\sum n_i b_i}{V}$$

Scattering Length Density (SLD)

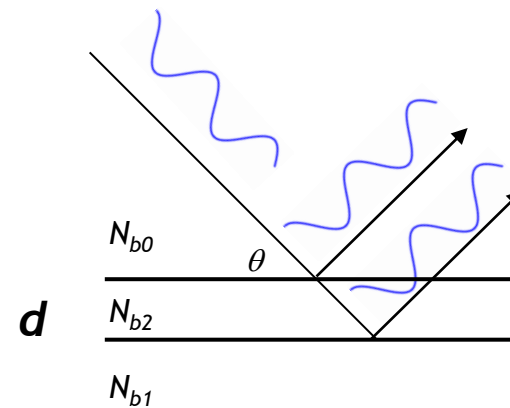
# SPECULAR REFLECTION FROM MULTILAYERS



More than 1 interface:

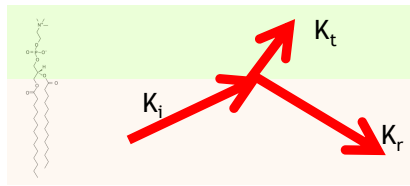


Newton, 1675



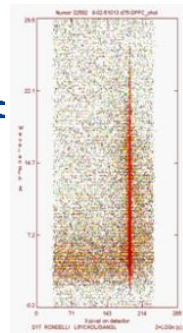
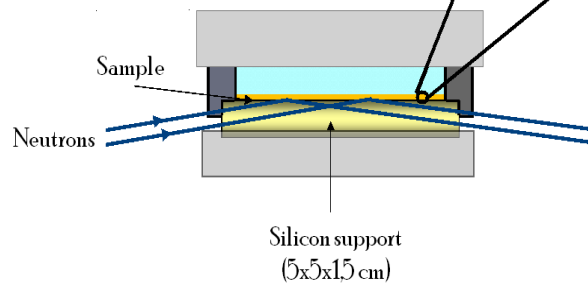
Interference depends on **thickness  $d$**  as well as on the refractive indexes  $N_{bi}$

# REFLECTOMETRY ON LIPID MEMBRANES

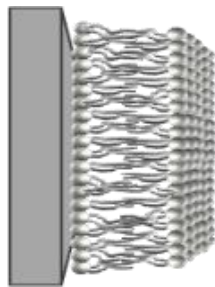
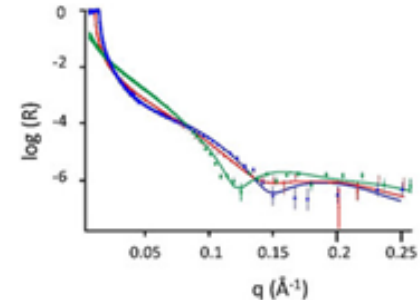


F. water  
E. heads  
D. chains  
C. chains  
B. heads  
A. water

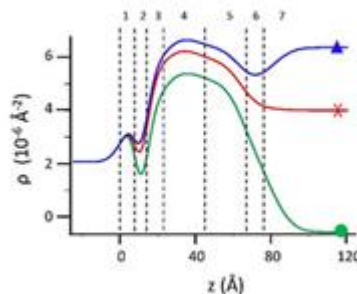
$$R \approx \left( \frac{16\pi^2}{q^4} N_b^2 \right) e^{-q_z^2 \sigma^2}$$



The Reflectivity spectrum obtained is given by the **interference** of the waves **reflected** from the top and bottom of each layer



## SAMPLE CROSS PROFILE



Information about the transverse structure of the sample, layer by layer:

**thickness, composition, compactness, roughness**

# WHY DIFFERENT RADIATIONS?

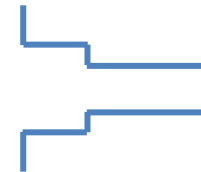
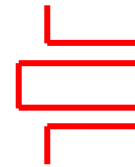
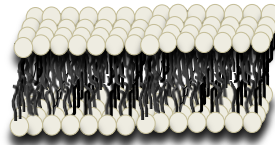
$$I(q) \div c M P(q) S(q) \text{contrast}^2$$

**X-ray**

Good contrast for SUGARS (high electron density)

X-rays

Neutrons

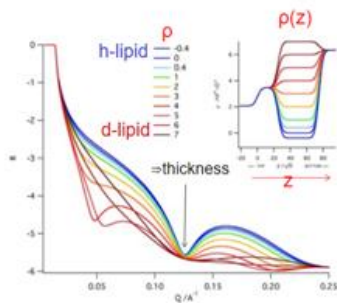


**Neutron**

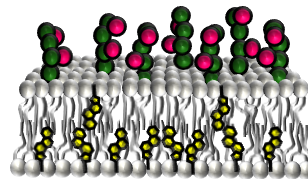
Hydrogen coherent scattering length :  $-3.74 \times 10^{-5} \text{ \AA}$   
Deuterium coherent scattering length :  $+6.67 \times 10^{-5} \text{ \AA}$



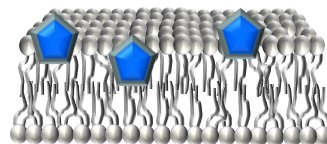
Playing with selective deuteration **protiated molecules can be evidenced in the deuterated phospholipid matrix**



H. Wacklin, ESS



Membrane components distribution

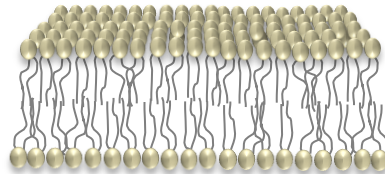


External interacting molecules distribution

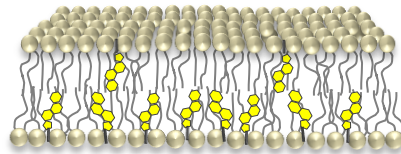


# MEMBRANE MODELS

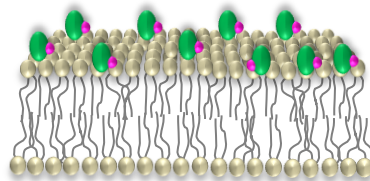
*phospholipids*



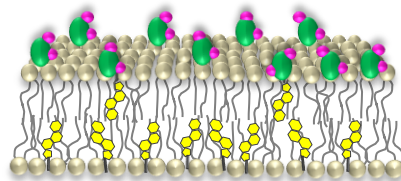
+  
*cholesterol*



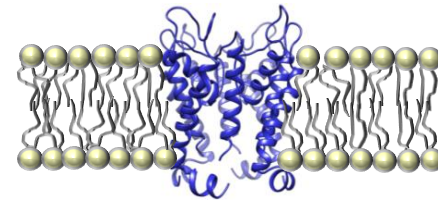
+  
*glycolipids*



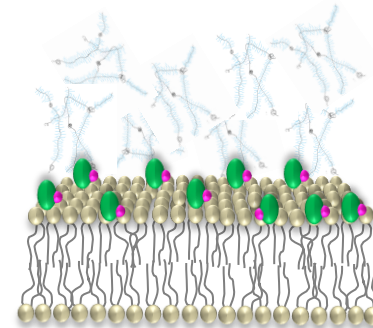
*raft models*



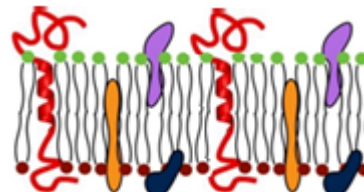
+  
*transmembrane protein (K<sup>+</sup> channel)*



+  
*gel forming proteins (mucin)*

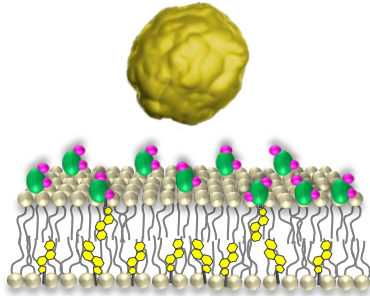


*from natural vesicles/membrane extracts*

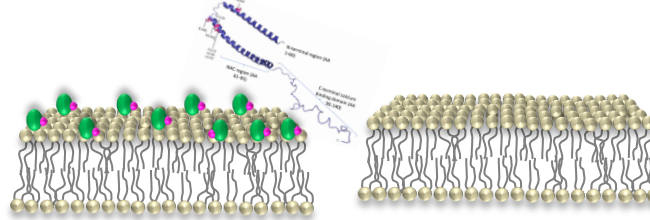


# WE CAN MODEL INTERACTIONS AT MEMBRANE SURFACE

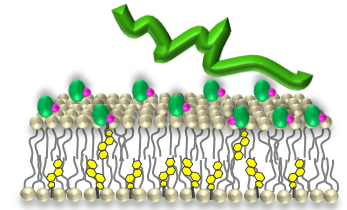
## Enzymatic digestion on membrane surface



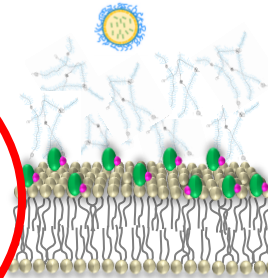
## $\alpha$ -synuclein interaction



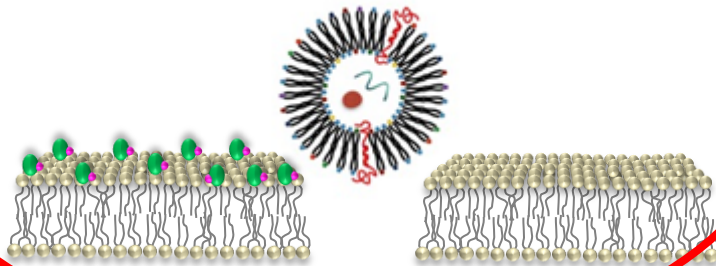
## A $\beta$ Peptide interaction



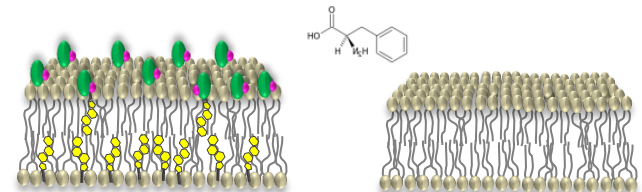
## Transmucosal delivery



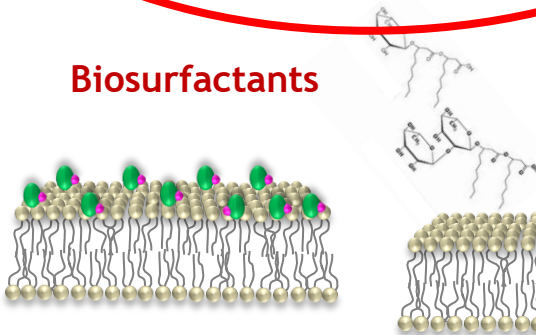
## Extracellular vesicles interaction



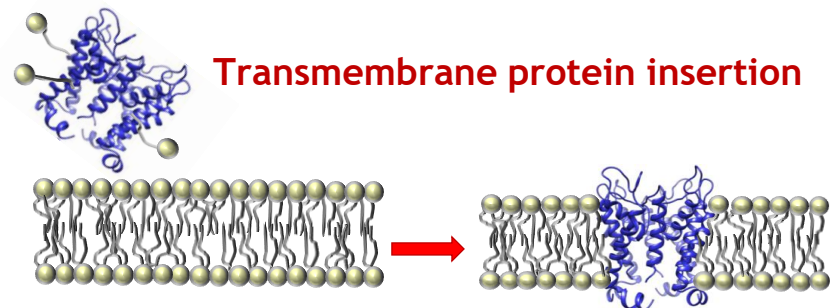
## Amino acid interaction



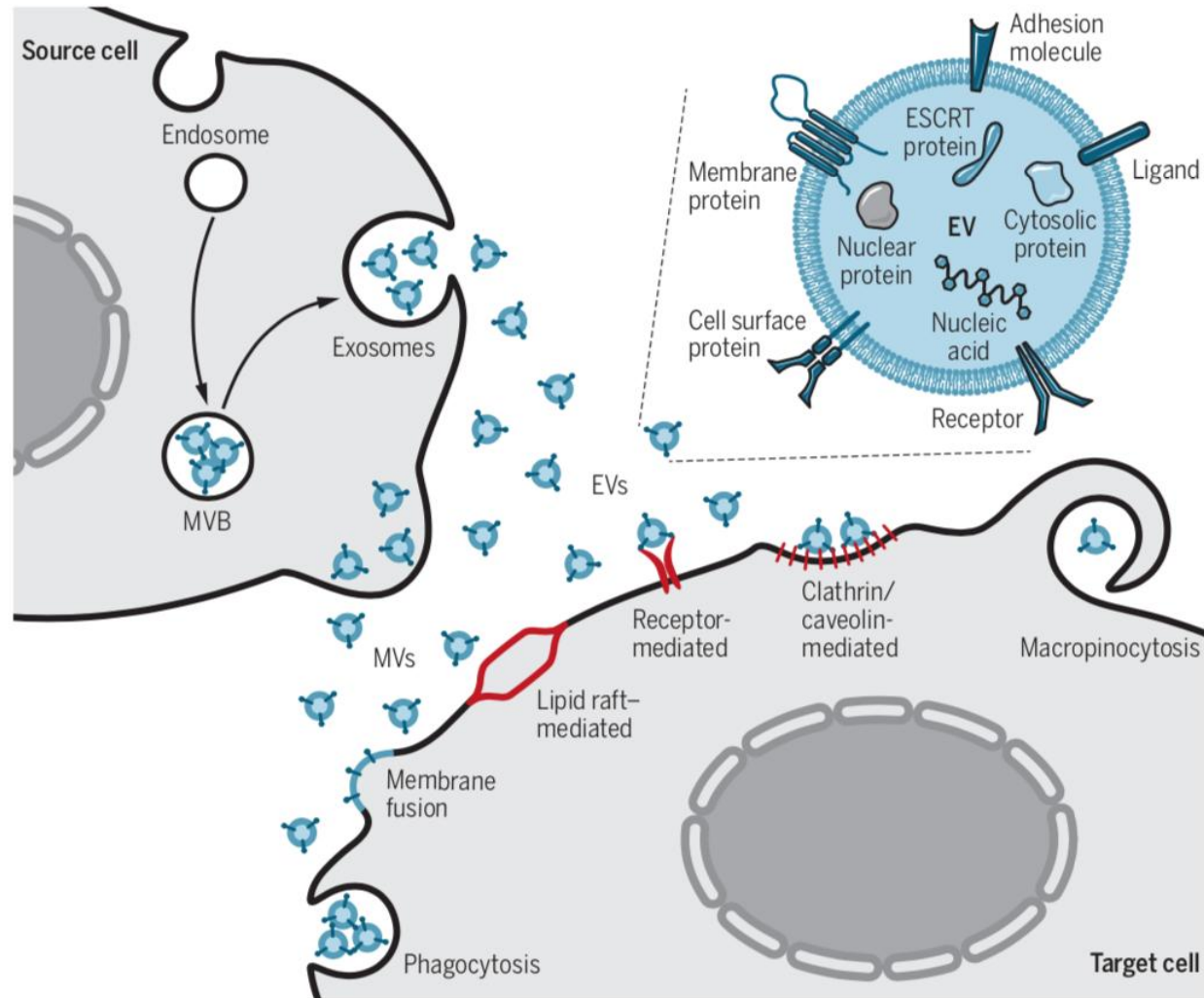
## Biosurfactants



## Transmembrane protein insertion



# EXTRACELLULAR VESICLES



Wiklander et al., Science Trans. Medicine (2019)

## GAPS

- ✓ Nanoscale spatio-temporal details on how different EVs interact with target cells
- ✓ Factors influencing the biogenesis and release of the molecular cargo

## KEY CONCEPTS

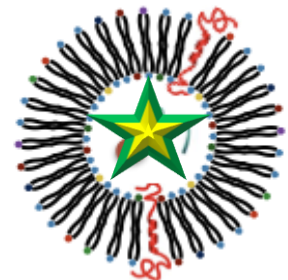
→ **Uptake** dynamics and mechanisms are tightly related to the **potency and function of EVs**

→ **Evs** play a key-role in **diseases spreading**



**Understanding the molecular basis of diseases**  
(eg. cancer, neuro-degenerative)

→ **devise of EVs-based therapies**

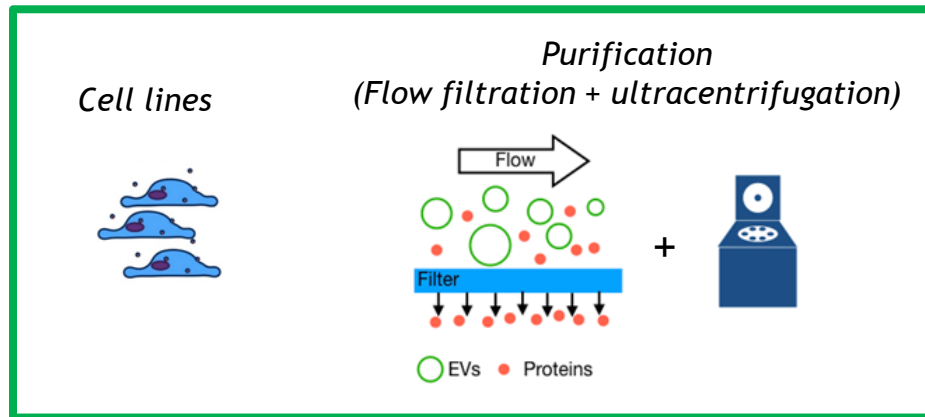


# OUR STRATEGY

## sEVs

Standardized protocols and Good Manufacturing Practice conditions to derive **highly stable vesicles** of **defined size** and **reproducible molecular profiles** from Umbilical Cord multipotent Mesenchymal Stem (Stromal) Cells (MSCs)

### 1. Isolation and Characterization of sEVs



- ✓ Nanoparticle Tracking Analysis
- ✓ Light Scattering
- ✓ Surface marker analysis
- ✓ Cryo-Electron Microscopy
- ✓ Small Angle X-Ray Scattering
- ✓ Small Angle Neutron Scattering
- ✓ Atomic Force Microscopy
- ✓ Neutron Reflectometry

### 2. Investigation of sEVs uptake mechanisms

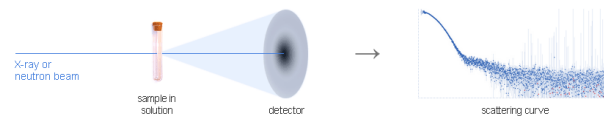
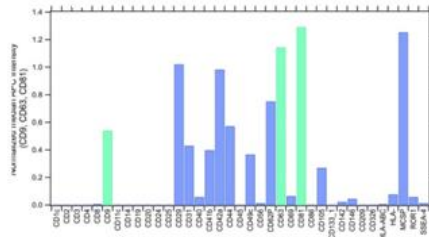
- ✓ Atomic Force Microscopy
- ✓ Small Angle Neutron Scattering
- ✓ Neutron Reflectometry



# sEVS CHARACTERIZATION

## Surface marker analysis

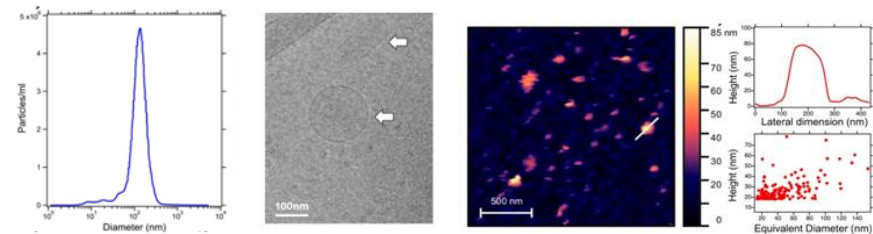
typical sEV/exosome markers (tetraspanins CD9, CD63 and CD81)



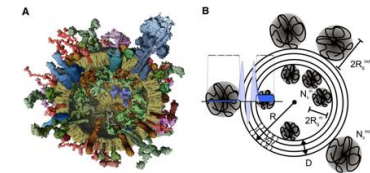
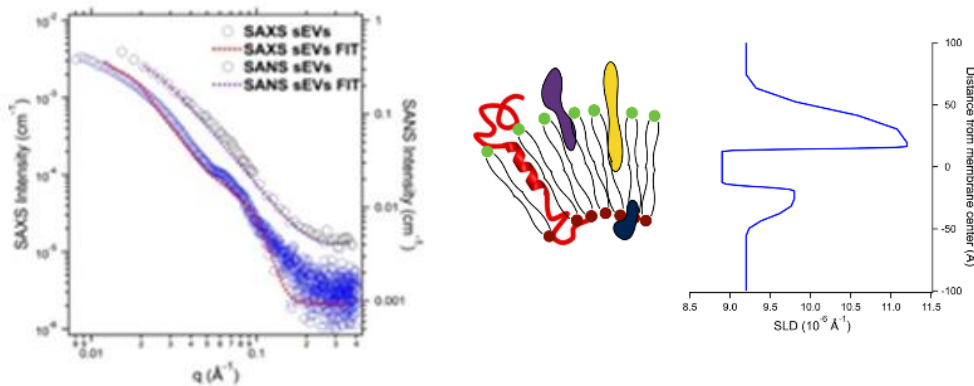
SAXS  
SANS

DLS  
NTA  
Cryo-EM  
AFM

particles sized less than 50 nm + sEVs sized ~ 100 nm



3-layered membrane model, accounting for the proteomic component extending in the extra-vesicular solution.

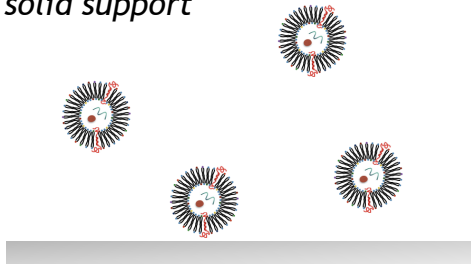


Castorph et al. Biophysical Journal (2010)  
(Synaptic vesicles)

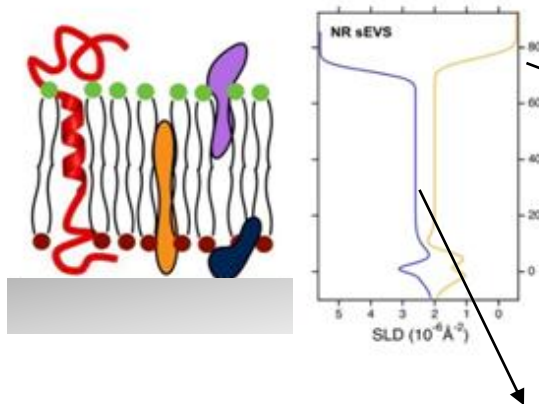
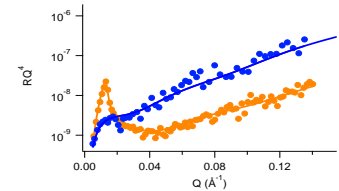
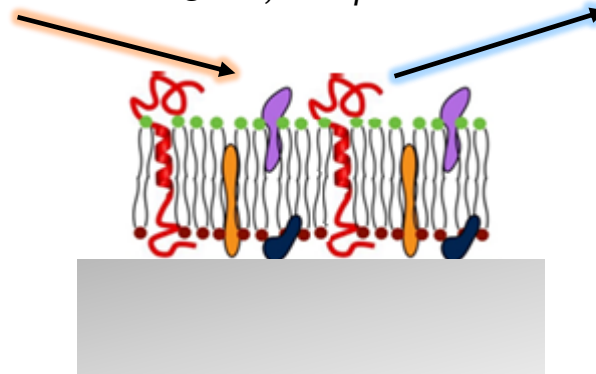


# NEUTRON REFLECTOMETRY ON sEVS-DERIVED SUPPORTED BILAYERS

Extracellular Vesicles fusion  
on solid support



Neutron reflectometry  
@BNC, Budapest



Thickness:  
 $6.9 \pm 0.2 \text{ nm}$

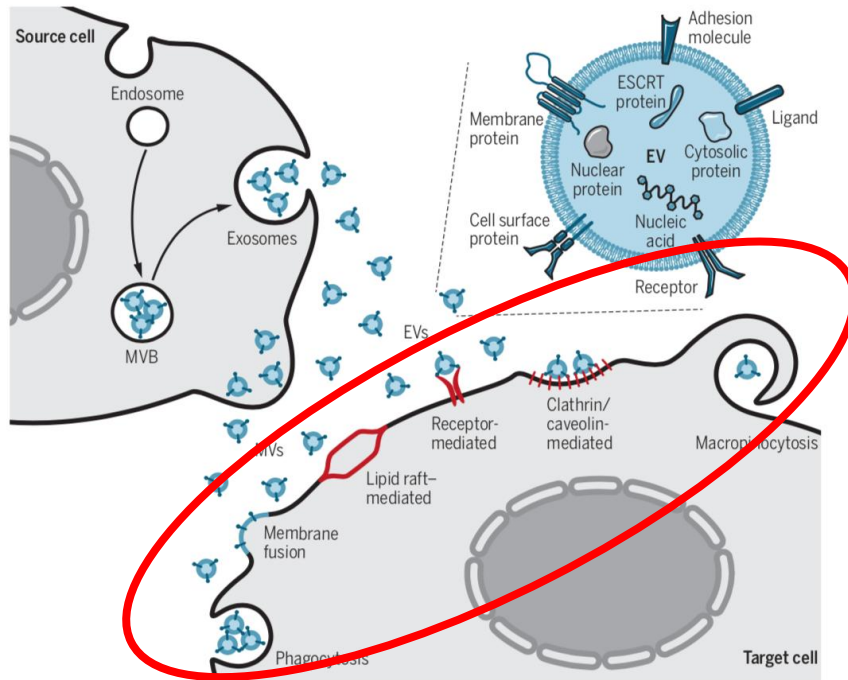
**single bilayer**  
containing molecules  
other than lipids, as  
**large proteins**

$\text{SLD } 2 \pm 0.2 \times 10^{-6} \text{ \AA}^{-2}$

**lipid : protein**  
**22 : 78**  
(by volume)

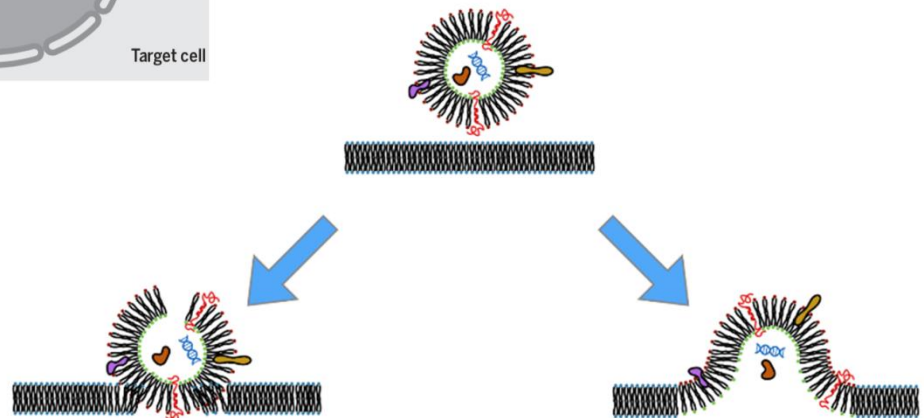
$$R \approx \left( \frac{16\pi^2}{q^4} \boxed{N_b^2} \right) e^{-q_z^2 \sigma^2}$$

# sEVS UPTAKE MECHANISMS



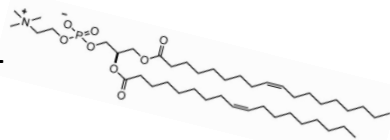
Wiklander et al., Science Trans. Medicine (2019)

Cargo release may  
be favoured or  
prevented

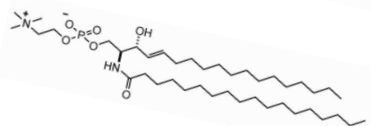


# PREFERENTIAL SITE OF INTERACTION

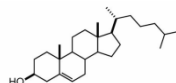
1,2-dioleoyl-sn-glycero-  
3-phosphocholine  
(DOPC)



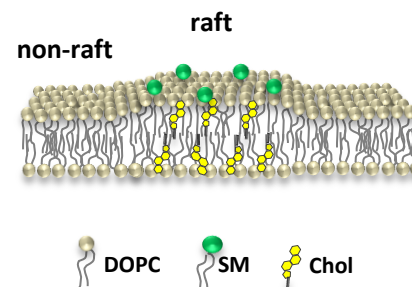
Sphingomyelin  
(SM)



Cholesterol  
(Chol)



DOPC:SM:Chol 2:1:0.15 mol

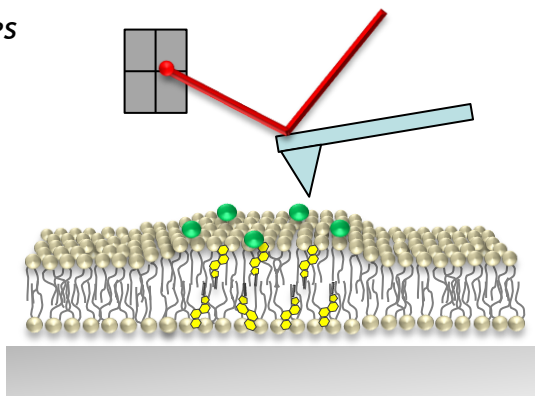


AFM

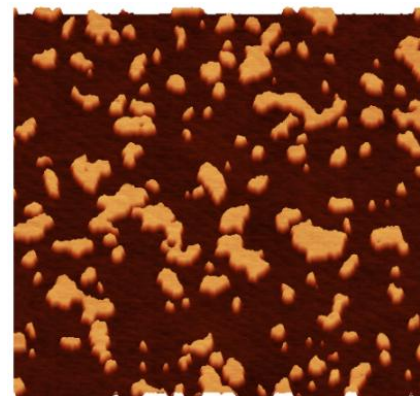
@ ELETTRA, Trieste

Fabio Perissinotto  
Pietro Parisse  
Loredana Casalis

Supported membranes  
by vesicle fusion



Lipid phase separation



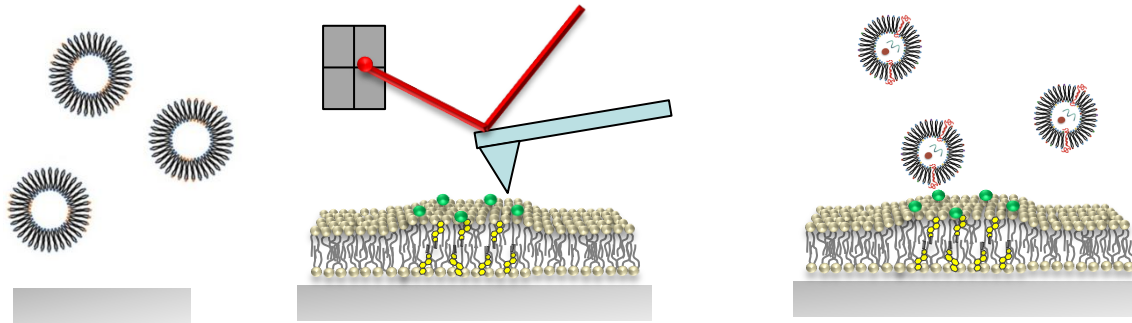
# PHASE BORDERS ARE DOCKING SITES

DOPC:SM:Chol (2:1:0.15)  
vesicles fusion

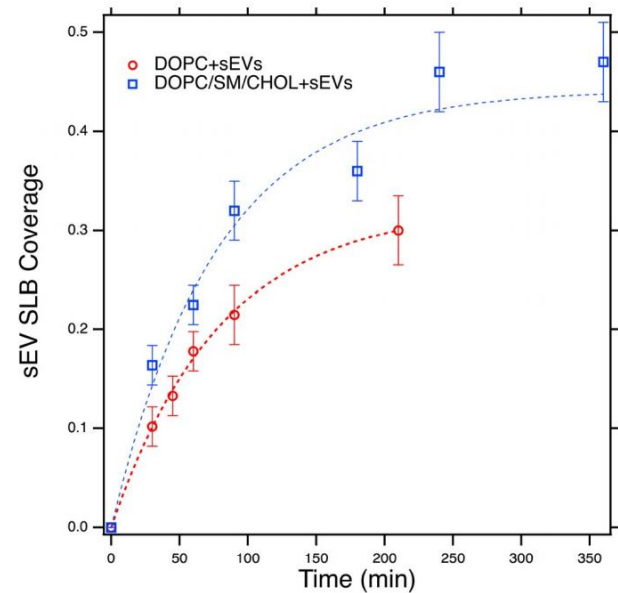
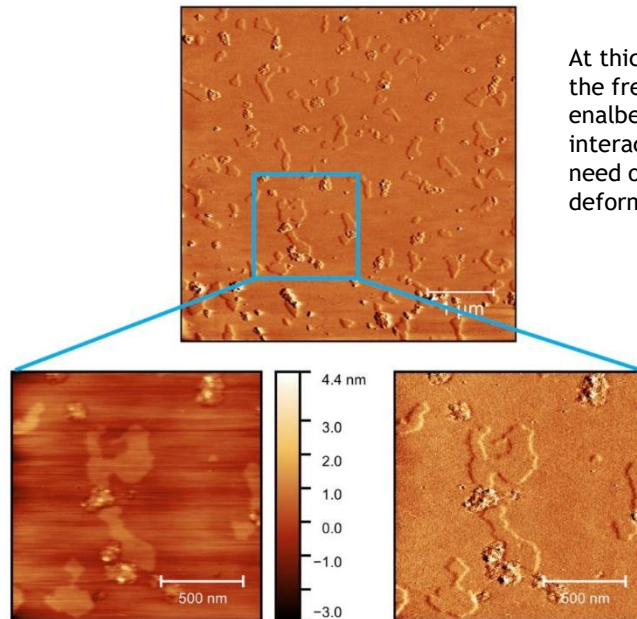
characterization

sEVs  
injection from bulk water

mixed system  
characterization



**SLB 5 minutes after addition of sEVs**



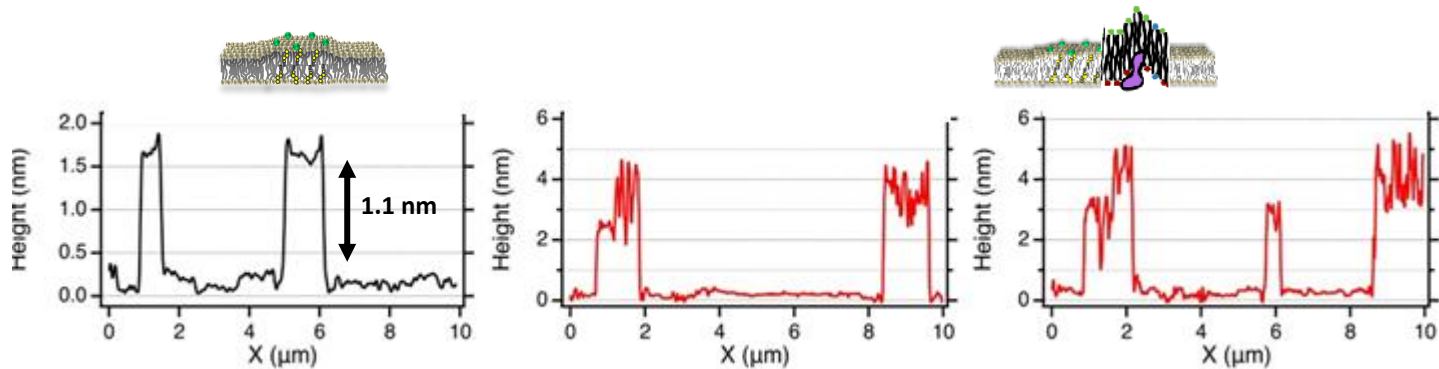
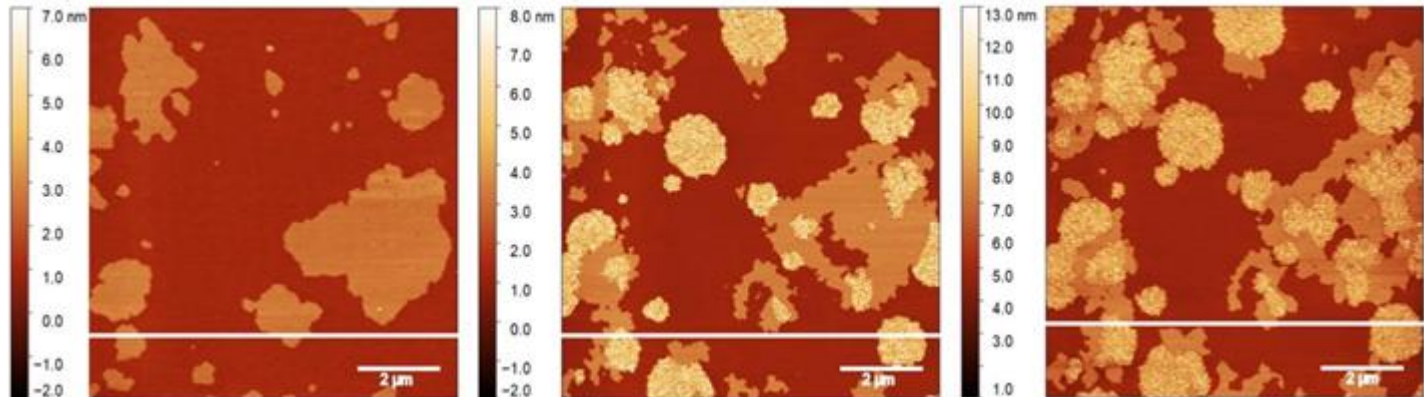
# PATCHES EXPANSION FAVOURED IN $L_d$ PHASE

+ sEVs

DOPC:SM:Chol

t = 30 min

t = 75 min



Patches protruding 3-4 nm above SLB



# STRUCTURAL DETAILS OF MIXING: NEUTRONS

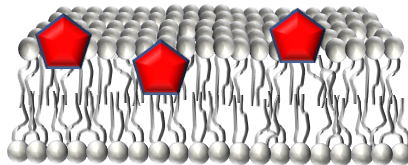
## Reflectometry

$$R \approx \left( \frac{16\pi^2}{q^4} N_b^2 \right) e^{-q_z^2 \sigma^2}$$

## Small Angle Scattering

$$I(q) \div c M P(q) S(q) (\Delta\rho)^2$$

Selective deuteration → H-bringing molecules can be evidenced in a deuterated phospholipid matrix



External molecules interaction and distribution within membrane leaflets



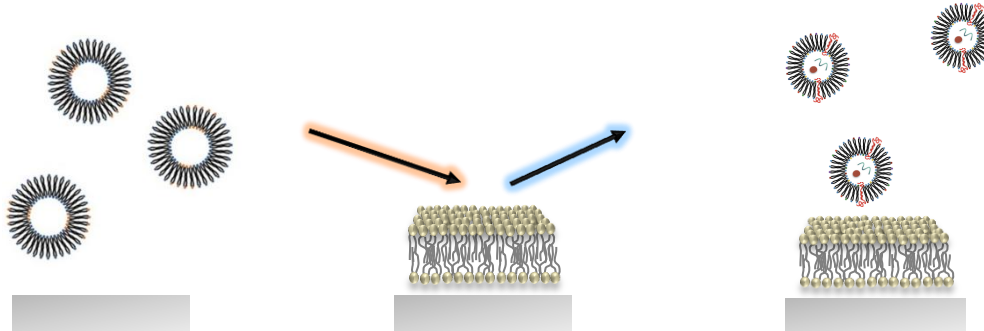
# NEUTRON REFLECTOMETRY

*dDMPC vesicles fusion*

*dDMPC characterization*

*sEVs injection from bulk water*

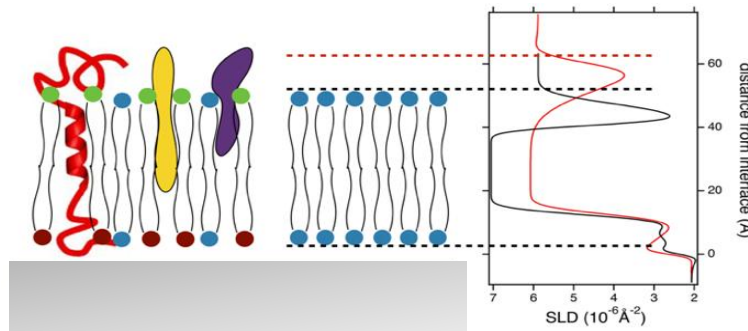
*mixed system characterization*



	AFM $\Delta Z$ (nm)	NR h (nm)
PC	$5.1 \pm 0.6$	$4.2 \pm 0.3$
PC+EVs	$6 \pm 2$	$5.4 \pm 0.3$
EVs	$9 \pm 3$	$6.9 \pm 0.3$

0.5nm water

?



- 20% volume penetration
- Change in contrast spans whole membrane thickness
- Asymmetric

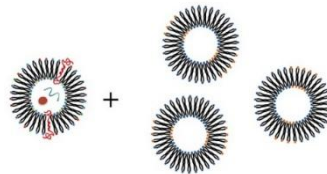
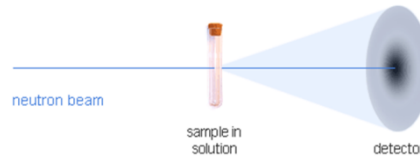
# SANS

sEVs : dDMPC Vs

1: 15000

1:3000

1:2700

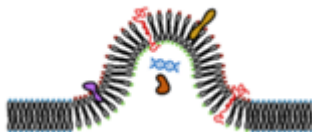
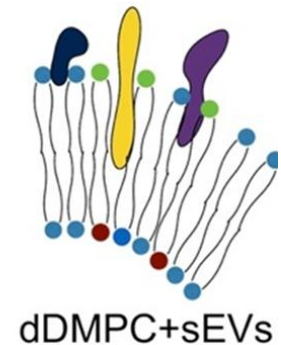
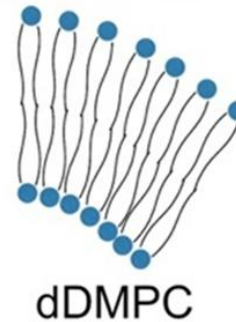
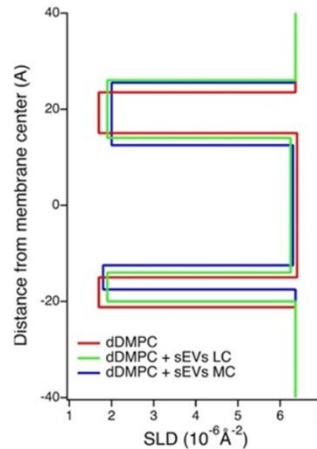
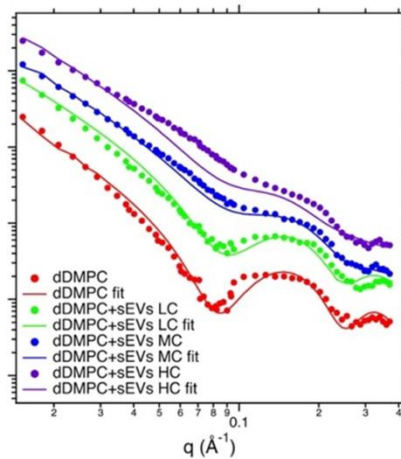


**FITTING MODEL**  
Multilayered spherical form factor

$$P(q) = \frac{\text{scale}}{V} F^2(q) + \text{background}$$

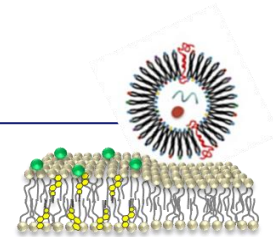
$$F(q) = \frac{3}{V_s} \left[ V_c(\rho_c - \rho_s) \frac{\sin(qr_c) - qr_c \cos(qr_c)}{(qr_c)^3} + V_s(\rho_s - \rho_{\text{solv}}) \frac{\sin(qr_s) - qr_s \cos(qr_s)}{(qr_s)^3} \right]$$

(Guinier, 1955)

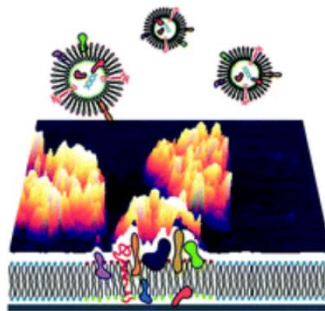


- Change in contrast spans whole membrane thickness
- Asymmetric

- ✓ Phase borders as docking sites, favouring sEVs uptake
- ✓ Separate processes can be distinguished
- ✓  $I_o$  borders granularity increases after fusion
- ✓ Patches expansion is favoured in  $I_d$  phase
- ✓ Final membrane is asymmetric



*F. Perissinotto & V. Rondelli et al.,  
Nanoscale (2021)*



The uptake process  
departs from the expected  
'simple' fusion



We can identify specific vesicle-cell uptake routes  
...eventually tunable for therapeutic needs

Miriam Grava - Sezione V - Comunicazioni

*DSC unveils new aspects of extracellular vesicles mixing with model membranes*

# Conclusion

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*Complex systems can be built up and characterized*

*X-ray and neutron small angle scattering and reflectometry are essential complementary techniques for cross-structural investigations down to the nanoscale of self assembled systems and of thick interfaces*

*Possibility to work in physiological conditions*

*Possibility for in-situ interaction studies*

*Perspectives in biology (and not only....) are numerous*

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