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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

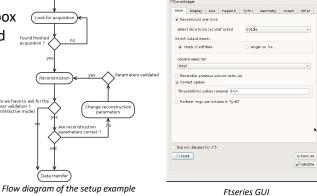
- Data validator : validate a dataset to move to the next box
- >Data list : allow user to define a list of scan to be treated
- Data transfer : transfer dataset from local buffers

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)