



The Use of Liposomes in the Assessment of Antimicrobial Activity of Membrane-Active Compounds

Christian Nehls

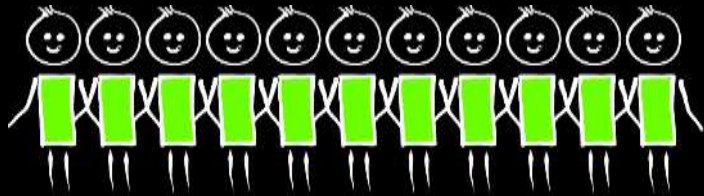
PRC-Webinar 10.12.2025



Antimicrobial peptides with therapeutic potential

Estimated deaths due to antimicrobial resistance

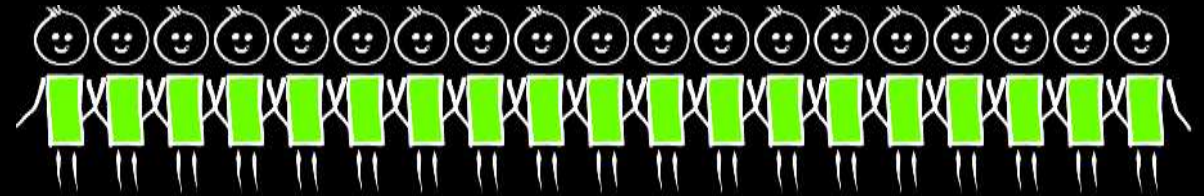
2021



1.14 Mio.

(4.71 Mio. associated)

2050



1.91 Mio.

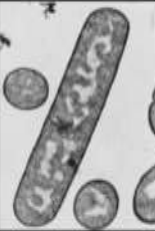
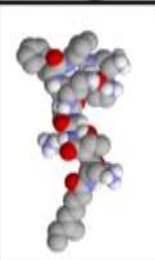
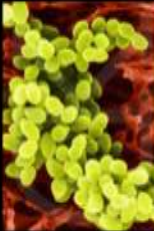
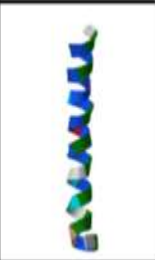
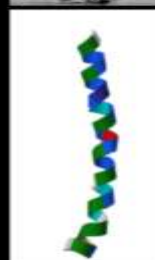
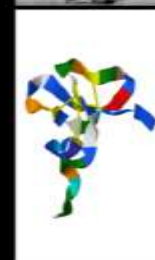
(8.22 Mio. associated)



= 100 000 estimated annual deaths

New treatment
options are needed

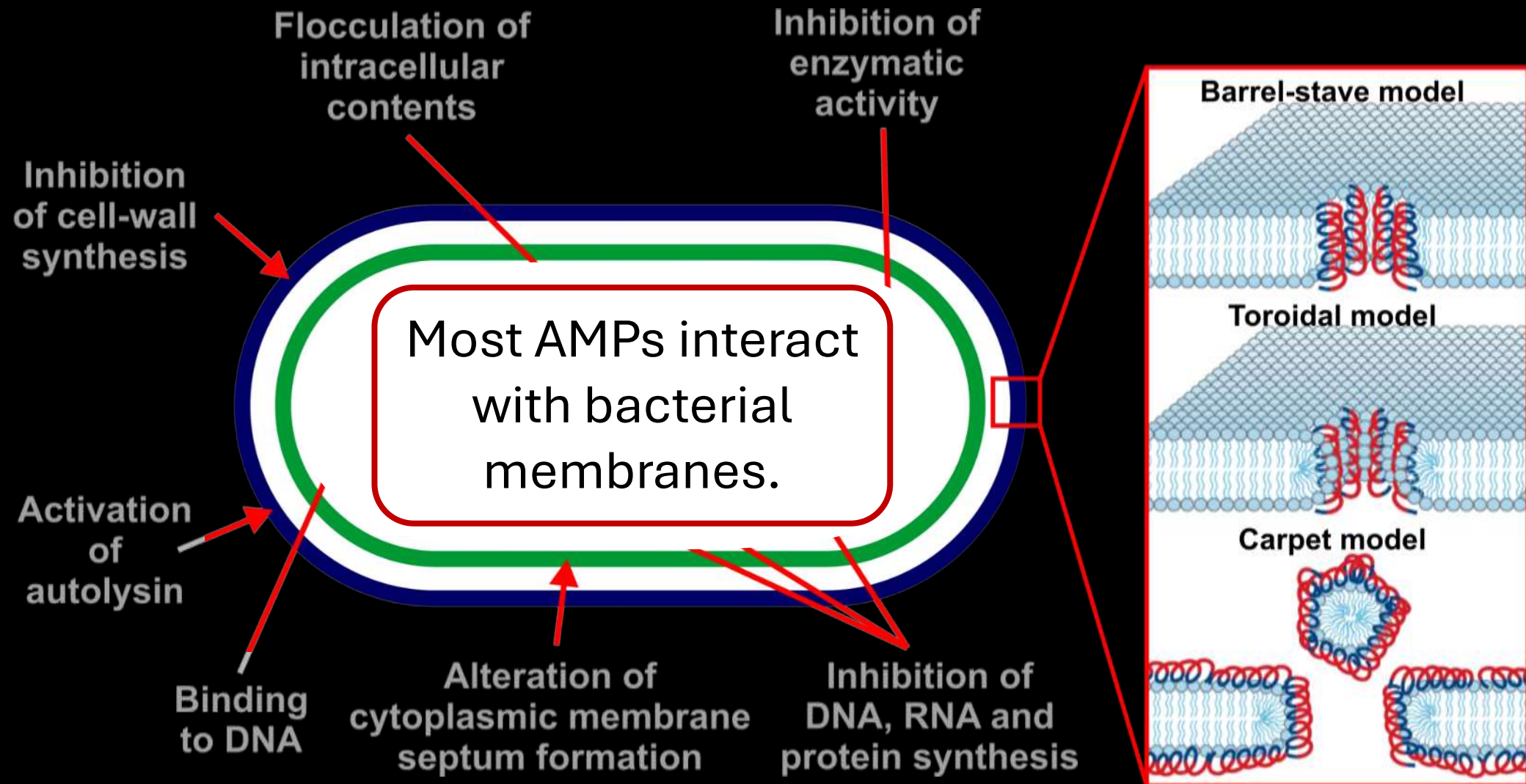
Occurrence of antimicrobial peptides (AMPs)

Polymyxin B  	Lactacin  	Amoebapore  	Hydramacin  	Caenopore  	Arenicin  
rCAP18  	NK-lysin  	hCAP18  	beta-defensins  	Psoriasin  	rBPI₂₁  

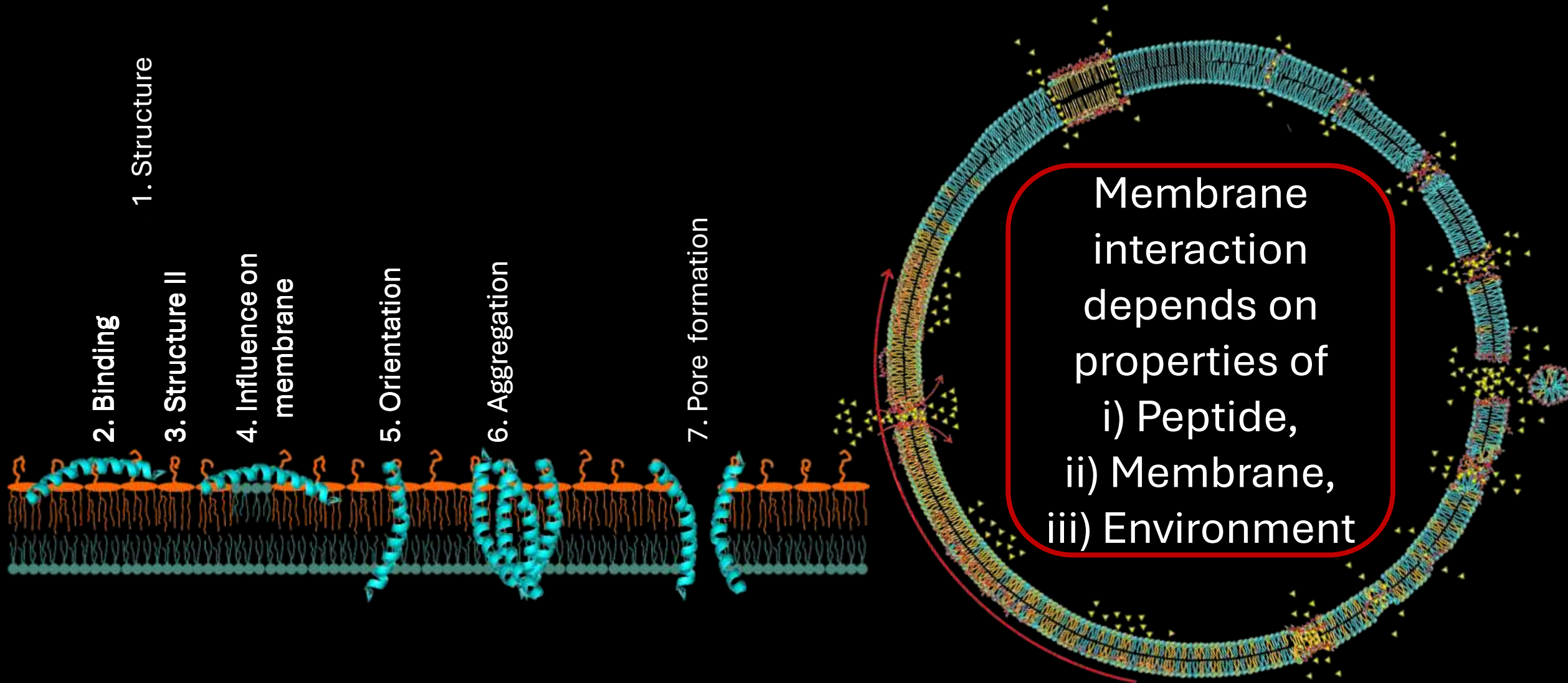
- Size
- Sequence
- Charge
- Dipole moment
- Flexibility
- Hydrophobicity
- Amphipathicity
- Conformation and structure

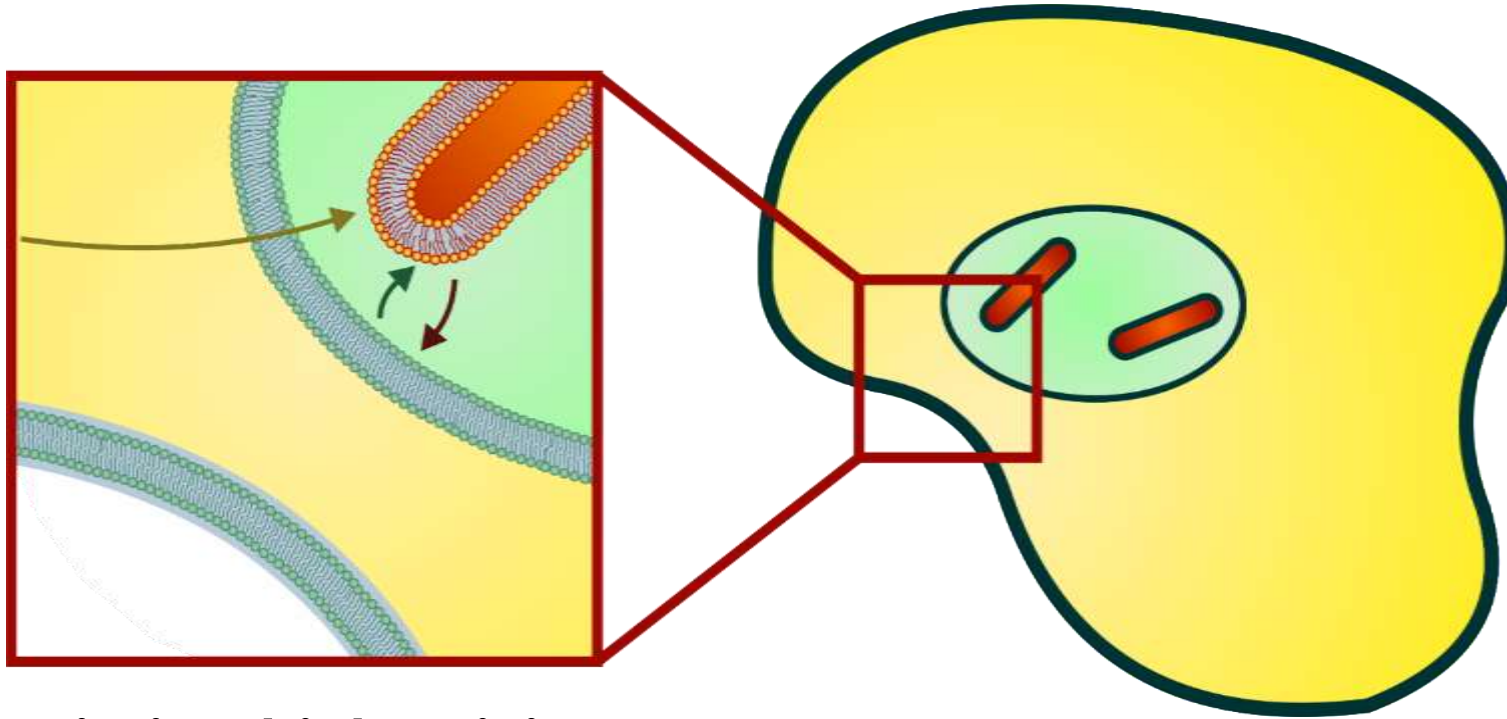
AMPs occur in a
large diversity

Modes of Action of AMPs



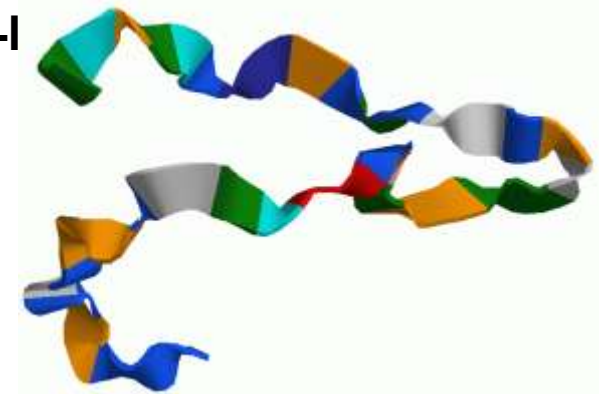
AMP membrane interaction





hBD-3

hBD-3-I

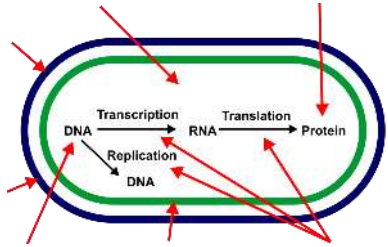


Antimicrobial activity:

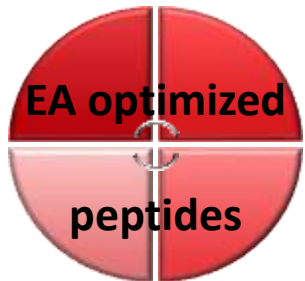
- I. Bacteriostatic: Lactoferrin (Nutrient deprivation)
- II. Bactericidal: Defensins (Membrane permeabilization)**
Cathelicidins (Membrane permeabilization)
- III. Hydrolysis: Lysozyme, ... (Carbohydrates, ...)



AMPs are a promising alternative to classical antibiotics with a much lower tendency for resistance induction.



Membrane interaction is prerequisite for any AMP-pathogen interaction.



Specificity of AMPs can be narrowed using directed optimization.



Where, who and what



Manor house from 1751



New NRC building from 2022



New lab building from 2024





CSSB
Centre for Structural
Systems Biology



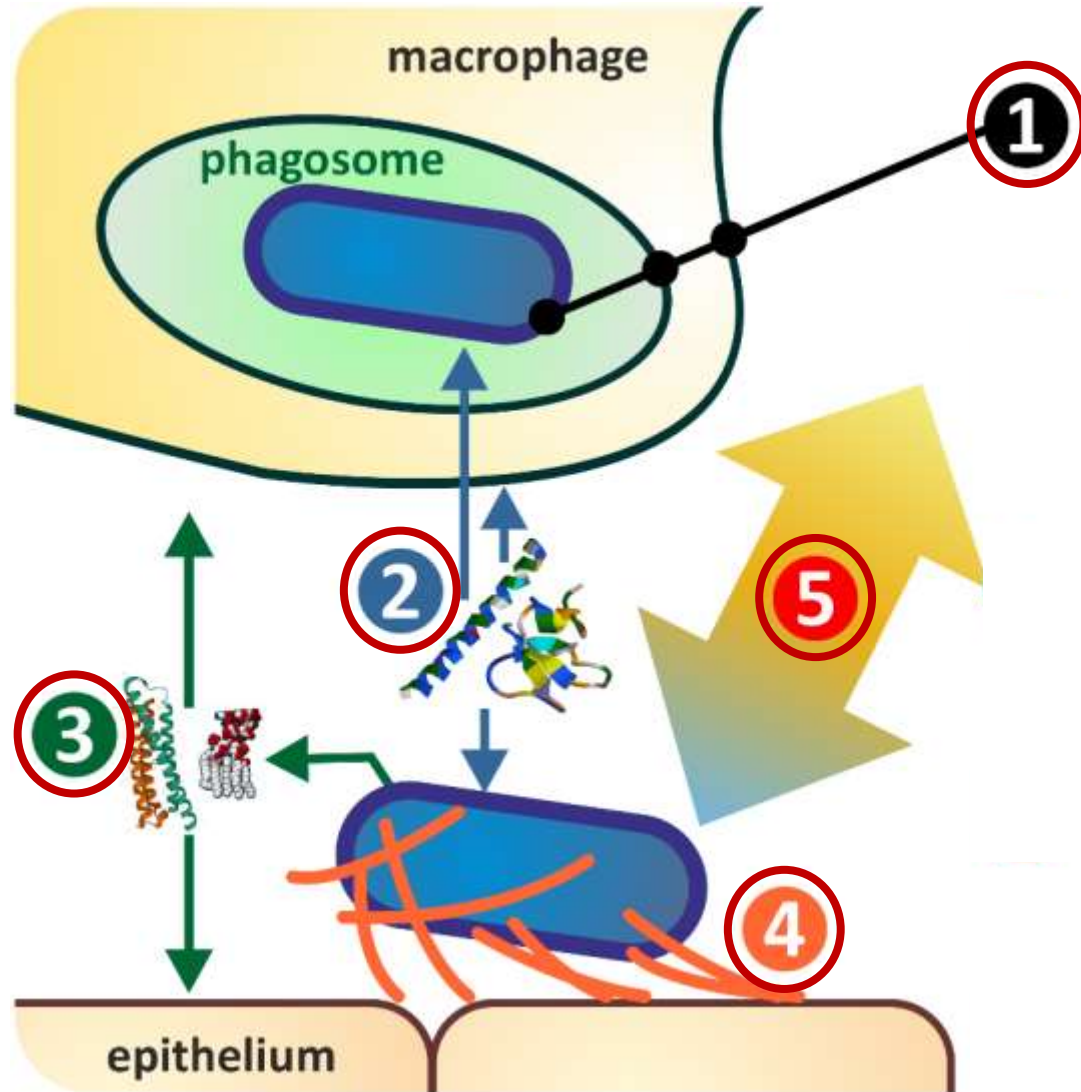
First CSSB Symposium
2015

Gutsmann lab @ Research Center Borstel



Team Microfluidics





To develop new therapies, we

investigate membrane properties and interactions in the context of lung disease.

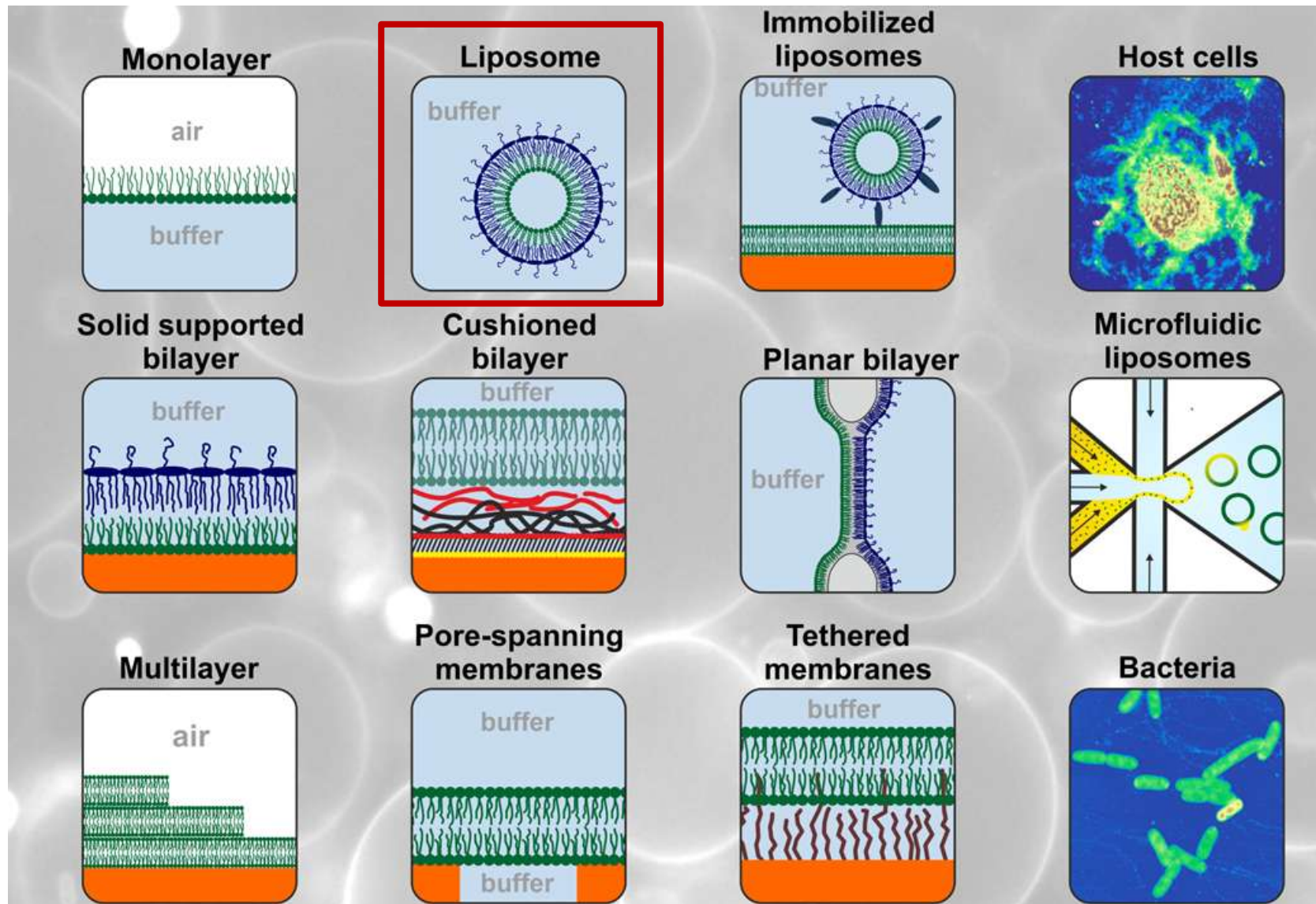
develop antimicrobials to combat antimicrobial resistance.

explain mechanisms of microbial toxin membrane activity.

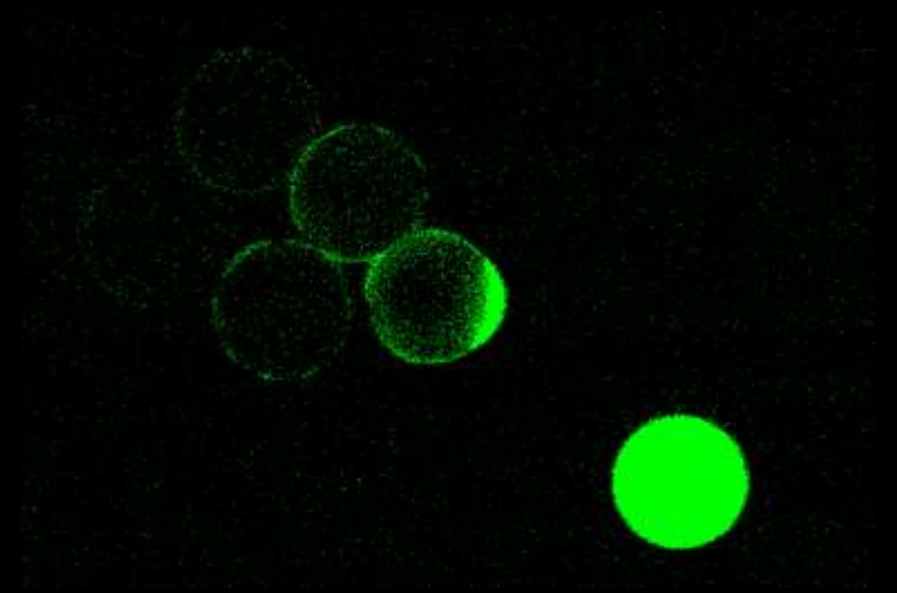
probe molecular adhesion mechanisms that underlie pathogen persistence.

generate, manipulate, characterize and detect bioactive aerosols.

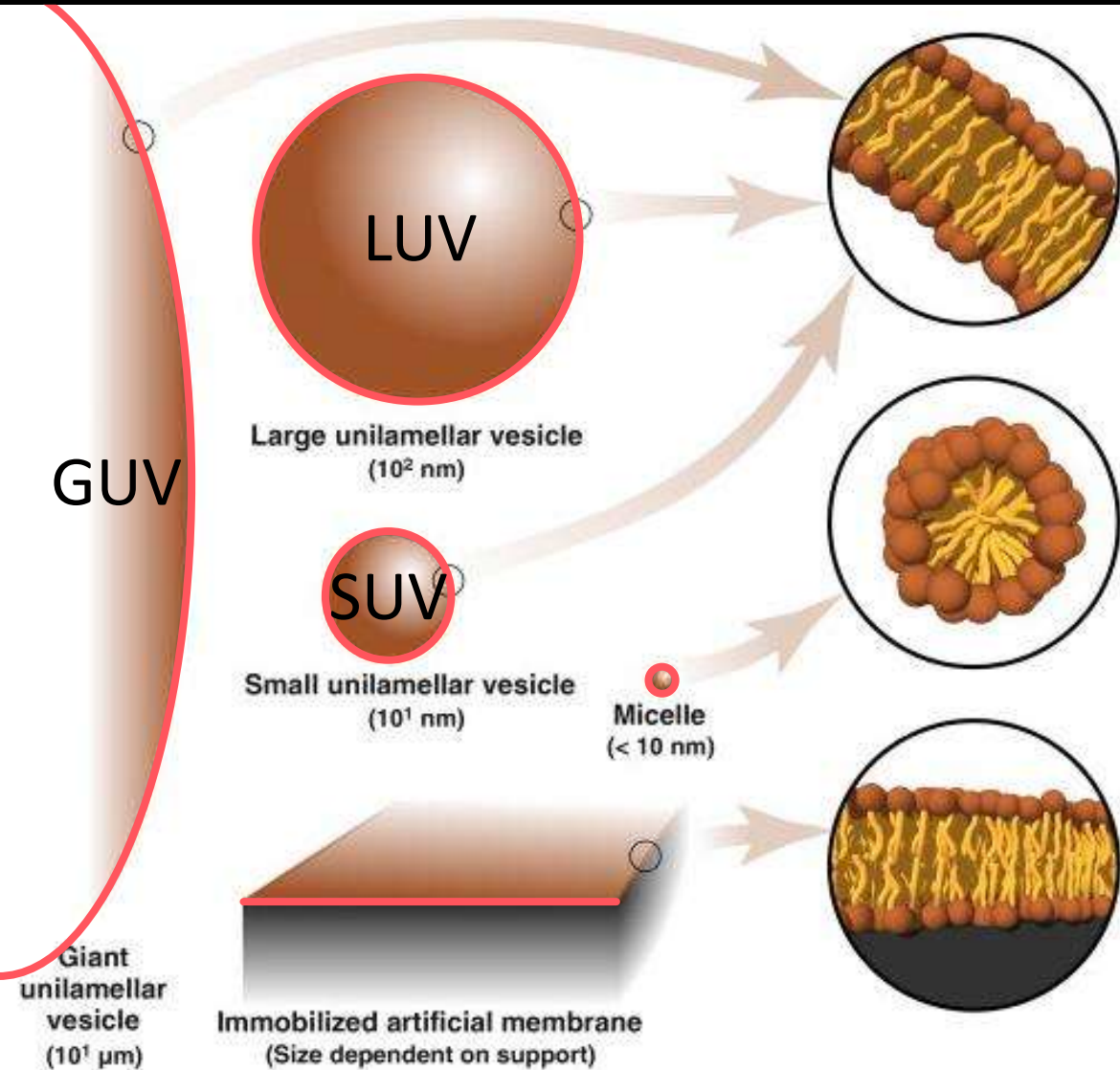
Model membrane systems



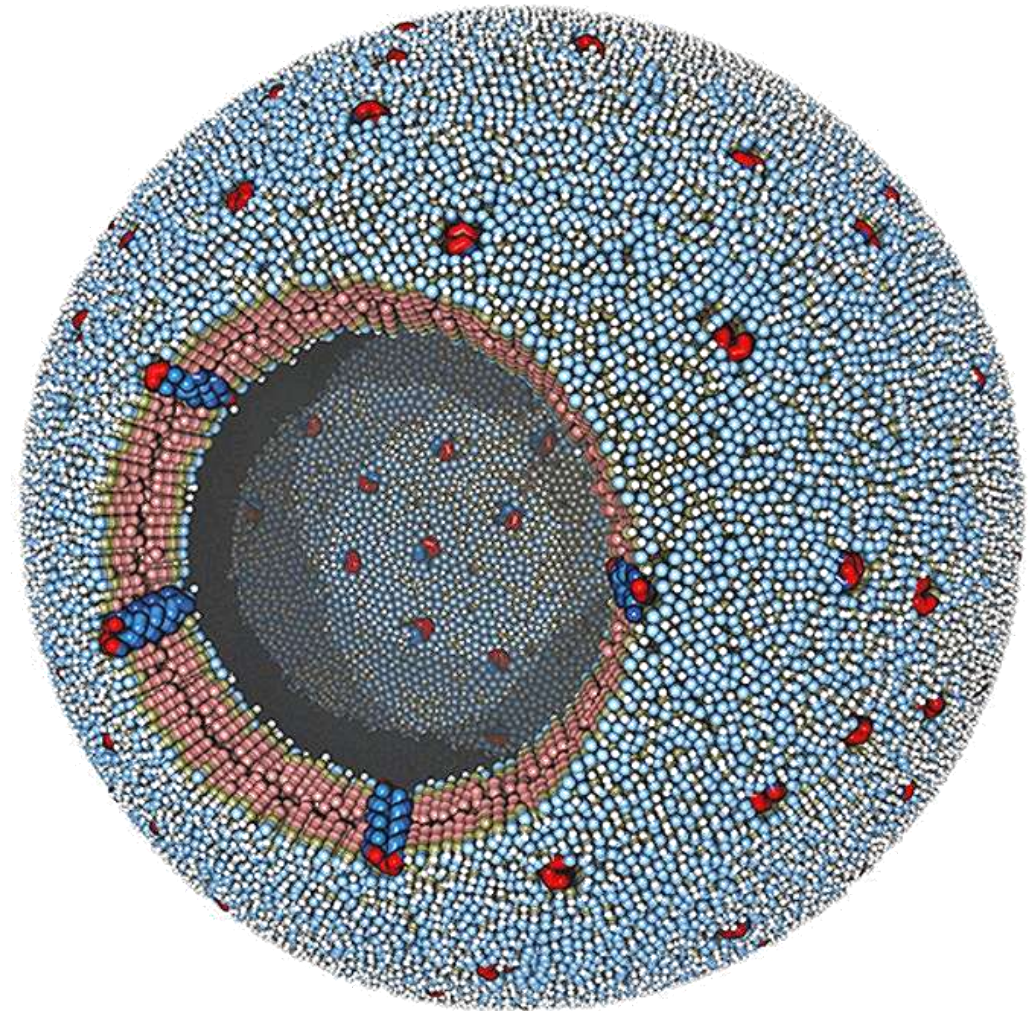
Liposome preparation



Size order of liposomes

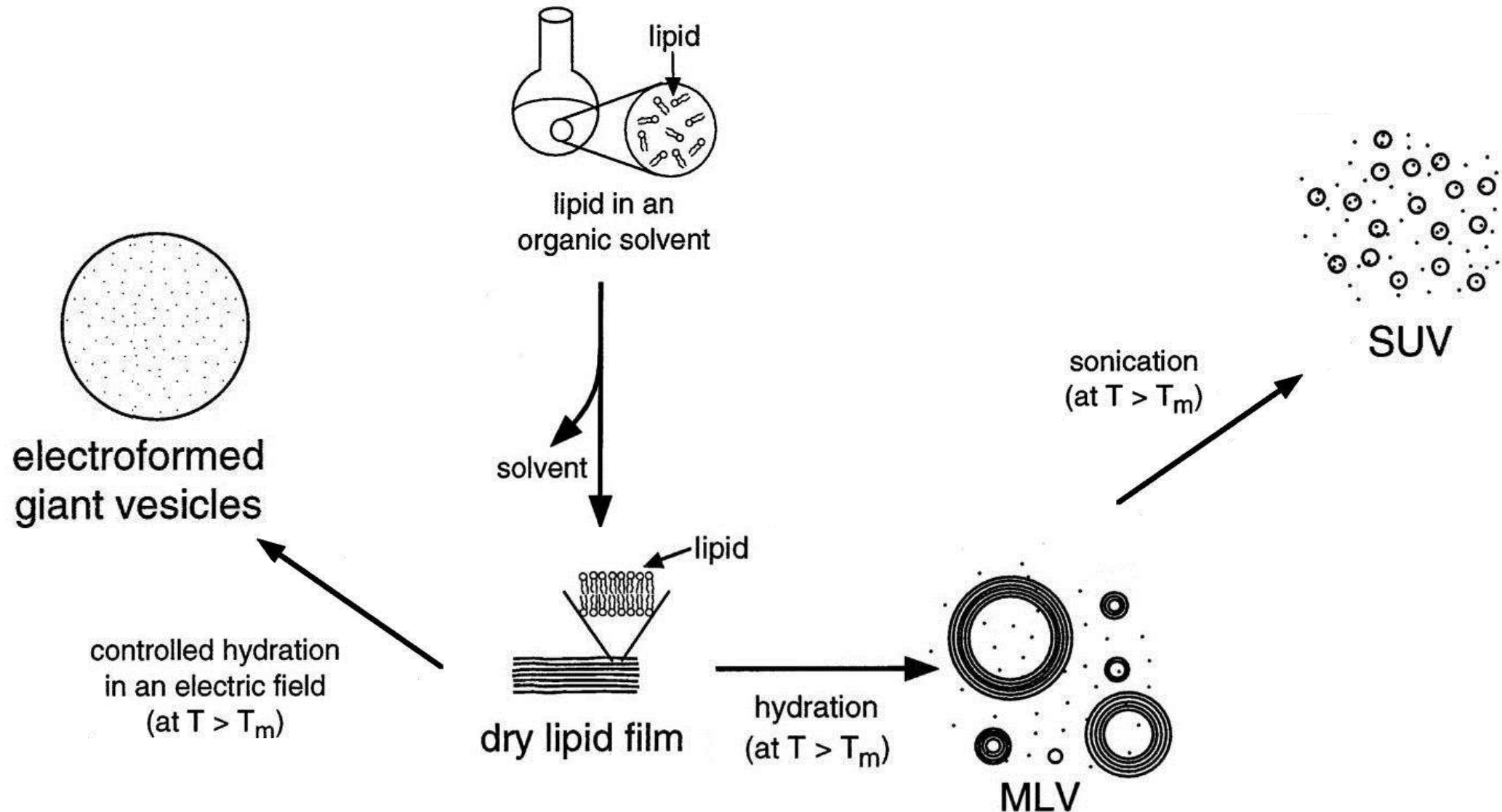


TRENDS in Pharmacological Sciences

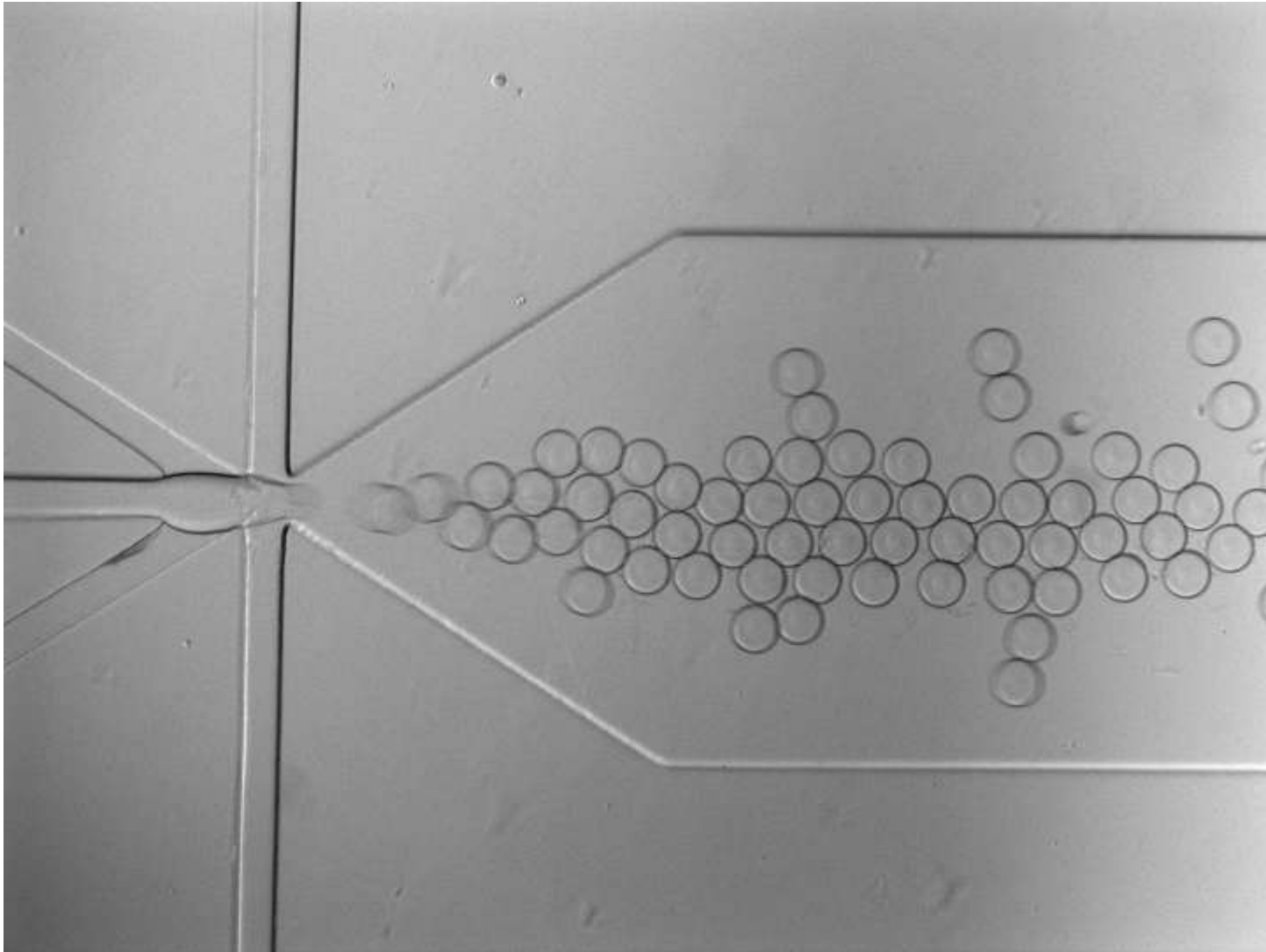


$d = 34 \text{ nm}, \approx 40.000 \text{ lipids}$

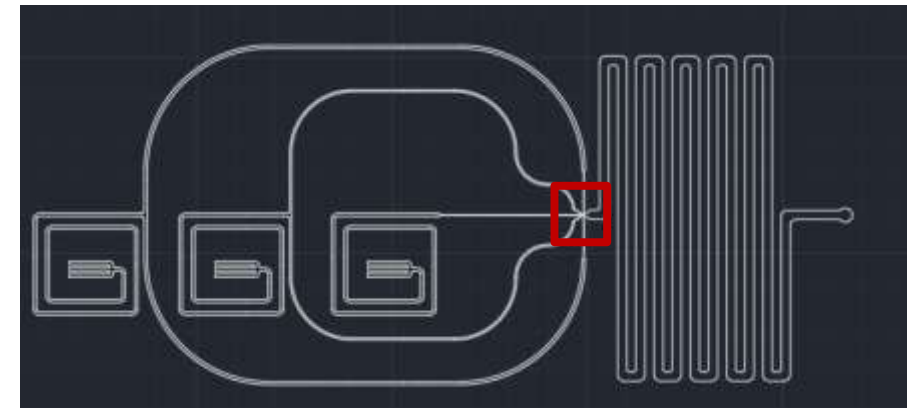
Conventional preparation of liposomes



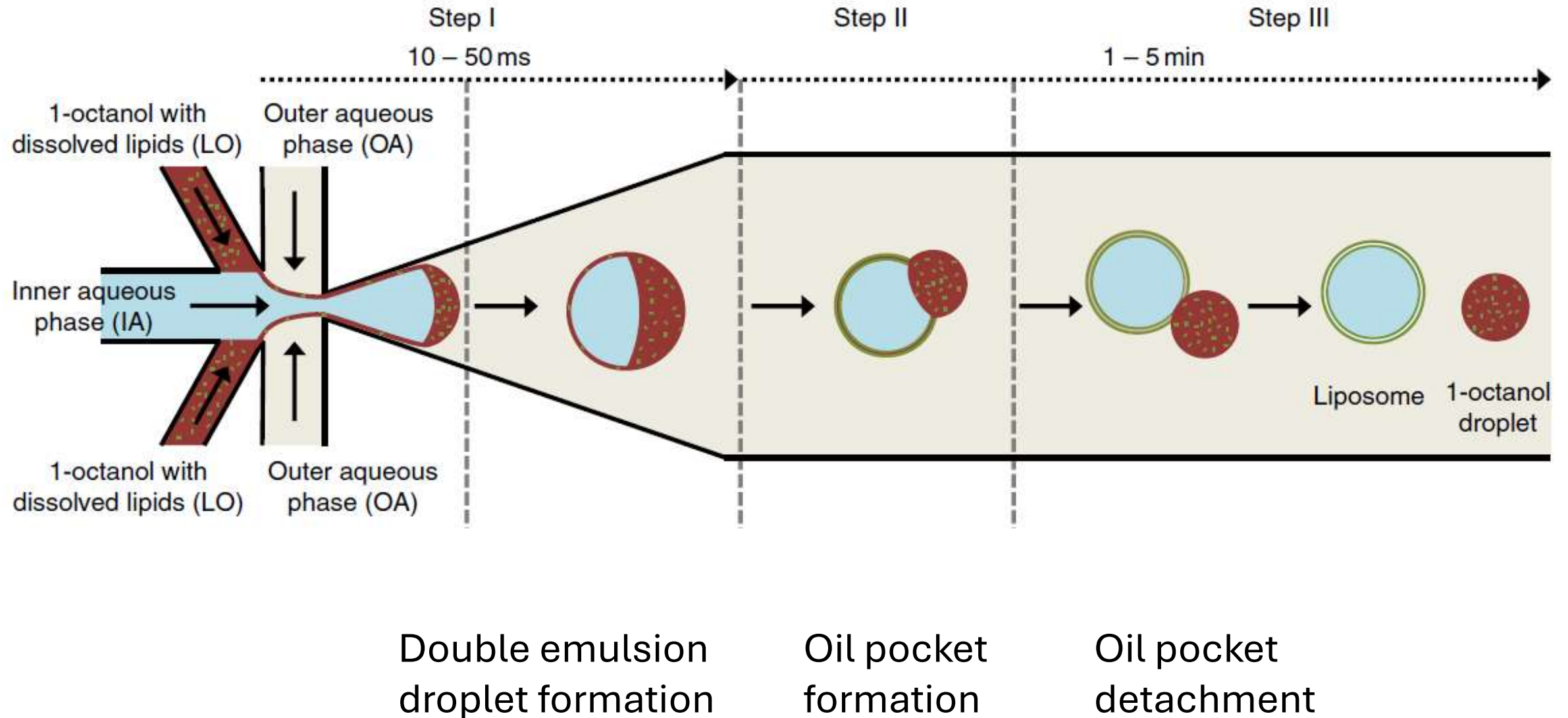
Formation of giant liposomes on microfluidic chip



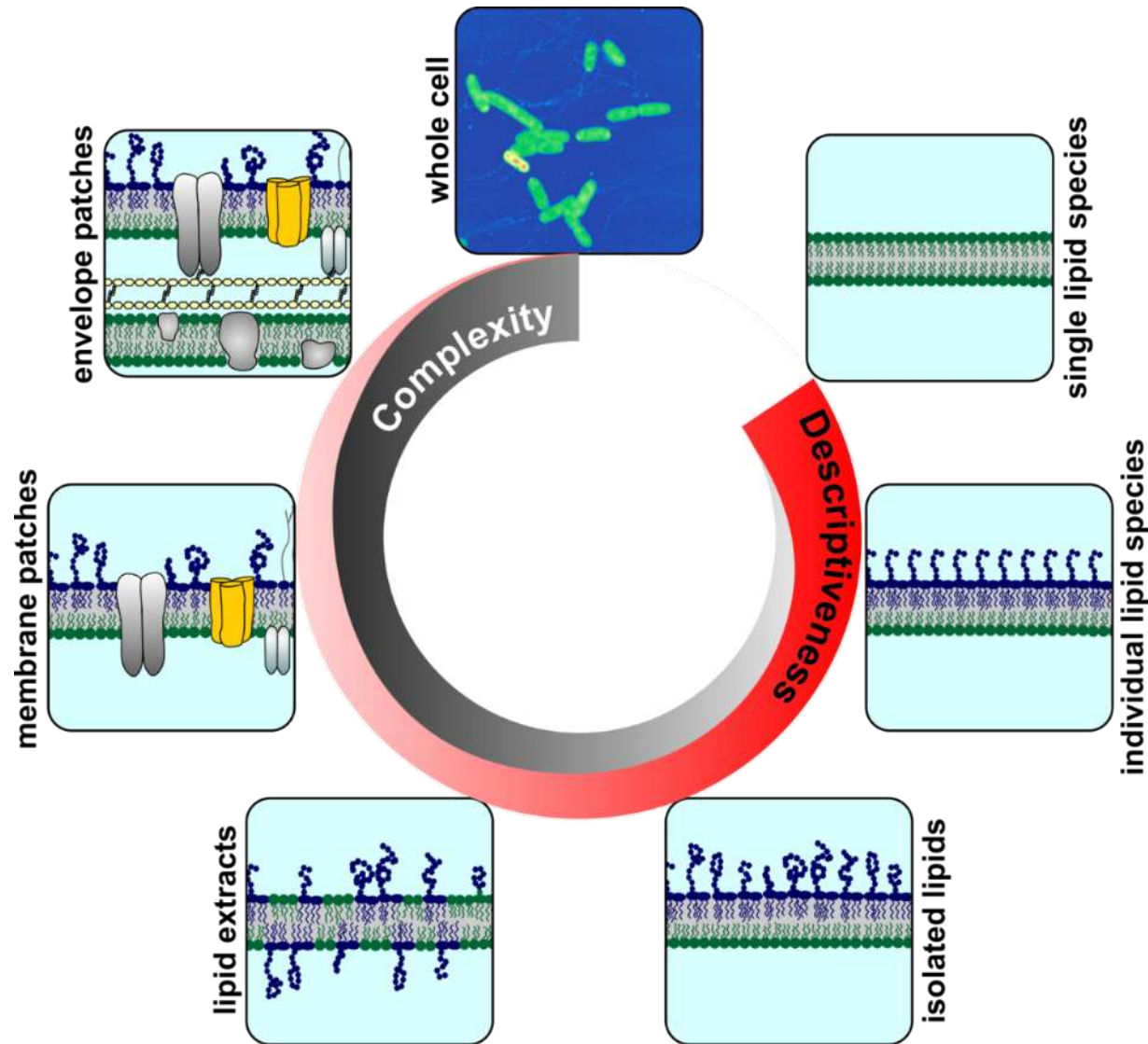
- High throughput
- Uniform size
- High encapsulation efficiency



Octanol-assisted liposome assembly on chip



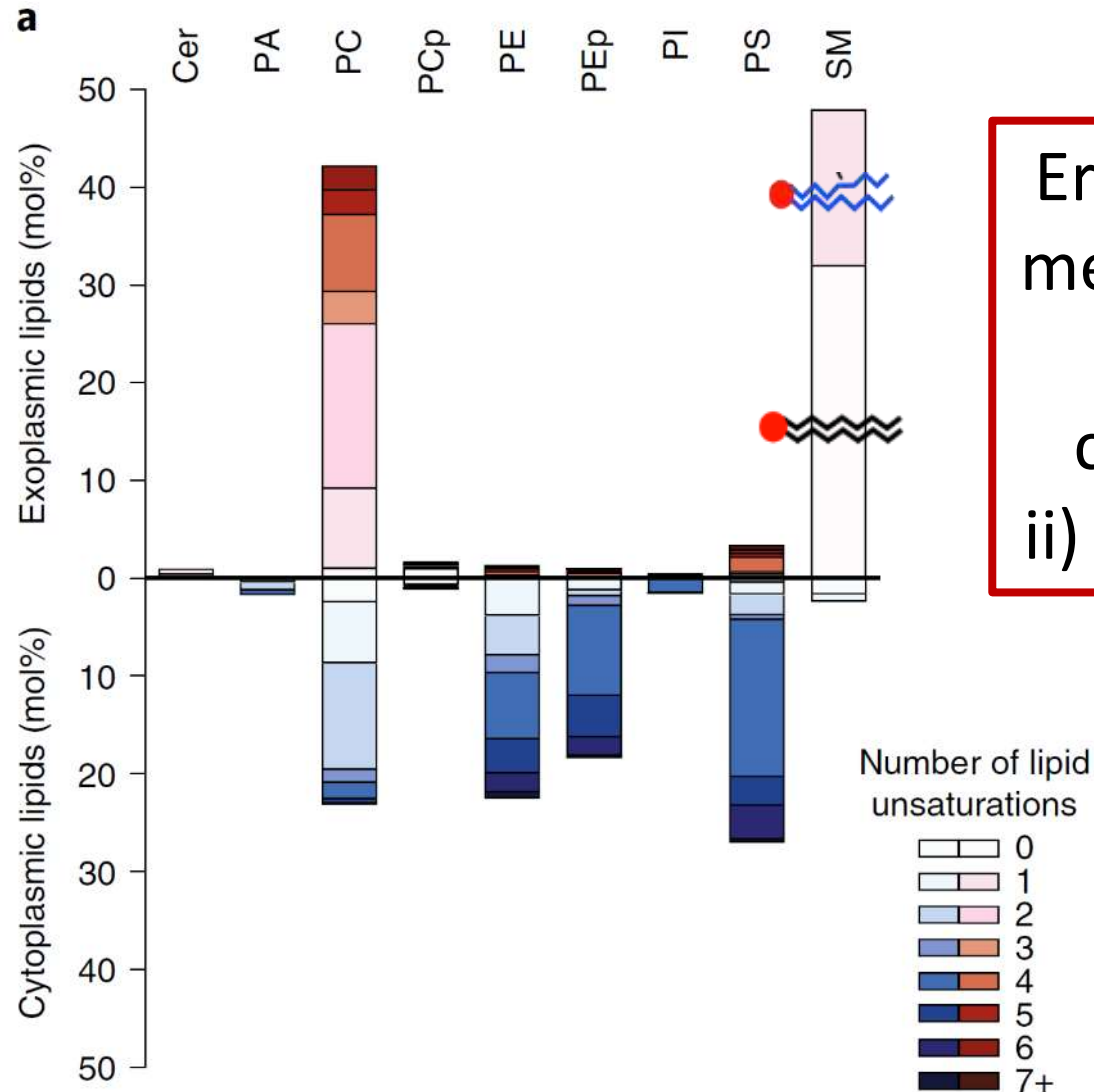
How complex should we make the system?



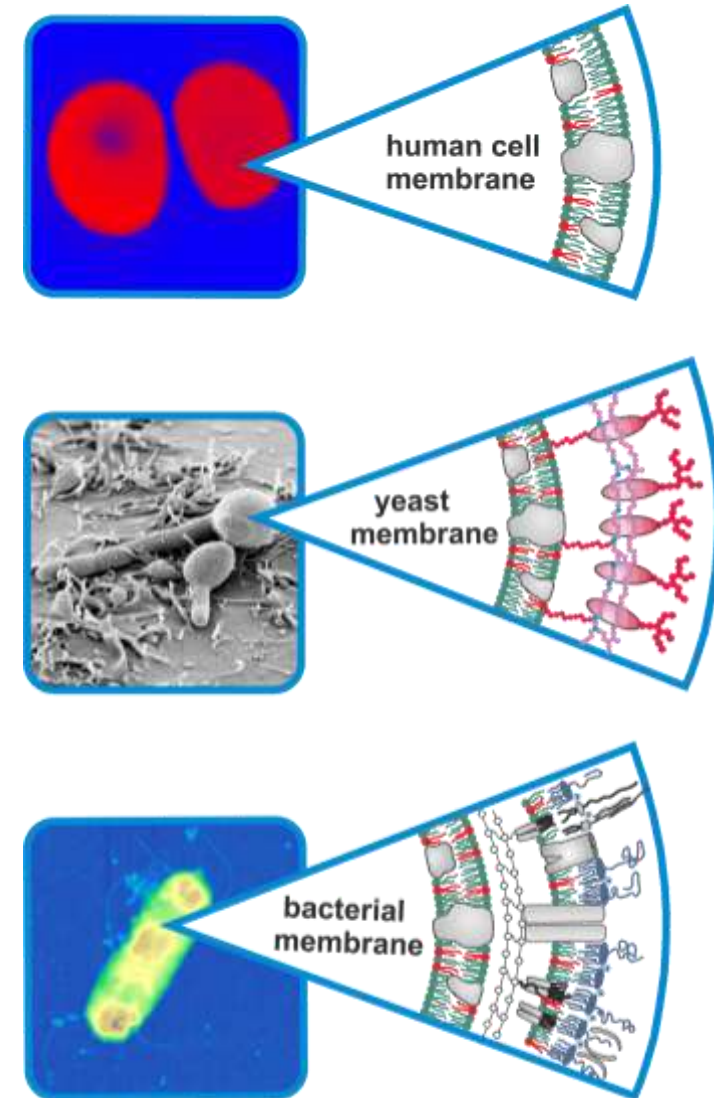
Successive complexity increase (bottom-up) adds factors for unspecific membrane interaction

Successive complexity decrease (top-down) can reveal causality for specific interaction

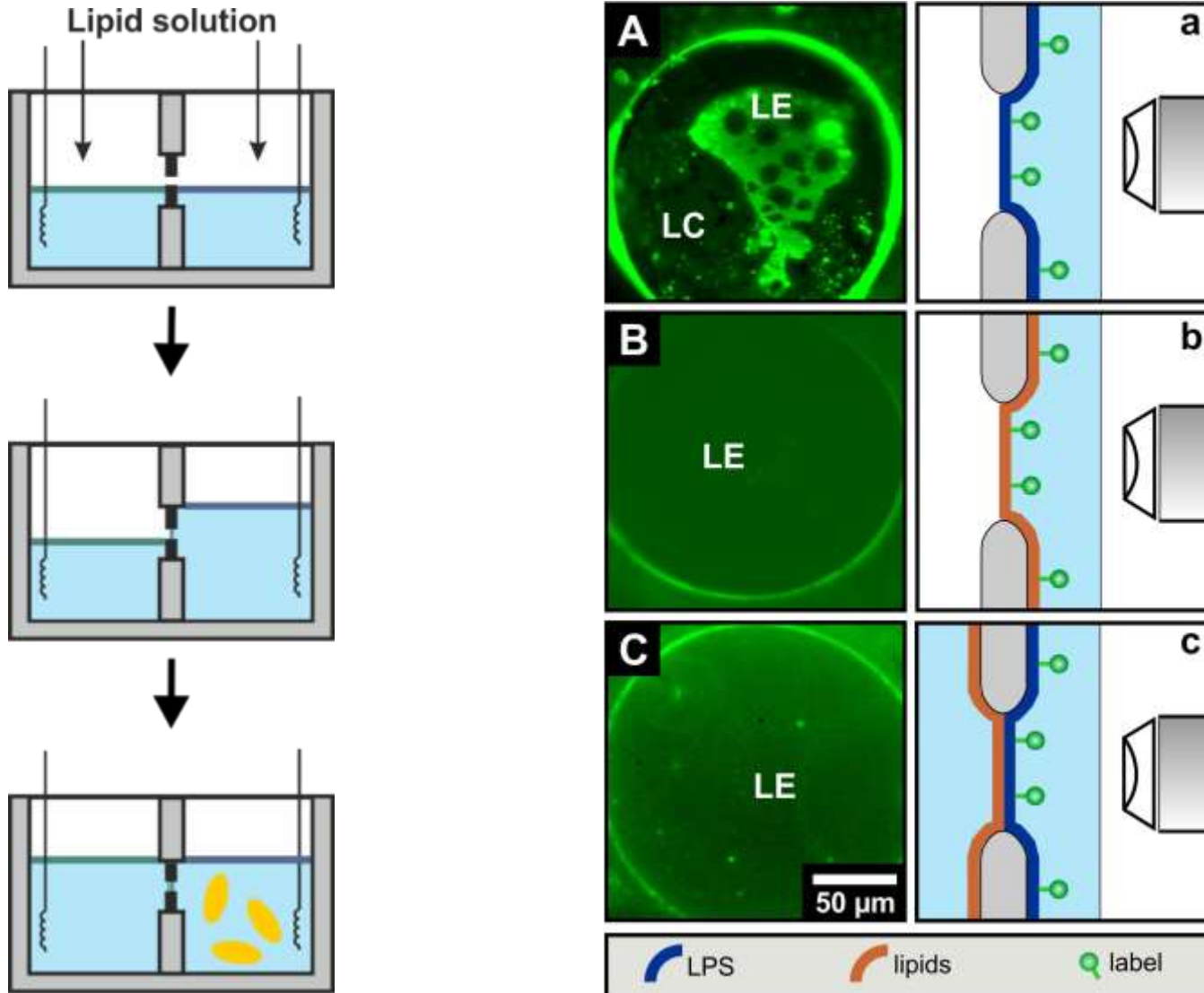
Which natural membranes are asymmetric?



Erythrocyte plasma membranes differ in
i) headgroup composition and
ii) lipid unsaturation

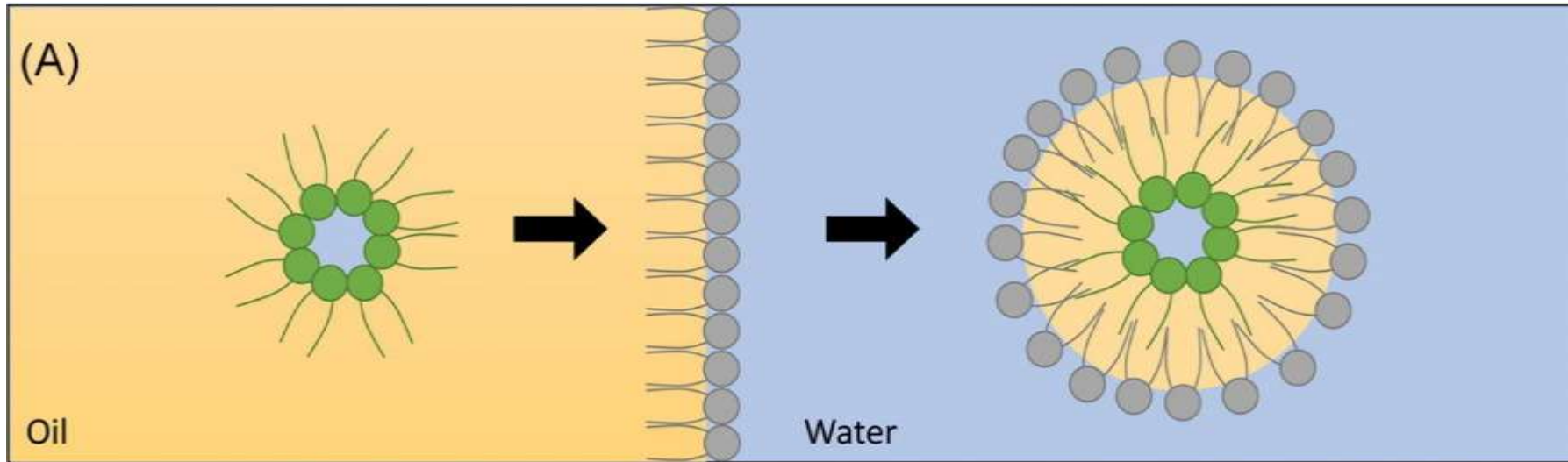


Montal-Mueller technique for asymmetric membranes

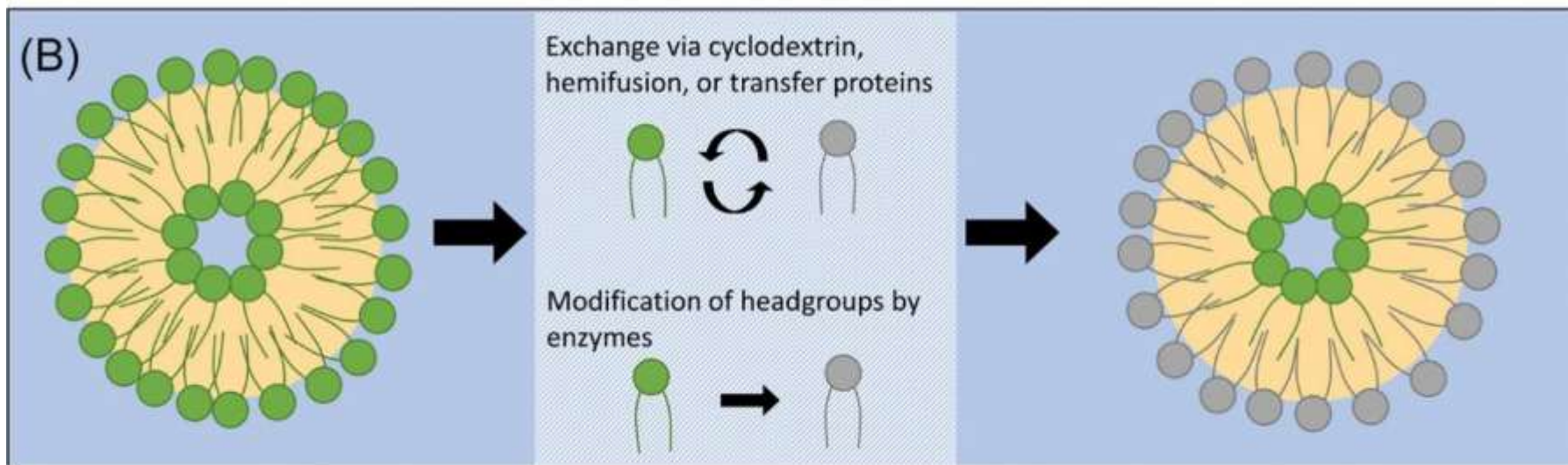


Asymmetric
membranes are more
than addition of two
monolayers

Formation of asymmetric liposomes

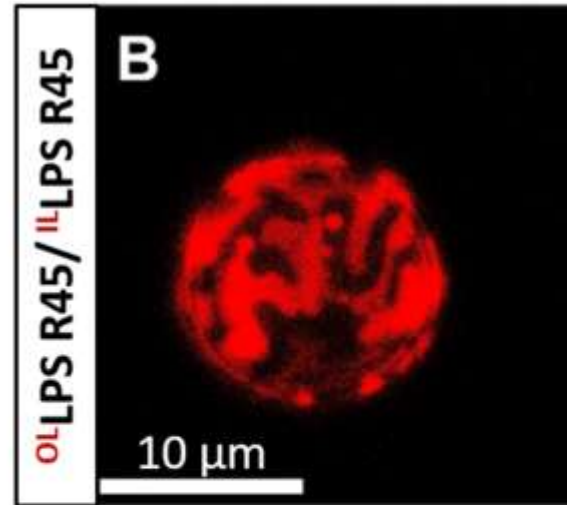
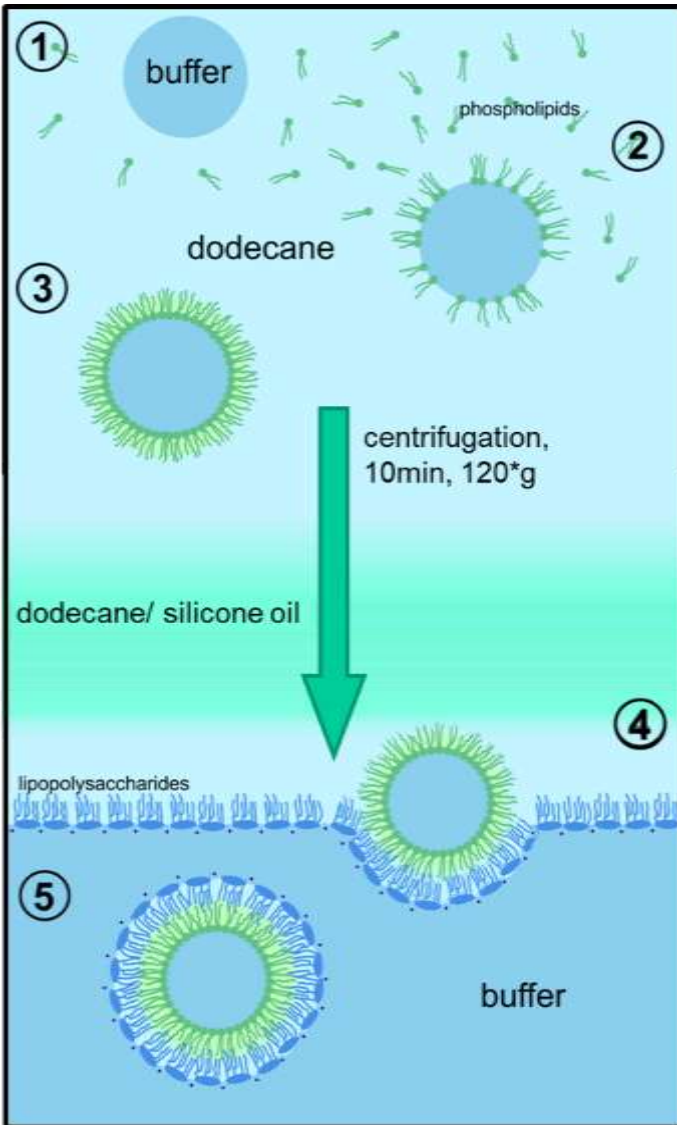


Layer-by-layer
assembly

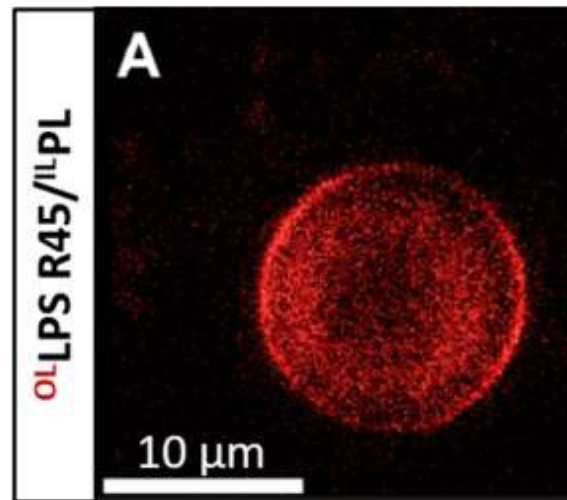


Outer leaflet
exchange

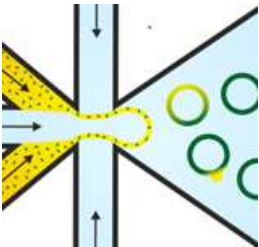
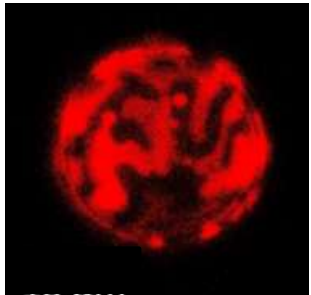
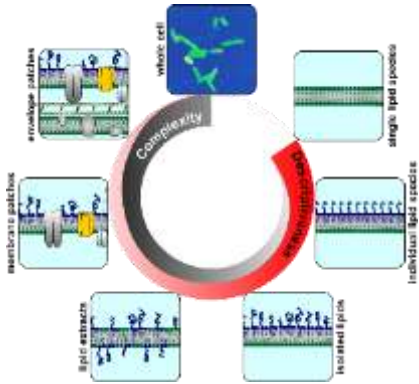
Trends in Biochemical Sciences



Phase transfer
derived liposomes
confirm relevance of
asymmetry



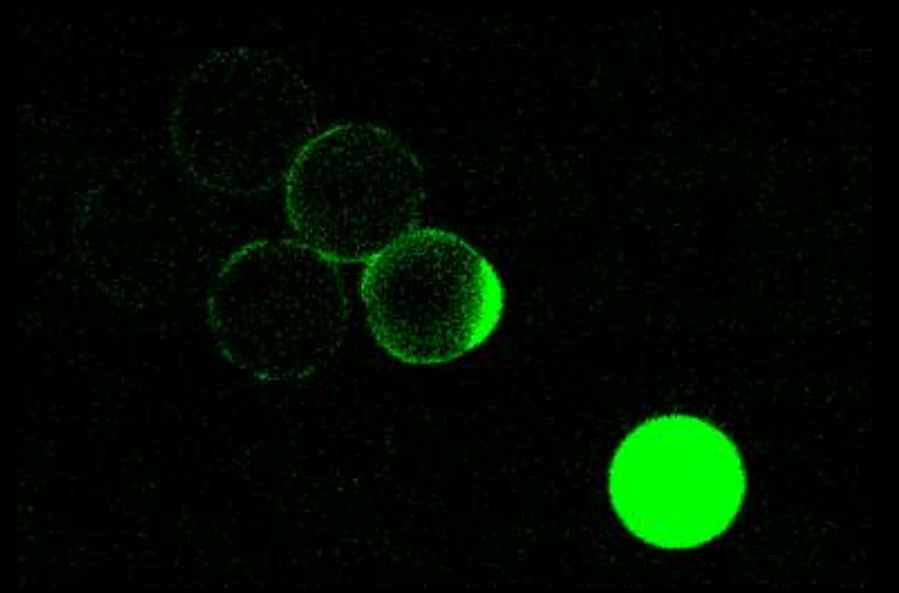
Phase transfer allows
formation of pure
lipopolysaccharide
liposomes



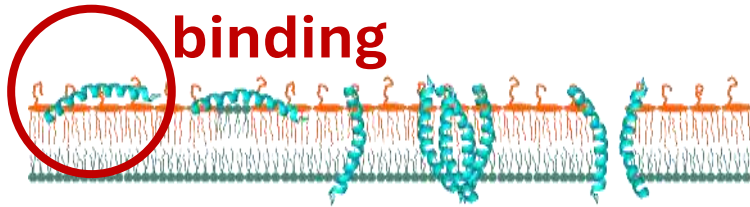
To understand interaction processes at membranes, the natural membrane complexity has to be reduced.

Membrane asymmetry is directly linked to biological function and has to be considered for model systems.

Microfluidic techniques allow homogeneous GUV generation with exceptional encapsulation efficiency.

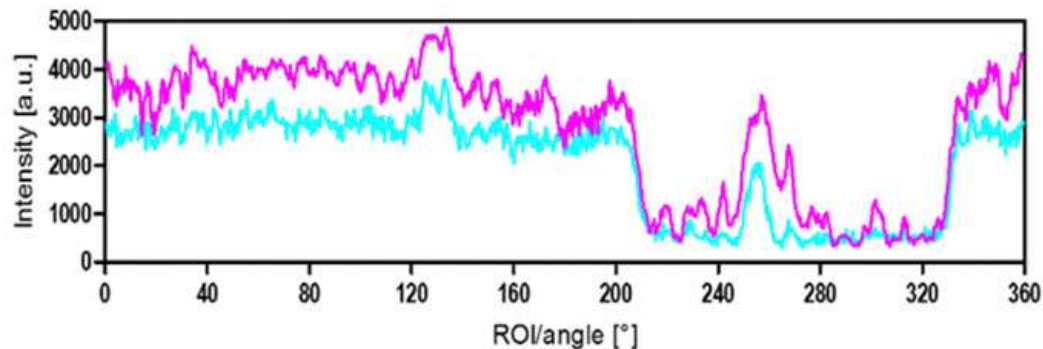
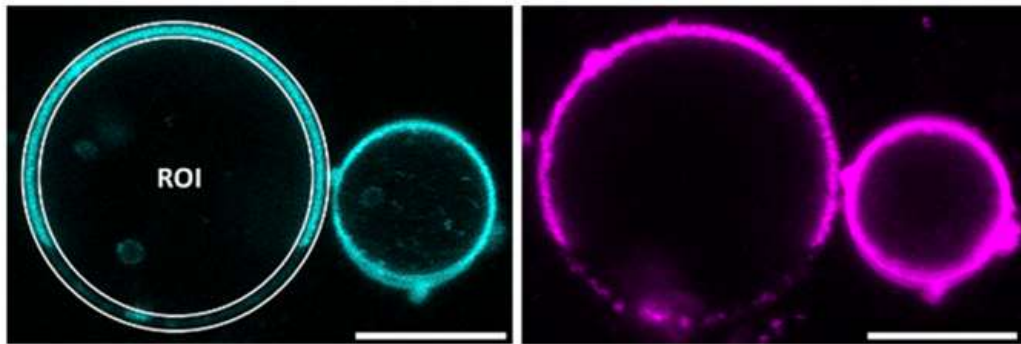


Evaluation of AMPs with liposome techniques

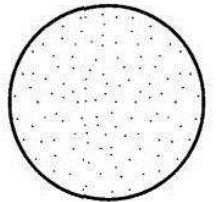
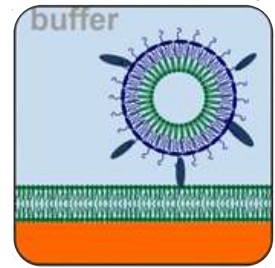
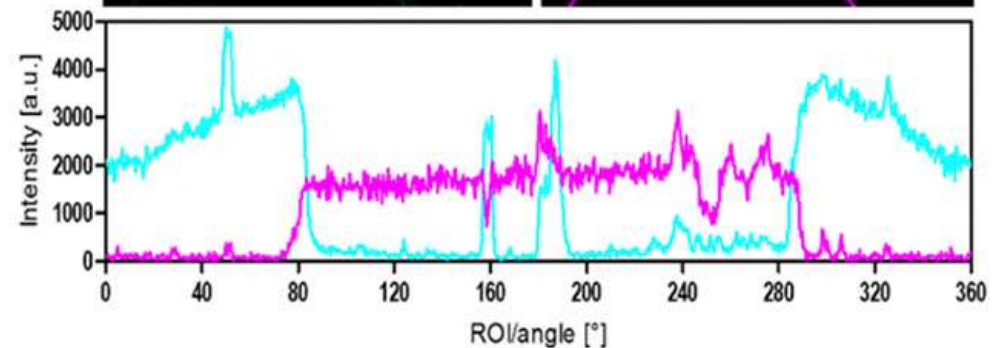
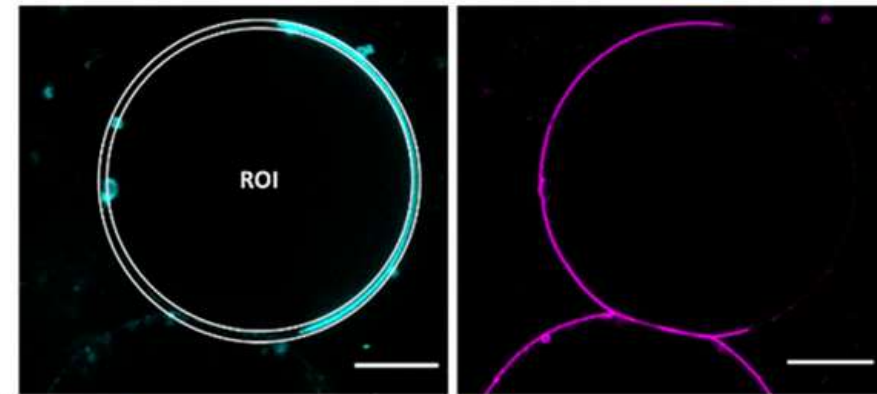


Specific binding to lipid domains
determines AMP mode of action

A — I_d -domain — LL-32

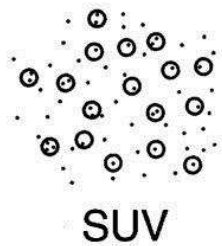
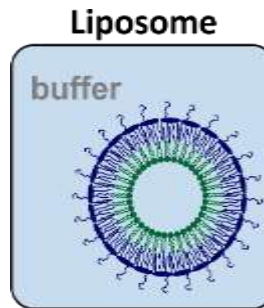
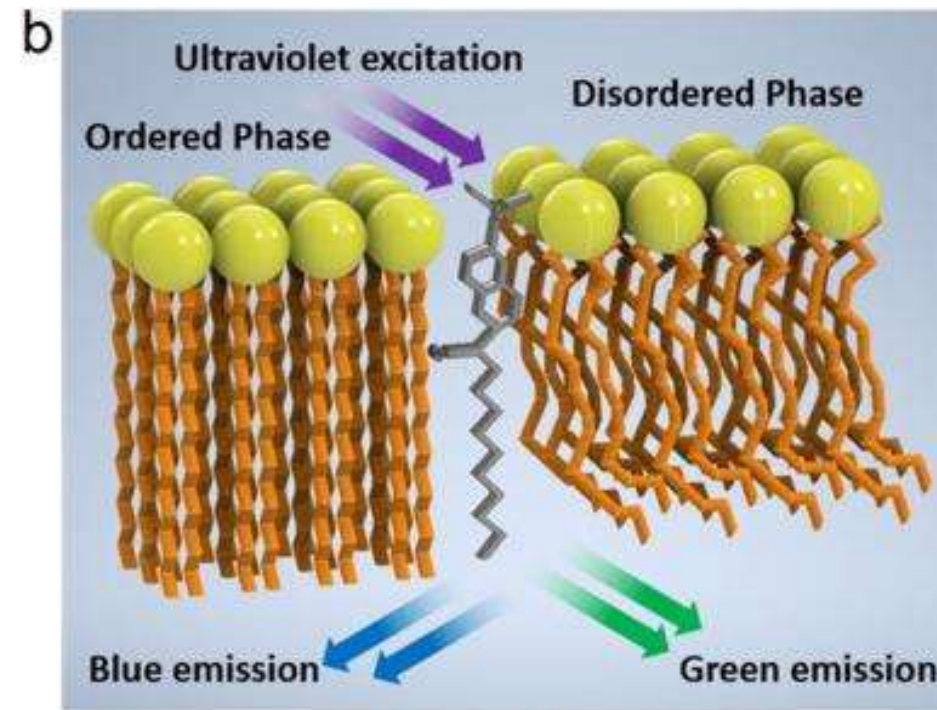
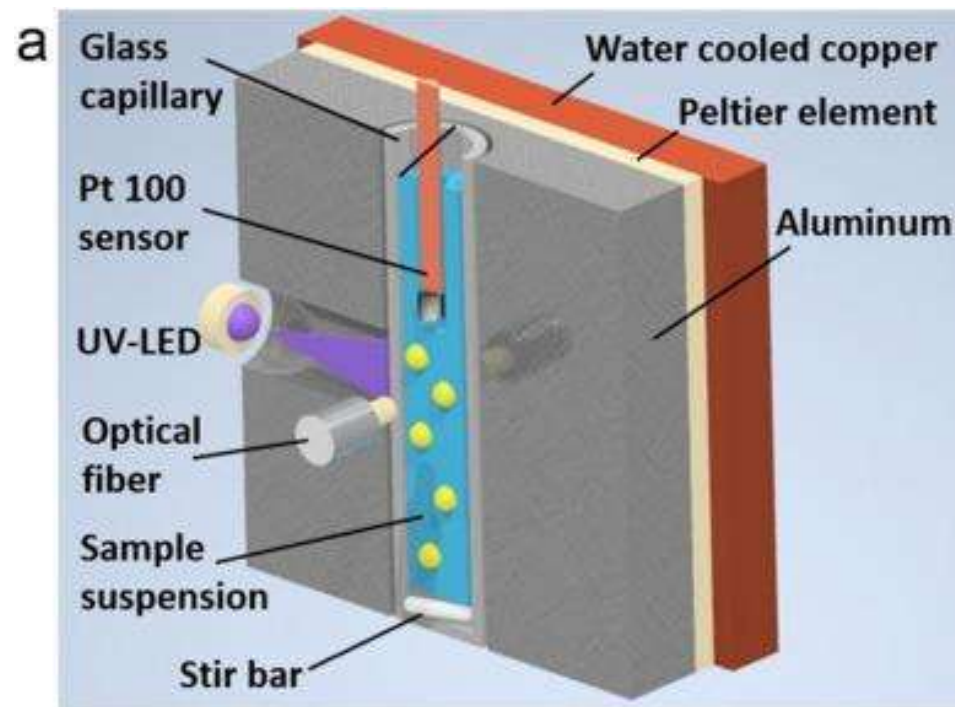
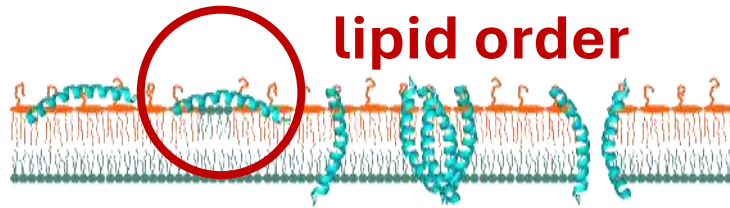


B — I_d -domain — PMB

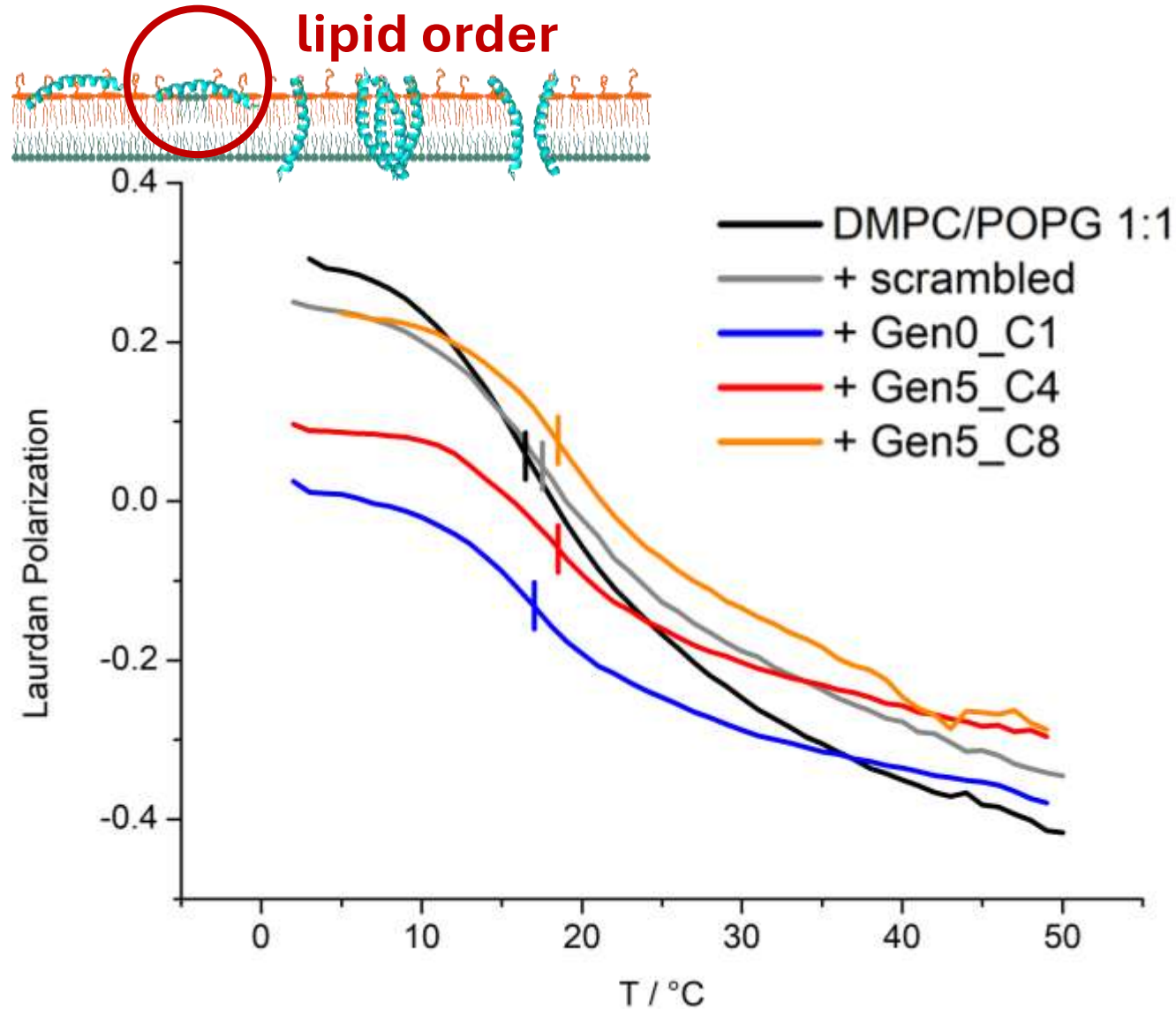


electroformed
giant vesicles

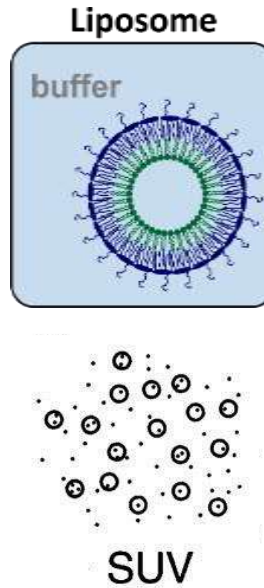
Lipid State Observer LISO on SUVs/LUVs



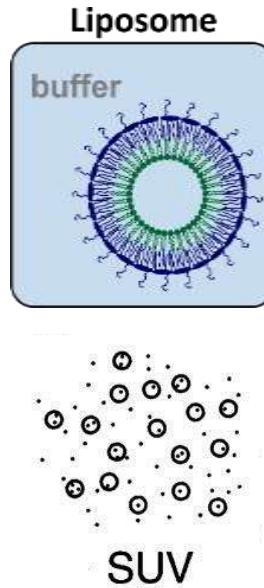
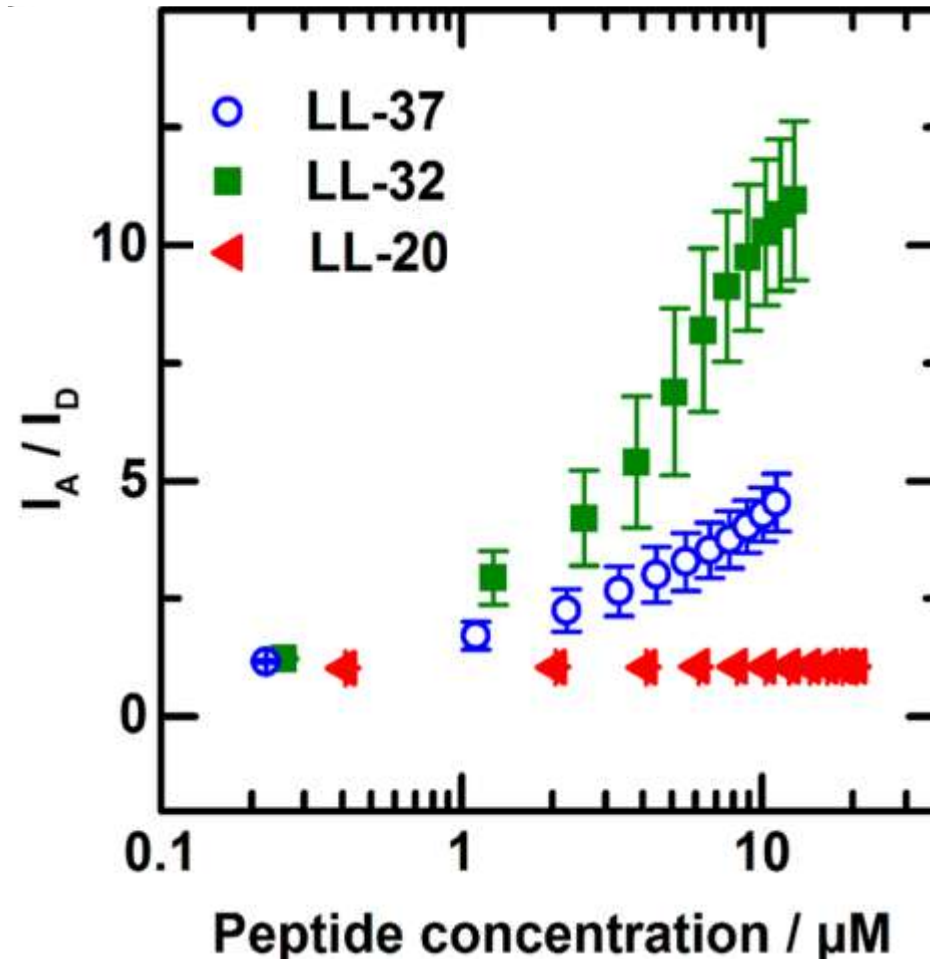
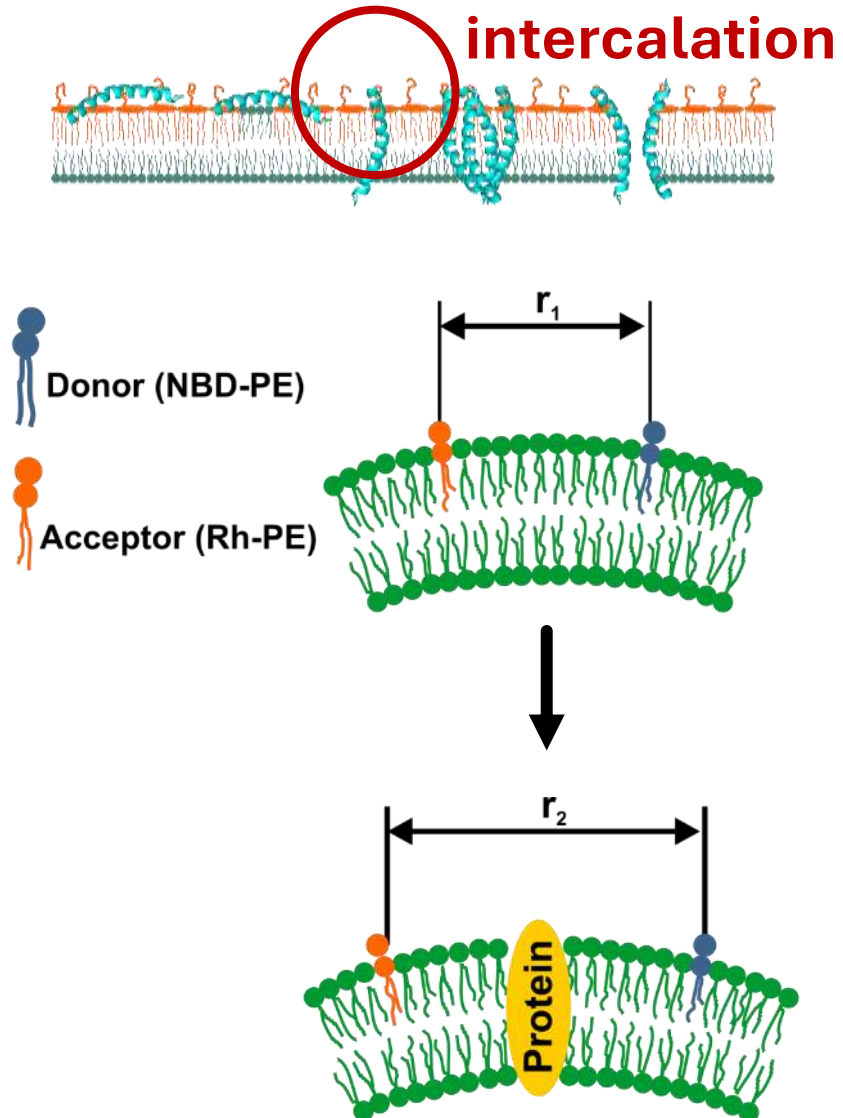
Lipid order and phase transition: LISO – Lipid State Observer



More successful EA
AMPs (Gen5)
i) level lipid order and
ii) shift phase
transition to higher T

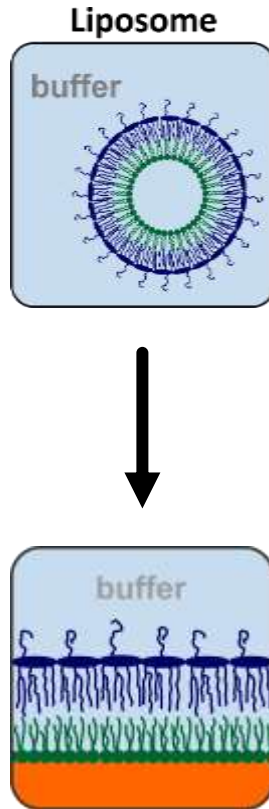
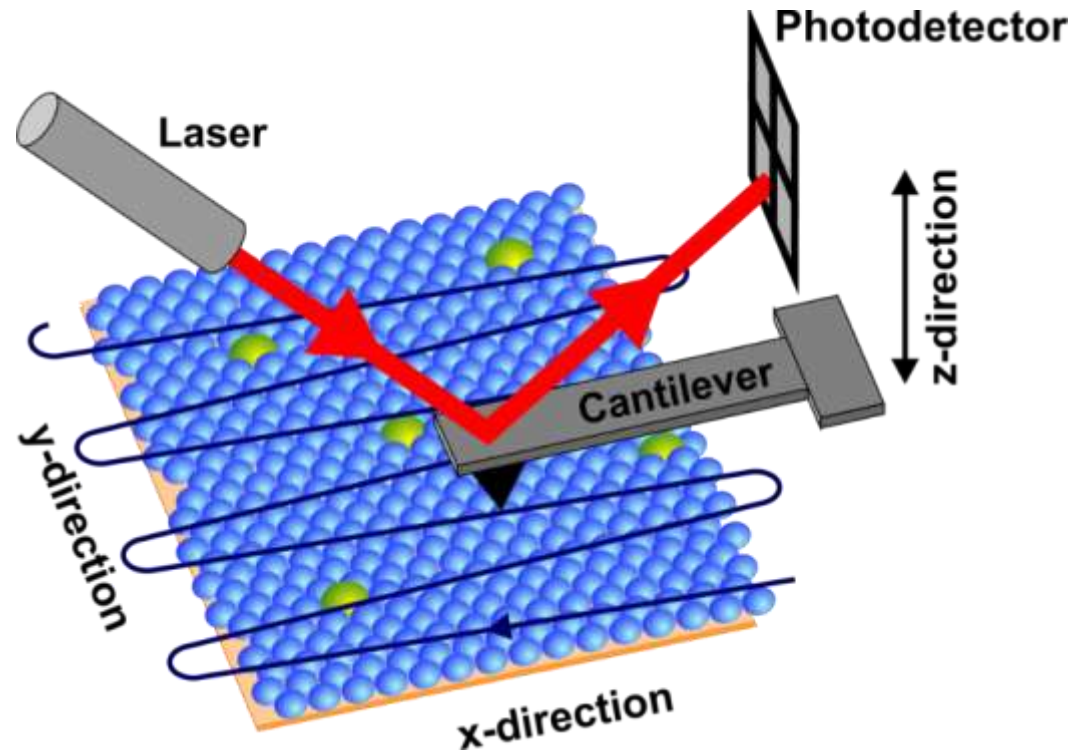
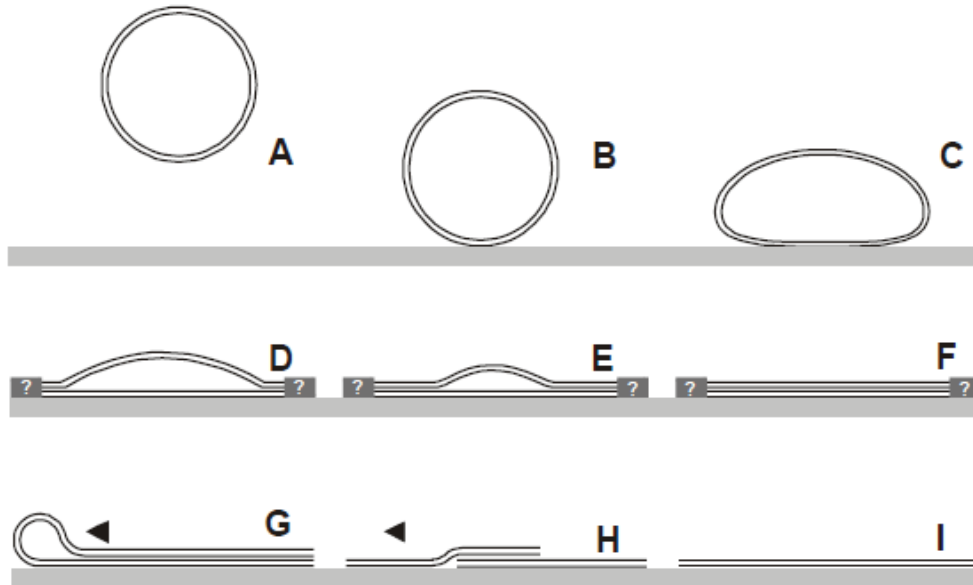
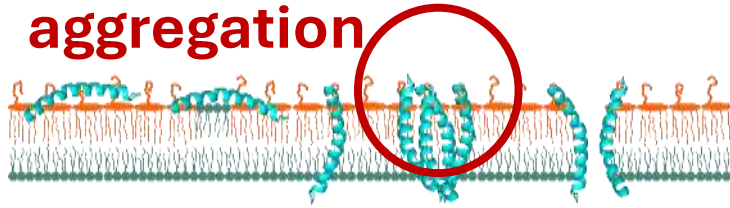


FRET spectroscopy on SUVs/LUVs

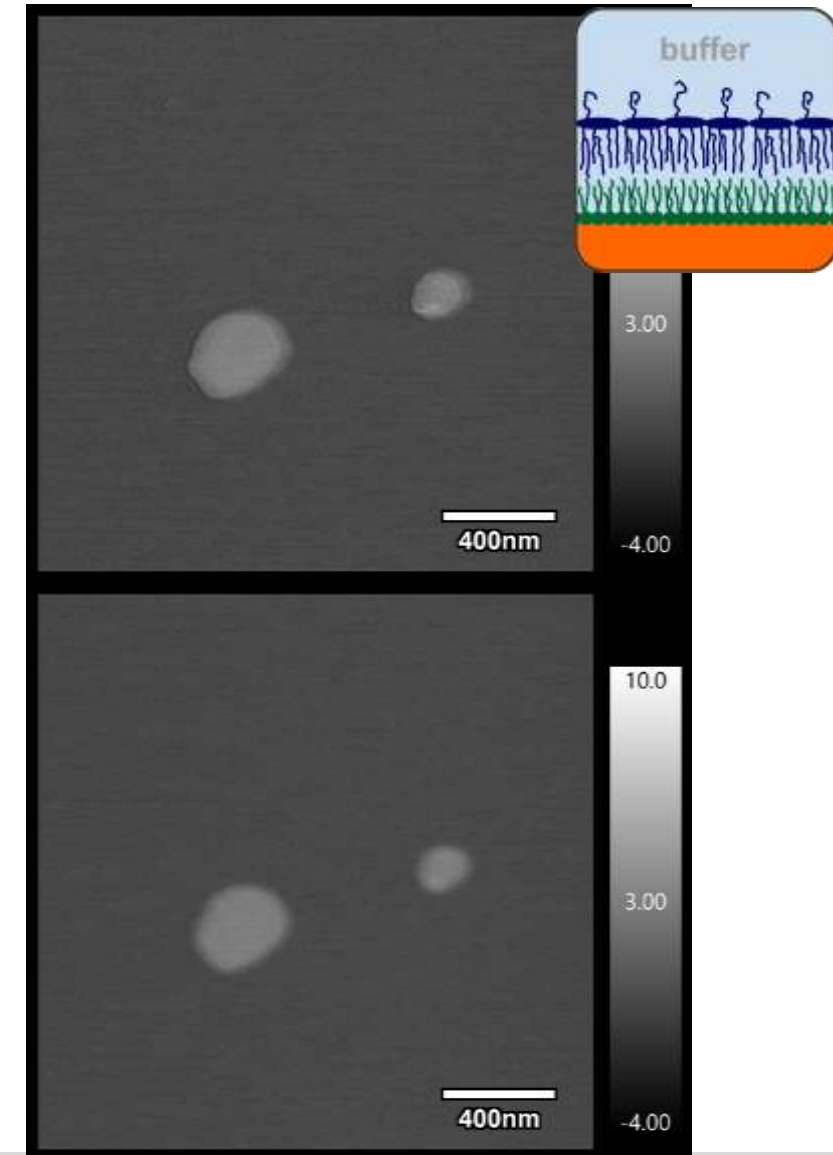
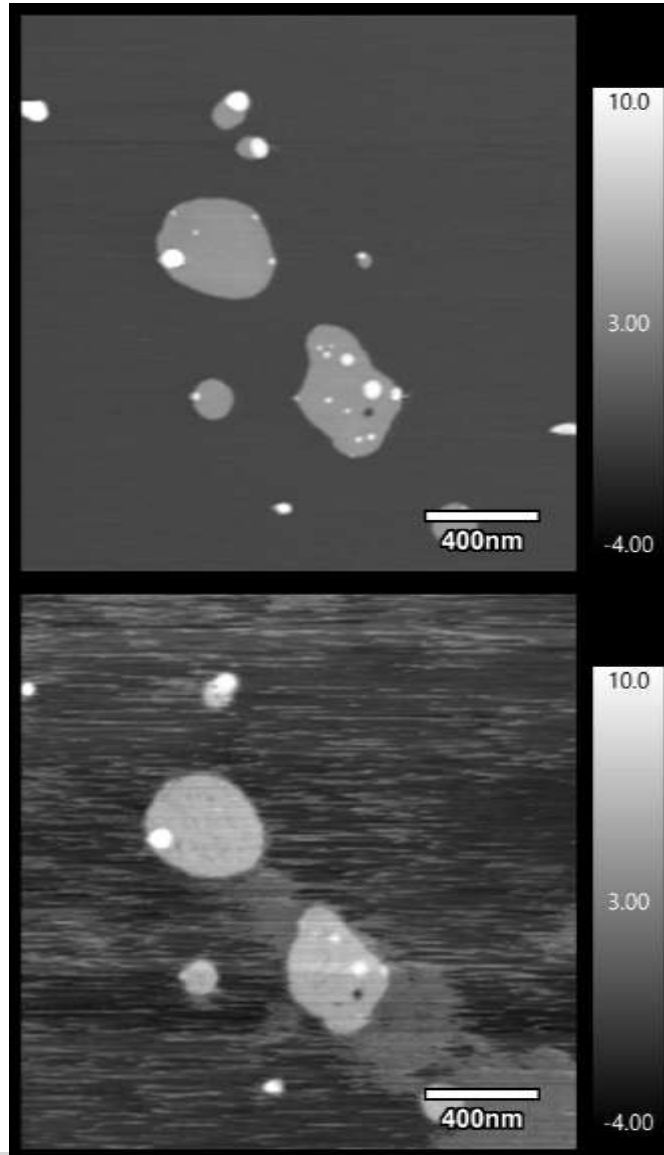
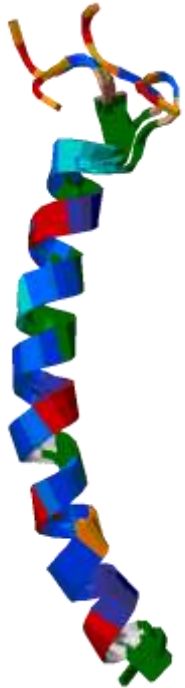


Partitioning
between bacterial
lipids depends on
VPRTES tail of
LL37

aggregation

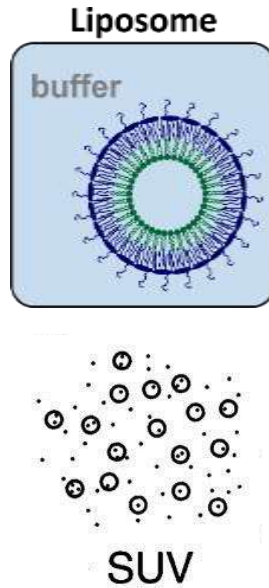
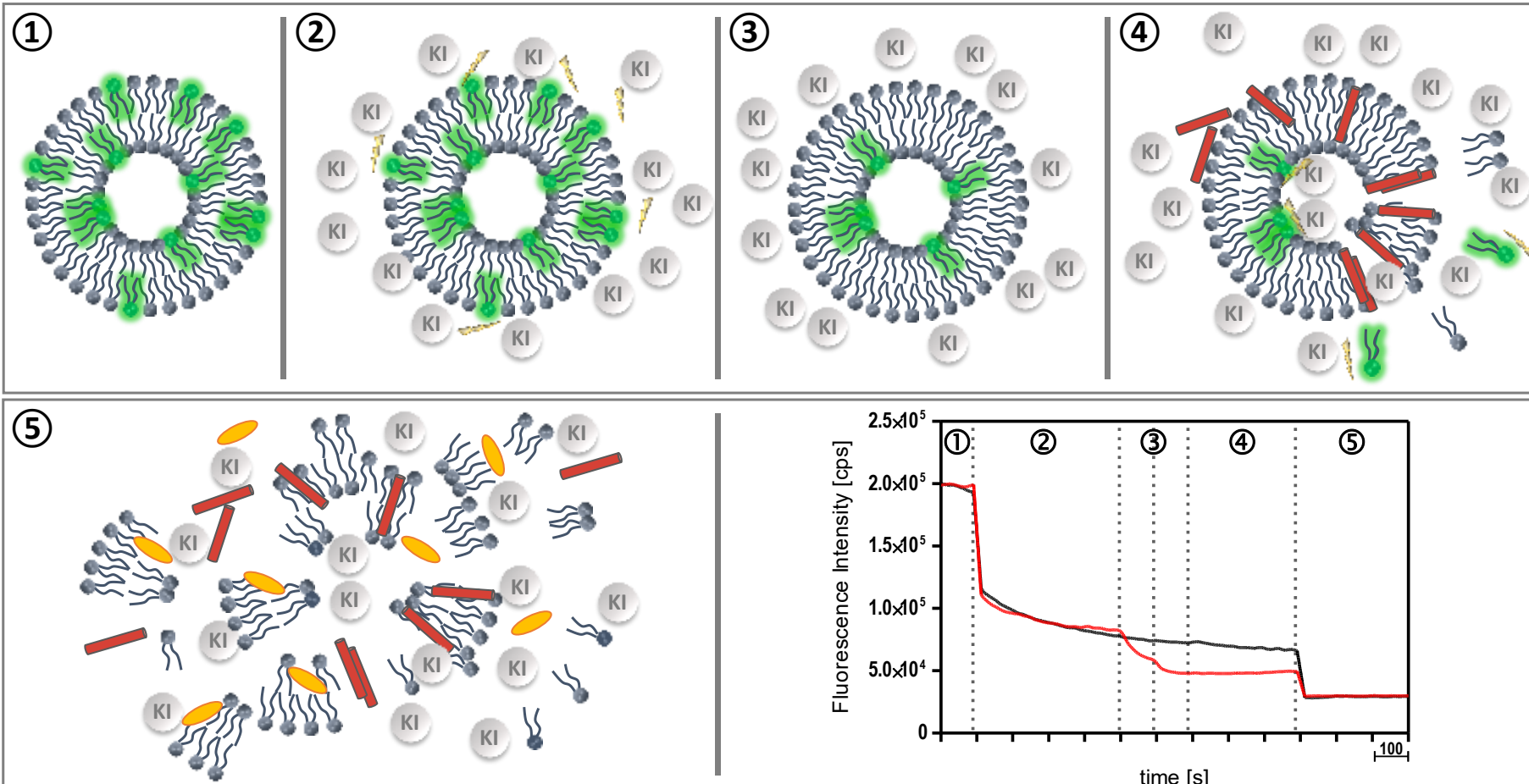
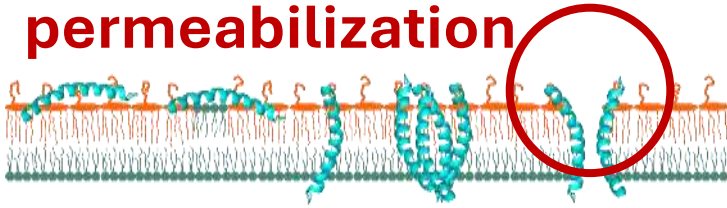


LL-32 and hBD3-l on PE+PG+TDM (1:1:2) bilayer



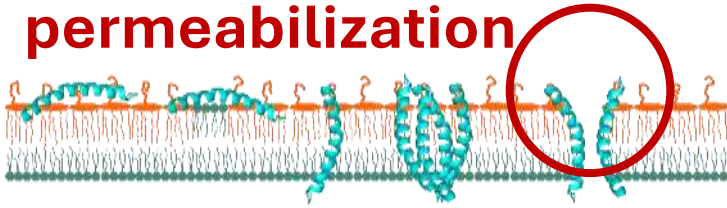
Potassium iodide fluorescence quenching assay

permeabilization

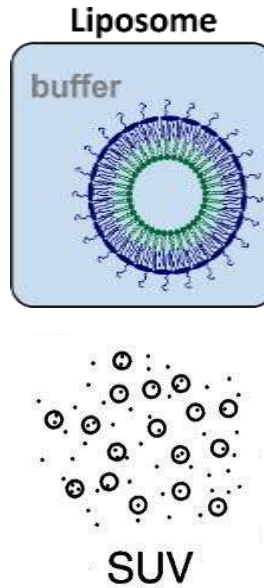
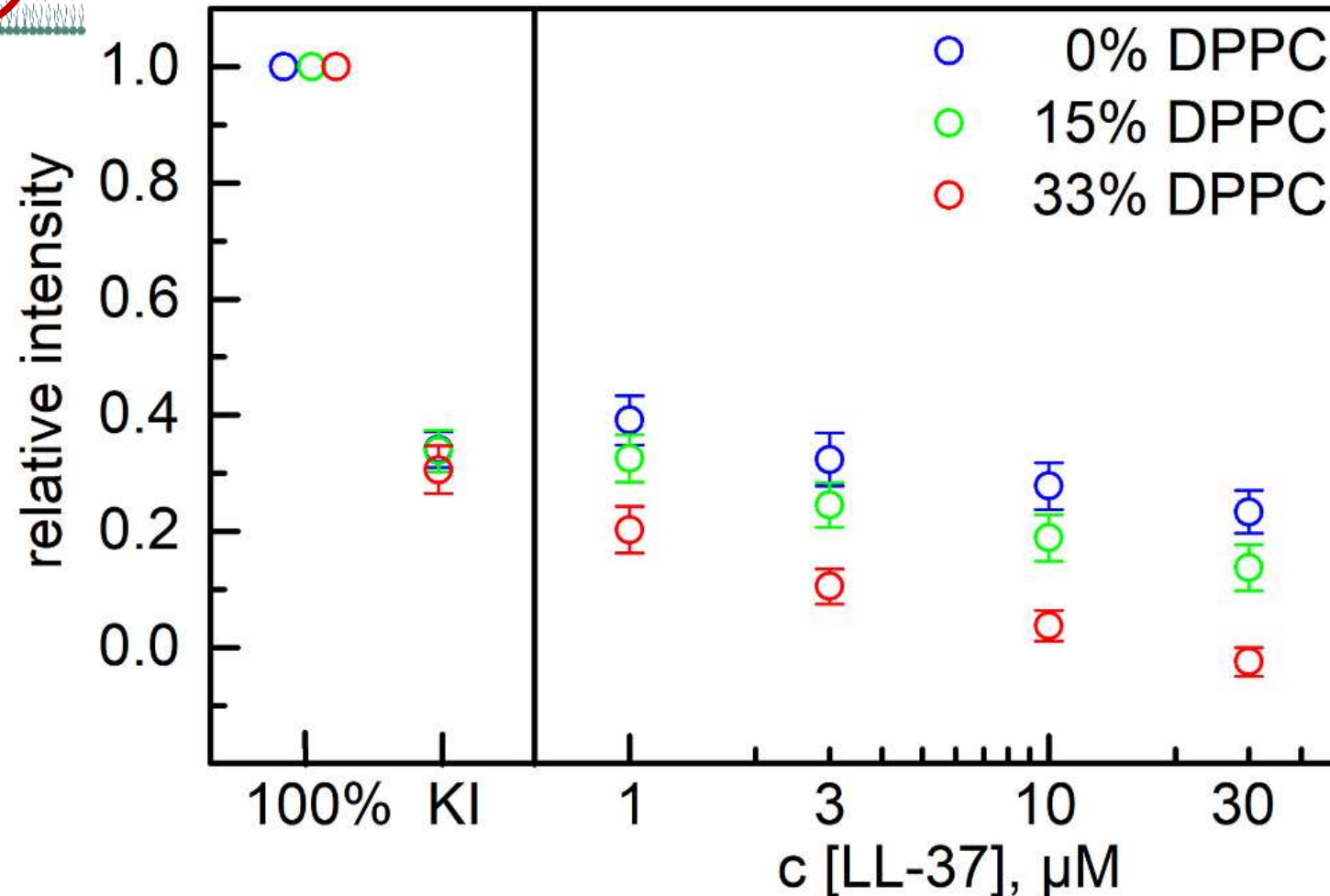


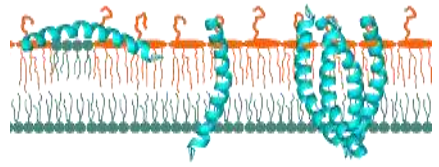
Decreasing the PC content as for resistant strain

permeabilization

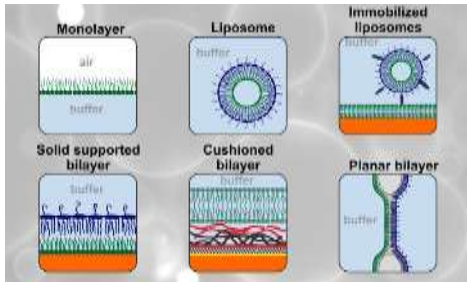


Resistance of
Legionella
micdadei towards
LL-37
depends on PC
content of
membrane

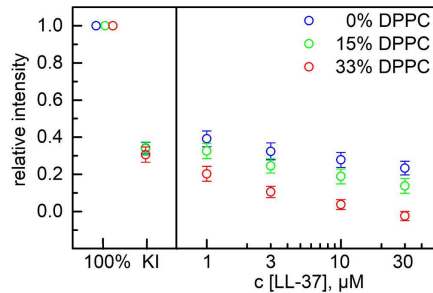




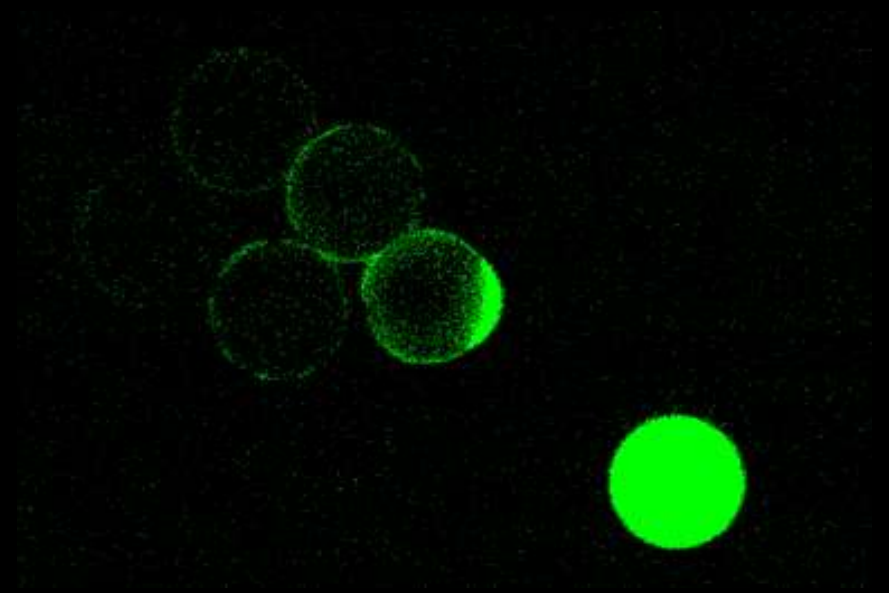
All critical membrane interaction steps can be analyzed using liposome techniques.



Only complementary comparison of different techniques and model systems provides full understanding.



Changes in lipid composition may dramatically modify the membrane interaction.



Microfluidic evaluation of AMPs



Phospholipid Research Center

Team Microfluidics

Janina Nandy
Research Center Borstel



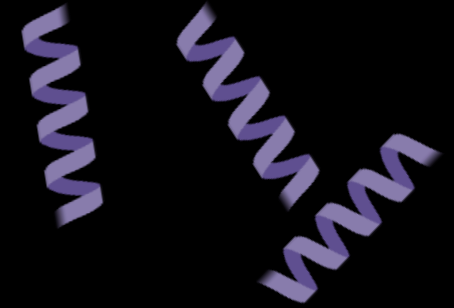
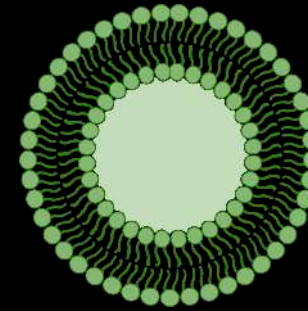
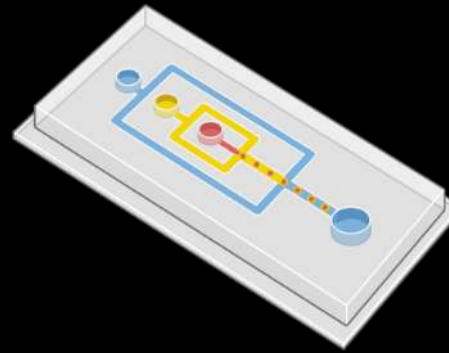
Rieke Bruns
Research Center Borstel



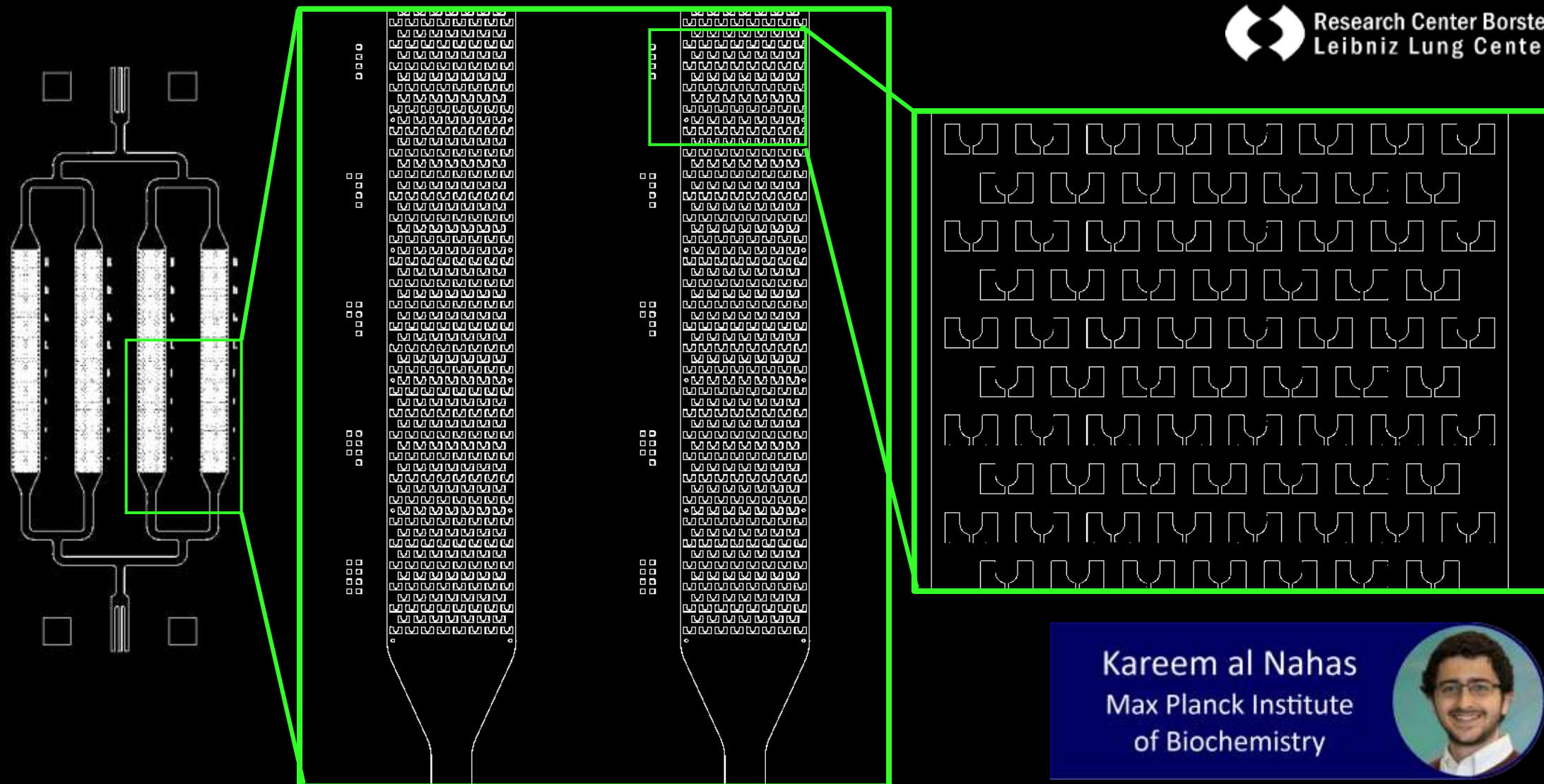
Marie Wetzel
Research Center Borstel



Franziska Kiesow
Research Center Borstel



**Standardized
prediction
of
antimicrobial
activity**



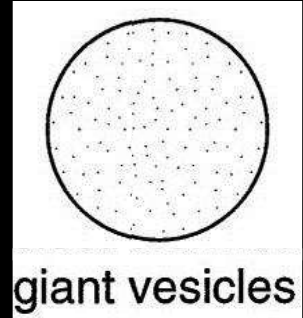
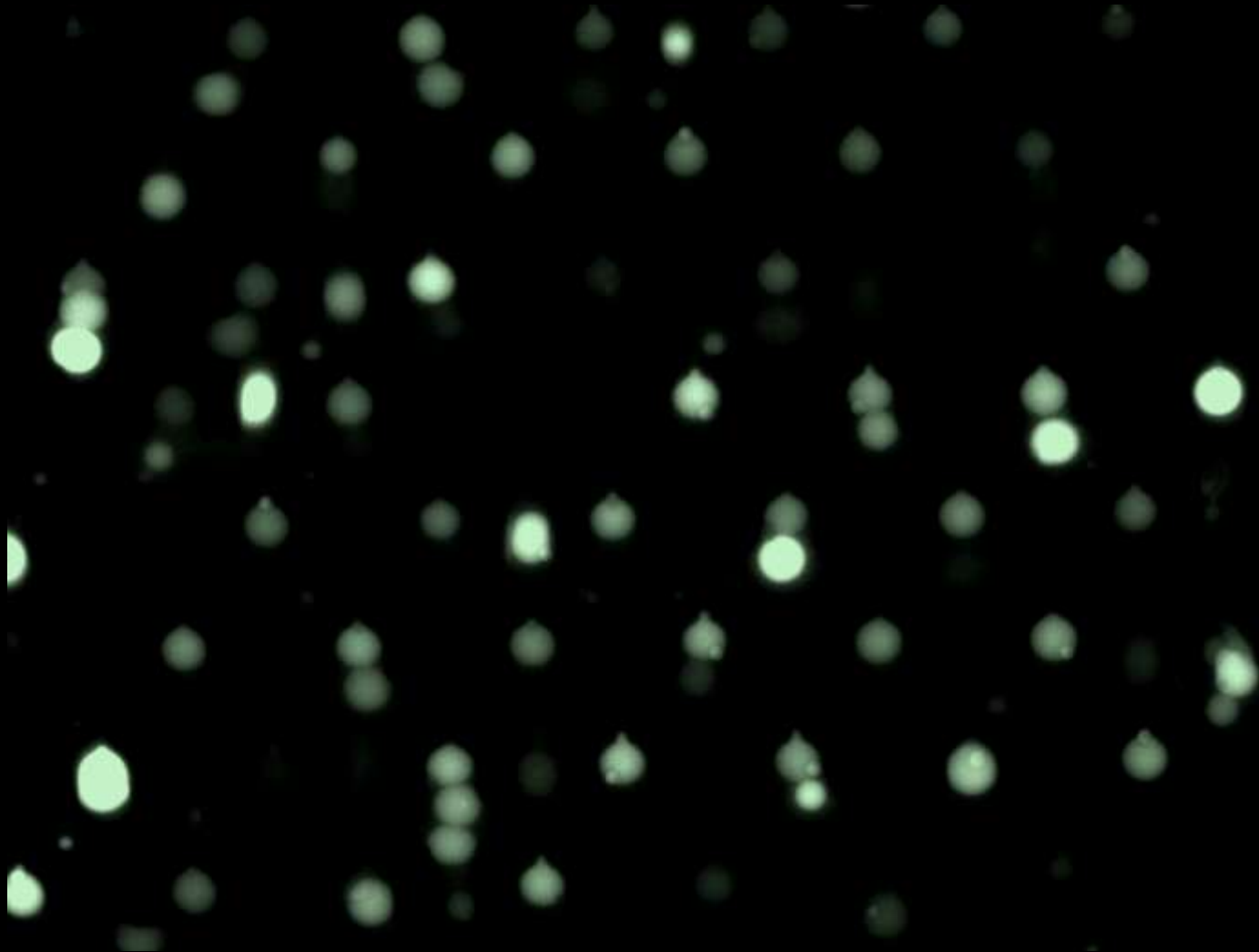
Kareem al Nahas
Max Planck Institute
of Biochemistry

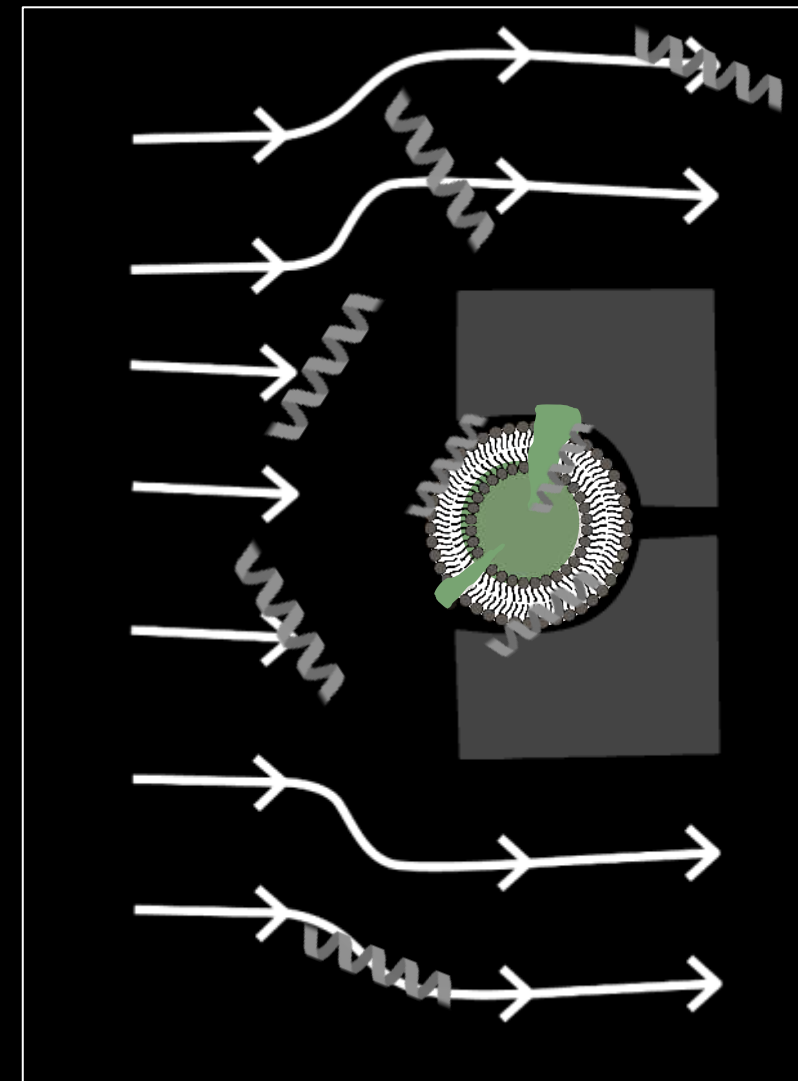
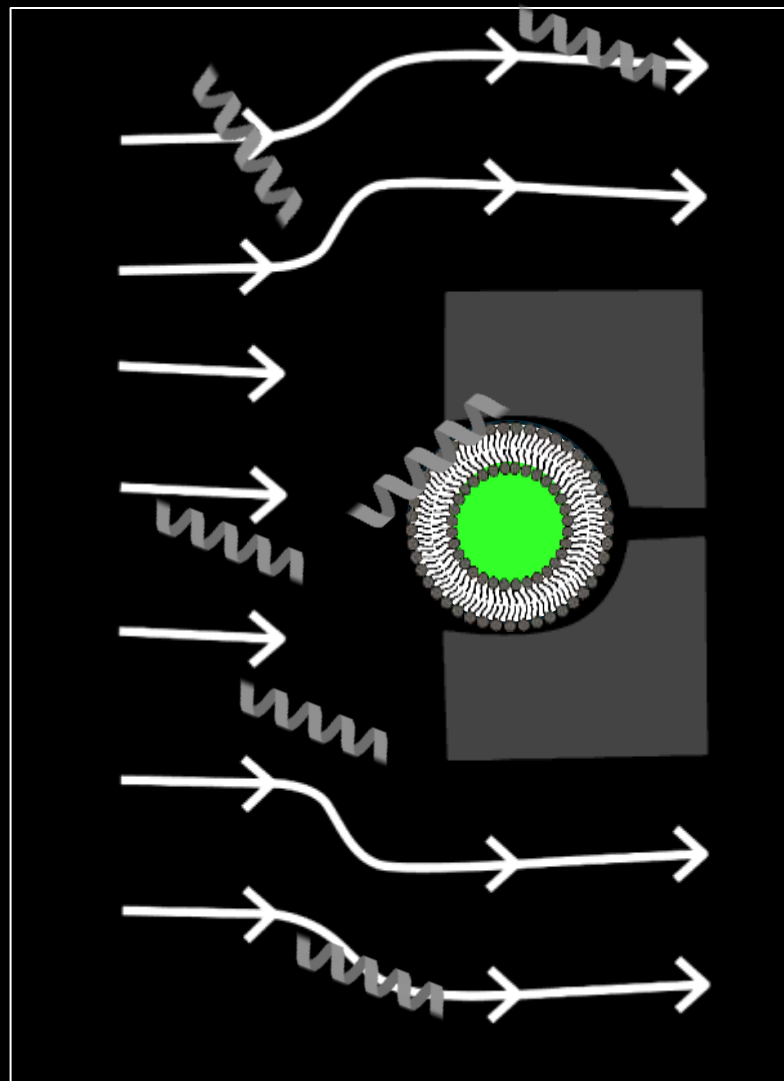
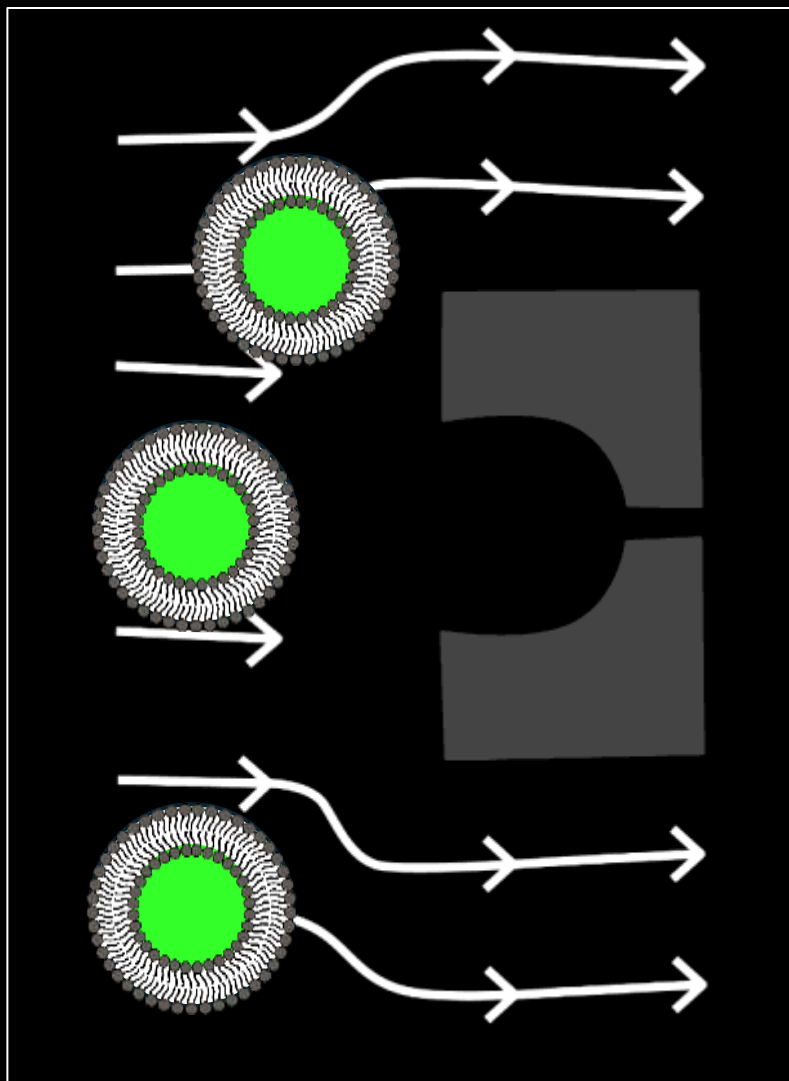


Marcus Fletcher
Imperial College London



Trapping hundreds of liposomes on chip

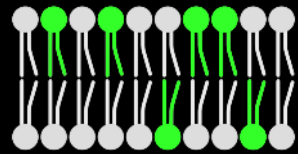




Time

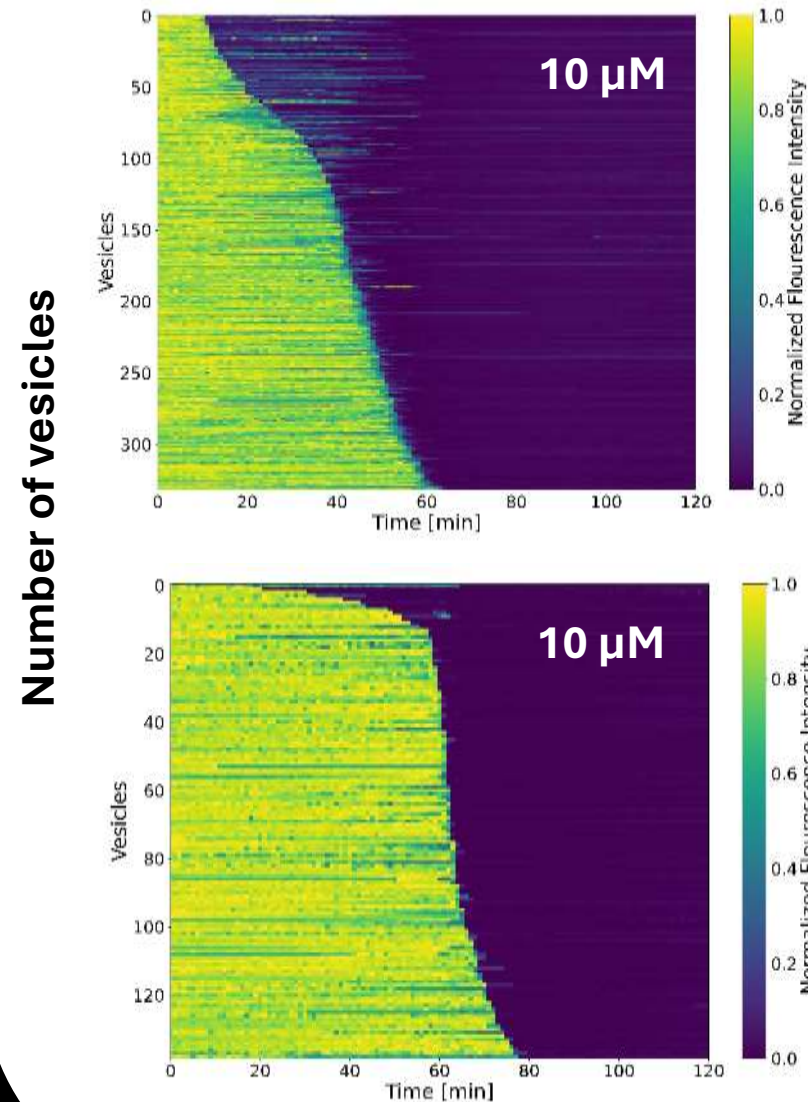
Proof of concept

Bacterial Model
DOPC:DOPG 1:1

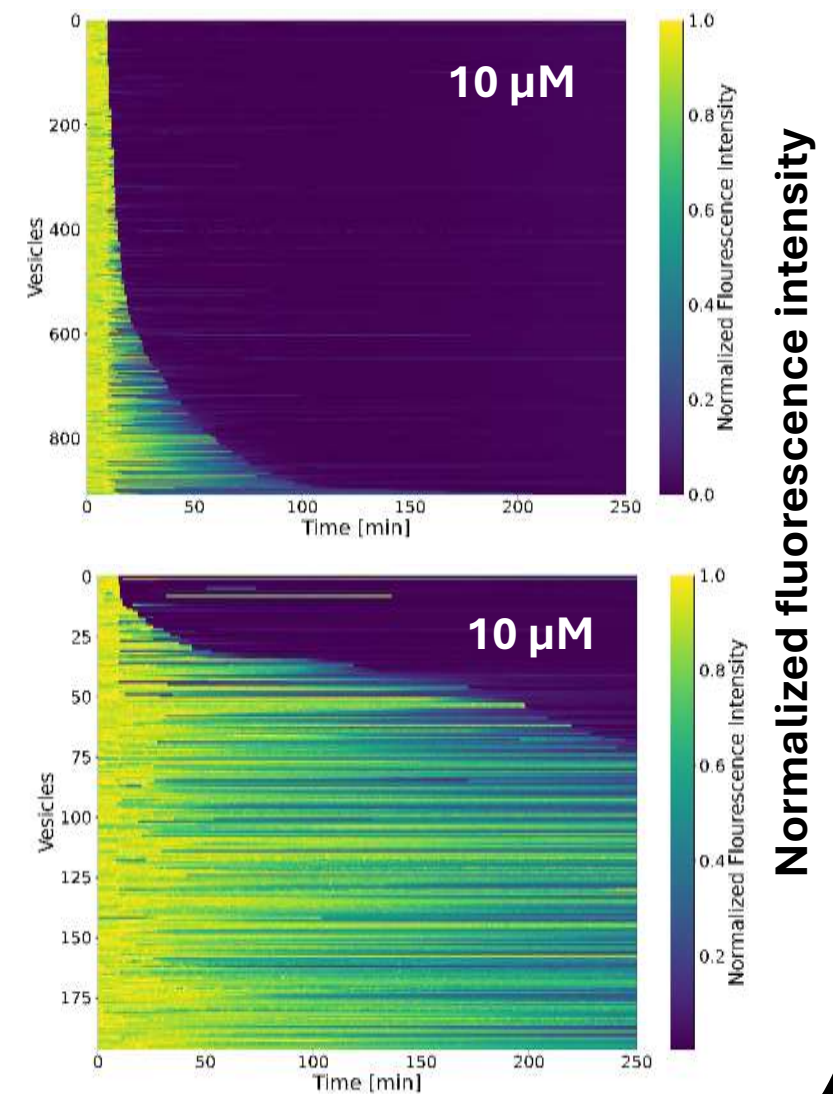


Dye release of
microfluidic trapped
liposomes provides
reasonable results

Melittin

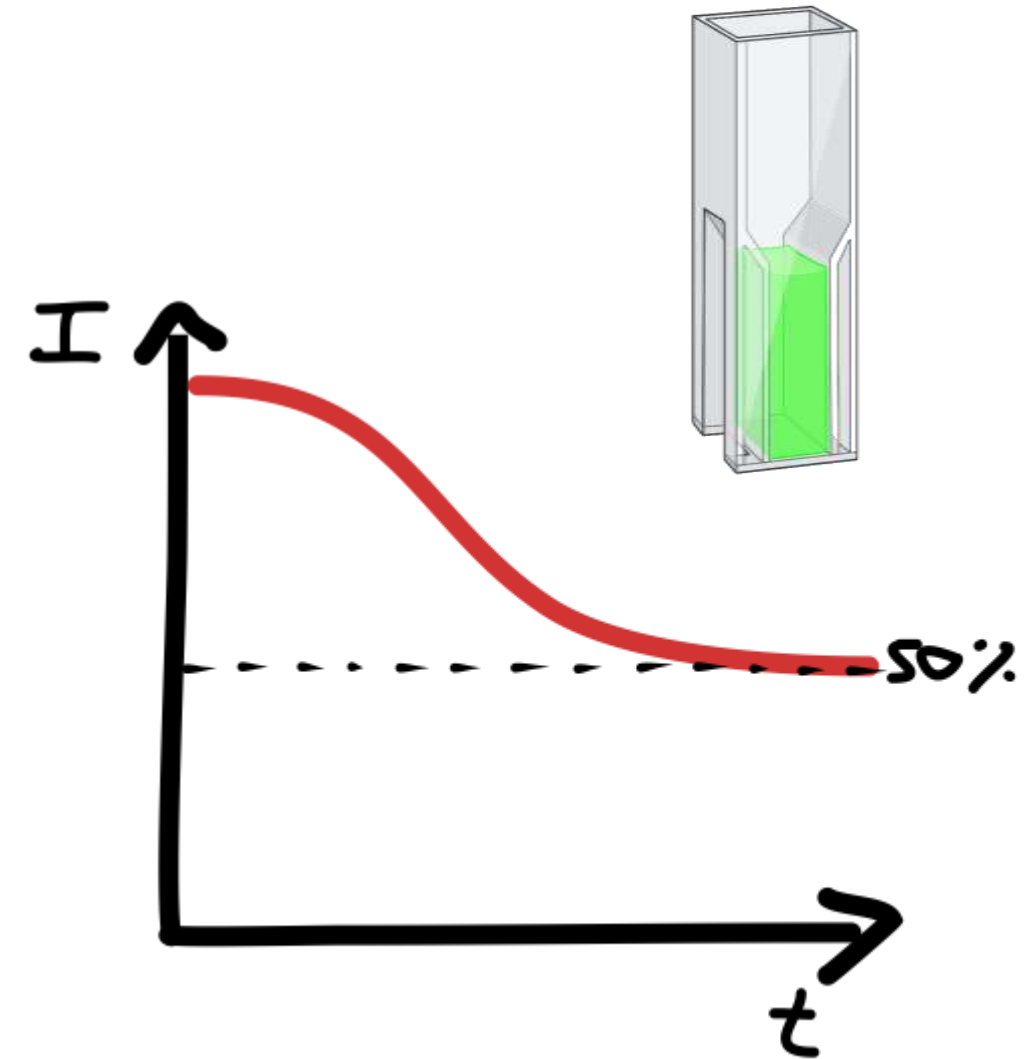


hBD3-l

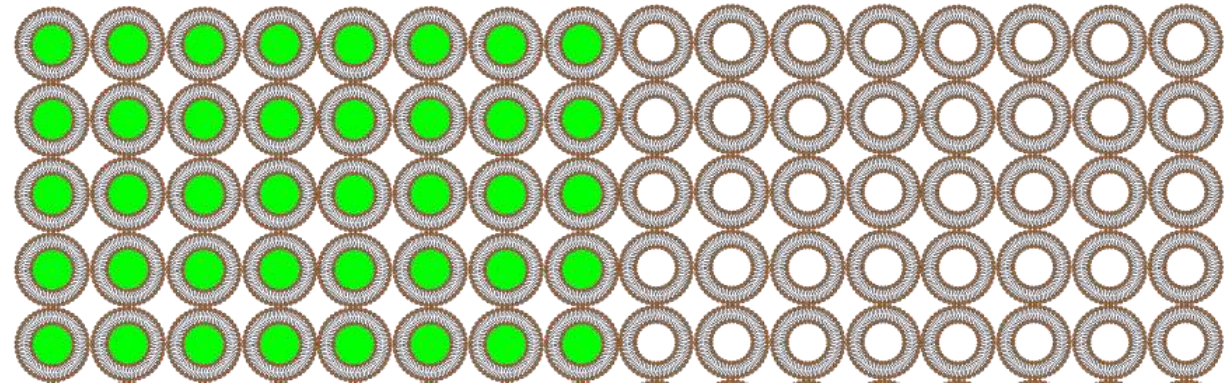


Time / min

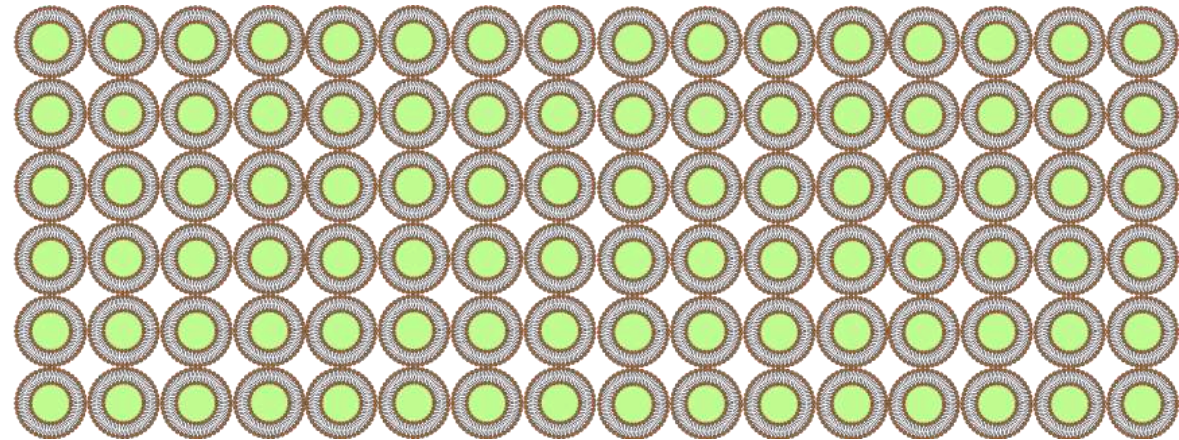
Our technique provides additional information...

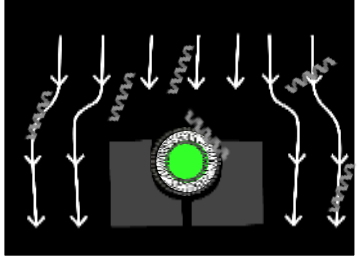


Half empty – half full **All-OR-NONE**

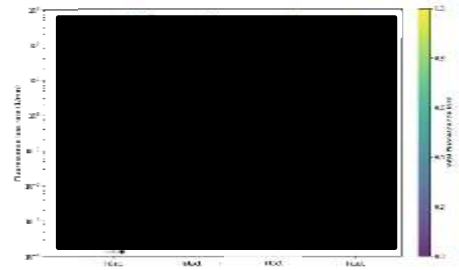


Partial release of dye **GRADED**

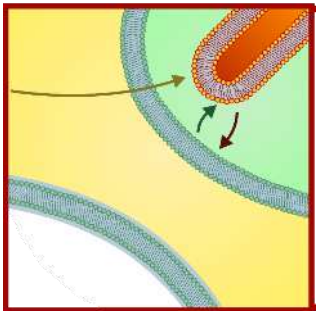




Microfluidic trapping bridges single liposome and bulk analysis.



Quantitative evaluation enables classification of mode of action continuum of membrane active compounds.



Membrane interaction is key to predict antimicrobial potential of AMPs (and other compounds).

Acknowledgement



Biophysics Group

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Other Groups

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Marcus Fletcher



Kareem Al Nahas



Naresh Yandrapalli



Ulrich Keyser



Marc Devocelle
John Connolly



Phospholipid Research Center



Federal Ministry
of Education
and Research

Contact



cnehls@fz-borstel.de

[https://fz-borstel.de/index.php
/christian-nehls](https://fz-borstel.de/index.php/christian-nehls)



Christian Nehls



@cnehls.bsky.social

International Membrane Biophysics Meeting Membranes as Biological Barriers

Drübeck Monastery | March 23th to 26th 2026

Confirmed Invited Speakers

Elina Ikonen, Helsinki, FI
Erdinc Sezgin, Karolinska, SE
Katia Cosentino, Modena, IT
Matti Javanainen, Tampere, FI
Luca Monticelli, Lyon, FR

Jesus Perez-Gil, Madrid, ES
Gerard Wong, Los Angeles, USA
Radek Šachl, Prague, CZ
Cecilia Clementi, Berlin, DE
Anna Taubenberger, Dresden, DE

James Saenz, Dresden, DE
Alf Honigmann, Dresden, DE
Frauke Gräter, Mainz, DE
Hans-Joachim Galla, Münster, DE

Registration

Early Bird Deadline: December 16, 2025
Registration and abstract submission will be possible:
November 03, 2023 - January 16, 2025

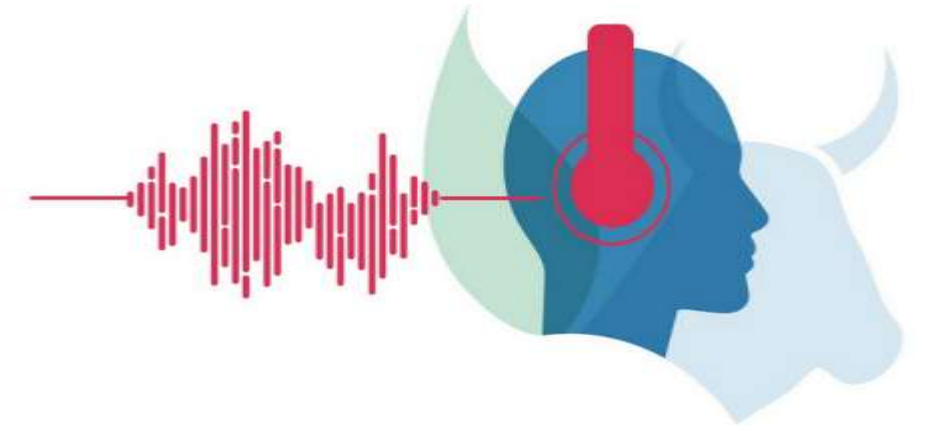


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Scientific organizers:

Christian Nehls, Borstel
Fabio Lolicato, Heidelberg
Ilpo Vattulainen, Helsinki



MIKROQBEN IM VISIER

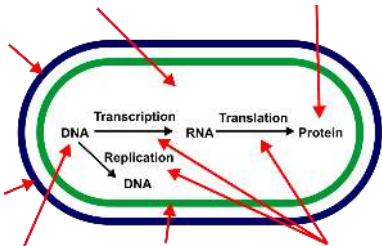
INFEKTIONEN VERSTEHEN, RESISTENZEN BESIEGEN



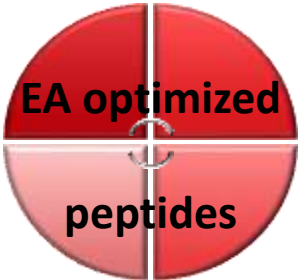
AMR



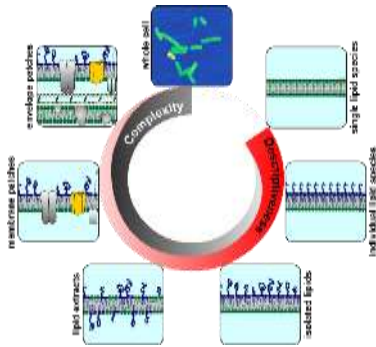
AMPs



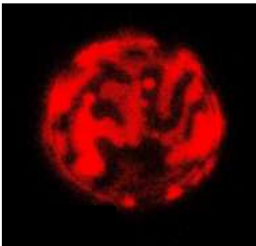
AMP optimization



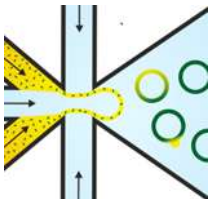
Model complexity



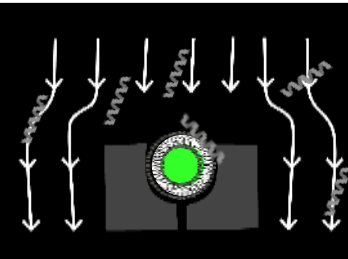
Asymmetriy



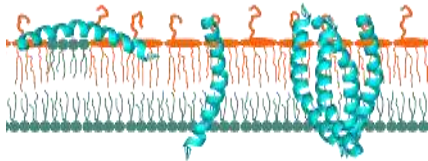
Microfluidics



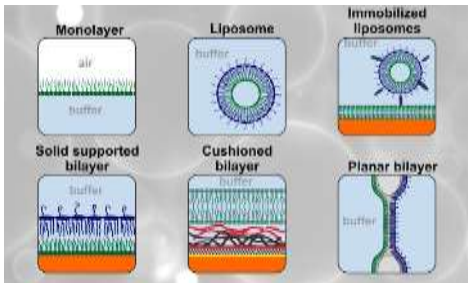
Microfluidic trapping



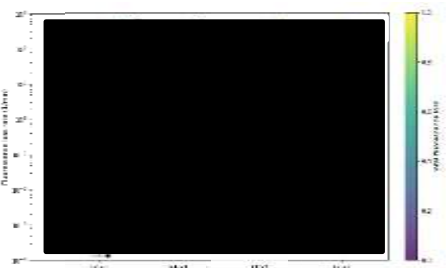
Membrane interaction



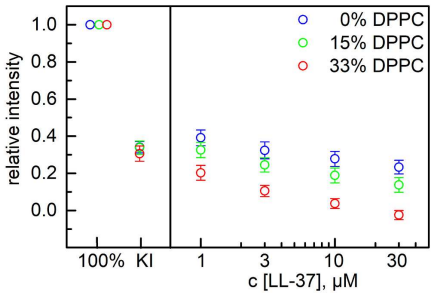
Membrane geometry



Graded - quantification



Resistance determinants



Membrane interaction

