

# dpdak

directly programmable data analysis kit

Gunthard Benecke  
HDRI-PNI  
27.02.2012



**Max Planck Institute  
of Colloids and Interfaces**

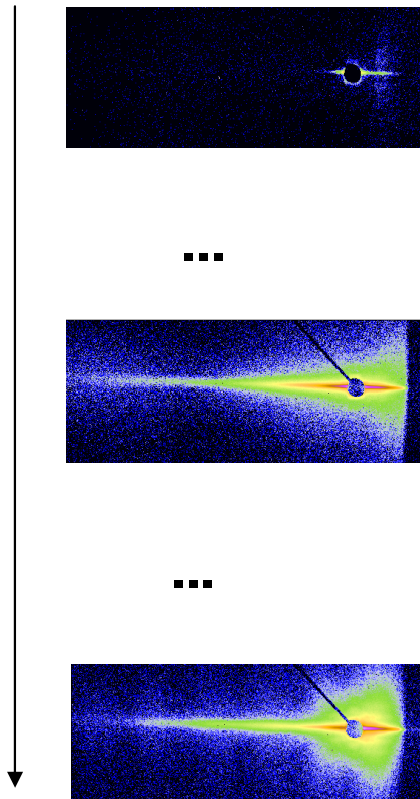
# Overview

- > Plugin framework for primary online analysis of SAXS/WAXS and GISAXS experiments
- > Developed in a DESY and MPIKG cooperation
- > Goals of the project:
  - fast analysis allowing to classify the data during the experiment
  - deeper analysis of data after experiemnt
  - extending functionality by plugins



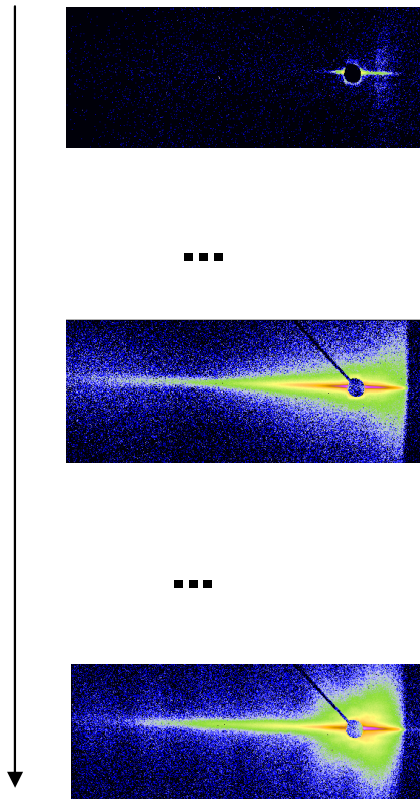
# Basic processing schema

## > Incoming Data

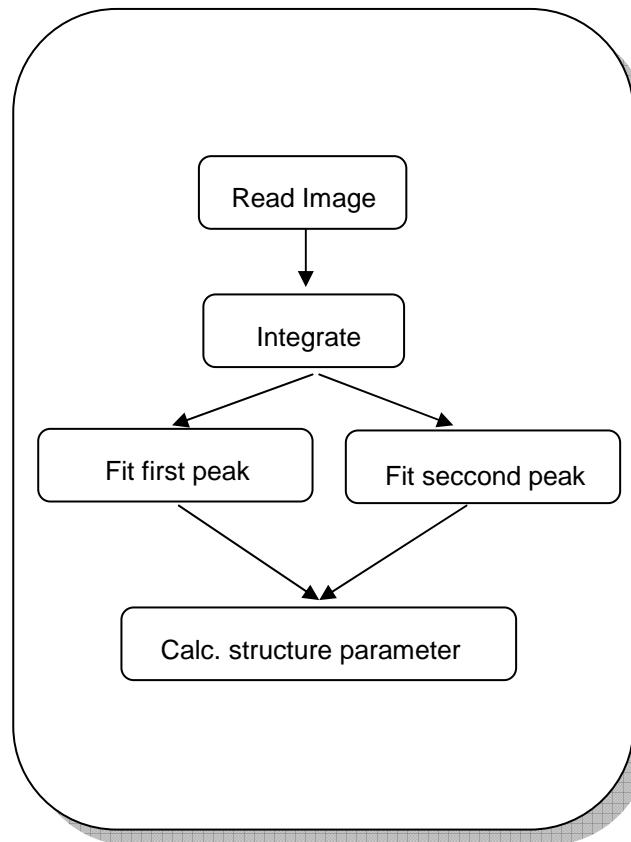


# Basic processing schema

## > Incoming Data

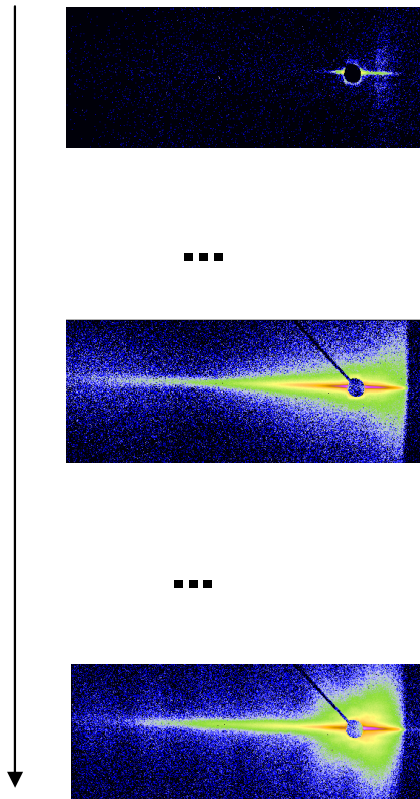


## > Data reduction

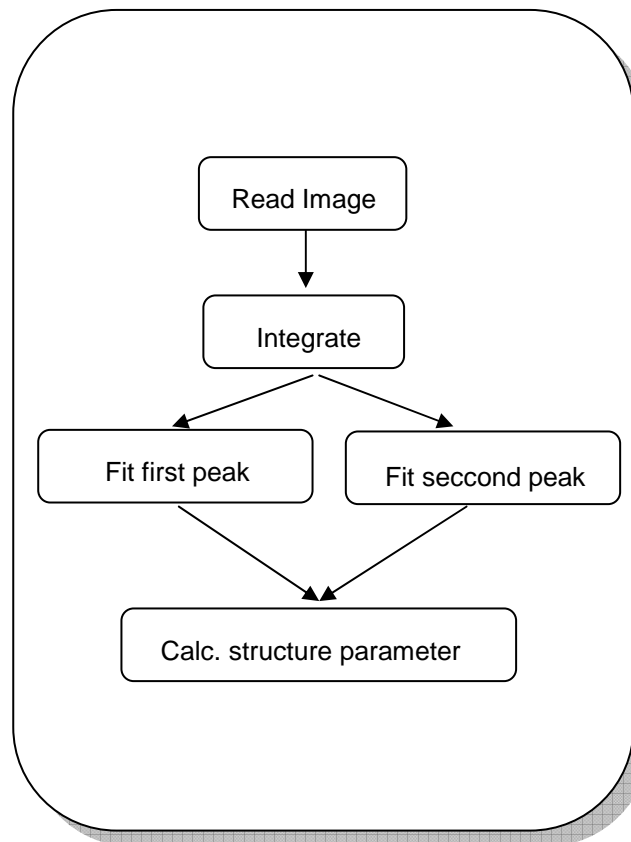


# Basic processing schema

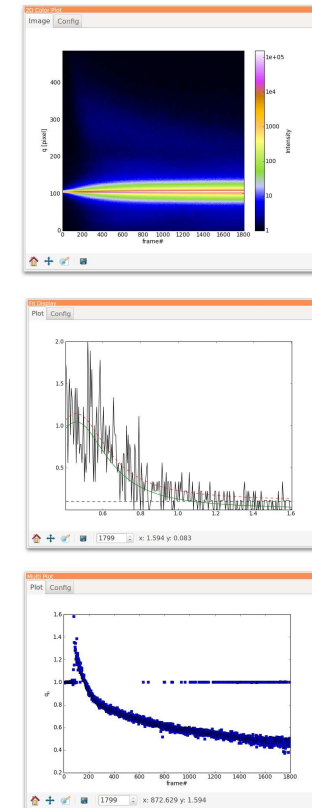
## > Incoming Data



## > Data reduction



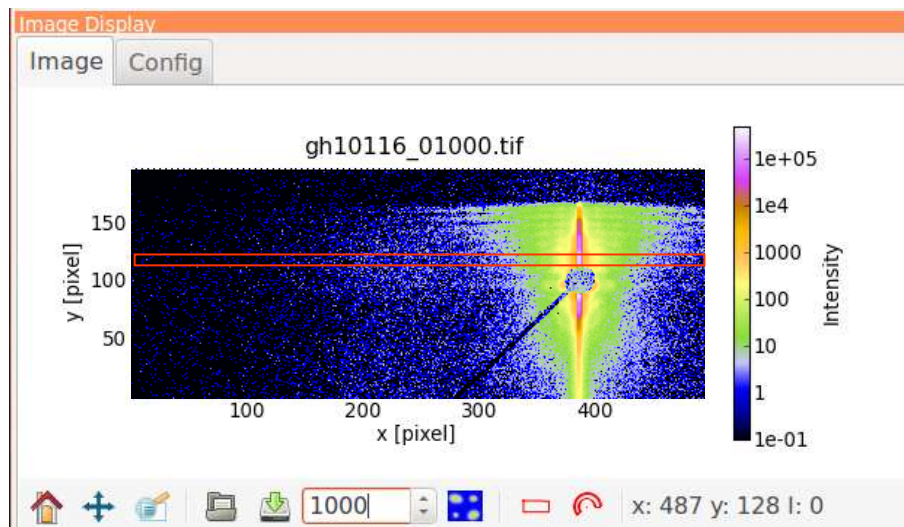
## > Display results



# Use Case I: Sputter Experiments

➤ Online (up to a frame every 20ms) plot of cuts show lateral structure

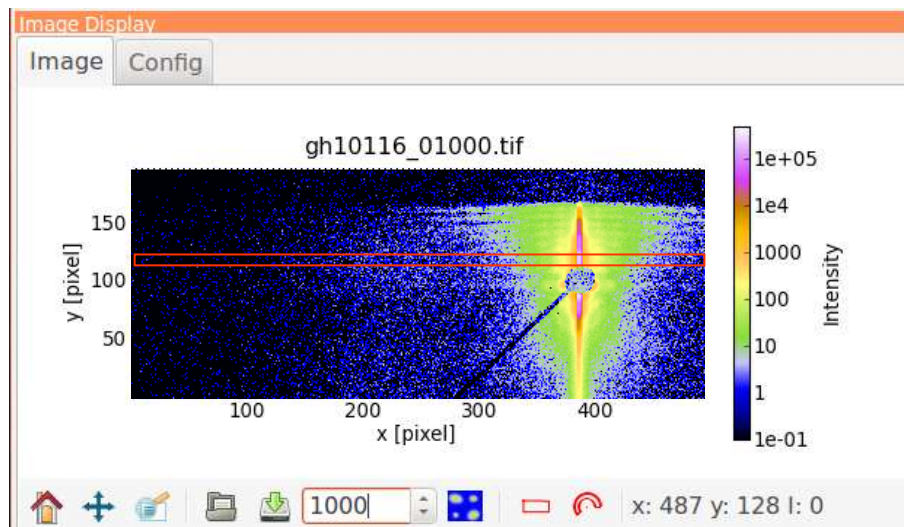
➤ Detector image with cut tool



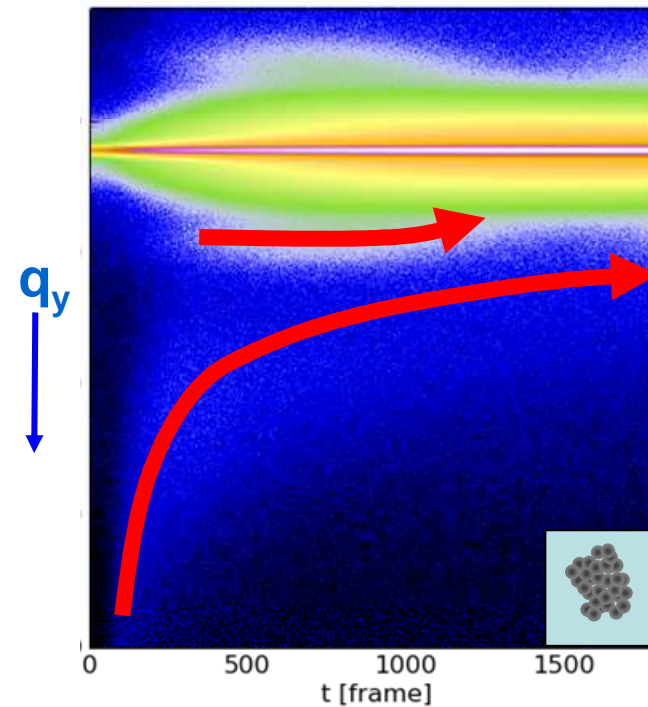
# Use Case I: Sputter Experiments

> Online (up to a frame every 20ms) plot of cuts show lateral structure

> Detector image with cut tool

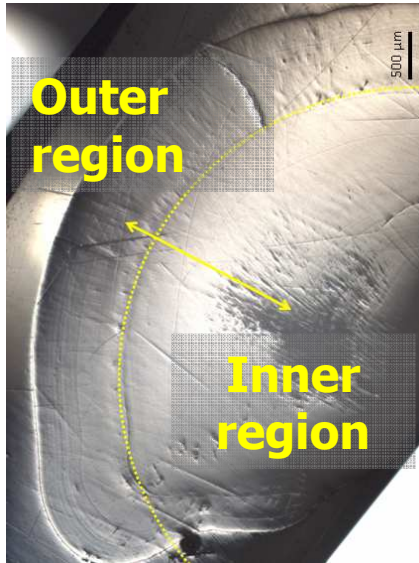


> Broad structure at ~30nm  
> Co nanoparticles, R(Co) growing



# Use Case II: Crayfish Gastrolith Nanostructure

## > Sample Data

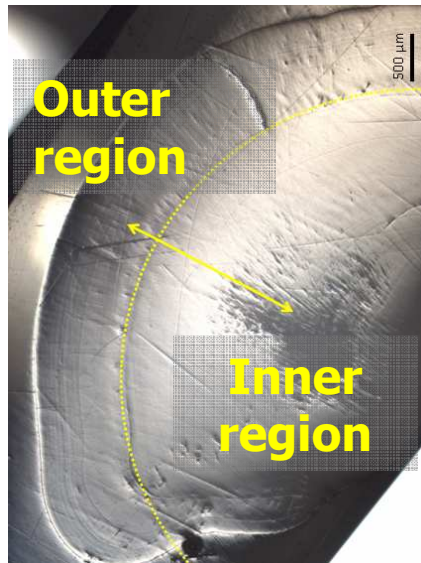


- 2D scan of 150 mccd detector files
- 10 samples scanned
- 20GB data



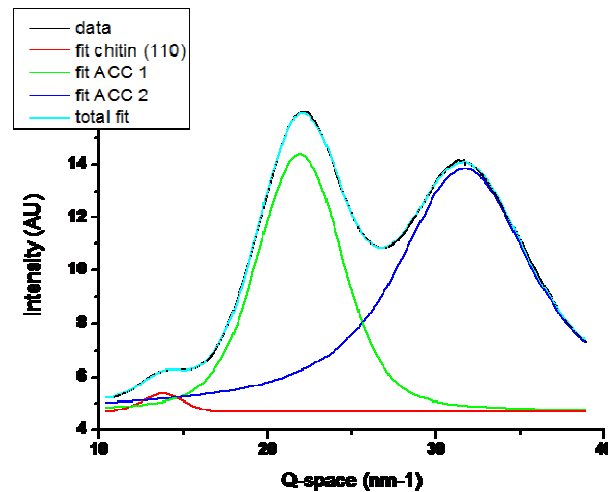
# Use Case II: Crayfish Gastrolith Nanostructure

## > Sample Data



- 2D scan of 150 mccd detector files
- 10 samples scanned
- 20GB data

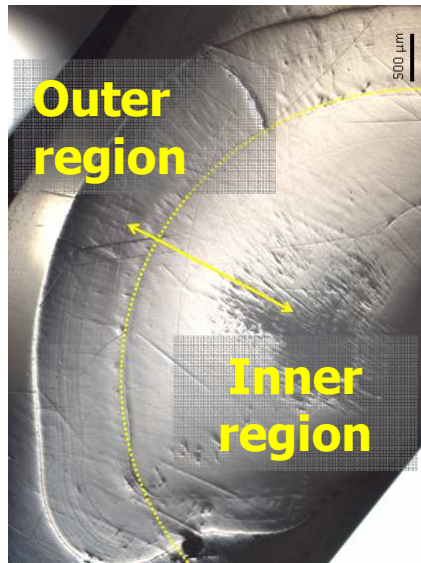
## > Processing



- integrating 2D data
- fitting chitin and ACC peaks
- calculate area of peaks and ratios

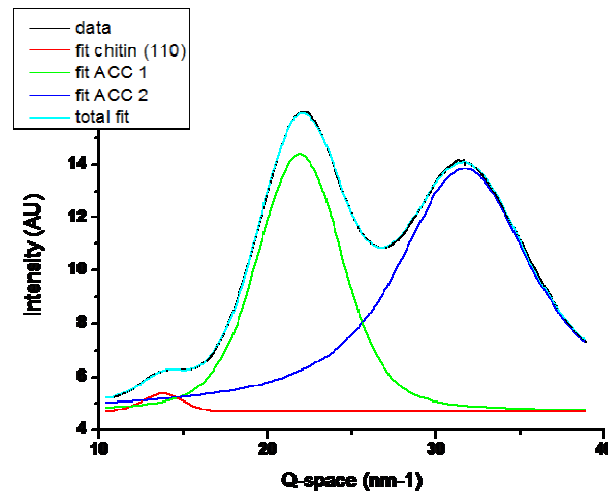
# Use Case II: Crayfish Gastrolith Nanostructure

## > Sample Data



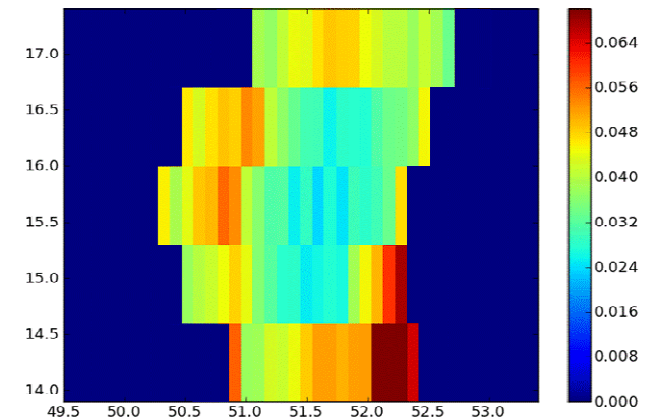
- 2D scan of 150 mccd detector files
- 10 samples scanned
- 20GB data

## > Processing



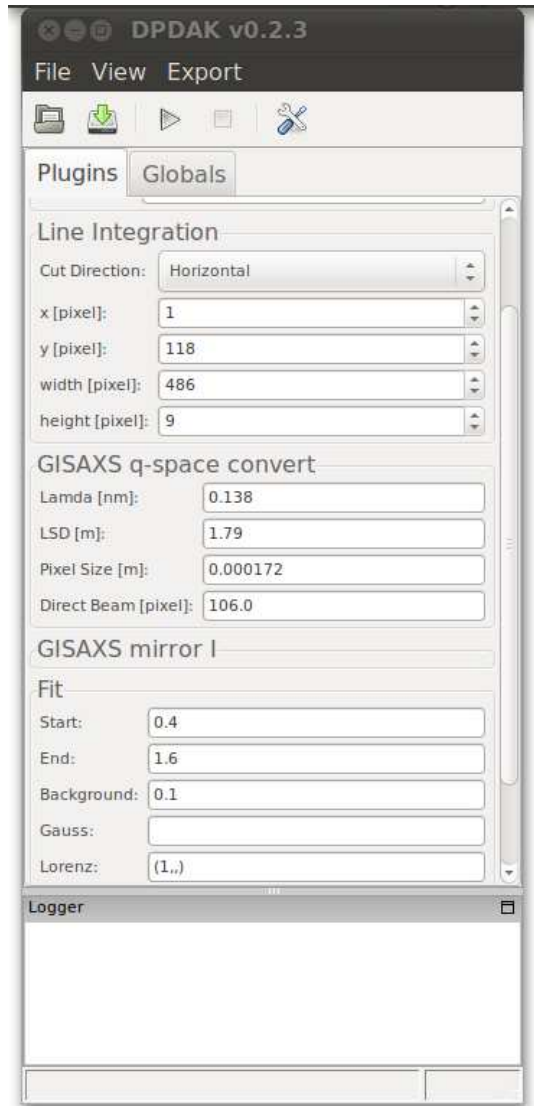
- integrating 2D data
- fitting chitin and ACC peaks
- calculate area of peaks and ratios

## > Result



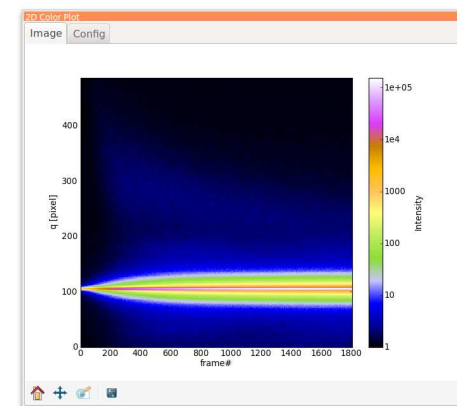
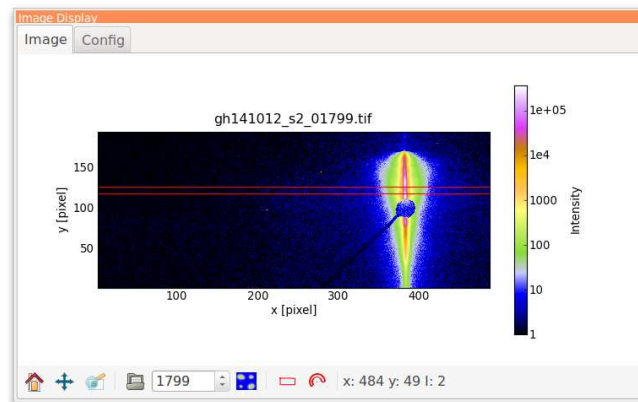
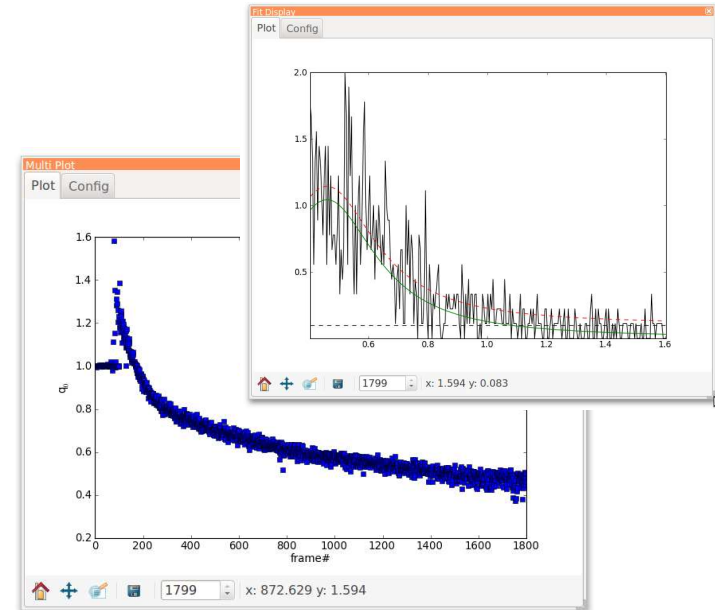
- 2D plots of ratios
- created online during the scan

# Screenshot of a analysis session



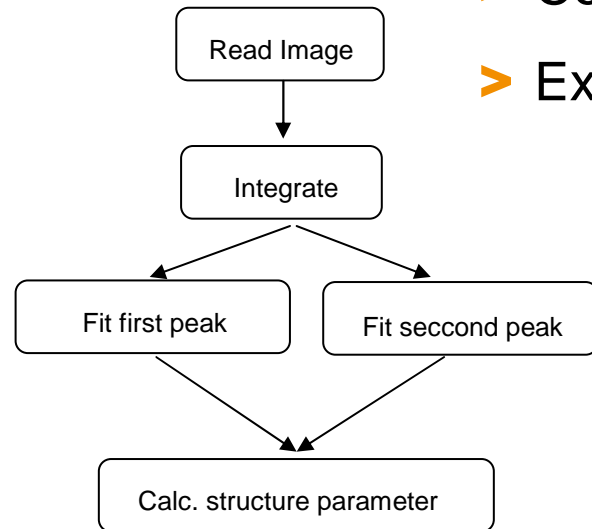
> Configuration window for plugins

> Displays for analysis results



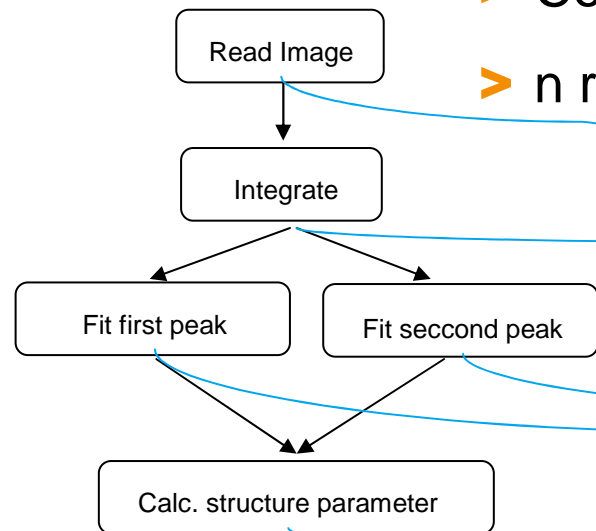
# Plugin Framework

- > Connected plugins for an analysis
- > Execute analysis for each data point in the dataset



# Plugin Framework

- > Connected plugins for an analysis
- > n runs for n datasets (images, etc.)

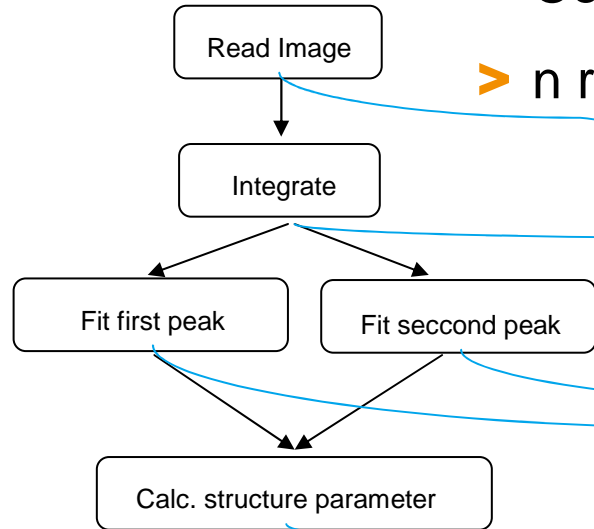


- > Plugin output written into a internal database

Out1	Out2	Out3	Out4	Out5
frame1	cut1	pos1	pos1	param1
frame2	cut2	pos2	pos2	param2

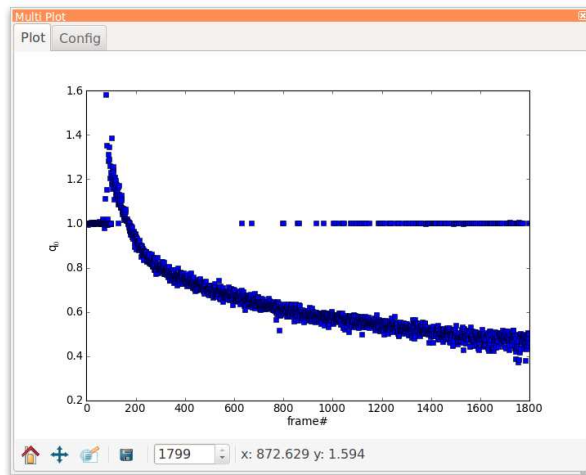
# Plugin Framework

- > Connected plugins for an analysis
- > n runs for n datasets (images, etc.)



- > Plugin output written into a internal database

Out1	Out2	Out3	Out4	Out5
frame1	cut1	pos1	pos1	param1
frame2	cut2	pos2	pos2	param2

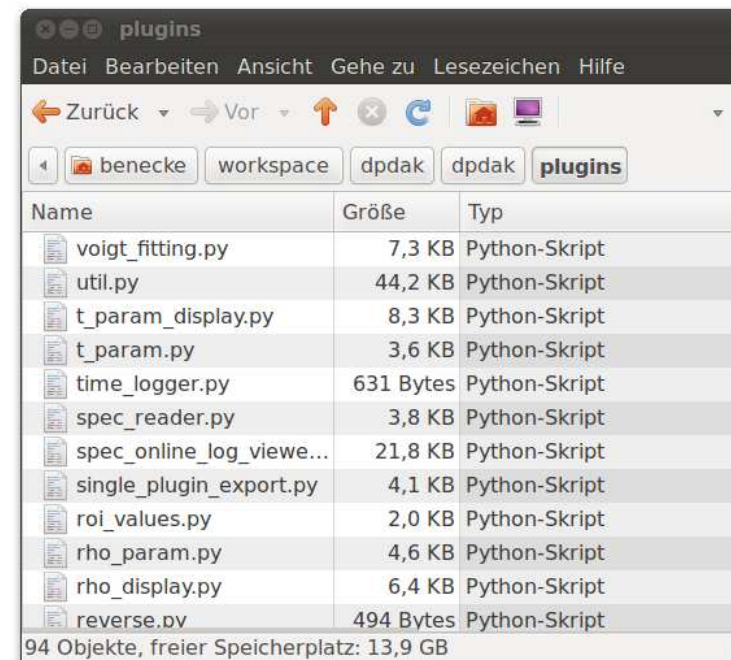


- > Display plugins read the database and visualize the results



# Plugins

- Plugins are python scripts
- Framework import plugin scripts on runtime
- Plugin interfaces for:
  - Data input & analysis (base plugins)
  - Visualisation (display plugins)
  - Export to different formats (export plugins)
- Base plugins are well defined:
  - Typed parameter
  - Typed inputs
  - Typed outputs



# Plugin Example

```
1 import numpy
2
3 import dpdak
4
5 class ROIValues(dpdak.BasePlugin):
6
7     def get_info(self):
8         info = dpdak.PluginInfo()
9         info.set_info('ROI Values', 'Gunthard Benecke', '1.0')
10
11         info.add_input('Image', dpdak.IMAGE)
12
13         info.add_output('Sum', dpdak.SCALAR)
14         info.add_output('Mean', dpdak.SCALAR)
15         info.add_output('Min', dpdak.SCALAR)
16         info.add_output('Max', dpdak.SCALAR)
17
18         info.add_parameter('x', dpdak.INTEGER, 0)
19         info.add_parameter('y', dpdak.INTEGER, 0)
20         info.add_parameter('width', dpdak.INTEGER, 10)
21         info.add_parameter('height', dpdak.INTEGER, 10)
22
23         return info
24
25     def process(self, counter, parameter, input):
26         x, y = parameter['x'], parameter['y']
27         width, height = parameter['width'], parameter['height']
28
29         image = input['Image'][y:y+height, x:x+width]
30         data = {}
31         data['Sum'] = image.sum()
32         data['Mean'] = sum / (image.shape[0] * image.shape[1])
33         data['Min'] = image.min()
34         data['Max'] = image.max()
35
36         return (dpdak.STATUS_OK, data)
```





# Features

## > Software

- Python + numpy, scipy, wx, matplotlib, PIL, h5py
- Windows (XP/Vista/7), Linux
- GNU GPL licence

## > Formats

- Detector images (tif, cbf, mccd, edf, ima, txt, ...)
- SPEC & Hasylab Online log file → **a standart data format would improve usability**
- Spreadsheet text files

## > Analysis

- Integration (azimuthal/radial) & cuts
- Basic fitting
- Structure parameter (Rho/T/L)

## > Visualisation of detector images, 2D maps of cuts and plots



# Contact

## > Web:

- <http://www.desy.de/~beneckeg/dpdak/>
- <http://dpdak.desy.de> (wiki, March 2012)

## > SVN:

- <https://svnsrv.desy.de/viewvc/dpdak/>
- open hoster (sourceforge, github, google google)

