# Handling different analysis workflows in a modular framework

**Malte Storm** 

Helmholtz-Zentrum Hereon, Institute of Materials Physics

deRSE 2024, Würzburg







## **Outline**

- Motivation of the scientific background
- Individual components of the solution
- Interaction of components and global solution architecture
- Conclusion



# **Motivation**

#### Helmholtz-Zentrum Hereon operates multiple X-ray experiments at the PETRA III storage ring (DESY, Hamburg)

- In situ nano diffraction (P03)
- Nano tomography (P05 EH1)
- Micro tomography (P05 EH2)
- High energy diffraction (P07 EH3)
- High energy micro tomography (P07 EH4)
- White beam diffraction (P61A)

In this talk: Focus on analysis solution for diffraction experiments





# **Exemplary samples**



TiN steel coating for tools Zeilinger et al. Scientific Reports 6 (2016), 22670



Tracheid wood cells Storm et al. J. Synchrotron Rad. 22 (2015), 267



In situ laser beam welding RWTH Aachen, 2022



FlexiStir in situ friction stir welding Blankenburg, PhD thesis, Kiel 2016



Modular analysis workflows, deRSE2024

# Motivation for new software solution

Political mandate to make experiments and analysis accessible for all domain scientists, not just experts in diffraction analysis

#### **Old analysis pipeline example:**





# Motivation for new software solution

Political mandate to make experiments and analysis accessible for all domain scientists, not just experts in diffraction analysis

#### **Old analysis pipeline example:** jupyter X Raw data inspection Data visualization: Data processing: Data calibration: (e.g. hdf5): python script, Excel, ... python script / gui pyFAI-calib2 python script / gui pydidas

# pydidas - Python Diffraction Data Analysis Suite

#### **Key requirements:**

- Easy to use and with graphical user interface
- Enable fast analysis during experiments
- Have a tool which is widely deployable and scalable

#### pydidas Python package with

- Graphical user interface
- Low-level access to all functionalities (e.g. for advanced scripting)
- Uses Qt for GUI, persistent configuration storage and communication





# **Pydidas workflow requirements**

#### **High flexibility required**

- different processing strategies required for
  - different analysis goals
  - different materials
  - different experimental conditions
- Generic functionality sufficient for routine experiments but specialist analysis methods required in some cases
   → implemention to extend generic functionality required
- $\rightarrow$  Workflow implemented based on a tree and plugins.





# **Plugins**

#### Plugins structure must be simple to allow non-experts to extend functionality

#### **Minimum requirements:**

- Classes must inherit from base classes for automatic discovery
- Some basic class attributes (plugin metadata)
- pre\_execute method
- execute method

#### Can be extended with e.g.

- Custom configuration widgets
- Storage of detailed intermediate results

```
plugin_type = PROC_PLUGIN
plugin_subtype = PROC_PLUGIN_IMAGE
basic_plugin = False
plugin_name = "Crop and bin image"
output_data_label = "Image intensity"
output_data_unit = "counts"
input_data_dim = 2
default_params = get_generic_param_collection(
    *param_keys: "Use_roi",
    "roi_xlow",
    "roi_xhigh"
    "roi_ylow",
    "roi_yhigh"
    "binning",
def __init__(self, *args: tuple, **kwargs: dict):
   super().__init__( *args: *args, **kwargs)
    _params = self.get_params(
        *param_keys: "use_roi", "roi_xlow", "roi_xhigh", "roi_ylow", "roi_yhigh", "binning"
    self._image_metadata = ImageMetadataManager(*_params)
def calculate_result_shape(self):
   if self._config["input_shape"] is None:
       raise UserConfigError(
            "No input shape has been given for the *CropAndBinData* plugin."
       )
    self._image_metadata.store_image_data(self._config["input_shape"], int, n_image: 1)
    self._image_metadata.update_final_image()
    self._config["result_shape"] = self._image_metadata.final_shape
def execute(self, data: Dataset, **kwargs: dict) -> tuple[Dataset, dict]:
    _roi = self._image_metadata.roi
    _binning = self.get_param_value("binning")
    if _roi is not None:
       data = data[_roi]
    if _binning > 1:
        data = rebin2d(data, _binning)
    return data, kwargs
```

class CropAndBinImage(ProcPlugin):



9

## **Plugins access**

Plugins are made accessible through a registry

- Plugin paths are managed through a Qsettings variable
- Registry scans all python files in plugin paths and stores classes which inherited from BasePlugin
- Registry offers access to
  - Plugin classes
  - Plugin names (also filtered by type)
  - Stored paths





# **Plugins access in the Ul**

#### Widget for browsing plugins

- Display all available plugins
- Search for plugins
- Display additional information about plugins

Available plugins:		
Search filter 🔇 Rese	t filter	HDF5 file series loader
<ul> <li>Input plugins</li> <li>Eiger scan series loader</li> <li>Fio MCA line scan series loader</li> </ul>		Name: HDF5 file series loader
Single frame loader HDF5 file series loader		Plugin description: Load data frames from Hdf5 data files.
<ul> <li>Generic processing plugins Center of mass 1d data</li> <li>Sum 1D data</li> <li>Sum 2D data</li> </ul>		This class is designed to load data from a series of hdf5 file. The file series is defined through the SCAN's base directory, filename pattern and start index.
<ul> <li>Processing plugins for image da Correct spline distortion</li> <li>Create dynamic data mask</li> </ul>	ta	The final filename is <scan base="" directory="">/<scan for="" hashes="" index="" name="" pattern="" subsituted="" with="">.</scan></scan>
Crop and bin image Mask image pyFAI 2D integration pyFAI azimuthal integration		The dataset in the Hdf5 file is defined by the hdf5_key Parameter. A region of interest and image binning can be supplied to apply directly to the raw image.
pvFAI azimuthal sector integration		Parameters



# **ProcessingTree and WorkflowNode**

- ProcessingTree: Controller class for nodes
- WorkflowNodes: Nodes to populate the ProcessingTree and which include a plugin

#### **ProcessingTree responsibilities:**

- Import and export
- Managing nodes (incl. addition / removal)
- Start of processing

#### WorkflowNode responsibilities:

- Handling of data transport
- Plugin execution
- Knows parent and children





# **Running Workflows**

- ScanContext and DiffractionExperimentContext handle experimental metadata
- WorkflowResultsContext stores results
- ExecuteWorkflowApp handles batch processing and result storage





# **Parallelization**

- Communications with worker processes through queues
- Communication between WorkerController and main ExecuteWorkflowApp with Qt's signals / slots
- Only memory pointers communicated, data shared through shared memory buffer



# **Pydidas results: Metadata**

#### Internal data handling:

 Data stored in subclassed np.ndarray to have metadata attached to raw data

#### **Exporting data:**

- Exports only in hdf5 file format (for now)
- All data axes are labelled and ranges are stored
- All processing metadata (including contexts) is stored in results file





# **FAIR principles**

- Pydidas is open source (GPL 3 license)
- Full architecture is designed for reusability:
  - Components designed with separation of concern in mind
  - Processing logic separated from UI
  - Dynamic UI: Composed of frames; frames usage can be modified in startup script
- To improve: Separate generic components from XRD-specific components to allow re-use with only necessary dependencies



# Conclusion

- Pydidas was develop with X-ray diffraction analysis in mind but framework versatile enough for other applications
- In production and used by users, feedback very positive
- Hereon is committed to maintaining and improving pydidas
- Goal is to build a larger community to extend functionality further



Thank you for your attention

**Any questions?** 

