

Overview

L8.1 – Introduction to Small Angle Neutron Scattering

L8.2 – SANS Instrumentation

EX8 – Virtual SANS Experiment

L9.1 – How to do a SANS Experiment

L9.2 – Small Angle Scattering Data Analysis

F9.3 – Applications of SANS

EX9 – Analysing Small Angle Scattering Data

How to Do a SANS Experiment

Andrew Jackson

NNSP-SwedNess Neutron School 2017, Tartu

Lecture L9.1

Planning an Experiment

- What is the question?
- Choosing samples
- Choosing an instrument
- Sample characterisation

As with any experiment, the question being asked must be carefully chosen.

SANS provides information about structure on the 1 to 100's of nm length scale

Is there contrast in the sample?

Do you need to use a deuteration scheme?

Can your system be studied as is, or does a model system need to be developed?

Planning an Experiment

- What is the question?
- Choosing samples
- Choosing an instrument
- Sample characterisation

Having identified the question, what samples are needed to answer that question?

This includes choices of concentration, deuteration, sample conditions (pH, temperature, pressure etc) and available sample amount.

Sample volumes for SANS are in the 0.1 to 1 ml range

Planning an Experiment

- What is the question?
- Choosing samples
- Choosing an instrument
- Sample characterisation

The choice of instrument is determined by:

- Required Q range
- Required beam flux
- Availability of access
- Availability of sample environment

To determine the requirements of Q range and flux, the scattering should be simulated.

Counting times are between minutes and hours per sample.

This requires some knowledge or expectation of what the sample structure will be.

The simulation can often be performed using the tools that will be used for data analysis.

Planning an Experiment

- What is the question?
- Choosing samples
- Choosing an instrument
- Sample characterisation

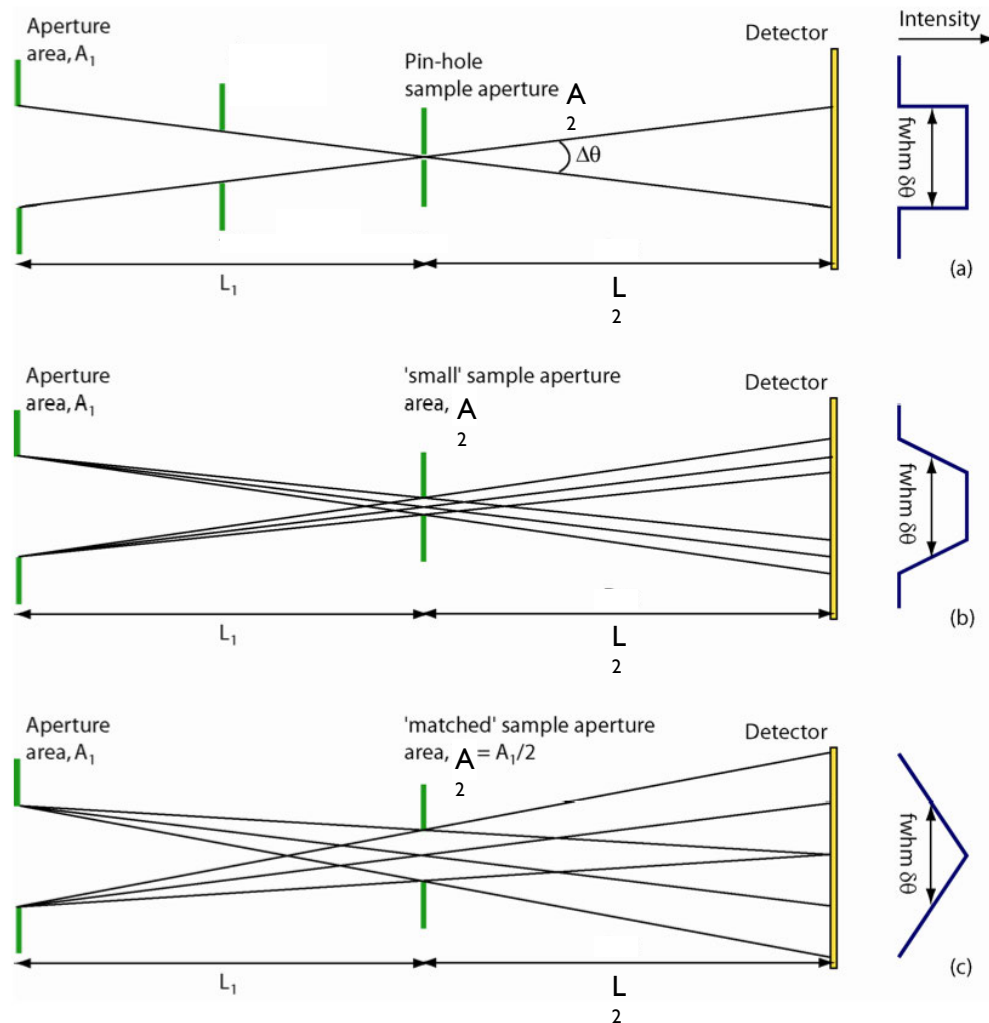
SANS is a relatively expensive technique that is uniquely capable for answering specific questions about nanoscale structure.

In order to make best use of SANS, the samples should be characterised with other techniques before planning and executing the SANS experiment.

Thus, for example, if light scattering or lab SAXS are available, these should be measured. Perhaps microscopy (light or electron) would be appropriate.

Bear in mind that these other techniques measure different aspects of the sample from SANS, but are all valuable information in being able to understand the SANS data.

Setting up the Instrument



The instrument scientist who is your contact at the scattering facility (“local contact”) will help you to determine the best instrument settings for your experiment.

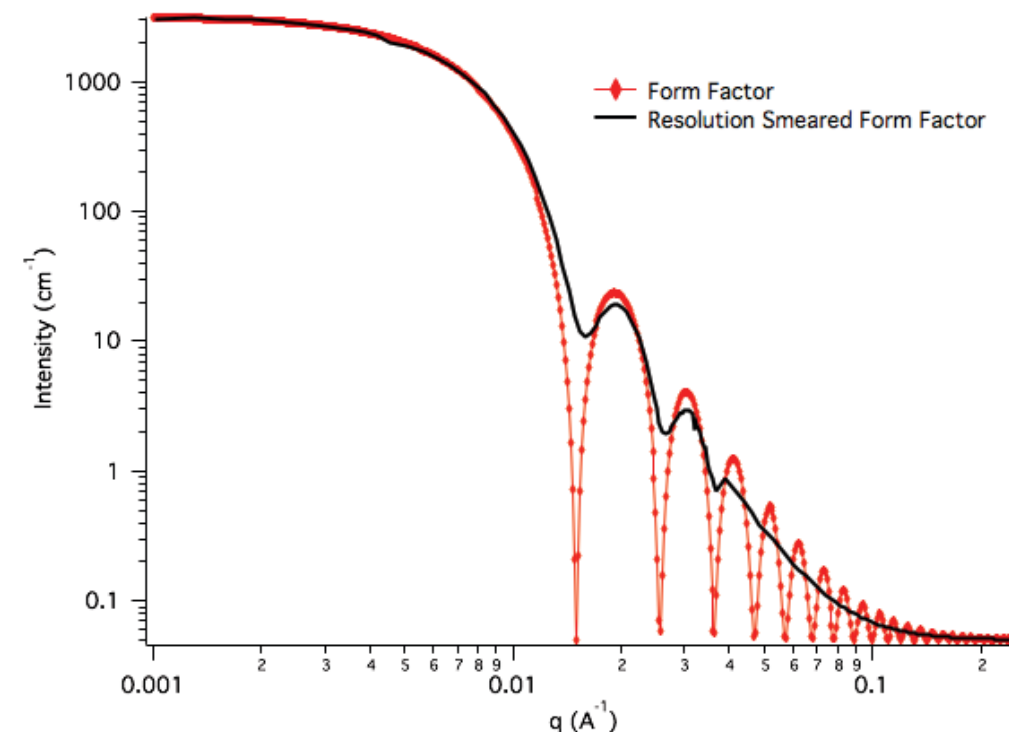
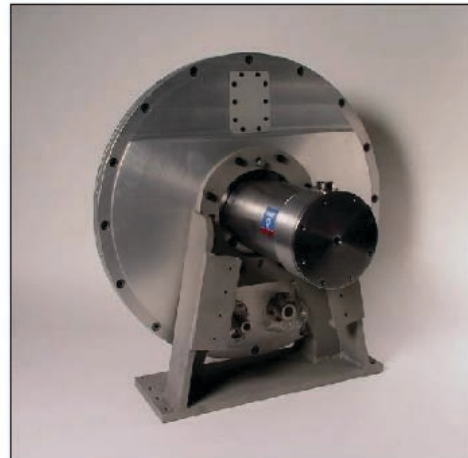
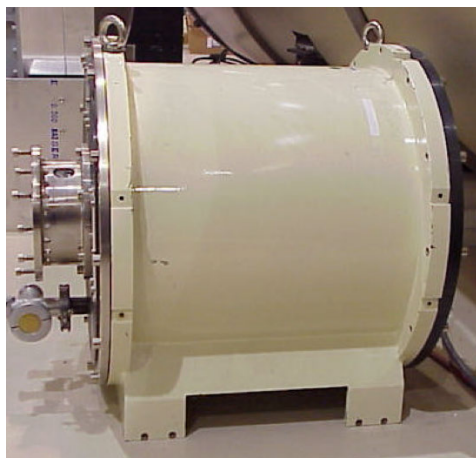
You need to choose:

Collimation length

Aperture sizes

Wavelength or wavelength range

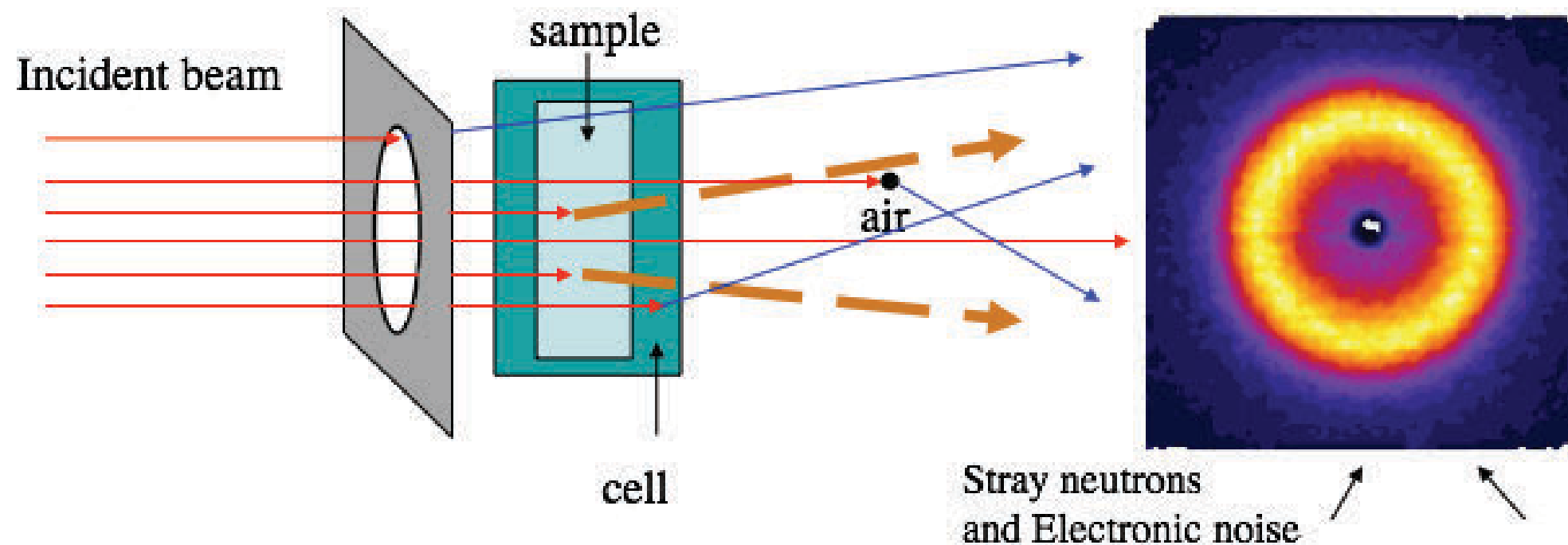
Detector position



Making a measurement

Contributions to counts on the detector:

1. Scattering from sample (what we want!)
2. Scattering from other than the sample (neutrons still go through sample)
3. Stray neutrons and electronic noise (neutrons don't go through sample)



$$I_{\text{meas}}(\mathbf{i}) = \Phi \, t \, A \, \varepsilon(\mathbf{i}) \, \Delta\Omega \, T_{\text{c+s}} [(\frac{d\Sigma}{d\Omega})_{\text{s}}(\mathbf{i}) \, d_{\text{s}} + (\frac{d\Sigma}{d\Omega})_{\text{c}}(\mathbf{i}) \, d_{\text{c}}] + I_{\text{bgd}} \, t$$

Making a measurement

$$\frac{d\Sigma}{d\Omega} \propto I(Q)_{measured}$$

$$I_{meas}(\mathbf{i}) = \Phi \, t \, A \, \varepsilon(\mathbf{i}) \, \Delta\Omega \, T_{c+s} [(\mathbf{d}\Sigma/\mathbf{d}\Omega)_s(\mathbf{i}) \, d_s + (\mathbf{d}\Sigma/\mathbf{d}\Omega)_c(\mathbf{i}) \, d_c] + I_{bgd} \, t$$

Φ = neutron flux on sample
 t = counting time for measurement
 A = sample area
 $\varepsilon(\mathbf{i})$ = detector element efficiency
 $\Delta\Omega$ = detector element solid angle

T_{c+s} = measured transmission of sample and holder
 d_s = thickness of sample
 d_c = thickness of cell
 I_{bgd} = stray neutrons and noise

We must make the necessary measurements:

- A. Scattering with sample in the neutron beam
- B. Scattering with an empty sample holder in the neutron beam
- C. Scattering with the sample position blocked by a neutron absorber
- D. The direct beam intensity with nothing in the neutron beam
- E. The direct beam intensity with the sample in the neutron beam
- F. The direct beam intensity with the sample holder in the neutron beam
- G. A measurement of the detector response variation (usually done by the facility before your experiment)

Your local contact for your experiment will make sure that these things are measured and the facility will provide the software necessary for you to leave with “reduced data” on “absolute scale” which is what you need to be able to perform an analysis and answer your scientific question.

What does it look like?



Image from ORNL

Two SANS Instruments @ HFIR reactor at Oak Ridge National Lab

What does it look like?

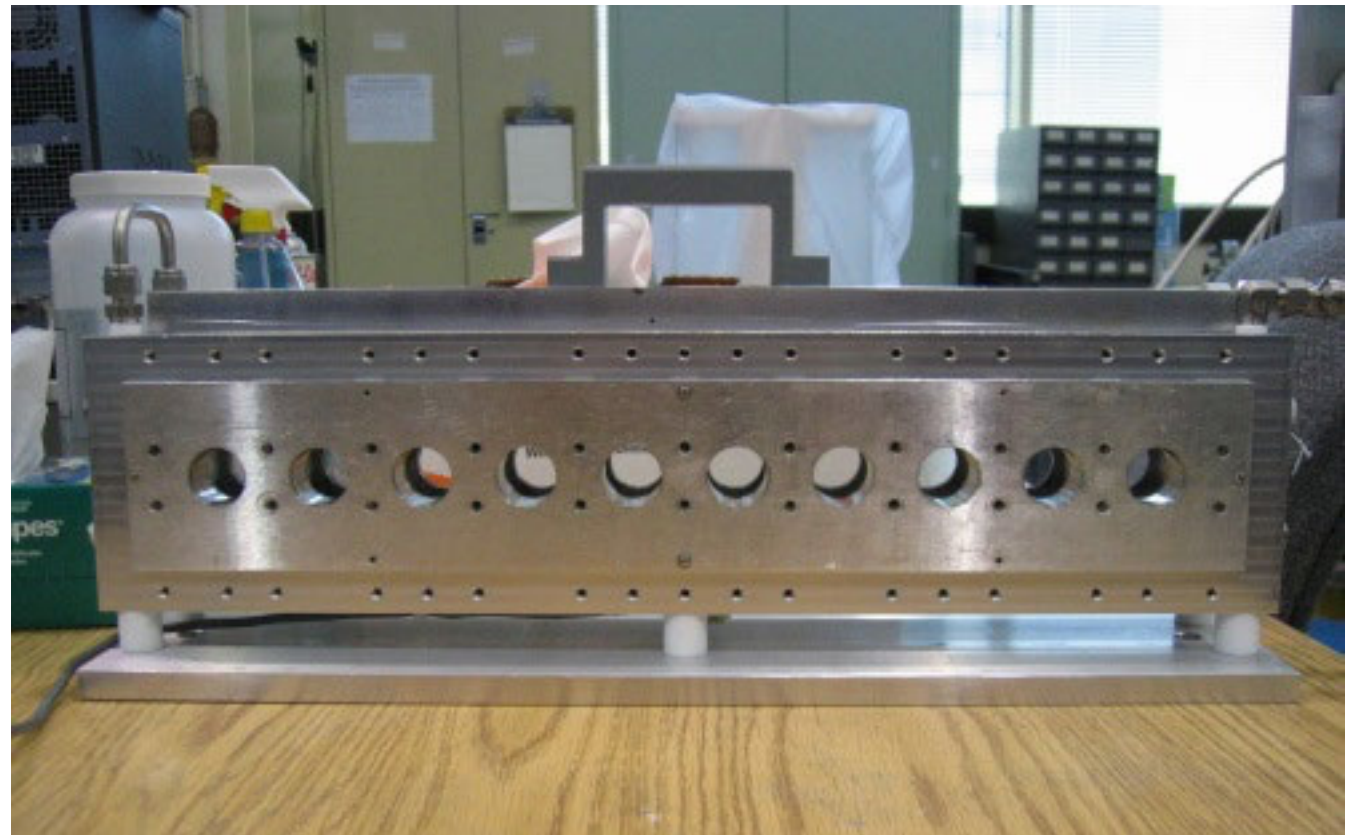
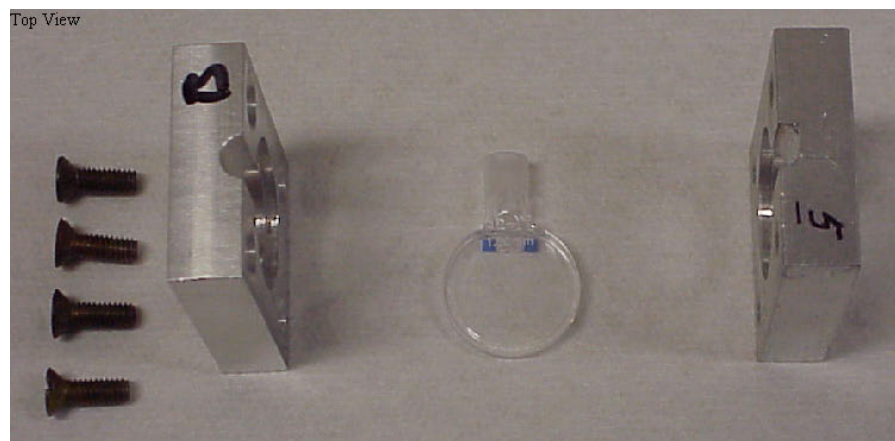


Image from ISIS/STFC

SANS instrument @ ISIS spallation neutron facility

What does it look like?

Sample Cells



Temperature Controlled Sample Changer

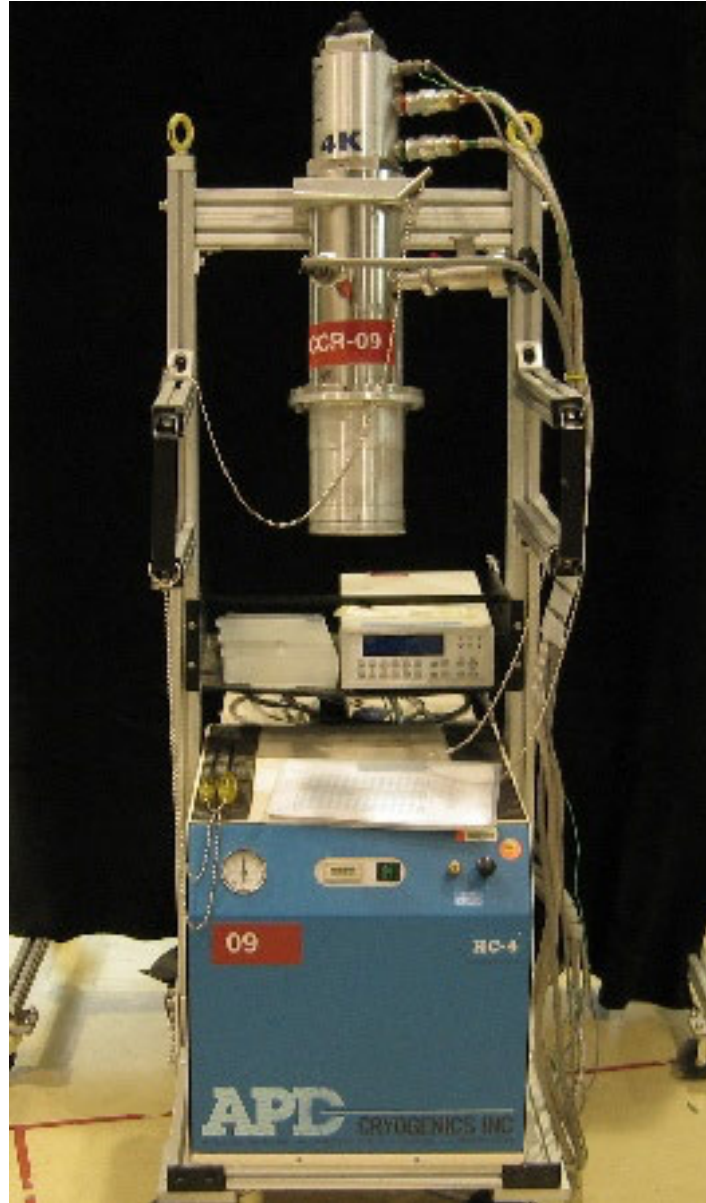
Images from NIST Center for Neutron Research

Sample environment is the various equipment that the sample is placed in - usually to apply a stimulus to the sample

What does it look like?



Rheometer



Closed Cycle Refrigerator



Humidity Chamber

Images from NIST Center for Neutron Research

Sample environment is the various equipment that the sample is placed in - usually to apply a stimulus to the sample

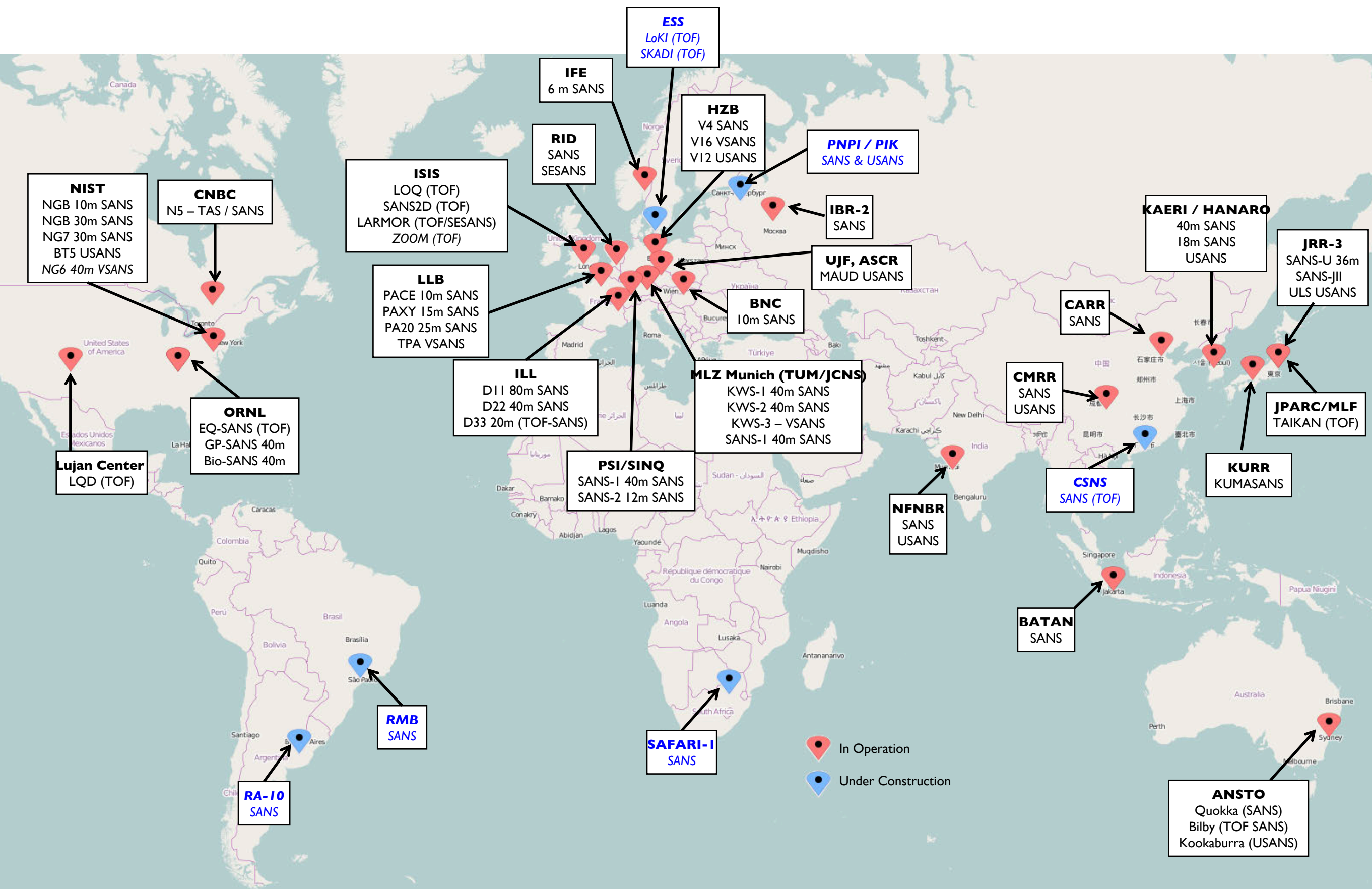
What does it look like?



Image from ISIS/STFC

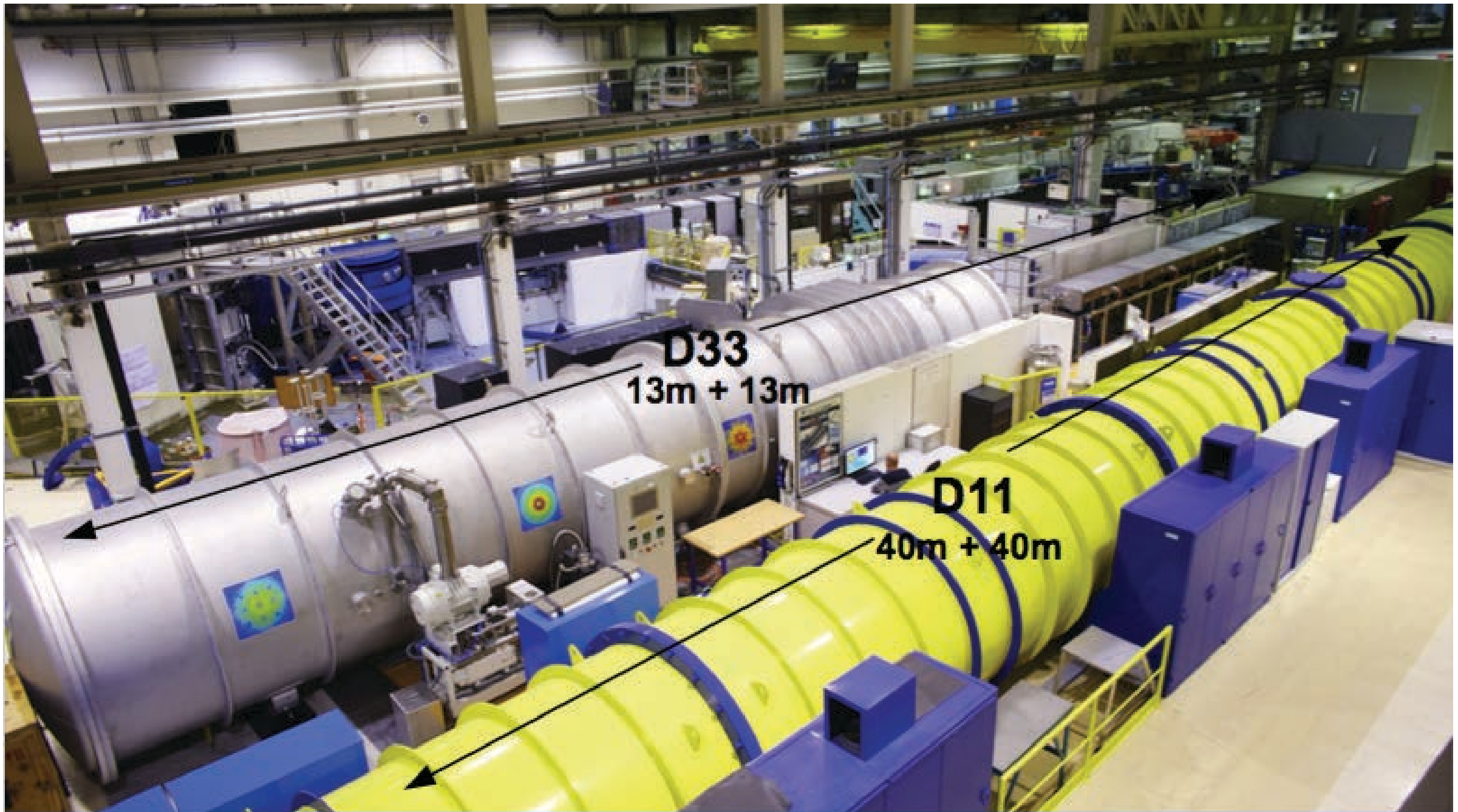
SANS sample position at SANS2D @ ISIS with 17T superconducting cryomagnet in place

SANS Instruments Around the World



SANS Instruments Around the World

ILL



SANS Instruments Around the World

D11

Monochromator	
velocity selector	
Eleanore/ Felicia (ASTRUM standard selectors)	$\Delta\lambda/\lambda = 9\%$ (FWHM)
incident wavelength	variable, $4.5 \leq \lambda [\text{\AA}] \leq 40$
max. wavelength at max. detector dist. (39 m)	$\lambda = 22 \text{ \AA}$
selector rotation normal to n-beam possible minimum wavelength at rotation -7 deg	-7 deg to + 7 deg $\lambda = 3.2 \text{ \AA}$
Collimation	
11 guide sections (computer controlled)	cross section (height x width) : 50 x 30 mm ² -> 50 x 45 mm ² over 6.5 m straight guide 50 x 45 mm ² over 28 m 50 x 45 mm ² -> 35 x 31.5 mm ² over 4 m
uncoated glass guides (SwissNeutronics)	
diverging "trumpet" + straight guide + focussing guide near sample position	
guide-to-sample distances	1.5, 2.5, 4, 5.5, 8, 10.5, 13.5, 16.5, 20.5, 28, 34, 40.5 m (12 discrete distances)
Attenuators	
choice between 3 cadmium sheets of different transmission (computer controlled) $T_s=3.461 \cdot 10^{-3}$ (att 1), $T_s=1.089 \cdot 10^{-3}$ (att 2), $T_s=3.524 \cdot 10^{-4}$ (att 3)	
Sample area	
flux at specimen at lowest resolution	$1 \cdot 10^8 \text{ n cm}^{-2} \text{ s}^{-1}$
typical sample size	10 x 10 mm ²
Detector	
sample-to-detector distances L	variable between 1.2 m and 39 m
momentum transfer range	$3 \cdot 10^{-4} \leq Q [\text{\AA}^{-1}] \leq 1$
detector type	³ He gas detector (CERCA)
area	96 x 96 cm ²
pixel size	7.5 x 7.5 mm ²
background	1 Hz on whole multidetector
detector deadtime	420 ns

ILL

D22

Monochromator	
velocity selector Anatole	$\Delta\lambda/\lambda = 10\%$ (standard)
wavelength	$4.5 < \lambda/\text{\AA} < 40$ (for $\Delta\lambda/\lambda = 10\%$)
Collimation	
8 guide sections	55 x 40 mm
source-to-sample distances / m	1.4, 2.0, 2.8, 4.0, 5.6, 8.0, 11.2, 14.4, 17.6, variable apertures at 19.1
Sample area	
maximum flux at sample (for $\Delta\lambda/\lambda = 10\%$)	$1.2 \times 10^8 \text{ n cm}^{-2} \text{ s}^{-1}$
typical sample size	10 to 300 mm ²
Detector	
distances	1.1 ... 17.6 m
rotation	-2° < 2θ < 22°
horizontal offset	-5 ... 50 cm
area	102.4 x 98 cm ²
pixel size	8 x 8 mm ²
maximum counting rate	5 MHz
electronic noise	2 Hz for the whole multidetector

D33

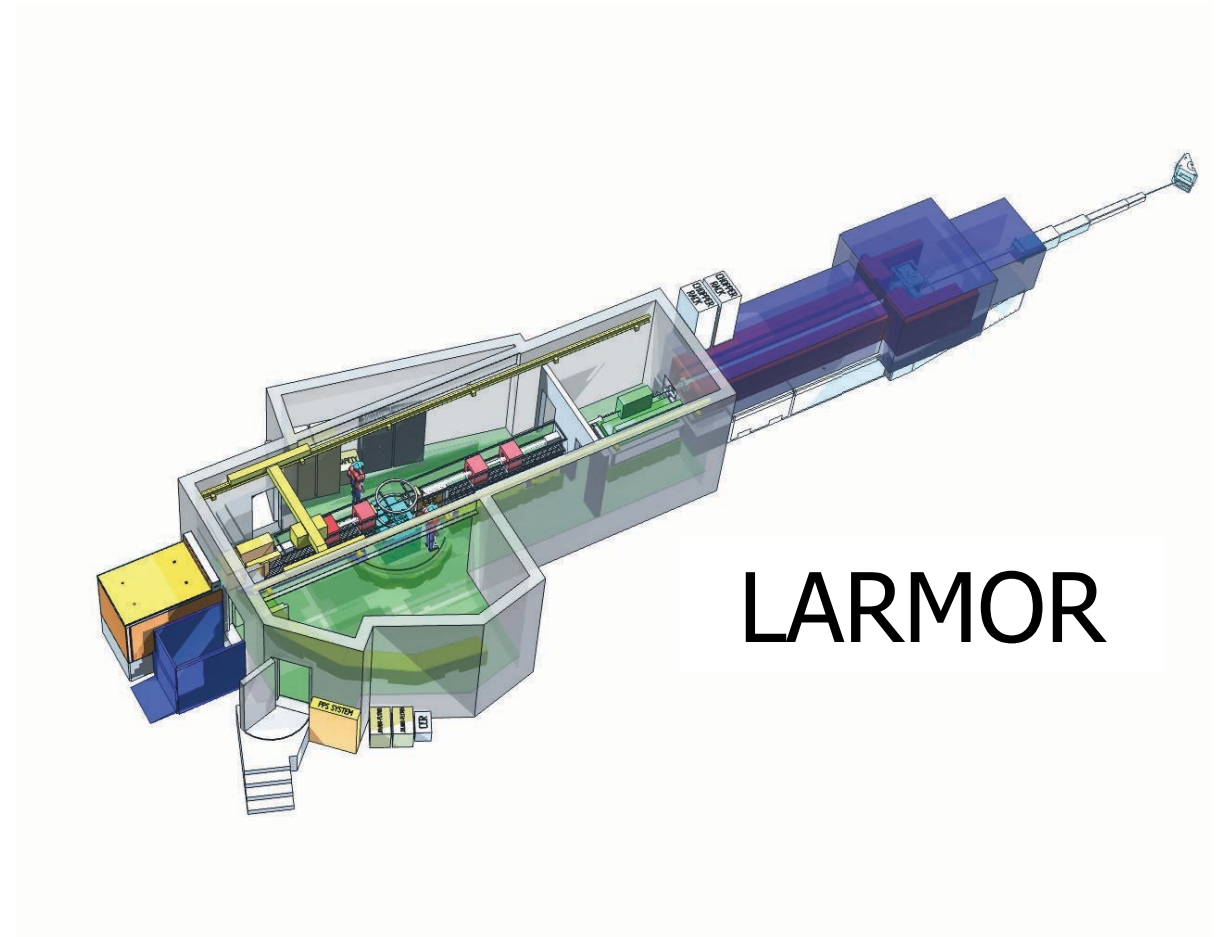
Wavelength Definition	
Monochromatic Mode:	
Velocity selector (Astrium)	$\Delta\lambda/\lambda = 10\%$ (standard)
Wavelength range	$4.5 < \lambda/\text{\AA} < 40$ (for $\Delta\lambda/\lambda = 10\%$)
Time-of-Flight (TOF) Mode:	
4-chopper system (Astrium)	
Wavelength cut-offs	14 Å and 20 Å
Wavelength resolutions	$\Delta\lambda/\lambda = 2\%$ to 26 % (depending on chopper pair & detector distance)
Collimation	
4 movable guide sections	2.5 m
Beam nose	2.8 m
Guide cross-section	30 x 30 mm
Source-to-sample distances (m)	2.8, 5.3, 7.8, 10.3, 12.8
Source apertures	30x50 mm (off-centre), 30x30 mm diameters: 5, 10, 20, 30 mm
Sample area	
Maximum flux at sample (for $\Delta\lambda/\lambda = 10\%$)	$4.1 \times 10^7 \text{ n cm}^{-2} \text{ s}^{-1}$
Brightness (flux / unit solid angle)	$3.57 \times 10^{11} \text{ n cm}^{-2} \text{ s}^{-1} \text{ strd}^{-1}$
Maximum sample dimensions	15 mm x 15 mm
Sample environments	Sample changer, Electromagnet, Cryostat, Cryomagnet, Furnace, Stopped-flow, Shear cell
Detectors	
Sample - Detector distances	1.2 ... 12.8 m
Detector 1 (rear):	
Single panel monoblock	640 x 640 mm
Pixel size	5 x 5 mm ² (128 x 128 pixels)
Maximum count rate	4 MHz (global) ; 3 kHz/pixel (local)
Detector 2 (front):	
4-panel monoblock	160 x 640 mm each panel
Pixel size	5 x 5 mm ² (32 x 128 pixels)
Maximum count rate	4 MHz (global) ; 3 kHz/pixel (local)

SANS Instruments Around the World

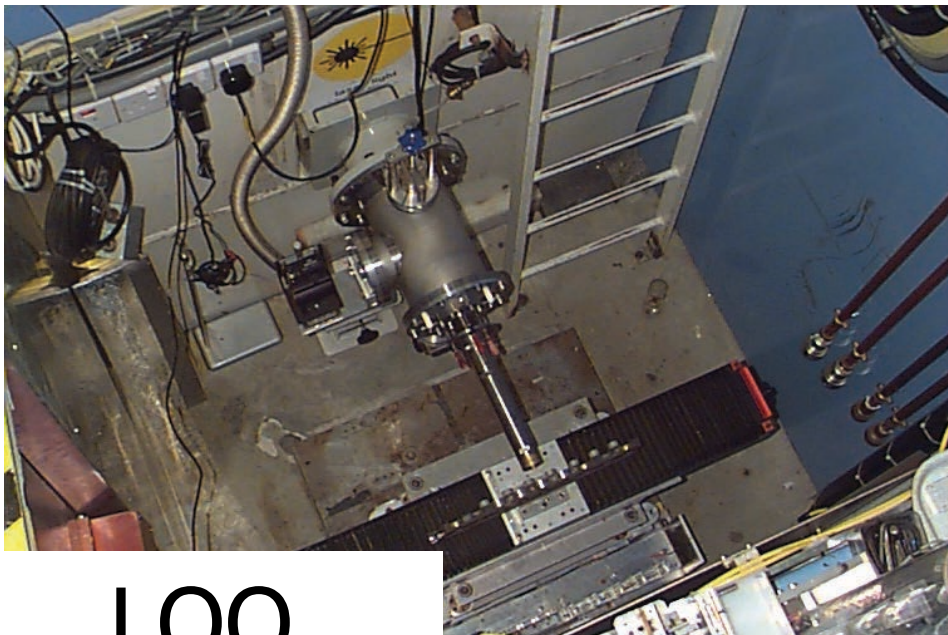
SANS2D



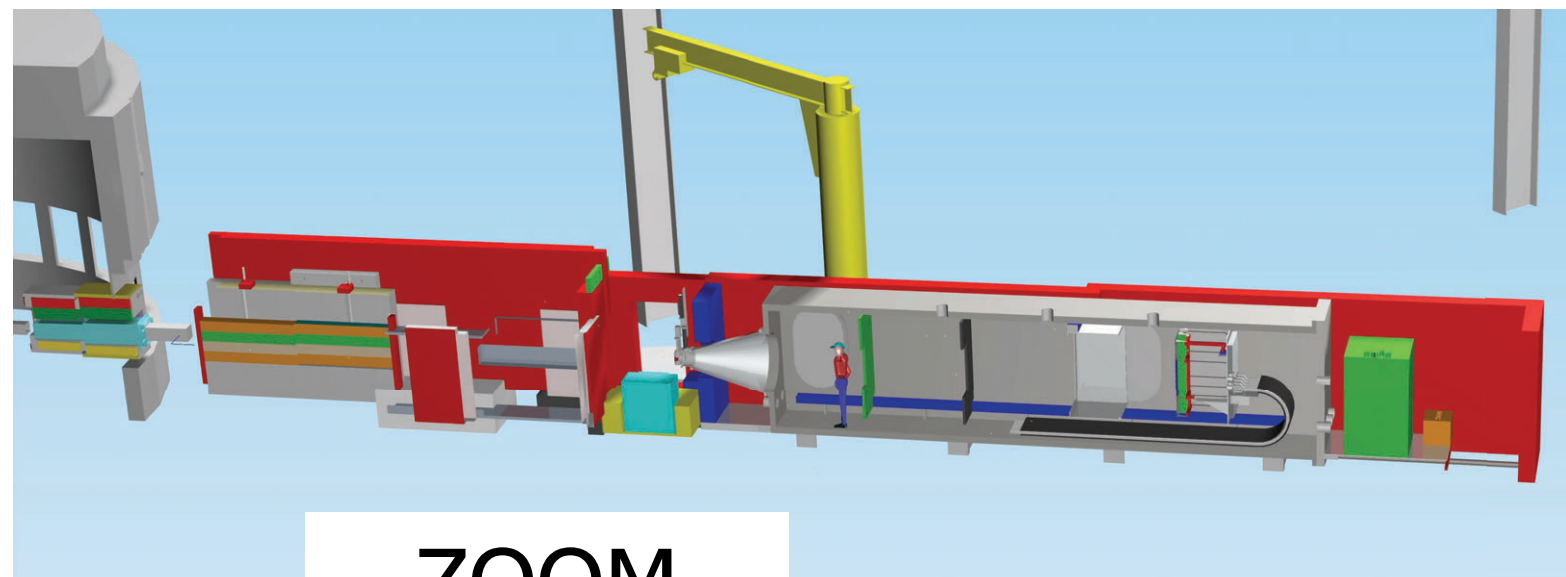
ISIS



LARMOR



LOQ



ZOOM

SANS Instruments Around the World

SANS2D



Incident wavelengths	2.0 - 14.0 Å at 10 Hz
Momentum transfer, Q	Depends on sample-detector distances and detector offsets: $Q_{\min} \sim 0.002 \text{ Å}^{-1}$, $Q_{\max} \sim 3 \text{ Å}^{-1}$

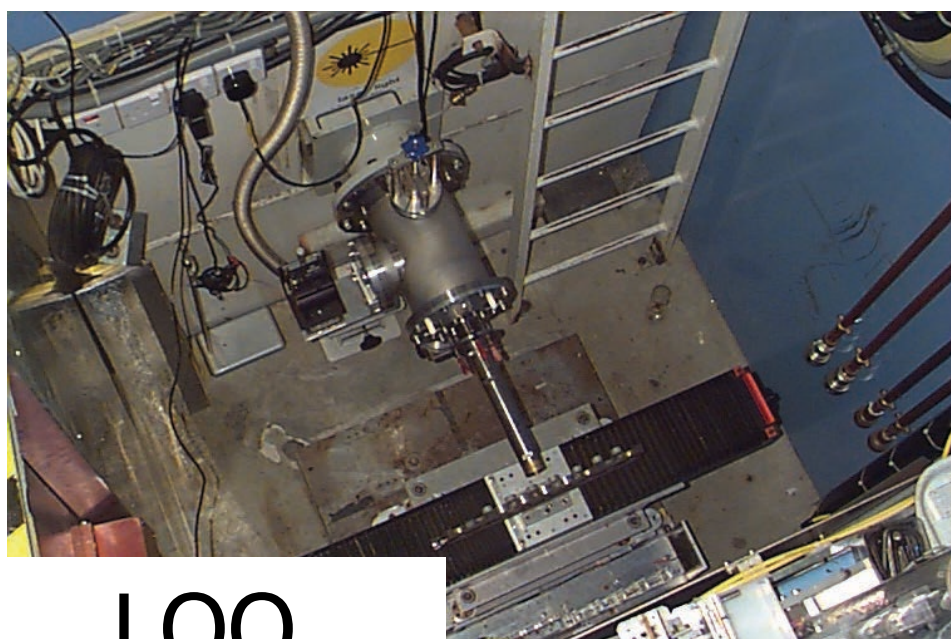
ISIS

Isis Beamline	E2, viewing the coupled cold moderator
Primary flight path	3Θ _c Ni super-mirror bender (to remove neutrons with wavelengths less than 1.5 Å), upstream scintillator monitor, overcount protection fast shutter, variable-opening counter-rotating disc chopper, 5 x 2 m moveable evacuated collimation sections (each with choice of Ni guide sections or plain pipe) for variable incident collimation, 5 moveable aperture strips (each with different size apertures), sample position scintillator beam monitor, sample aperture strip, final collimation tube.
Sample position	Around 19 m from moderator. Side access. No height restriction. Beam is approximately 1 m above base plate. Crane access possible (SWL <5000 Kg). Sample transmission scintillator monitor on motorised rack. Provided with water, helium and electrical services.
Beam size at sample	Defined by sample aperture strip and final collimation. Up to 15 mm diameter is possible, typically 8 mm diameter.
Neutron flux at sample	Dependent on collimation, accelerator performance and target type. Typical time-averaged flux is <i>currently estimated</i> to be >10 ⁶ cm ⁻² s ⁻¹ (ISIS TS2 at 10Hz, 40 uA 800 MeV proton beam, tantalum target).
Secondary flight path	Evacuated tank containing detectors.
Detectors	Two ³ He-CF ₄ filled ORDELA "area" detectors. Active area of each is 96.5 cm x 96.5 cm with 5 mm resolution. The detectors can be moved in the vacuum tank both along the beam (to vary sample-detector distance between 2 and 12 m) and to give a sideways offset (to extend the Q range at a given detector distance) of up to 1200 mm. The front detector can also be rotated to face the sample. Detector mapping under software control.

SANS Instruments Around the World

ISIS

Incident wavelengths	2.2 - 10.0 Å at 25 Hz, 2.2 - 6.7 Å or 6.3 - 10.0 Å at 50 Hz
Momentum transfer, Q	0.006 - 0.24 Å ⁻¹ (main detector) 0.15 - 1.4 Å ⁻¹ (high-angle bank)
Dynamic range in Q	40 (on main detector), 230 (simultaneous use of all detectors)



LOQ

Isis Beamline	N5, viewing the 25 K liquid hydrogen (lower) moderator.
Primary flight path	Soller supermirror bender (24 mrad, to remove neutrons with wavelengths less than 2 Å), upstream scintillator monitor, aperture dial No 1, variable-opening (2 - 126 degrees) disc chopper, frame overlap mirror (removes neutrons with wavelengths greater than 12 Å), 3 m evacuated flight tube, sample position scintillator beam monitor, aperture dial No 2, final collimation tube.
Sample position	Around 11.1 m from moderator. Approximate size is 0.4 m (parallel to beam) by 1.5 m. No height restriction. Beam is approximately 0.63 m above base plate. Crane access possible (SWL 1000 Kg). Sample transmission scintillator monitor on motorised rack. Provided with water, helium and electrical services. Secondary, top-loading, in-vacuum sample position with limited access and services around 12.5 m from moderator (giving an approximate Q range of 0.01 - 0.34 Å ⁻¹).
Beam size at sample	Defined by aperture No 2 and final collimation. Between 2 - 20 mm diameter. Typically 8 mm diameter.
Neutron flux at sample	Dependent on collimation, ISIS accelerator performance and target type. Typical time-averaged flux is 2x10 ⁵ cm ⁻² s ⁻¹ (ISIS TS1 at 40Hz, 160 uA 800 MeV proton beam, tantalum target).
Secondary flight path	Evacuated tank to main detector.
Detector	3He-CF ₄ filled ORDELA "area" detector 15.15 m from moderator. Active area is 64 cm x 64 cm with 5 mm resolution. Detector mapping under software control. External, annular, high-angle, scintillator "area" detector bank 11.6 m from moderator.

Summary

Careful planning is needed to get the most information from a SANS experiment

Processing the data requires knowledge of some instrument specific values and calibrations – these will be provided by the facility.

So, choice of SANS instrument is driven by the needs of the experiment in terms of **Q-range, resolution and sample environment**

Questions?

Small Angle Scattering Data Analysis

Andrew Jackson

NNSP-SwedNess Neutron School 2017, Tartu

Lecture L9.2

What is SANS Data Analysis?

$$\frac{d\Sigma}{d\Omega} = \frac{N}{V} \frac{d\sigma}{d\Omega} = \frac{1}{V} \left| \int_V \rho(\vec{r}) e^{i\vec{q} \cdot \vec{r}} d\vec{r} \right|^2$$

“Rayleigh-Gans Equation”

Thus, inhomogeneities in $\rho(\vec{r})$ give rise to small angle scattering

Aim of data analysis is (usually) to extract information about the structure of the system from the scattering data.

This means recovering information about $\rho(r)$ from $I(Q)$

SANS Data Analysis

Model Independent

We can use an approximation from Guinier

$$I(Q) = I(0)e^{-\frac{(QR_g)^2}{3}}$$
$$\ln(I(Q)) = \ln(I(0)) - \frac{R_g^2}{3}Q^2$$

to obtain the radius of gyration of the scattering objects assuming particulate scatterers and $S(q) = 1$.

Similar approximations can be made to get the cross section of cylinders or the thickness of disks. Various other model independent approaches exist to extract information from the data without a scattering model.

Indirect Fourier Transform

Since we are missing the phase information as a result of the differential cross section being related to the square of the amplitude of the fourier transform, we cannot simply take the fourier transform of our data to get back to $\rho(r)$. Thus we must use an indirect method.

A popular implementation of this method is found in the *ATSAS suite* of software from Prof. Svergun's group. *SasView* also has an implementation of this method.

Model Dependent

We calculate the form and structure factors for a given scattering system and compare that with the measured scattering data. The model is fitted to the data to obtain the parameters that describe the scattering. We can simultaneously fit multiple contrasts to be able to study complex structures.

The software we will be using for this course is called *SasView* (<http://www.sasview.org>) and is being jointly developed by NIST, ILL, ISIS, SNS, ANSTO and ESS. Other software packages for this kind of analysis include the *NIST Igor Macros* developed at the NCNR and *SasFit* developed at the Paul Scherrer Institute.

Ab-initio Structure Generation

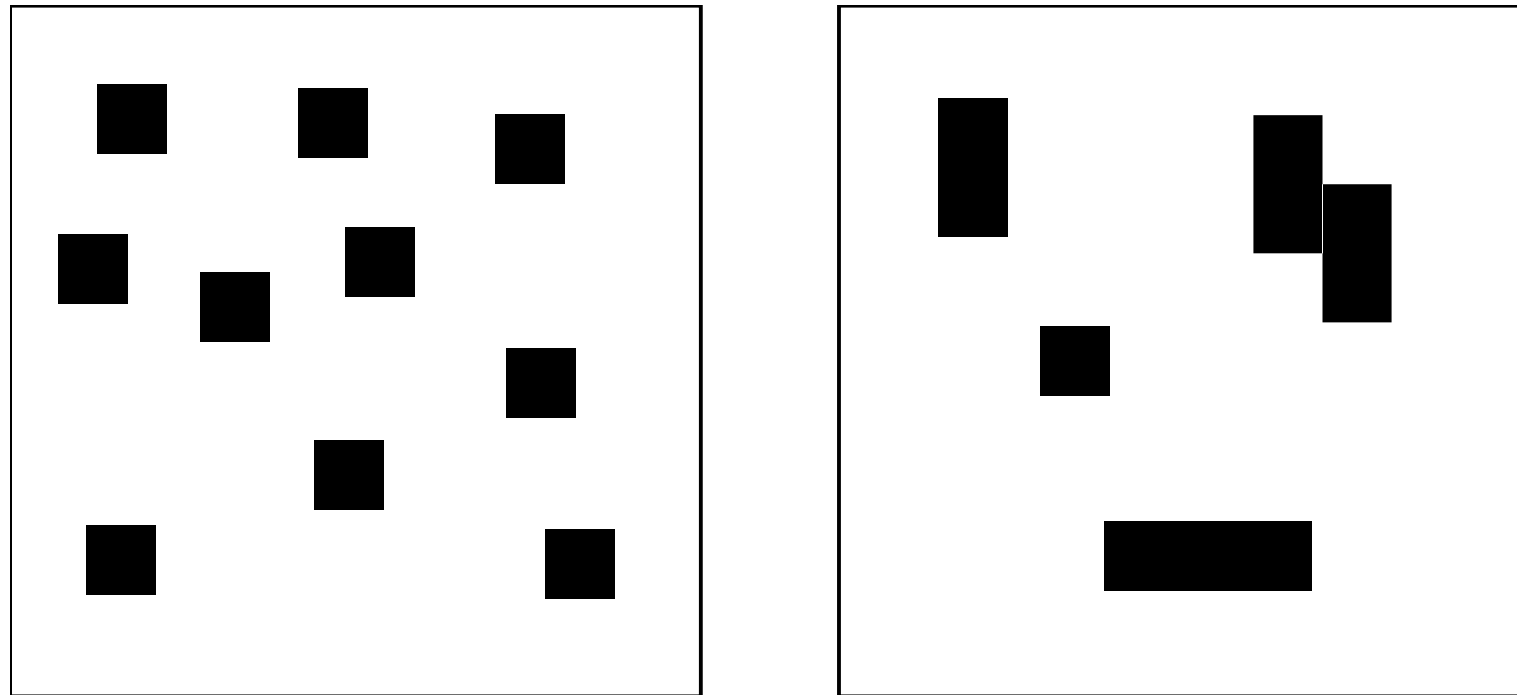
An approach that is popular for bio-macromolecules in solution is to generate a structure from many sub-resolution spheres and calculate what the scattering would be. That is then compared with the data and the spheres redistributed. This is repeated until agreement is found.

The *ATSAS suite* is the primary example of software using this method

Model Independent

Scattering Invariant

Porod showed that the total small angle scattering is invariant, irrespective of how the matter is distributed.



Two systems where the contrast and volume fraction are the same, but the distribution of matter is different. Both are 10% black and 90% white.

$$Q = \int \frac{d\Sigma}{d\Omega}(\mathbf{q}) d\mathbf{q}$$
$$= (2\pi)^3 (\rho(\mathbf{r}) - \bar{\rho})^2$$

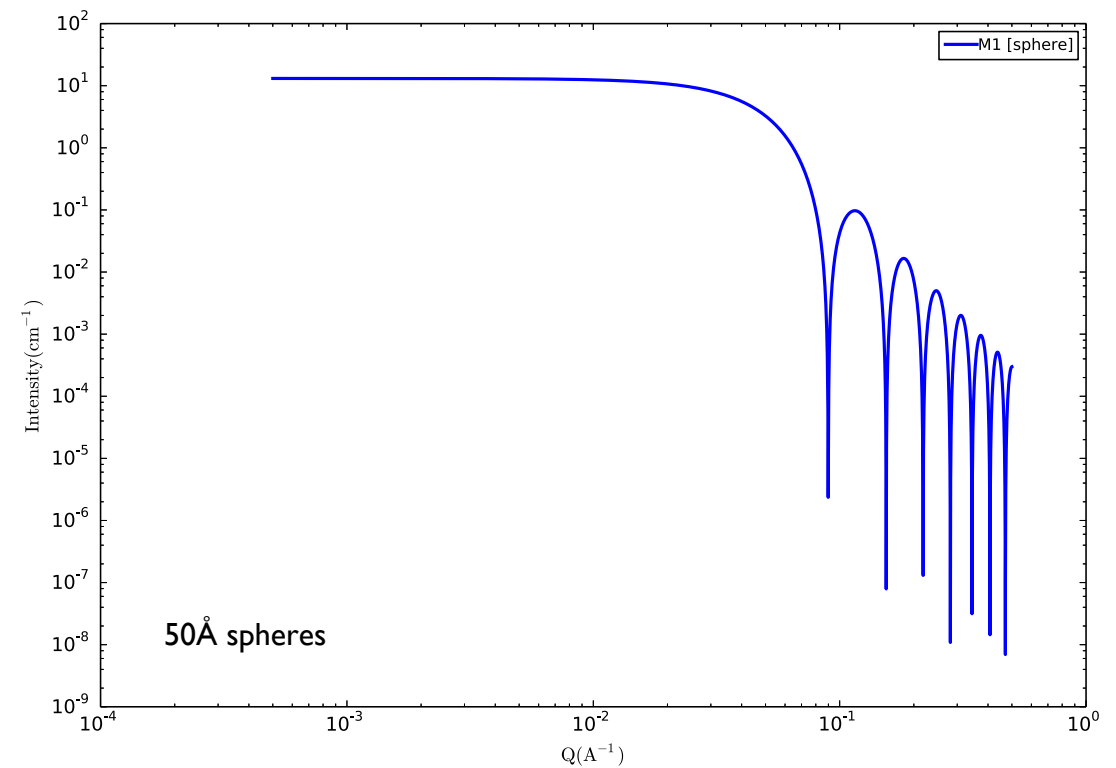
$$\frac{Q}{4\pi} = Q^* = 2\pi^2 \phi_1 (1 - \phi_1) (\rho_2 - \rho_1)^2$$

Model Independent Guinier Plot

We can use an approximation from Guinier¹
for spherical/globular objects

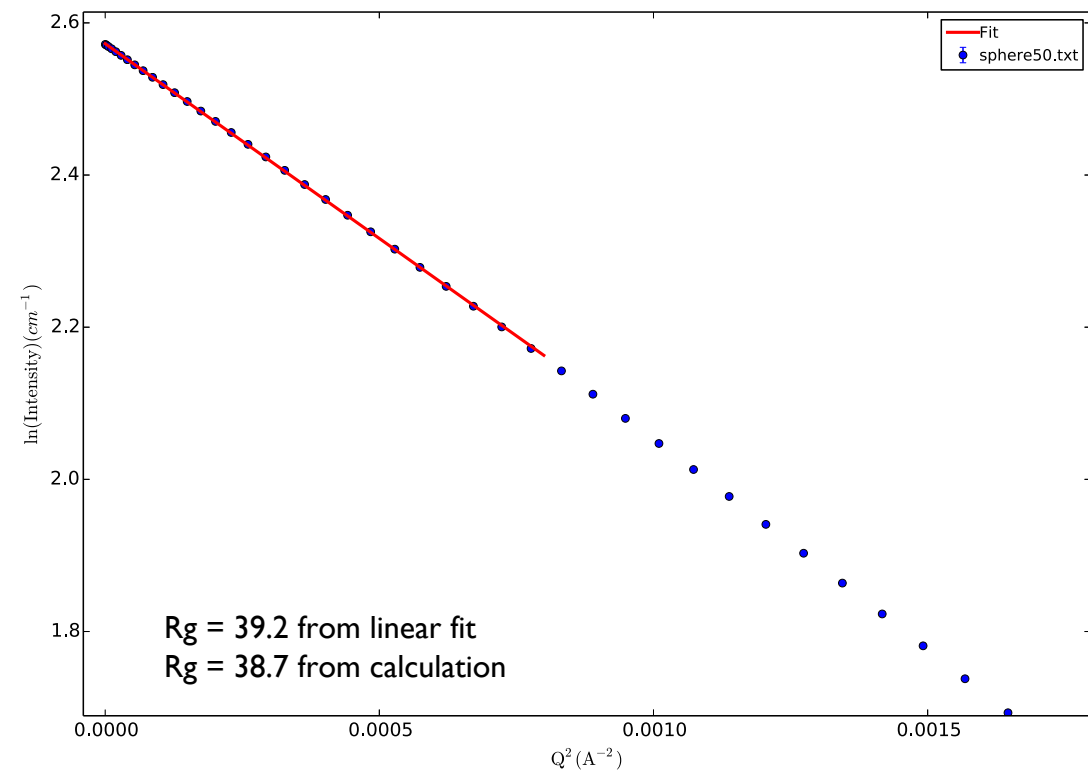
$$I(Q) = I(0)e^{-\frac{(QR_g)^2}{3}}$$

$$\ln(I(Q)) = \ln(I(0)) - \frac{R_g^2}{3}Q^2$$



Perform fit for $y(x) = ax + b$

Parameter a	-512	+/-	7.37e-15
Parameter b	2.57	+/-	2.25e-18
Chi2/dof	5.78e+26		
Maximum range (linear scale)	Min: 0	Max: 0.0283	
Fit range of $x^2(2)$	0	0.0008	
$I(q=0)$	13.1	+/-	2.95e-17
R_g [Å]	39.2	+/-	2.82e-16
$R_g \cdot Q_{min}$	0		
$R_g \cdot Q_{max}$	1.11		



Model Independent

Guinier Plot

For non spherical objects, similar results exist.

In the case of rod-like and disk-like or lamellar objects, the overall radius of gyration is given similarly to spherical objects from a plot of $\ln[I(Q)]$ vs Q^2 :

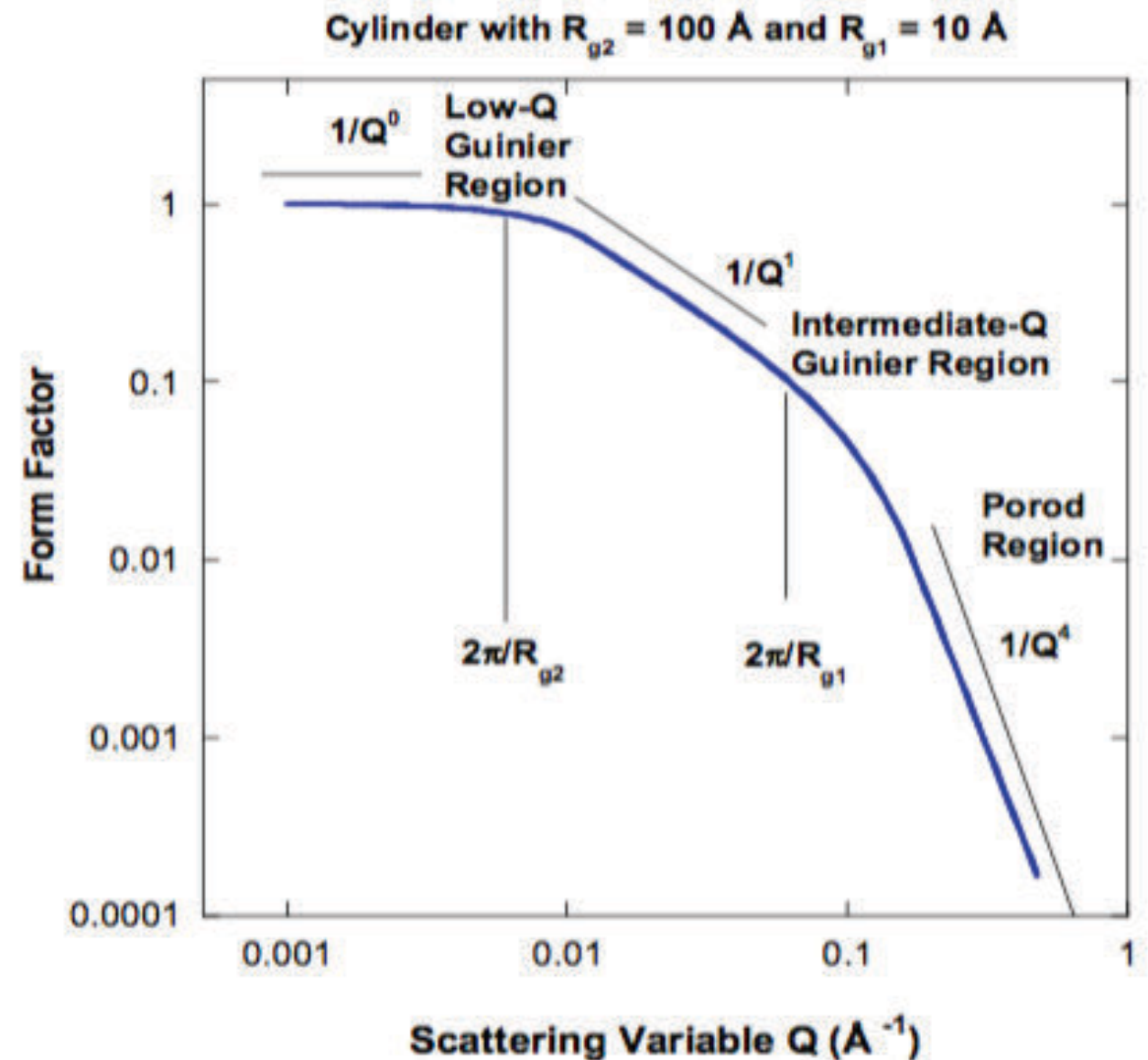
$$I(Q) = I(0) \exp\left(-\frac{Q^2 R_g^2}{3}\right) \quad \text{where } R_g^2 = \frac{L^2}{12} + \frac{R^2}{2}$$

There is also the intermediate region that gives the “cross sectional R_g ” from a plot of $\ln[QI(Q)]$ vs Q^2 :

$$I(Q) = \frac{I(0)}{Q} \exp\left(-\frac{Q^2 R_g^2}{2}\right) \quad \text{where } R_g^2 = \frac{R^2}{2}$$

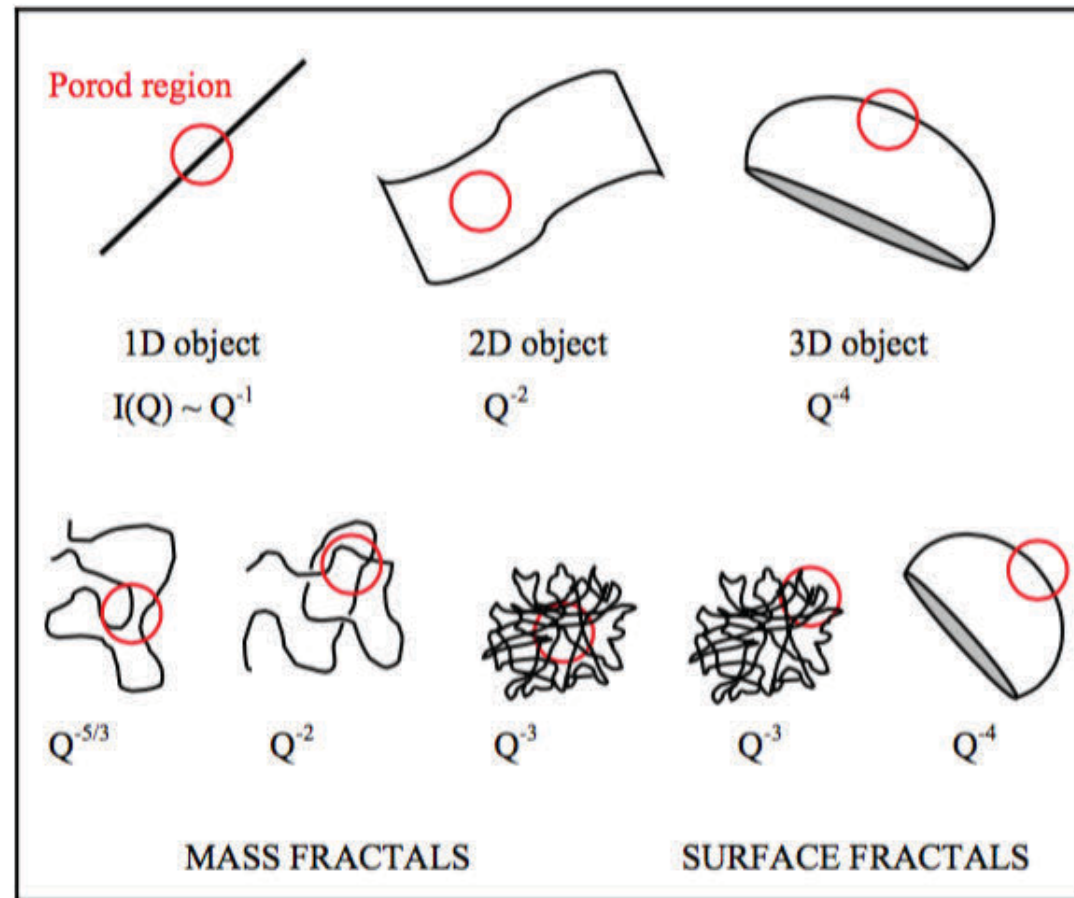
In the case of disk-like or lamellar objects, the intermediate region gives the “thickness R_g ”:

$$I(Q) = \frac{I(0)}{Q^2} \exp\left(-\frac{Q^2 R_g^2}{1}\right) \quad \text{where } R_g^2 = \frac{T^2}{12}$$



Model Independent

Porod Region and Porod's Law

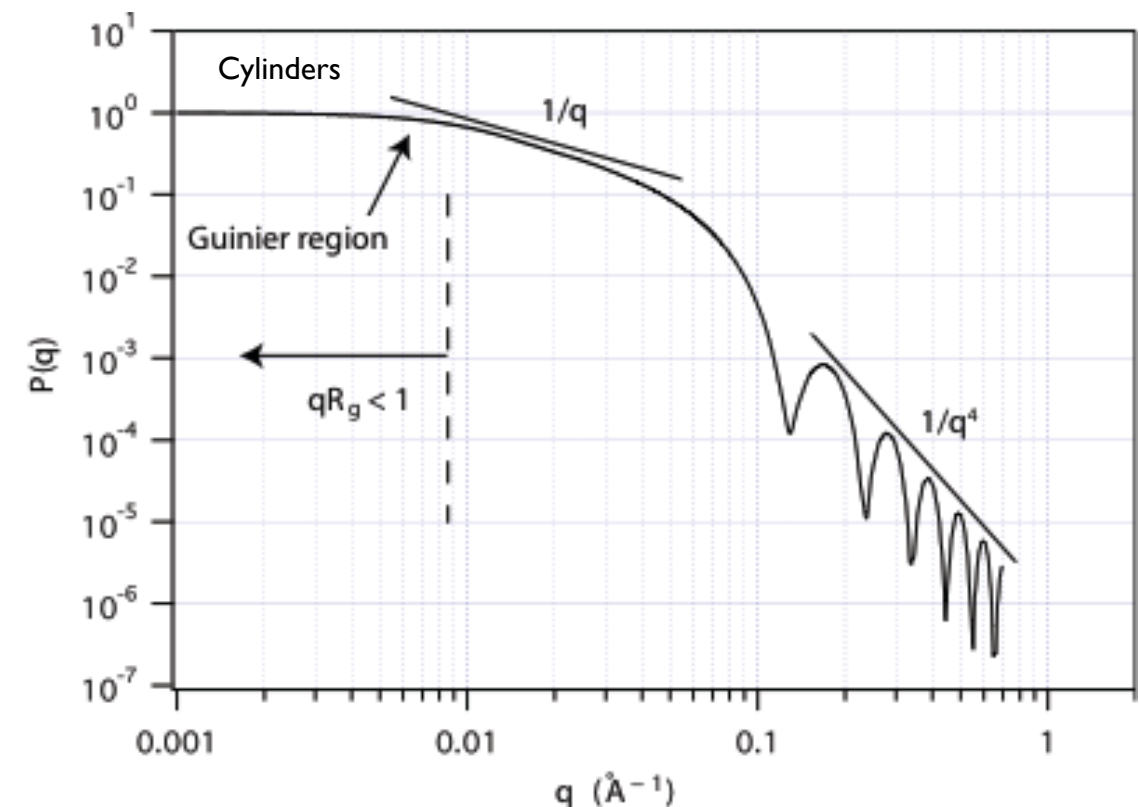
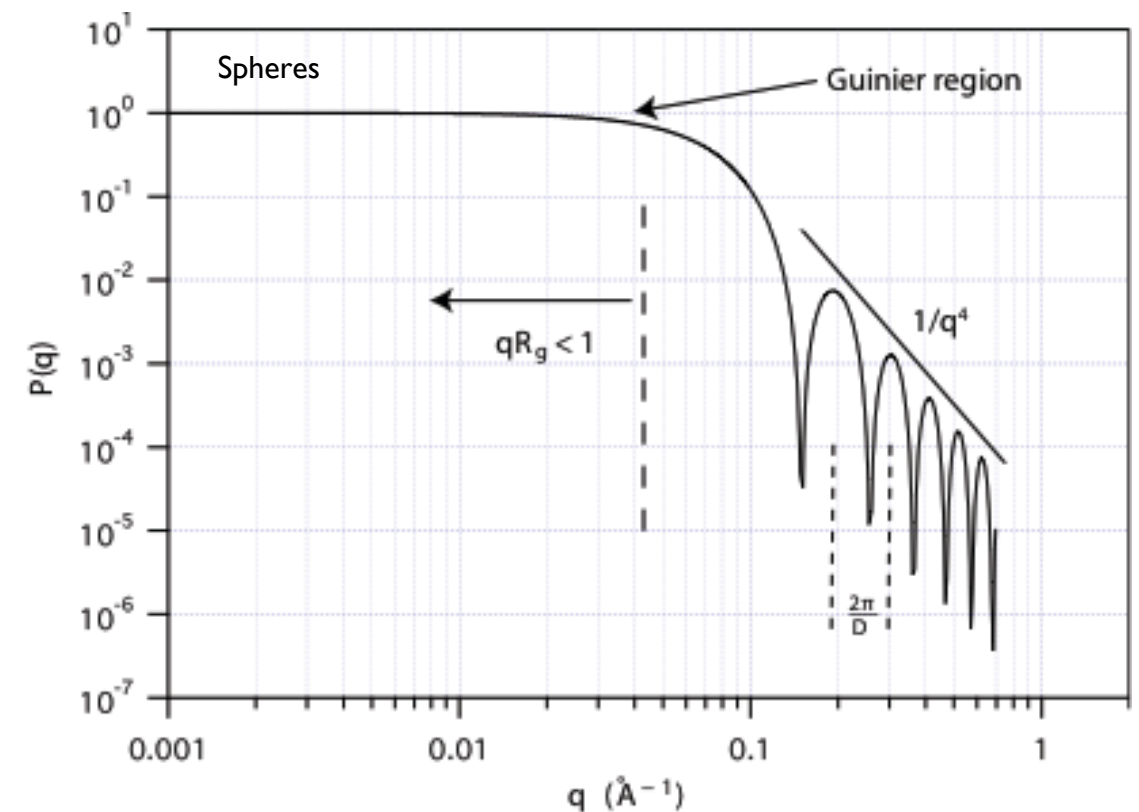


From "The SANS Toolbox" by B. Hammouda

The slope of a log-log plot in a region where the size being examined is smaller than the scattering object is called the "Porod region" and gives information about the local structure.

Porod's Law gives us the surface to volume ratio from the scattering invariant for systems with sharp interfaces:

$$\frac{\pi}{Q^*} \cdot \lim_{q \rightarrow \infty} (I(q) \cdot q^4) = \frac{S}{V}$$



Form and Structure Factors

Lots of form and structure factors have already been calculated



Advances in Colloid and Interface Science
70 (1997) 171–210

ADVANCES IN
COLLOID AND
INTERFACE
SCIENCE

Analysis of small-angle scattering data from colloids and polymer solutions: modeling and least-squares fitting¹

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Abstract

Analysis and modeling of small-angle scattering data from systems consisting of colloidal particles or polymers in solution are discussed. The analysis requires application of least-squares methods, and the basic principles of linear and non-linear least-squares methods are summarized with emphasis on applications in the analysis of small-angle scattering data. These include indirect Fourier transformation, square-root

Form and Structure Factors

Lots of form and structure factors have already been calculated

NIST Center for Neutron Research

NCNR Home Instruments Science Experiments Sitemap

SANS & USANS Data Reduction and Analysis

Data Analysis Using SasView

The recommended tool for analyzing small angle scattering data

Data Reduction Using Igor

An upgrade of the reduction and analysis software is available to our users in order to plan better experiments currently possible.

- All of the SANS and USANS Reduction and analysis software is available.
- Installation package: [NCNR SANS package](#)
- Installation instructions: [Install Instructions](#)
- Watch the installation movie: [Install SASfit](#)
- A Quick Start guide to the package is here ([see the movies](#))

What's New?

- Manuals are included in the download package
 - [SANS Reduction Help File](#) (PDF)
 - [USANS Reduction Help File](#) (PDF)
 - [Data Analysis Help File](#) (PDF)

Visit the main page

SASfit MANUAL

Overview
Get SASfit
Recent changes
Random page
Help

Toolbox
What links here
Related changes
Special pages
Permanent link

Print/export
Categories

Overview

SASfit is a curve fitting program for small-angle scattering curves, mostly for biological chemistry. It features more than 200 particle models and is designed to fit multiple curves simultaneously.

Contents [hide]

- About
- A User guide for SASfit
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About

SASfit calculates integral structural parameters like radius of gyration, scattering invariant, Porod constant. Furthermore it can fit size distributions together with several form factors including different structure factors.

Additionally an algorithm has been implemented, which allows to simultaneously fit several scattering curves with a common set of (global) parameters. This last option is especially important in contrast variation experiments or measurements with polarised neutrons. The global fit helps to determine fit parameters unambiguously which by analyzing a single curve would be otherwise strongly correlated.

The program has been written to fulfill the needs at the Small Angle Neutron Scattering Facility at the Paul Scherrer Institute. The numerical routines have been written in C whereas the menu interface has been written in Tcl/Tk. There are binary packages for Windows, Linux and MacOS (see [Get SASfit](#) and the

SasView for Small Angle Scattering Analysis

A SAS Community Project launched from the NSF DANSE effort

home about SasView links/downloads meetings/workshops people faq help

SasView is a Small Angle Scattering Analysis Software Package, originally developed as part of the NSF DANSE project under the name SansView, now managed by an international collaboration of facilities. Feedback and contributions are welcome and encouraged.

Download The Latest Version of SasView

User Mailing List

Subscribe to the user mailing list (sasview-users@lists.sourceforge.net).

News:

- July 2015: SasView v3.1.0 released Jul 14, 2015.
- February 2015: [SasView Code Camp III](#) being held at ESS DMSC in Copenhagen, Denmark, February 11th - 20th
- September 2014: New optimization package project, BUMPS - with several optimization options, integrated into trunk for beta testing.
- May 2014: SasView v3.0.0 released May 21, 2014.
- April 2014: [SasView Code Camp II](#) held at ISIS/Diamond from March 31 through April 6, 2014.
- October 2013: SasView team was working on a version 3.0 release.
- April 2013: The first [SasView developers workshop](#) was held at NIST in Gaithersburg Maryland April 2-7, 2013
- November 2012: SasView was presented at the international [Small Angle Scattering meeting](#) in Sydney Australia, Nov. 18-23, 2012.
- November 2012: The first official release under the [SasView](#) label is launched. This marks the official graduation from a DANSE incubator project to a long term collaborative maintenance and development project

screenshot of multiple/global fitting

diamond OAK RIDGE National Laboratory NIST NEUTRONS FOR SCIENCE ISIS Science & Technology Facilities Council ESS UFR

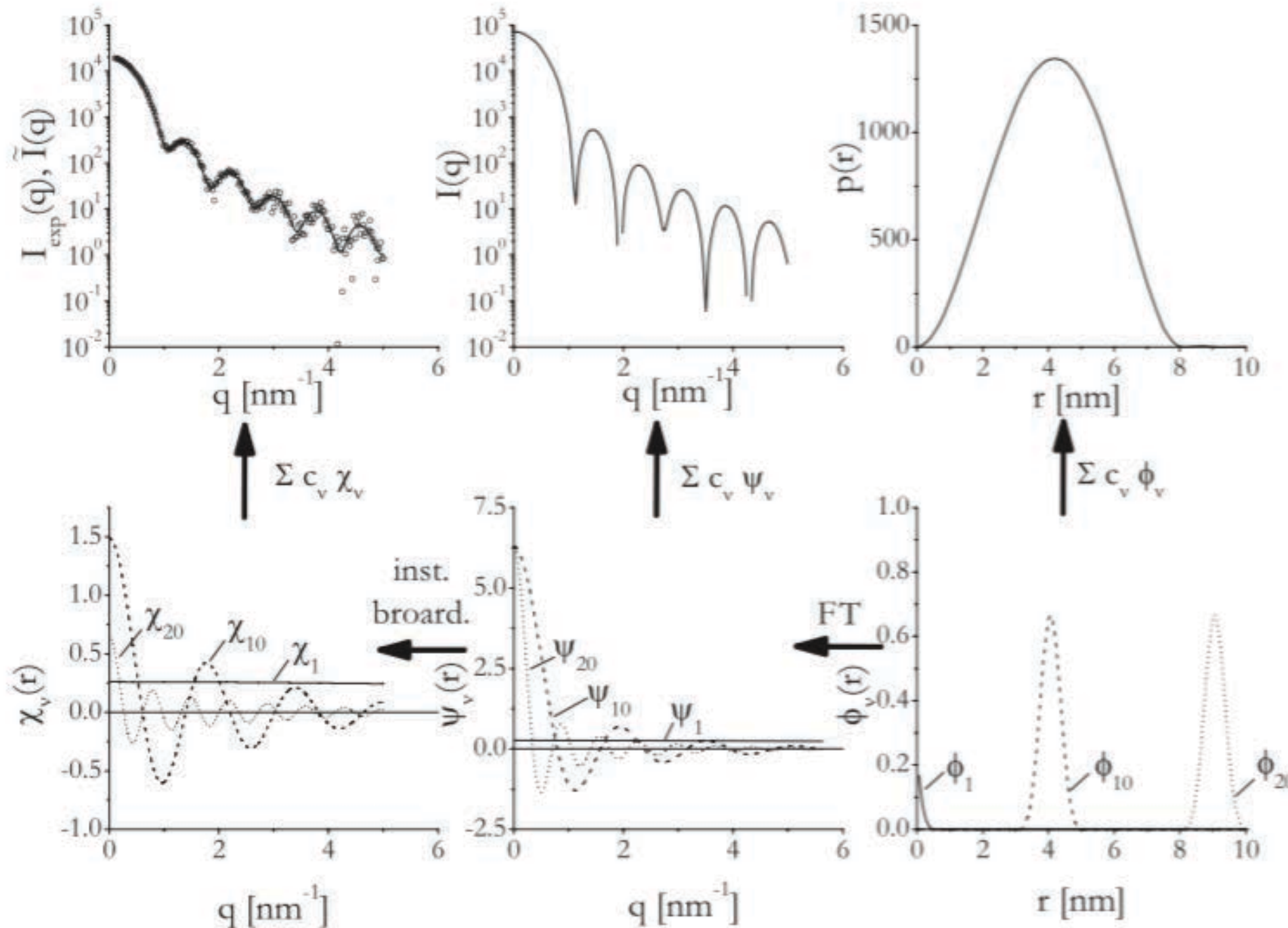
SasView : <http://www.sasview.org>

SASFit : <https://www.psi.ch/sinq/sansii/sasfit>

NIST Igor : http://ncnr.nist.gov/programs/sans/data/red_anal.html

... and coded into software.

Indirect Fourier Transform



Glatter, O. (1977) New Method for Evaluation of Small Angle Scattering Data, *Journal of Applied Crystallography*, 10, 415–421.

Fritz, G., & Glatter, O. (2006) Structure and interaction in dense colloidal systems: evaluation of scattering data by the generalized indirect Fourier transformation method *New Journal of Physics*, 18, S2403–S2419

Ab-Initio Structure Generation

Data analysis software ATSAS 2.7.1

A program suite for small-angle scattering data analysis from biological macromolecules

Data processing

[PRIMUS](#) - manipulations with experimental 1D SAS data

[GNOM](#) - indirect transform program that evaluates the particle distance distribution function $p(r)$

[Data manipulation and analysis tools](#) - AUTORG, ALMERGE, DATGNOM, DATPOROD etc.

Ab initio methods

[DAMMIN](#) - *ab initio* shape determination using a dummy atom model

[DAMMIF](#) - rapid shape determination

[GASBOR](#) - reconstruction of a protein structure by a chain-like ensemble of dummy residues

[MONSA](#) - shape determination using a multiphase dummy atom model

Rigid body modelling

[SASREF](#) - modelling of multisubunit complexes

[BUNCH](#) - modelling of multidomain proteins against multiple data sets

[CORAL](#) - modelling of multidomain protein complexes against multiple data sets

[MASSHA](#) - interactive modelling of atomic structures and shape analysis

[GLOBSYMM](#) - rigid body modelling of symmetric oligomers

Mixtures and flexible systems

[OLIGOMER](#) - volume fractions of mixtures with known scattering intensities from the components

[MIXTURE](#) - modelling of multicomponent systems

[EOM](#) - Ensemble Optimization Method for flexible proteins

[SREFLEX](#) - flexible refinement of high-resolution models combining SAXS and NMA

PDB oriented tools

[CRY SOL](#) - X-ray scattering patterns from known hi-res structures

[CRYSON](#) - neutron scattering patterns from known hi-res structures

[SUPCOMB](#) - superimposes one 3D structure onto another

[DAMAVAR](#) - align *ab initio* models, select the most typical one

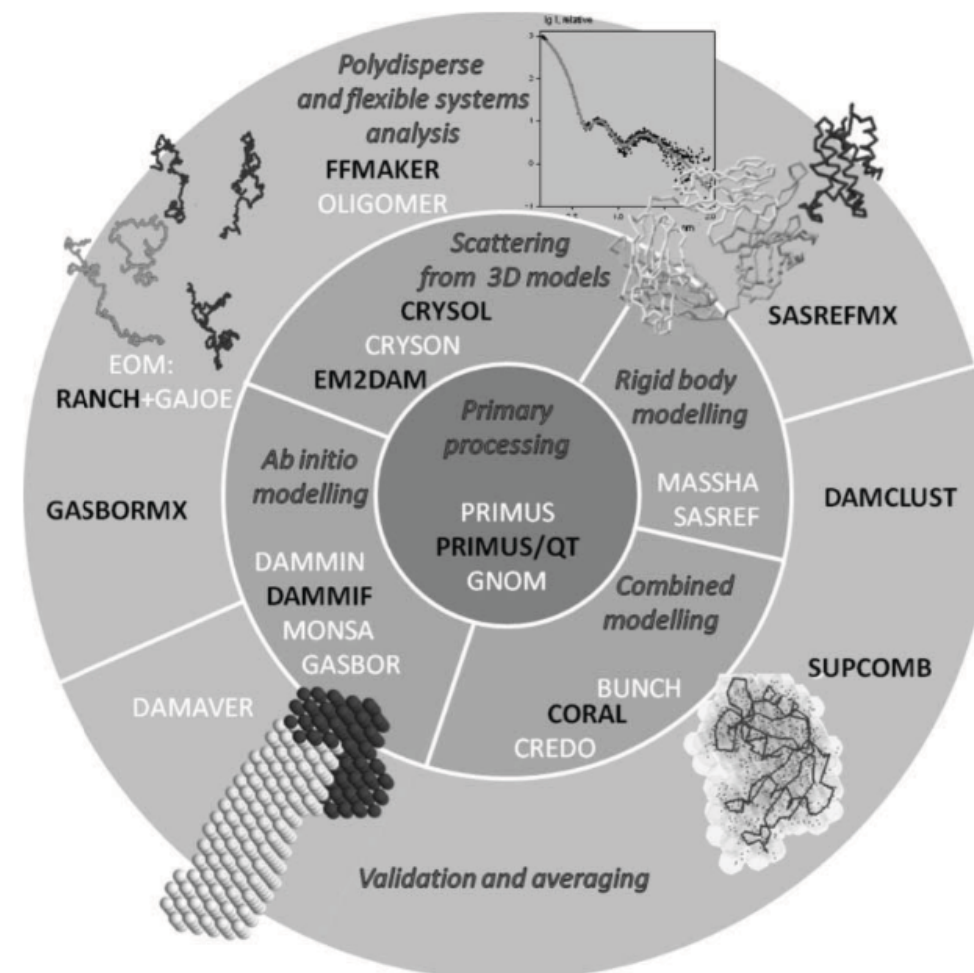
Manuals

If you use ATSAS please cite:

Petoukhov, M.V., Franke, D., Shkumatov, A.V., Tria, G., Kikhney, A.G., Gajda, M., Gorba, C., Mertens, H.D.T., Konarev, P.V. and Svergun, D.I. (2012)

[New developments in the ATSAS program package for small-angle scattering data analysis](#)

J. Appl. Cryst. 45, 342-350 © International Union of Crystallography [DOI](#)



Philosophy of Data Analysis

1) Look at the data ...

- Trends
- Shape

... do they match your expectation?

... if not, why not?

2) Extract model free information ...

- R_g (Guinier)
- Slopes (Porod)
- $I(0)$

3) Consider model fitting ...

- Which model?
- If model doesn't fit, don't just add more parameters!
- How much information is in my data?

4) Consider Indirect Fourier Transform ...

- Can give hints as to why model isn't fitting
- If possible use deconvolution to SLD profile
- Use as a basis for developing parameterised model fit

5) Consider Ab Initio methods ...

- Do you have a known structure to compare to?

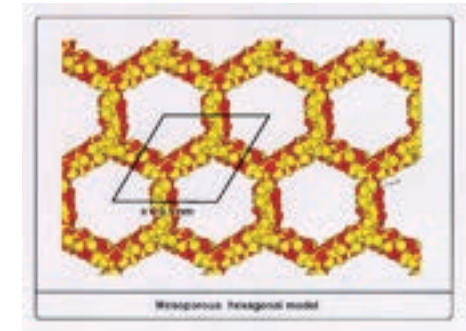
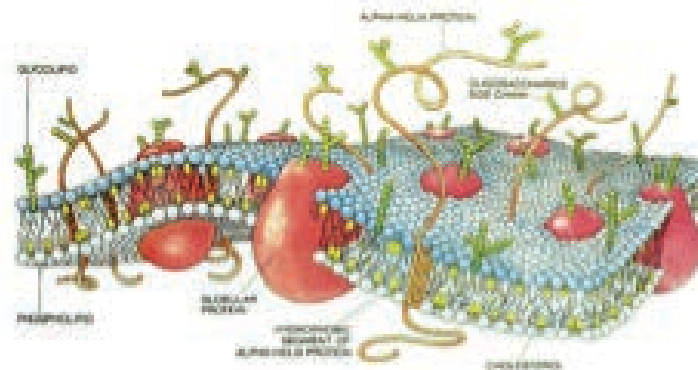
Applications of SANS

Andrew Jackson

NNSP-SwedNess Neutron School 2017, Tartu

Lecture L9.3

SANS is Versatile



Mesoporous structures

Biological structures (membranes, vesicles, proteins in solution)

Polymers

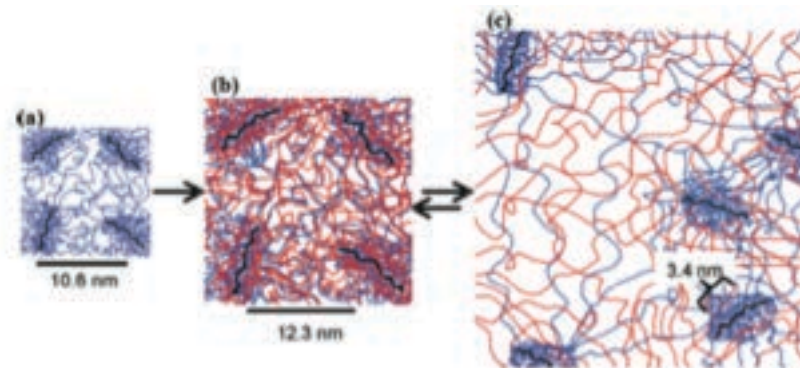
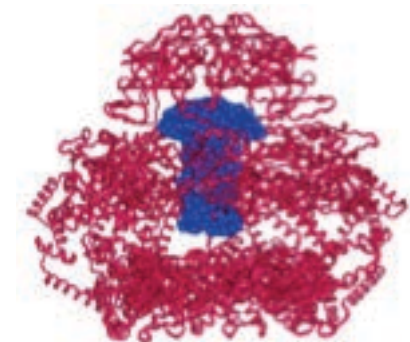
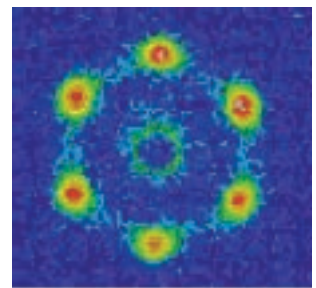
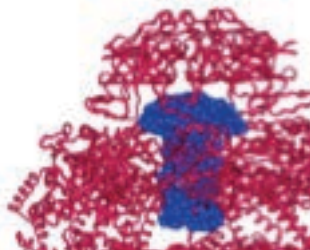
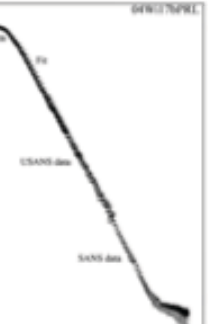
Colloids and surfactants

Magnetic films and nanoparticles

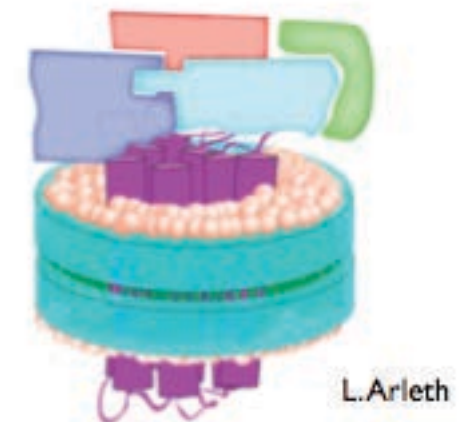
Superconductors

Voids and Precipitates

Geology



Waters et al (2011) Macromolecules 44 5776

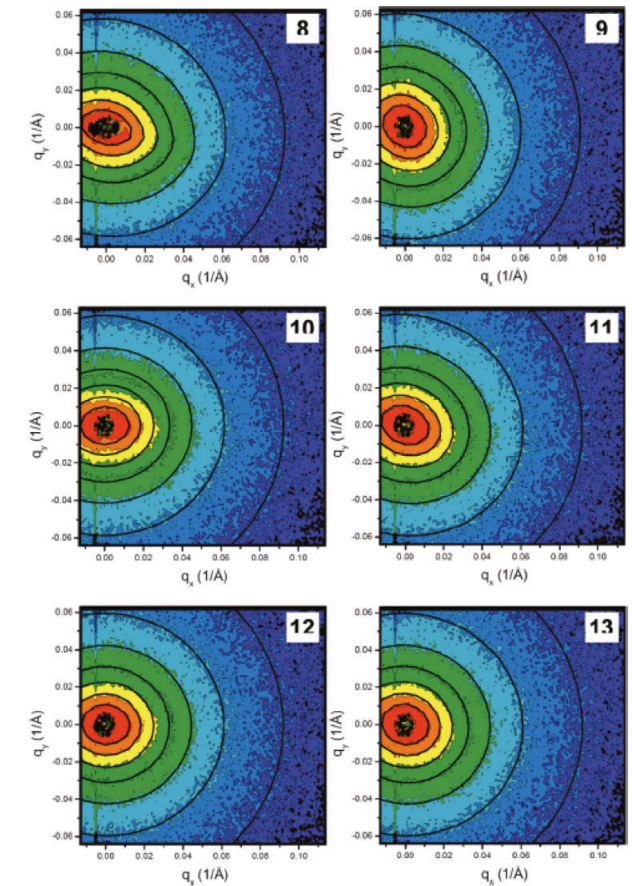
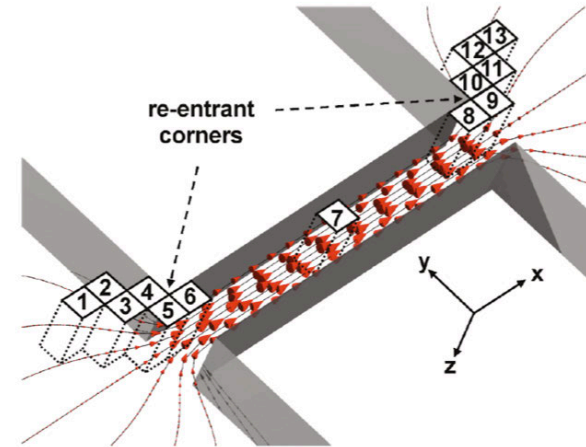


Fluid Flow

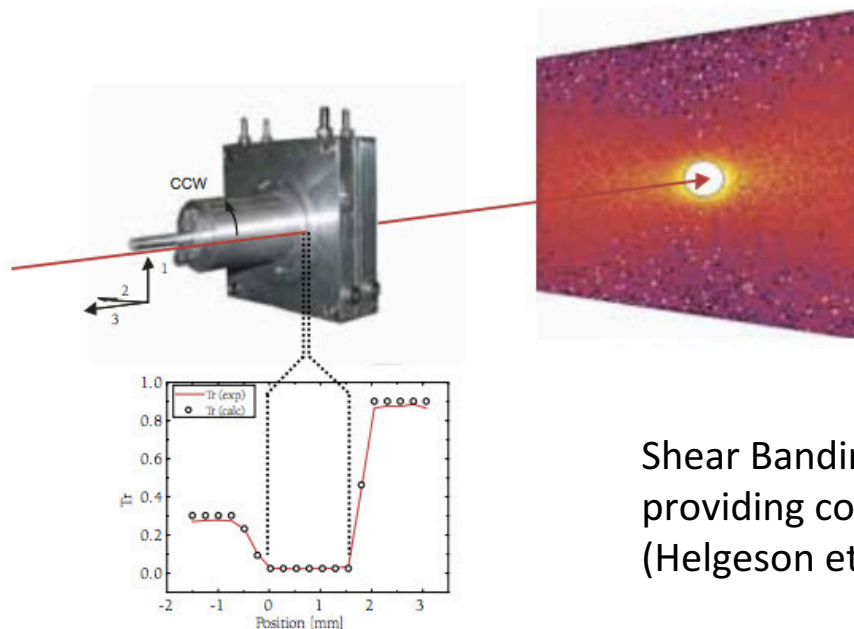
The flow of **complex fluids** through **complex geometries** is relevant to many industrial processes including polymer processing and oil recovery.

Microfluidic devices are increasingly used for chemical and pharmaceutical discovery, production and processing.

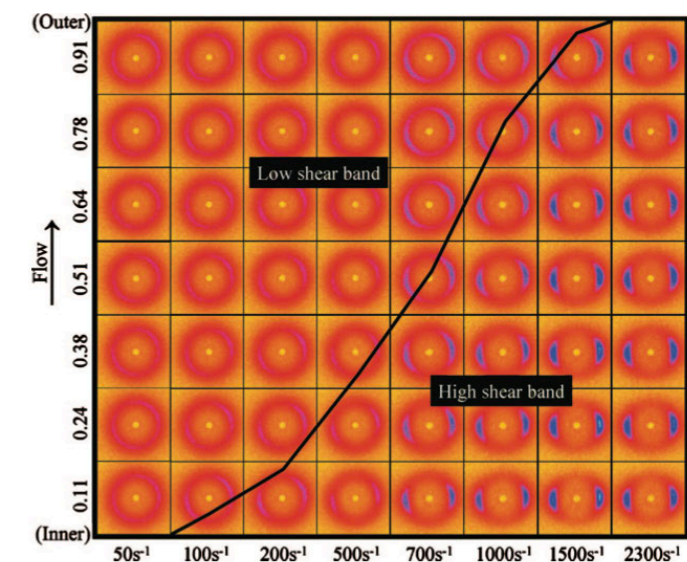
There is a need to understand **structural effects of flow** both for practical purposes and to compare with fluid flow models.



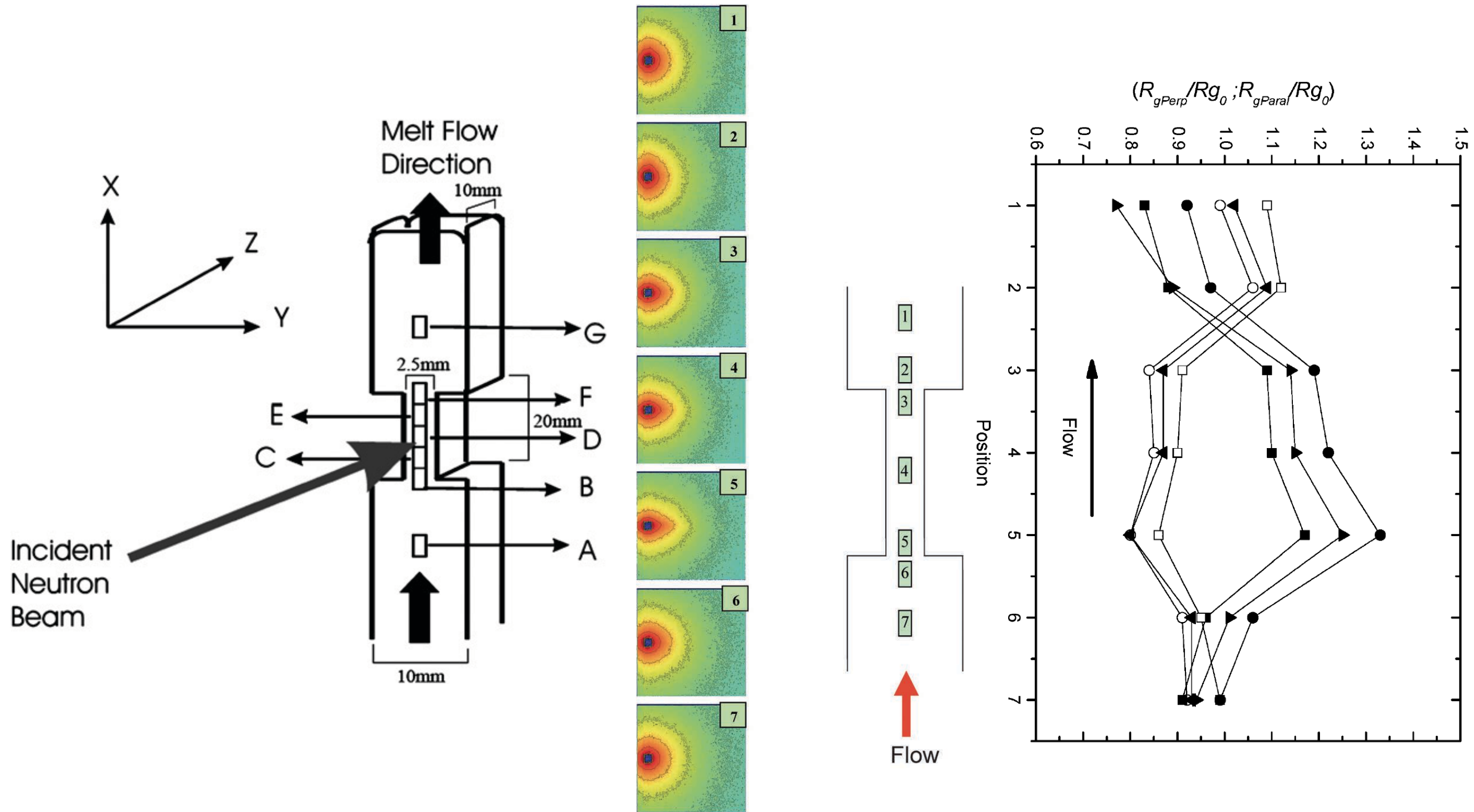
Measuring the deformation of polymer chains allowing development of new models of polymer flow (Clarke et al. (2010) *Macromolecules* 43, 1539)



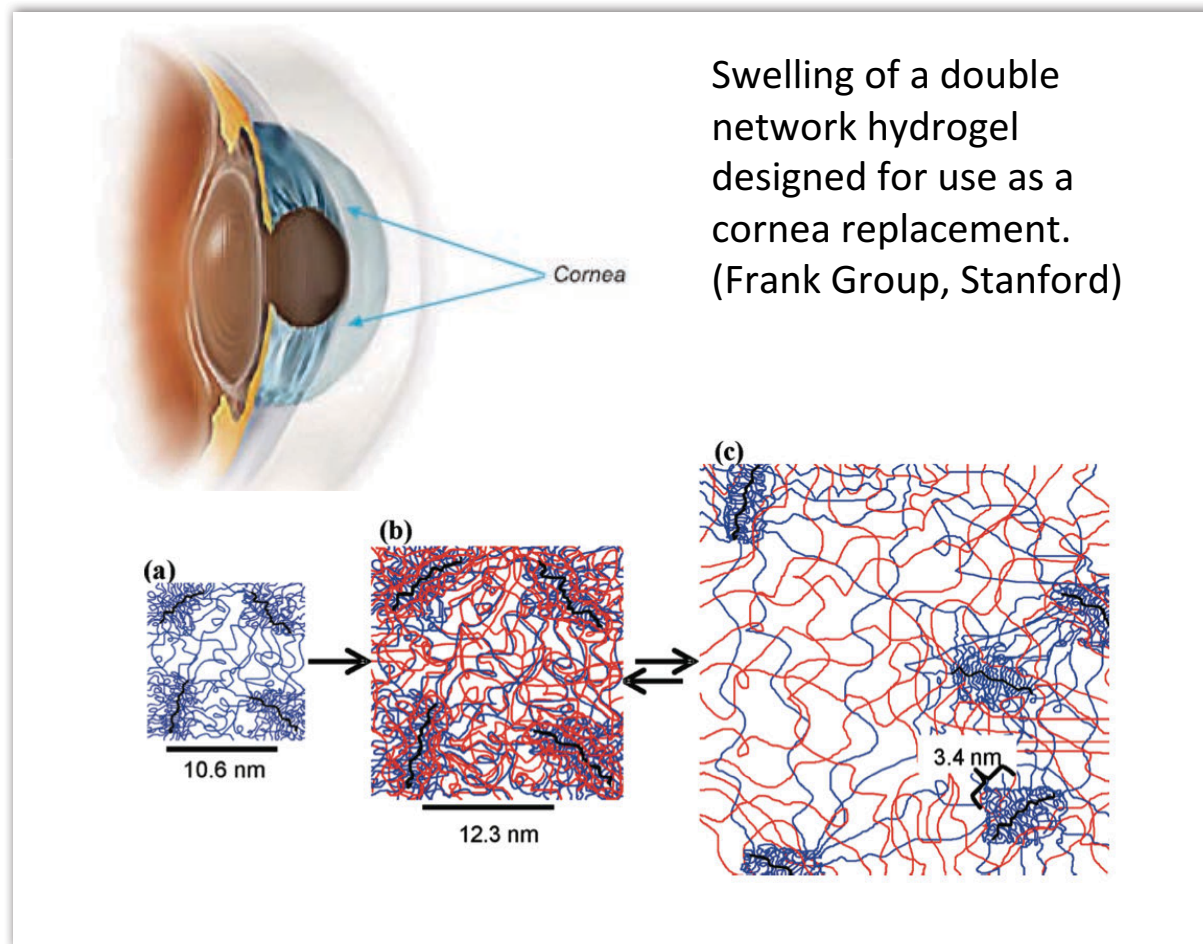
Shear Banding in CTAB wormlike micelles providing confirmation of rheological model. (Helgeson et al. (2009) *J. Rheol* 53, 727)



Flow mapping in polymer melts

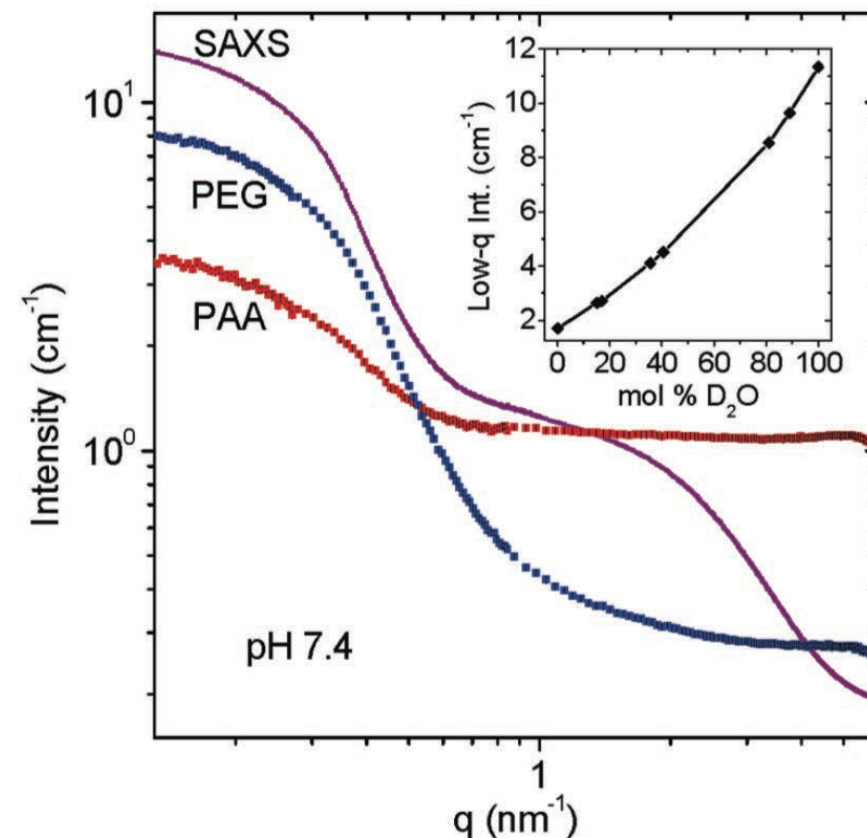
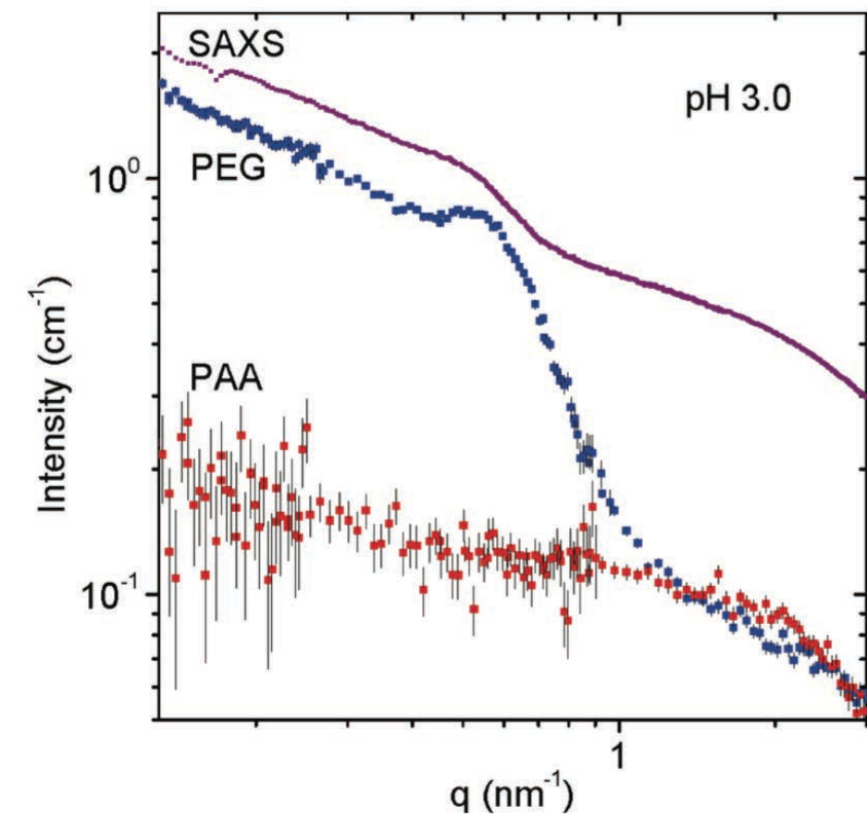


Double Network Hydrogels

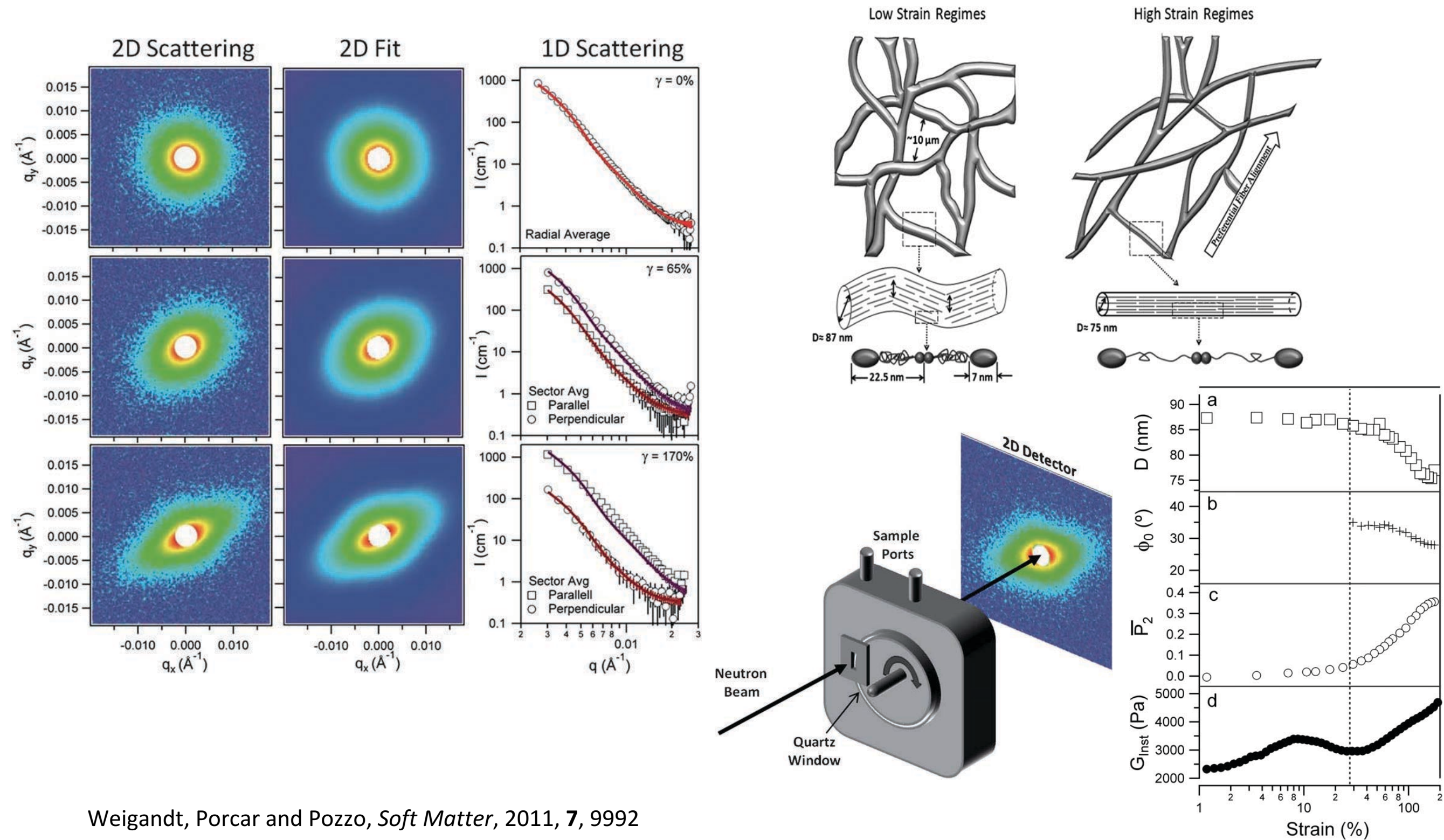


Double network hydrogels provide strength and resilience together with high water content.

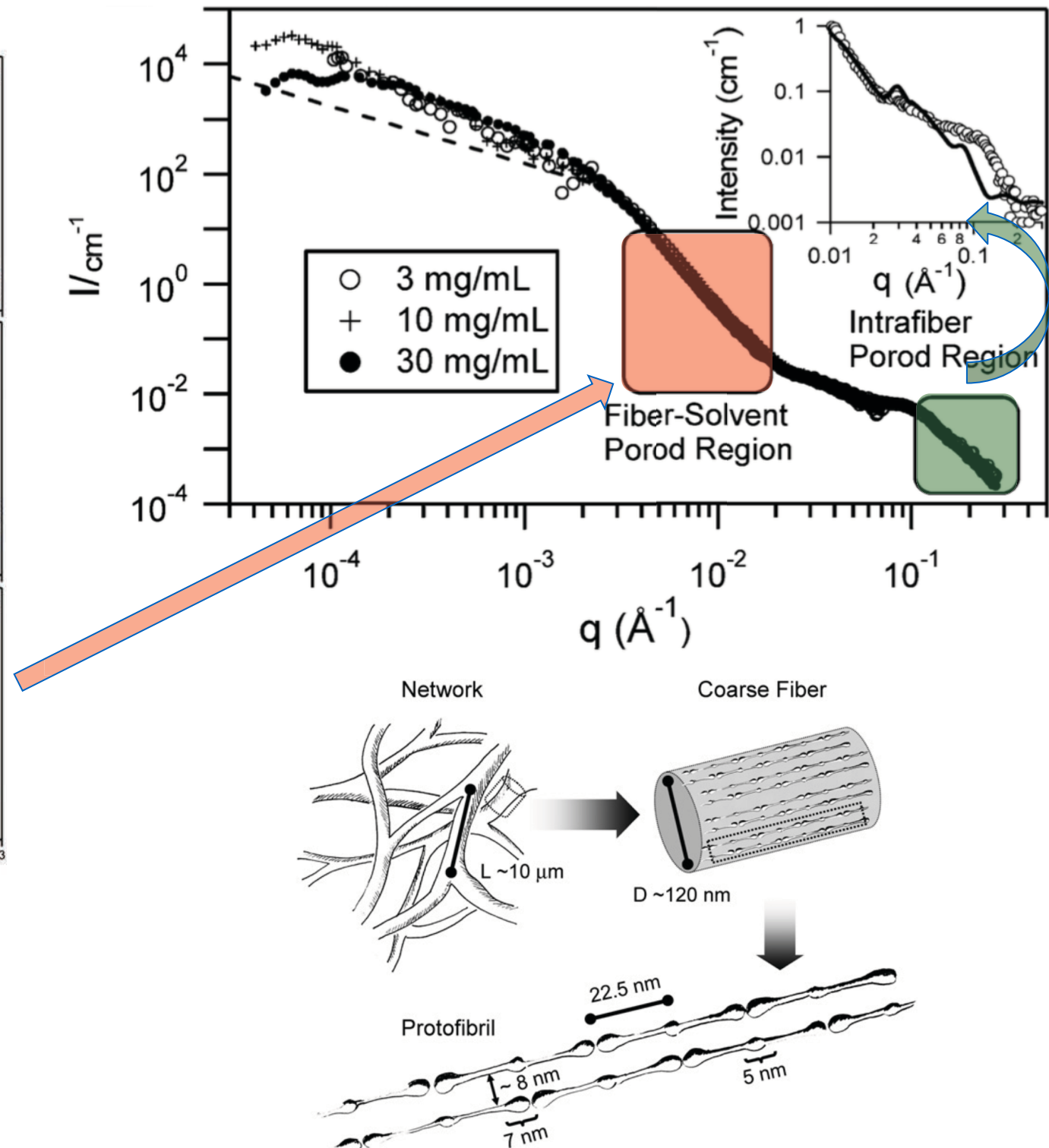
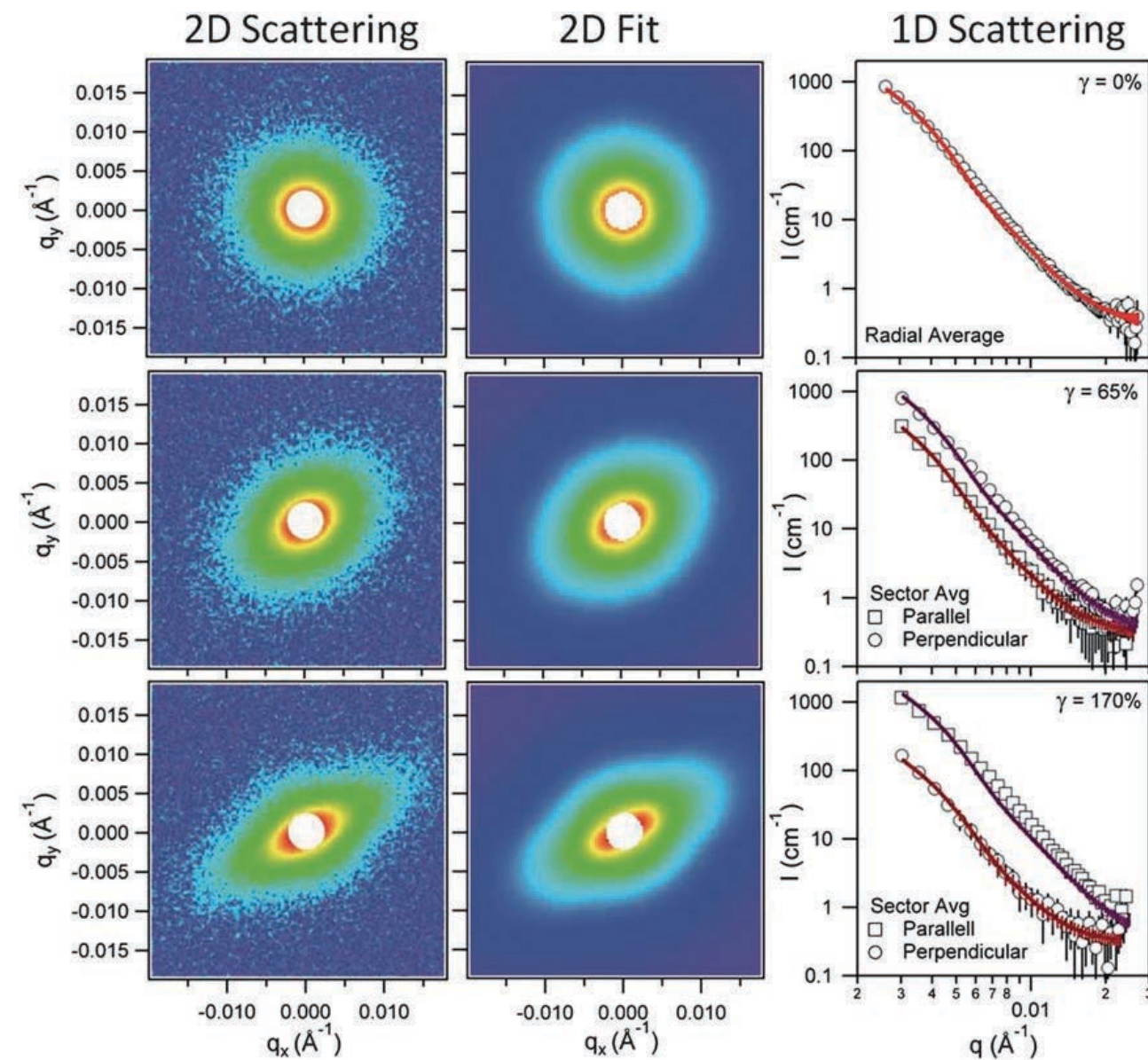
Neutrons provide the structure of each component in the presence of the other. But contrast matching can lead to weak scattering signals and incoherent scattering can be a problem.



Structure of gel networks under shear



Structure of gel networks under shear



Weigandt, Porcar and Pozzo, *Soft Matter*, 2011, **7**, 9992
 Weigandt, Pozzo and Porcar, *Soft Matter*, 2009, **5**, 4321

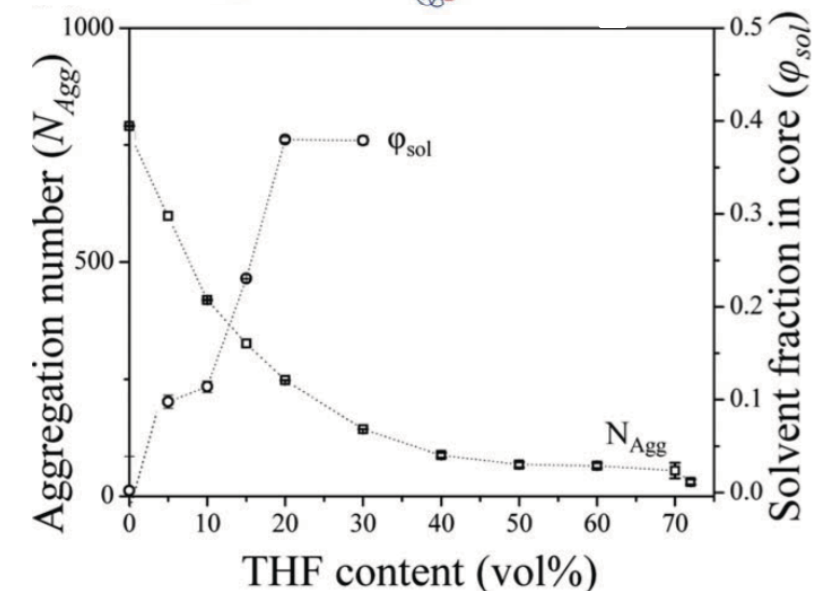
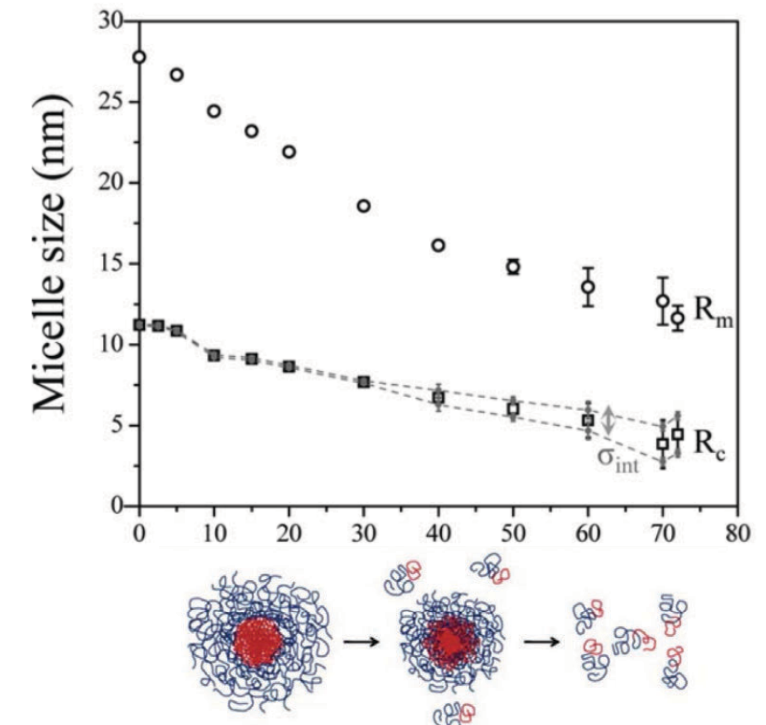
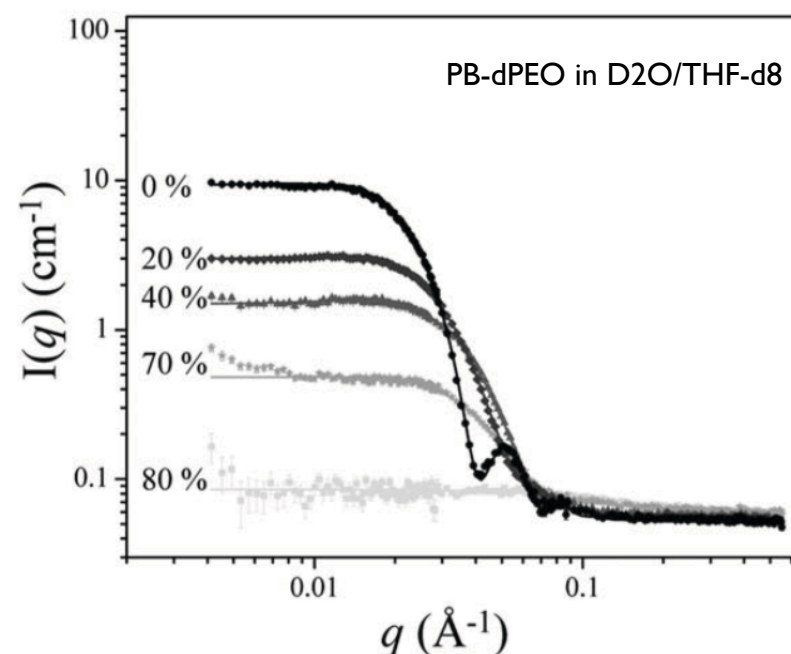
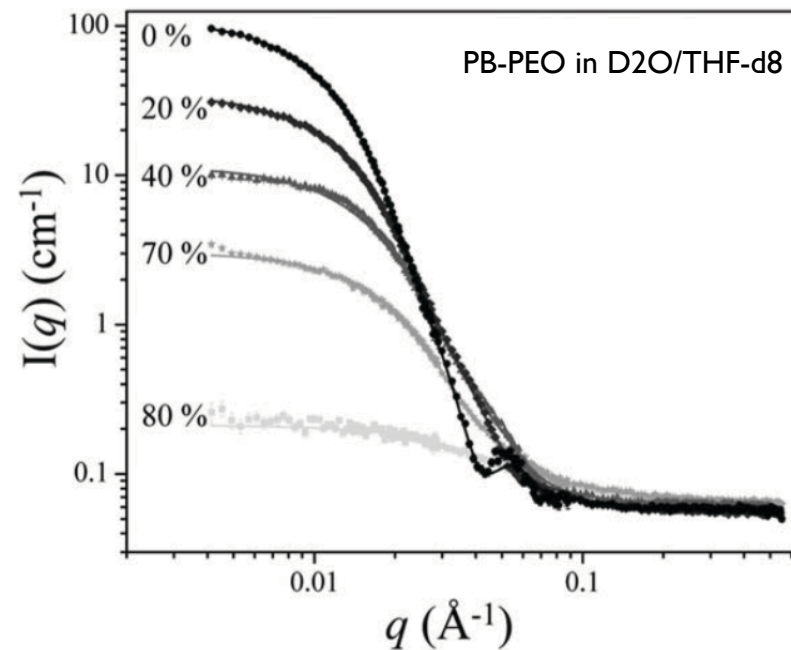
Block Copolymers

Amphiphilic block copolymer assemblies are good candidates for targeted drug delivery applications due to their synthetic versatility, stability, and ability to encapsulate therapeutic molecules

Solvent interactions can be used to tune the structure of micelles formed by block copolymers – here poly(1,2-butadiene-*b*-ethylene oxide) [PB-PEO].

Contrast variation (solvent and polymer) SANS revealed the structural variation seen as the water-THF ratio was varied.

This control of structure allows for micelles to be designed to hold specific drugs through functionalization of the butadiene block.



Stopped-flow SANS - in-situ vesicle formation

Disc to vesicle transition: 50-100ms shots repeated 10-25 times (1ms mixing time)

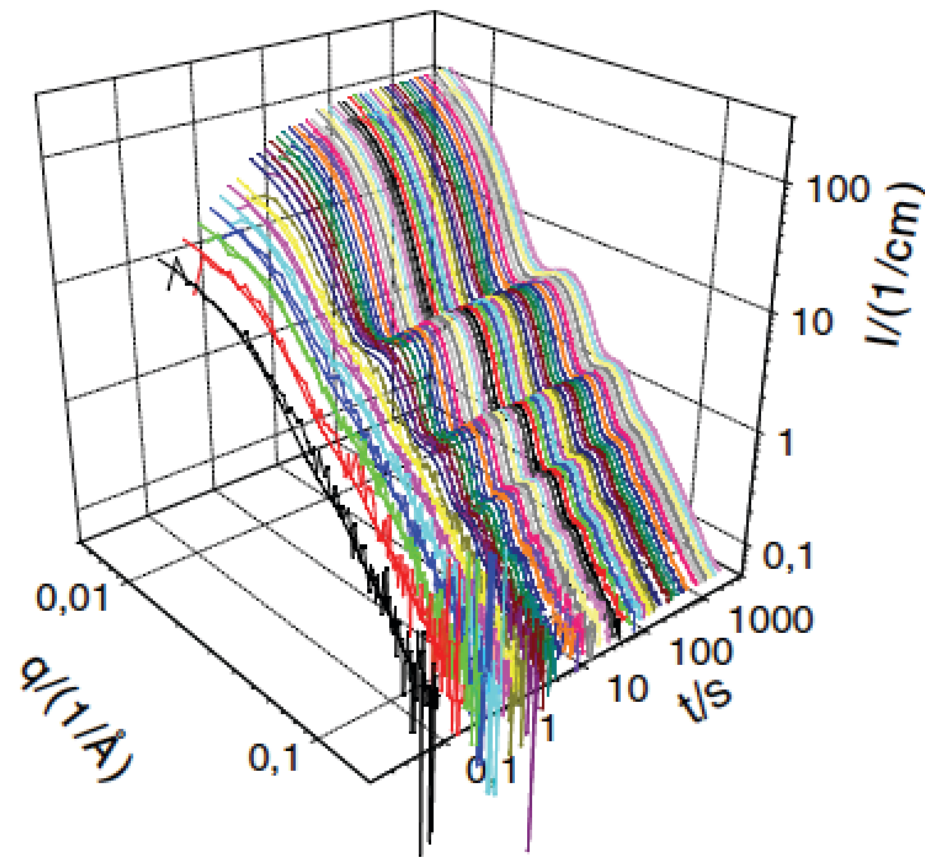


Fig. 6 Time-resolved SANS spectra for the case of mixing 50 mM TDMAO with 50 mM LiPFOS to yield a final composition with $x(\text{TDMAO})=0.525$ mixing ratio at 25 °C (fits are included as *solid lines*)

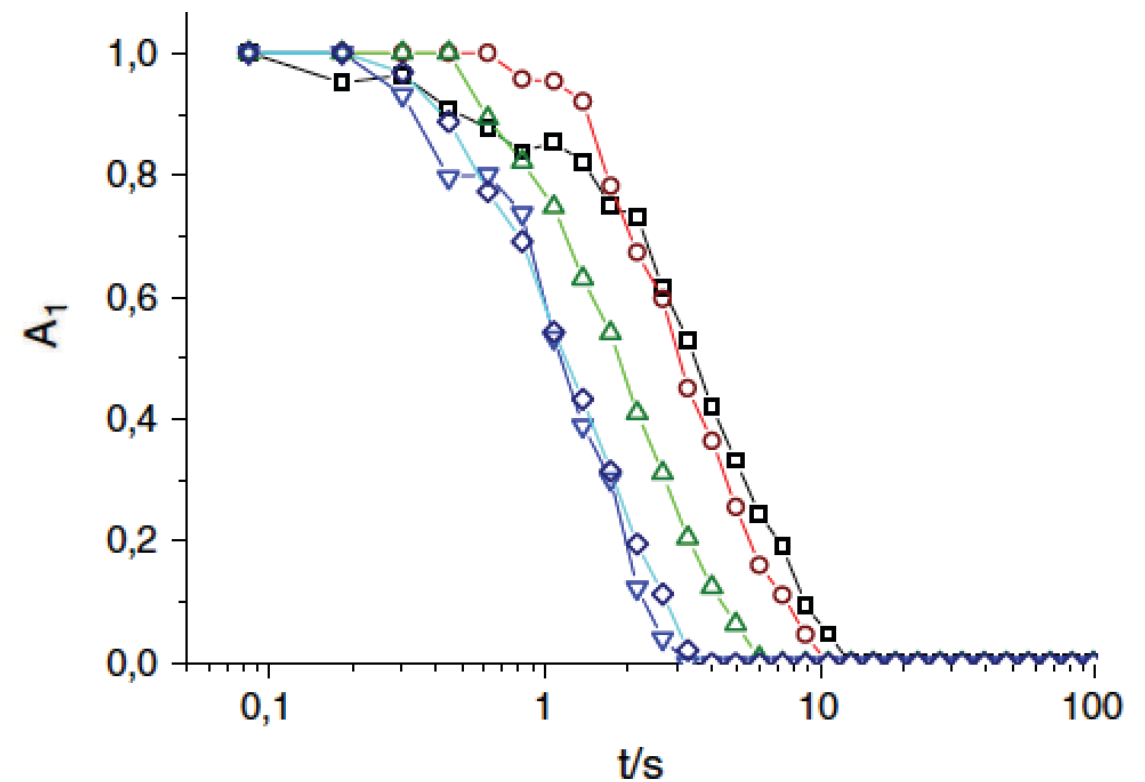
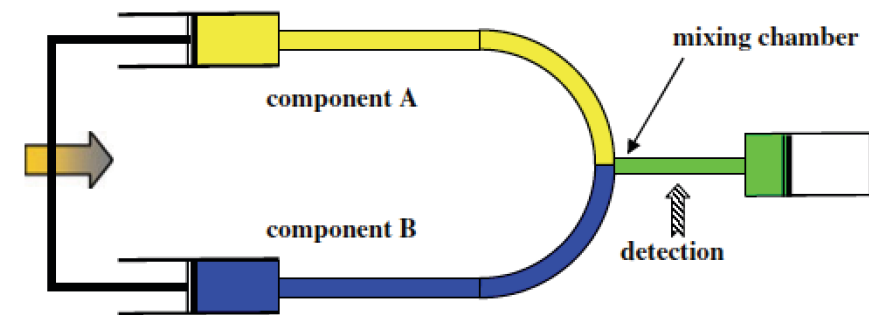
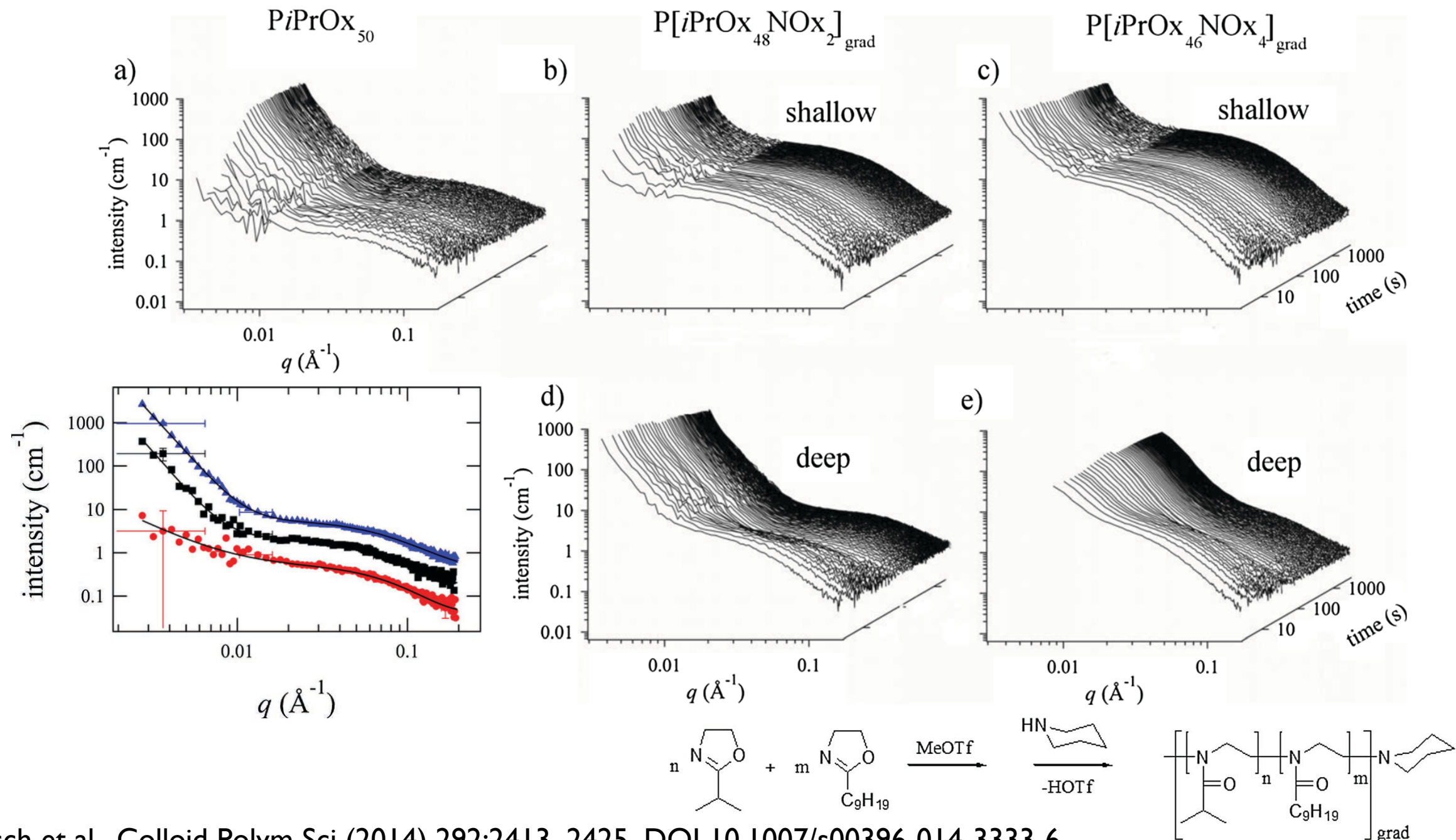


Fig. 9 The relative content A_1 of disks as a function of time in the mixtures with vesicles for various mixing ratios of TDMAO/LiPFOS at 25 °C; $x(\text{TDMAO})$ in the mixtures was *open square*, 0.5; *open circle*, 0.525; *open up-pointing triangle*, 0.55; *open down-pointing triangle*, 0.6; *open diamond*, 0.65

T-jump TR-SANS on poly(2-oxazoline) gradient copolymers

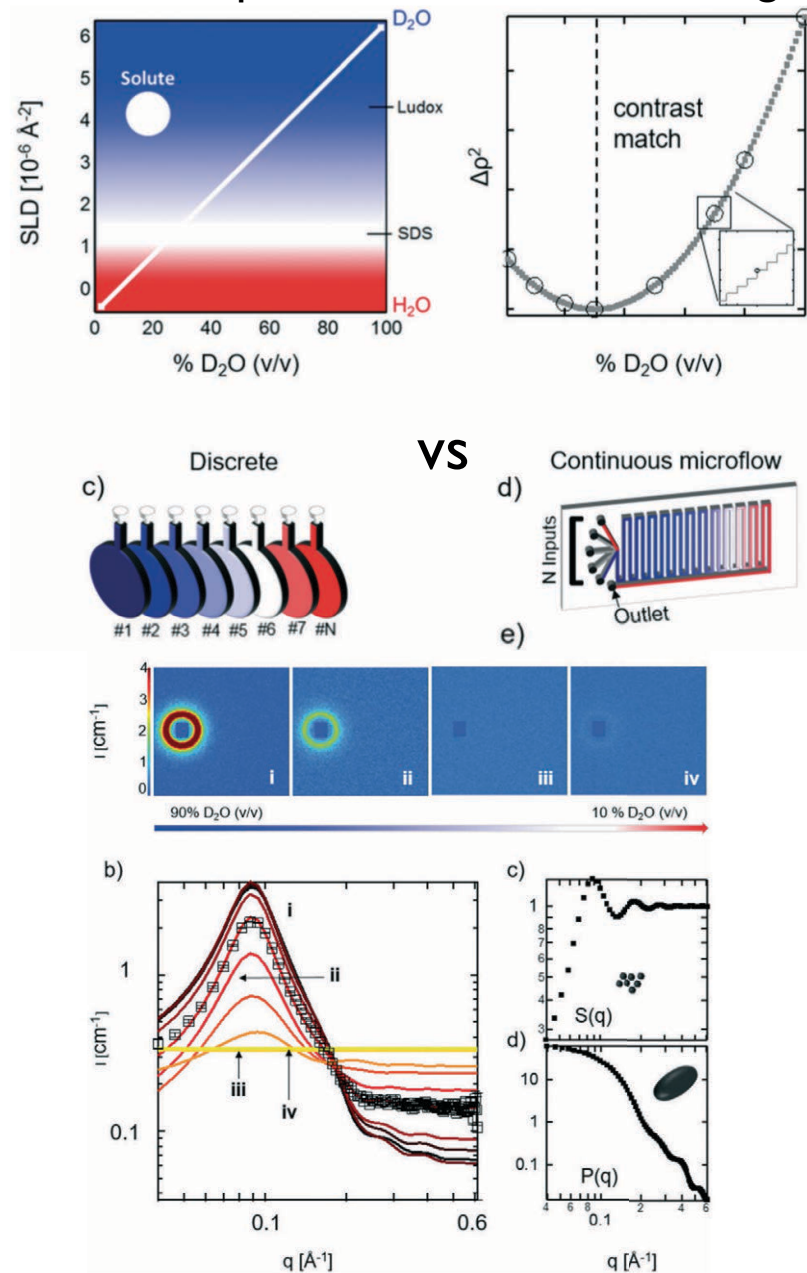
Sample injection into pre-heated cuvette.



Microfluidic SANS

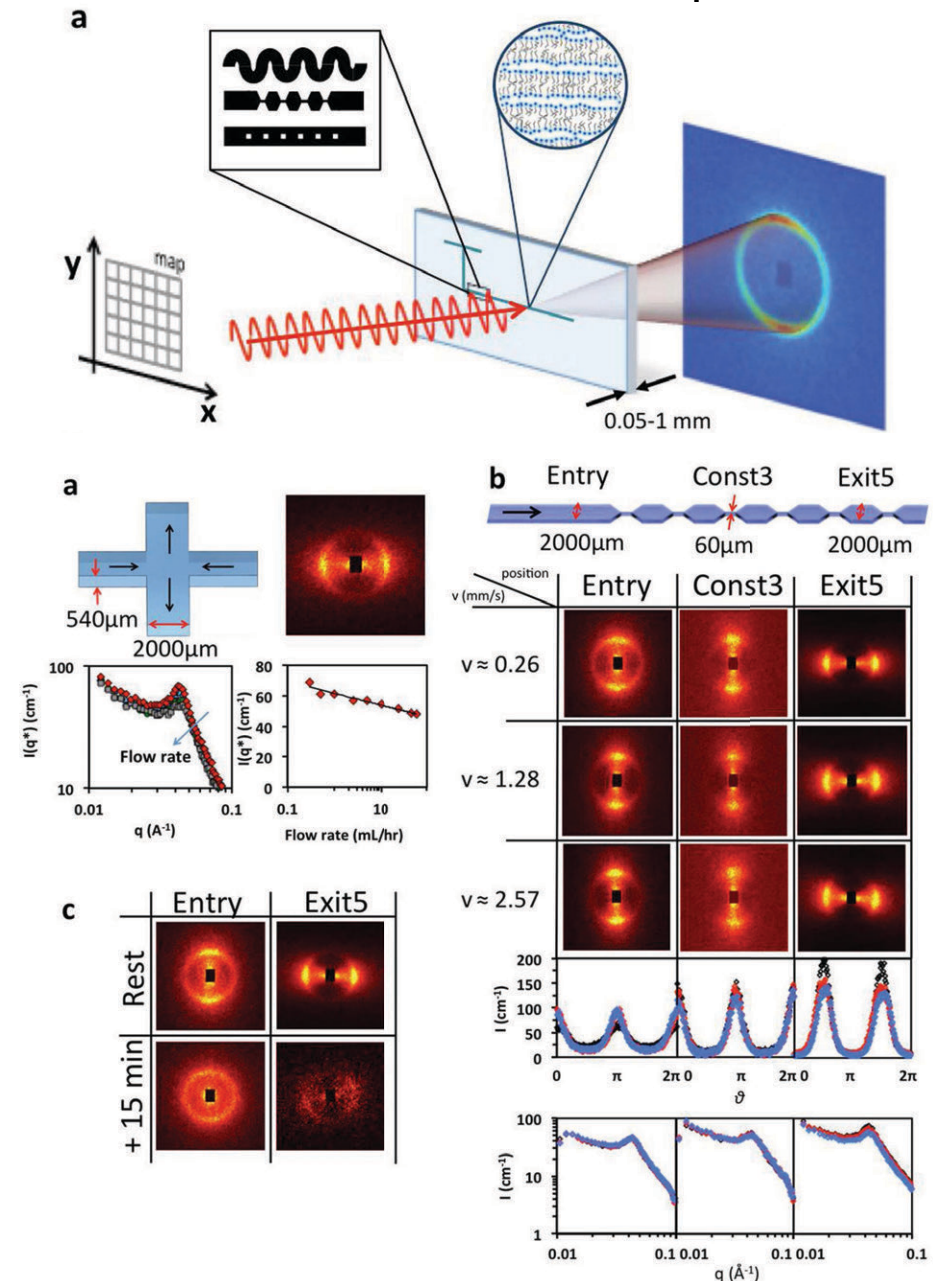
High Throughput Mixing & Tailored Flow Geometry

400 μm channels, 12 mm beam avg. over 20 channels
5s acquisition, continuous mixing



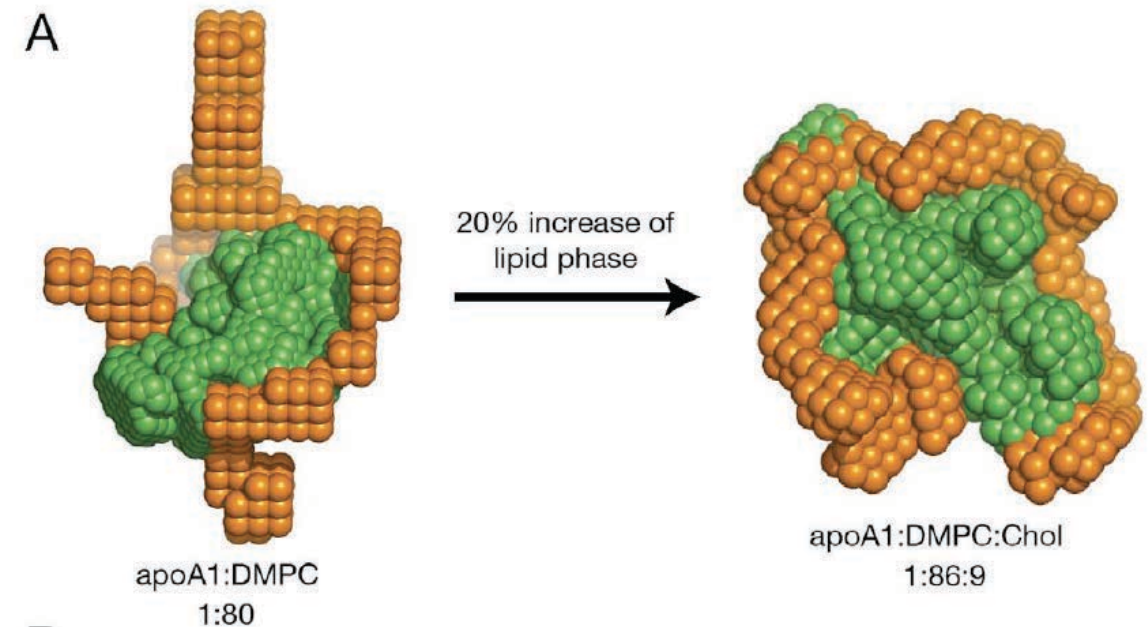
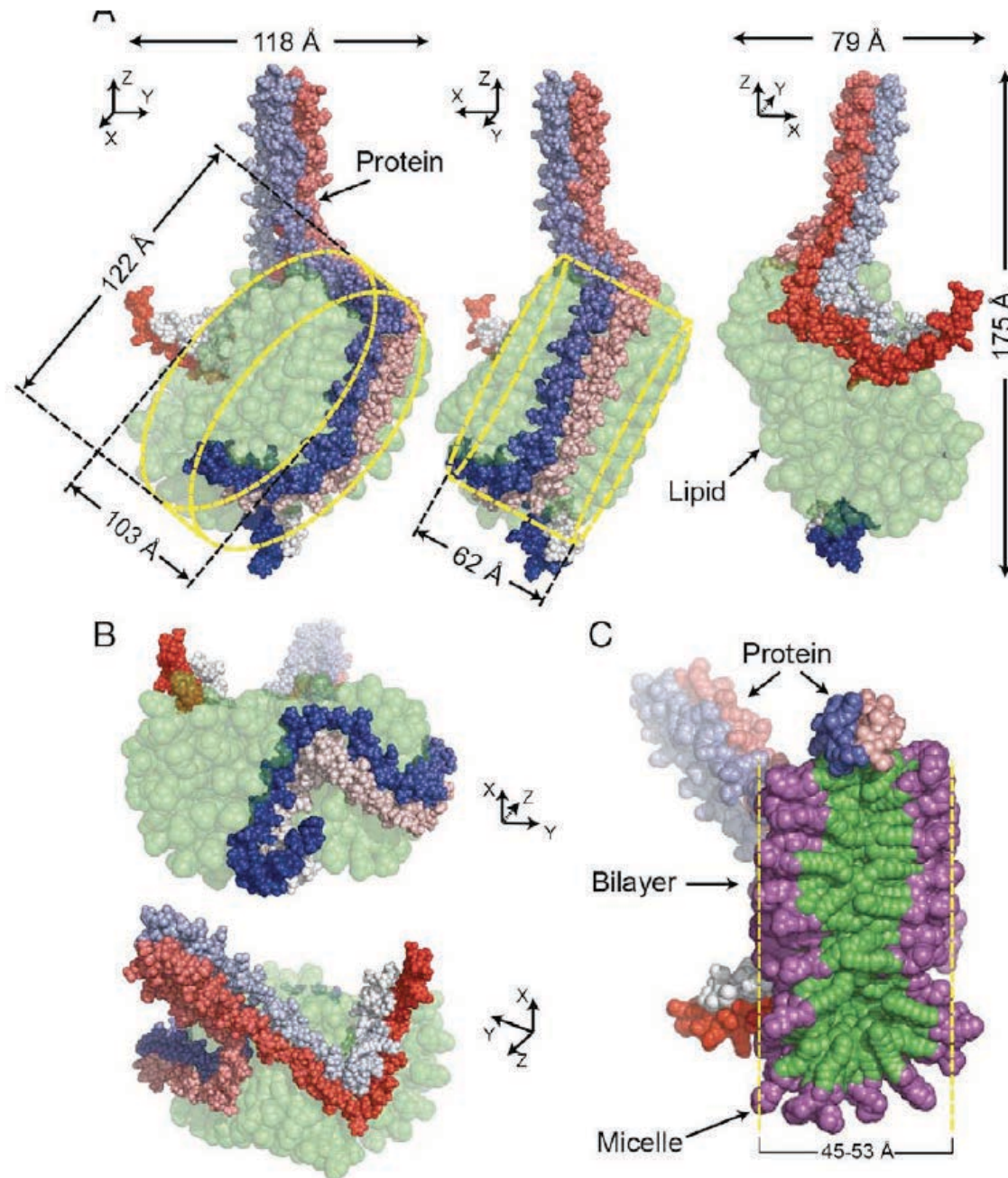
Adamo, M., Poulos, A. S., Miller, R. M., Lopez, C. G., Martel, A., Porcar, L., & Cabral, J. T. (2017). *Lab Chip*, 17(9), 1559–1569.

60 μm channels, 500 μm beam
0.01 – 0.3 \AA^{-1} – 1s – 5 min. acquisitions



C.G. Lopez, T. Watanabe, A. Martel, L. Porcar, J.T. Cabral, Scientific Reports, 5 (2015) 7727.

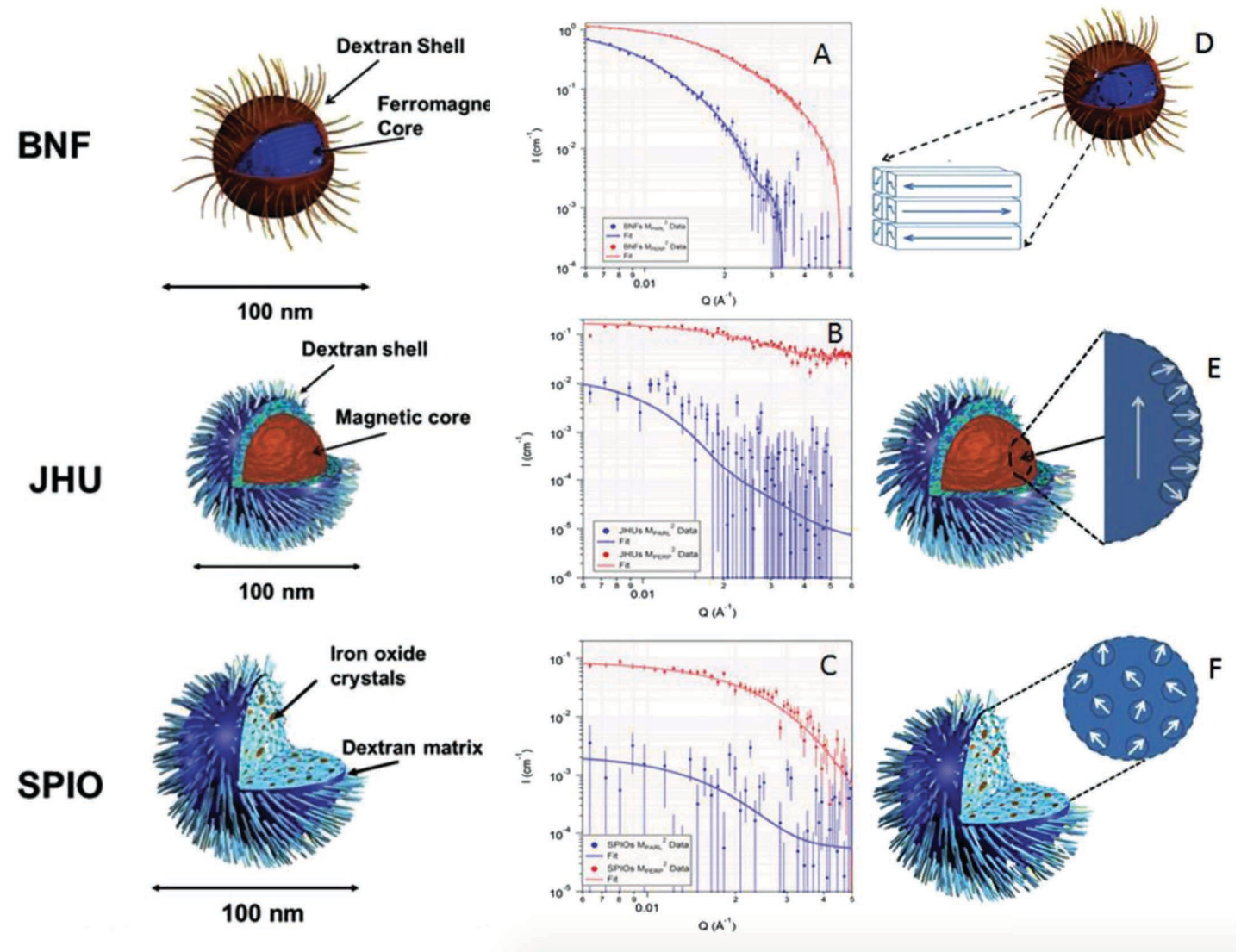
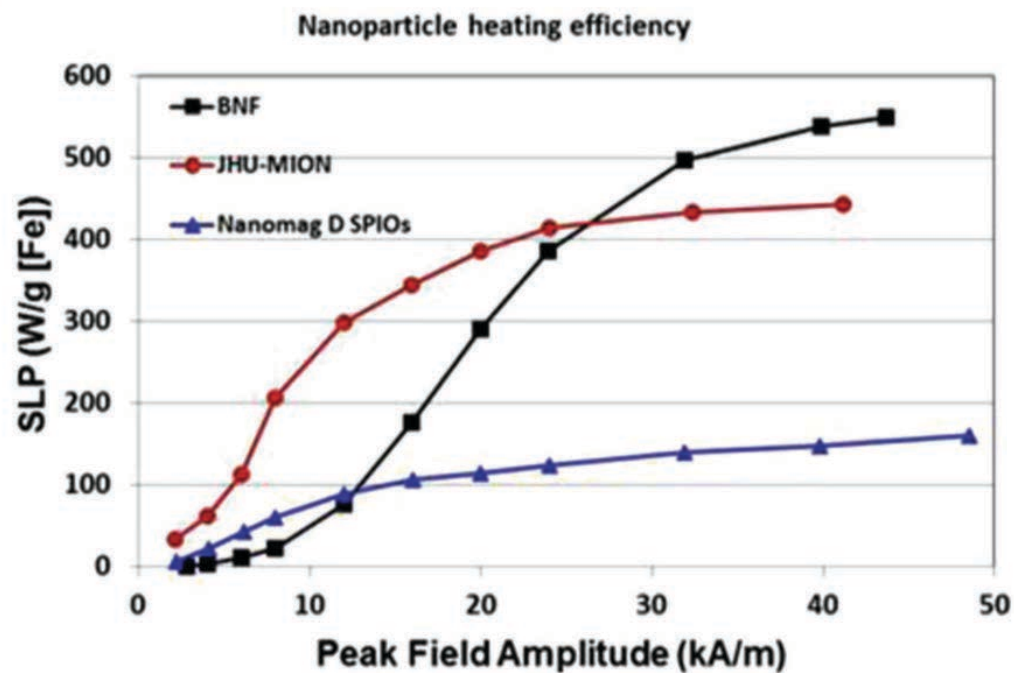
Small Angle Neutron Scattering of High Density Lipoprotein in solution



The low resolution structure of nascent high density lipoprotein with and without cholesterol reveals a mechanism for particle expansion

Hyperthermia Treatment of Cancer

- Use magnetic nanoparticles with alternating magnetic field to locally heat tumour
- Structure of the nanoparticles has strong impact on heating response
- Use polarized neutron scattering to understand these differences



Summary

SANS is a versatile method for studying structure on the nano- to micro- scale

A wide range of scientific areas can be studied

The use of contrast variation methods is key to maximising the information from
SANS

Sample environment equipment for SANS is varied and may need to be designed
or modified for your experiment

Questions?