

Data Analysis of Hierarchical Systems

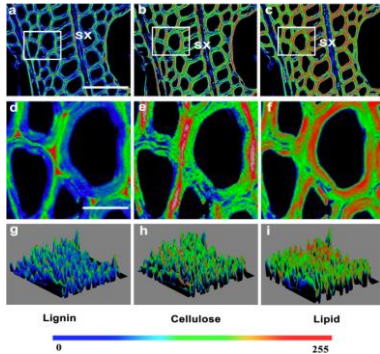
Sai Venkatesh Pingali

pingalis@ornl.gov

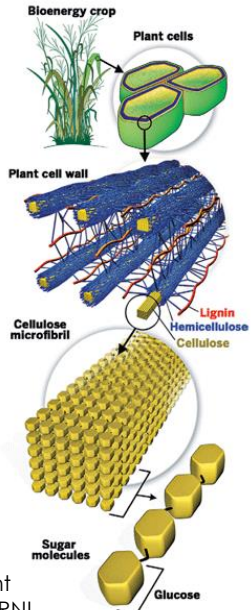
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Example of Hierarchical Systems Around us

Plant Cell Wall

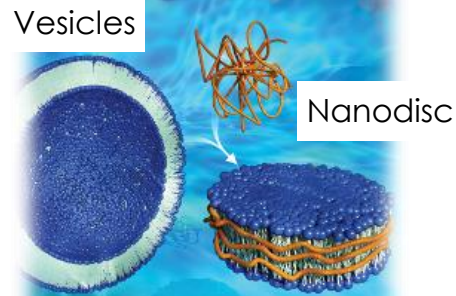


Plant Methods **17**, 29 (2021).

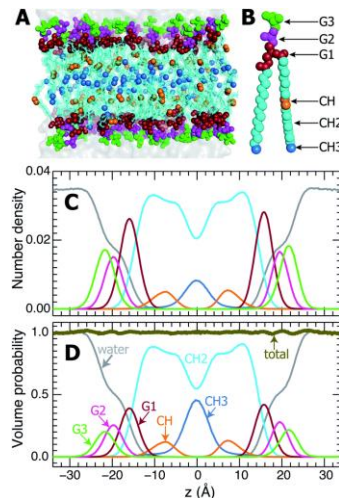


Genome Management
Information Systems/ORNL

Lipid Systems

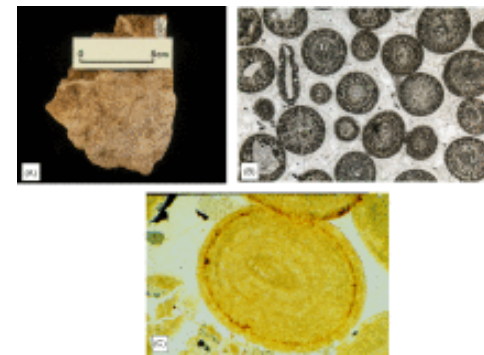
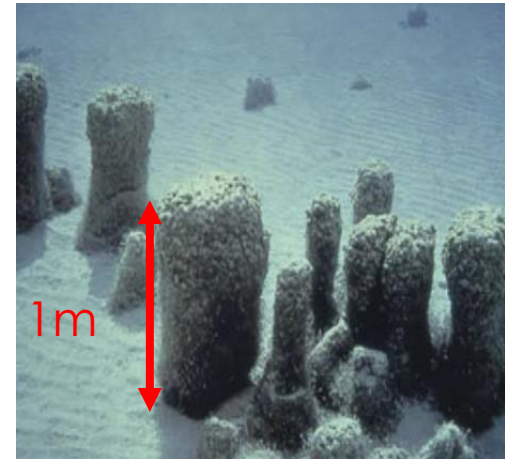


ACS Appl. Bio Mat. **4**, 4760–4768 (2021)



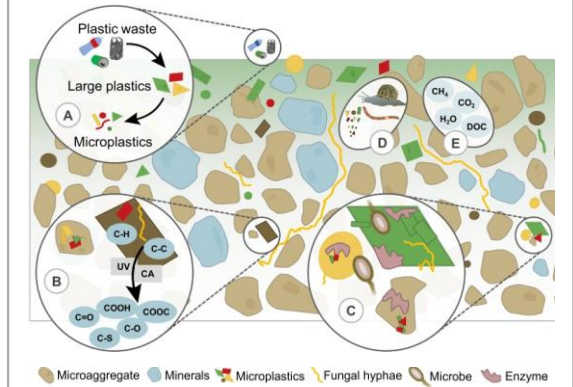
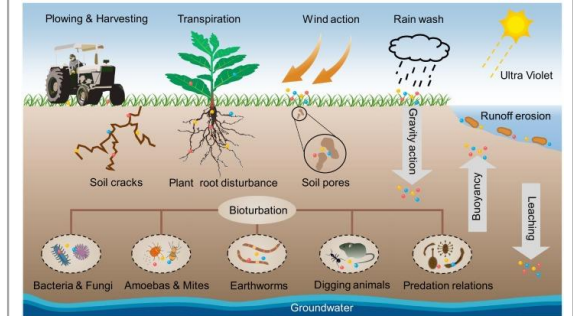
Soft Matter **10**, 3716–3725 (2014)

Sedimentary Rocks



Encyclopedia of Geology 382-394 (2021)

Plastic/Soil Composites

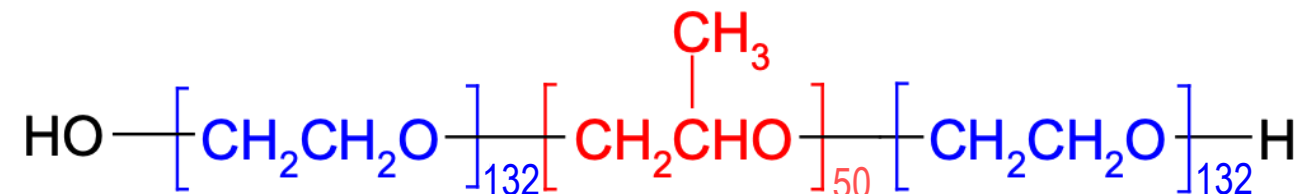


Applied Soil Ecology **176**, 104486 (2022)

Data Analysis Approaches Available

- Model Fit –
Triblock Copolymer Structures in different Solvent Conditions
 - Debye Coil
 - Solid Sphere
 - Triblock Copolymer Micelle StructureLipid Vesicles
 - Core-shell (or multi-shell)
- Shape-independent Fit - Guinier/power-law)
Plant Cell Wall (powder sample state)
 - Unified Fit (Combination of Guinier and power-law functions)
- Composite Fit (A combination of Models and Shape-independent Approaches)
Plant Cell Wall (Aligned state)
 - Cylinder Form Factor combined with Guinier and/or power-law functions

Amphiphilic Polymers - Drug Delivery

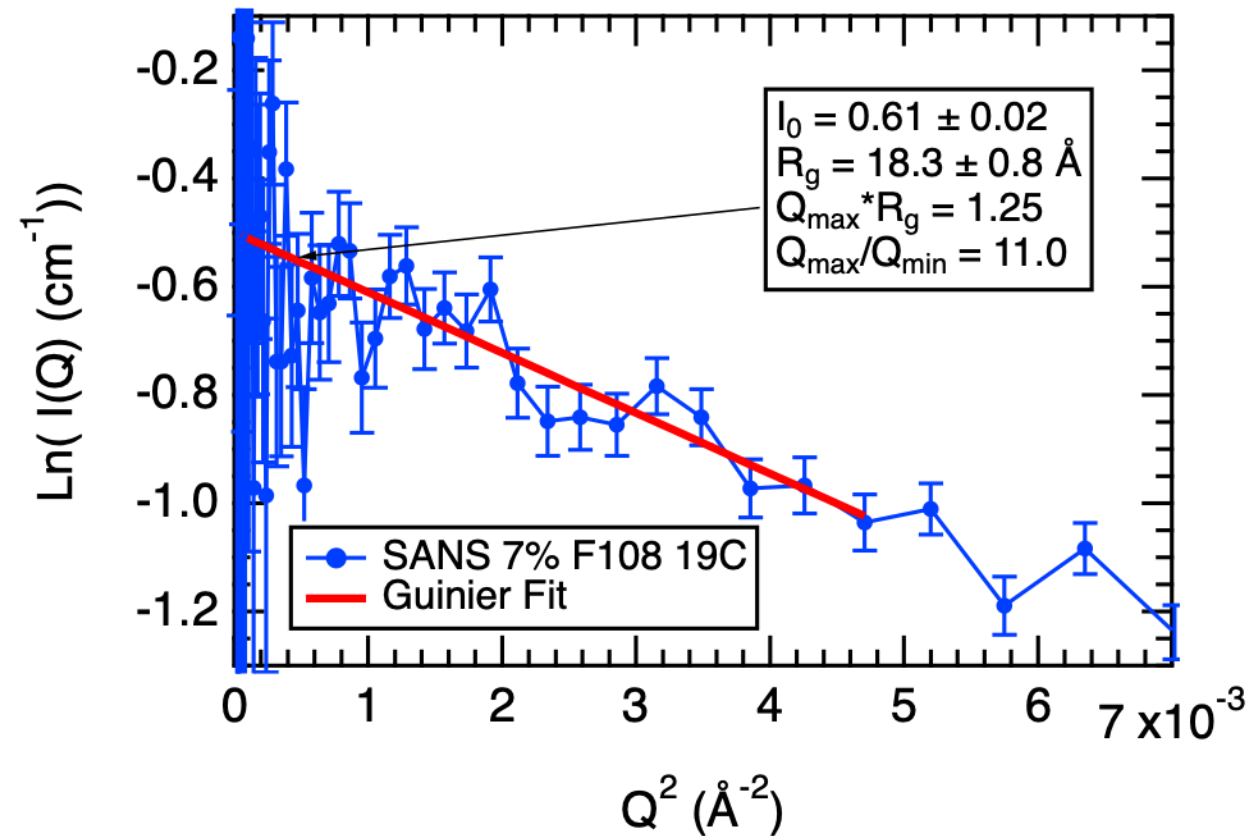
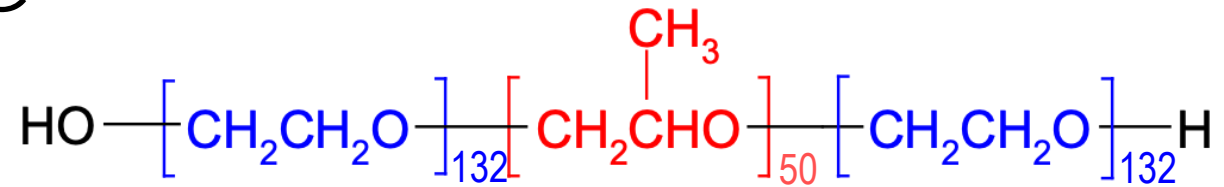
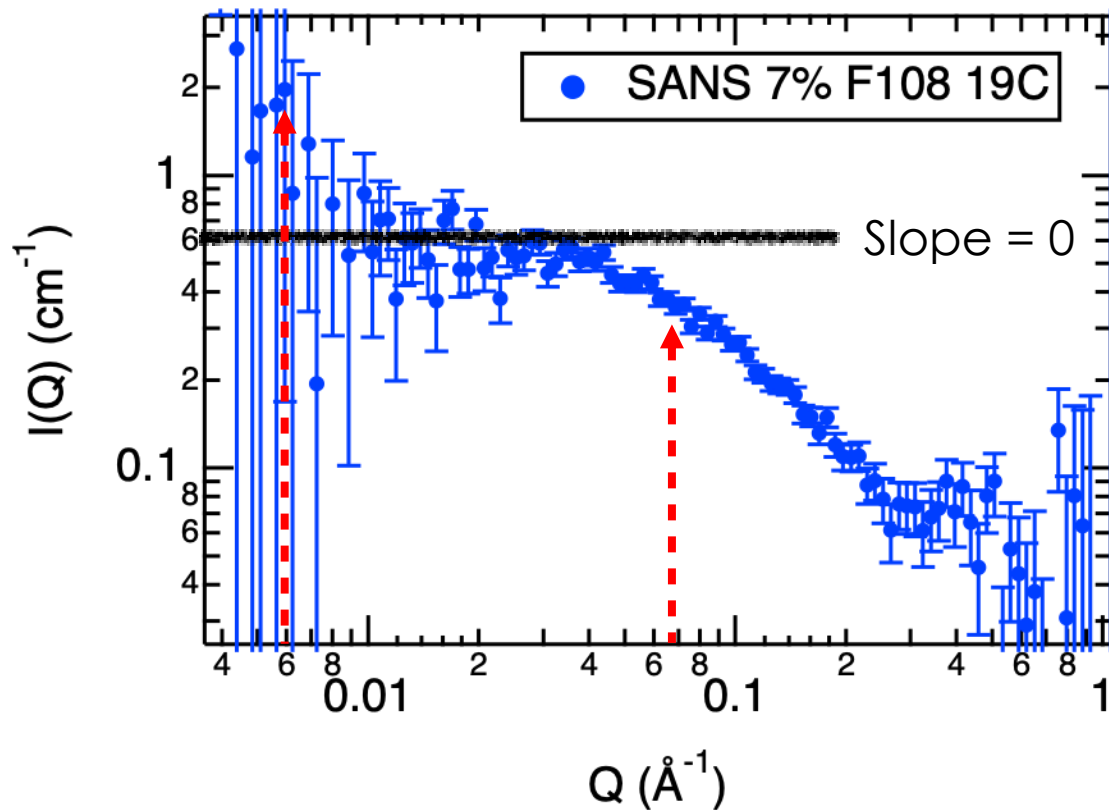


Pluronic F108

Hydrophilic
Hydrophobic

- Pluronic F108 7% solution in 0.1M sodium carbonate
- Molecular Weight (Mw) is 16400 Da
- Structure at room temperature (19 °C)

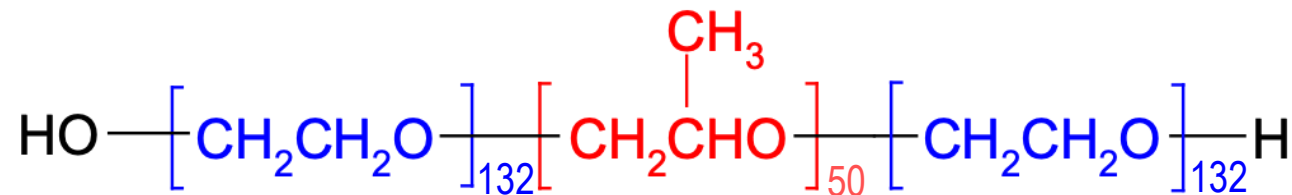
Guinier Analysis 7% F108 at 19 °C



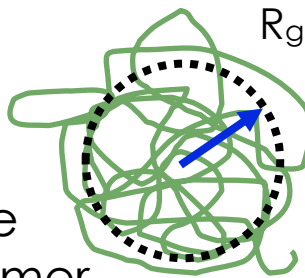
- Guinier analysis results

- Globular particle and $R_g \sim 18.3 \text{ \AA}$
- $Q_{\text{max}} * R_g < 1.3$
- Q-range of straight line fit, $Q_{\text{max}}/Q_{\text{min}} \sim 2.0+$

Model Fits



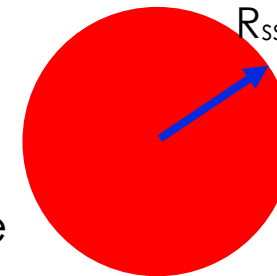
Debye Flexible
Gaussian Polymer



$$I(Q, R_g) = SF \frac{2[\exp(-z) - 1 + z]}{z^2} + I_{bkg}$$

$$z = Q^2 R_g^2$$

Solid Sphere



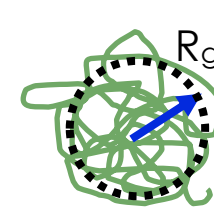
$$I(Q, R_{ss}) = SF \left(\frac{3[\sin(z) - z\cos(z)]}{z^3} \right)^2 + I_{bkg}$$

$$z = QR_{ss}$$

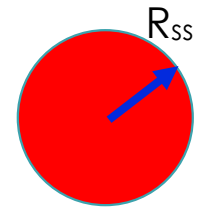
- Models

- Debye Flexible Gaussian Polymer (random conformations)
- Solid Sphere (when low degree of solvent penetrates the particle)

Debye Flexible Gaussian Polymer vs. Solid Sphere

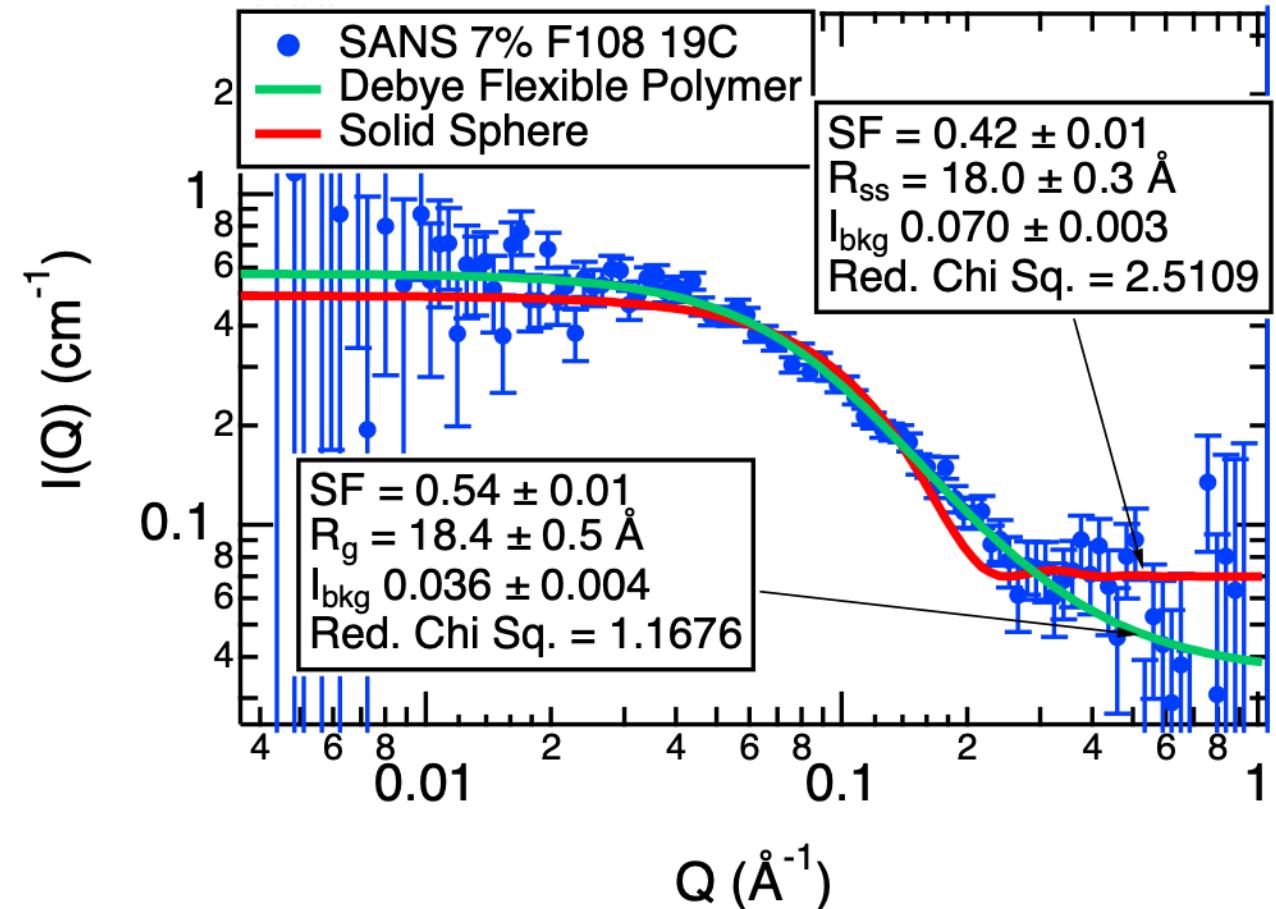


Debye Flexible
Gaussian Polymer

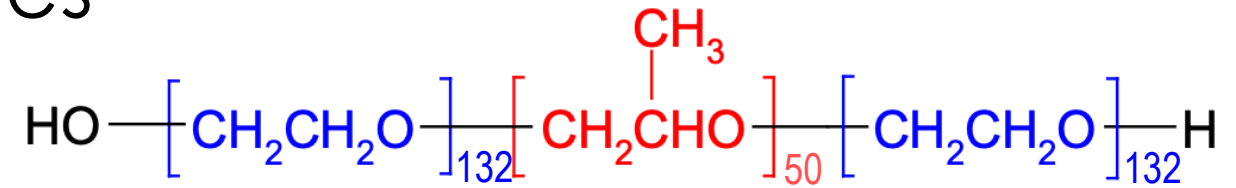


Solid Sphere

- F108 dissolved as individual polymers in solution of sodium carbonate
- High-Q region is highly sensitive to the surface morphology of the particle.
- **Debye flexible Gaussian polymer** function (High-Q PL exponent = 2) fits better than **solid sphere** (High-Q PL exponent = 4)

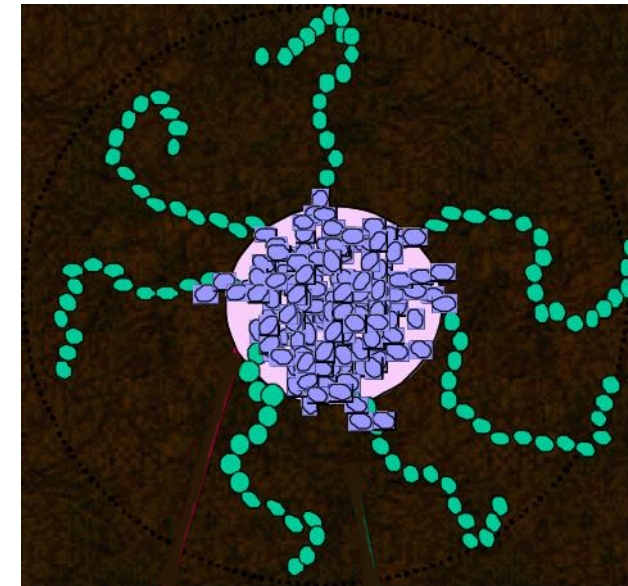


Self-Assembled Suprastructures

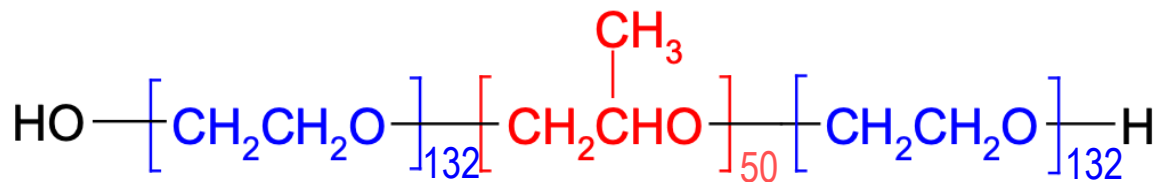


- Central core consists of the hydrophobic block of F108
- The tentacles are the hydrophilic blocks of F108
- Particle model
 - Spherical particle core for hydrophobic block
 - Debye Flexible Gaussian Polymer chain for hydrophilic blocks.
 - Interaction term- chain-chain and chain-Core
- Structure factor
 - Percus-Yevick structure factor (hard sphere approximation to enable developing an analytical expression)

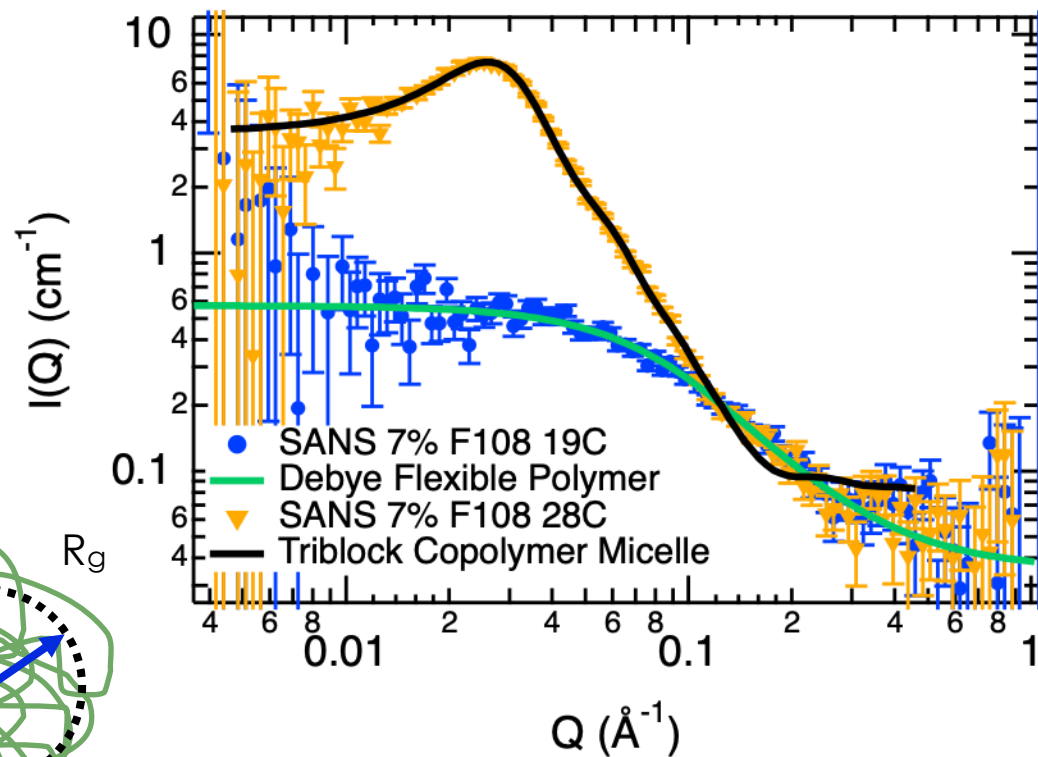
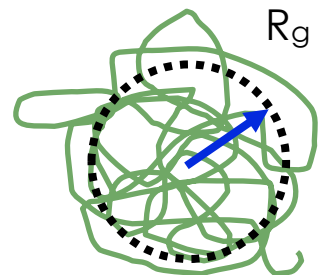
Hydrophilic - PEO corona
Hydrophobic - PPO core



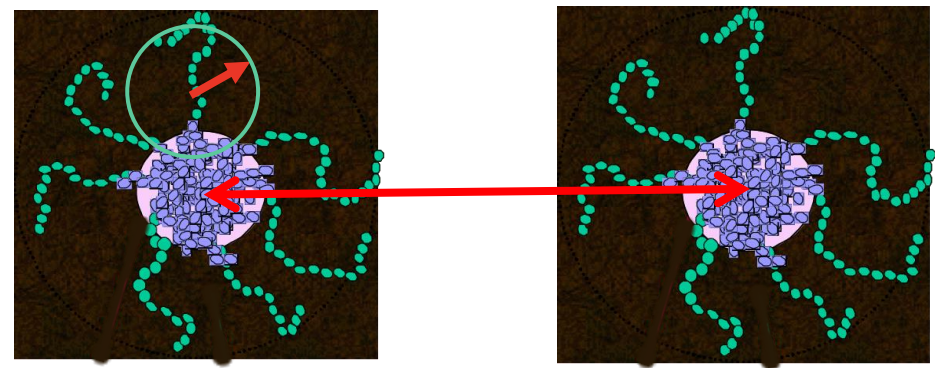
SANS data fit



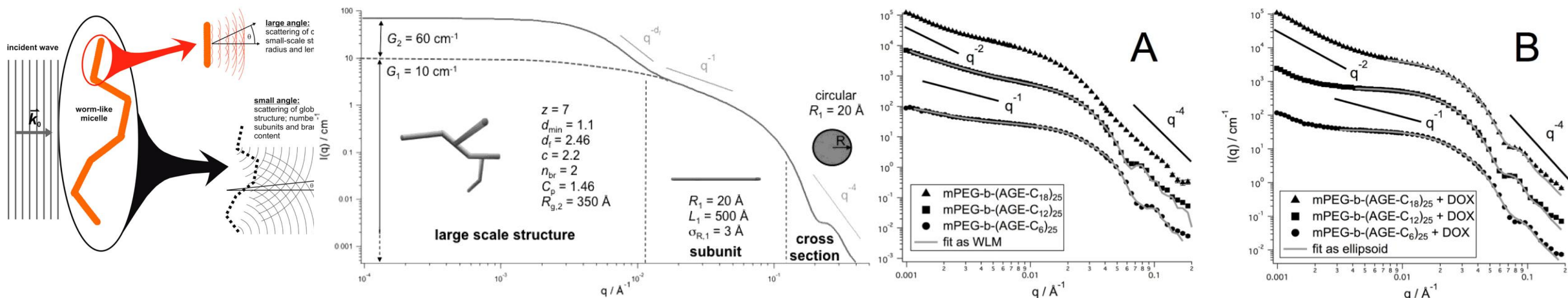
- Debye Flexible Gaussian Polymer Model
 - R_g (Å) 18.4 ± 0.5
- Triblock Copolymer Micelle Model
 - Aggregation (#) 13.1 ± 0.2
 - R_{core} (Å) 18.8 ± 0.3
 - R_{corona} (Å) 25.9 ± 0.6
 - Core Overlap Factor, d (#) 0.67 ± 0.02
 - Volume Fraction (#) 0.173 ± 0.003
 - Micelle-Micelle distance (Å) 209 ± 3
 - Particle diameter (Å) 72 ± 2
 - = $2 \times [R_{\text{core}} + (d \times R_{\text{corona}})]$



Hydrophilic - PEO corona
Hydrophobic - PPO core



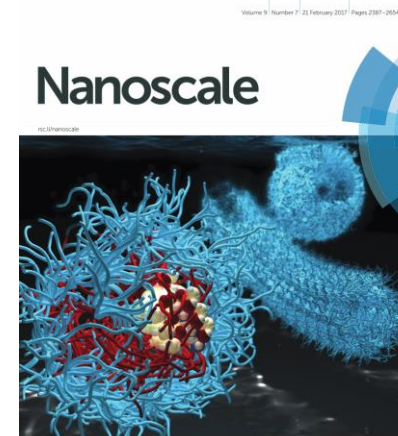
Drug release from Worm-like Micelles



- Amphiphilic polymers as drug delivery for Cancer drug-DOX
- Worm like Micelles Structure is hierarchical
- Worm like Micelles transitions to Ellipsoidal Structure impeding efficient drug delivery

Le Dévédec, F., Her, S., Vogtt, K., Won, A., Li, X., Beaucage, G., Allen, C. **Nanoscale (2017) 9, 2417-2423**

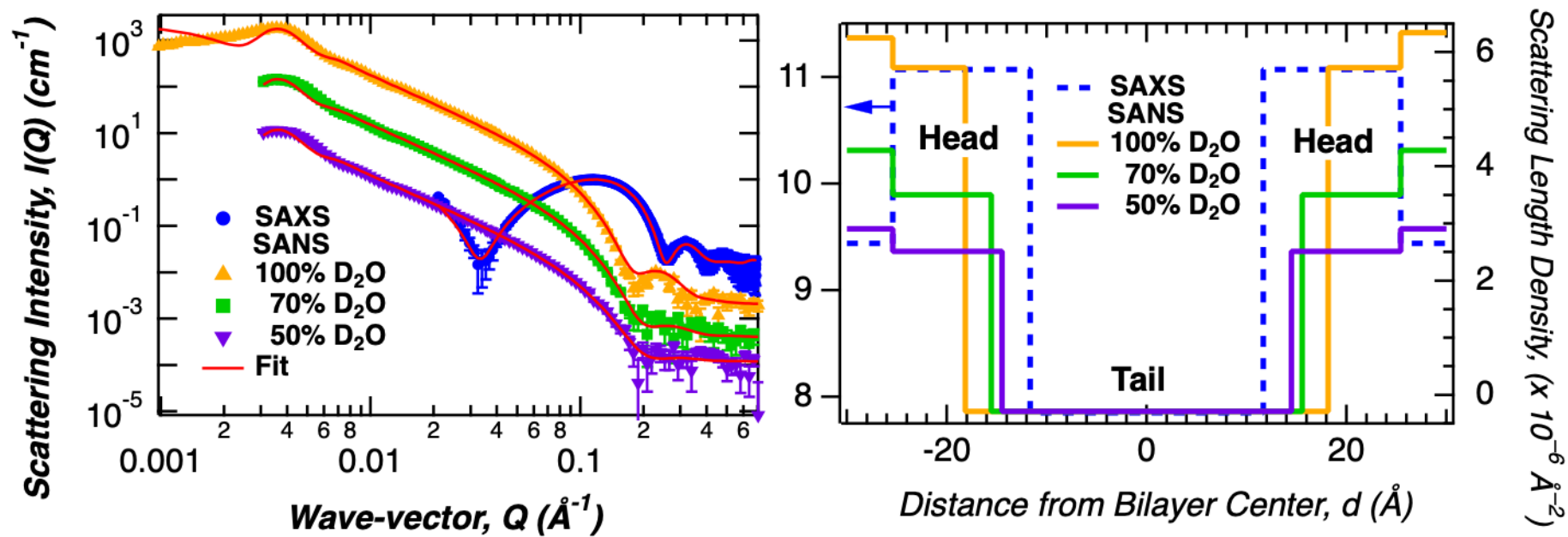
Vogtt, K., Beaucage, G., Weaver, M., Jiang, H. **Langmuir (2015) 31, 8228-8234**



Lipid Vesicles

- Slab Model
 - Core-multishell (Lipid Tail, Lipid Head, Solvent)
- SPD Model
 - Parsing of Lipid Molecule
 - Lipid Tail into 3 regions
 - Lipid Head into 3 regions
 - Solvent

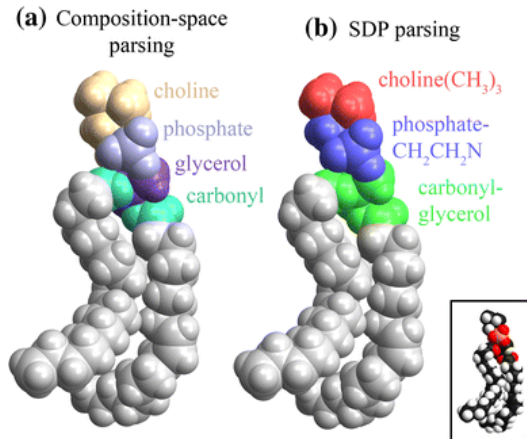
Contrast Variation improves data interpretation



- Core-multishell Model
- Choice of solvent D_2O % is important to ensure head group region can be defined well

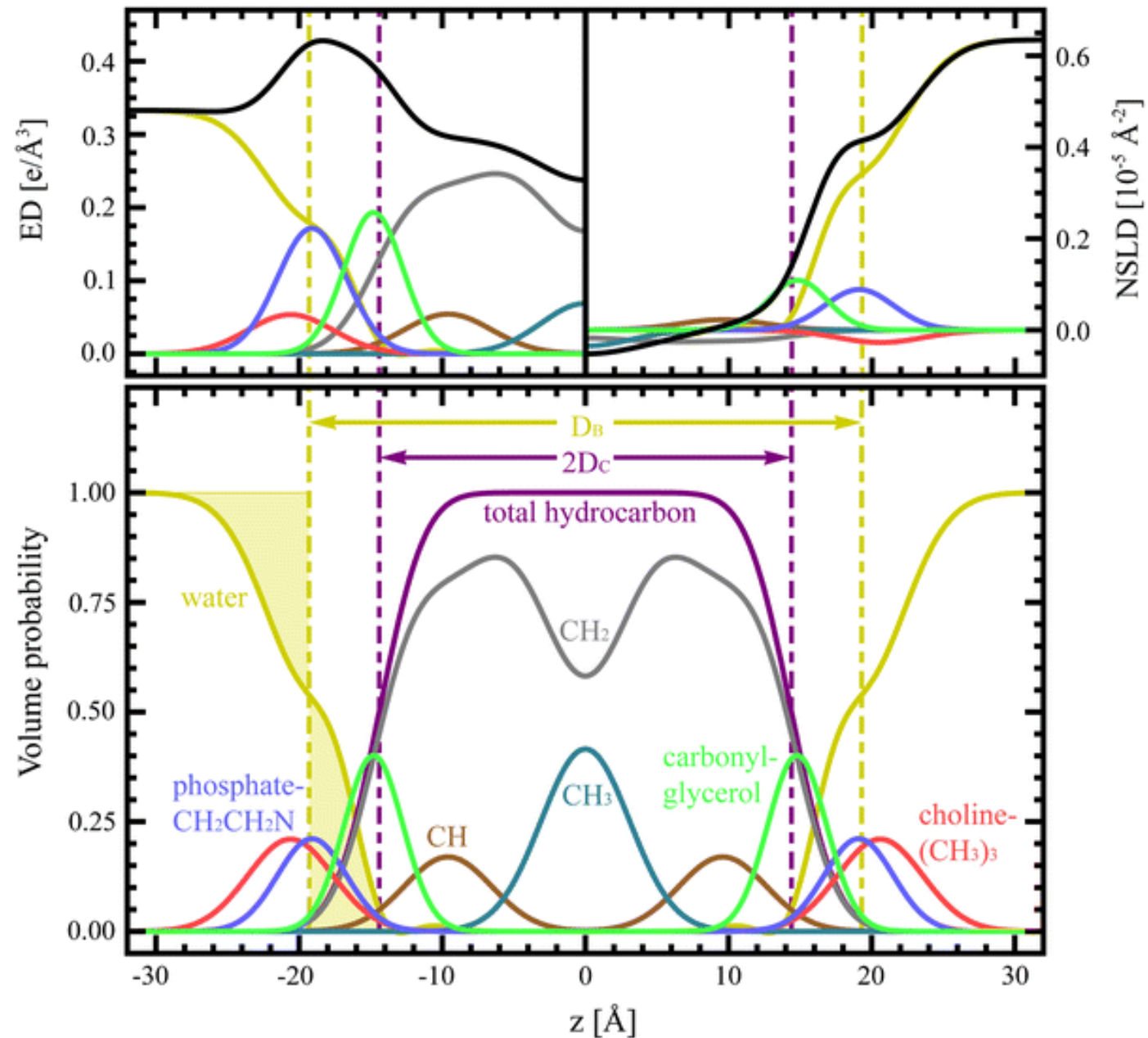
Smith, M.D., Pingali, S.V., Elkins, J.G., et al. **Green Chemistry** (2020) 22, 8278-8288

SDP Model



- Parsing different for SDP model
- To resolve several components, simultaneously fit 4-5 contrast conditions

Herberle, F., Pan, J., Standaert, R.F., et al. *Eur Biophys J* (2012) 41, 875-890



Plant Cell Wall

- Unified Fit Model (isotropic or random data)
- Composite (Form Factor + Power-law)

Fractal Slopes Interpretation

Fractal Type

Power-law Slope (Range)

Mass	- 1.0 to - 3.0	Bulk
Surface	- 3.0 to - 4.0	
Porod	- 4.0	
Diffuse-scattering	- 4.0 to - 5.0	

Mass-fractal structures	Power-law slope	
Polymers		
Polymer in good solvent	-1.5	
Self-avoiding walk (linear swollen polymer)	-1.67	
Random walk (polymer in theta solvent)	-2.0	
Swollen branched polymer	-2.0	
Randomly branched ideal polymer	-2.29	
Non-equilibrium growth processes		
Multiparticle diffusion limited aggregate	-1.8	
Percolation cluster	-2.5	
Diffusion limited aggregate	-2.5	
Low-dimensional objects (apparent mass fractals)		
Randomly distributed rods	-1.0	
Randomly distributed lamellae or platelets	-2.0	

Beaucage, G. *J Appl. Cryst.* (1995) 28, 717-728

Beaucage, G. *J Appl. Cryst.* (1996) 29, 134-146

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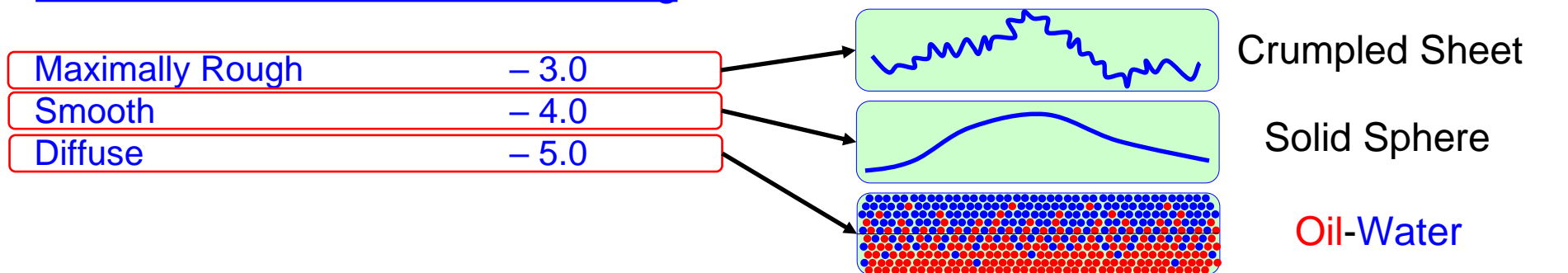
Beaucage, G. *J Appl. Cryst.* (1995) 28, 717-728

Beaucage, G. *J Appl. Cryst.* (1996) 29, 134-146

Fractal Slopes Interpretation

<u>Fractal Type</u>	<u>Power-law Slope (Range)</u>	
Mass	- 1.0 to - 3.0	} Interface/ Surface
Surface	- 3.0 to - 4.0	
Porod	- 4.0	
Diffuse-scattering	- 4.0 to - 5.0	

Surface-fractal Power-law Scattering

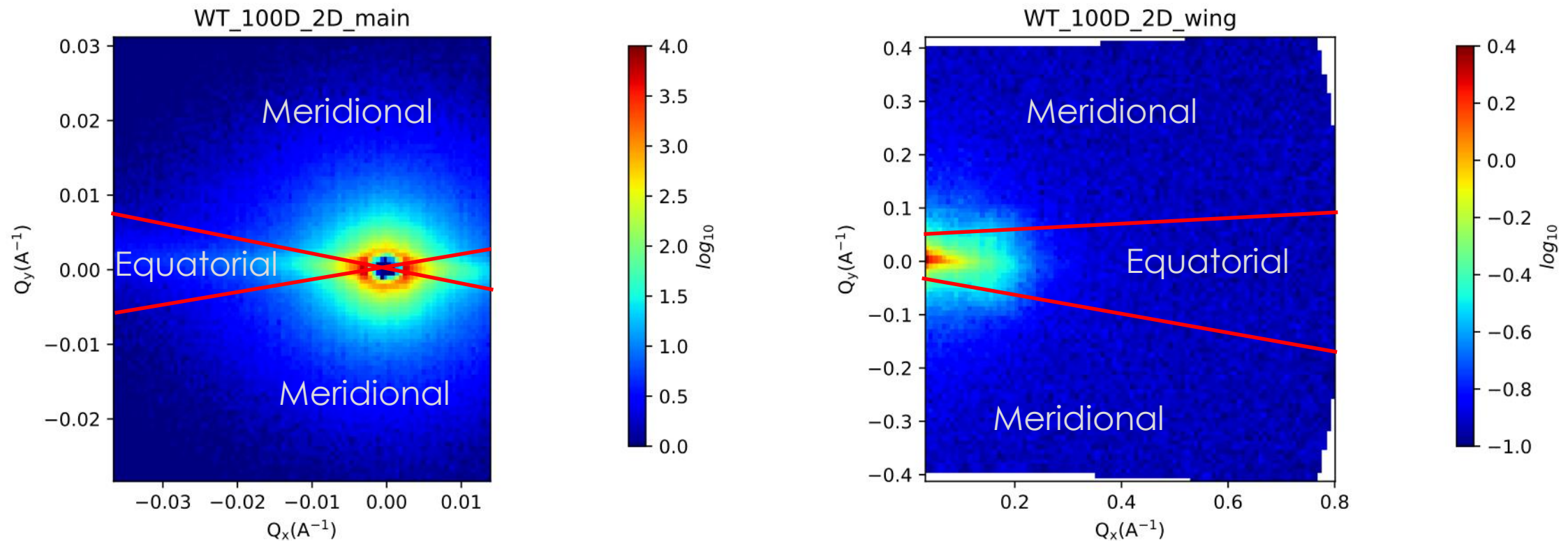


Beaucage, G. *J Appl. Cryst.* (1995) **28**, 717-728
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Plant Cell Wall

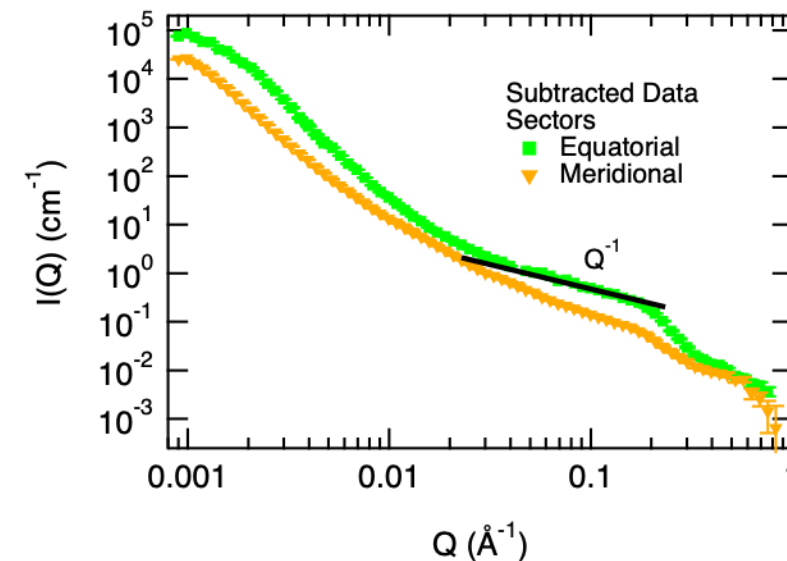
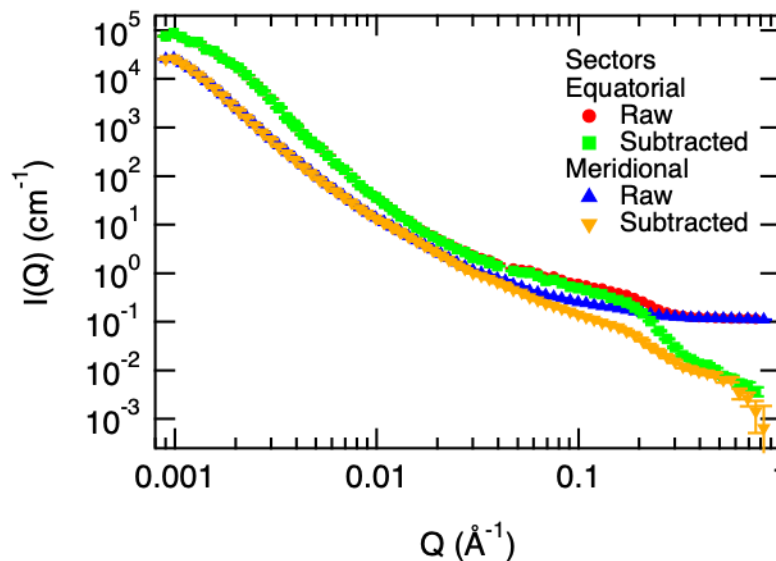
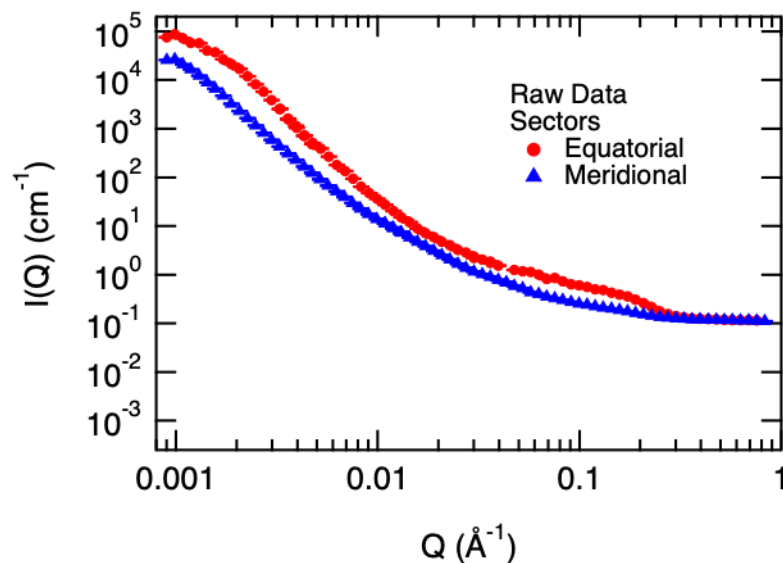
- Unified Fit
- Composite Model Fit

Anisotropic Data - Plant Cell Wall



- Divide into sectors - Equatorial and Meridional

Anisotropic Data - Plant Cell Wall



- Solvent Scattering is the same
- Clear differences in structure based on direction of scatter

Questions?

Facility Acknowledgment Statement

- A portion of neutron scattering research presented as examples in this introduction used resources at the High Flux Isotope Reactor or Spallation Neutron Source, DOE Office of Science User Facilities, operated by the Oak Ridge National Laboratory.
- The Bio-SANS of the Center for Structural Molecular Biology at the High Flux Isotope Reactor is supported by the Office of Biological and Environmental Research of the U.S. DOE.