SASVIEW

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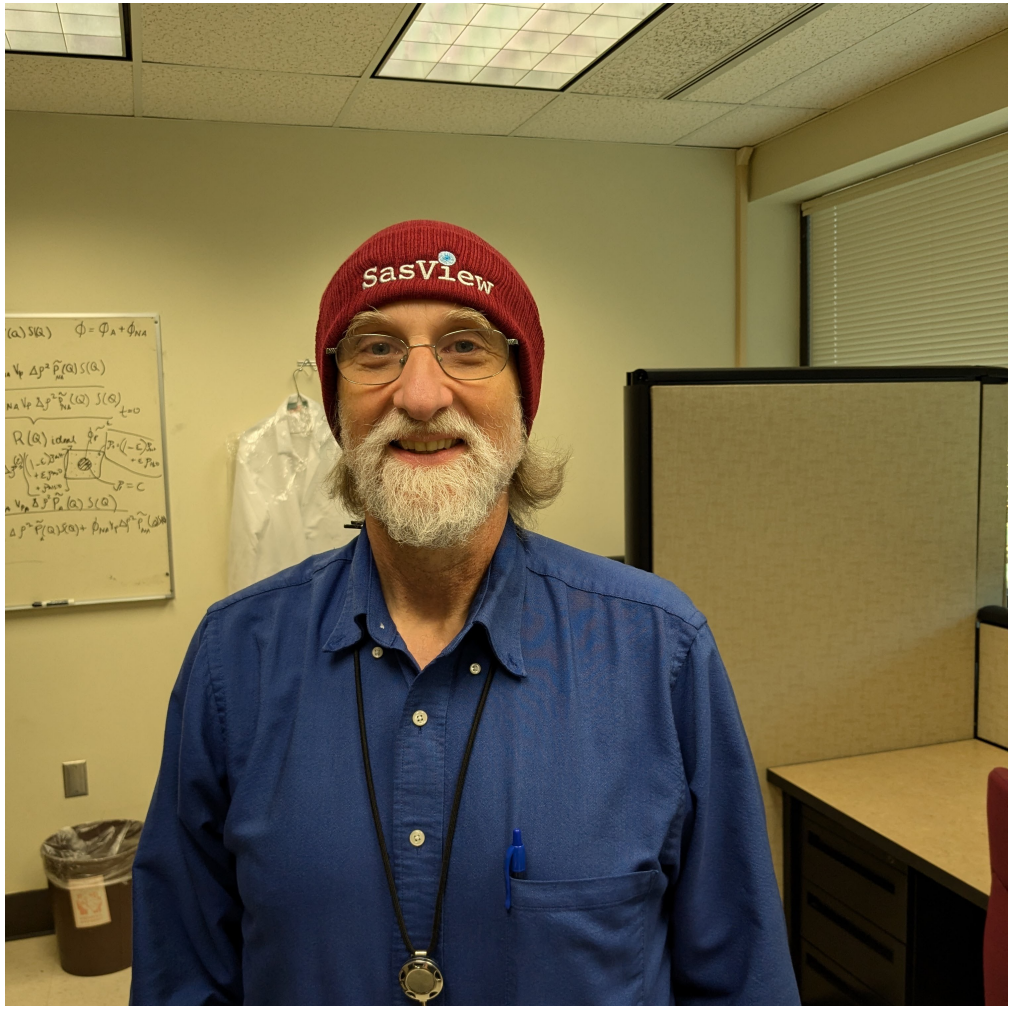
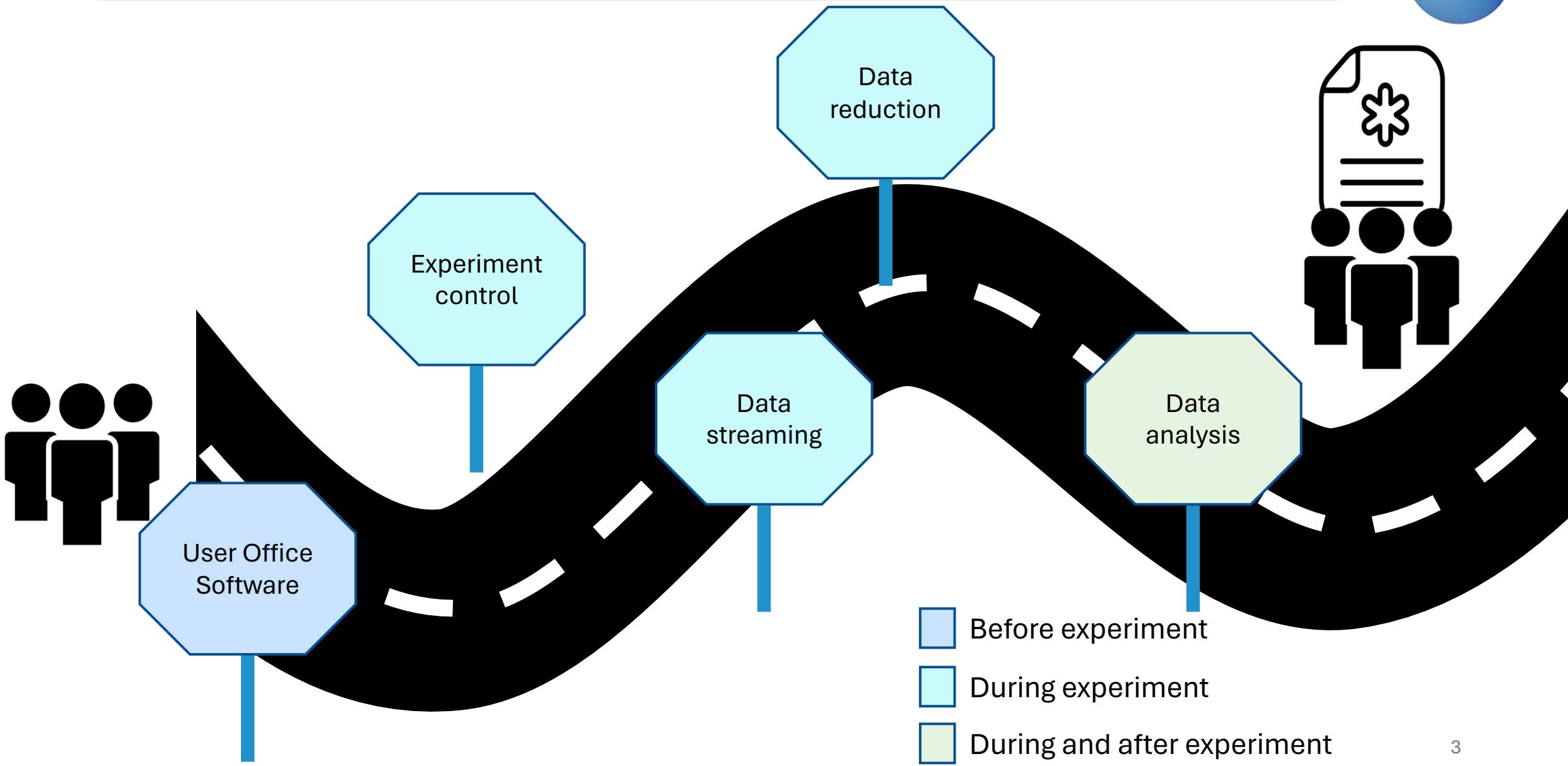
I will change the hat now…



Paul Butler, NIST

User Journey at Large-Scale Facility



Data analysis – facts of life

Data analysis is a part of the processing workflow

Reduced data on the disks is useless

User expertise is critical for successful data analysis

Facilities should support data analysis software but shouldn’t fully drive the process

->We need open and collaborative software tools to succeed!

The SasView approach

An open, collaborative, community development

- Community driven project
- Supported by 10 facilities, 40 contributors (~15 active at any one time)
- Small leadership team: Paul Butler (NIST), Steve King (ISIS), Wojciech Potrzebowski (SciLifeLab, formerly ESS)
- Biweekly calls
- Regular SasView camps and hackathons

- Hosted on github: https://github.com/SasView/sasview

Pictures from Code Camps: V, VI, VIII, X

www.sasview.org

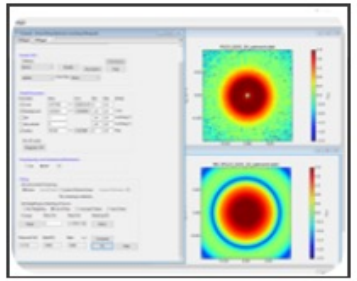
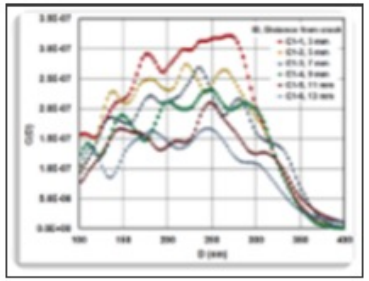
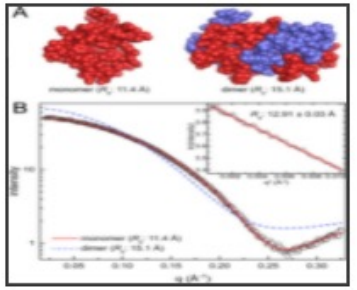
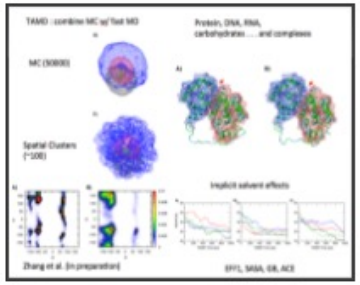
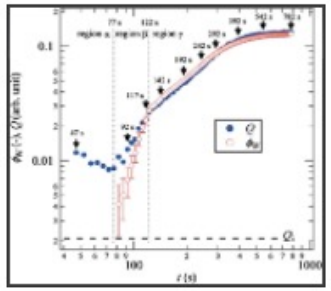
SasView collaboration

- Those who bring the resources (time, effort, funds) choose what to work on

- but cannot break existing experiences ...

Different approaches to SAS data analysis

Not in SasView scope

Model-Fitting Methods	Real-Space Methods	~~Ab Initio Methods~~	~~MC/MD Methods~~	Other Methods
				
Example of 2D model-fitting using the SasView application	*Cavity size distributions in a steel weldment as derived from SANS 10.1179/1743284714Y.0000000577*	*Ab-initio modelling of polcalcin constrained by SAXS 10.1002/pro.3376*	*MC & TAMD modelling of proteins constrained by SANS 10.1016/j.jmgm.2017.02.010*	*Time evolution of the invariant during crystallisation of P4MP1 10.1038/pj.2012.204*

https://www.isis.stfc.ac.uk/Pages/SANSdataanalysisOverview.aspx

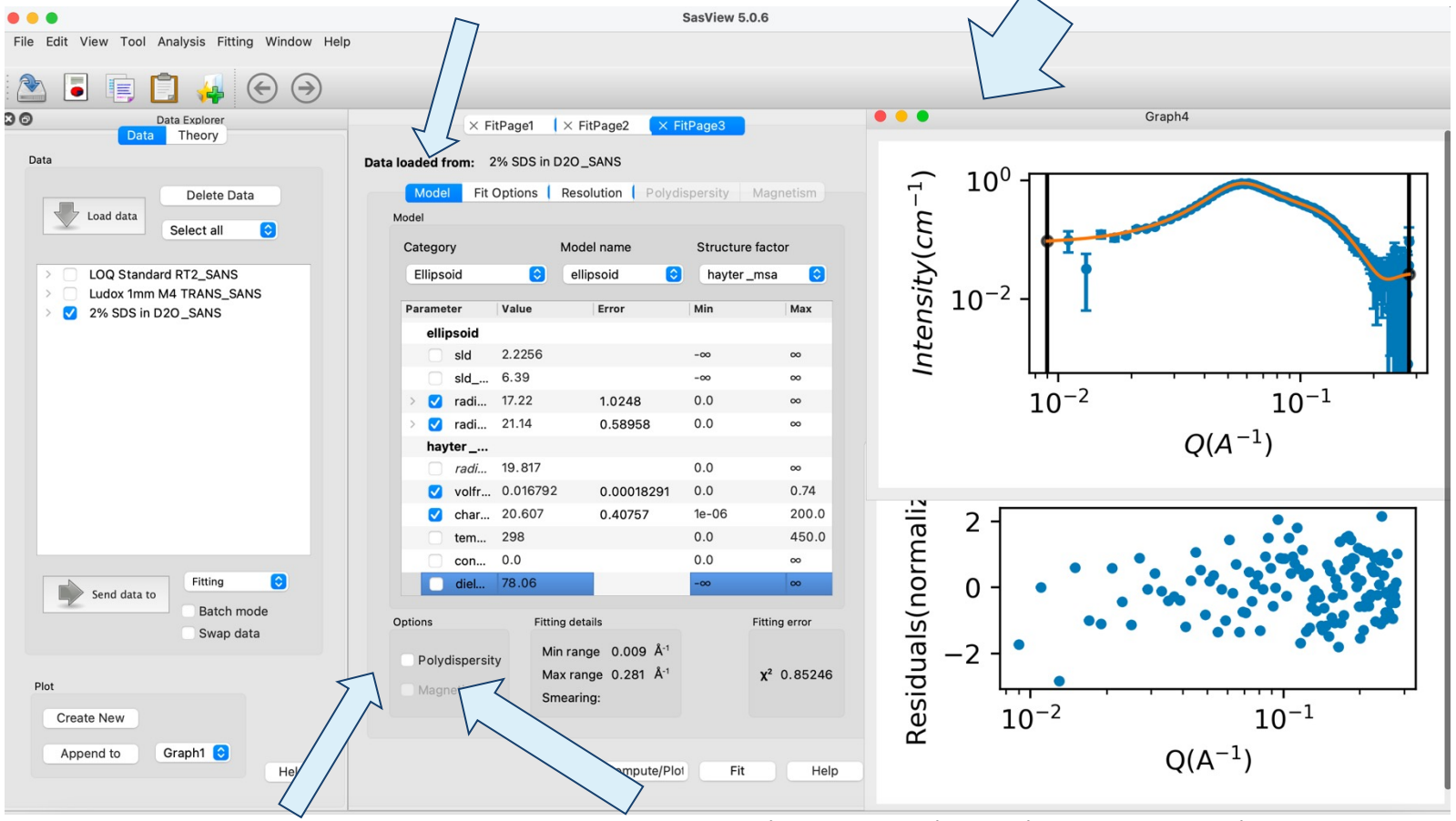
70+ models to explain data

Wide choice of built-in models (> 70)
F(Q), S(Q) & F(Q)*S(Q)

Single, batch and simultaneous
1D and 2D fitting

Data management
Common data formats supported, including NXCansas & *cansas1D*

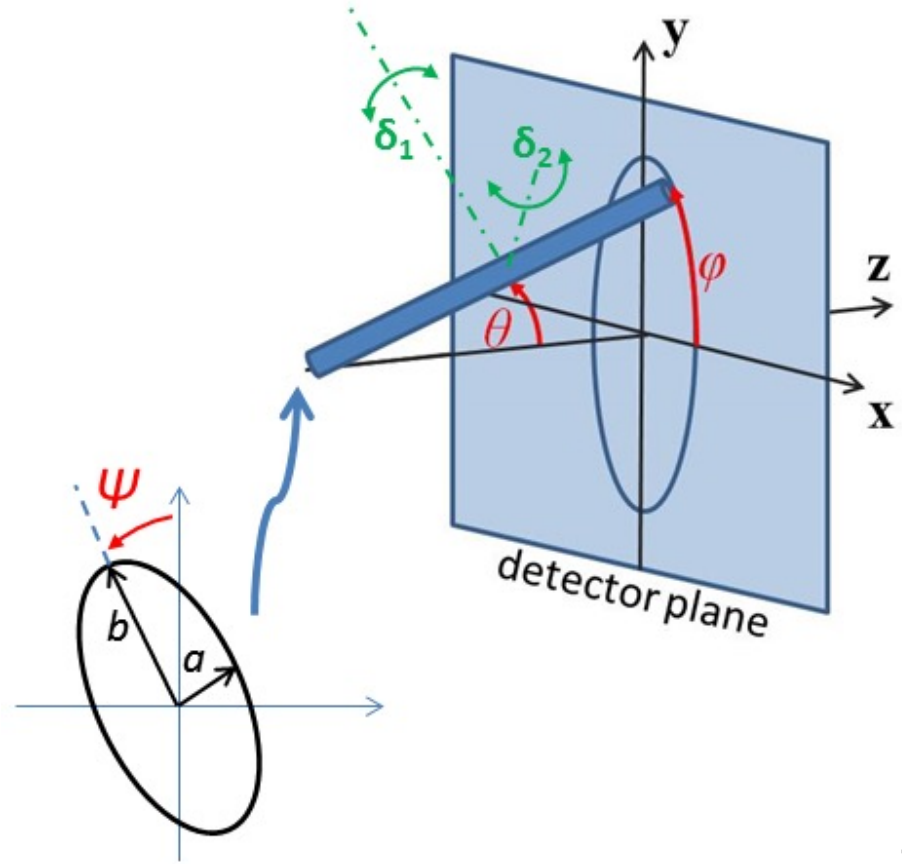
Analysis Tool Choice
&
Plotting

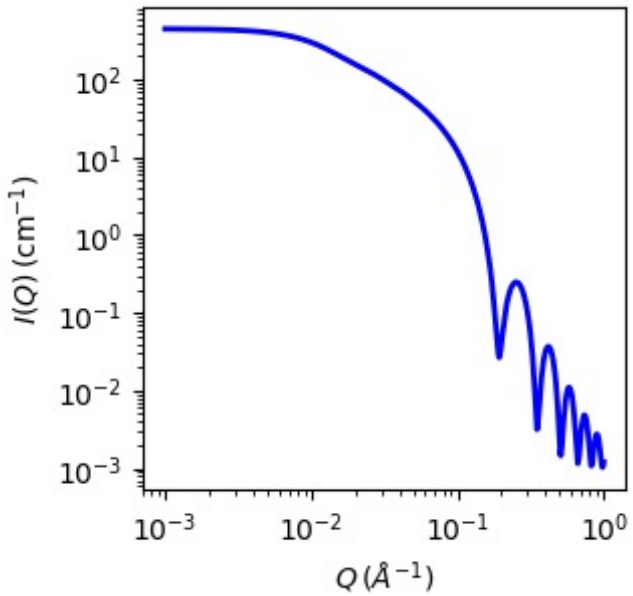


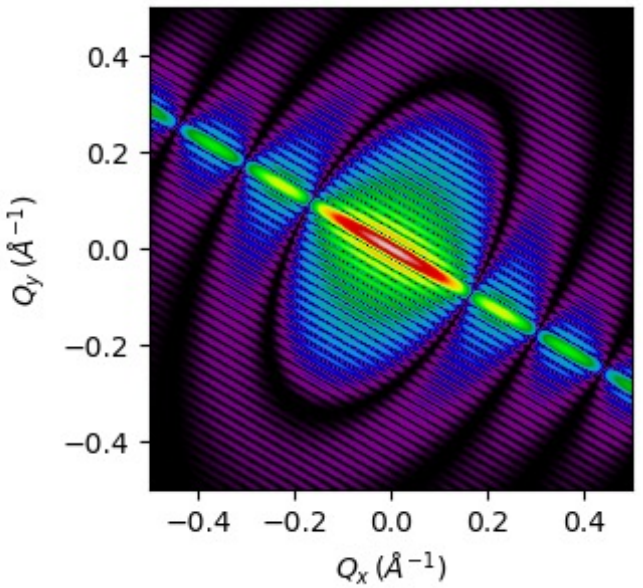
Polydispersity (choice of distribution and distribution parameters)

Resolution smearing (pinhole and slit)
Automatically from data or provide parameters

2D fitting for oriented or magnetic particles



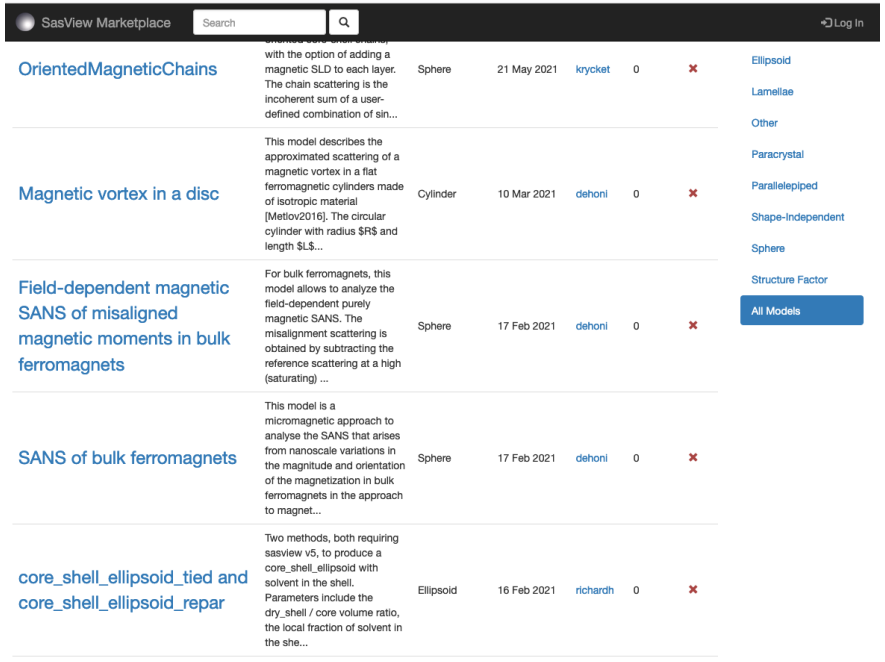


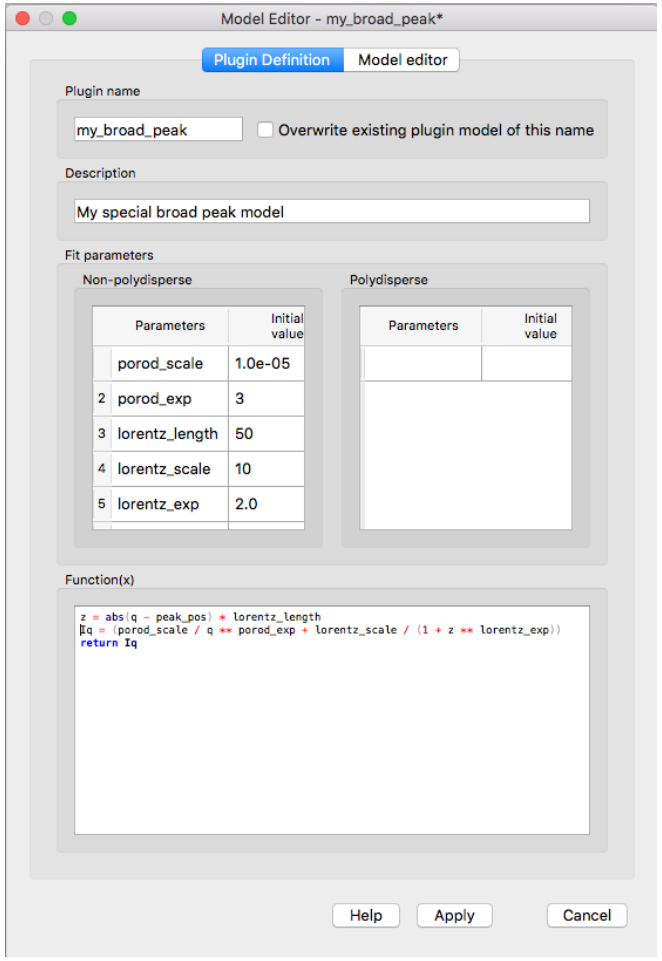


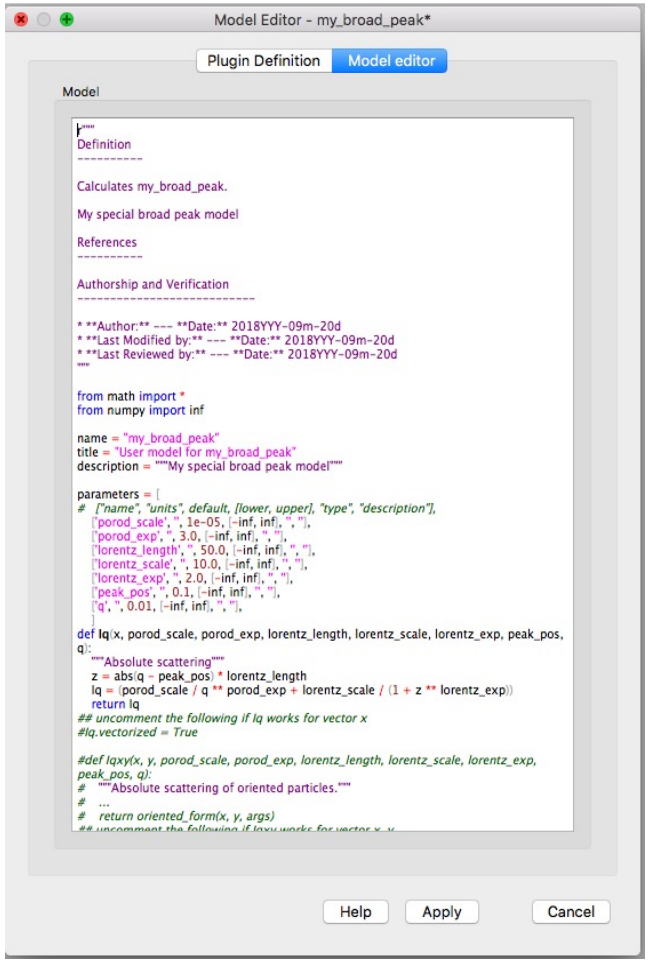
1D and 2D cylinder model

Plugin models

- Dedicated editor
- Syntax and performance testing
- Directly available in SasView ecosystem
- Community-developed models can be deposited to marketplace: https://marketplace.sasview.org/



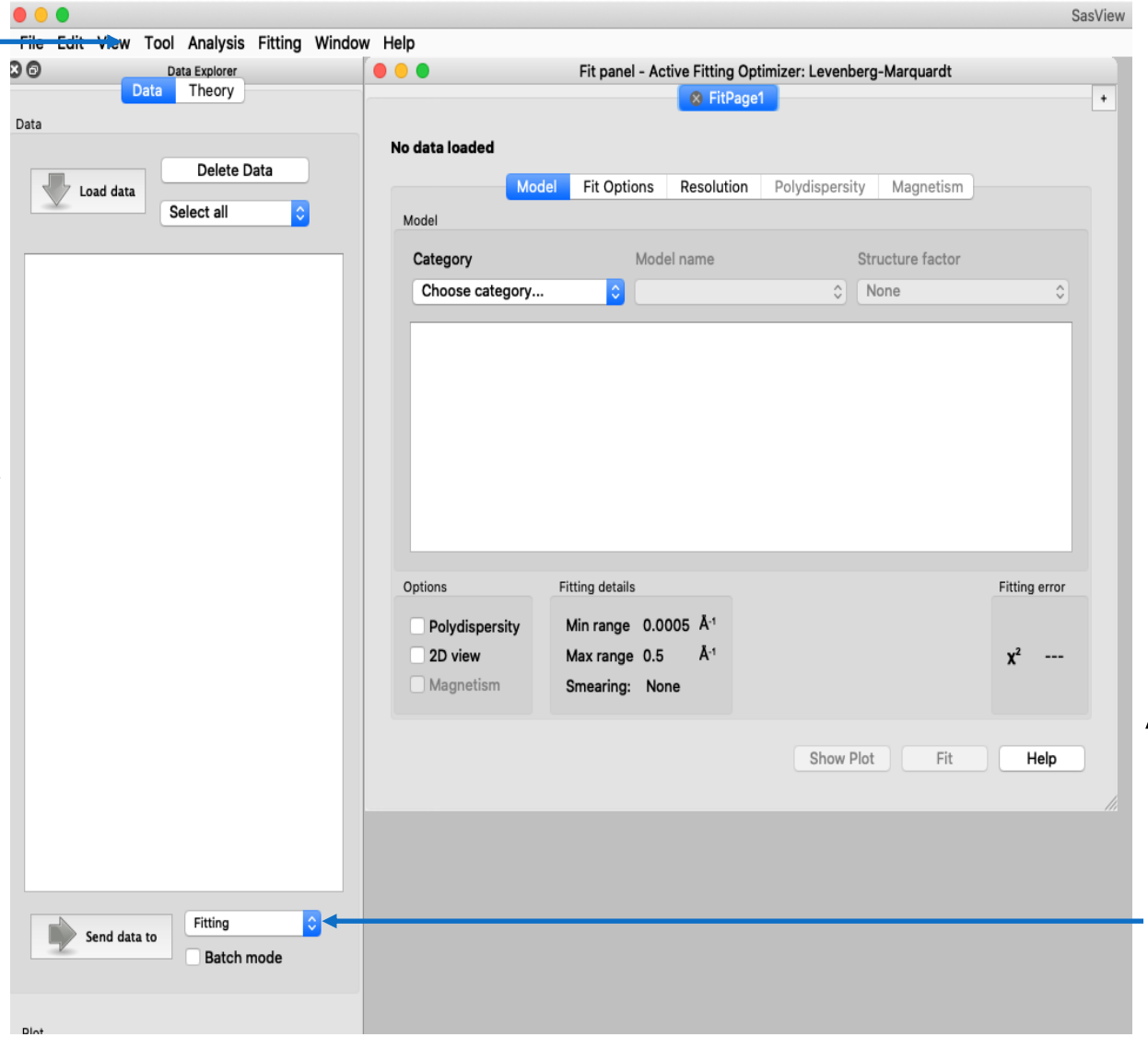




Other SasView functionality

Tools

- Data Operation
- SLD calculator
- Density/Volume calculator
- Slit Size Calculator
- Kiessig Thickness Calculator
- Q Resolution Estimator
- Generic Scattering calculator
- Orientation Viewer
- Python Shell/Editor
- Image Viewer
- File Converter



Analysis

- Fitting
- Invariant
- Pr Inversion
- Correlation Function

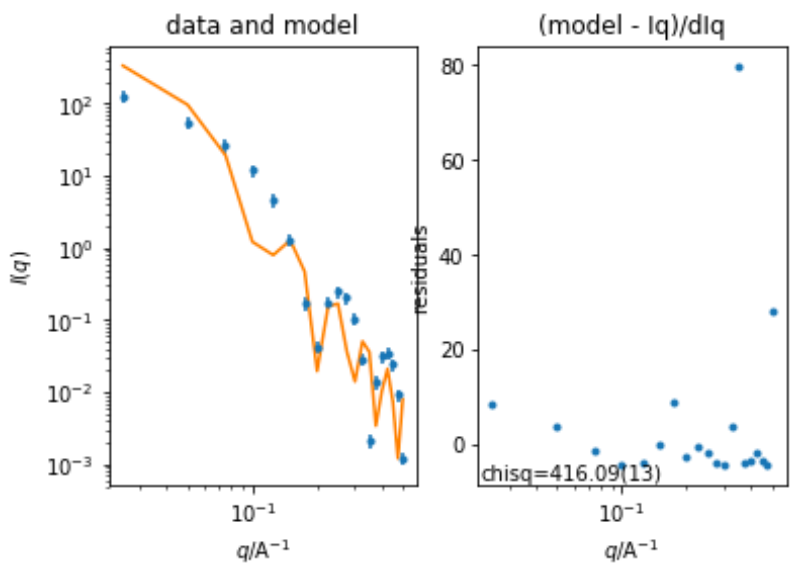
Beyond Graphical User Interface (GUI)

- Useful for batch jobs and reproducibility
- Scripts can be run on computer cluster

```
import pylab
from bumps.names import *
from sasmodels.core import load_model
from sasmodels.bumps_model import Model, Experiment
from sasmodels.data import load_data

from bumps.fitters import fit
from bumps.formatnum import format_uncertainty

test_data = load_data('cyl_400_20.txt')
kernel = load_model('cylinder')

test_data.dy = 0.2*test_data.y

pars = dict(radius=35,
            length=350,
            background=0.0,
            scale=1.0,
            sld=4.0,
            sld_solvent=1.0)
model = Model(kernel, **pars)

# SET THE FITTING PARAMETERS
model.radius.range(1, 50)
model.length.range(1, 500)

M = Experiment(data=test_data, model=model)
problem = FitProblem(M)
print("Initial chisq", problem.chisq_str())
problem.plot()
pylab.show()

result = fit(problem, method='amoeba')
print("Final chisq", problem.chisq_str())
for k, v, dv in zip(problem.labels(), result.x, result.dx):
    print(k, ":", format_uncertainty(v, dv))
problem.plot()
pylab.show()
```

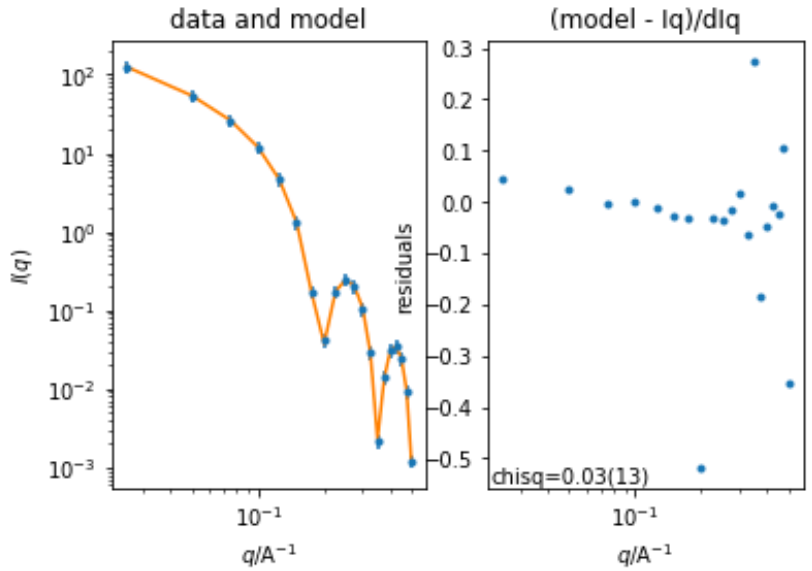
Before fit

```
Initial chisq 416.09(13)
```



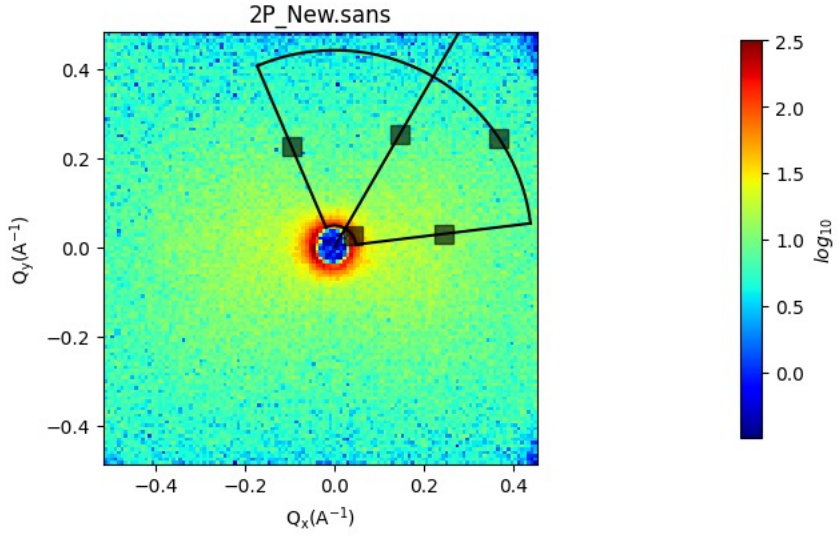
After fit

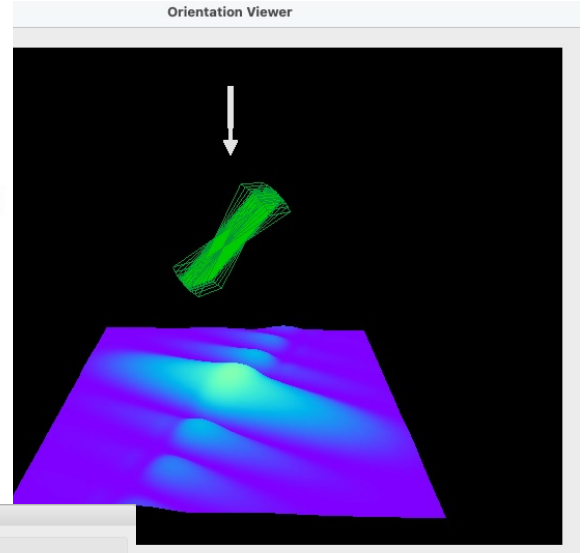
```
Final chisq 0.03(13)
length : 464.9(55)
radius : 19.977(64)
```

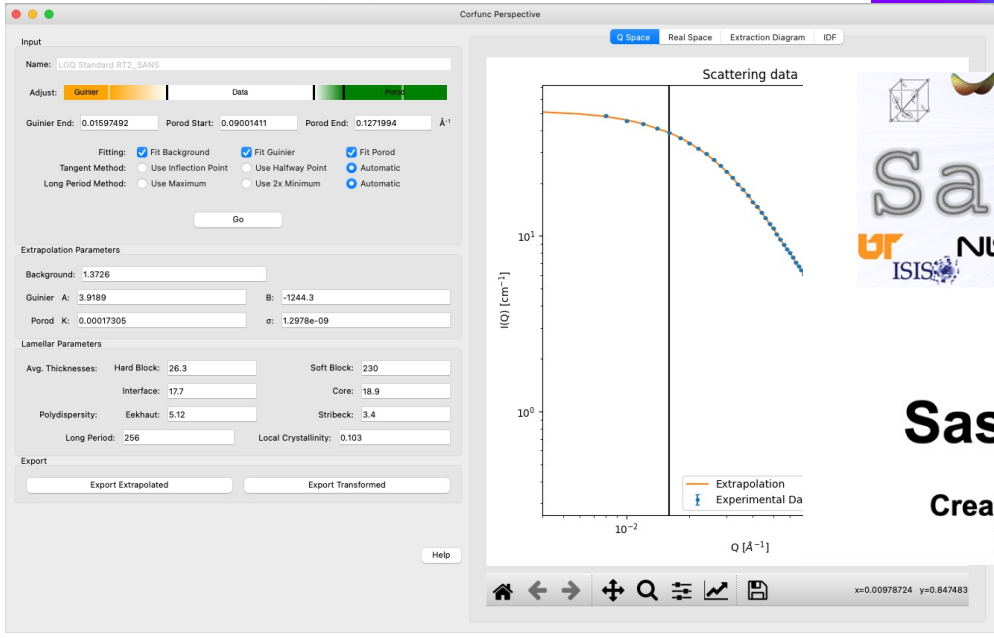


SasView 6.0.0-beta release

- Orientation viewer available
- Corfunc perspective refactored
- Simultaneous fitting allows for a weighting scheme
- Preferences panel with display and plotting options
- Improved label handling on plots
- Residuals plots refactored
- PDB reader refactored
- Wedge slicer added
- Sasdata package separated
- Custom Model writing tutorial







SasView Tutorials

Creating Custom Fitting Models in SasView Version 5.x

https://github.com/SasView/sasview/releases/tag/v6.0.0-beta-2

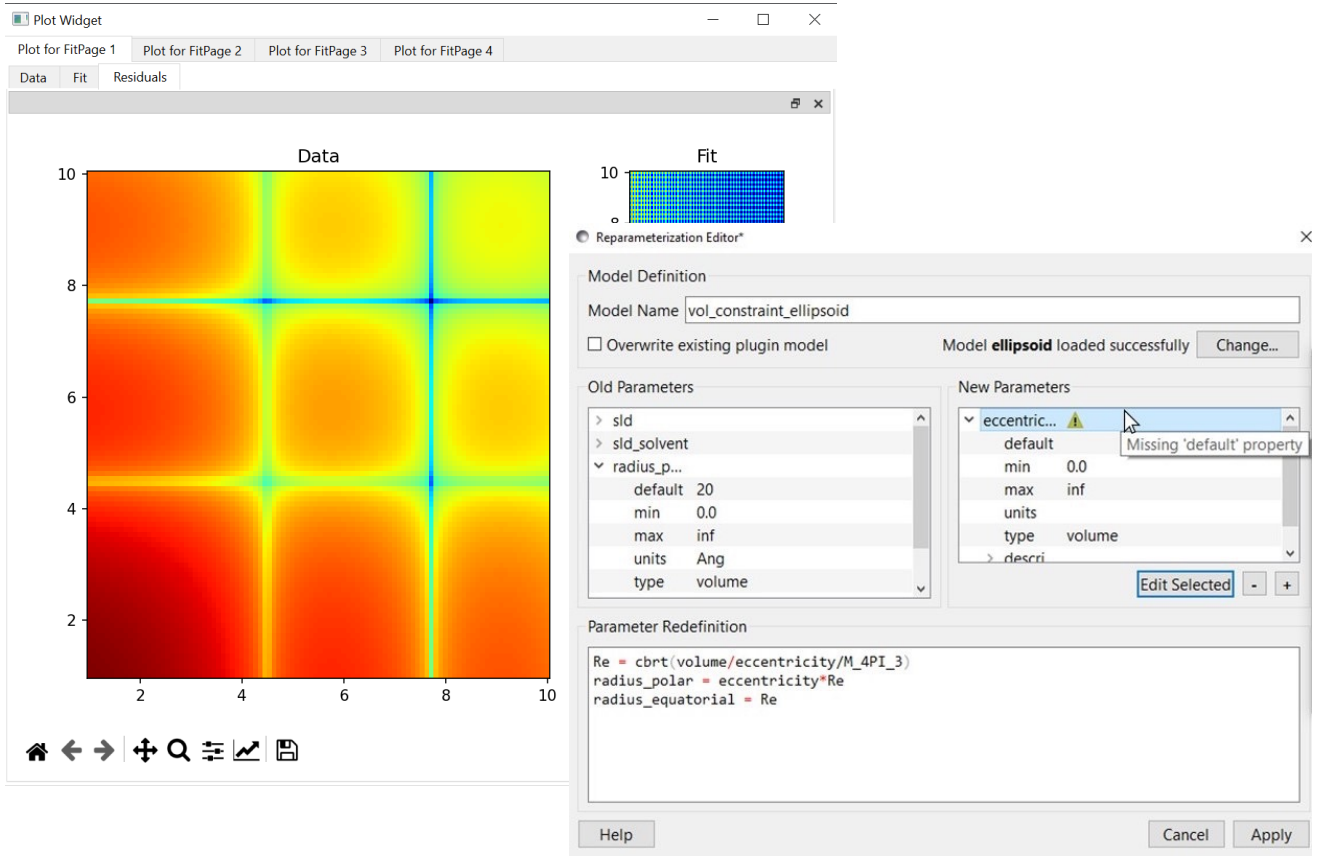
Examples of contributions

- Magnetic SANS community workshop
- Student projects
- Incorporating external software tools

Contributor camp, University of Delaware, January 2024

Student projects

- Improvements in:
  - Plotting
  - Data structure
  - Documentation
  - Plugin Editor
  - Slicers
  - Generic Scattering Calculator
- Web-based API
- Pore size distributions
- Simultaneous SAXS and SANS fitting



Iestyn Cadwallader-Jones, Nouhalia Agouzal, Dorian Lozano, Julius Karliczek (ILL), Ellis Hewis, James Crake-Merani, Ruben Lopes (ISIS), Alex Zheng (NIST), Brayden Miller (NIST), Xael Shan (NIST), Anita Zhang (Princeton), Kristian Lytje (Arhus University)

Magnetic SANS workshops

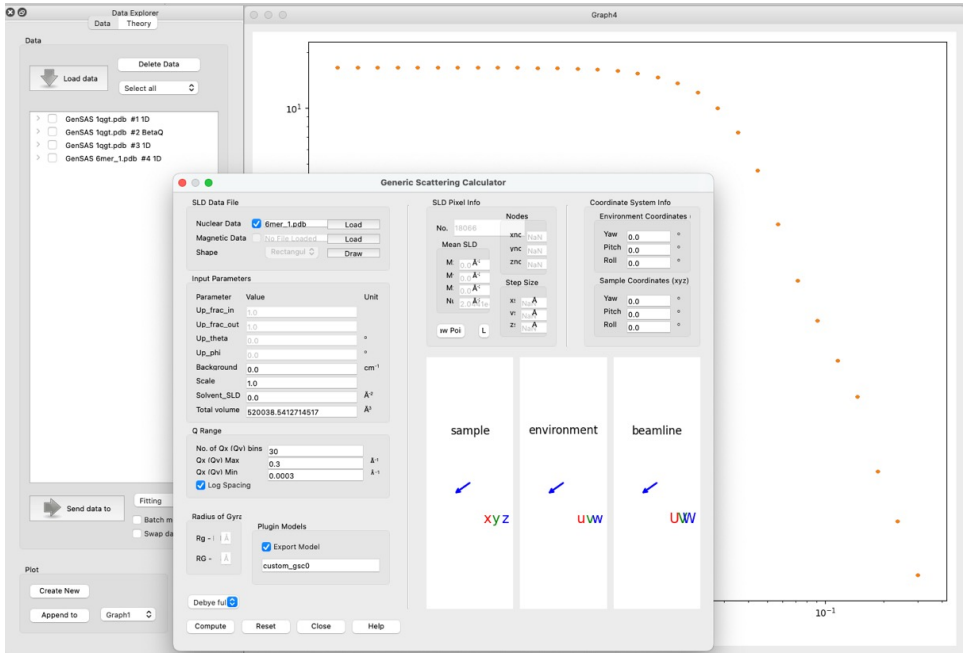
- 2-day in-person workshop in 2022
- 1-day follow-up hybrid workshop in 2023
- Participants from universities and facilities
- Gather requirements for Magnetic SANS data analysis
- A number of GitHub issues created
- Ongoing work on slicers and porting MuMag functionality (**Michael Adams**, Univ. of Luxembourg)

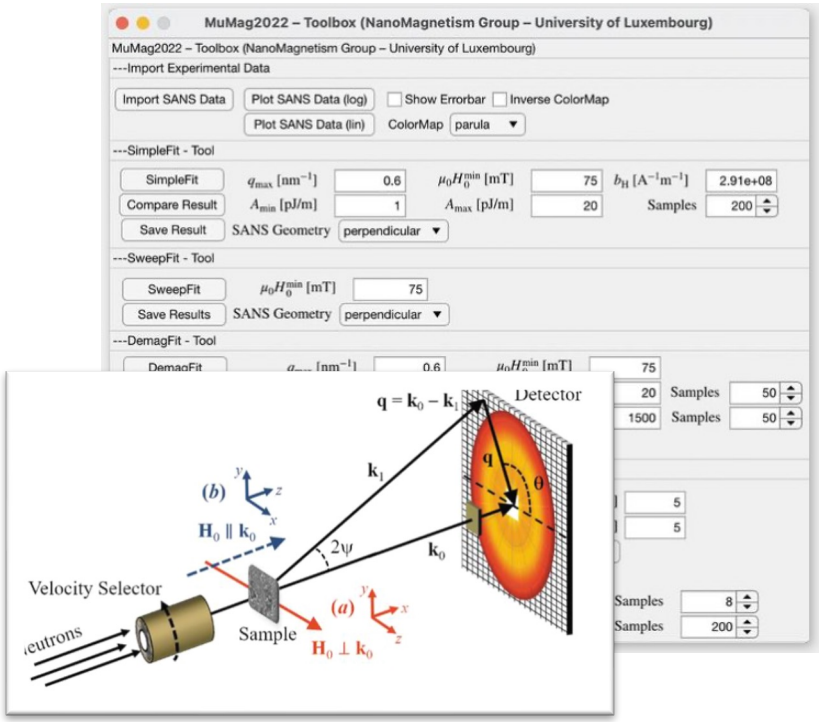
https://www.tandfonline.com/doi/full/10.1080/10448632.2022.2126691

Credit: Annika Stellhorn (ESS)

https://www.linxs.se/events/2023/1follow-up-magnetic-sans-data-and-software-h4crj

Incorporating external tools





Interacting bio-molecules

- Generate scattering profile from PDB file
- Save as a custom/plugin model
- Use for fitting with structure factors (including beta approximation)
- PDB reader and engine refactored

K. Lytje and J. S. Pedersen Acta Cryst. (2024). D80, 493-505

Analyzing ferromagnets

- MuMag analyzes unpolarized total SANS data within the approach-to-saturation regime
- Matlab based GUI converted to SasView GUI

M. P. Adams, M. Bersweiler, E. M. Jefremovas and A. Michels J. Appl. Cryst. (2022). 55, 1055-1062,

How to get involved?

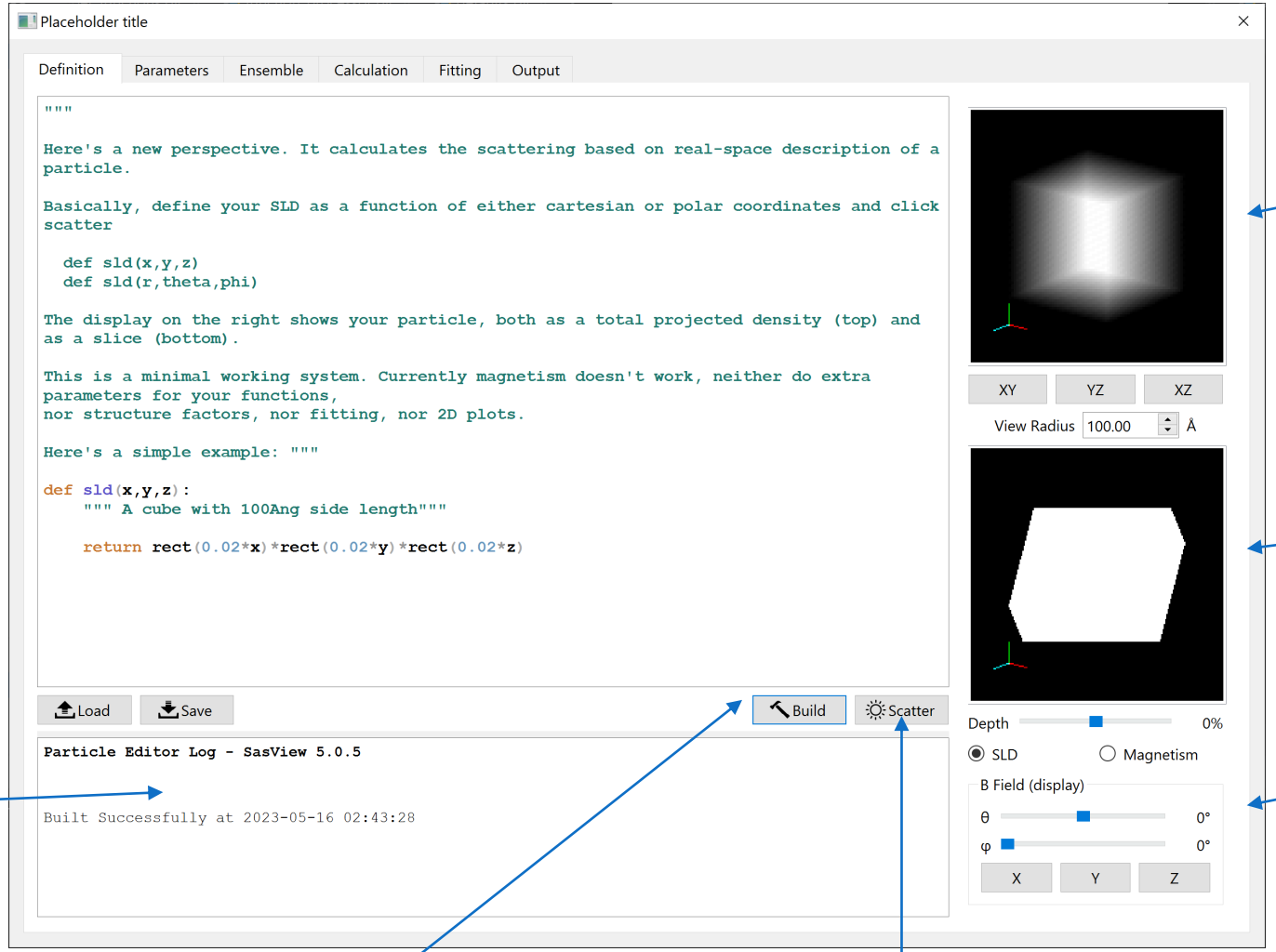
- Build up community/working group (like Magnetic SANS)
- Student projects
- Integrate own software libraries
- 5yr road map
- Project ideas (related to LOKI):
  - Rheo-SANS
  - SEC-SANS data treatment (ATSAS equivalent)
  - Improvements in scattering calculator for biomolecules
  - TOF resolution function (together with CanSAS)

www.sasview.org/contribute

Questions?

Thank you!

Features beyond 6.0.0 - particle editor



Define functions sld and (optionally) magnetism

sld / magnetism "x-ray" projection

sld / magnetism cross section

Feedback on code and calculations

Magnetic field controls for display

"Recompile" and update display

Compute scattering