



Società Chimica Italiana
Divisione di Didattica
Chimica

VI SCUOLA NAZIONALE DI DIDATTICA DELLA CHIMICA
"GIUSEPPE DEL RE"
XIII SCUOLA DI RICERCA EDUCATIVA E DIDATTICA CHIMICA
"ULDERICO SEGRE"
17 – 30 novembre 2021



Per le diverse scale

Gerarchia e dinamica dei processi di self-assembly

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The Bottom-up Approach

Self-assembly

Definition: The spontaneous and reversible association of molecular species to form larger, more complex supramolecular entities according to the intrinsic information contained in the components.

Equilibrium methods

The final structure must be:

- reasonably stable at room temperature
- weakly enough bound so that the system can explore the large number of configurations needed to find the desired configuration of lowest free energy.

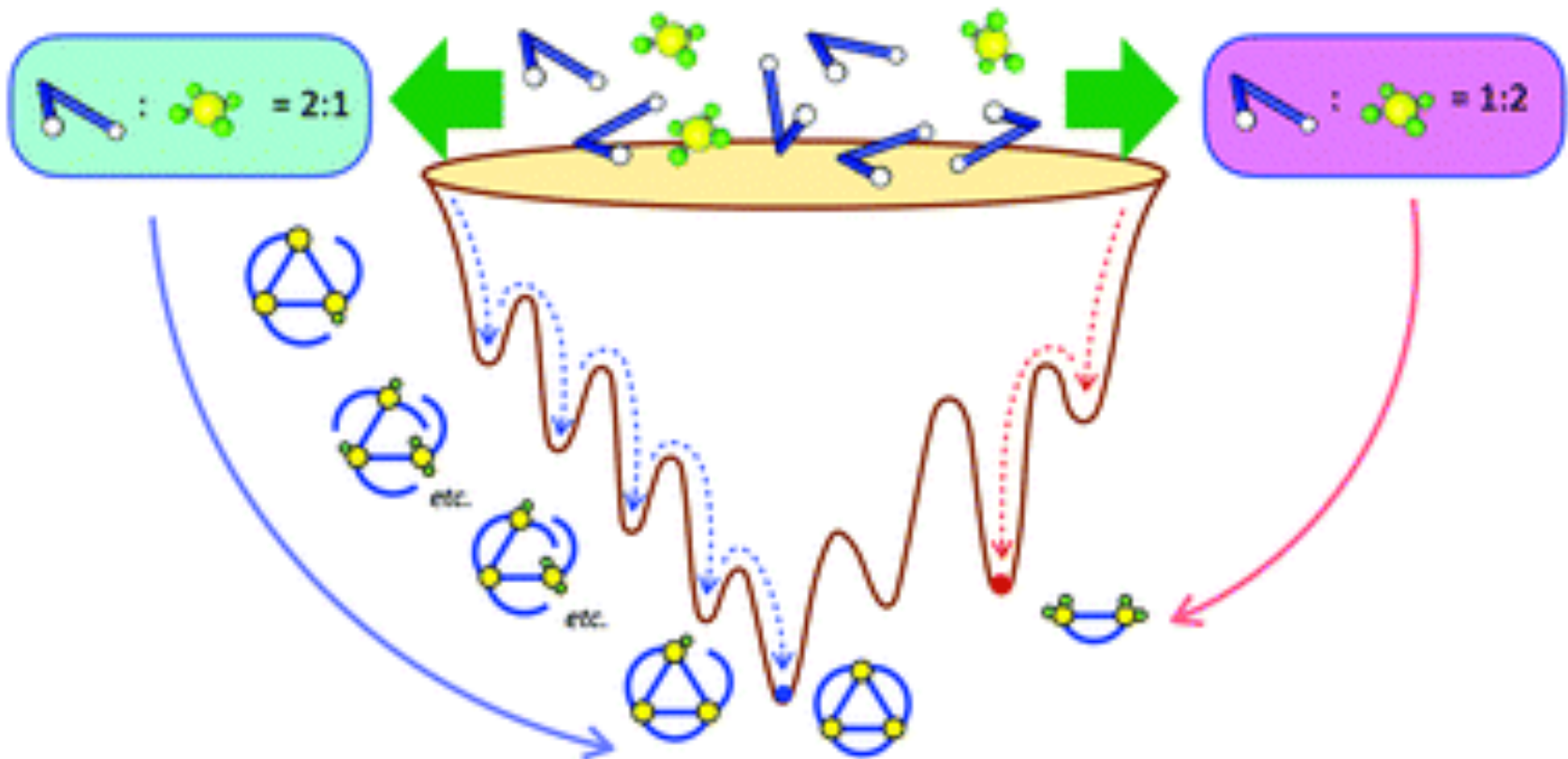
The most stable structure in a given environment originates from a delicate balance between entropy (at a given temperature) and binding enthalpies.

- **Self-healing:** Due to the reversible nature of the assembly process, self-assembling systems are able to correct ‘mistakes’ during assembly and gradually work their way towards the most thermodynamically stable product.
- The entropic penalty is somewhat offset by the release of solvent molecules that were previously interacting with the binding areas of the assembly components.

Entropy driven aggregation!

Kinetic trapping

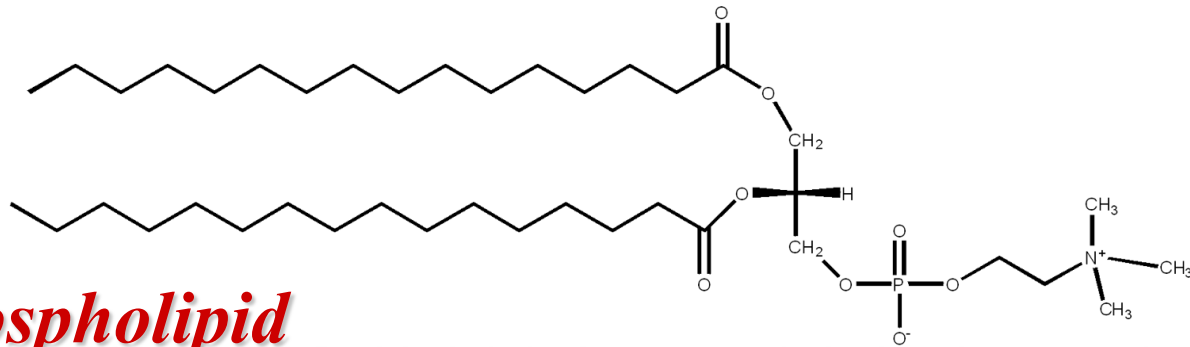
Trapping the system in some non-equilibrium configuration.



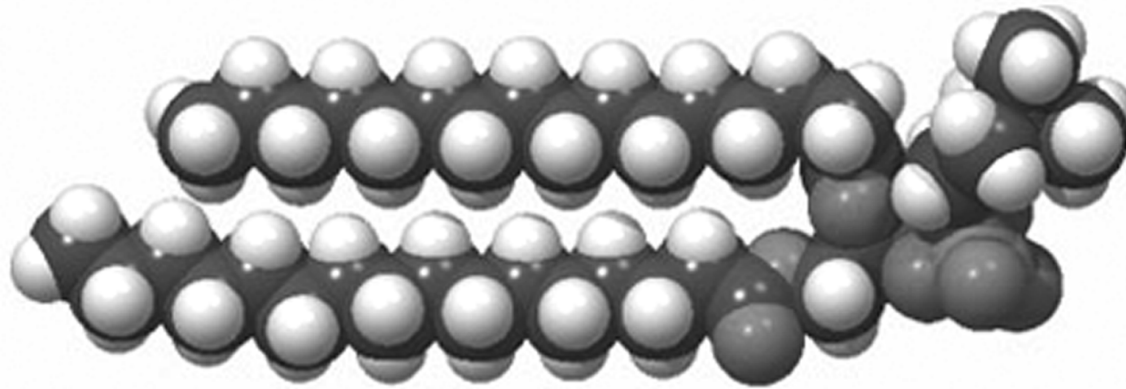
Self-assembly driving force

- **electrostatic, Van der Waals, HB interactions**
- **π - π interaction**
- **comparative solvation of monomers and aggregates**
- **Aggregation is an interfacial phenomenon**
 - **size and molecular shape**

Amphiphiles



Phospholipid

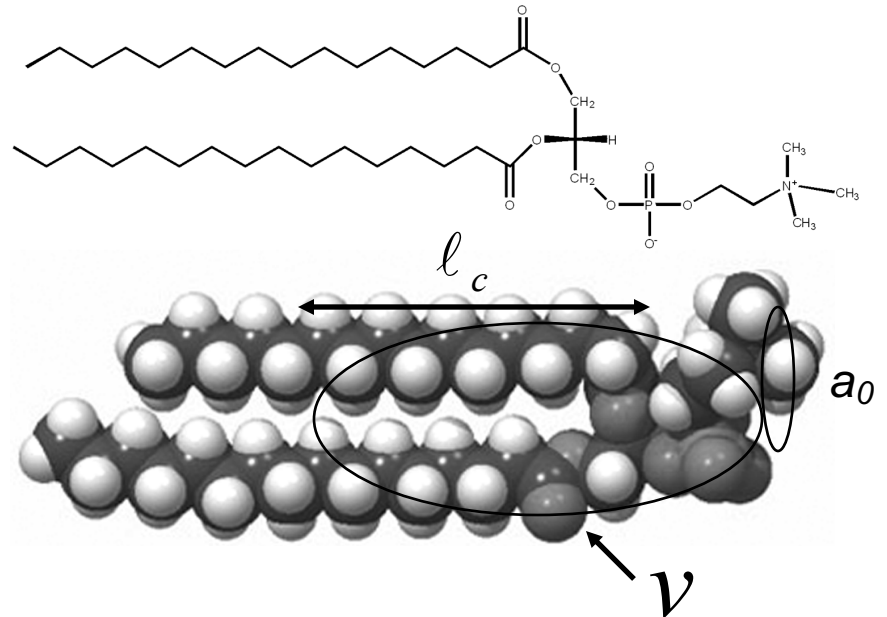
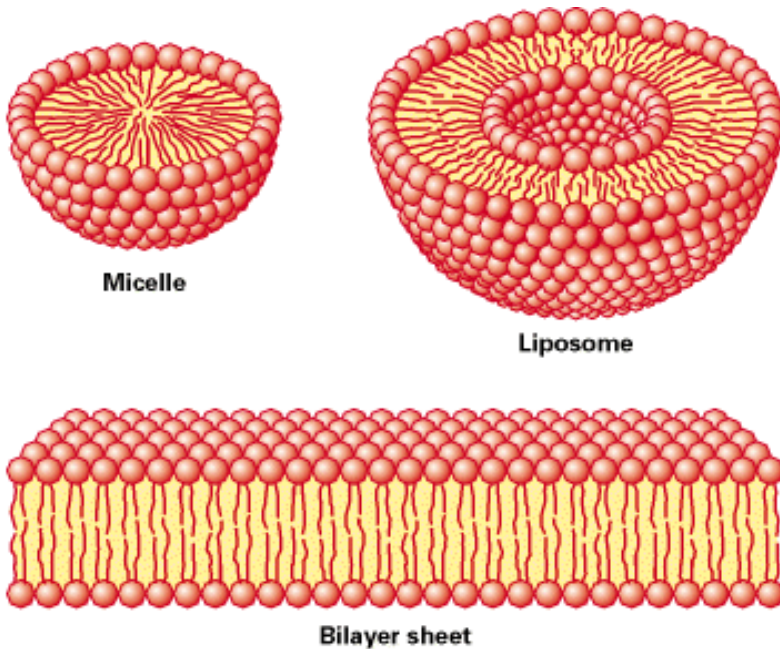


Polar head

Hydrophobic tail

Shape of aggregates

Packing effects depend on geometry



l_c = critical length of the hydrocarbon chain (maximum effective length)

v = volume occupied by the hydrocarbon chain

a_0 = optimal area of the head group

Packing factor

$$\frac{v}{a_0 l_c}$$

A dimensionless shape factor determining the aggregate geometry.

$$\frac{v}{a_0 l_c} < \frac{1}{3}$$

Spherical micelles

$$\frac{1}{3} < \frac{v}{a_0 l_c} < \frac{1}{2}$$

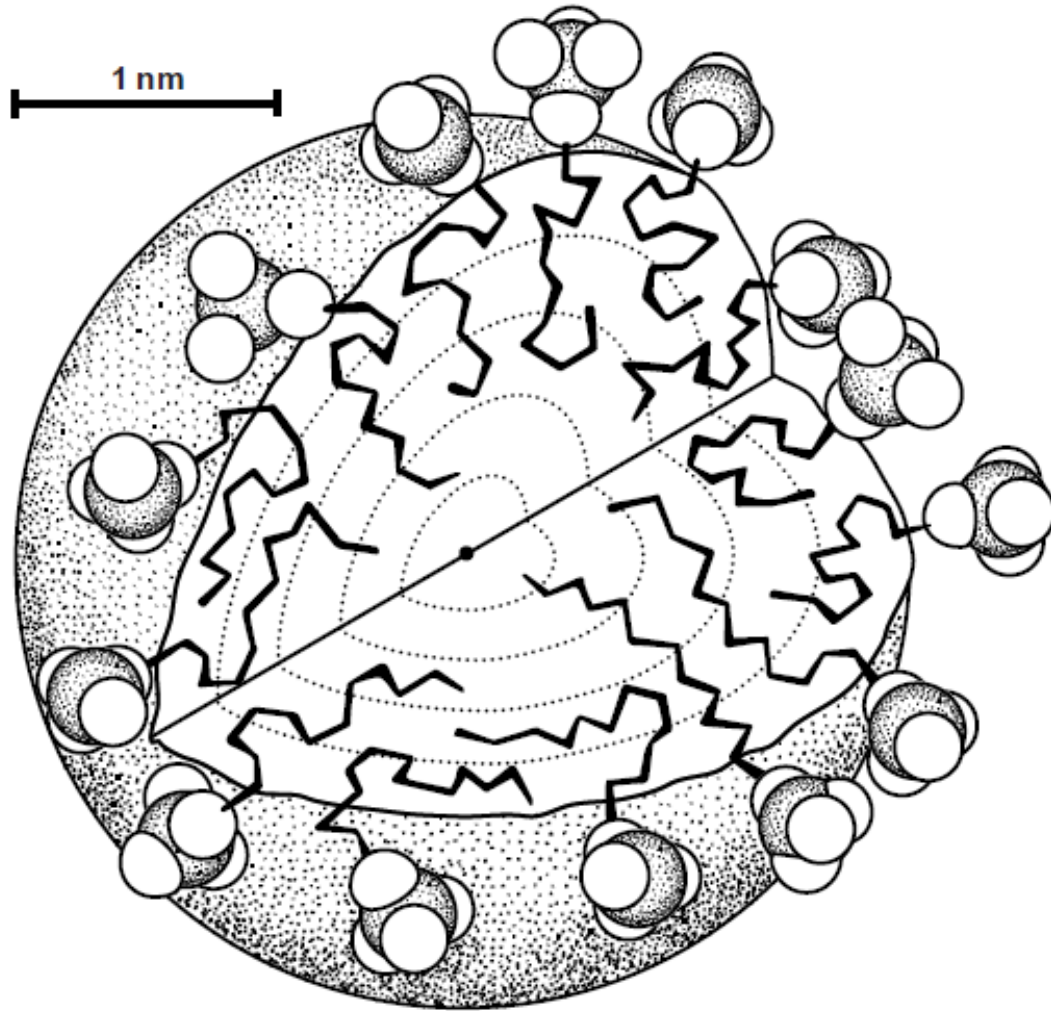
Non-spherical micelles

$$\frac{1}{2} < \frac{v}{a_0 l_c} < 1$$

Vesicles or bilayers

$$\frac{v}{a_0 l_c} > 1$$

Inverted cones

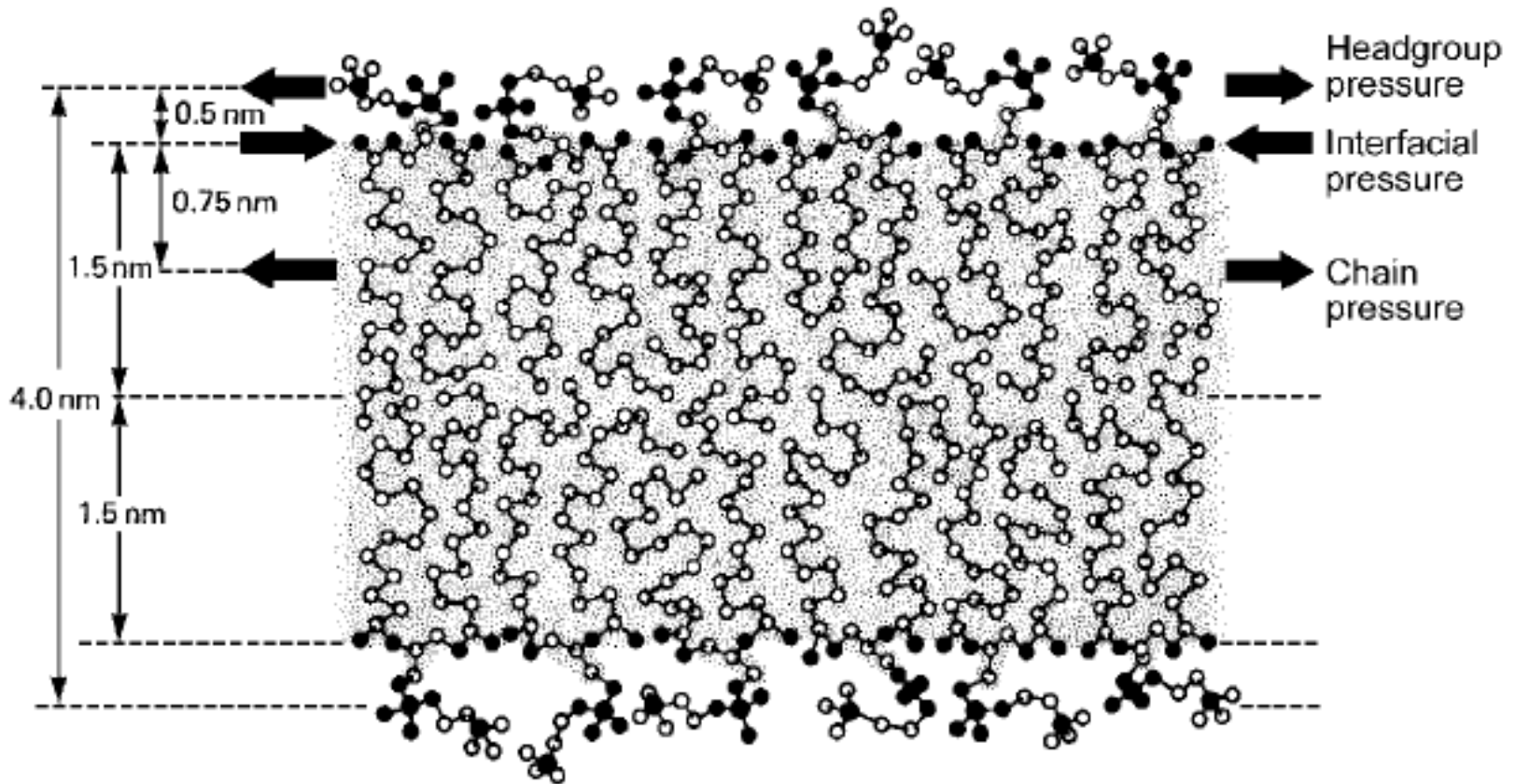


SDS micelles ($N=60$).

Note that all segments of the chain spend an appreciable proportion of time near the micelle surface.

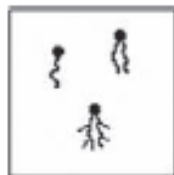
Bilayers

$$\frac{v}{a_0 l_c} \approx 1$$

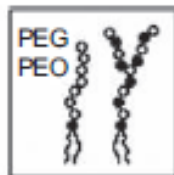


Phosphatidyl choline (PC)

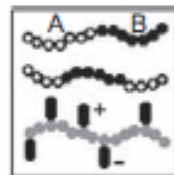
MESOPHASES



Single, double and branched chains



PEGolated and peptide amphiphiles



Di- and tri-blocks, poly-ampholytes



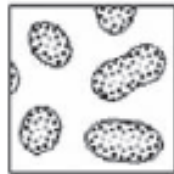
Micelle (spherical)



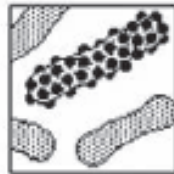
Micelle (spherical)



Single molecule micelle



Micelle (prolate)



Micelle (cylindrical or rod-shaped)



Micelle (threadlike)



Ordered 'hexagonal' cylinders



Bicontinuous 3-armed junctions



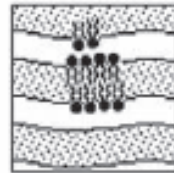
Overlapping bicontinuous 'phases'



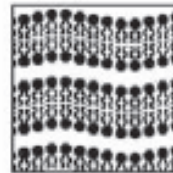
Bicontinuous 4-armed junctions



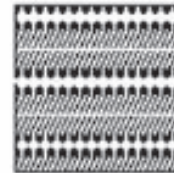
Bicontinuous 6-armed junction



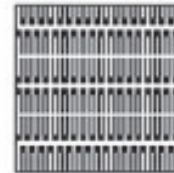
Lamellar bilayers (L_{α} phase)



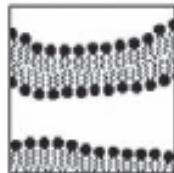
Lamellar (rippled) phase



Lamellar bilayers (L_{β} phase)



Lamellar bilayers (L_{γ} phase)



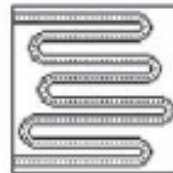
Isolated fluid state bilayers



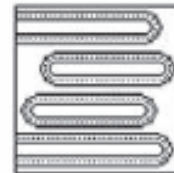
Vesicle



Bilayer-rod transition



Folded single bilayer



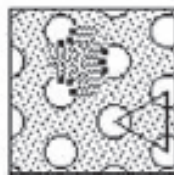
Platelets



Inverted micelle in oil



Inverted branched rod micelles



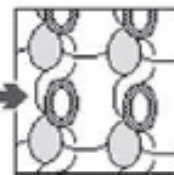
Inverted hexagonal phase



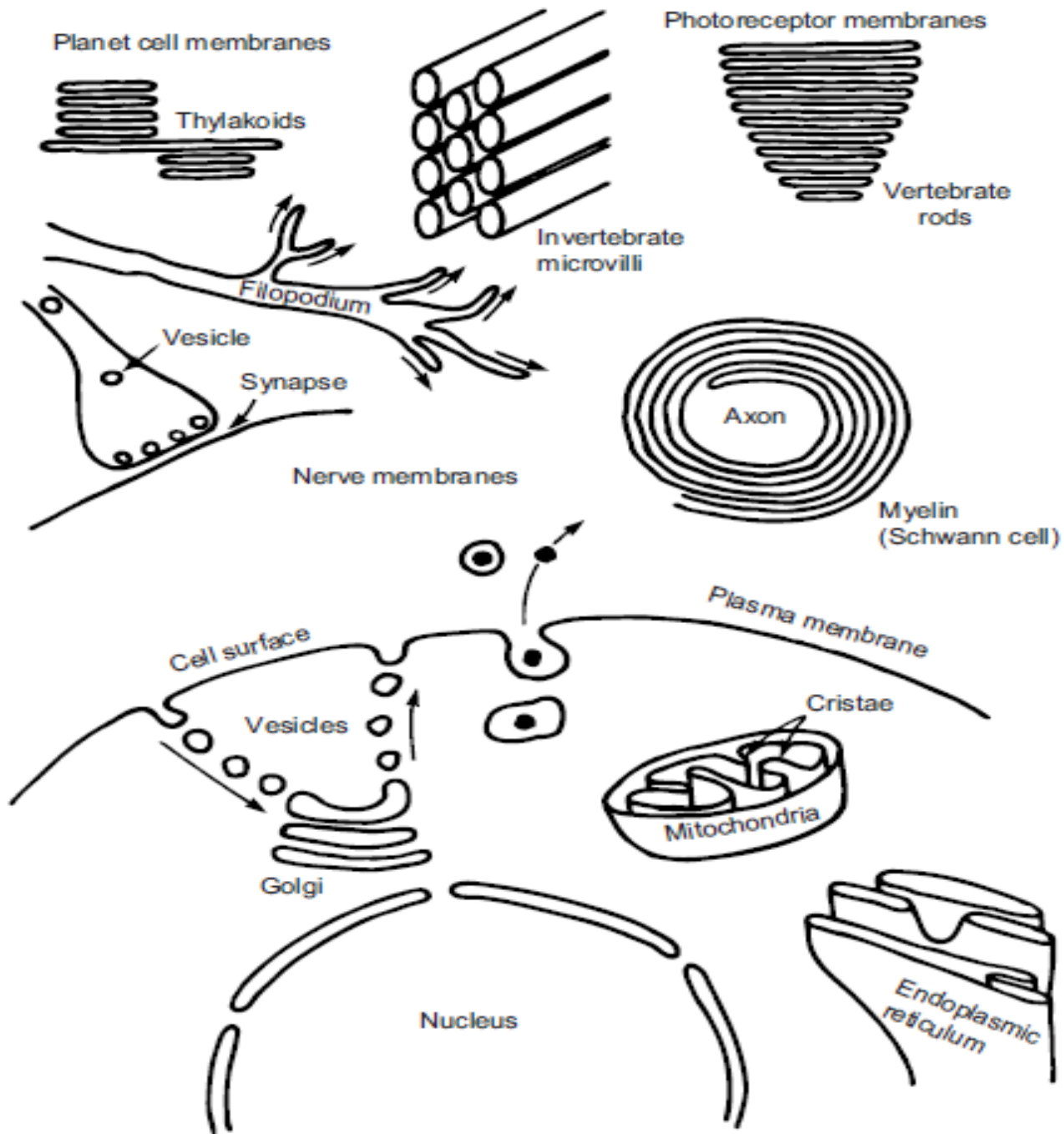
Tricontinuous 4-armed junction



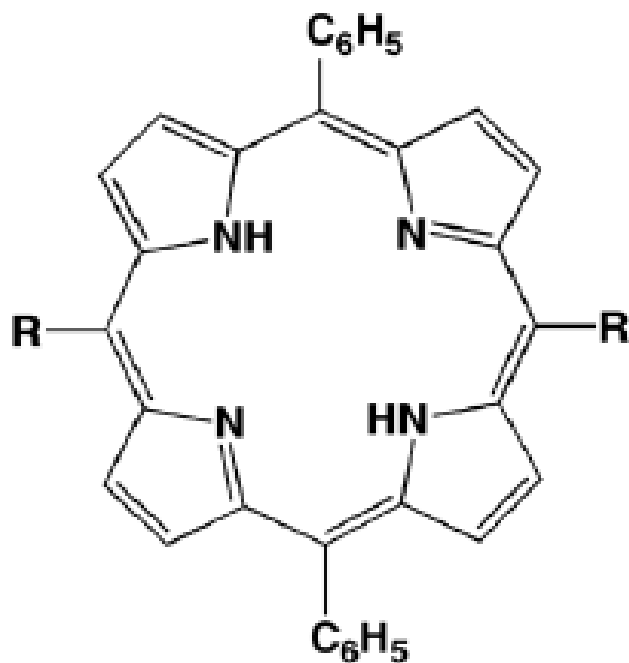
Tricontinuous 6-armed junctions separating cell interiors from exteriors



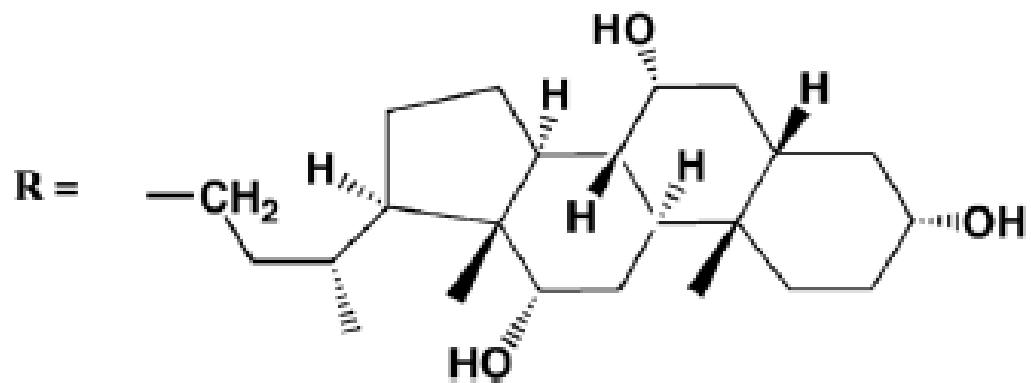
Tricontinuous tubules (microvilli)



Chiral self-assembly



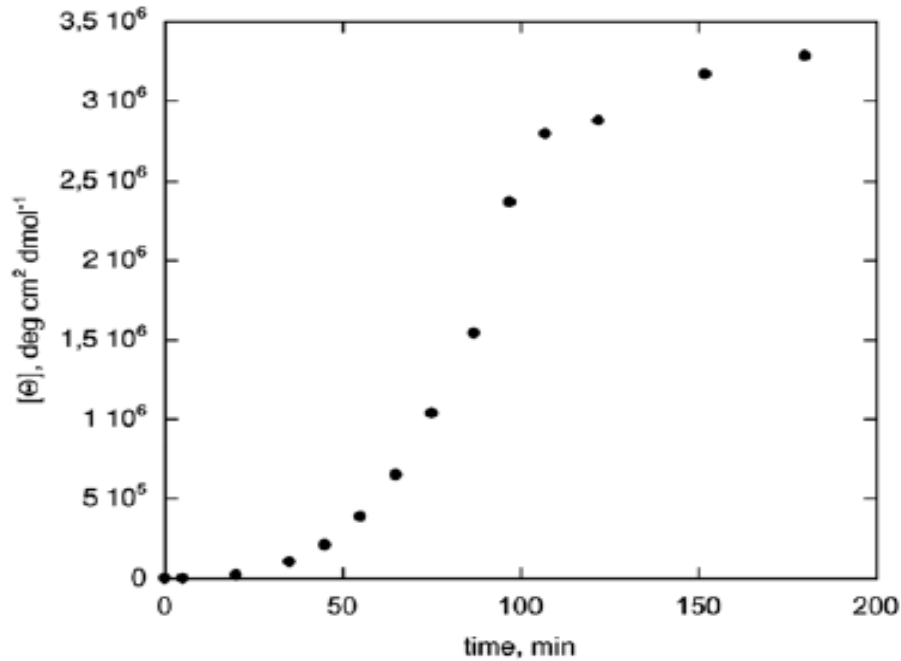
Donato Monti



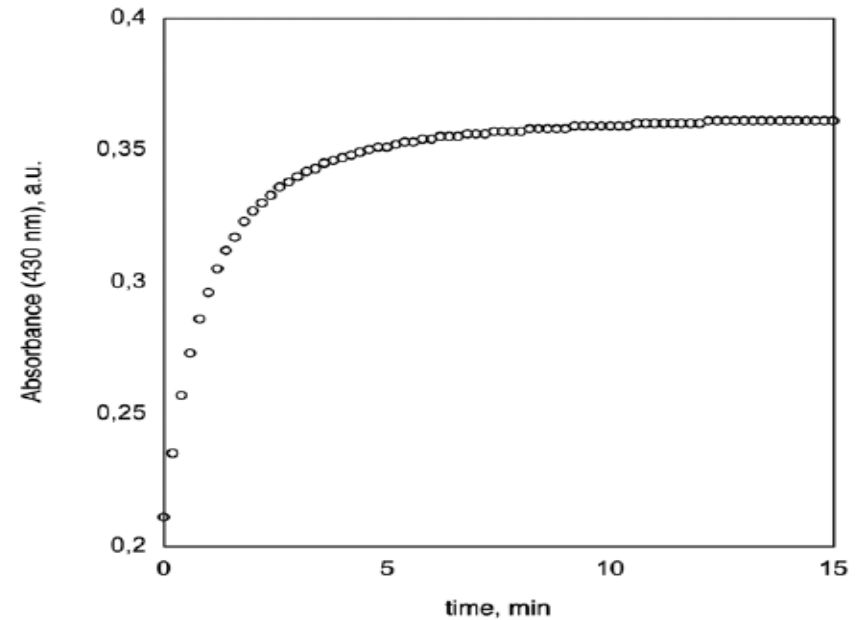
Cholic acid

Concentration-dependent aggregation

3 μM

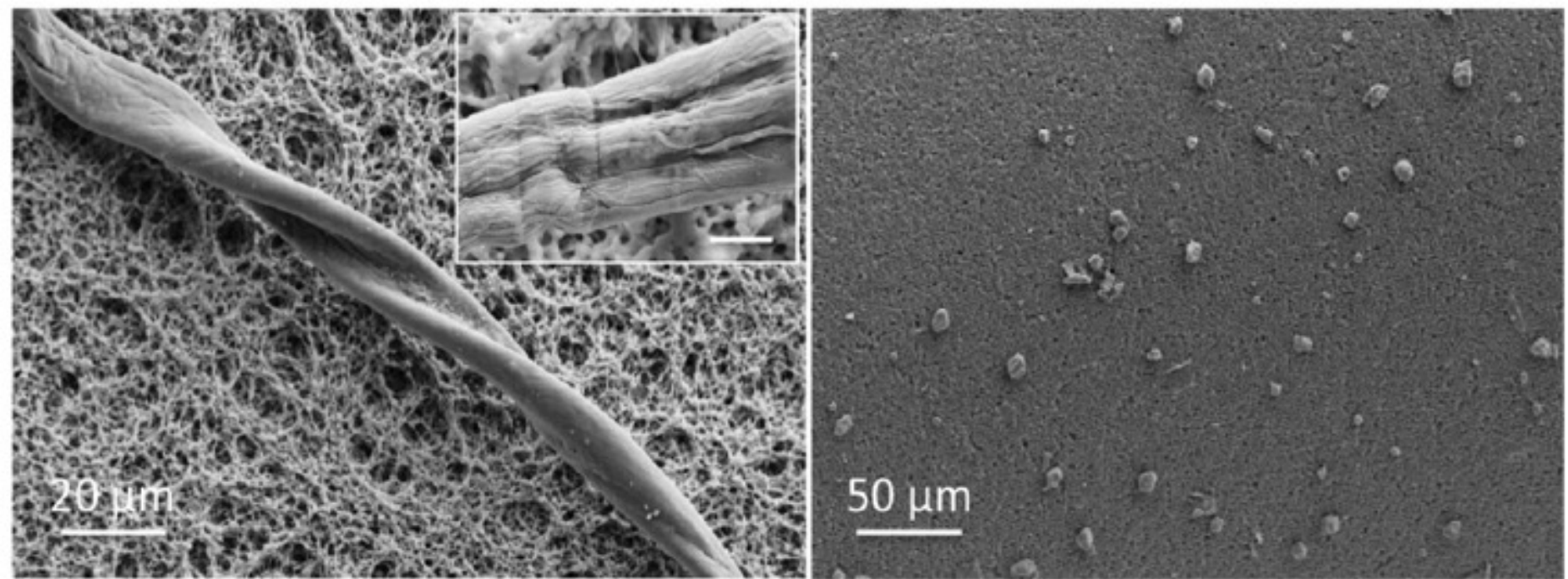


10 μM



in DMA/H₂O 58/42 (v/v)

Imaging Steroid Porphyrins superstructures



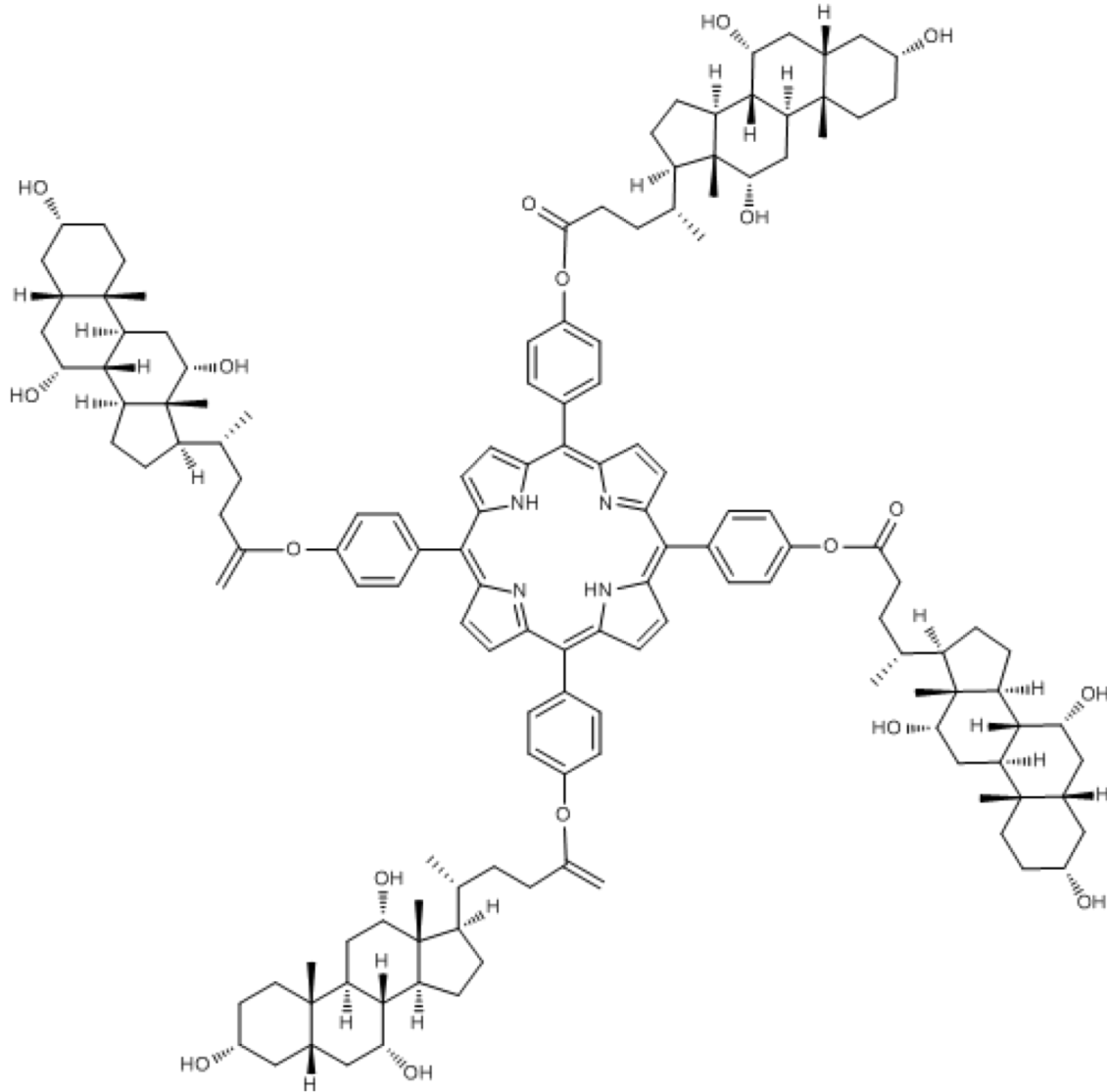
3 μM in DMA/H₂O 58/42 (v/v)

Entangled fibrils!

10 μM in DMA/H₂O 58/42 (v/v)

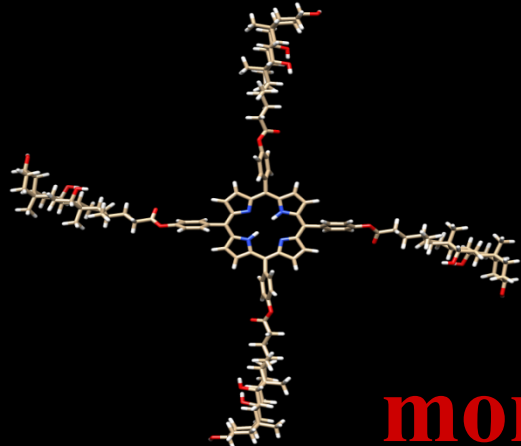
Globules!

Tetrasteroid porphyrins

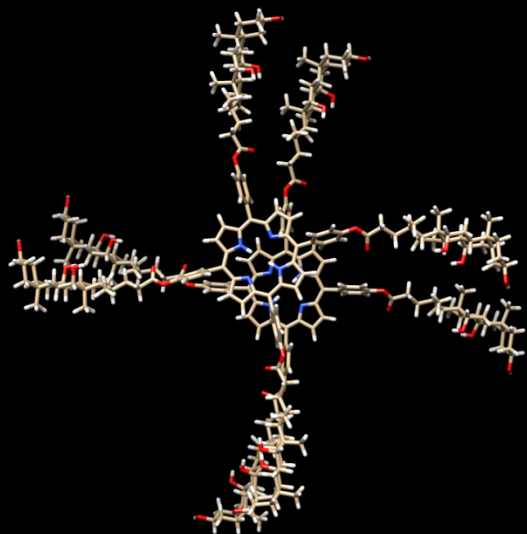


TSPc

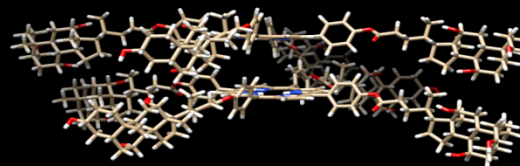
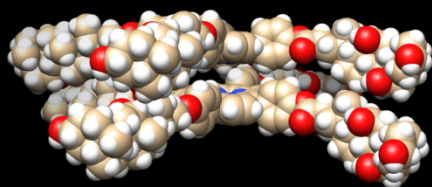
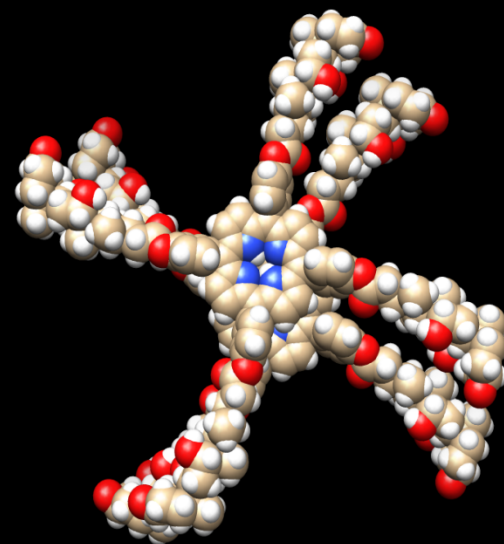
TSPc



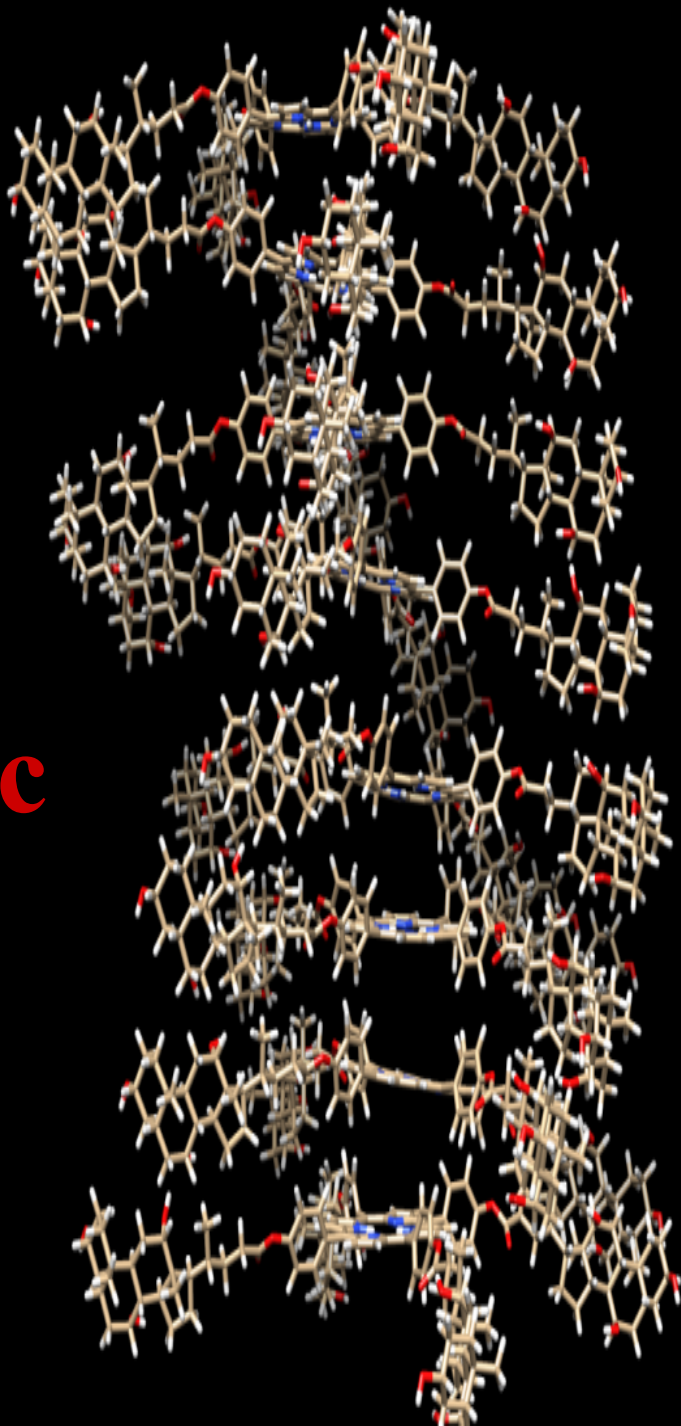
monomer



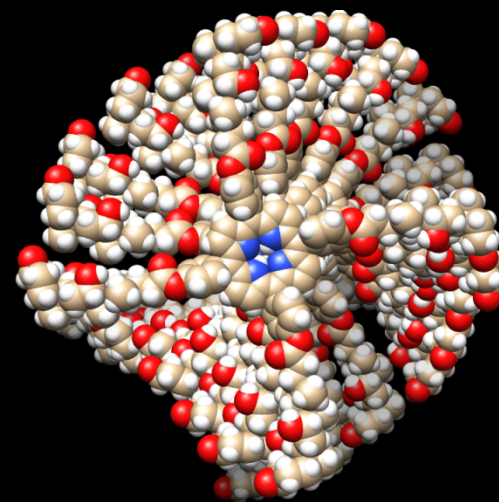
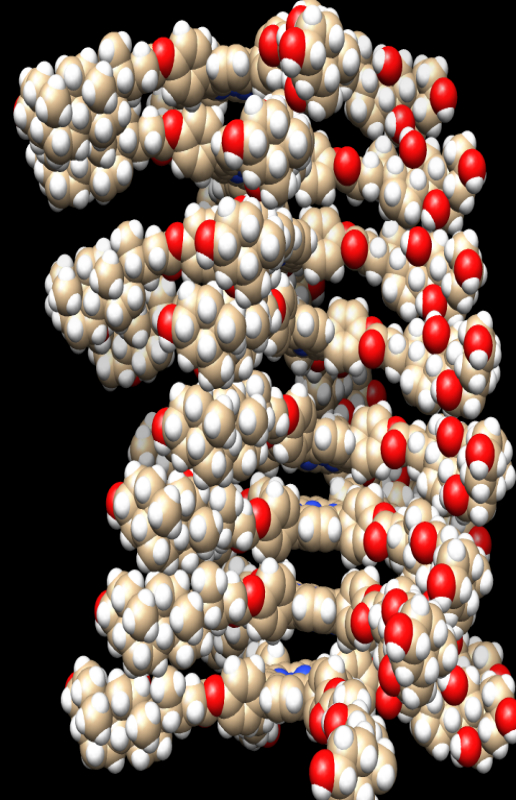
dimer



TSPc

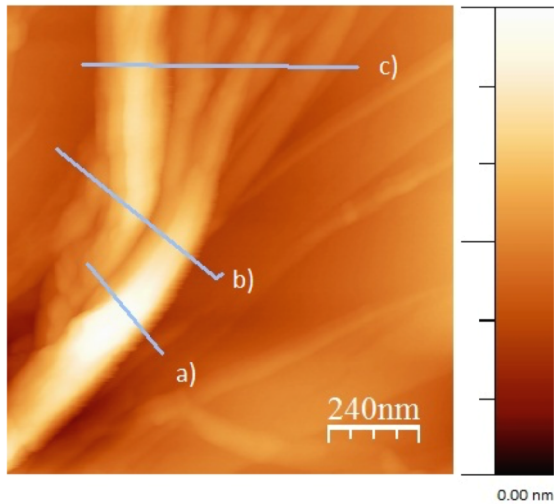
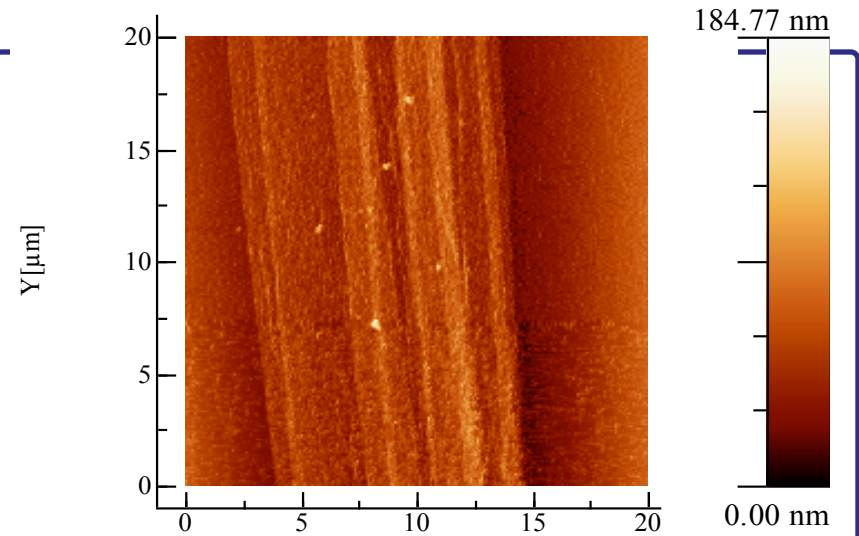
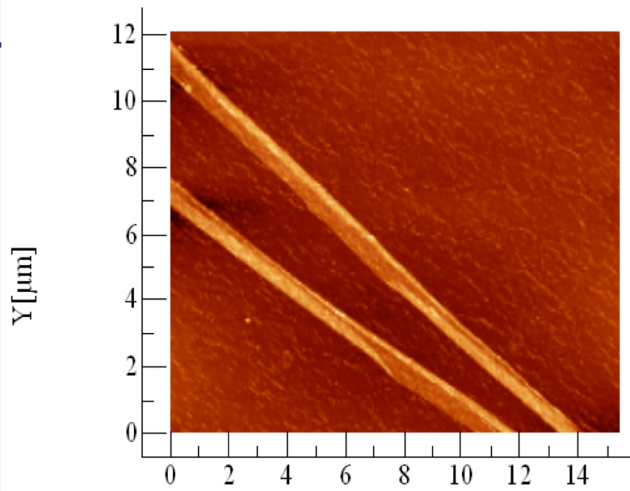


octamer

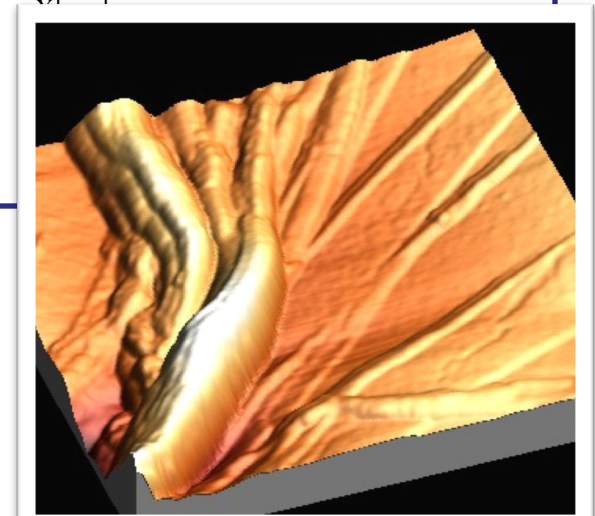


Atomic Force Microscopy

TSP: LB deposition on silicon wafer

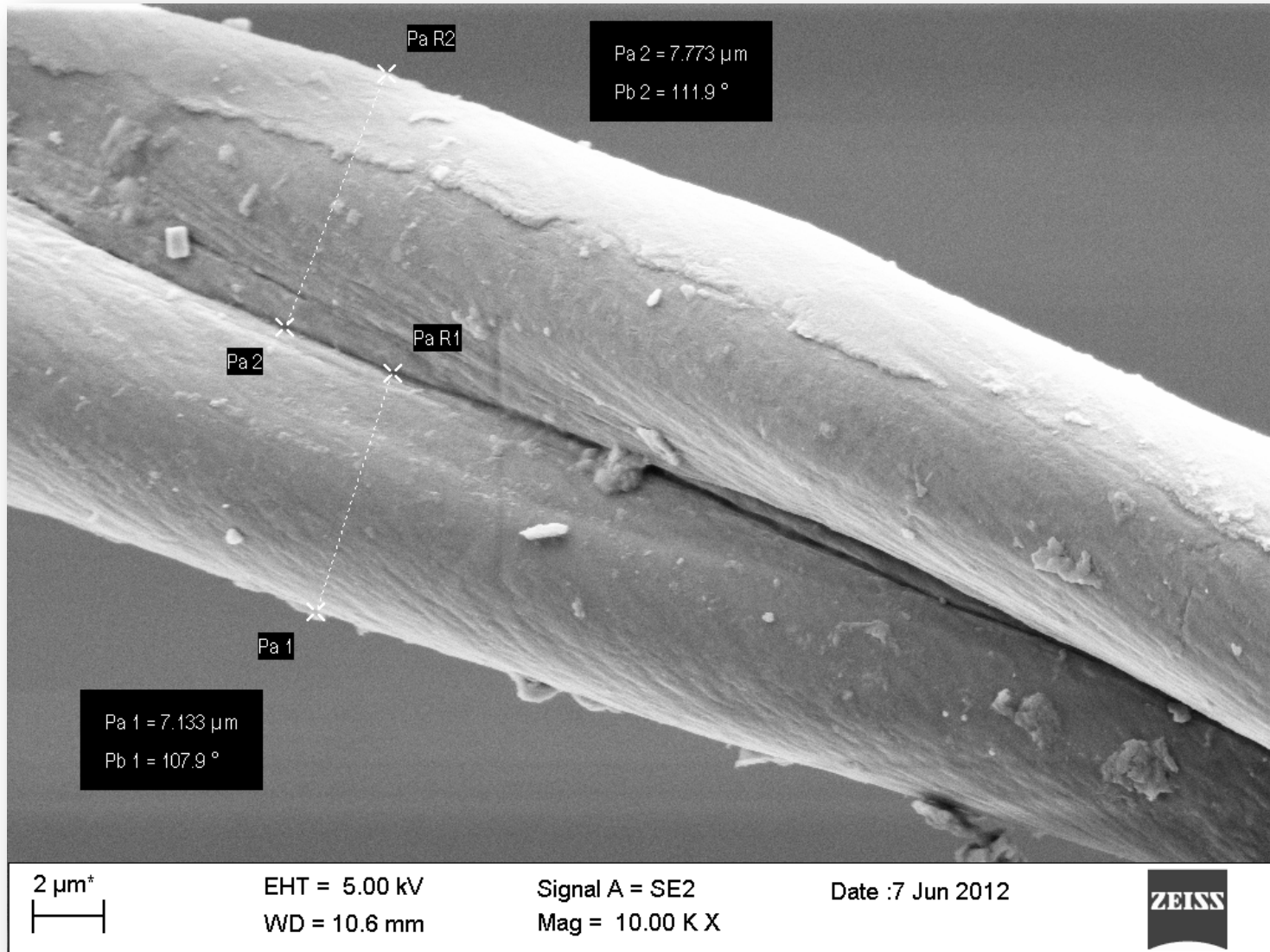


Diameter: 130 ± 30 nm
Thickness: 80 ± 20 nm



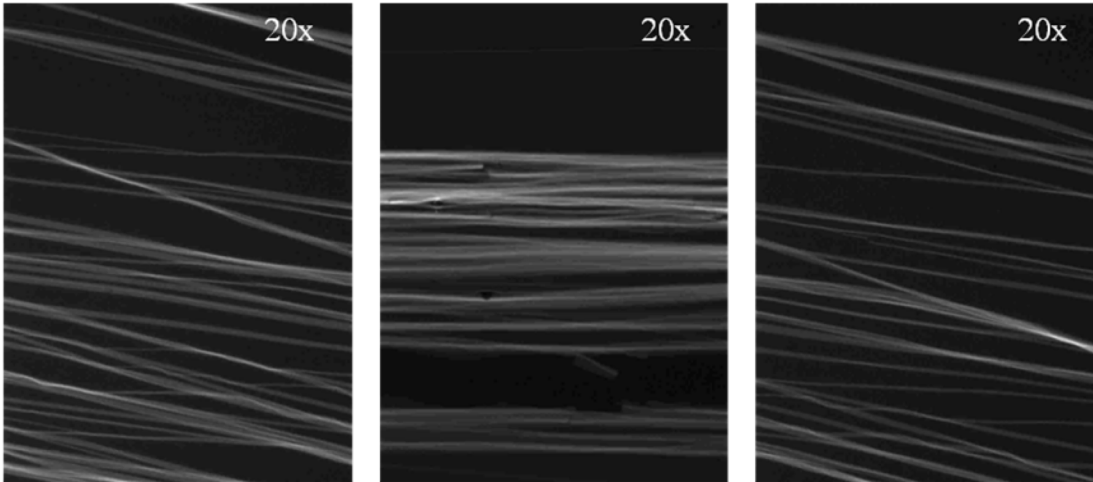
Scanning Electron Microscopy

TSPc multilayer on glass

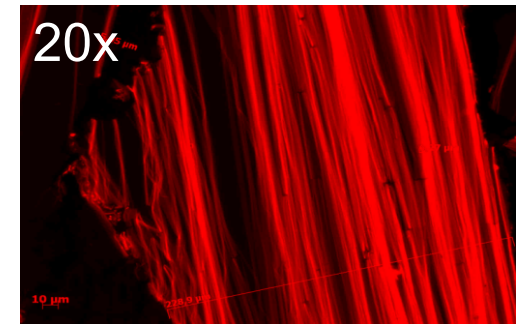
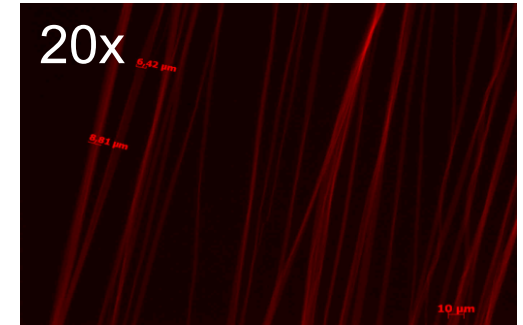


Fluorescence Microscopy of LB films

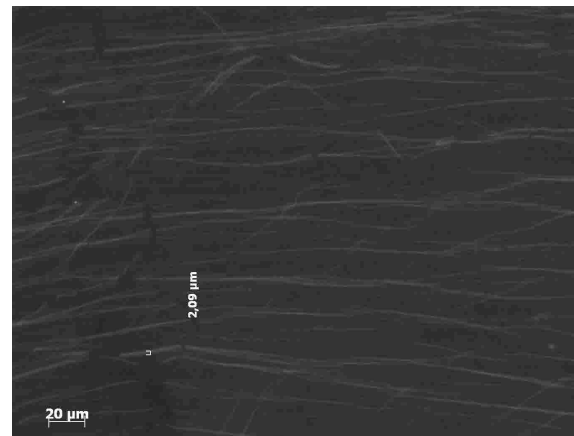
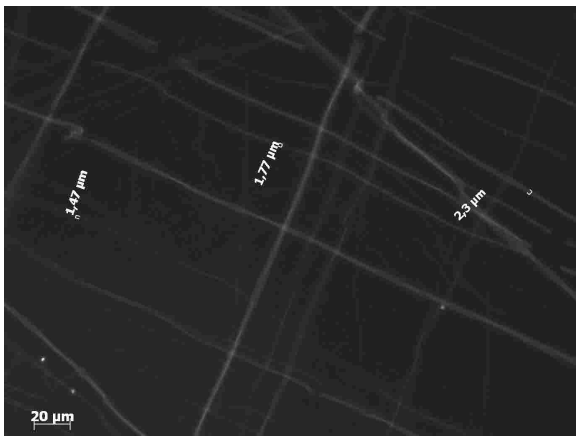
TSP (multilayer)



Fibrils of millimetric length



TSPc (multilayer)

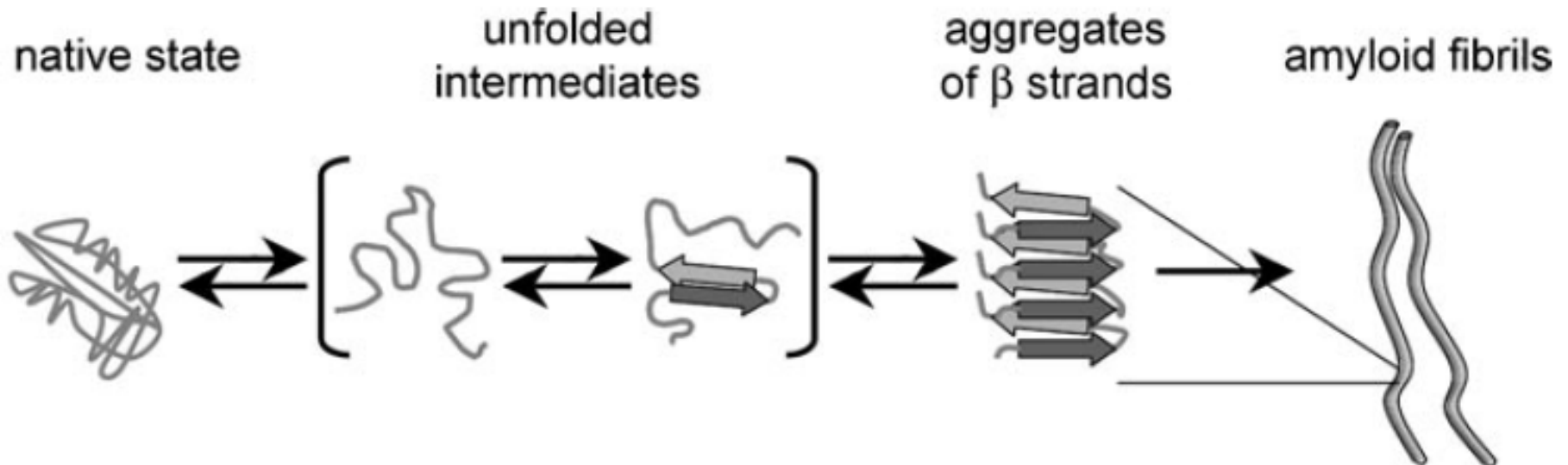


Thickness:

$2.4 \pm 0.6 \mu\text{m}$

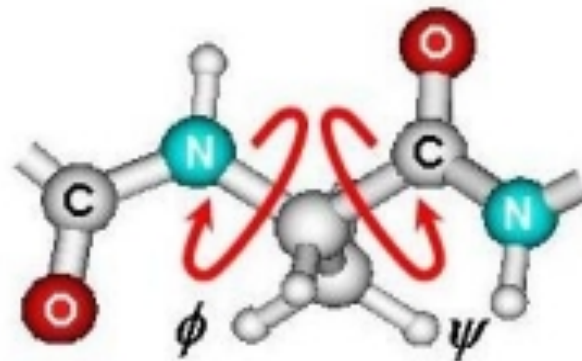
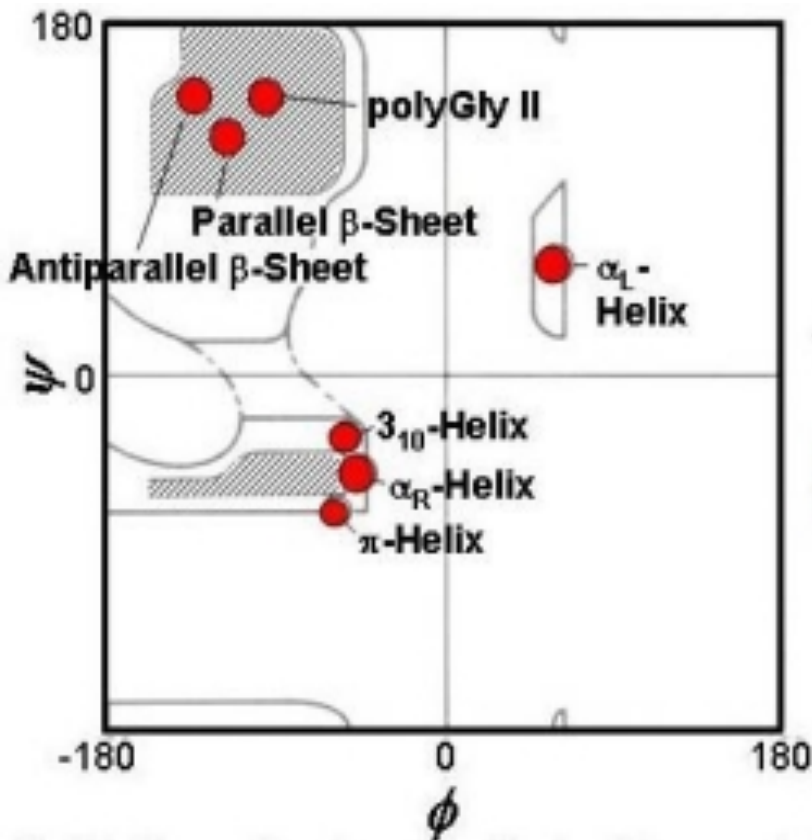
Hierarchical self-assembly

A self-assembly process that can be broken down into distinct steps that can not proceed until the preceding step is complete.



Peptide Foldamers

‘Foldamers are molecules that have well-defined and predictable folding properties in solution.’

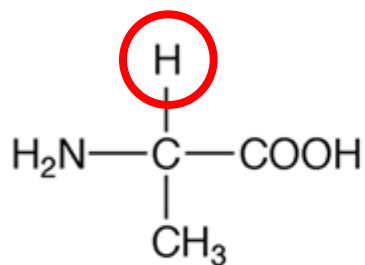
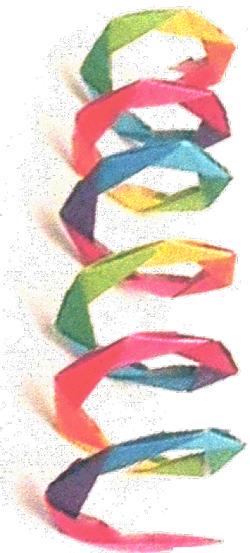


Claudio Toniolo

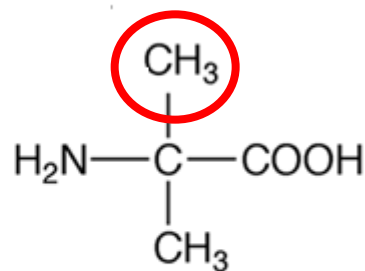


Fernando Formaggio

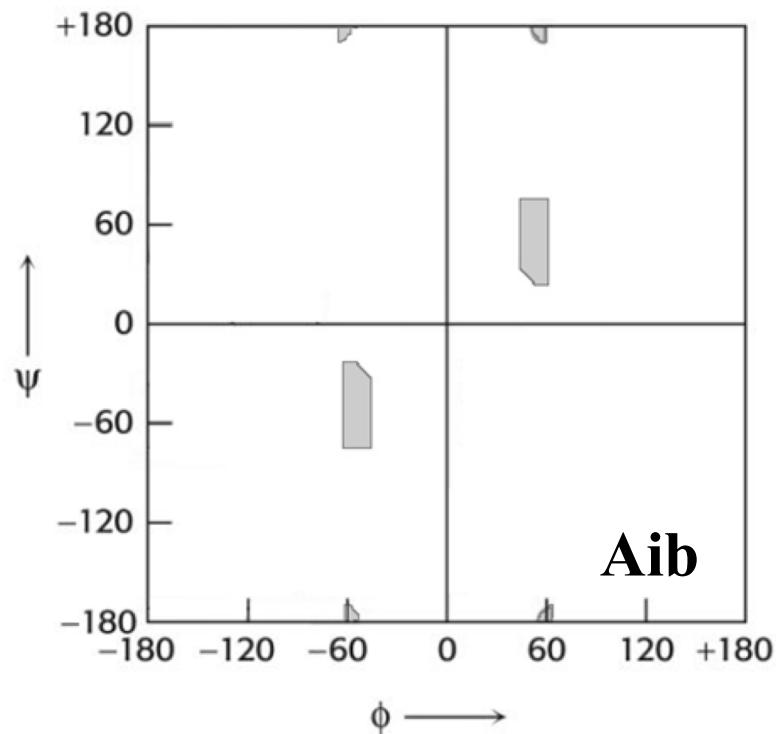
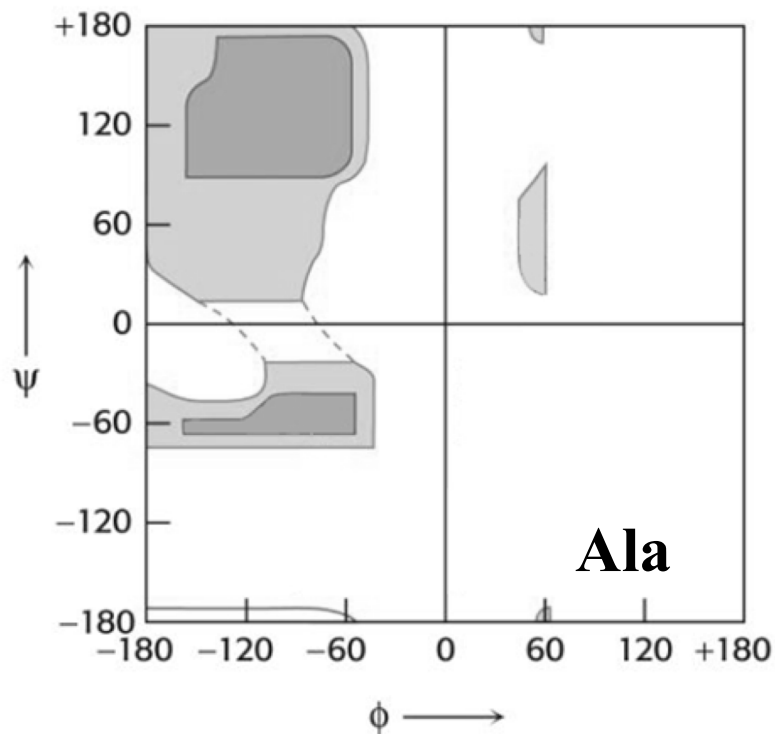
$C^{\alpha,\alpha}$ -disubstituted aminoacids



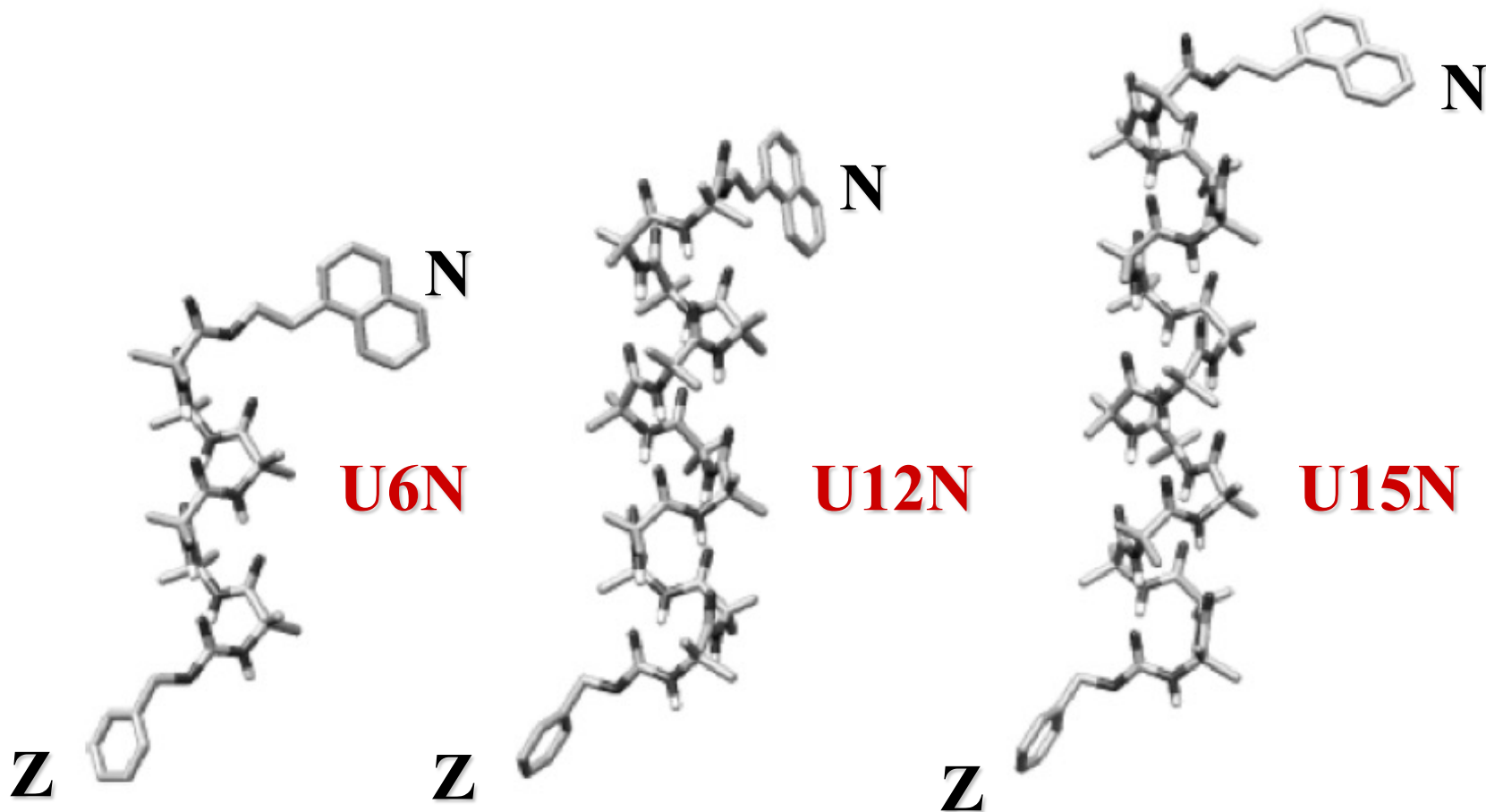
Ala



Aib



Aggregation of conformationally constrained model peptides



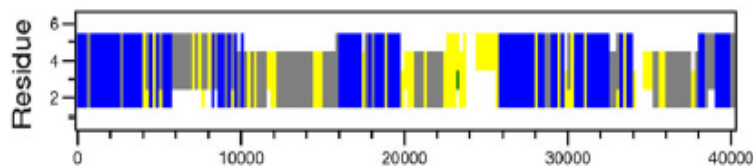
U = Aib; N = Naphthyl; Z = Benzyloxycarbonyl

Molecular Dynamics

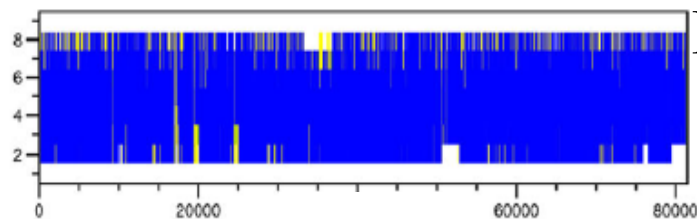


Gianfranco
Bocchini

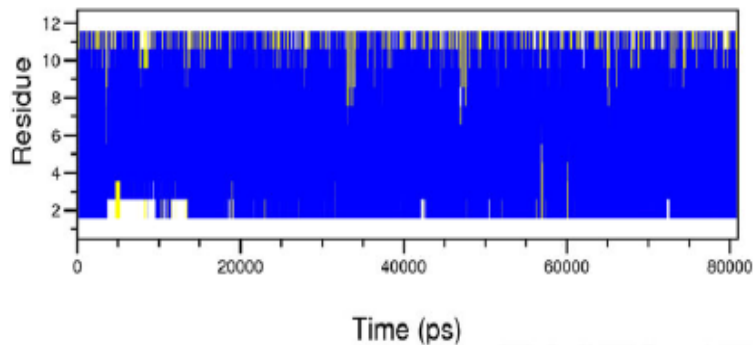
U6N



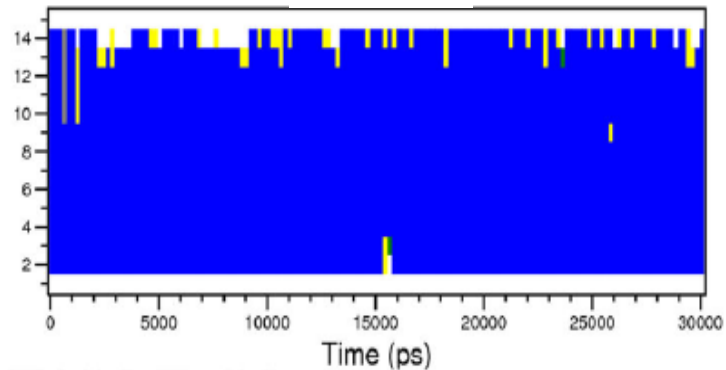
U9N



U12N



U15N

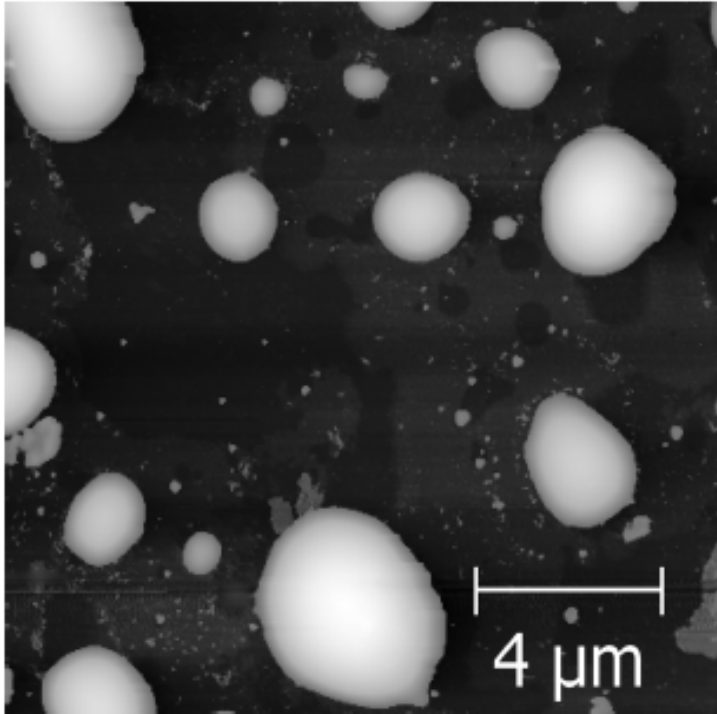


□ Coil ■ Bend ■ Turn ■ α -Helix ■ 3_{10} -Helix

For the longer homologues of the series ($n \geq 9$), unfolding events during the simulation time were not detected.

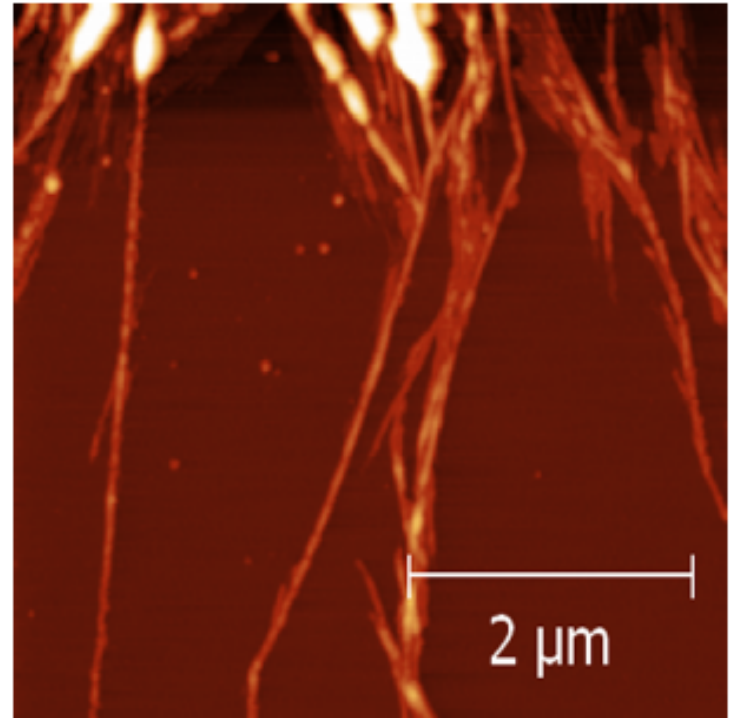
Atomic Force Microscopy

10 μM 70/30 v/v MeOH/H₂O solutions on mica



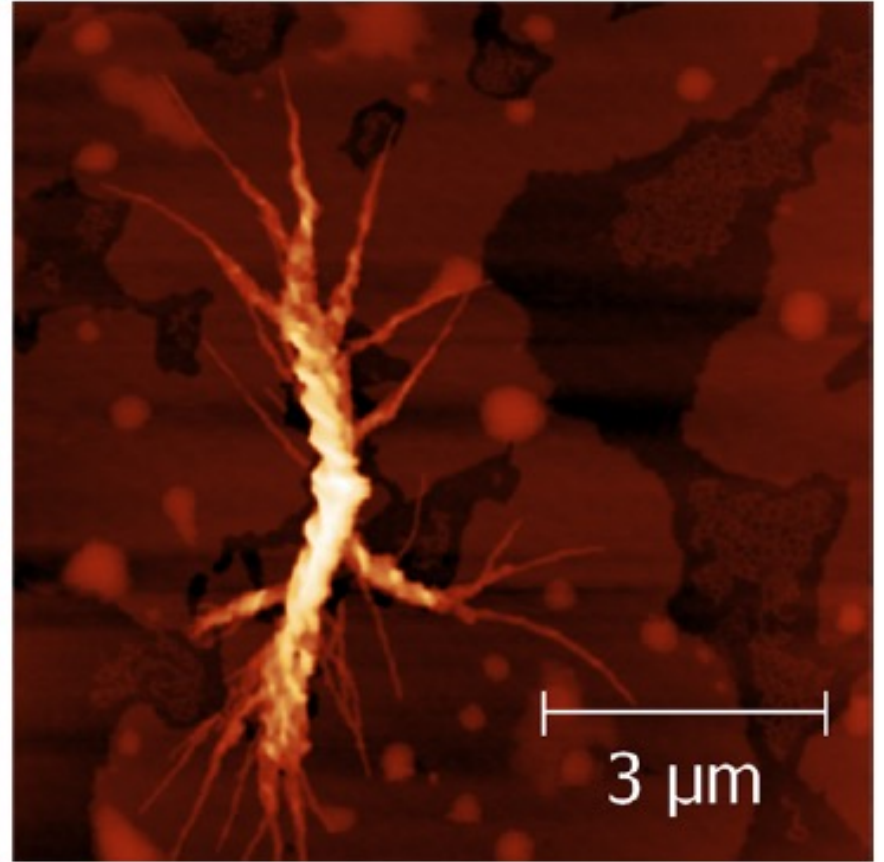
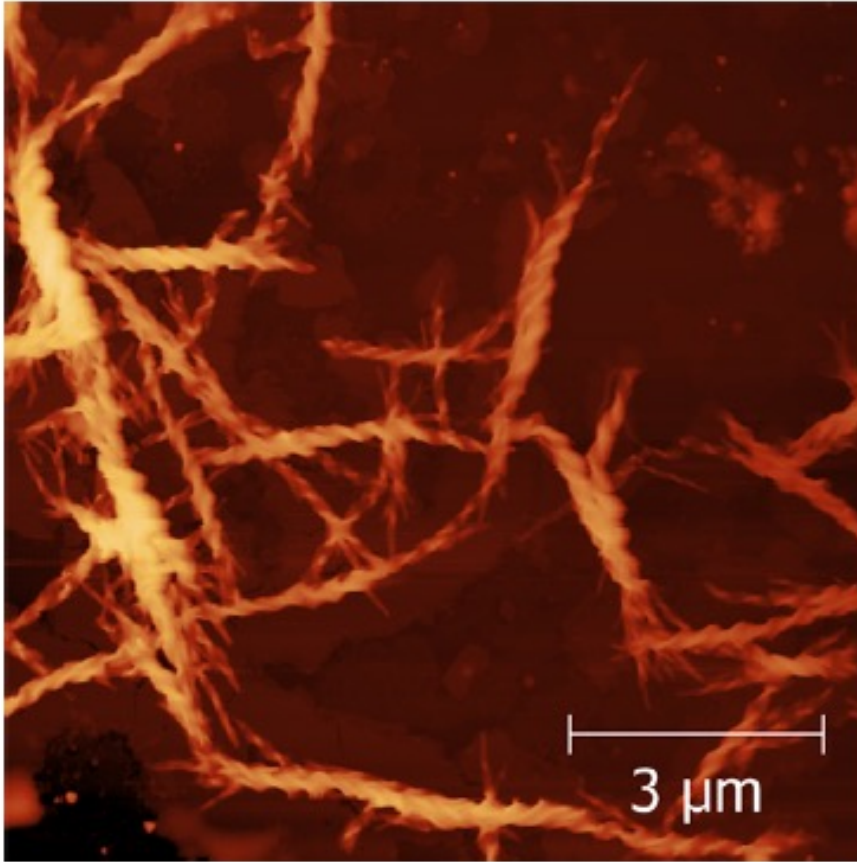
U6N

Hydrophobic effects predominate



U15N

Secondary structure drives hierarchical self-assembly



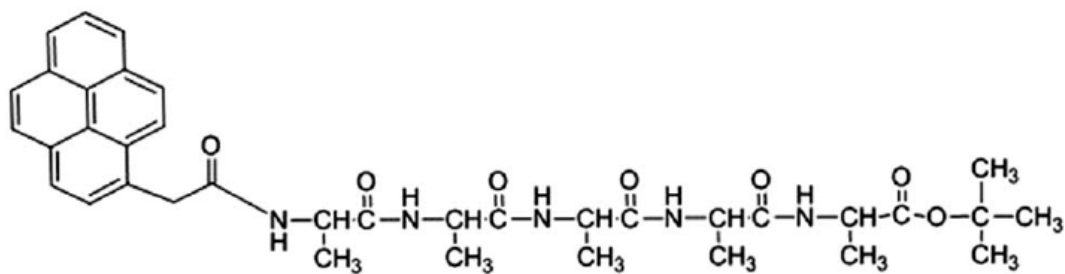
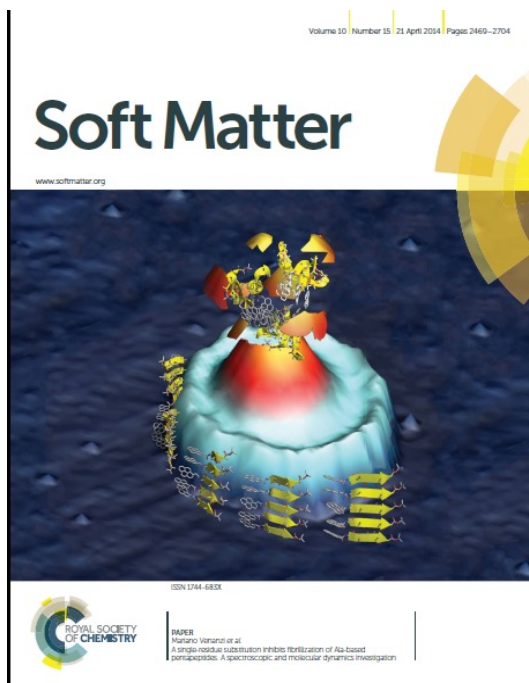
U12N

(from 125 μM 70/30 v/v MeOH/H₂O deposition solution)

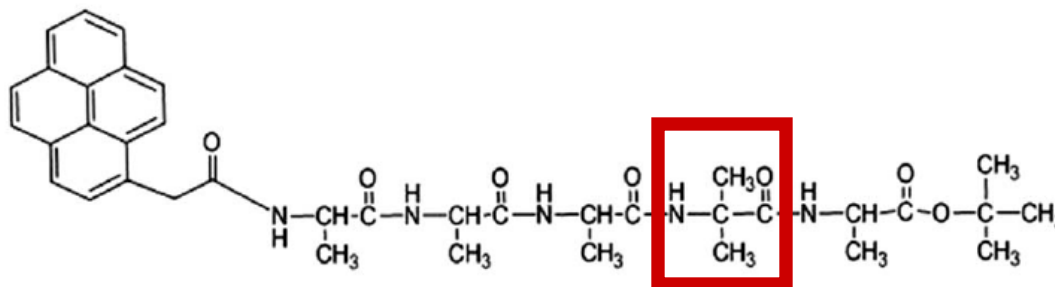
Cite this: *Soft Matter*, 2014, 10, 2508

A single-residue substitution inhibits fibrillization of Ala-based pentapeptides. A spectroscopic and molecular dynamics investigation†

Mario Caruso,^a Emanuela Gatto,^a Ernesto Placidi,^b Gema Ballano,^c Fernando Formaggio,^c Claudio Toniolo,^c David Zanuy,^d Carlos Alemán^{d,e} and Mariano Venanzi^{*a}



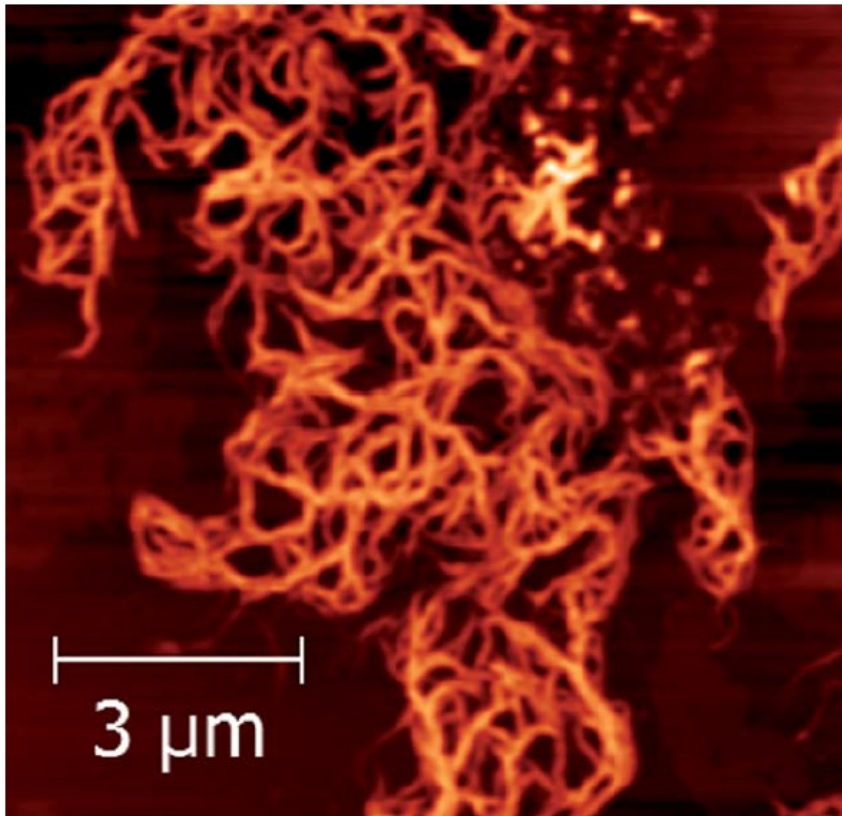
Py-CH₂-CO-(L-Ala)₅-OtBu (PyA5)



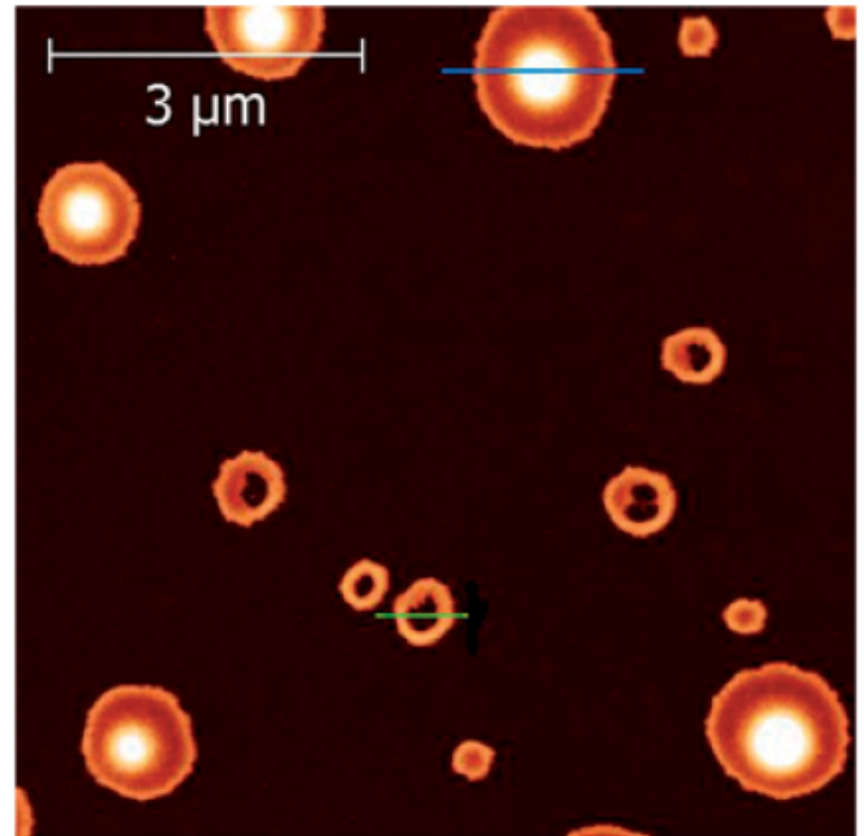
Py-CH₂-CO-(L-Ala)₃-Aib-L-Ala-OtBu (PyA3UA)

Atomic Force Microscopy

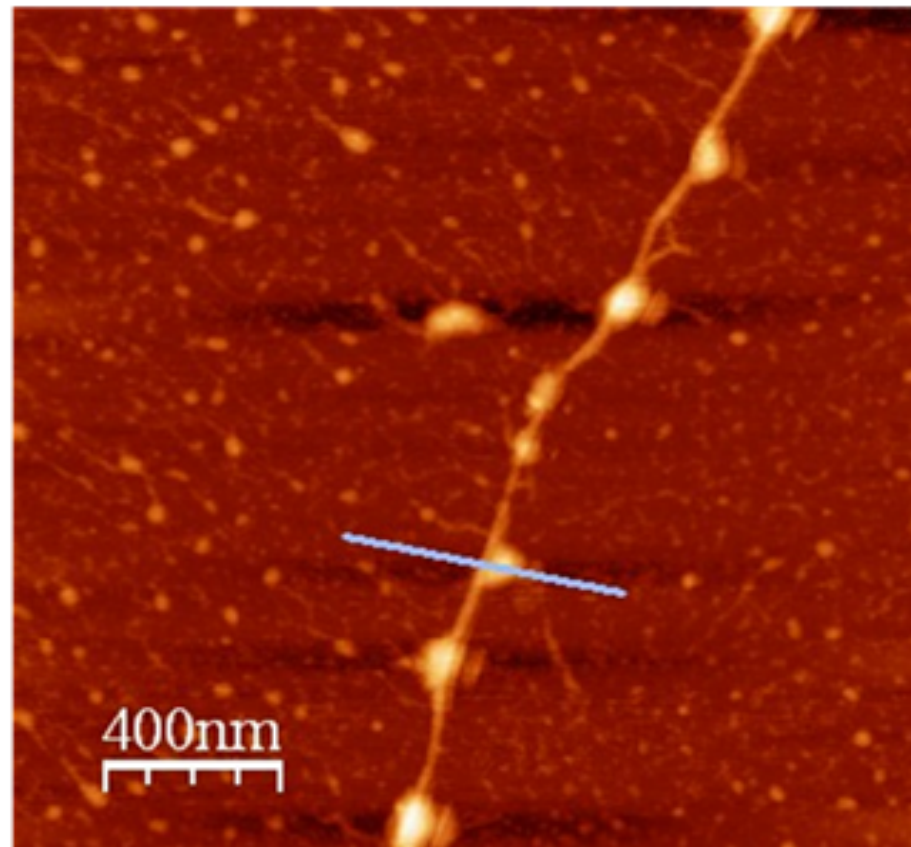
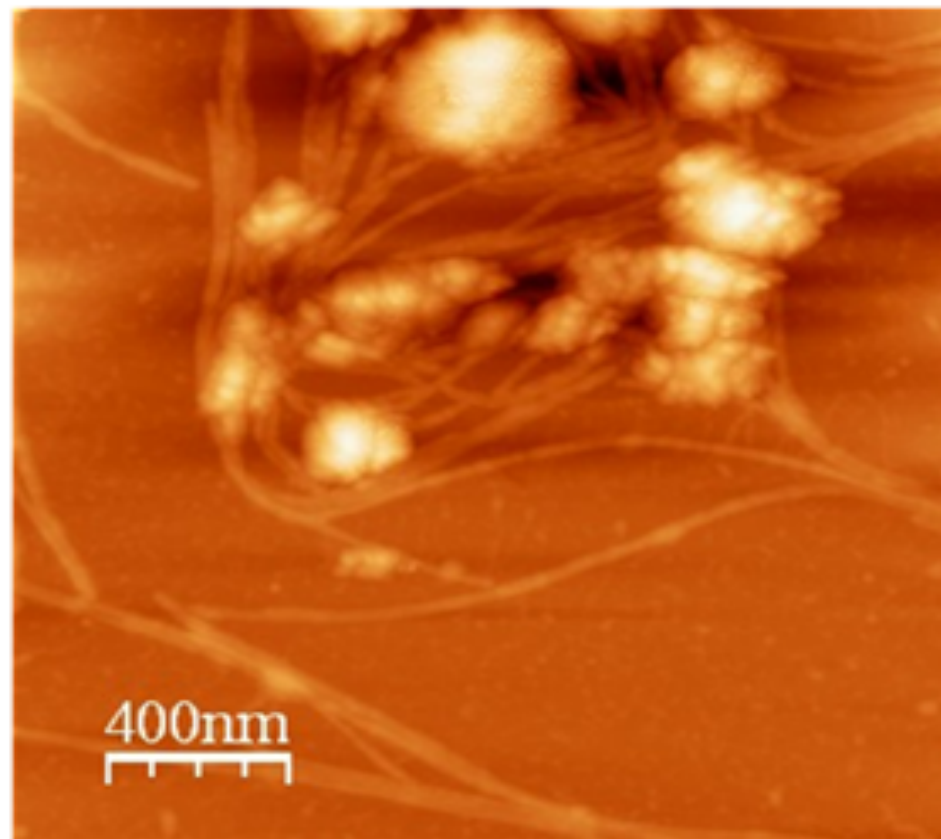
PyA5 on mica



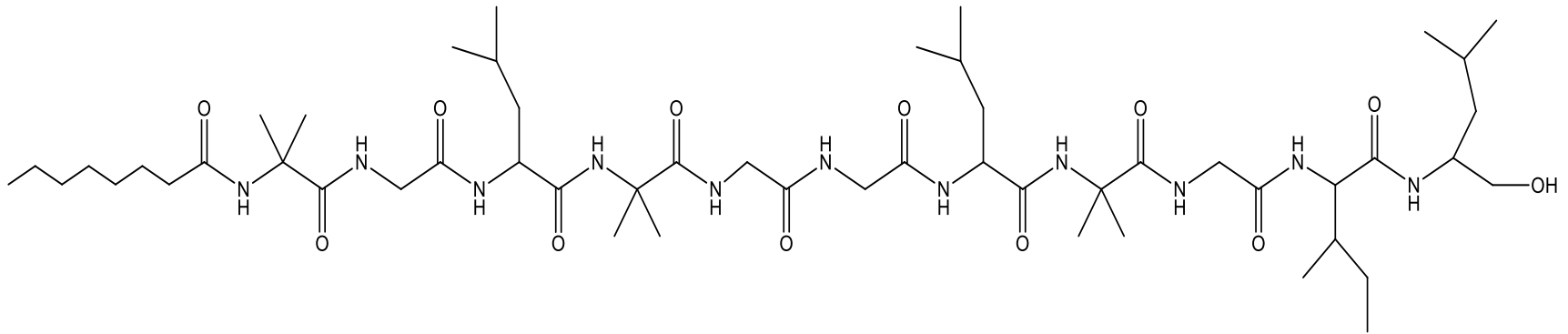
PyA3UA on mica



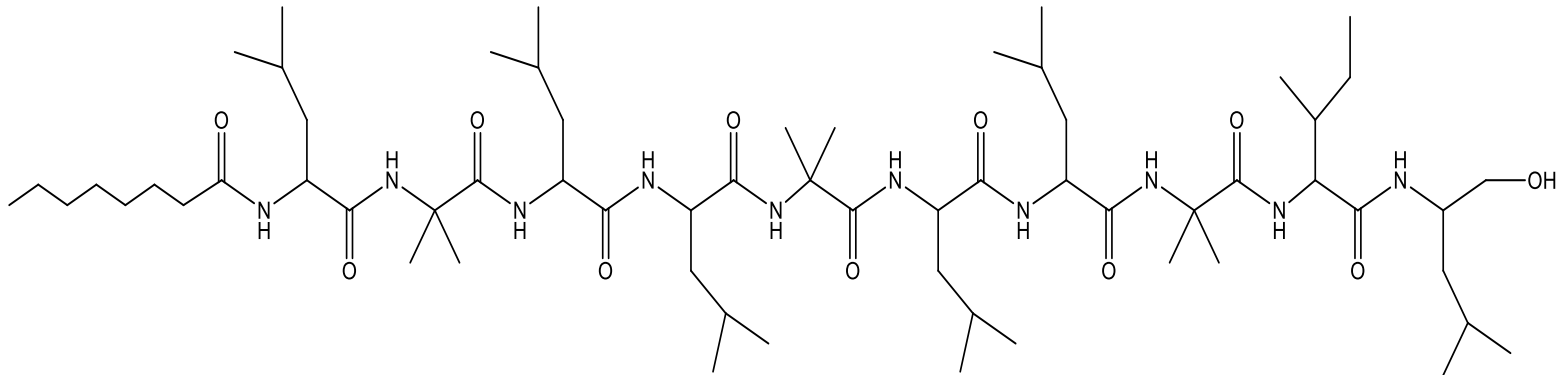
PyA5 LB films on mica



An antimicrobial peptide: Trichogin GA IV



Oct-Aib-Gly-Leu-Aib-Gly-Gly-Leu-Aib-Gly-Ile-Lol
(TrGA)



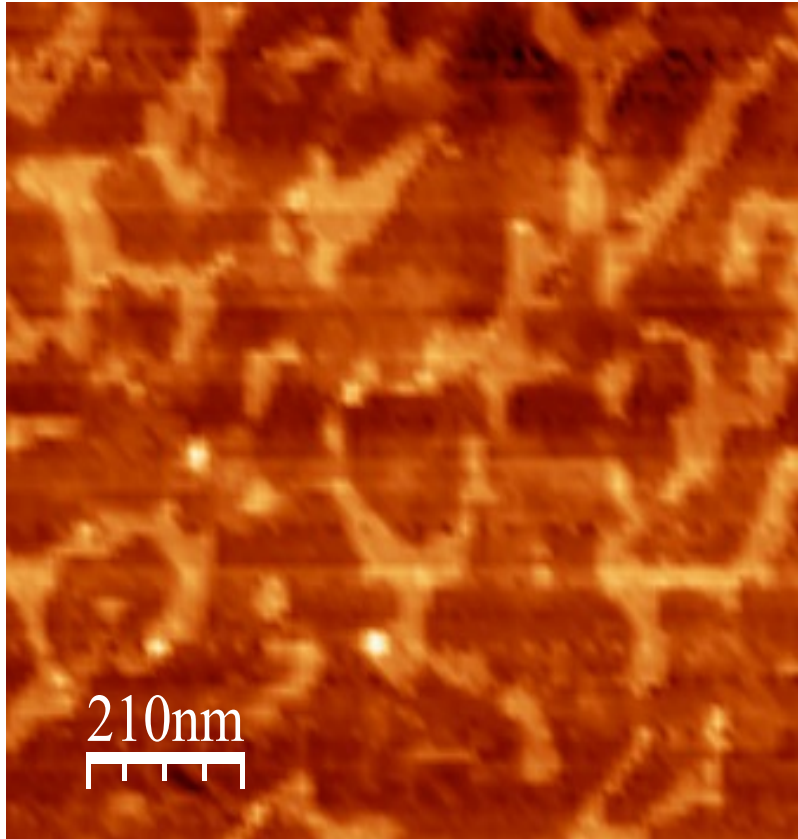
Oct-Leu-Aib-Leu-Leu-Aib-Leu-Leu-Aib-Ile-Lol
(TrGAR)

Marta De Zotti
(University of Padova)

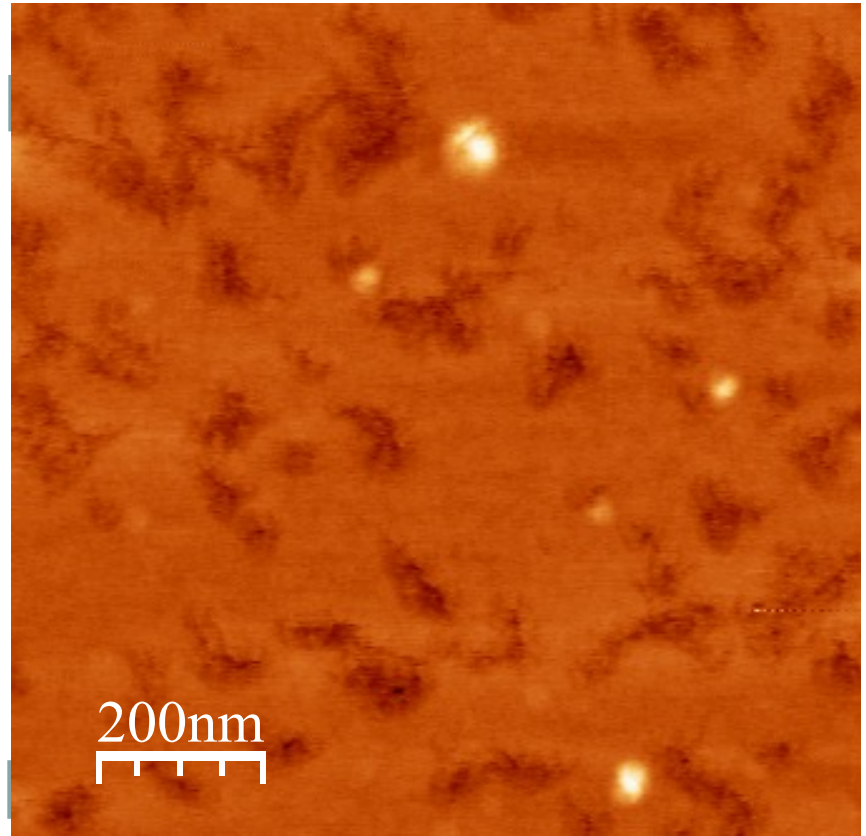
TrGA

LB films

Graphite (hydrophobic)



Mica (hydrophilic)



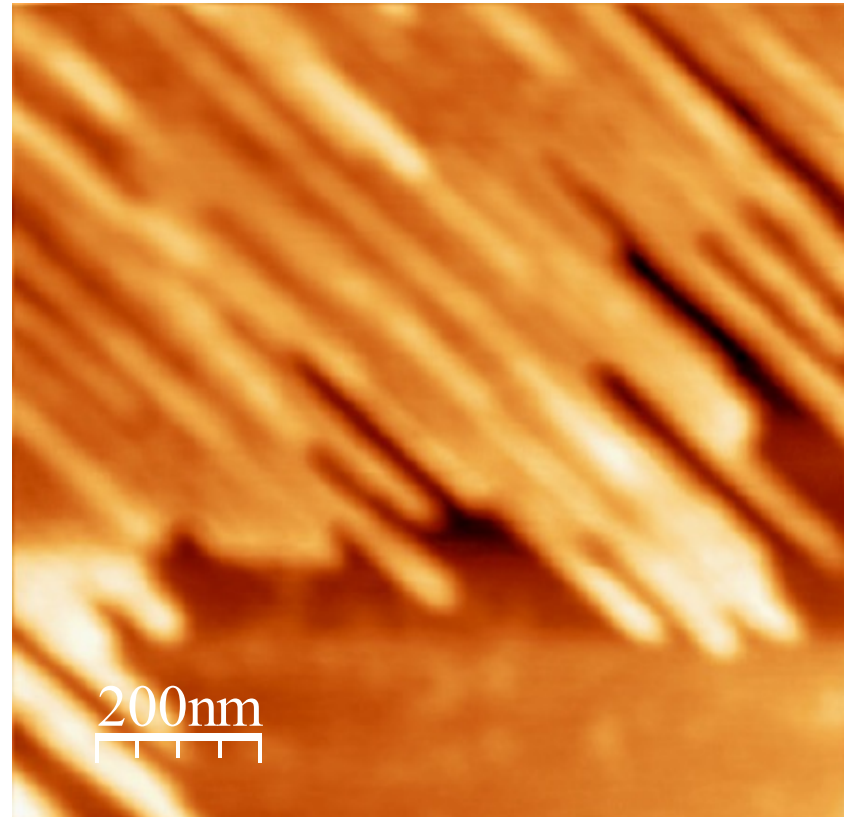
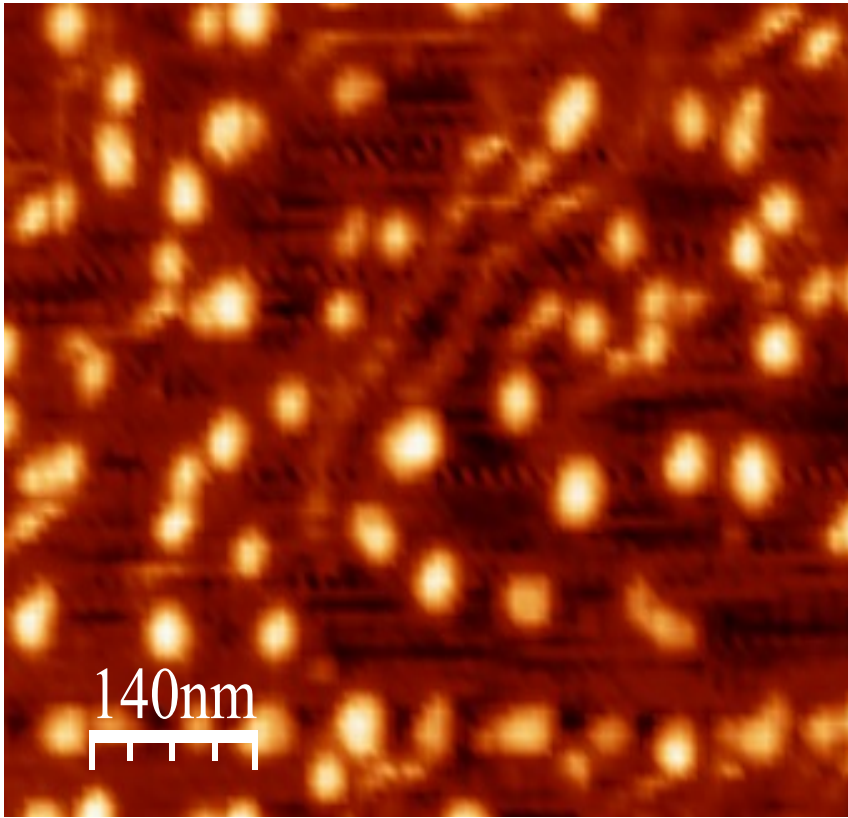
at 40 mN/m surface pressure

TrGAR

LB films

Graphite

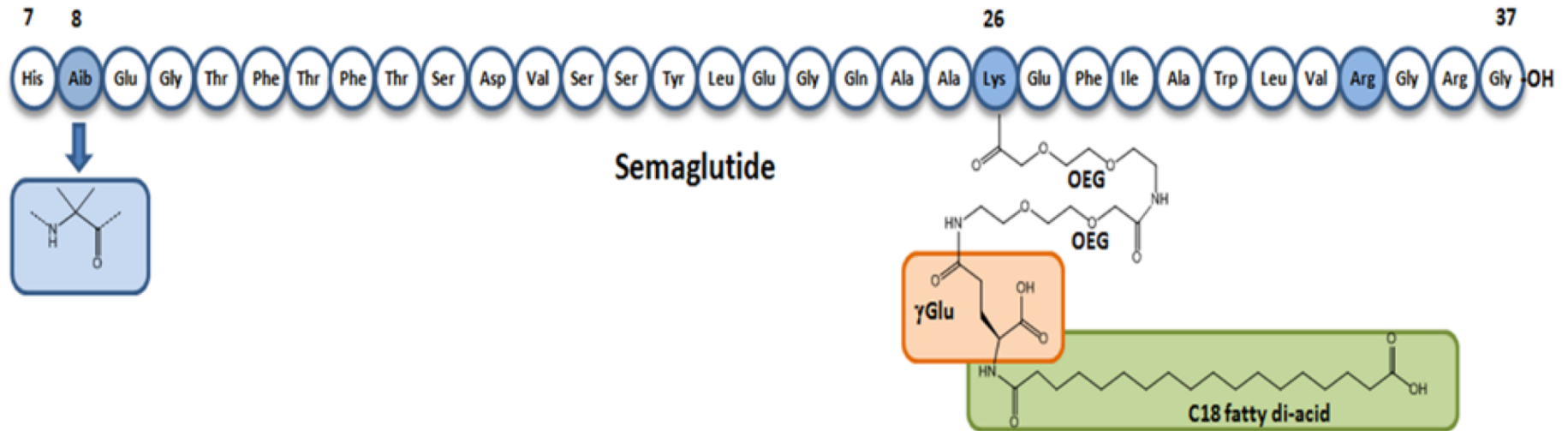
Mica



at 20 mN/m surface pressure

A therapeutic peptide

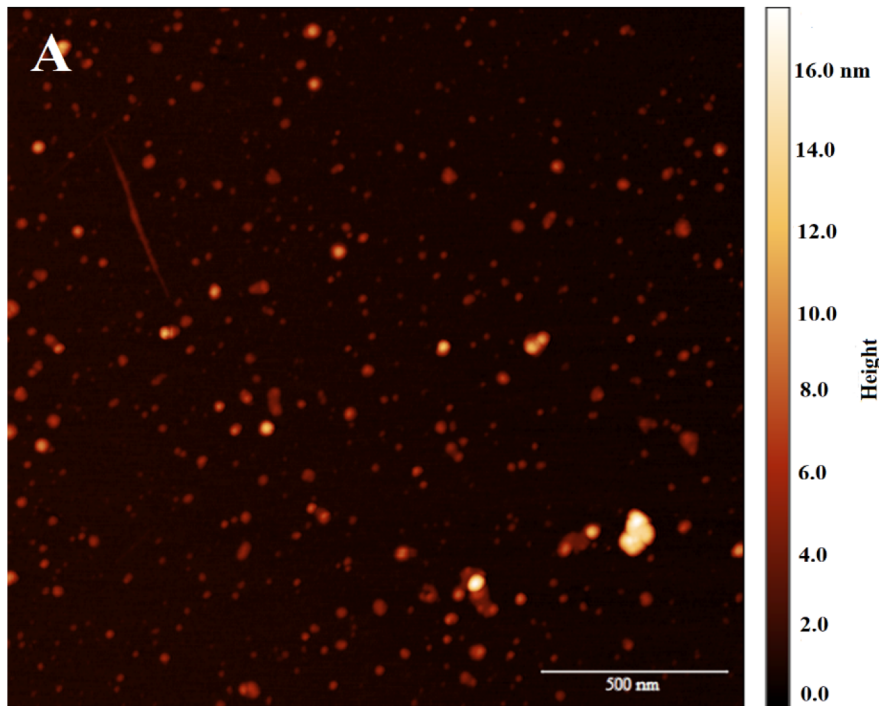
Semaglutide, a GLP-1 analog (IRBM)



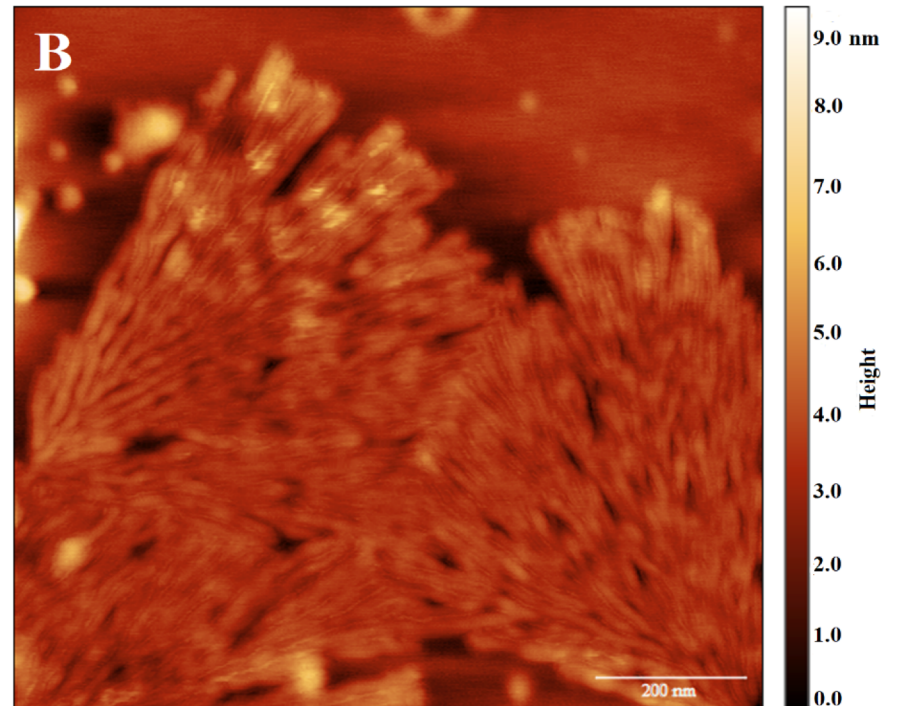
A spectroscopic and molecular dynamics study on the aggregation process of a long-acting lipidated therapeutic peptide: the case of semaglutide.

Venanzi M. et al. *Soft Matter* 2020, 16, 10122-10131

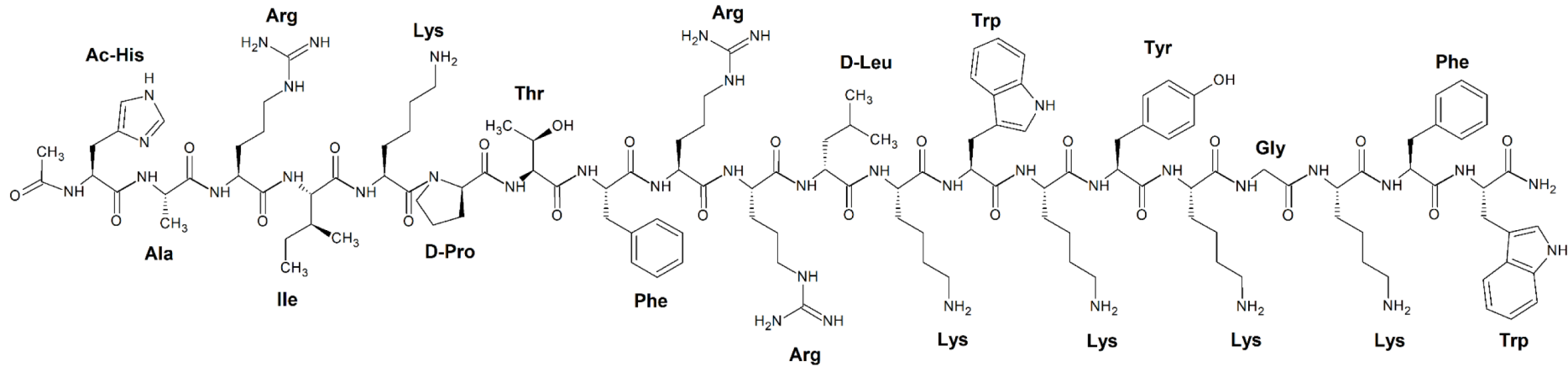
A: fresh solutions



B: aged solutions



A therapeutic peptide: CIGB552



Ac-HARIK(dP)TFRR(dL)KWKYKGKFW-NH₂

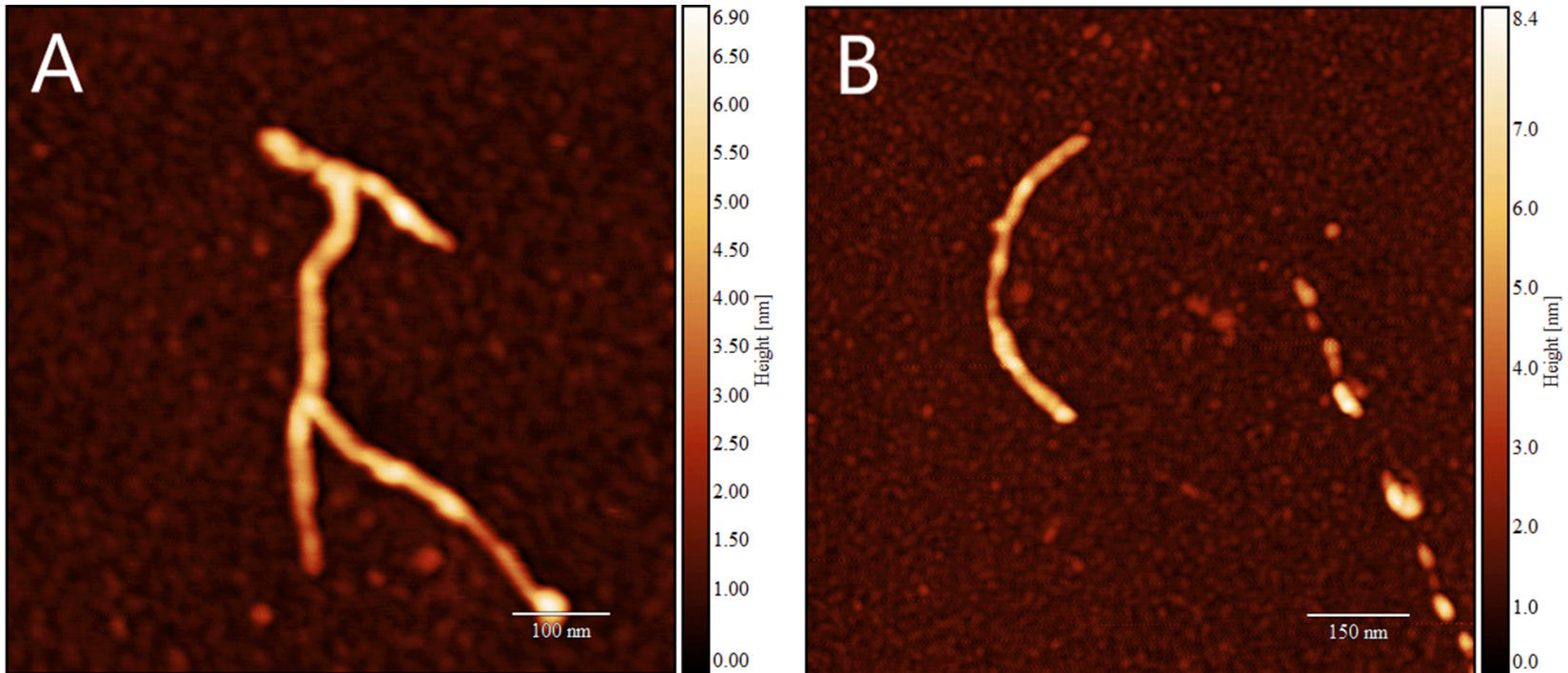
Antitumor activity was demonstrated in mouse cells.

Formulation matters! A spectroscopic and molecular dynamics investigation on the peptide CIGB552 as itself and in its therapeutical formulation

M. Savioli et al. J. Pept. Sci. 2021, <https://doi.org/10.1002/psc.3356>

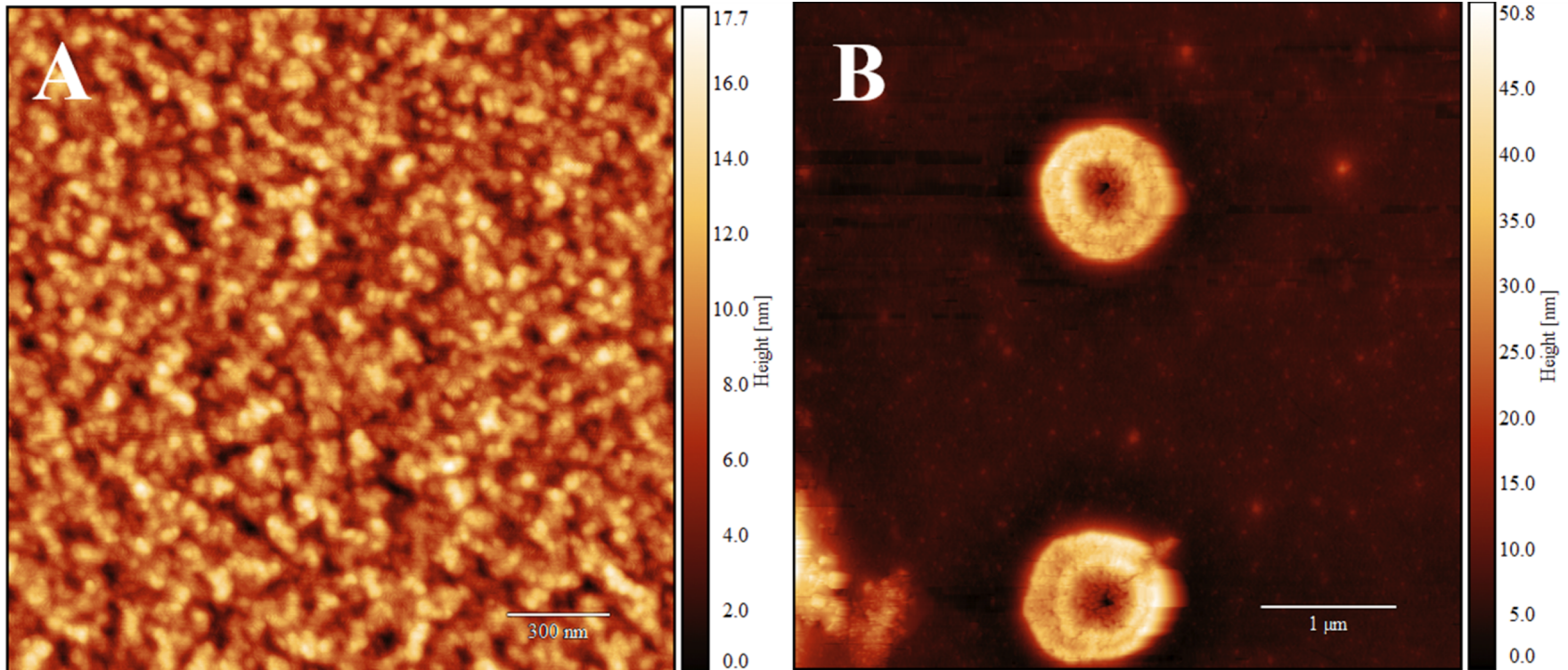
Atomic Force Microscopy

CIGB552



Filaments of lengths comprised between 20 and 32 μm and width of 2 μm were observed. B: globular structures nucleating the growth of peptide fibrils.

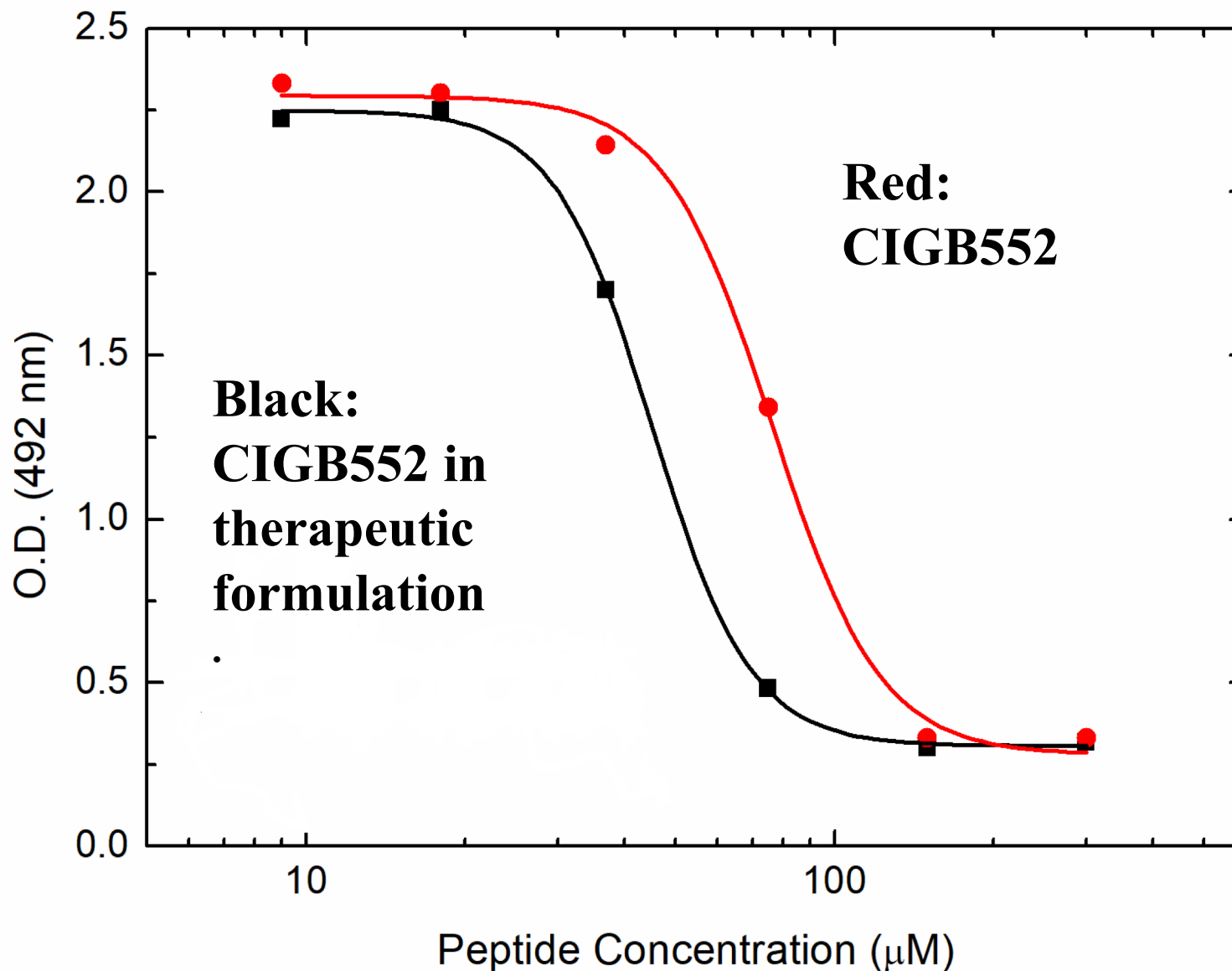
CIGB552 in therapeutic formulation



A: peptide nanoglobules (diameters between 10 - 50 nm)

B: micrometric annular structures (outer diameter: 1 μm; inner diameter: 60 nm).

Cell viability assay on H-460 tumour cells



The still open question is:

***Do peptide nanostructures play a specific
therapeutic role?***

Take home messages

Controlling self-assembly	Response
Thermodynamic , kinetic and spatio-temporal aspects should be considered (systems chemistry)	✓
Structure and dynamics properties of molecular building blocks matter!	✓
The morphology of nanostructures can be tuned by molecular engineering (hierarchical self-assembly)	✓
Nanostructures show emergent properties at each level	✓