



# Fiji workshop 2025

- Day 1:
  - Image data visualization in Fiji
- Day 2:
  - Quantitative image analysis in Fiji
- Day 3:
  - One-on-one analysis support with your own image data

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ABIF

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https://github.com/ABIF-McGill/ABIF-Fiji-Workshop-2025

# Day 1: Data visualization in Fiji

- Today's goals
  - Get acquainted with Fiji!
  - LUTs, contrast
  - Composite images
  - Intro to scripting!
  - Look at data from 3D images
  - Look at data from timelapse images
  - Images for publications

# Fiji is awesome! But it has some drawbacks...

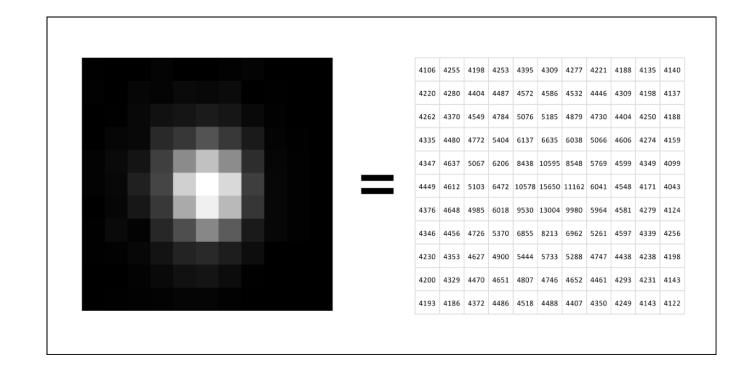


- Why should you use Fiji?
  - Free open-source software
  - Huge development community
  - Librairies to open nearly any microscope image (!)
  - Automation and scripting with multiple programming languages (!)

- When should you use something else than Fiji?
  - More powerful computation
  - Better 3D visualization
  - Many deep learning tools
  - Instrument-specific processing

### Digital images

- A digital image is simply a table of numbers
- Each number represents the intensity value of a pixel
- The intensity value is proportional to the amount of light collected at that location
- Software such as Fiji simply displays each number as a certain shade or colour



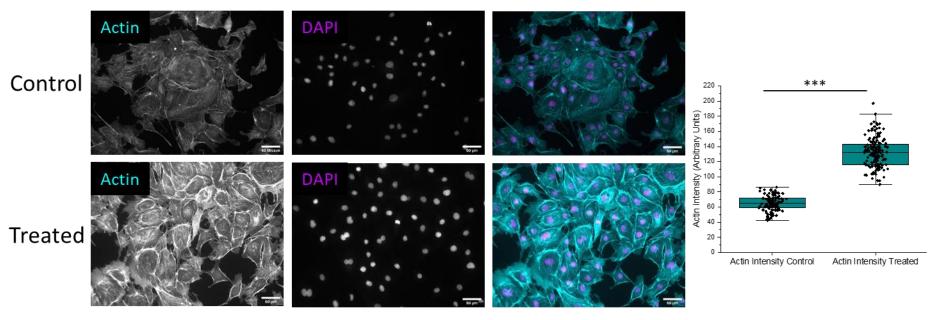
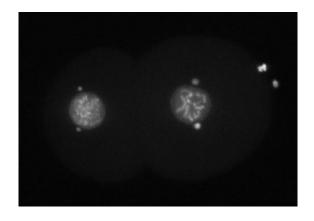
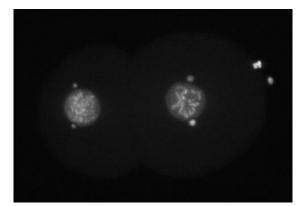
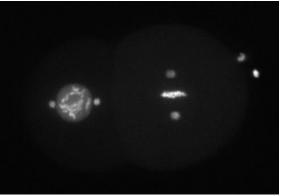


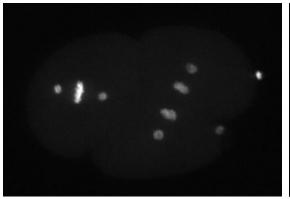
Figure 1: Images of CHO-K1 cells stained with Phalloidin Alexa Fluorophore 488 and stained with the nuclear probe DAPI. Images were flat field corrected using images of a fluorescent plastic slide imaged on the same microscope. Images were collected on an upright Zeiss Axioskop with a EC PlanNeoFluar 20x/0.5 NA objective lens with an AxioCam ICm1 camera. DAPI was imaged using a DAPI cube and an exposure time of 50 ms. Actin was imaged with a FITC cube and a 150 ms exposure. Brightness, gamma and contrast were adjusted to visualize the actin features well. The scale bar is 50  $\mu$ m. Box plot of actin intensity for control (n=110) and treated (n=149) cells. T-test was done with a two sample t-test, unequal variance, two tailed. \*\*\* corresponds to P<0.001.

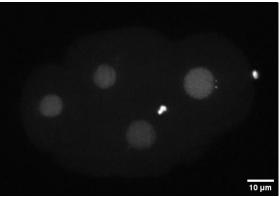


C. elegans embryo
eGFP-H2B
eGFP-gamma-tubulin
Max projection movie
One z-stack per minute acquisition
4 frames-per-second playback





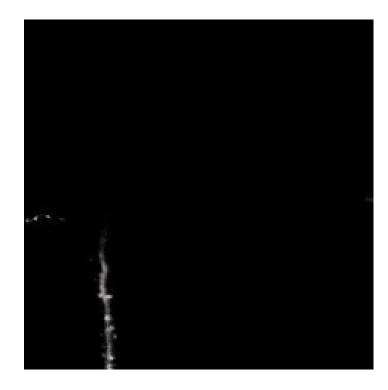




0 minutes 2 minutes

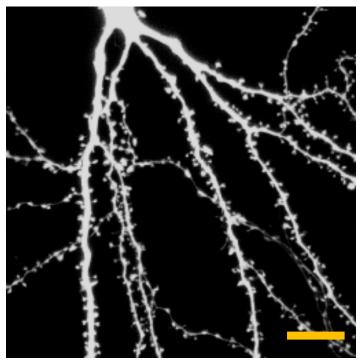
4 minutes

8 minutes



Imaris demo image

Pyramidal cell z-stack 69 z-slices 15 frames-per-second playback

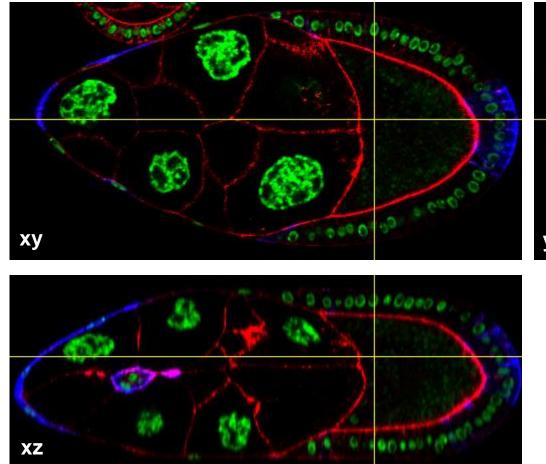


Maximum intensity projection (Fiji) Scalebar: 10 um



Imaris demo image

Drosophila egg Chamber
3 channels x 98 z-slices
16 frames-per-second playback



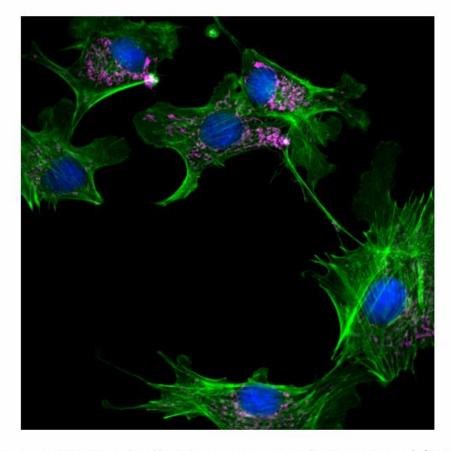
Orthogonal slice views

### Exercise 1 – raw vs composite RGB images



demo\_DAPI\_Phalloidin\_Mitotracker\_001.tif

Raw image, drag-and-drop in powerpoint.



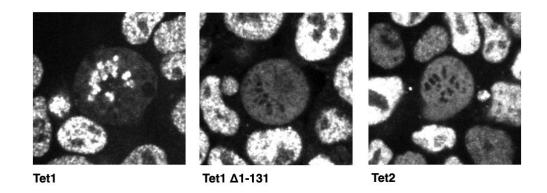
demo\_DAPI\_Phalloidin\_Mitotracker\_001.tif (RGB).tif

Composite RGB image generated with Exercise 1 macro, drag-and-drop in powerpoint.

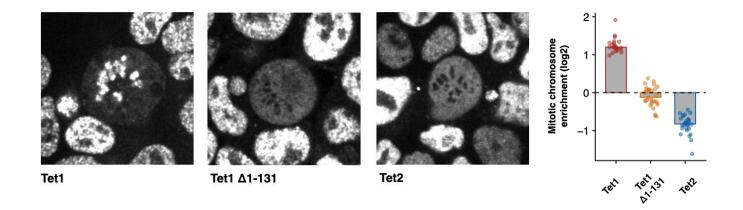
# Day 2: Quantitative imaging analysis in Fiji

#### • Goal:

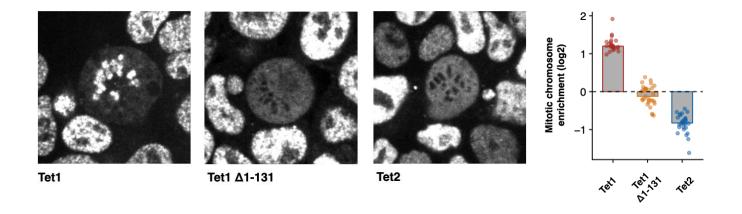
Why quantitative analysis?



Why quantitative analysis?



Why quantitative analysis?



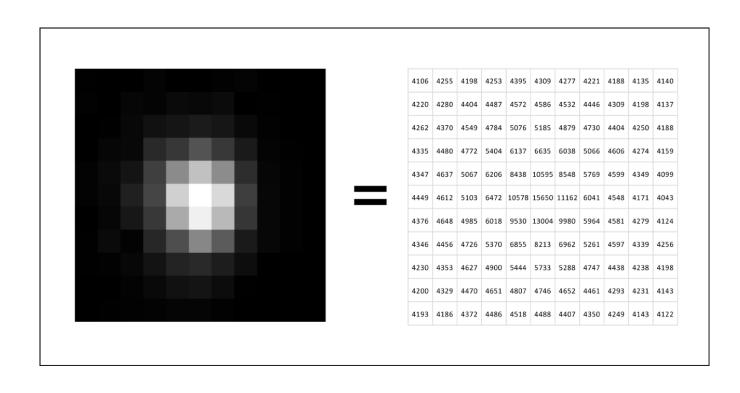
- Measured quantity (space, time, intensity etc)
- Increases robustness of observations
- Elucidate patterns that are not easily visible by eye



Better questions, better science!

#### Images can be quantitative:

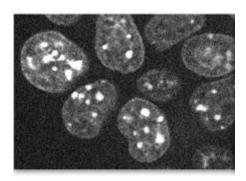
- A digital image is simply a table of numbers
- Each number represents the intensity value of a pixel
- The intensity value is proportional to the amount of light collected at that location



Value at each pixel = **Signal + Background + Noise** 

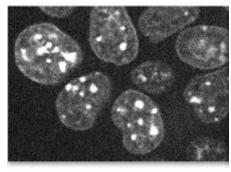
#### • Goal:

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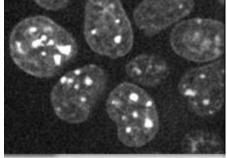


Open an image

#### • Goal:

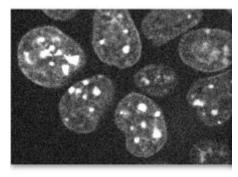


Open an image

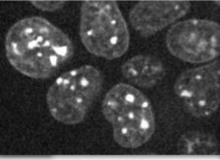


Processing such as filtering

#### • Goal:



Open an image

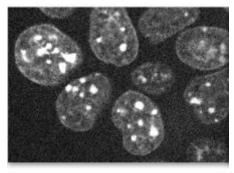


Processing such as filtering

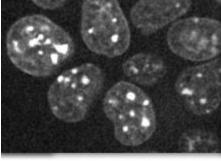


Segmentation or object detection

#### • Goal:



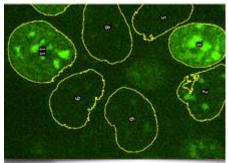
Open an image



Processing such as filtering



Segmentation or object detection



Data extraction from segmented objects

- -Number of objects
- -Shape / size
- -Intensity within object...

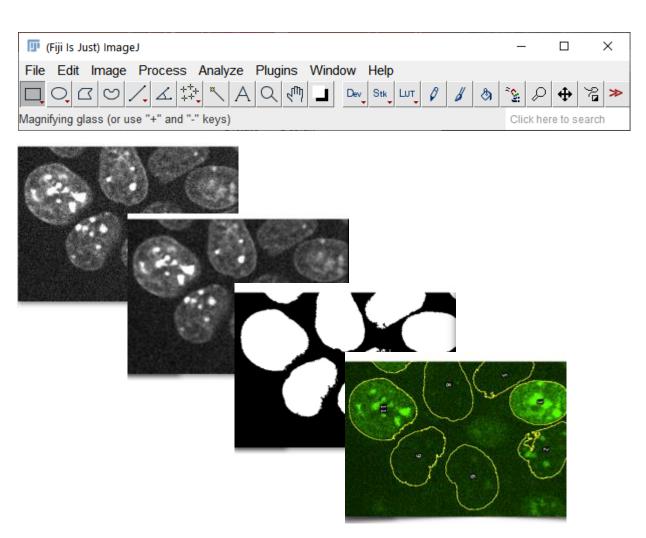
#### • Goal:



- Why Fiji?
  - Free open-source software
  - Huge development community
  - Librairies to open nearly any microscope image (!)
  - Automation and scripting with multiple programming languages (!)

#### • Today:

- Load images in Fiji, get familiarised with the program
- Processing using convolutional filters
- Segmentation with thresholding
- Data extraction
  - ...plotting?



```
for (i = 0; i < numROI; i++) {
    roiManager("Select", i);
    Stack.setChannel(2);
    run("Measure");
}</pre>
```

```
for (i = 0; i < numROI; i++) {</pre>
     roiManager("Select", i);
     Stack.setChannel(2);
     run("Measure");
```

```
for (i = 0; i < numROI; i++) {</pre>
     roiManager("Select", i);
     Stack.setChannel(2);
     run("Measure");
```

first, "build a sequence of integers going from the Start Value, to the End value, with an increment of 1"

```
for (i = 0; i < numROI; i++) {</pre>
     roiManager("Select", i);
     Stack.setChannel(2);
     run("Measure");
```

```
first, "build a sequence of integers going from the Start Value, to the End value, with an increment of 1" --- in the case of numROI being 5: 0, 1, 2, 3, 4
```

```
for (i = 0; i < numROI; i++) {</pre>
     roiManager("Select", i);
     Stack.setChannel(2);
     run("Measure");
```

first, "build a sequence of integers going from the Start Value, to the End value, with an increment of 1" ----in the case of numROI being 5: 0, 1, 2, 3, 4,

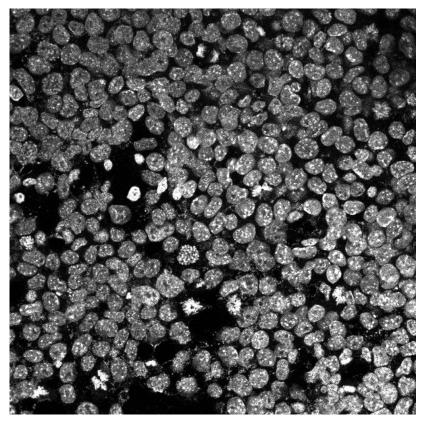
"Run the { content of the loop } replacing i with the first integer of the sequence.

Then run the { content of the loop } replacing i with the second integer in the sequence... " and so on...

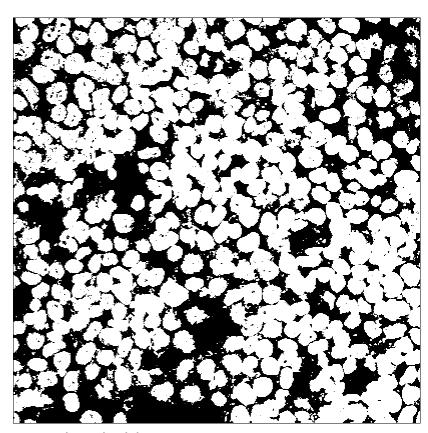
```
for (i = 0; i < numROI; i++) {
      roiManager("Select", i);
      Stack.setChannel(2);
      run("Measure");
```

```
i = 0
roiManager("Select", i);
Stack.setChannel(2);
run("Measure");
i = 1
roiManager("Select", i);
Stack.setChannel(2);
run("Measure");
i = 2
roiManager("Select", i);
Stack.setChannel(2);
run("Measure");
i = 3
roiManager("Select", i);
Stack.setChannel(2);
run("Measure");
. . .
```

• Sometimes objects are too close for adequate thresholding

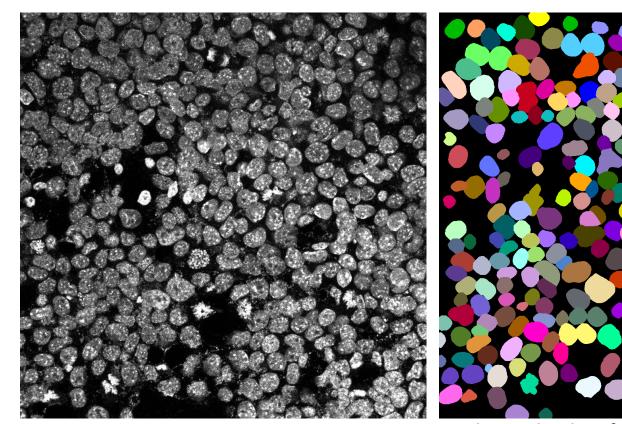


DAPI image



Otsu threshold image

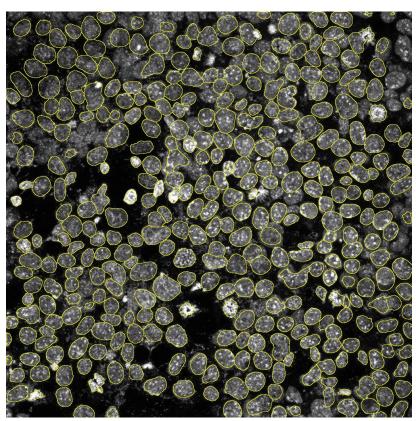
• Machine learning tools trained on cell images can help distinguish close objects



DAPI image

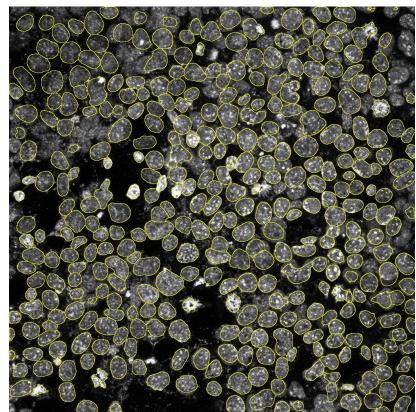


Nuclei masks identified with *Cellpose* 

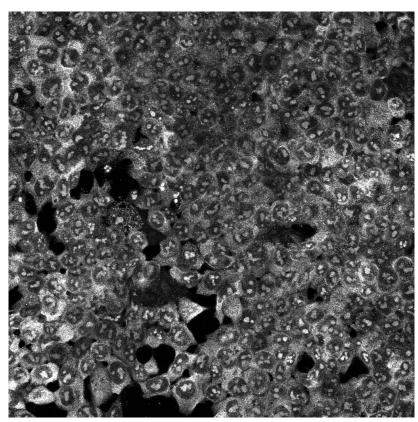


DAPI image + mask outlines

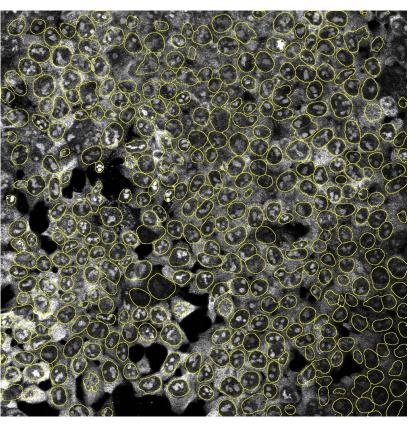
• Using the DAPI masks, we can then quantify intensities of other stainings in the nucleus



DAPI image + nucleus mask outlines

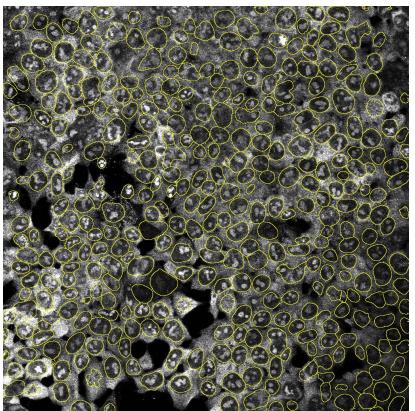


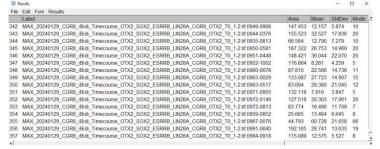
LIN28-GFP image

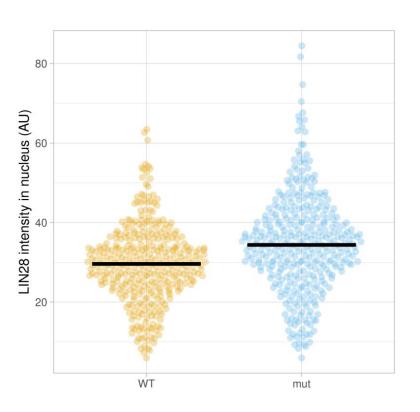


LIN28-GFP image + nucleus mask outlines

• Using the DAPI masks, we can then quantify intensities of other stainings in the nucleus







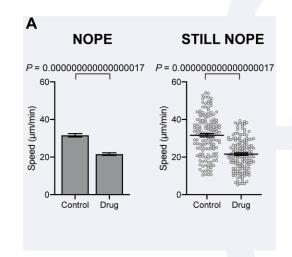
LIN28-GFP image + nucleus mask outlines

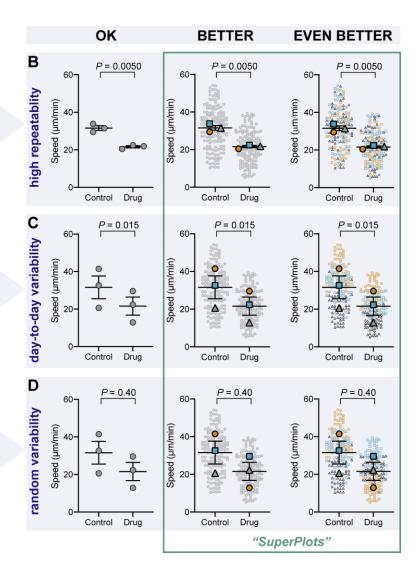
# Plotting resources

# SuperPlots: Communicating reproducibility and variability in cell biology

Samuel J. Lord<sup>1</sup>, Katrina B. Velle<sup>2</sup>, R. Dyche Mullins<sup>1</sup>, and Lillian K. Fritz-Laylin<sup>2</sup>

- SuperPlots provides guidelines for presenting experimental data.
- Best to show all data points and colour code them per experimental replicate.
- For statistics the *n* value should be the number of experiments.





### Plotting resources

#### PlotsOfData - Plots all Of the Data

