

PAUL SCHERRER INSTITUT



Anders Kaestner :: Imaging scientist :: Paul Scherrer Institut

Open source tools for analyzing neutron imaging data

ILL-ESS topical meeting, ~~Grenoble~~ Online, 14-15 October, 2020

# About me and my roles

Instrument scientist

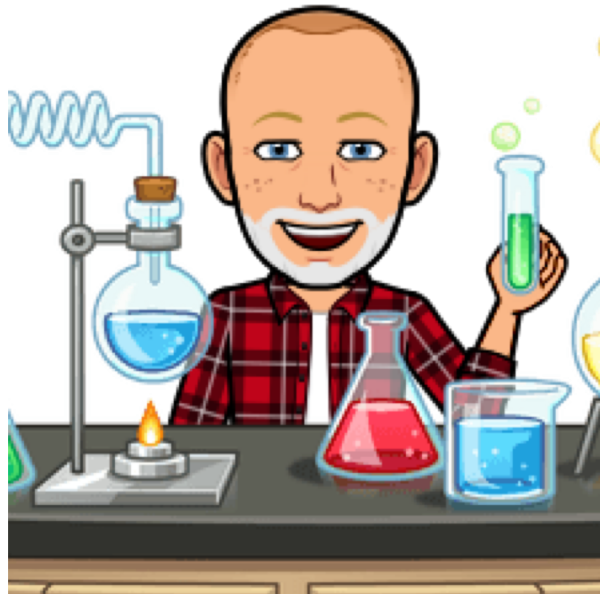


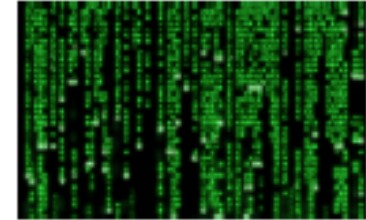
Image processing & software developer





# Introduction

# What happens after an imaging experiment?



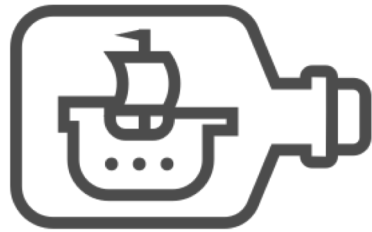
Gigabytes. . .

Image acquisition is not the end of the experiment

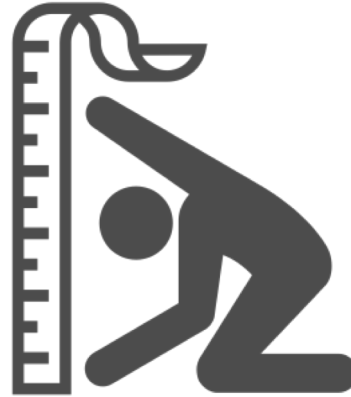
Collaboration with



# What do we want to know?



Find hidden objects



Dimensions



Counting objects

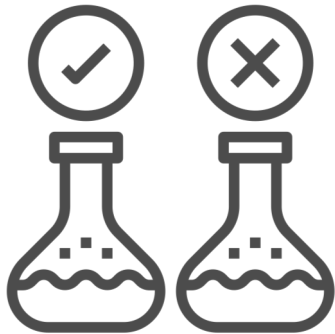


Mixing ratios

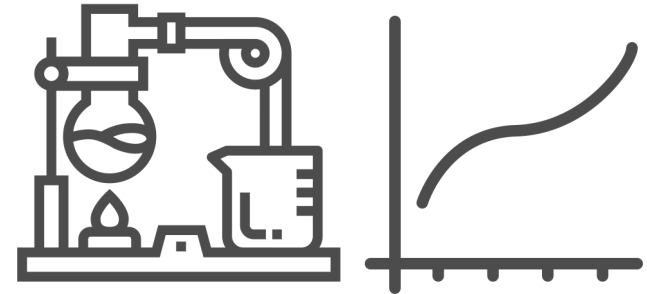
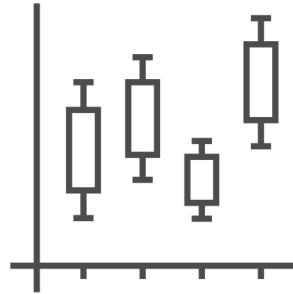


Track changes

# What do we *really* want to know?



Compare treatments



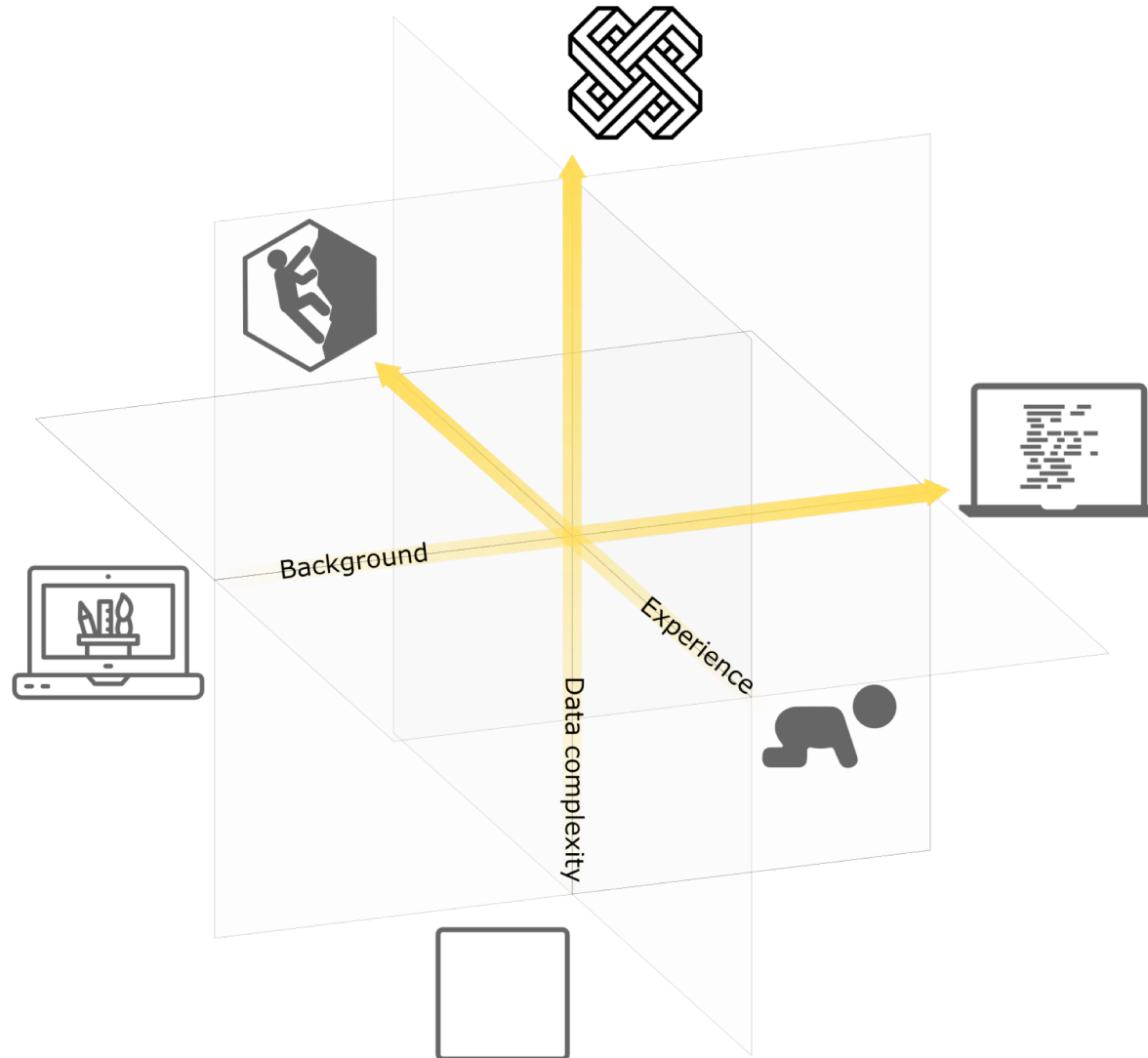
Understand and model processes



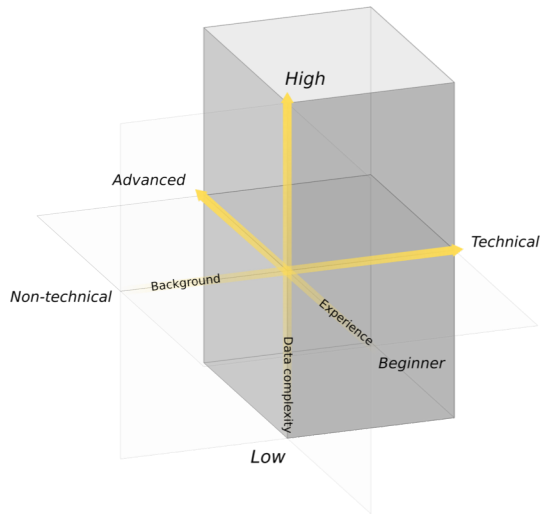
Inspect components

# Data evaluation fitness of users

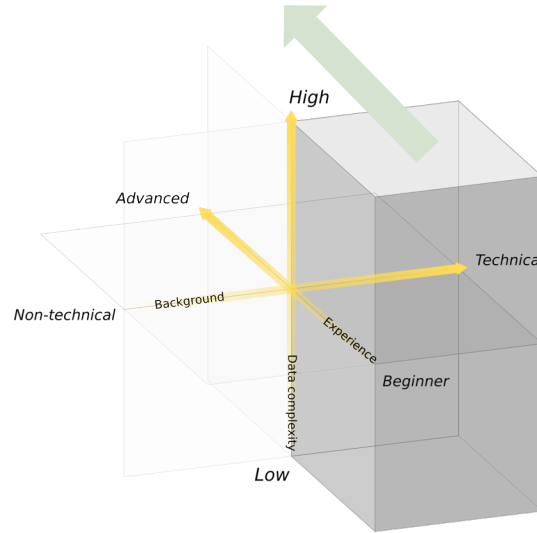
Users of neutron imaging have very different background and experience



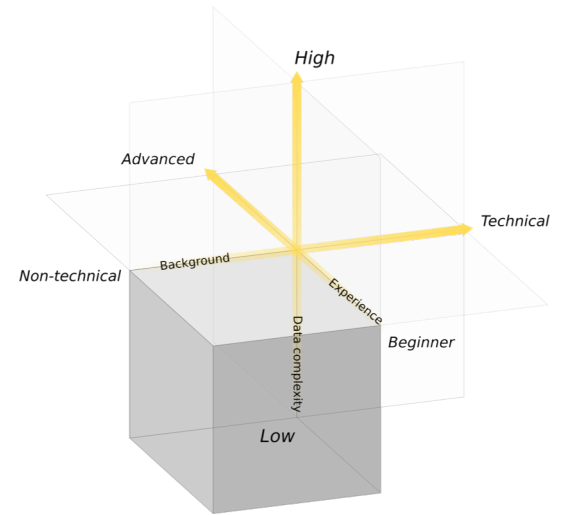
*Beamline scientist*



*Early stage PhD student*



*Cultural heritage users*



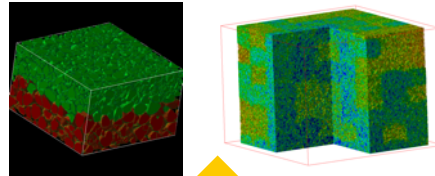
Serving people with different background and needs is a great challenge



# How to analyze the data

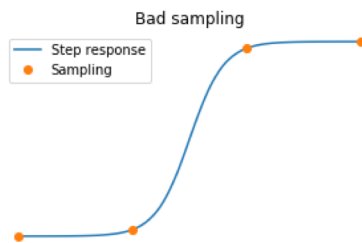
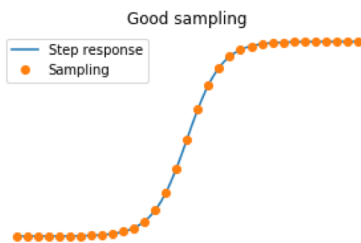
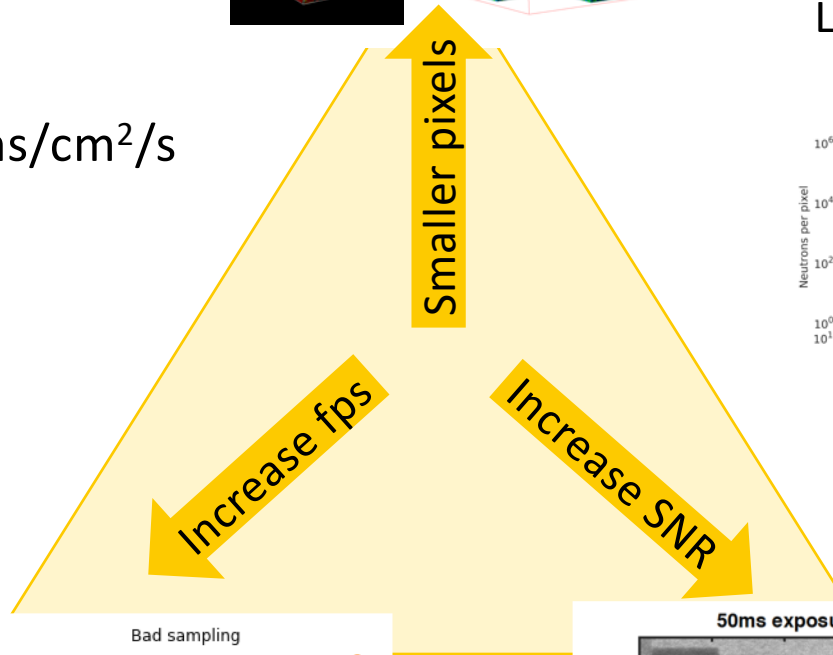
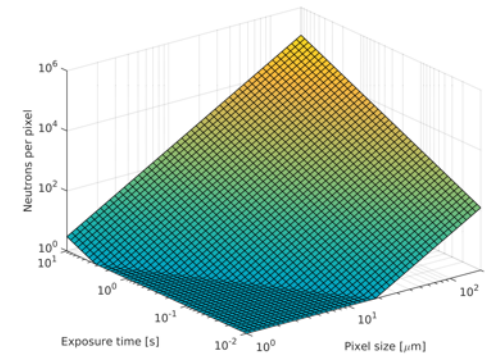
# Challenge of neutron imaging and the analysis

## Feature size vs resolution

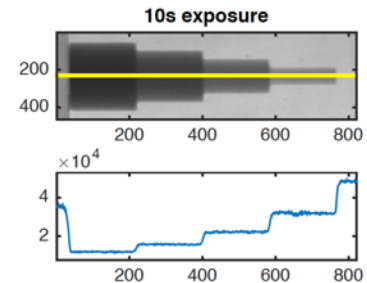
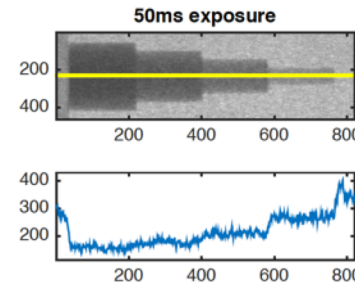


About  $10^7$  neutrons/cm<sup>2</sup>/s

## Limitation: Neutron flux



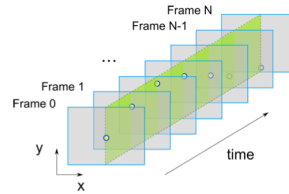
## Process speed vs. Frame rate



## Contrast vs noise

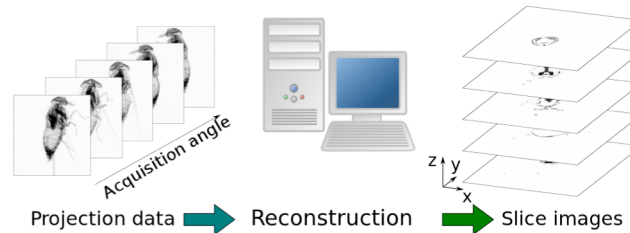
Data processing can be a very labor- and computationally intense task

Basic processing



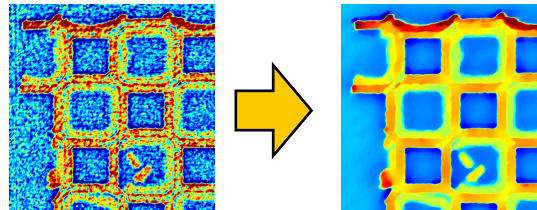
Minutes, hours

CT reconstruction



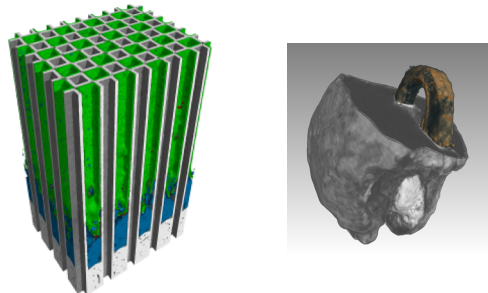
Minutes, hours

Image processing



Hours

Visualization  
Analysis

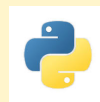
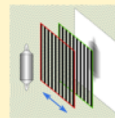


Hours, days

# Synergies in the analysis workflow

Most experiments do (some of these):

- Basic processing
  - Normalization
  - Artifact removal
  - Scattering correction
  - Denoising
- Transformations
  - CT reconstruction
  - nGI reduction
  - Bragg-edge fitting
- Geometric transformations
  - Stitching
  - Registration



Experiment specific analysis

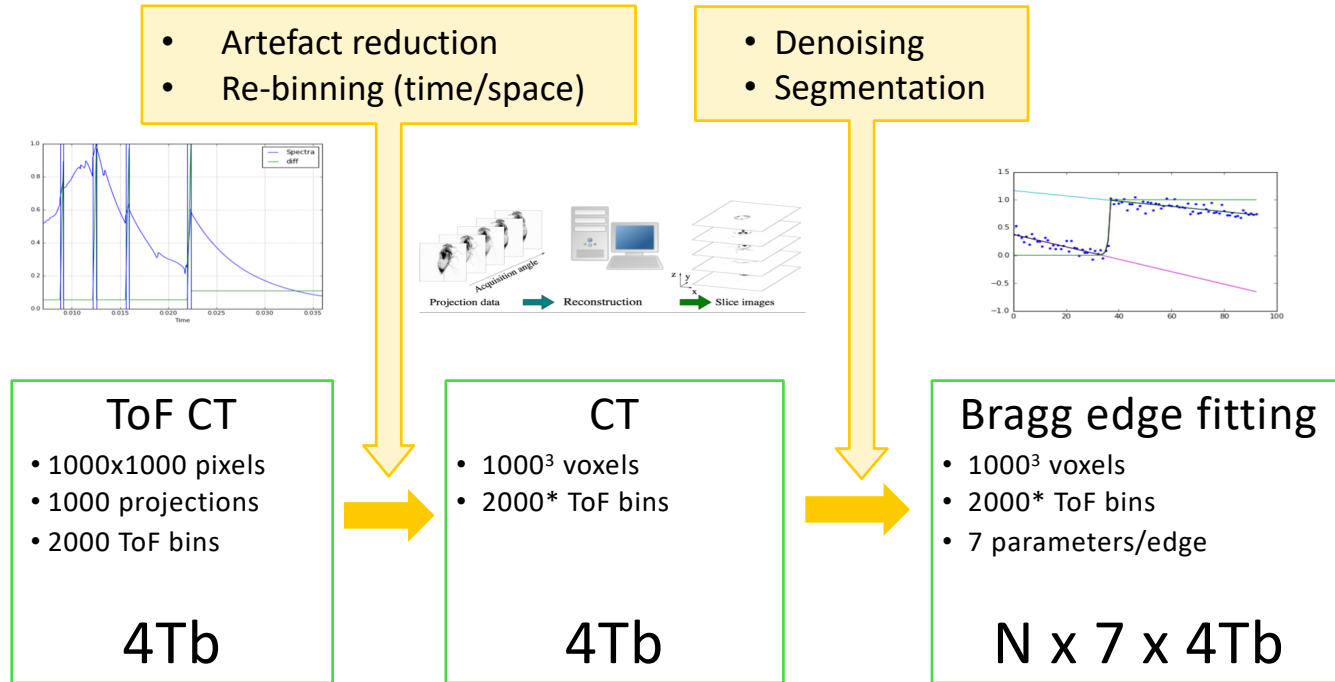
- Segmentation/Classification
- Modelling
- Feature analysis
  - Dimensions
  - Counting
- Etc



→ Template workflows



# New challenges: 4D imaging



- CLI tools
- Python
- Mantid
- KipTool

- MuhRec
- (Octopus)
- Python/Matlab
  - ASTRA
  - TomoPy

\* 2000 volumes is a worst case scenario.

- RITS
- iBeatles
- ToFImaging (under development)

\* 2000 volumes is a worst case scenario.

# Scalability – Computing infrastructure needed

Laptop



Pre-evaluation  
Training sessions

Small data

Workstation



Most evaluation  
Explore data

Mid sized data

Cluster



Mass evaluation

Full sized data

Software needs to be scalable to support different infrastructure

# Data management – The FAIR principle

Findable

Accessible

Interoperable

Reproducible

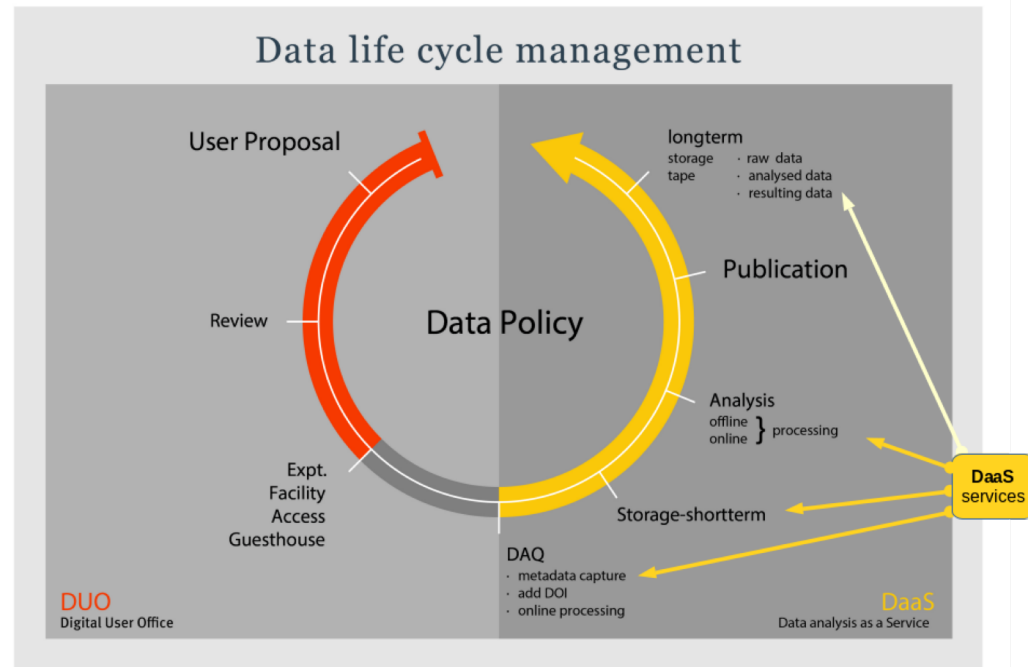
A requirement from funding agencies

- Recording meta data
- Electronic log books  
(no more paper and glue)



- Storage
- Data identifiers

- Common and efficient data formats



# Developing analysis software



# Developing software for neutron imaging

*Neutron imaging: The final frontier – These are the experiments by many scientists – Their many year mission – To explore strange new worlds – To seek out new samples and new processes*

*... To boldly go where no man has gone before.*



## Why?

- Explore new scientific ideas.
- Flexibility and experimental creativity.
- No commercial options available.
- To provide open source alternatives.

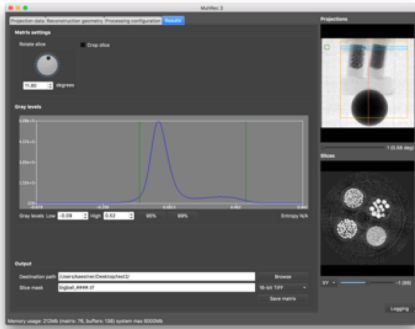
## How?

- Scripting languages
- High performance languages
- Existing platforms/from scratch

GUI applications are great for specific tasks that require interaction.

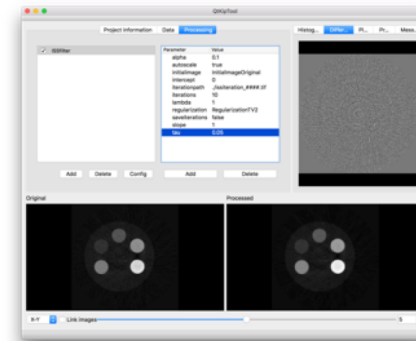
## MuhRec

CT reconstruction tool



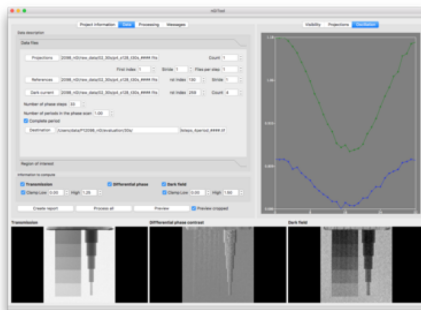
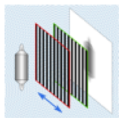
## KipTool

General processing tool for 3D images



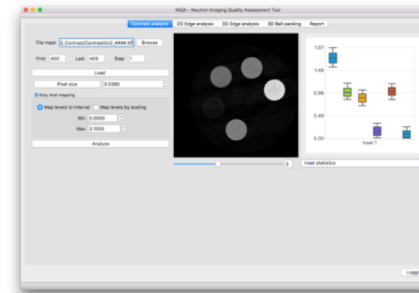
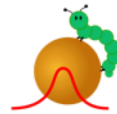
## nGI tool

Reduction of phase stepping scans

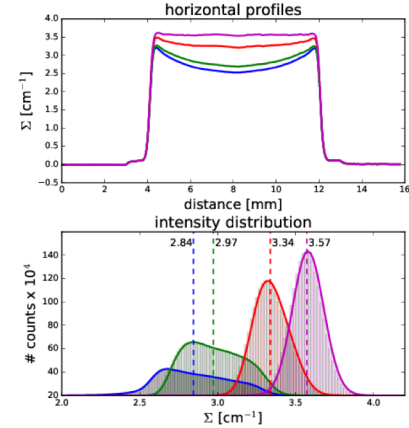
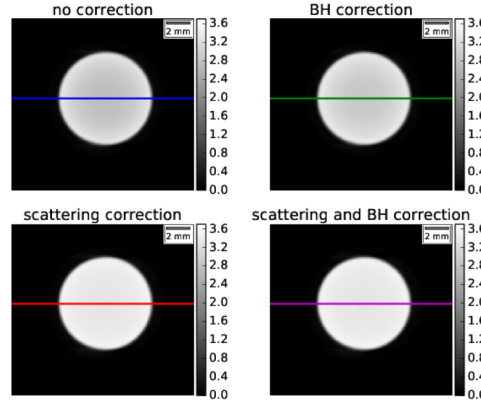
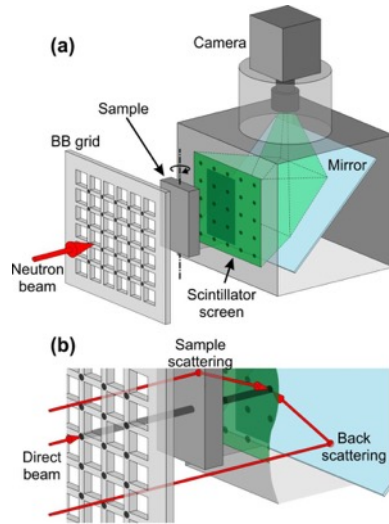


## NIQA tool

Analysis of IAEA QA samples



# The effect of scattering and its correction



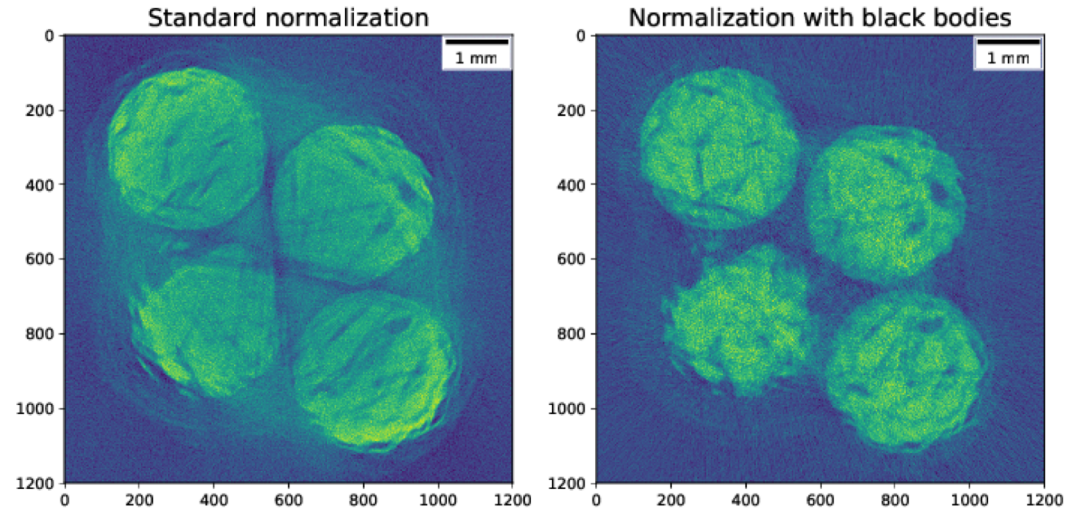
— no correction  
— BH correction  
— scattering correction  
— scattering and BH correction

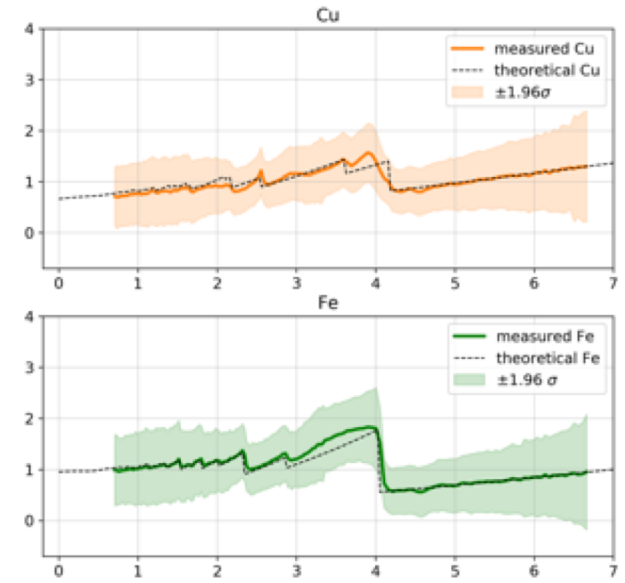
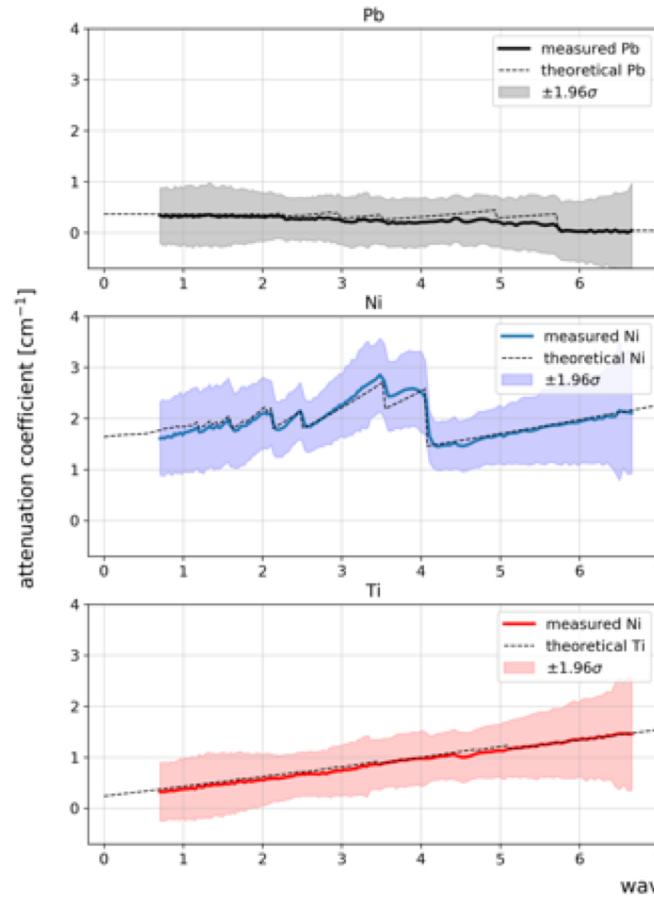
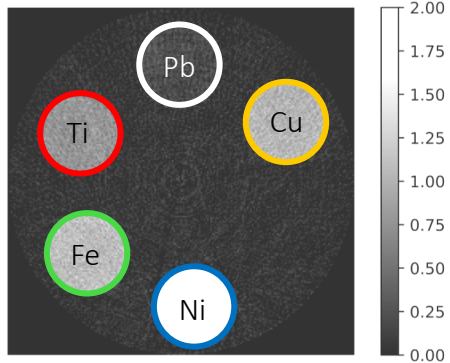
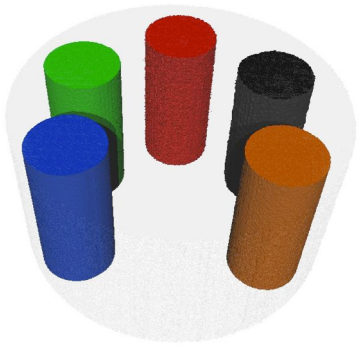
## Correction using revised normalization

$$\frac{\tilde{I}_n}{\tilde{I}_{OB}} = \frac{I_n}{I_{OB}} \cdot \frac{D(I_{OB})}{D(I_n)}$$



$$\frac{\tilde{I}_n}{\tilde{I}_{OB}} = \frac{I_n}{I_{OB}} \cdot \frac{D(I_{OB})}{D(I_n)} = \frac{I_n^* - I_{DC} - I_{n,BB}^S \frac{D(I_n^* - I_{DC})}{D(I_{OB}^* - I_{DC}) - (1 - \frac{1}{\tau_{BB}}) I_{n,BB}^S \tau_{BB}}}{I_{OB}^* - I_{DC} - I_{BG,BB}^S \frac{D(I_{OB}^* - I_{DC})}{D(I_{OB,BB}^* - I_{DC}) - (1 - \frac{1}{\tau_{BB}}) I_{BG,BB}^S \tau_{BB}}} \cdot \frac{D(I_{OB}^* - I_{DC})}{D(I_{OB,BB}^* - I_{DC}) - (1 - \frac{1}{\tau_{BB}}) I_{BG,BB}^S \tau_{BB}} \cdot \frac{D(I_n^* - I_{DC})}{D(I_{n,BB}^* - I_{DC}) - (1 - \frac{1}{\tau_{BB}}) I_{n,BB}^S \tau_{BB}}$$





Scripting is the choice for mass production and increased repeatability.

## Development focus

- Time-of-flight tool box (Bragg-edge fitting is central)
- Provide workflow notebooks
  - Experiment setup
  - Time series analysis
  - Mass reconstruction
  - ...
- Provide an interface to C++ algorithms
  - Spot cleaning
  - Denoising
  - Scattering correction
- Interoperability with multi-dim array module SCIPP (<https://scipp.github.io/>)

# Analysis using Jupyter notebooks

## Notebooks

- Provide some interaction
- Are less intimidating than writing pure code
- Allow to document the analysis steps

The screenshot shows a JupyterLab notebook titled "TimeSeriesNormalization.ipynb". The interface includes a file browser on the left, a menu bar (File, Edit, View, Run, Kernel, Tabs, Settings, Help), and a main content area. The content area contains the following text and code:

**Normalize**

The time series must be normalized using Beer-Lambert law

$$I = I_0 e^{-\int_L \mu dx}$$

before any further analysis. In practice this looks like if you set the `neg log=True`

$$\int_L \mu dx = \frac{D_0}{D} \cdot \frac{I_{sample} - I_{DC}}{I_0 - I_{DC}}$$

where  $D$  and  $D_0$  are scalars computed as the average of the doseROI area.

```
[ ]: nsample = amg.normalizeImage(sample, ob, dc, neg log=True, doseROI=[200, 20, 400, 100])
```

```
[ ]: N = 5 # number of images to show
```

```
if nsample.shape[0] < N :
    N = nsample.shape[0]
```

```
fig, ax = plt.subplots(1, N, figsize=(15, 6))
ax = ax.ravel()
```

```
for idx in range(N) :
    amg.imshowPercentile(sample[idx], ax[idx], factor=1.96)
```

**Observe relative changes**

If you want to compare the changes between different sample conditions, you have the following condition

- Initial image  $I_{T=0} = I_0 e^{-L_{media} \mu_{media}}$
- Image at time  $T$   $I_{T=T} = I_0 e^{-(L_{H_2O} \mu_{H_2O} + L_{media} \mu_{media})}$

The status bar at the bottom indicates "Python 3 | Idle", "Mode: Command", and "Ln 1, Col 1 TimeSeriesNormalization.ipynb".

- Agile development
  - Iterative development
  - Collaboration
- Using repositories
  - New issues in repository branches
  - Maintain stable master through merge reviews
- Issue tracking (New features, Improvements, Fixing bugs)
- Frequent releases (2 x year)
- Automated testing and builds
- Online documentation (wiki)



# Some technical details about the development

## Main coding language



In particular: C++11

- XCode (Mac)
- MSVC15 (Windows)
- g++ (Linux)

## GUI Library



Version: 5.15 LTS

- Cross platform
- Qmake/CMake
- QTest

## Python bindings



Python 3 + PyBind11

- *Embedded scripting*
- Call libs from python
- *Tutorials*
- *Typical work flows*

## Repository: GitHub

<https://github.com/neutronimaging>



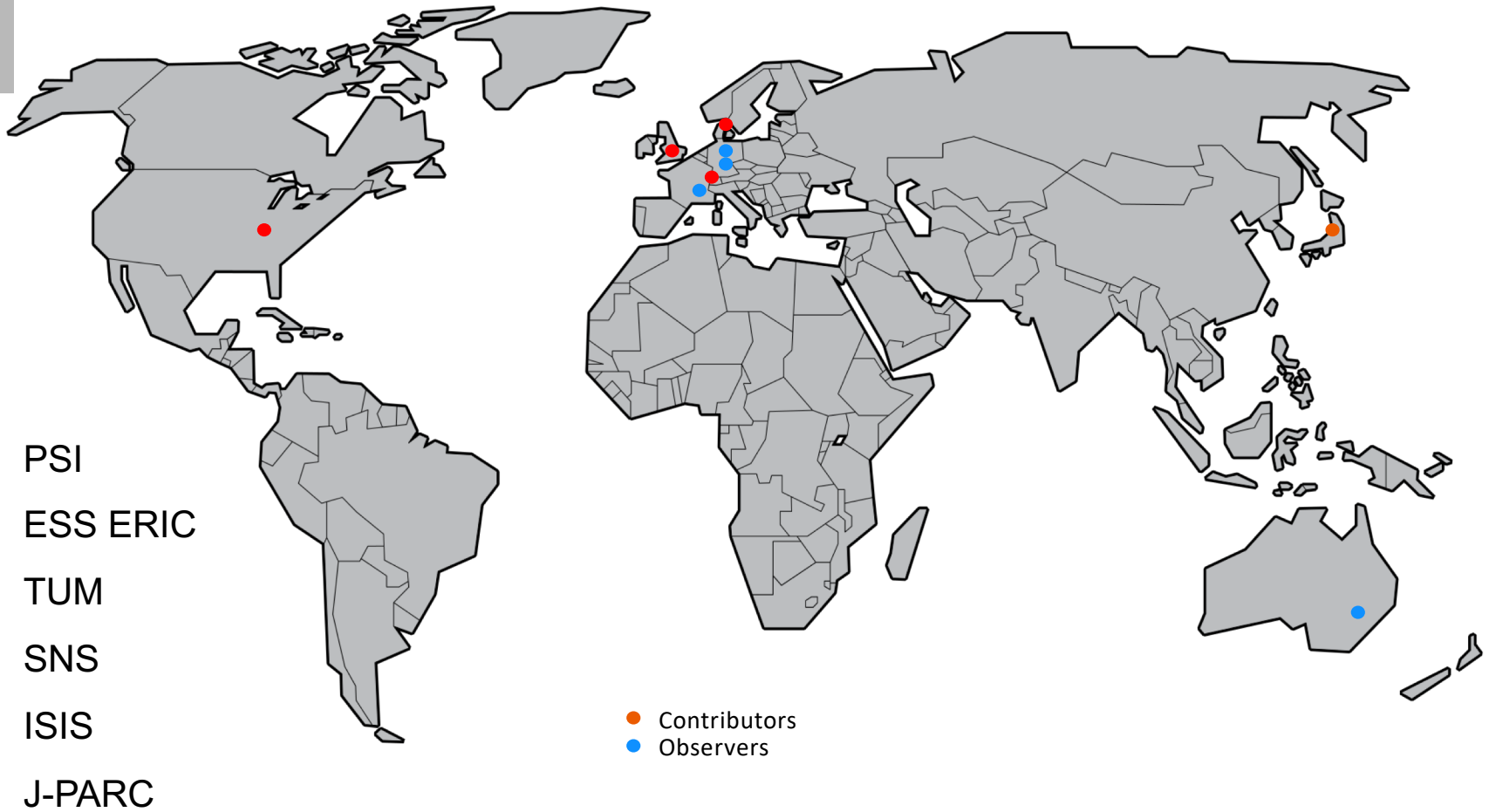
- Development history
- Issue tracking
- Documentation (Wiki)

## Build server: Jenkins



- Automated builds
- Nightly build installers





Recent new collaboration efforts with TUM and ILL

## The analysis of imaging data should be ...

- ... done in a reproducible way.
- ... open source to promote collaboration.
- ... developed with modern techniques.

We welcome new partners to join.

<https://neutronimaging.github.io/>



# Acknowledgements

## Development team

- Chiara Carminati
- Matteo Busi

## User feedback

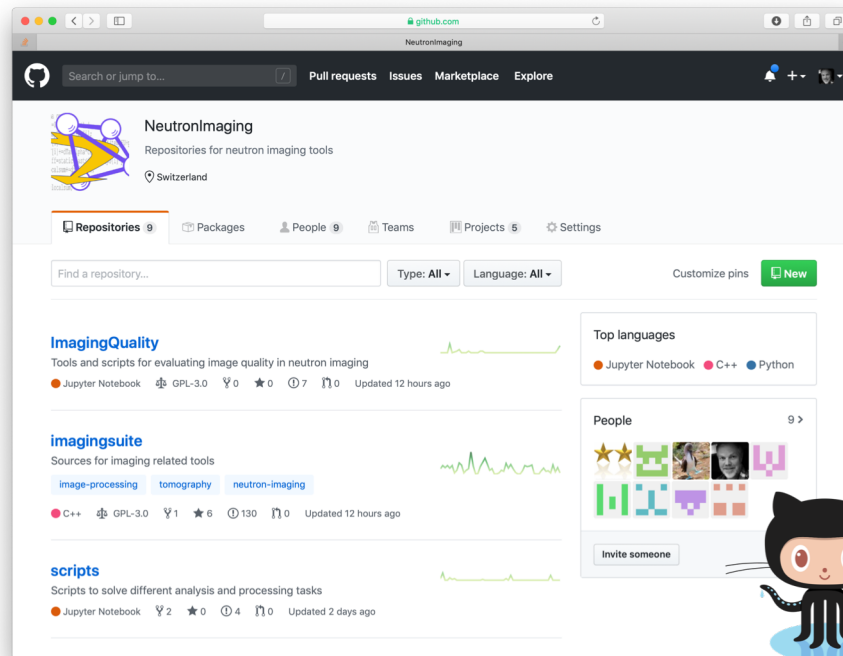
- AMG
- Experiment users

## Funding

ESS - DMSC



All developed code shall be open source

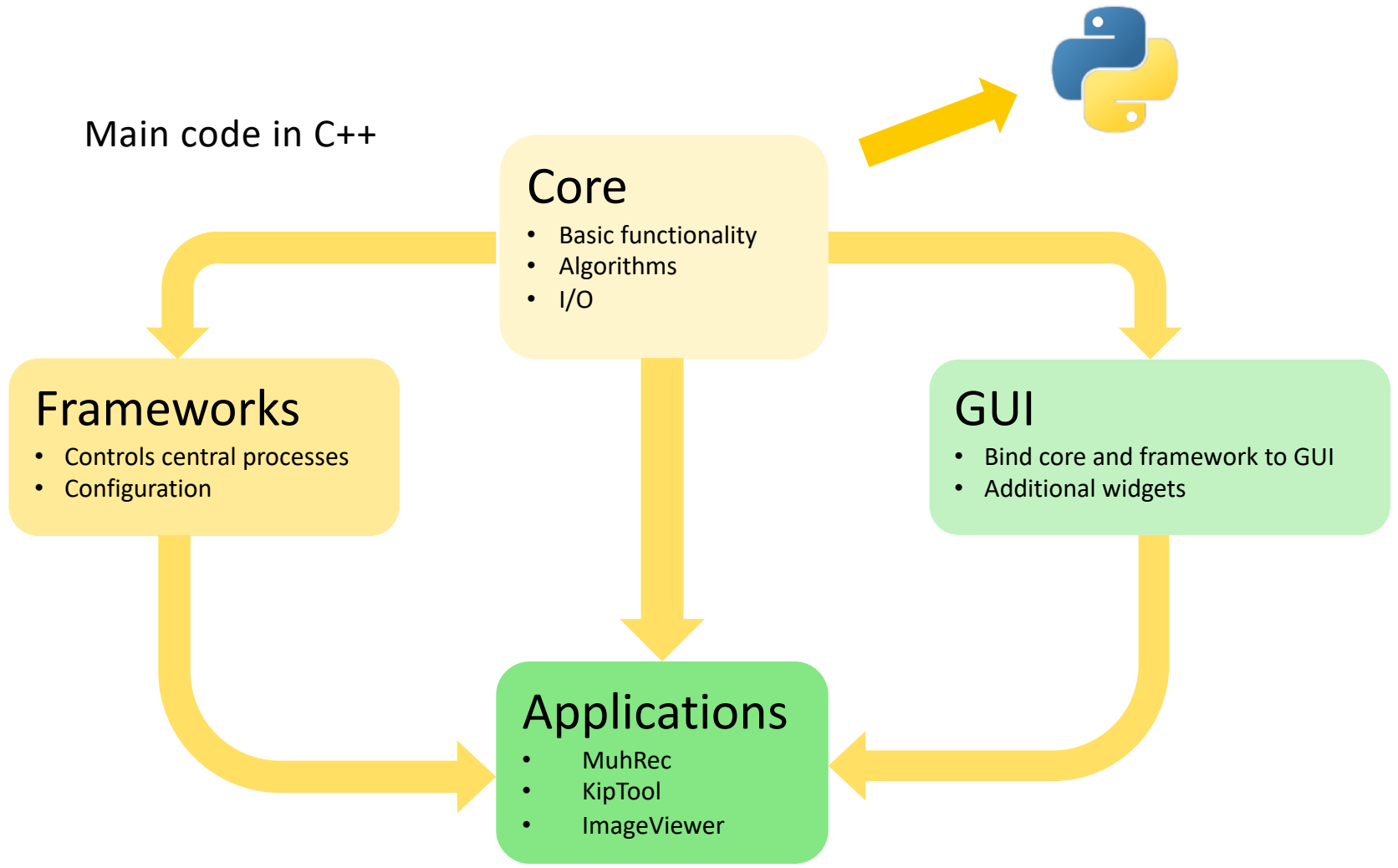


Done

- Most of the code is transferred
- License GPL 3.0
- Issue tracking (GitHub)
- Submit issue link in apps
- Transfer remaining code
- Prepare build scripts
- Setup automated build
- Prepare build instructions
- Documentation on Wiki

Source repository <https://github.com/neutronimaging>

# Design – Strict separation per purpose

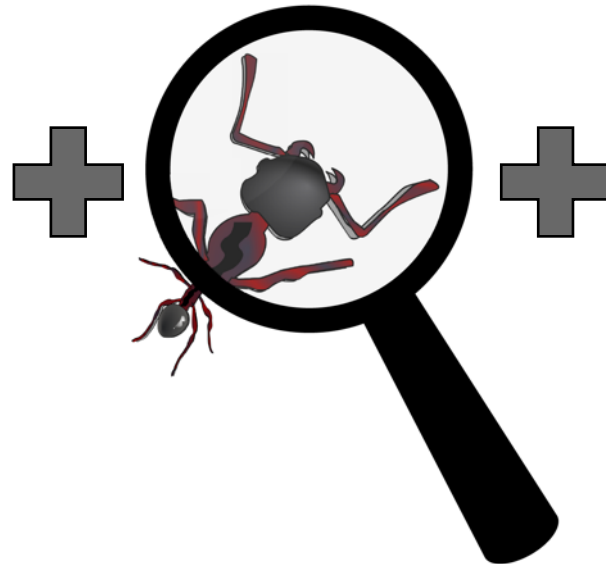


Our experiment users want ...

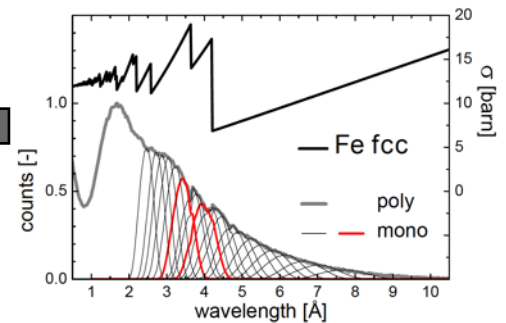
Fast acquisition



High resolution



Monochromatic



But the number of neutrons is limited...

## Data quality

- Limited number of observations
- Low SNR
- Less sharp than X-rays

*long experiment times*

*few detected neutrons per pixel*

*detection principle*

## Data structure

- Single modality
- Temporal dependency
- Multi modality
- Multi wavelength
- Tensor valued

*traditional transmission imaging*

*time series*

*combining images from different sources*

*material response, mixes*

*magnetism, strain, diffraction*

Investigations often only involve few or even single samples ...

Is generalization possible?