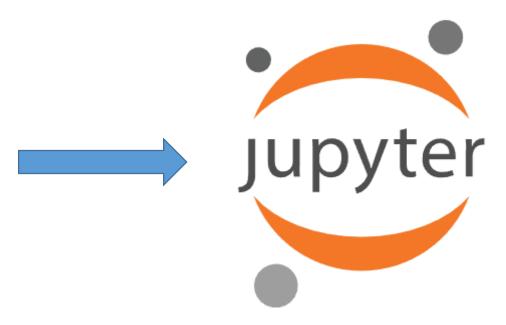




# From Assistant to Notebooks







With material from: Robert Haase

#### Getting started



 Follow this blog post to install python, mamba and devbio-napari: https://t.ly/0BXB





https://t.ly/DVsyu



### Download sample data and notebooks:

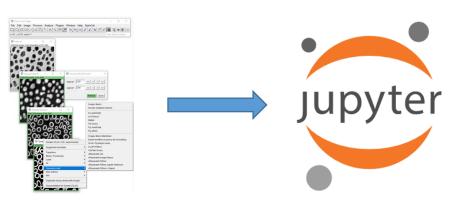


https://t.ly/qlma





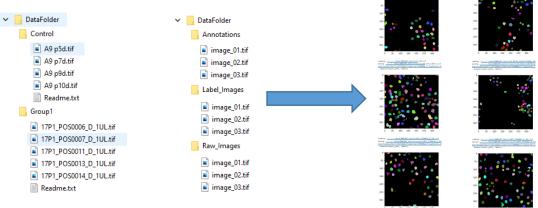
Exporting workflows as notebooks



Python programming basics

Processing folders of images









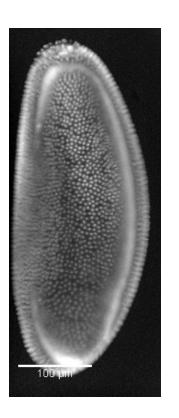
Load data

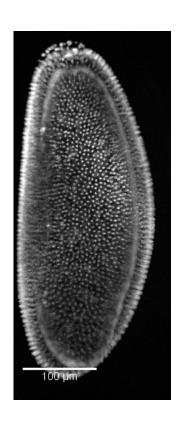
Preprocessing

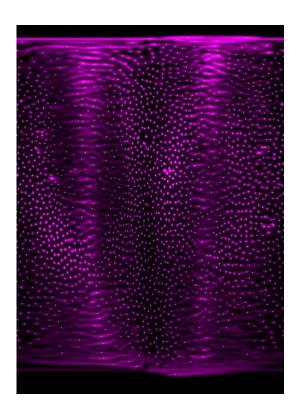
Transformation

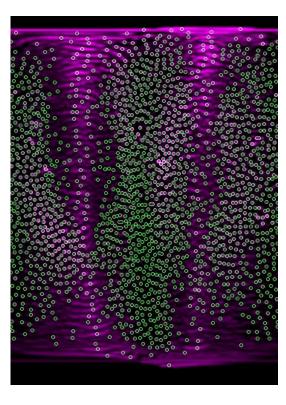
Segmentation

Save data









#### Why Notebooks?



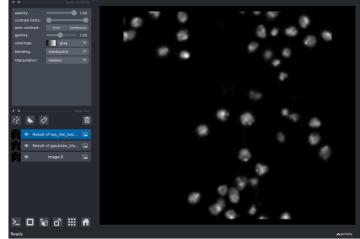
- A great way to document your analysis workflow
  - Understandable
  - Reproducible
  - Publishable
  - Reusable don't forget to add a license: https://choosealicense.com/
- Helps automate your workflow (some programming required)
  - Saves time
  - Improves reproducibility and repeatability
  - Reduces bias (but you still need to be careful when you set up the workflow)

#### top hat box

Background was removed using a top-hat algorithm with a symmetric, box-shaped footprint of 10 by 10 pixels.

This algorithm treats objects that are significantly larger than the footprint as background and smaller objects as signal.





#### threshold otsu binarization

The image was binarized using a threshold determined by using Otsu's method [1]



## Fiji/clij: Set up a workflow



Make x, y, z – projections match biological axes



Special thanks to Elisabeth Kugler!



Elisabeth Kugle @KuglerElisabeth

□ GeFor... - □ ×
200x100 pixels; RGB; 78K
GeForce RTX 2080 TI
0.0 / 11.0 GB

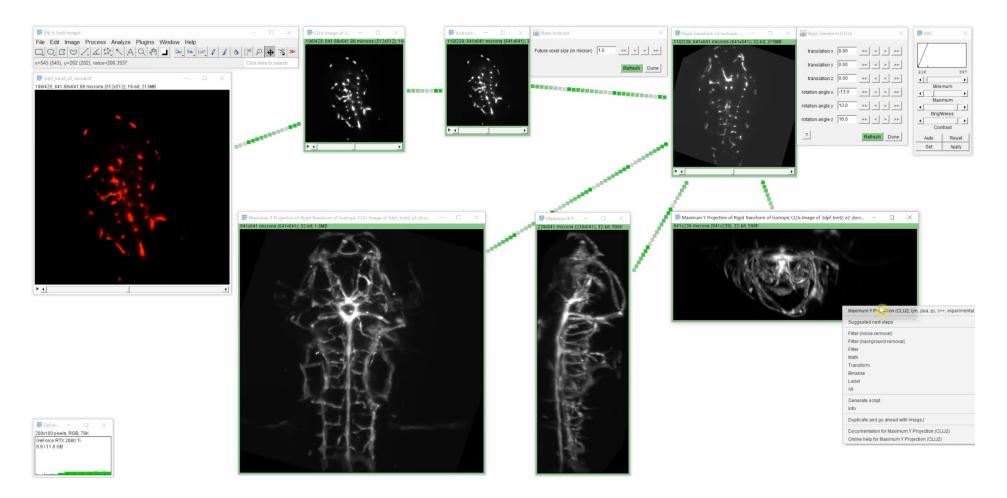
Image data source: Elisabeth Kugler; labs of Tim Chico and Paul Armitage, The University of Sheffield (UK)" <a href="https://zenodo.org/record/4204839#.X8DCRGj7Q2w">https://zenodo.org/record/4204839#.X8DCRGj7Q2w</a>



#### Fiji/clij: Generate code



After setting up the workflow, generate code!



Special thanks to Elisabeth Kugler!



Elisabeth Kugle @KuglerElisabeth

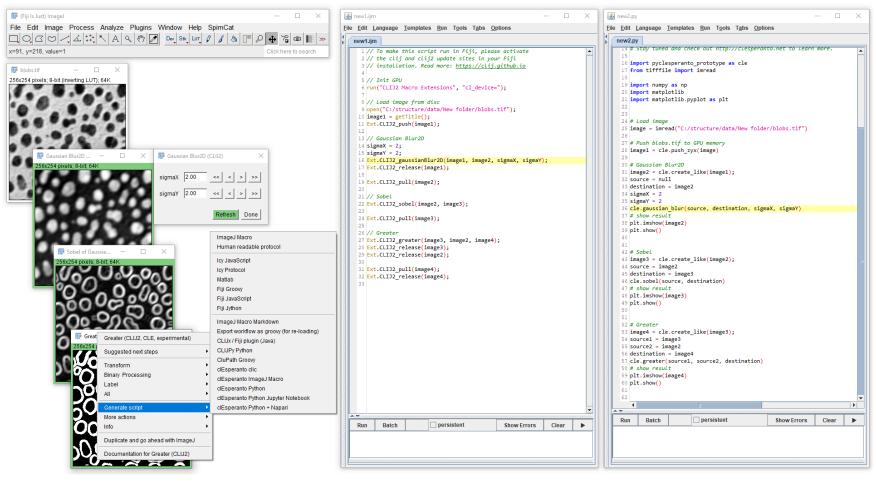
Image data source: Elisabeth Kugler; labs of Tim Chico and Paul Armitage, The University of Sheffield (UK)" https://zenodo.org/record/4204839#.X8DCRGj7Q2w



#### Choosing a programming language



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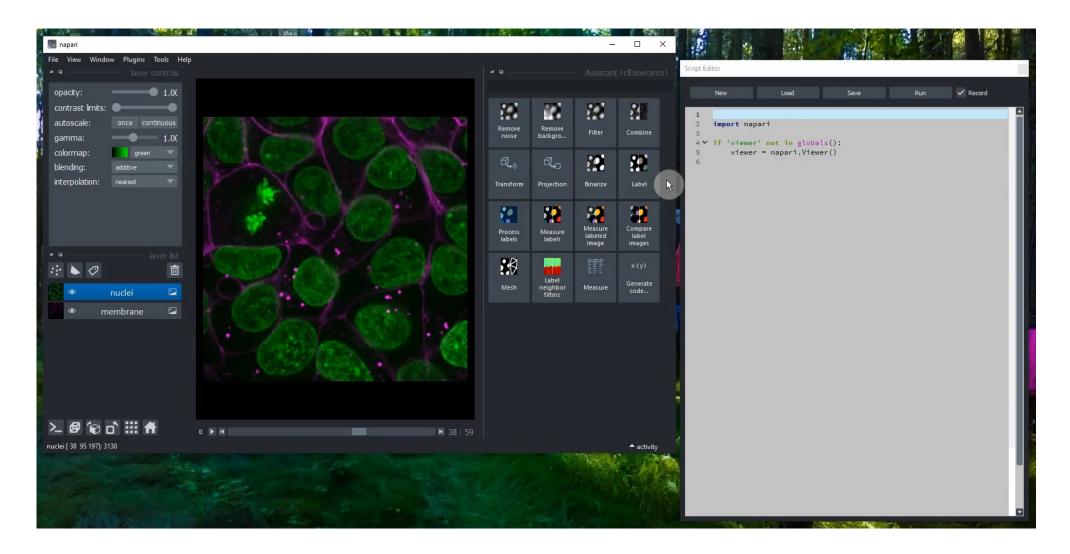


- Do you have access to existing scripts from others (and does the license allow you to use them)?
- Do you need other tools/packages (e.g. a deep learning python package)?
- Personal preference



#### Create a workflow with Napari-Assistant

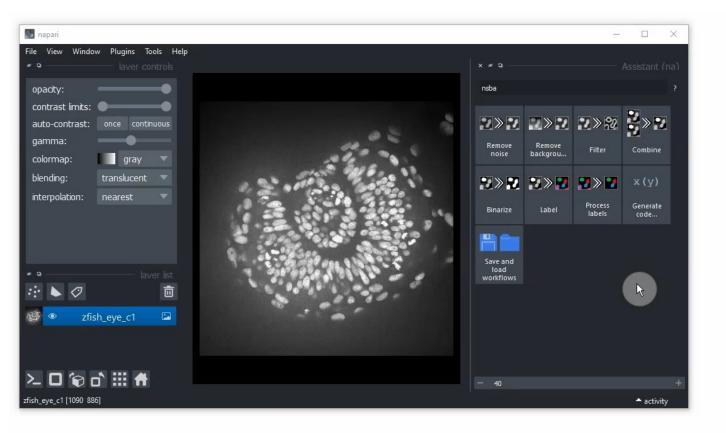




#### Export code to Jupyter Notebooks



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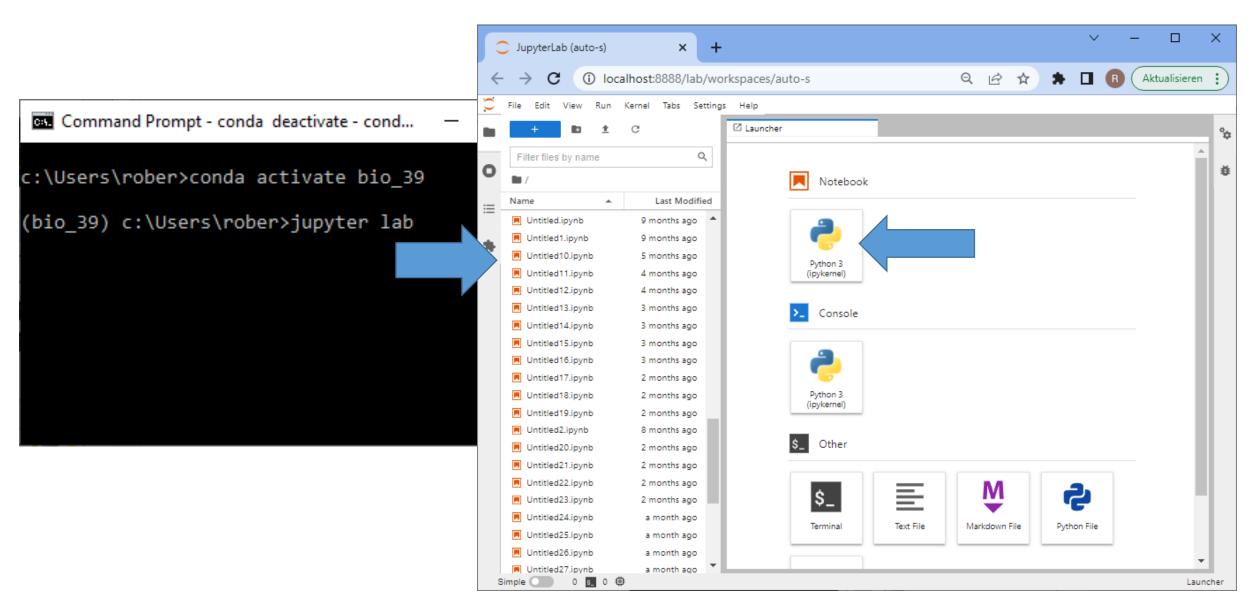
https://github.com/haesleinhuepf/napari-assistant

Image data source: Mauricio Rocha Martins, Norden lab, MPI CBG (now at IGC Oeiras)



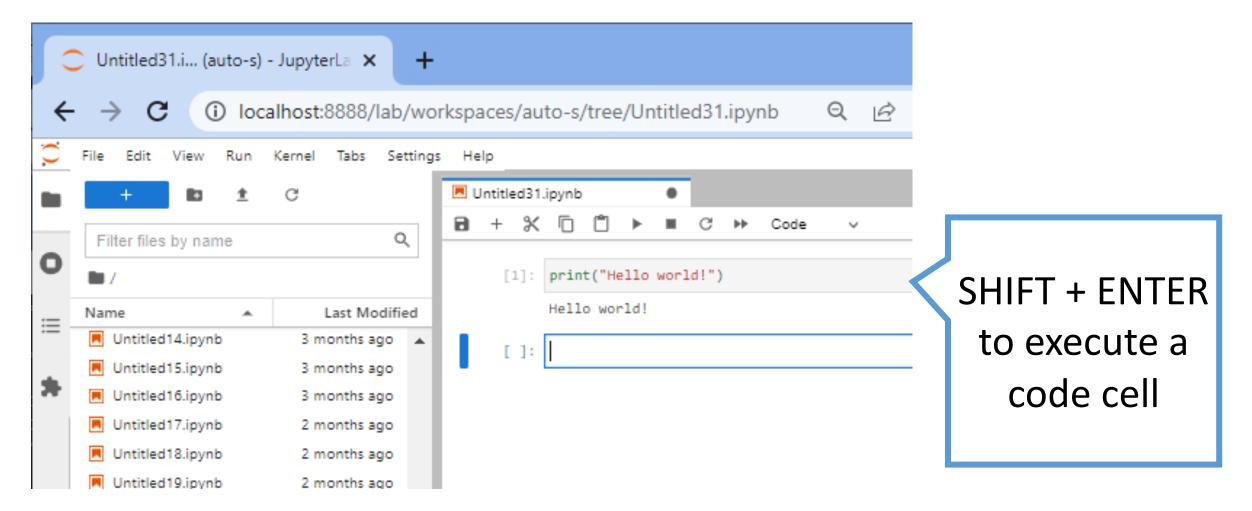
#### Open the notebook in Jupyter lab





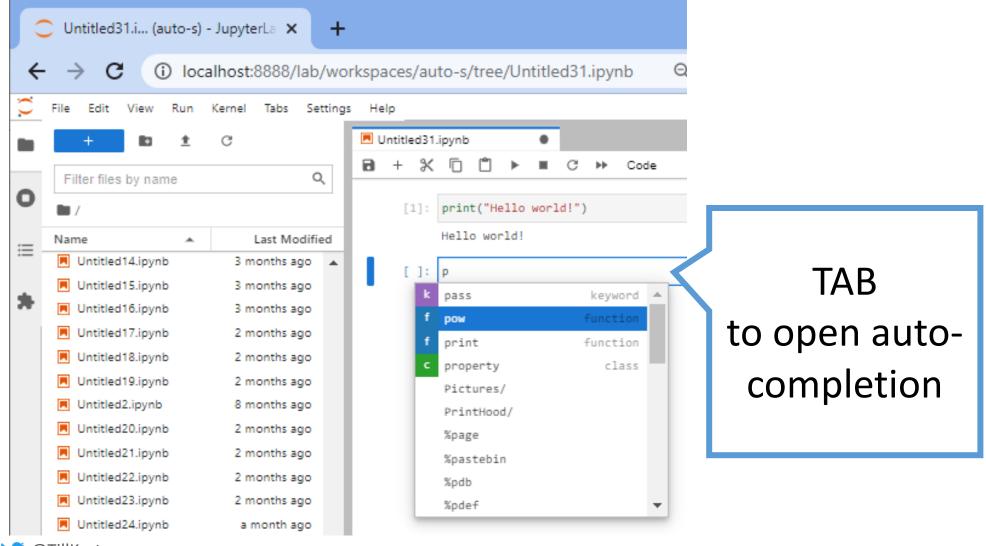


Execute code cell-by-cell and see results instantaneously





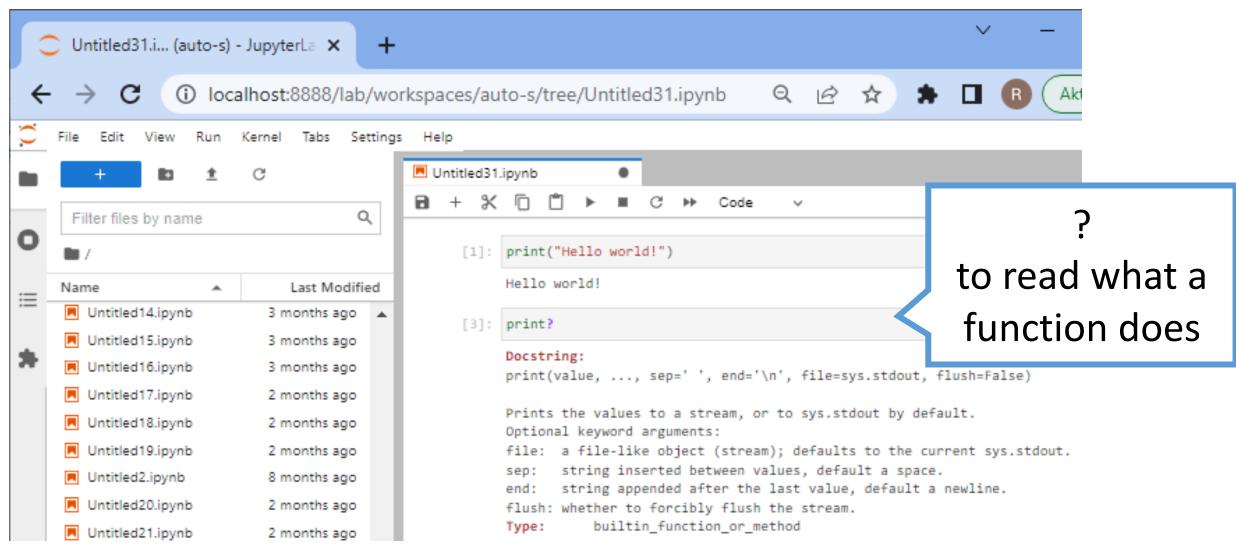
Context-specific help, auto-completion



TillKorten

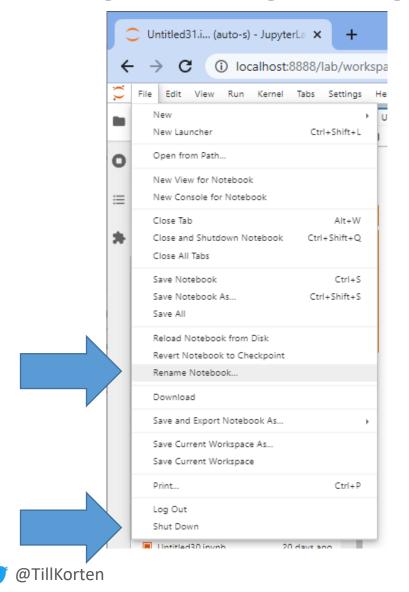


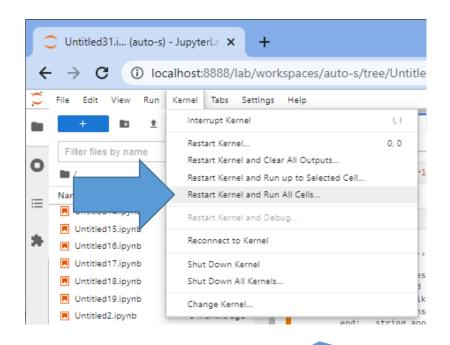
Help / "docstrings"





Saving / renaming / closing





Enforcing a "clean" execution state is important for ensuring reproducibility and repeatability



• Example/exercise notebooks online: <a href="https://t.ly/DVsyu">https://t.ly/DVsyu</a>

