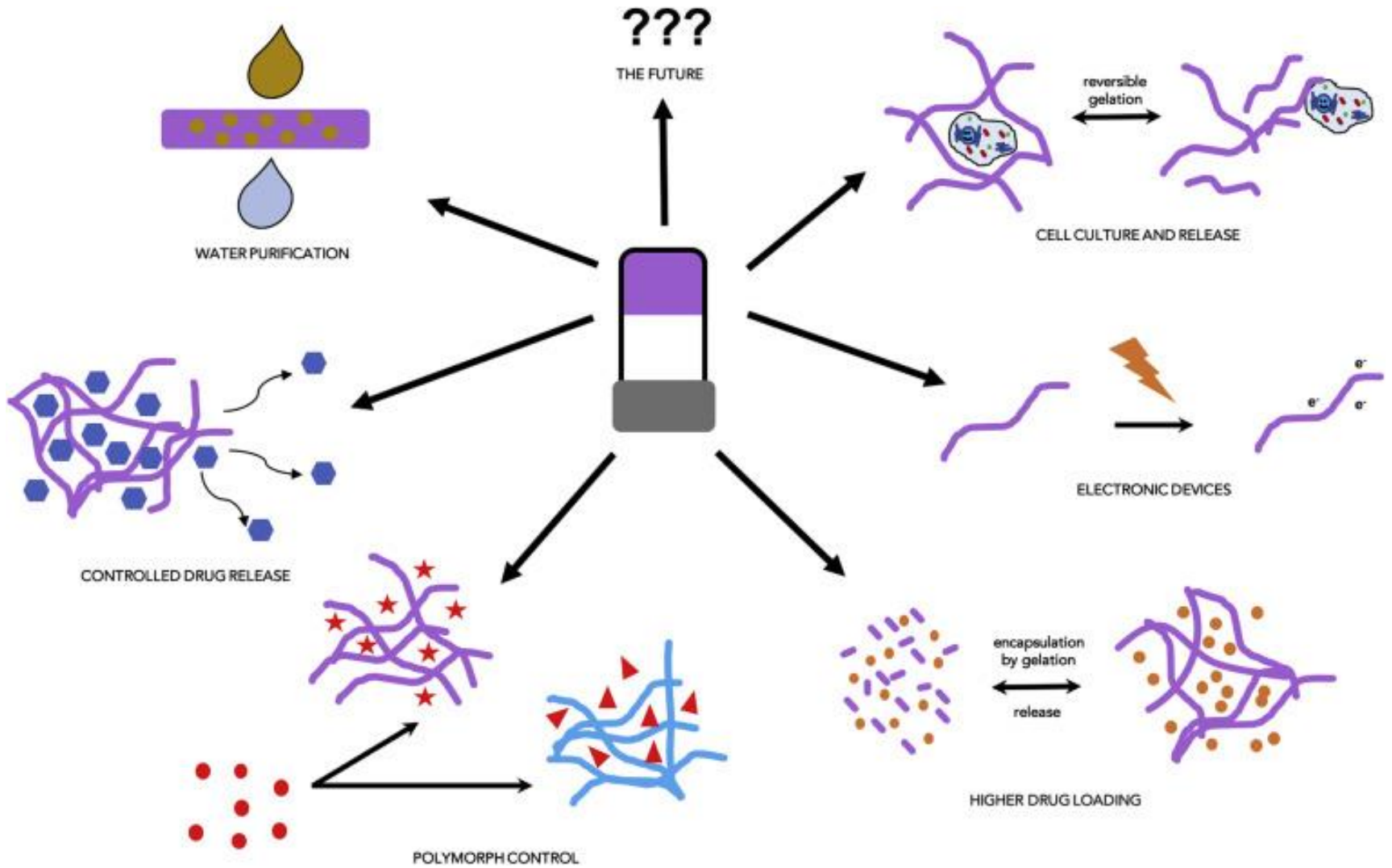
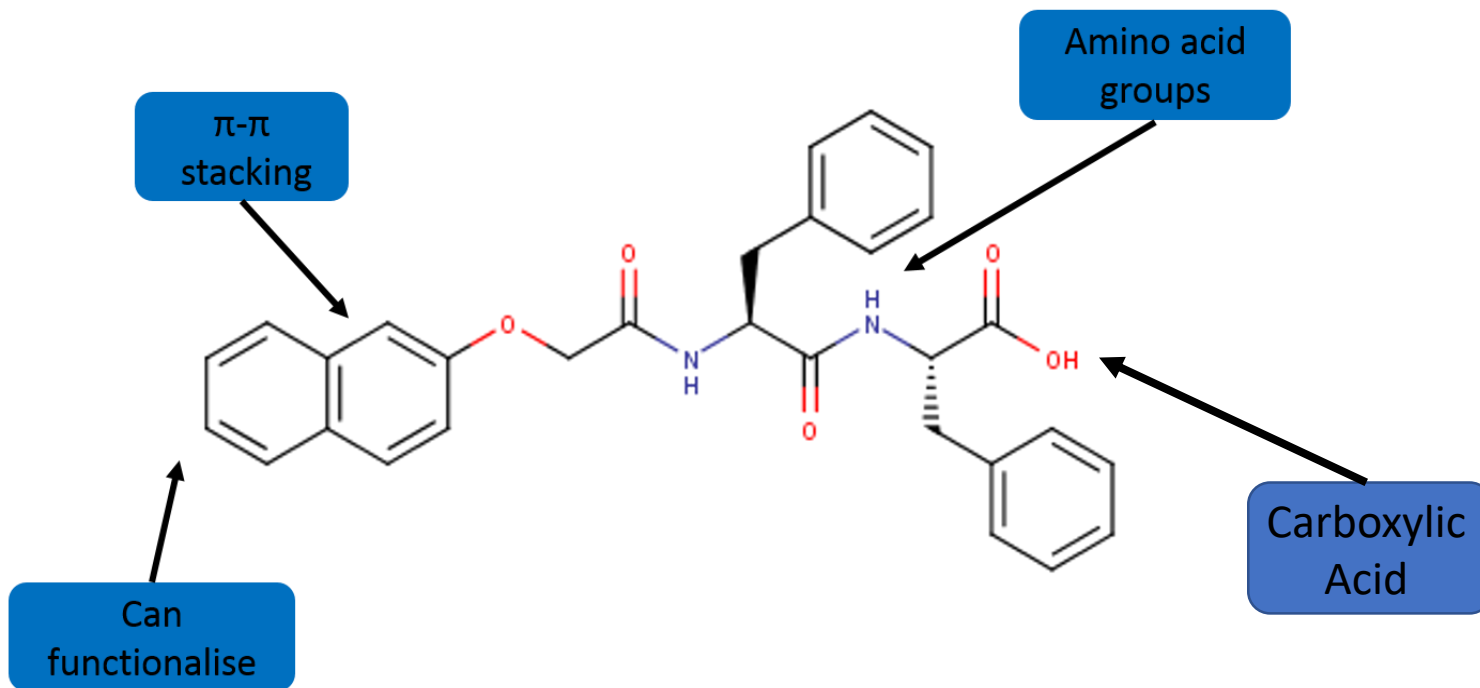


Self Assembled Soft Matter: Using SAXS

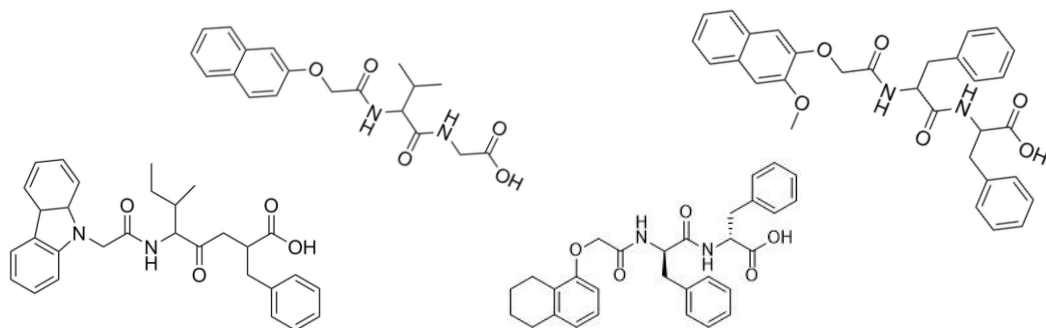
Kate McAulay, Ana Mari Fuentes Caparrós, Lisa Thomson and
Dave Adams.

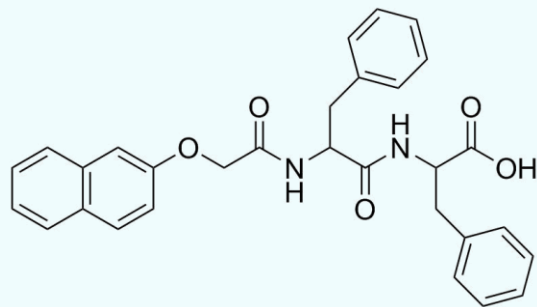




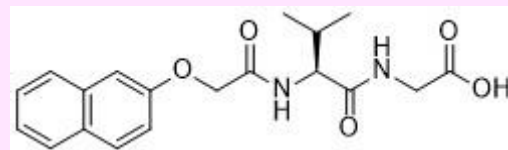
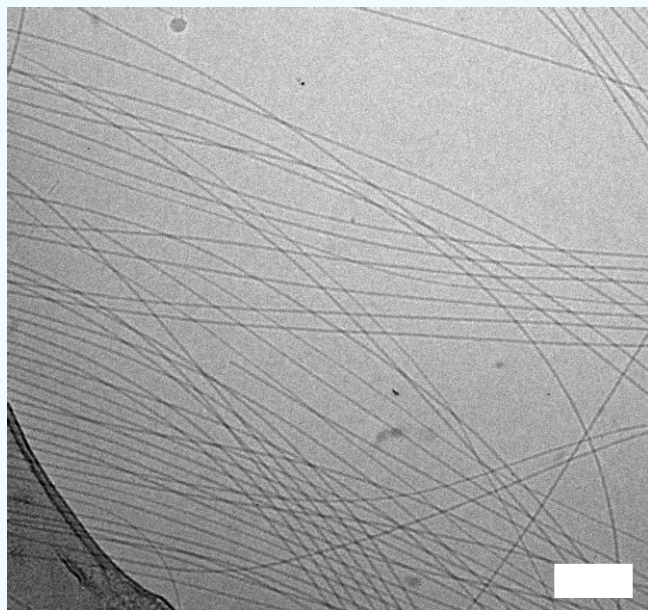


2NapFF Solutions at pH 11:

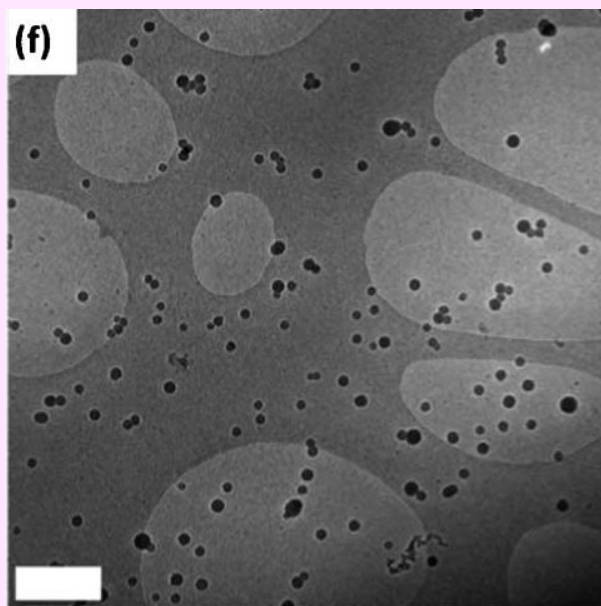


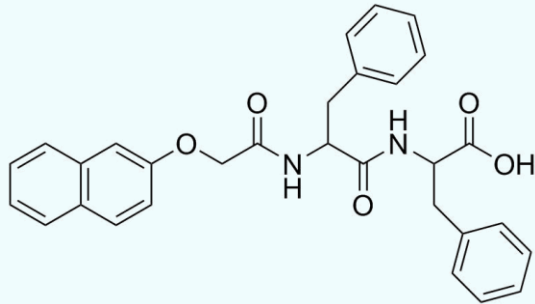


**Very hydrophobic = anisotropic,
persistent structures
(worm-like micelles)**



**Less hydrophobic = less
persistent structures
(spherical micelles/aggregates)**

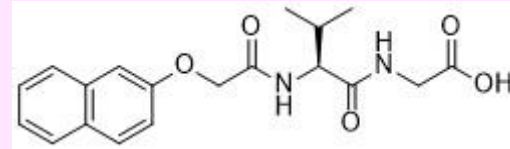




**Very hydrophobic = anisotropic,
persistent structures
(worm-like micelles)**

**SAXS/SANS: Highly scattering
at high pH**

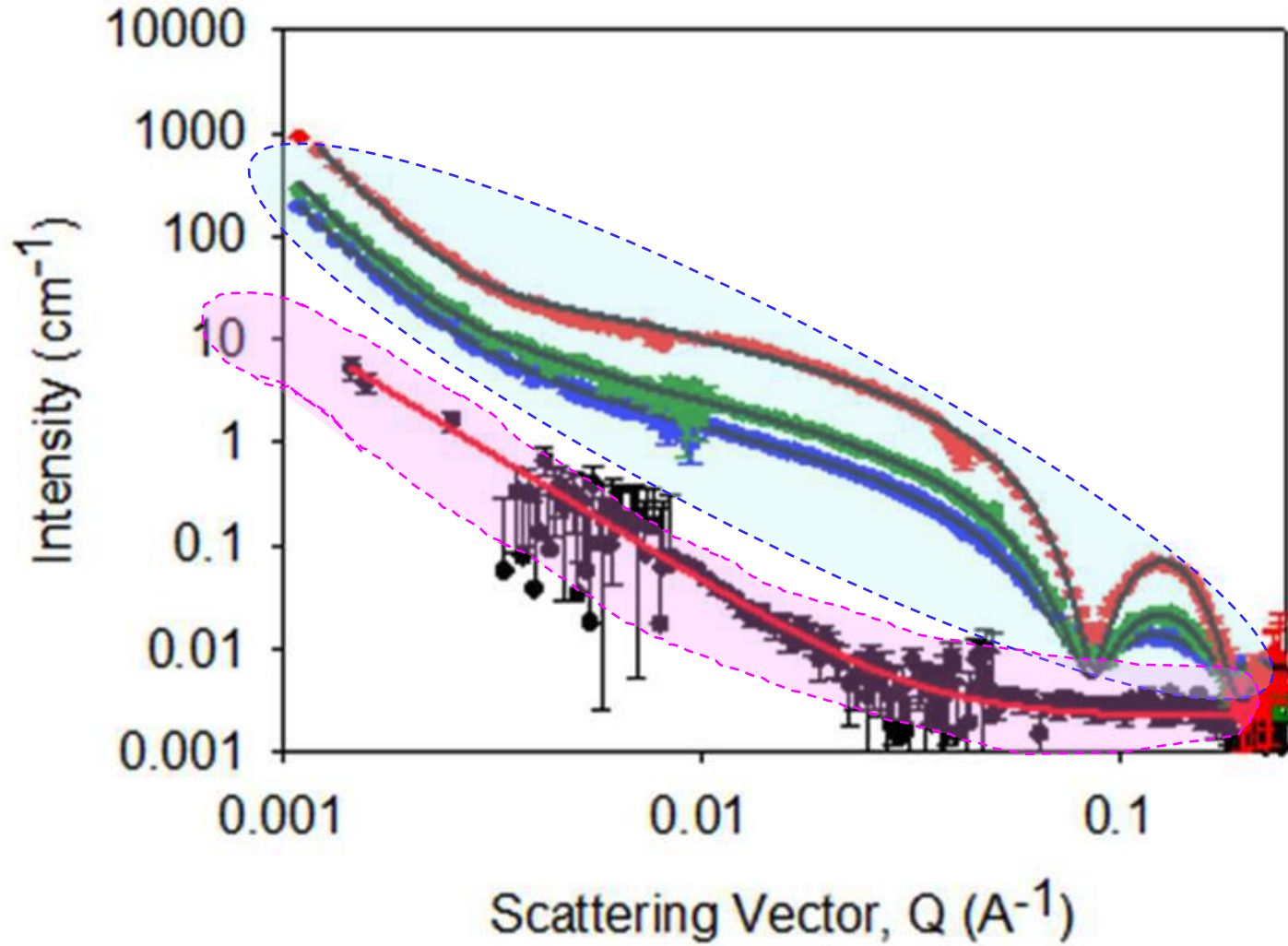
**Normally fit to flexible cylinder
or hollow cylinder**



**Less hydrophobic = less
persistent structures
(spherical micelles/aggregates)**

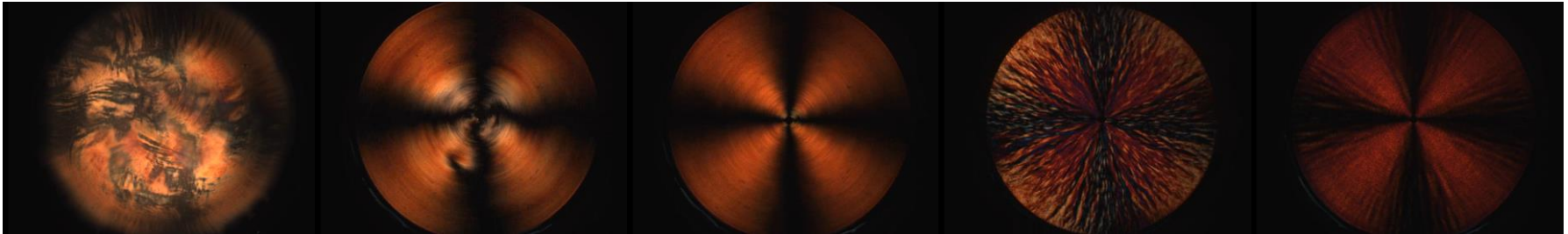
**SAXS/SANS: Weakly
scattering at high pH**

**Generally fit to a power law
only.**

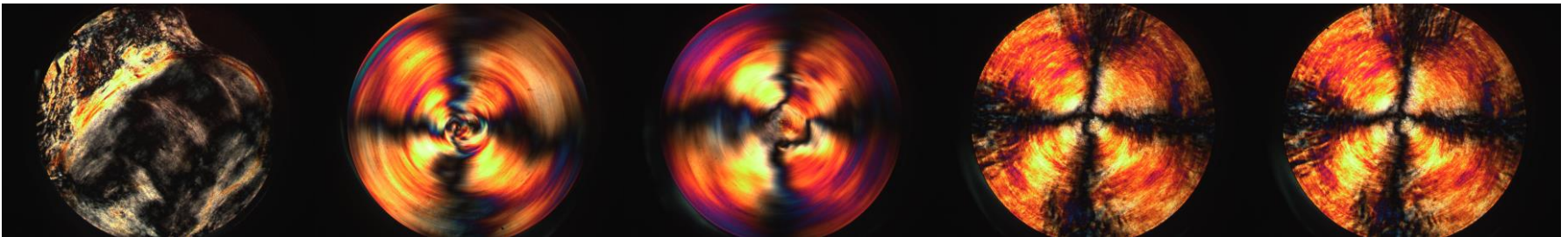


No Shear Under Shear After Shear
10 s 60 s 30 s 360 s

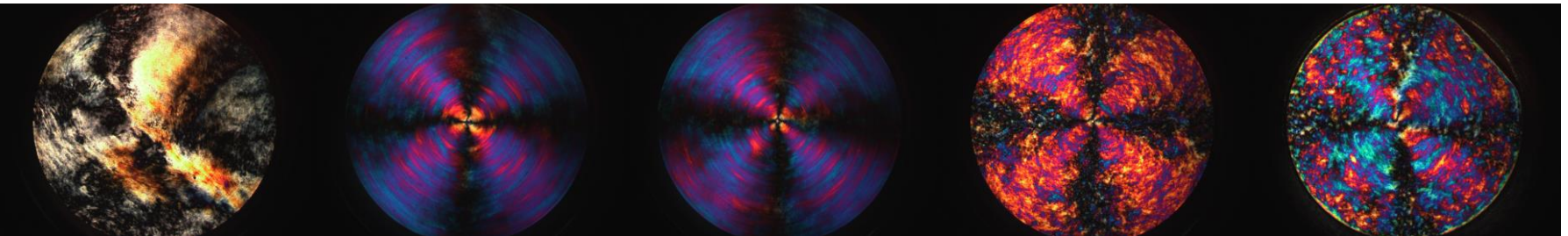
(1)



(2)



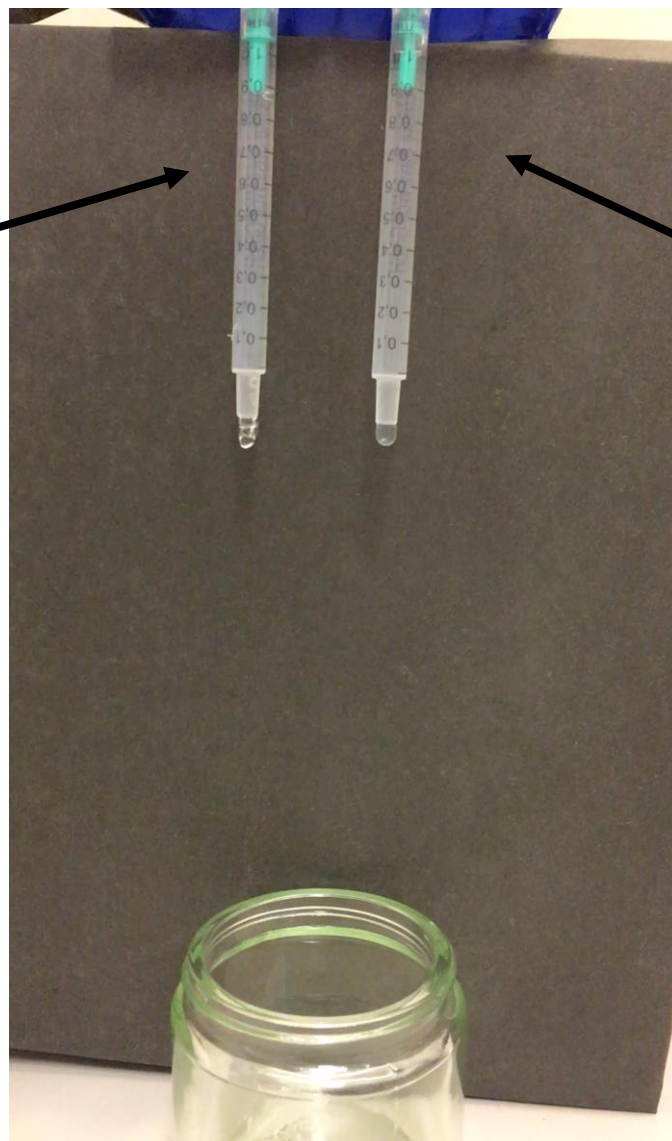
(3)



Can put the rheometer in the beam!

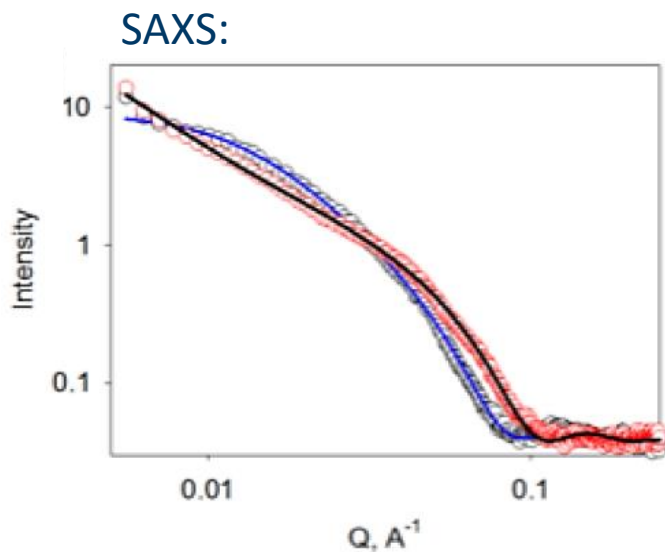
Before Heating

More viscous



After Heating

Less Viscous



Before Heating:

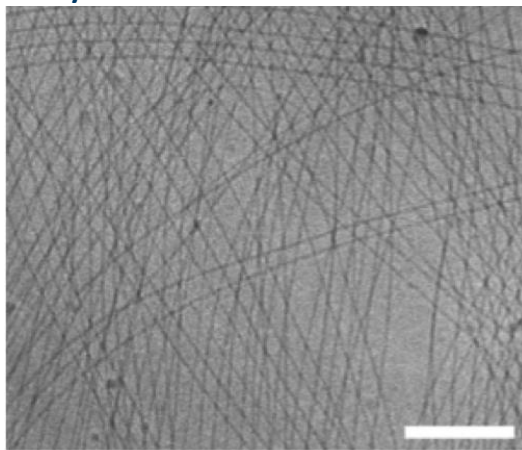
Flexible Cylinder

Radius $40.5 \pm 0.2 \text{ \AA}$

Kuhn length $63.0 \pm 2.4 \text{ \AA}$

Length $949.6 \pm 5.6 \text{ \AA}$

Cryo-TEM:



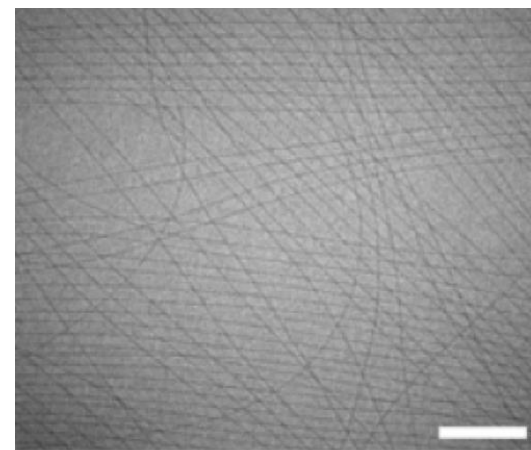
After Heating:

Flexible Cylinder

$34.0 \pm 0.1 \text{ \AA}$

Kuhn length of $339.7 \pm 3.4 \text{ \AA}$

Length of $4976.4 \pm 232.8 \text{ \AA}$

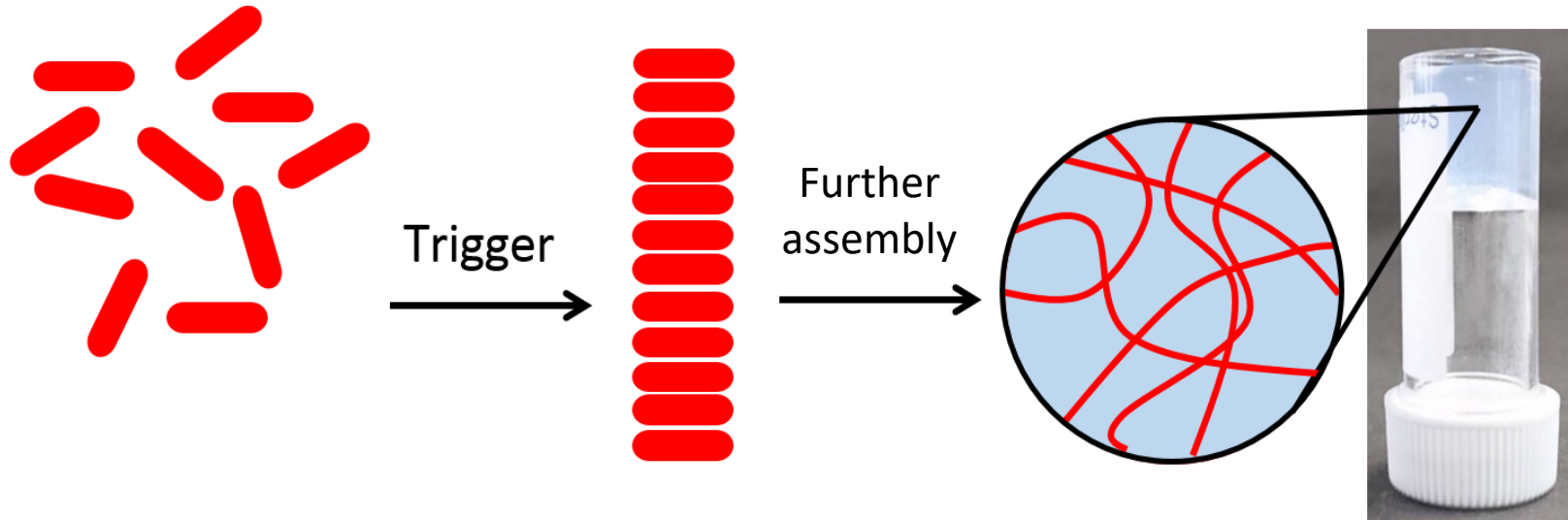


In both cases, the absolute length is beyond the resolution of the fit,

The length increases after the heat/cool cycle.

The radius decreases after the heat/cool cycle.

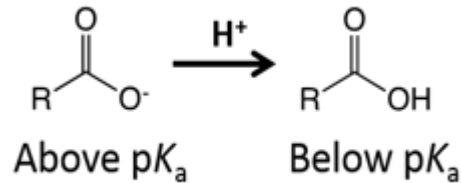
The Kuhn length increases after the heat/cool cycle so the flexibility decreases.



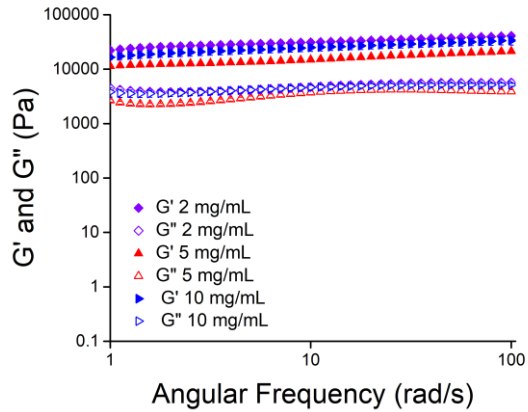
LMWG
in solution at
High pH

pKa = pH
Assembly into 1-D fibre
structures

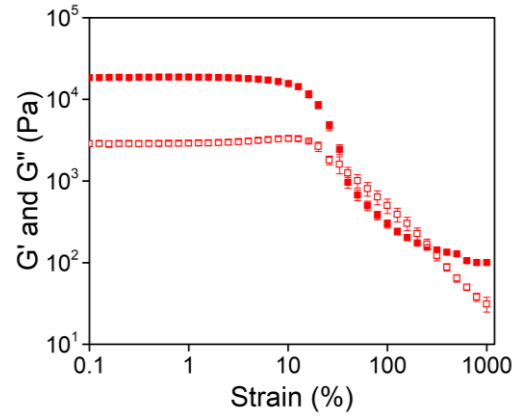
Fibre 3-D network
giving a self-supporting
gel



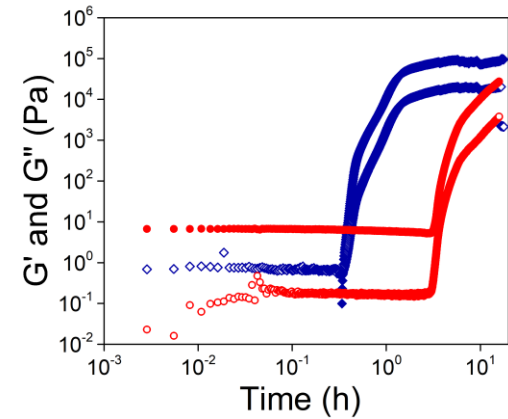
Frequency Sweep



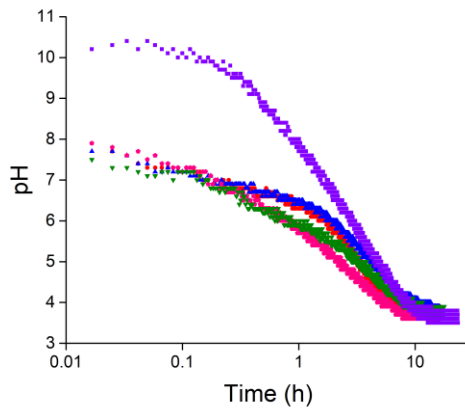
Strain Sweep:



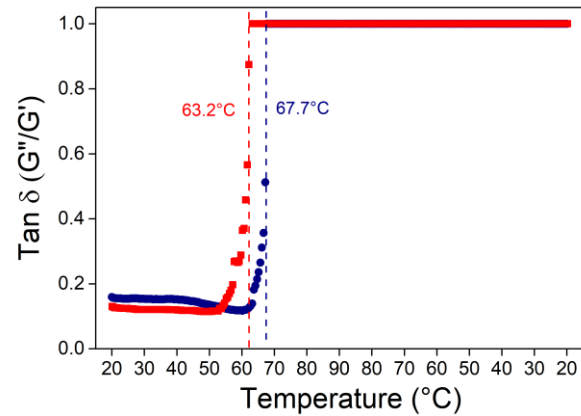
Time Sweep:



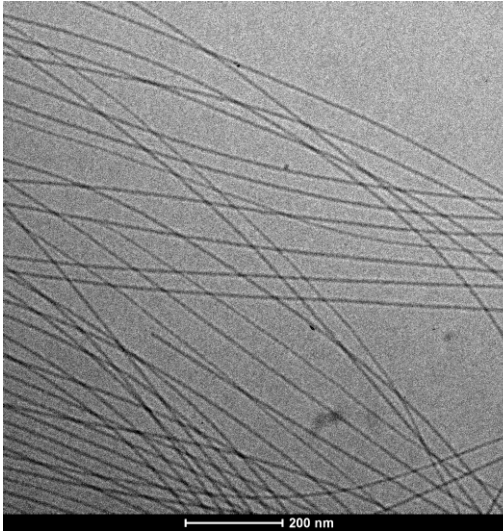
pH Measurements:



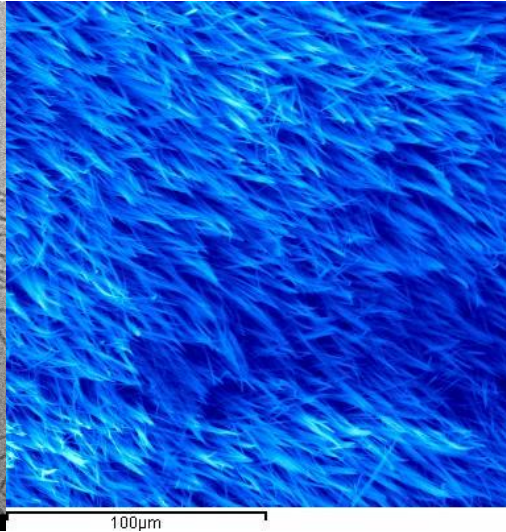
Heat/Cool:



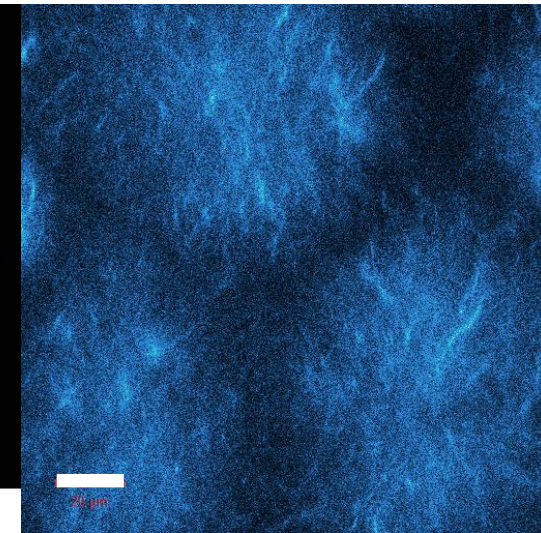
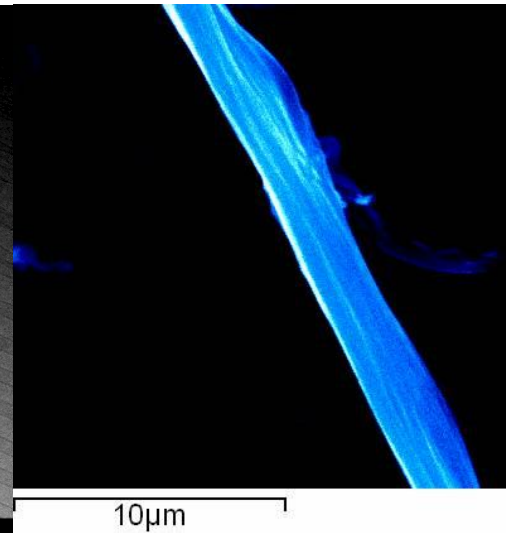
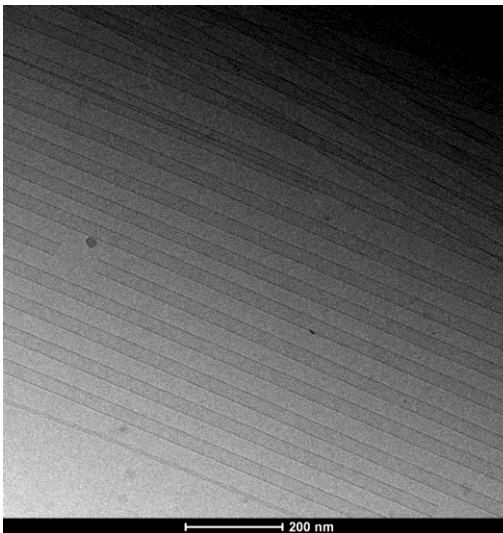
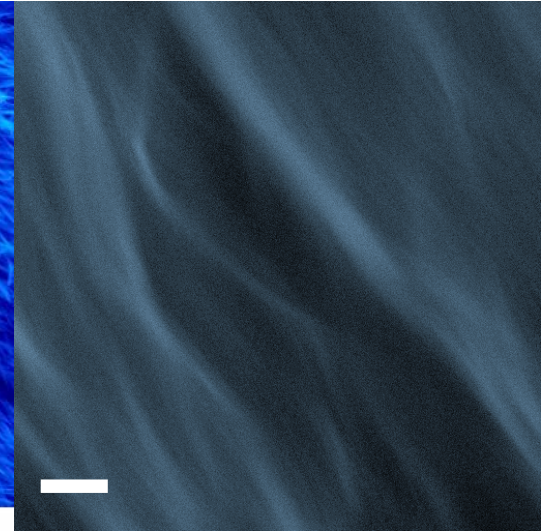
Cryo-TEM



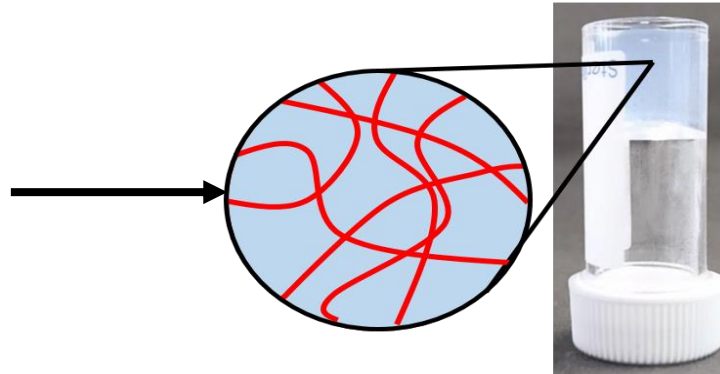
SEM



Confocal Microscope

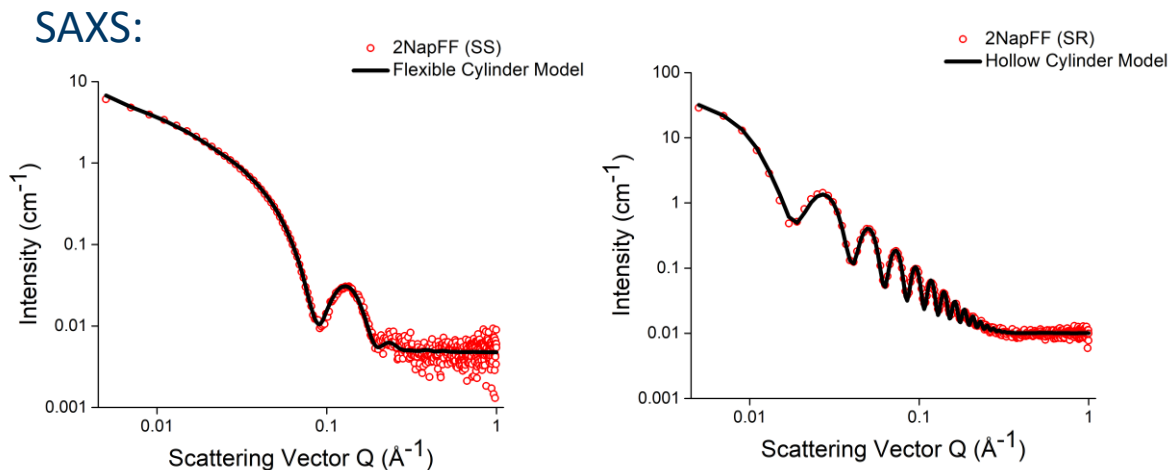


Small angle scattering
can be used to probe
the primary fibres.

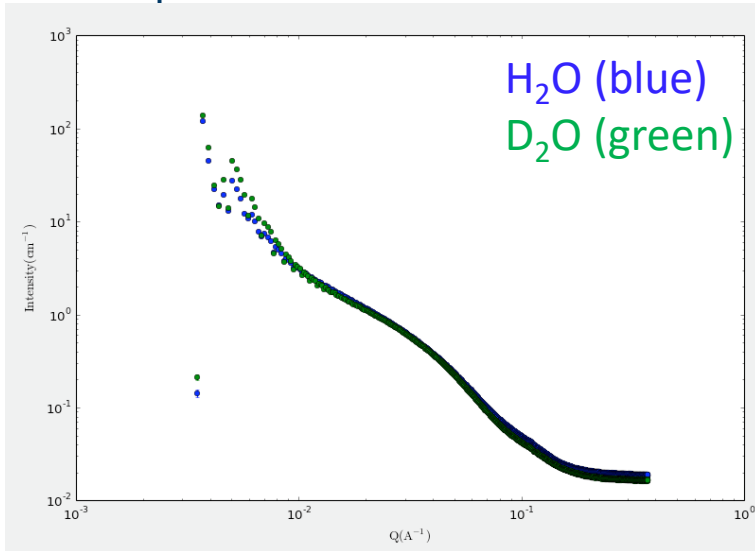


The fit provides information as to the radius of the structure, the length, persistence, Kuhn length *etc.*

LMWGs, the best fits usually are best to a cylinder, a flexible cylinder, a hollow cylinder, or some other long, anisotropic structure.



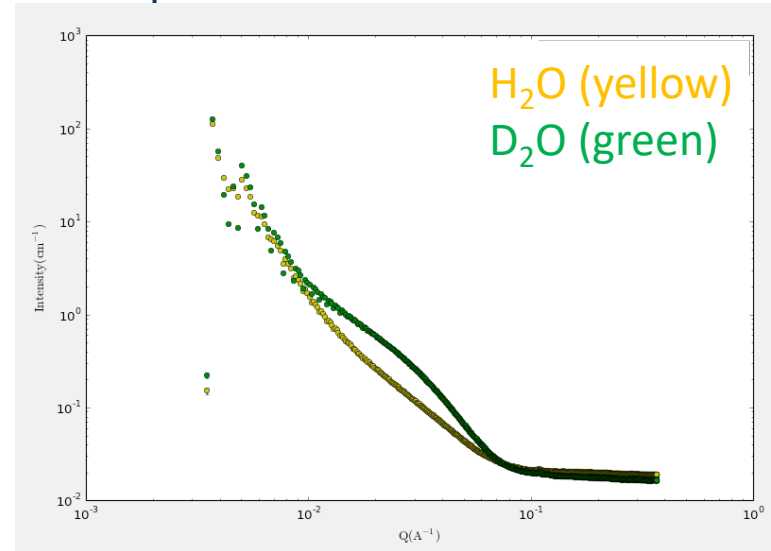
BrNapAG



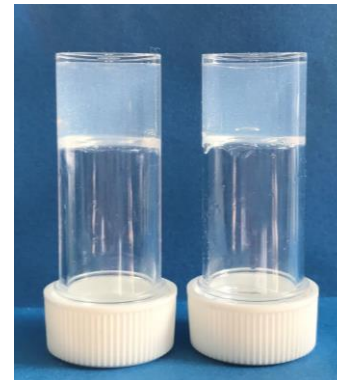
Very similar scattering.

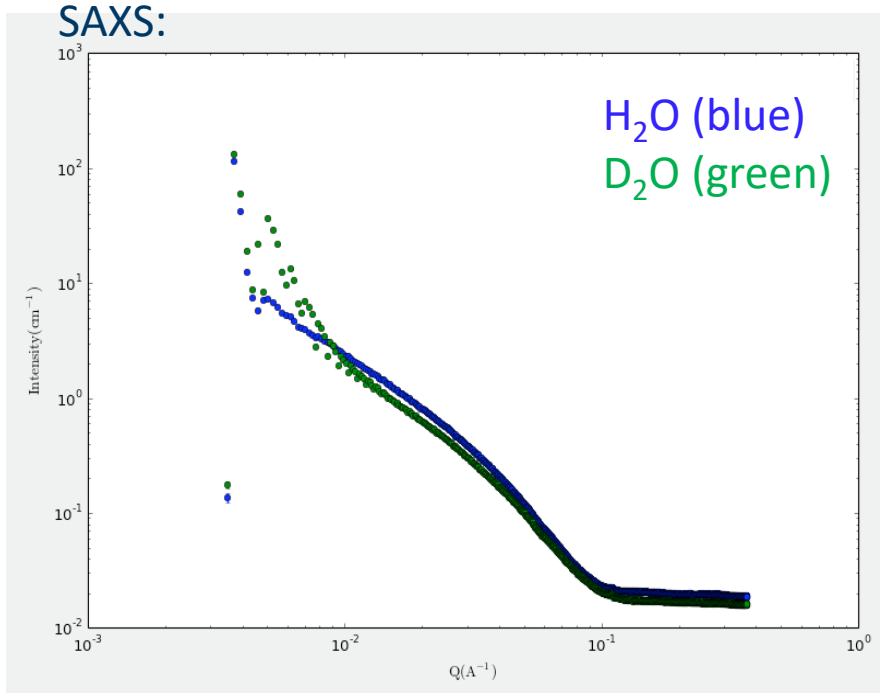
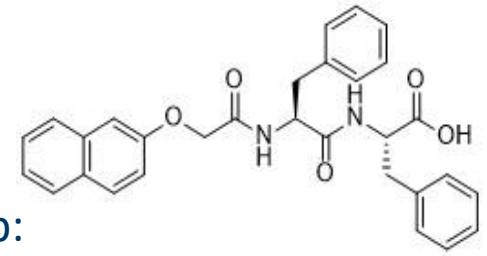


BrNap1F

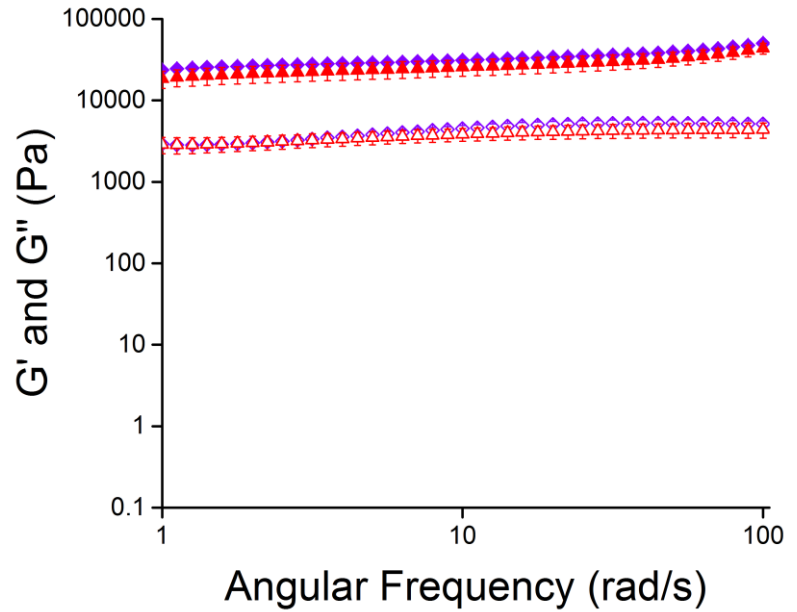


Differences in scattering.



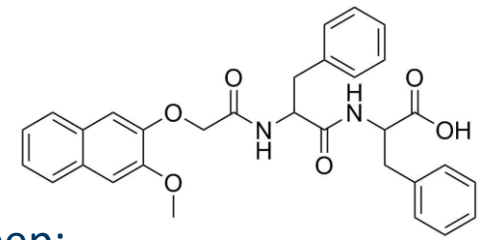


Frequency Sweep:

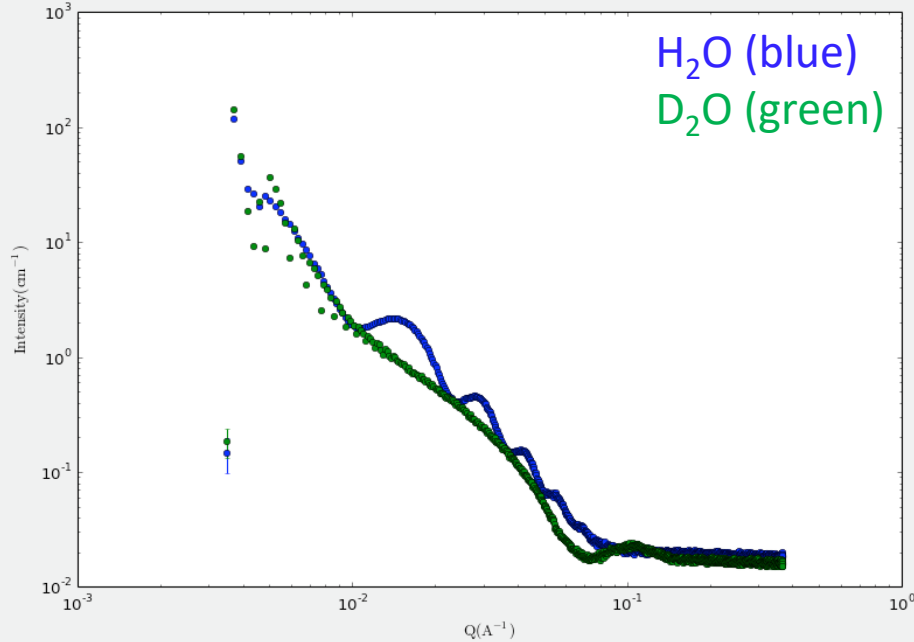


H_2O (purple) and D_2O (Red)
closed symbols G' and open G''

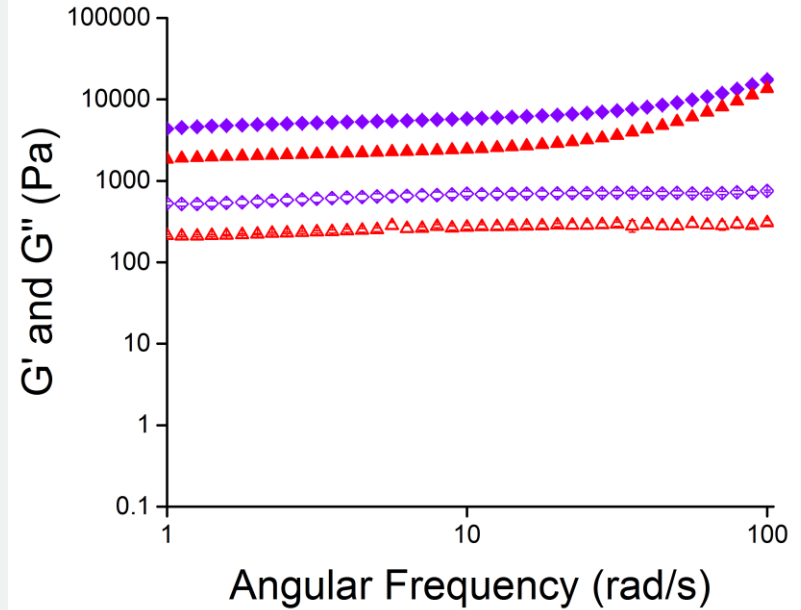
- Very similar in general as expected from previous data.



SAXS:

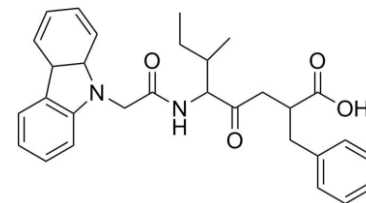


Frequency Sweep:

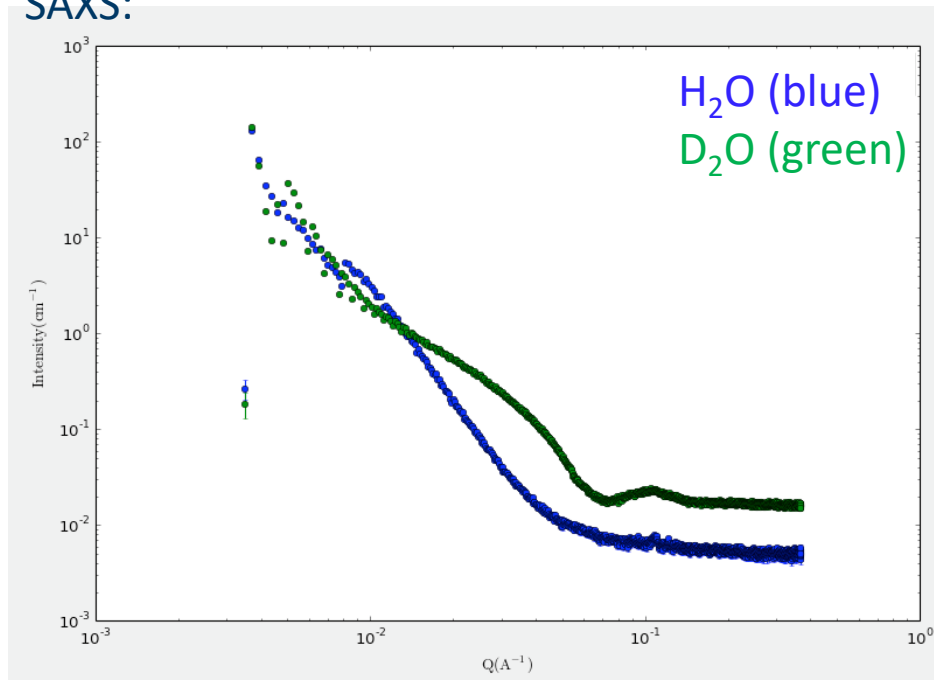


H₂O (purple) and D₂O (Red)
closed symbols G' and open G''

- Different structures AND different rheology data!
- H₂O fits to a hollow cylinder with a radius of around 3.6 nm and a thickness of 1.6 nm
- D₂O fits to a hollow cylinder with a radius of around 21 nm and a thickness of 7.6 nm



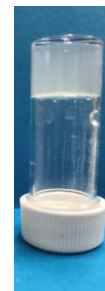
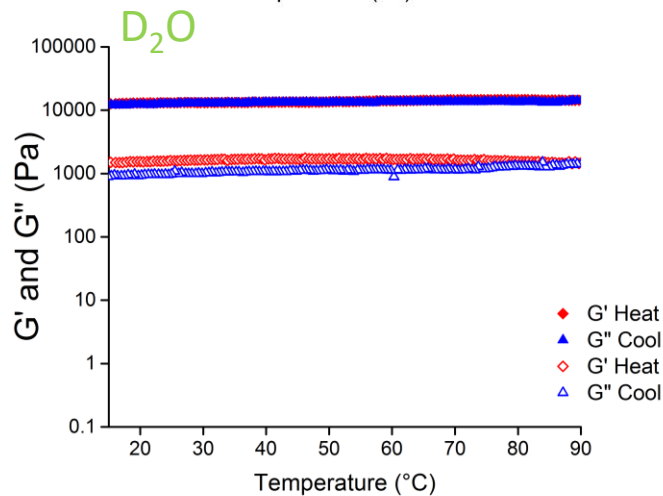
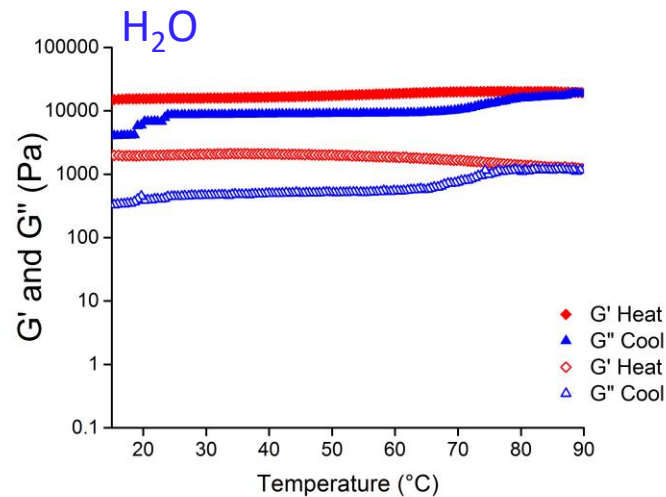
SAXS:

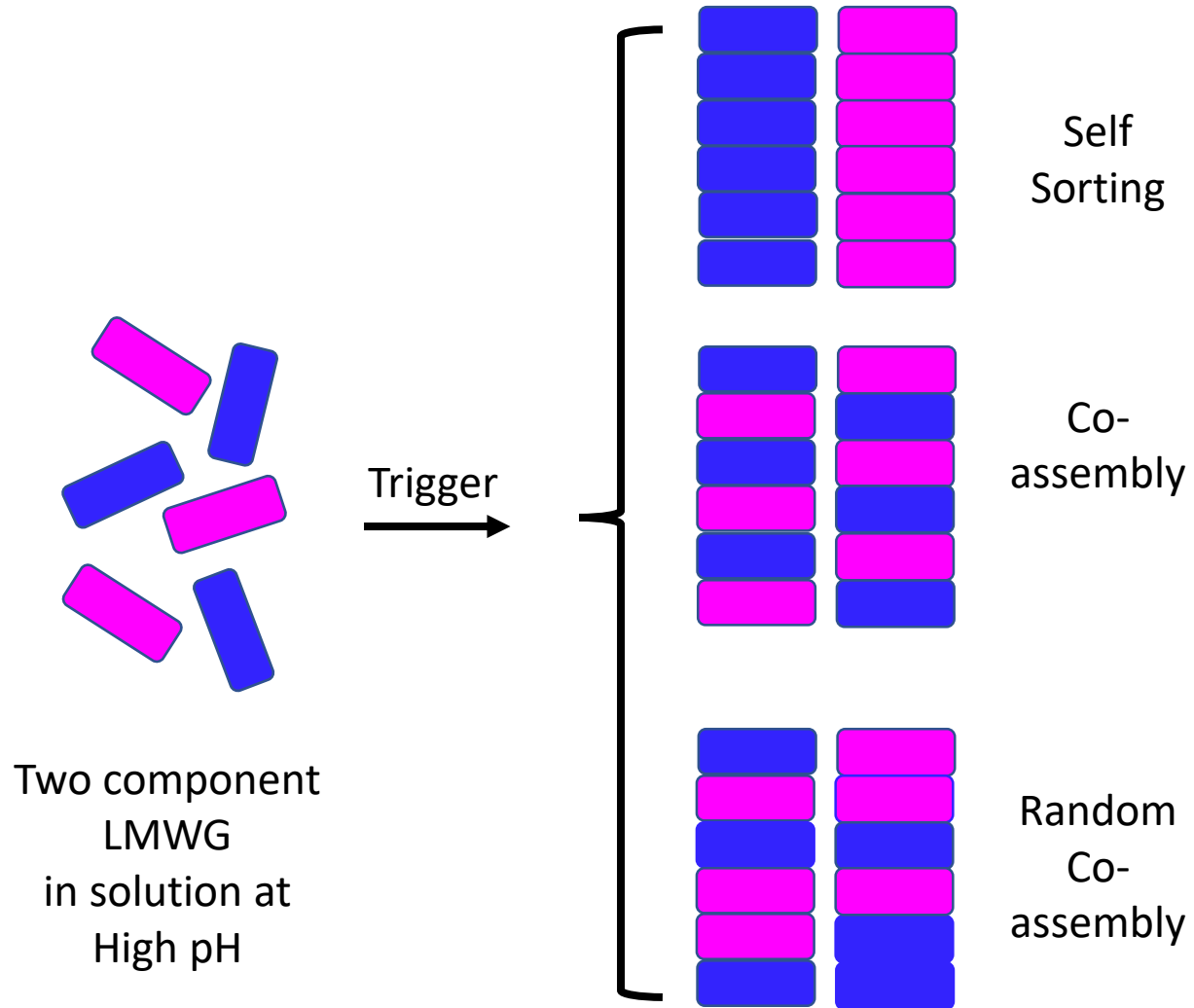


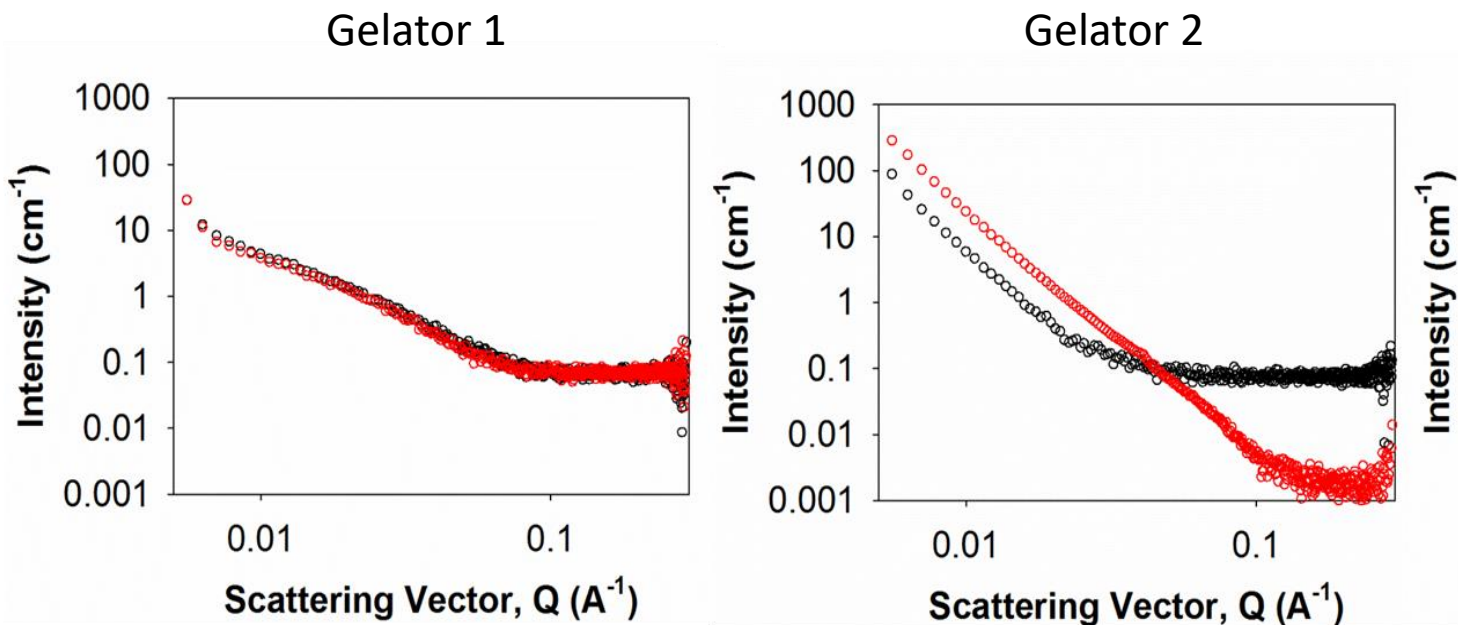
Significant difference!

- H_2O fits to a polydisperse flexible cylinder with a radius of around 16 nm.
- D_2O fits to a hollow cylinder with a radius of around 3 nm and a thickness of 1 nm.

Temperature Sweep:



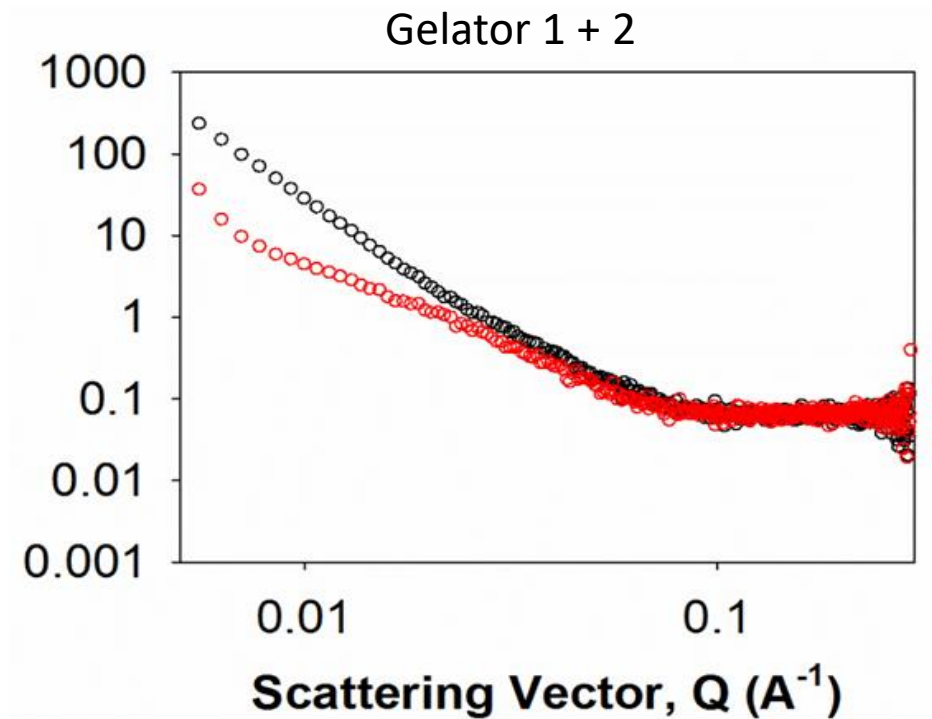




SAXS data before (black) and after (red) annealing.

For **Gelator 1** the data fits to a flexible cylinder model with a radius of 4.3 ± 0.07 nm.

The baseline for **Gelator 2** drops after annealing as there is crystallisation and a small amount of precipitation, meaning that there is less sample in the beam after annealing.



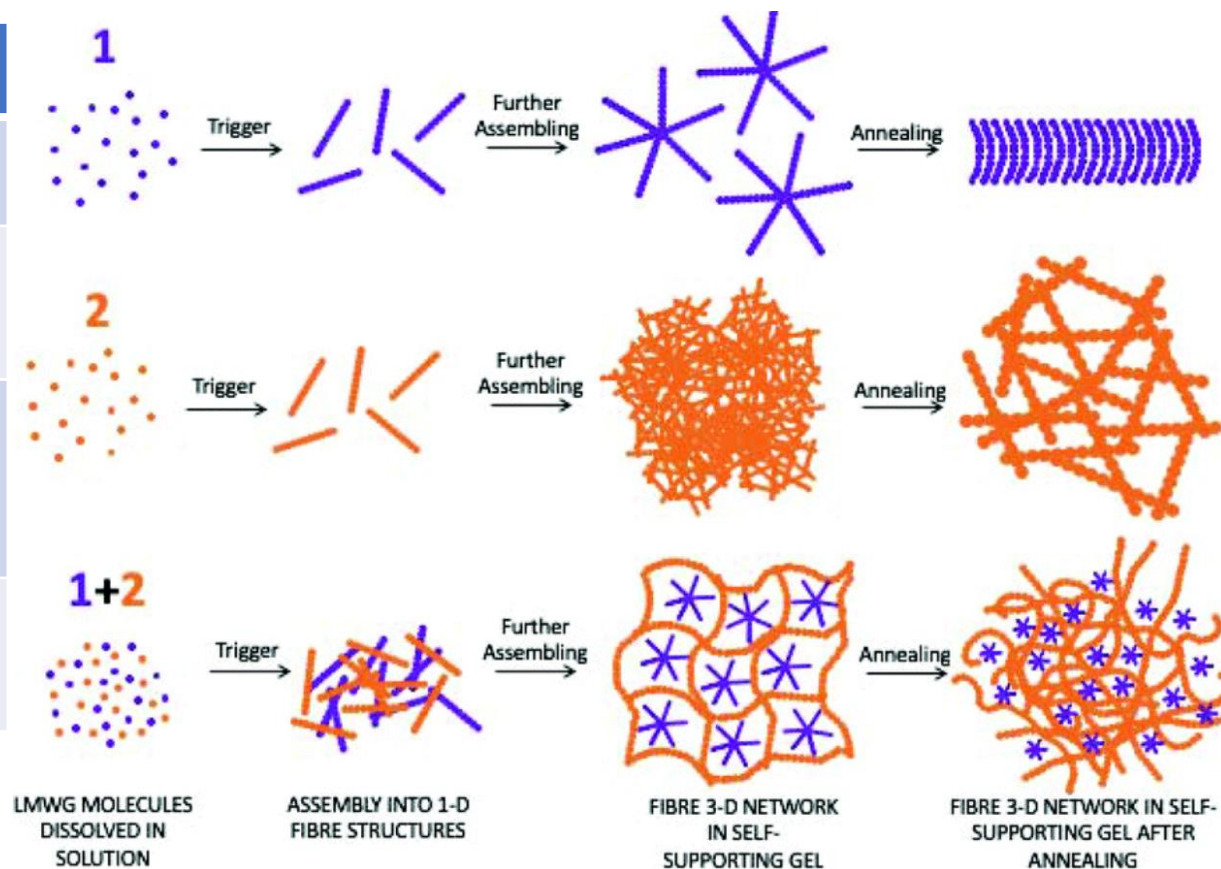
SAXS data
Before (black)
After (red)
annealing.

For the multicomponent system, the SAXS data is different before and after annealing.

Before heating, the data are reminiscent of that of **2**

After annealing, the data are very similar to that of **1**

| Gelator | Gradient | Description |
|--------------------|----------|--|
| 1 before and after | -2 | mass fractal network |
| 2 before | -4 | objects too large to be resolved by SAXS |
| 1+2 Before | -3 | mass fractal with a tight interconnected structure |
| 1+2 After | -2 | mass fractal network |



Conclusions

- We are using SAXS to look at the self-assembled LMWG structures in solutions and gels.
- Can investigate structural information without modifying the samples.
- Don't need to use D₂O.
- Relate these structures to the physical and mechanical properties of the samples.

- Professor Dave Adams
- Ana Mari Fuentes Caparrós
- Lisa Thomson

- Adams Group
- Bart Dietrich

- Annela Seddon

- Hao Su
- Honggang Cui

- Nikul Khunti
- Nathan Cowieson

