TIM2020 Workshop

The presentation and a PDF version of the workshop are available at https://downloads.openmicroscopy.org/presentations/2020/TIM-Munsingen/

Software versions used for this workshop:

- OMERO: 5.6.0
- OMERO.web: 5.6.2
- OMERO.insight: 5.5.9
- OMERO.insight-ij: 5.5.9
- OMERO.iviewer: 0.9.1
- OMERO.figure: 4.2.0
- OMERO.mapr 0.4
- OMERO.parade: 0.2.1
- OMERO.FPBioimage: 0.4.0
- OMERO training scripts: 0.7.3
- OMERO training notebooks: 0.7.2
- OMERO-downloader: 0.2.1
- OMERO.matlab-5.5.3
- OMERO R gateway: 0.4.8
- omero-metadata: 0.5.0
- omero-guides: 2020.02.27
- Bio-Formats: 6.3.1
- Fiji/ImageJ: 2.0.0-rc-69/1.52p
- Matlab R2019a
- Ilastik 1.3.2

Summary

Import
- Cover the various import options
OMERO core concepts
- Data management - Metadata
- Search
- Viewer -3D Viewer
Data mining using OMERO.parade
OMERO figure
Analysis with 3rd party tools
- Analysis with Fiji: manual
- Analysis with Fiji: scripting
Server side scripts
- Analysis in OMERO using Cell Profiler (optional, depending on time)

Programme

Import

In this section we will cover the various import options such as the import with or without data transfer and synchronous vs. asynchronous.

Desktop client install and import

For these 2 workflows shown in the workshop, see
https://omero-guides.readthedocs.io/en/latest/upload/docs/import-desktop-client.html and

Command line import, bulk import, in-place import (for your information only)

These import sections not covered in the workshop can be found at

OMERO core concepts

Data management and cooperation


Viewing images (OMERO.iviewer)


Annotate data and filter using annotations


Search


Viewing images (3D viewer: OMERO.FPBioimage)

OMERO parade

Data mining using OMERO.parade on Projects and Plates

OMERO figure

Fast creation of publication figures using OMERO.figure
See https://omero-guides.readthedocs.io/en/latest/figure/docs/omero_figure.html

Analysis
This part constitutes the core of the training and we will explore the different means OME provides to interact with image and non-image data and how to best integrate these into your workflows.

Analysis with Fiji

- Analysis with Fiji: Java
  - Fiji client side: manual Analysis via UI
  - Fiji client side: scripting: Groovy and Macro
  - Fiji: Analysis in the cloud: Java and Macro

For setup of the Fiji plugin see
For the walkthrough in this workshop, see the four Fiji chapters
https://omero-guides.readthedocs.io/en/latest/fiji/docs/threshold_scripting.html and

Server side analysis

Server-side scripts (python)
For further information about how to write and manage server-side scripts see
Analysis with CellProfiler (shown depending on time)

- Analysis with CellProfiler: Python
  - Analysis in the cloud: Python and using CellProfiler API
See for all CellProfiler workflows
Note that https://mybinder.org/ will be used for CellProfiler setup as described in
https://github.com/ome/omero-guide-cellprofiler

Export (for your information only)

Analysis in R (for your information only)
See for R analysis

Analysis with Ilastik (for your information only)

- Analysis with Ilastik: Python
  - Manual Analysis via UI
  - Analysis in the cloud: Python
See for both setup and workflows

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Optional analysis (for you information only)
See for Python scripts (for your information only)
See for Java scripts (for your information only)