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A library offering tools to automate acquisition and reconstruction process for Tomography

Break down data treatment into atomic operations that can be associated Orange3 add-on: user friendly canvas, check input and output of each tool,



existing widgets. •Official: https://github.com/biolab/orange3 •Fork used for tomwer: https://github.com/payno/orange3 silx (plot, core functions) •https://github.com/silx-kit/silx Octave fastomo3 script for reconstruction Documentation : http://www.edna-site.org/pub/doc/tomwer/dev/





File Edit View Widget Options Help boxStackViewer last received other Data ..... Visualize /vg-unused/payno/datasets/id17\_christian/new/lbto\_95\_WT\_8weeks\_1\_| Select folder data watche ٩ silx Scans Radios T tomwer 🗙 🎯 🌖 al Y Median 💾 🐨 🛛 📐 — 丨 🔨 🛩 W: 垩 2000 1500 data data list § 1000 500 Friendly user interface to design personalized workflows 1000 1500 2000 2500 -500 Ó 500 Widget displaying results of a reconstruction and asking to the user if he want to validate or not the reconstruction. User can also ask for some modification on the reconstruction parameters Columns X: 1833.969 Y: 2060.013 Data: data transfer 0 0 file : lbto\_95\_WT\_8weeks\_1\_slice\_0043.edf i ±9 # **T** \ II

## Set of tools:

- Tomwer GUI setup example
- Data watcher : detect finished acquisition and data ready to be reconstructed.
- Ftseries reconstruction : launch reconstruction of the acquisition according to user input
- > Data viewer : acquisition and reconstruction viewer

License: MIT Python > 3.4

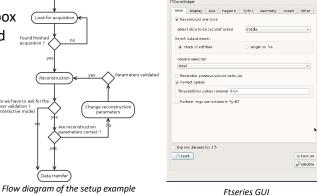
- Data validator : validate a dataset to move to the next box
- >Data list : allow user to define a list of scan to be treated
- Data transfer : transfer dataset from local buffers

## **Status**

 Release 0.1 in September 2017 Tested on ESRF Tomography beamlines •Set of unit test to secure development

## **Perspectives**

•Adapt current tools for Holotomography and Nano-imaging •Deal with scan series and existing post-processing. Add more feature



Monitor workflow status by means of a graylog interface (www.graylog.org)