

EUROPEAN SPALLATION SOURCE



DMSC STAP Updates from DRAM

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TORBEN NIELSEN

October 2022

Agenda



- 1 DRAM
- 2 Data reduction updates
- 3 Modelling updates
- 4 Data analysis updates

DRAM

Data Reduction, Analysis and Modelling





Simon Heybrock



Neil Vaytet



Jan-Lukas Wynen



Piotr Rozyczko



Simon Ward



Andrew Sazonov



Peter Willendrup



Mads Bertelsen



Justin Bergmann

Scope

The DRAM group is responsible for providing the data reduction, analysis and modelling software for all instruments at ESS.

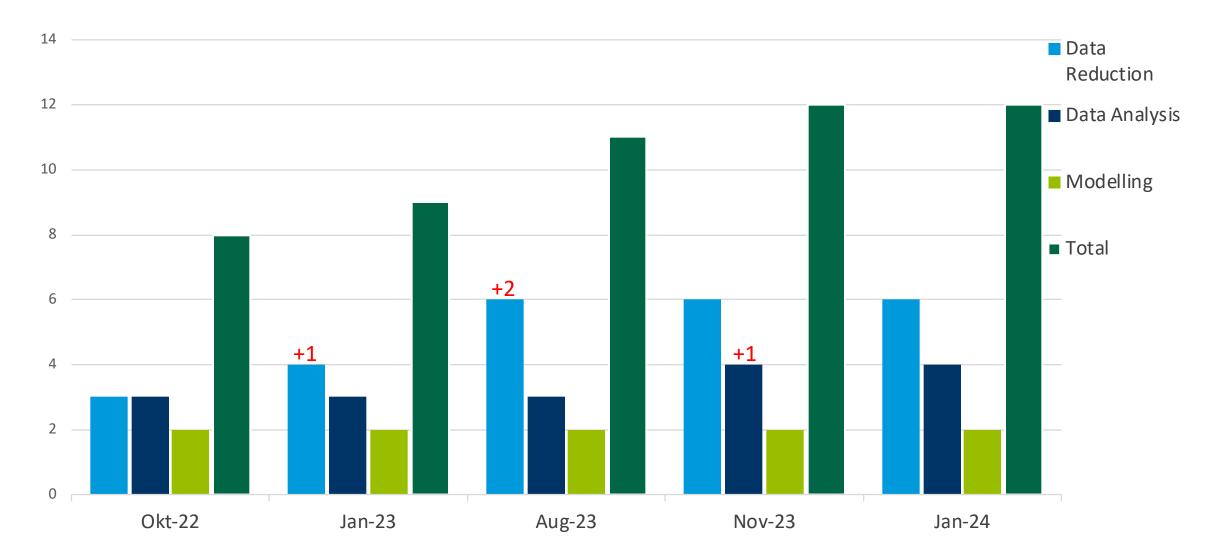
- □3 teams (9 persons)
 - ☐ Data Reduction (scipp)
 - □ Data Analysis (SasView, SpinW, EasyScience, external collaborations)
 - Modelling (McStas++, pan-learning.org, NMX)

DRAM - staff profile

P0/P6 – spring 2022

S

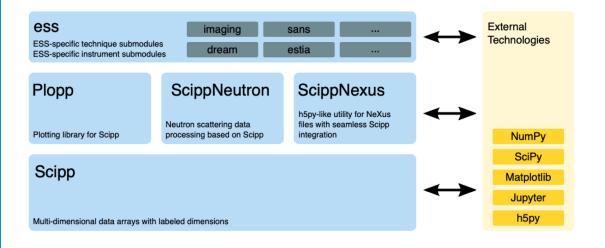




scipp

scipp, scippneutron, scippnexus, plopp and ess





- ☐ Close collaboration between scipp and IDS team for defining data reduction workflows in scipp
- ☐ Joint meetings, pair-programming, and lively discussion on a dedicated Slack channel.

- □ scipp continues to be developed with input and requirements from instrument data scientists.
- ☐ Good progress
- ☐ Lots of features added & issues fixed

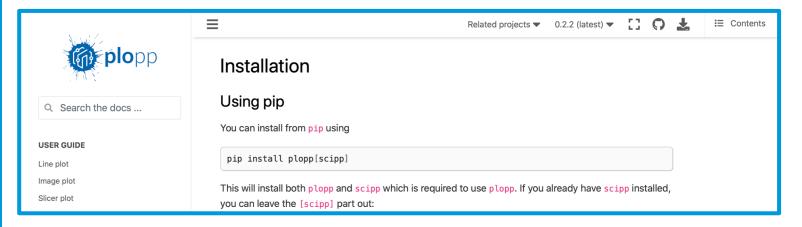




Documentation on https://scipp.github.io



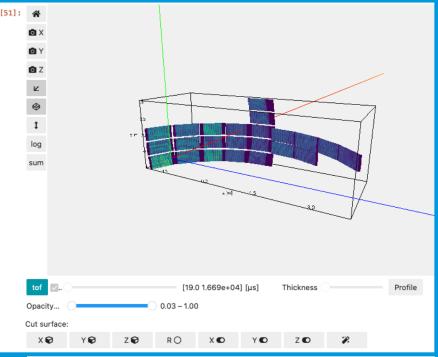
 scippnexus: An h5py-like utility for NeXus files with seamless Scipp integration

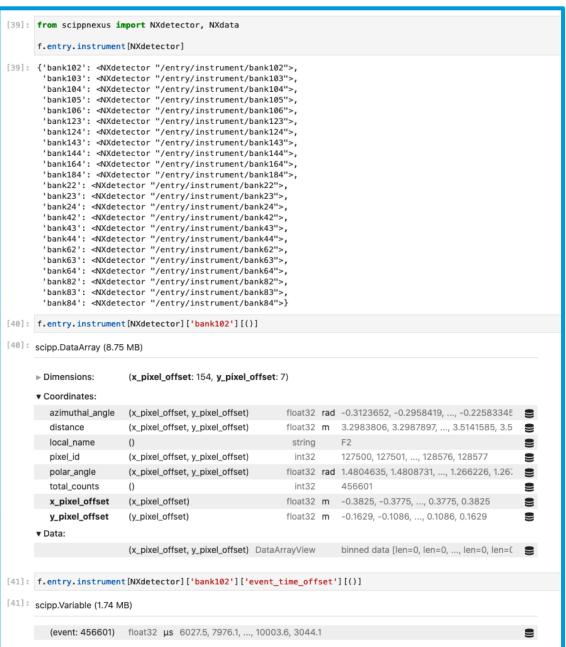


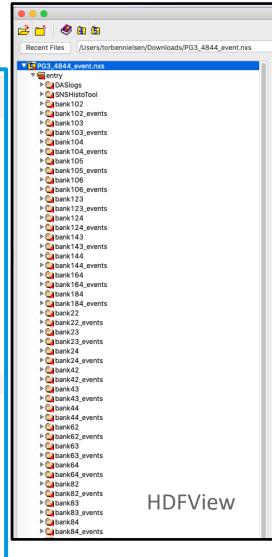
□ plopp: plotting library for scipp

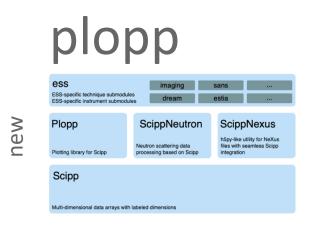
scippnexus

- POWGEN data
- ☐ PG3_4844_event.nxs









Scipp vs Plopp

This notebook aims to illustrate the differences between the current (old) Scipp plotting, and the new Plopp package.

Background

See ADR0013 for more details.

The current plotting functionality automatically resamples 2-D (or higher) data as well as binned data of any dimension. There were three main drivers/motivations for this:

- We frequently deal with data with "long" dimensions, such as "pixel" or "spectrum" dimensions, which may exceed a million.
- We often deal with "ragged" data, i.e., data with 2-D coordinates that originate after coordinate transformations.
- We frequently plot binned data. Bins are often much larger than the target plot resolution, so histogramming must be done. This is combined with "long" dimensions and potentially high resolutions along other dimensions (exceeding 1000 or even 10000).

In the past we have therefore taken it as a given that resampling is a non-optional requirement. However, we have since then experienced a multitude of problems with this, ranging from simple bugs to complex and hard to maintain code. More importantly, the resulting plots are unpredictable and not faithful to the actual data.

One of the most compelling conclusions was:

"Scientifically any plot generated with resampling is not usable, e.g., for publications."

New requirements

From scipp#2475

- 1. Faithful representation of data
- 2. Not tied to plotting/widget library
- 3. Updatable data
- 4. Both interactive and static figures
- 5. Customizable figures
- 6. Plots should still be visible in saved notebooks
- 7. API for adding custom tools with callback (or similar) ESENTATION TITLE/FOOTER

Issues with automatic resampling [3]: print(da.min().value, da.max().value) -0.9999902065507035 0.9999886166646338 [4]: s = sc.plot(da, title='Scipp') p = pp.plot(da, title='Plopp') ipw.HBox([s._to_widget(), p.to_widget()]) Scipp [4]: Plopp 0.006 0.75 0.004 40 40 0.50 - 0.002 30 0.25 yy [m] yy [m] 0.000 0.00 20 20 -0.25-0.002-0.5010 10 -0.004-0.75

10

15

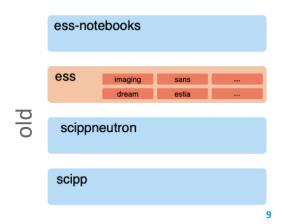
20

xx [m]

25

30

35



5

10

15

20

xx [m]

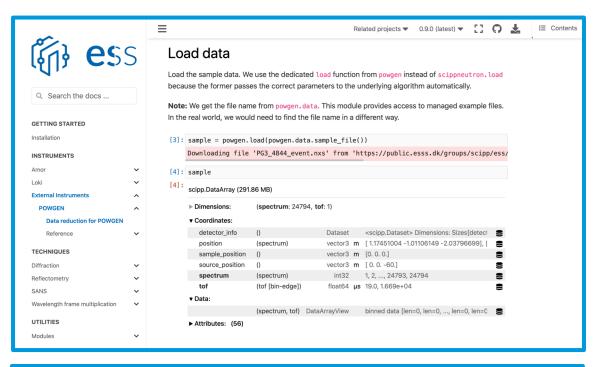
25 30

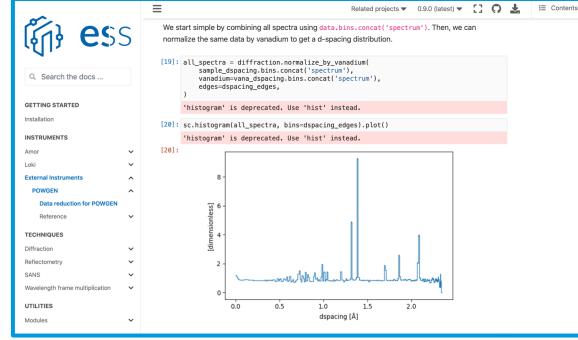
35

scipp / IDS

Data reduction workflow - - see ess package

- Description of the highlights include the ess package being extended for powder diffraction support such that reduction workflows are now available directly for the users on the online documentation pages.
- Data reduction for POWGEN
- This notebook shows a basic reduction workflow for powder diffraction for the SNS POWGEN instrument. It serves mainly to develop and present routines for powder diffraction and will eventually be removed in favor of a workflow for DREAM at ESS.
- See ess docs on <u>io pages</u>





scipp



A bunch of releases

scipp: #10: v0.16.4 - v0.16.3 - v0.16.4 - v0.16.2 - v0.16.1 - v0.16.0 - v0.15.1 - v0.15.0 - v0.14.1 - v0.14.0 - v0.13.1 (April 01 2022)

scippneutron: #9: v.0.9.4 - v.0.9.3 - v.0.9.2 - v.0.9.1 - v.0.9.0 - v.0.9.4 - v.0.8.0 - v.0.7.0 - v.0.6.0 - v.0.5.3 (30 March 2022)

ess: #7: v0.9.0 - v0.8.0 - v0.9.0 - v0.7.1 - v0.7.0 - v0.6.0 - v0.5.0 - v0.4.0 (Feb 28 2022)

scippnexus: **#21**: v0.3.2 - v0.3.1 - v0.3.2 - v0.3.0 - v0.2.1 - v0.2.0 - v0.1.8 - v0.1.7 - v0.1.6 - v0.1.5 - v0.1.4 - v0.1.3 - v0.1.2 - v0.1.1 - v0.1.0 - v0.0.9 - v0.0.8 - v0.0.7 - v0.0.6 - v0.0.5 - v0.0.4 (31 March 2022)

plopp: #8:: v0.3.0 - v0.2.2 - v0.2.1 v0.2.0 - v0.1 - v0.03 - v0.0.2 - v0.0.1 (25 Aug 2022)

mpltoolbox: #7: v0.1 - v0.0.8 - v0.0.7 - v0.0.6 - v0.0.5 - v0.0.4 - v0.0.3 (11 Jul 2022)

#62 - releases

Modelling

ess

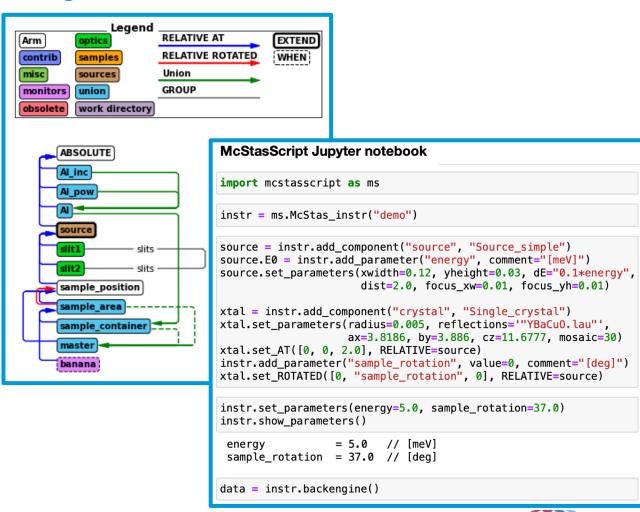
McStas Python API with tools, diagrams and widgets

New developments

- Diagrams of instrument objects
- Improved error checking
- Compatibility with McStas 3.X
- Interactive quiz system in jupyter notebooks
- Dynamic splits of instruments using MCPL

In collaboration with PaNOSC

Instrument database with API





Modelling: workshops and conferences



Events attended since april

Participated in following events

- DMSC McStas workshop
- McStas training days (May-July 2022)
- ICNS Argentina
 Neutron Instrumentation and Innovation award for Mads Bertelsen
- PaNOSC Summer school
- ESS-ILL User meeting
- PaNOSC WP 5 sprint (online)



McStasScript



Modelling overview

mcstasscript: #10: 0.0.51, 0.0.50, 0.0.49, 0.0.48, 0.0.46, 0.0.44, 0.0.43, 0.0.42, 0.0.41, 0.0.40 (March 22 2022)

guide_bot: #10: 0.0.27 - 0.0.26 - 0.0.25 - 0.0.24 - 0.0.23 - 0.0.22 - 0.0.21 - 0.0.20 - 0.0.19 - 0.0.18 - 0.0.16 - 0.0.15 (Feb 16 2022)

Data Analysis

Updates since last STAP



For easyScience core:

- □extension for multiple datasets/fit functions fitting
- □IO rework of easyCore for speedup in XML encoding, and hence a more responsive GUI.
- ■Started work on asymmetric peaks fitting in easyDiffraction

For easyDiffraction:

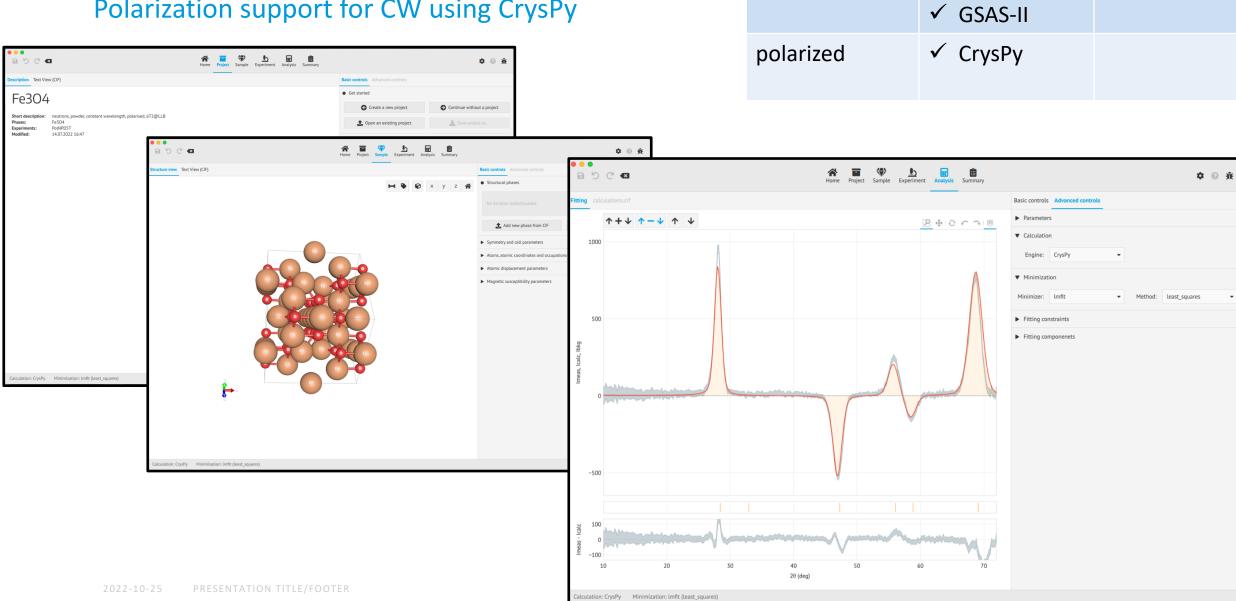
- ☐ easyDiffraction v0.8.4 was released
- □polarized support in the GUI
- ☐ changing how experimental data are handled
- ☐ starting working on file-less interface to CrysFML
- ☐ Improved CrysPy interface

For easyReflectometry:

easyReflectometry v0.0.5 was released surfactant Layer in GUI was added general bug fixes and minor improvements

easyDiffraction 0.8.4

Polarization support for CW using CrysPy



easyDiffraction

un-polarized

CW

✓ CrysFML

✓ CrysPy

TOF

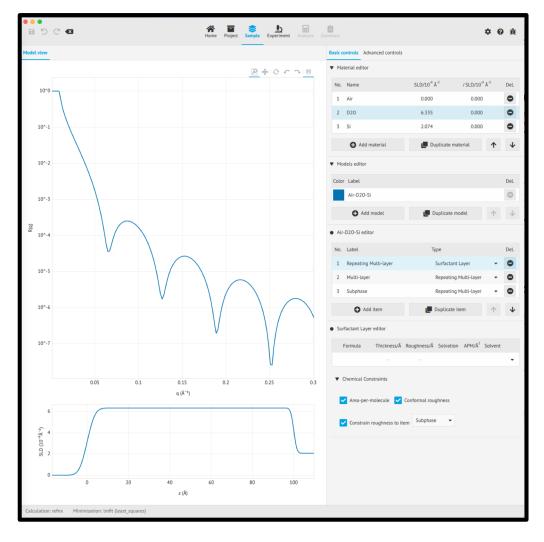
✓ CrysPy

easyReflectometry

v0.0.5

- ☐ surfactant Layer in GUI was added
- ☐general bug fixes and minor improvements
- ☐ Python notebooks



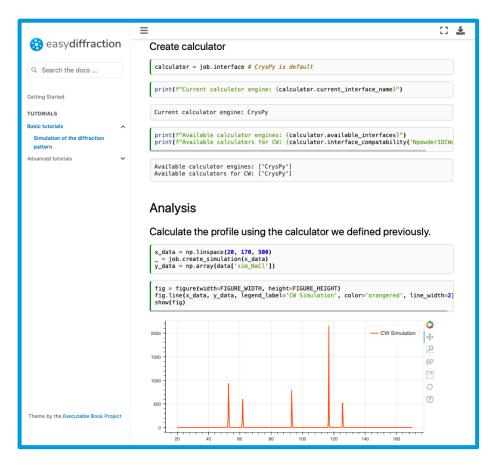


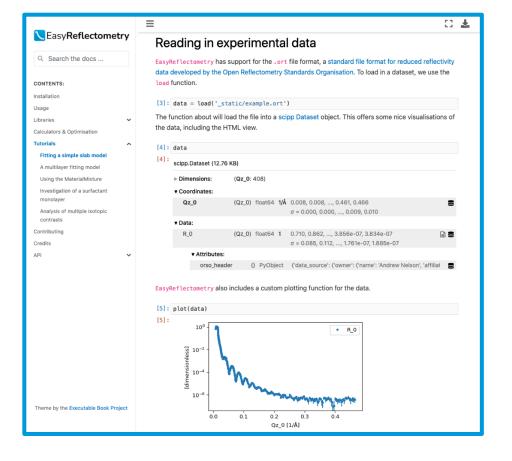
Documentation

https://docs.easydiffraction.org/lib/



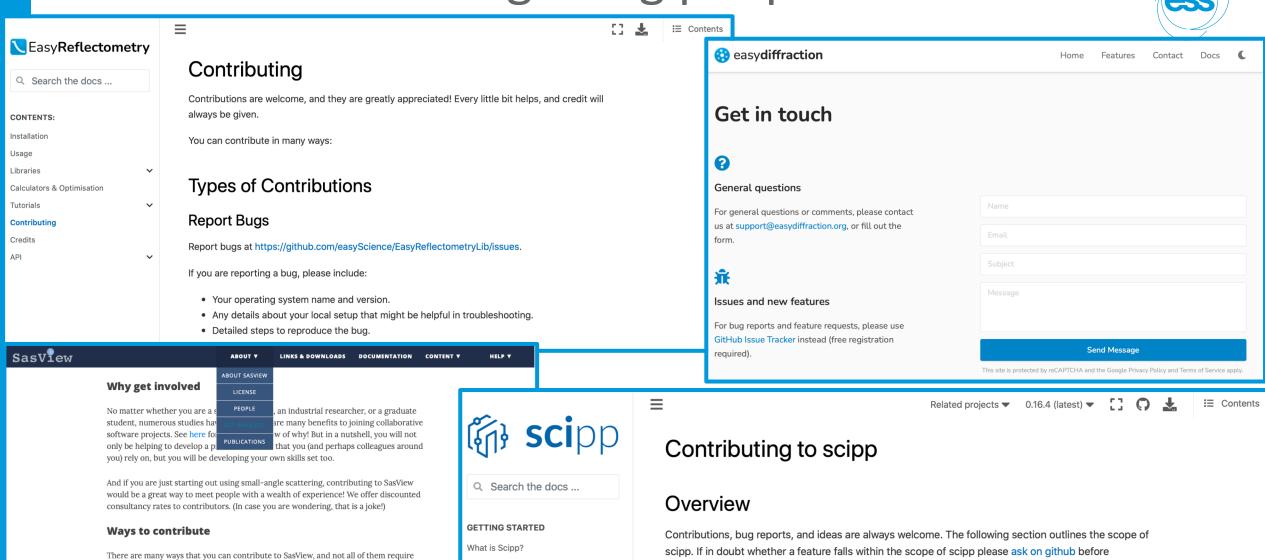
https://docs.easyreflectometry.org





Collaborations – getting people onboard





implementing functionality, to reduce the risk of rejected pull requests. Asking and discussing first is

generally always a good idea, since our road map is not very mature at this point.

Installation

Quick start

Frequently Asked Questions

coding!

We need contributors that can..

Respond to queries posted to help@sasview.org;Teach others how to get the best from SasView;



Finish presentation