



Introduction to Bioimage Analysis with Fiji/ImageJ

Pasteur NEUBIAS Course on Bioimage Analysis

Early Career Investigators Track

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Who has done image analysis
before?

Who has used Fiji/ImageJ before?

What are your backgrounds?

Plan for today

I/ Bioimage Analysis (BIA)

II/ Fiji

III/ Fundamentals

IV/ Image ethics

V/ Quantification

Hands-on 1: Count nuclei and determine which ones are green

Hands-on 2: Quantify the number of red spots per nucleus

VI/ Intro to automation

Plan for today

I/ Bioimage Analysis (BIA)

What do you mean, you want to « quantify » things in images?

Piloting experiments all the way

The typical BIA workflow

Software: the usual suspects

II/ Fiji

Flavors of ImageJ

Installing and updating

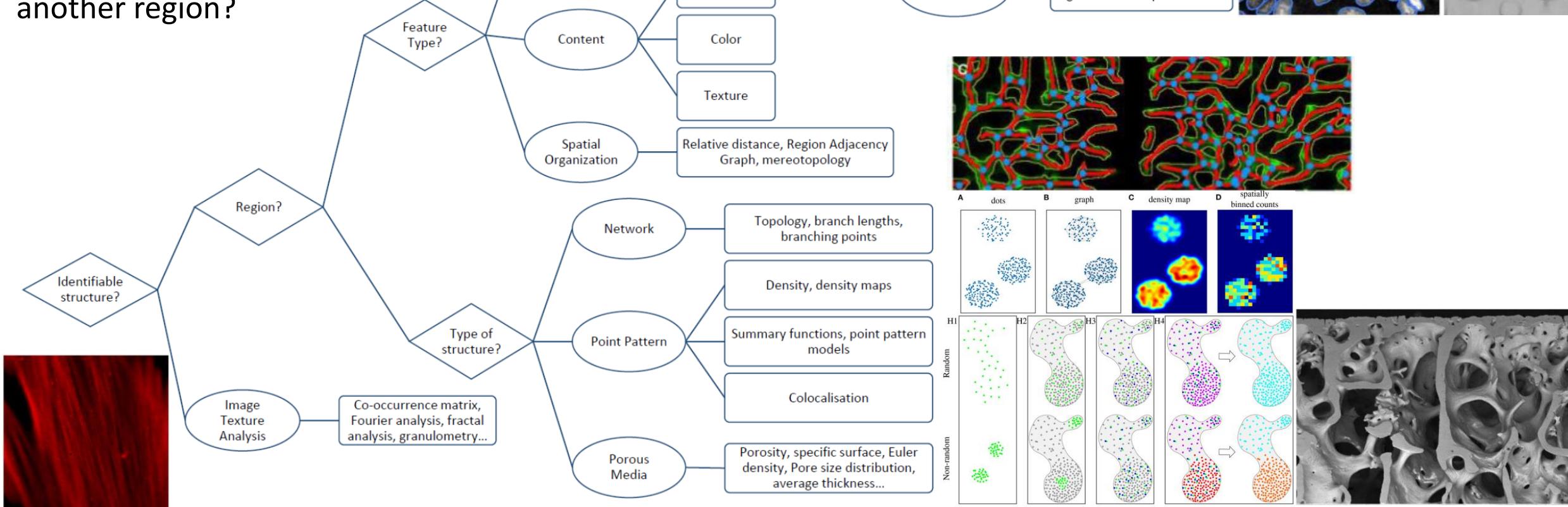
Plugins

Opening images

I/ Biolimage Analysis (BIA): What do you mean when you say you want to quantify things in a biological image?

I/ BIA: Typical things you'd want to measure in biological images

Object counts? Density per region? Shape comparison? Surface %? Intensity? Object class? Diameter? Distance to another region?



Fazeli, Elnaz, Robert Haase, Michael Doube, Kota Miura, et David Legland. « From Cells to Pixels: Bridging Biologists and Image Analysts Through a Common Language », 2024.

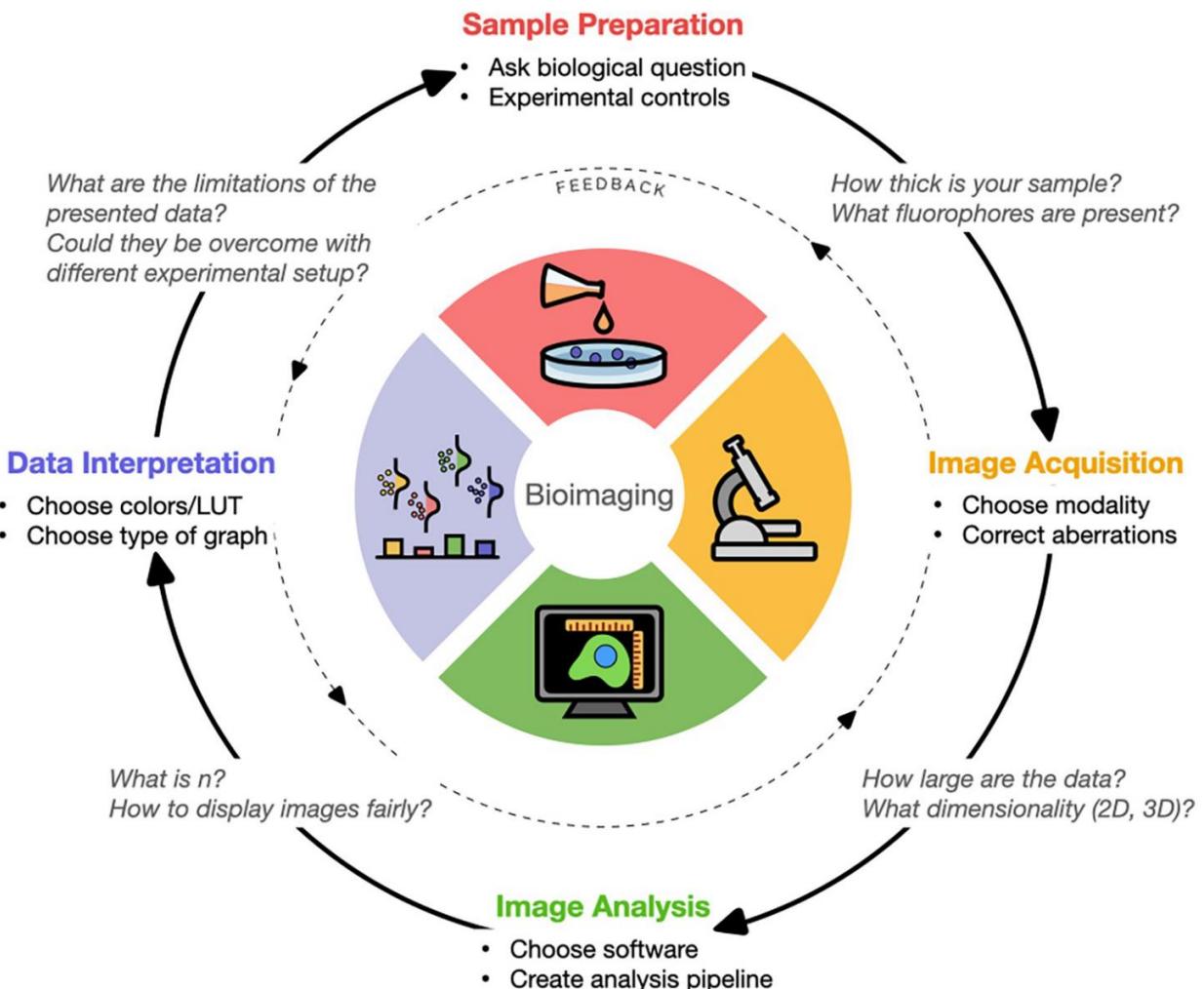
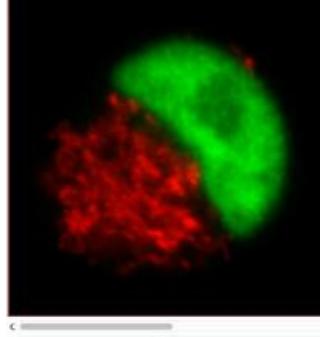
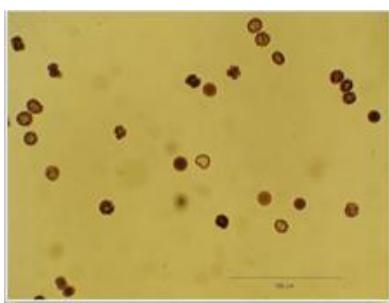
I/ BIA: Pilot your experiments all the way!

It is a great idea to pilot your experiment all the way to the analysis and visualization!

“Do I need to segment surfaces? Individual objects?”

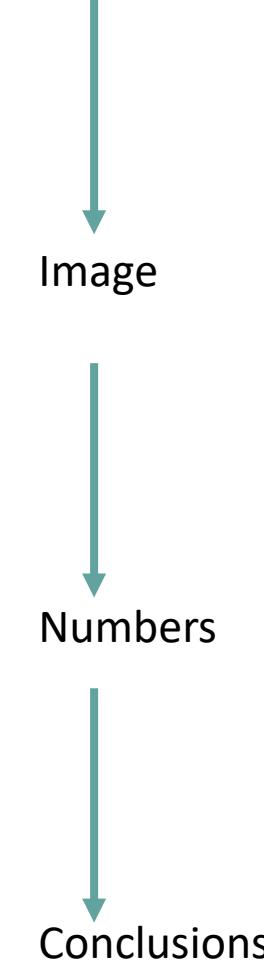
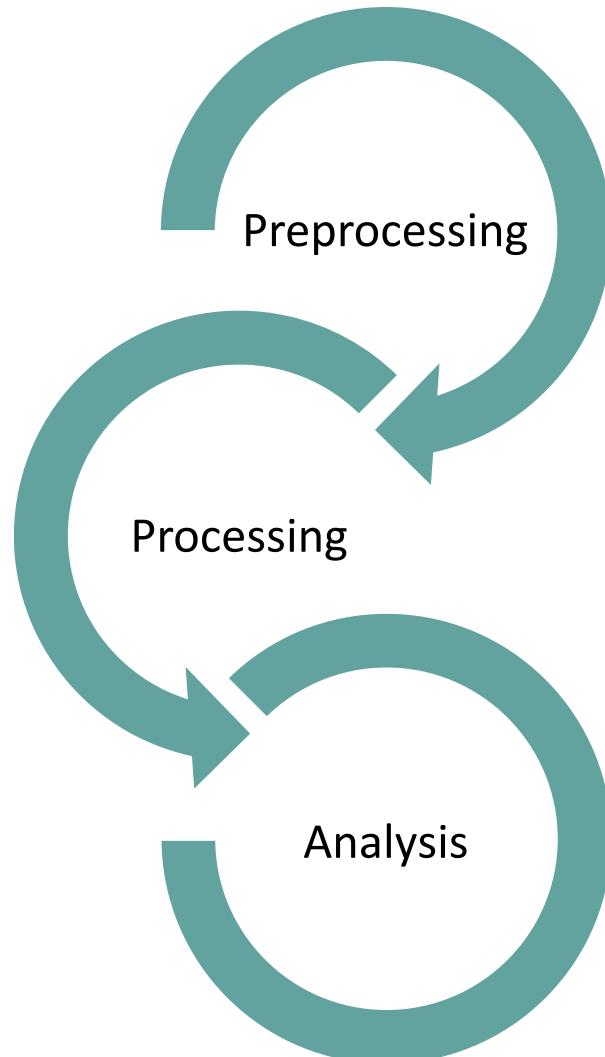
“What type of control do I need?”

“What type of microscopy am I going to use? Is there a marker that could help?”



Senft, Rebecca A., Barbara Diaz-Rohrer, Pina Colarusso, Lucy Swift, Nasim Jamali, Helena Jambor, Thomas Pengo, et al. « A Biologist's Guide to Planning and Performing Quantitative Bioimaging Experiments ». *PLOS Biology* 21, n° 6 (27 juin 2023): e3002167. <https://doi.org/10.1371/journal.pbio.3002167>.

I/ BIA: Typical bioimage analysis workflow



From pixels to numbers
to statistics to biological
conclusions:
“information
condensation”

I/ BIA: Software: the usual suspects

Open source



Fiji/ImageJ

This workshop



Icy



QuPath



Ilastik



Napari

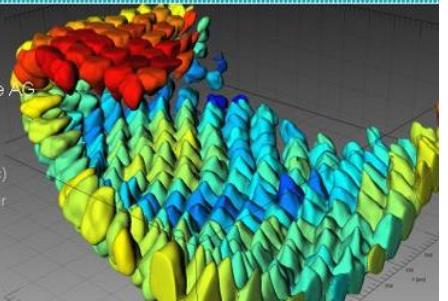


Commercial

IMARIS®

Imaris
7.2.1 [Dec 22 2010]
Build 22330 for Win32
Copyright 1993-2010 Bitplane AG
All rights reserved

European Patent EP20050009577
United States Patent 7382374
Using XML parser expat. Copyright (c)
1996, 1999, 2000, Trai Open Source
Software Center Ltd and Clark Cooper
This software uses libraries from the
FFmpeg project under the LGPLv2.1



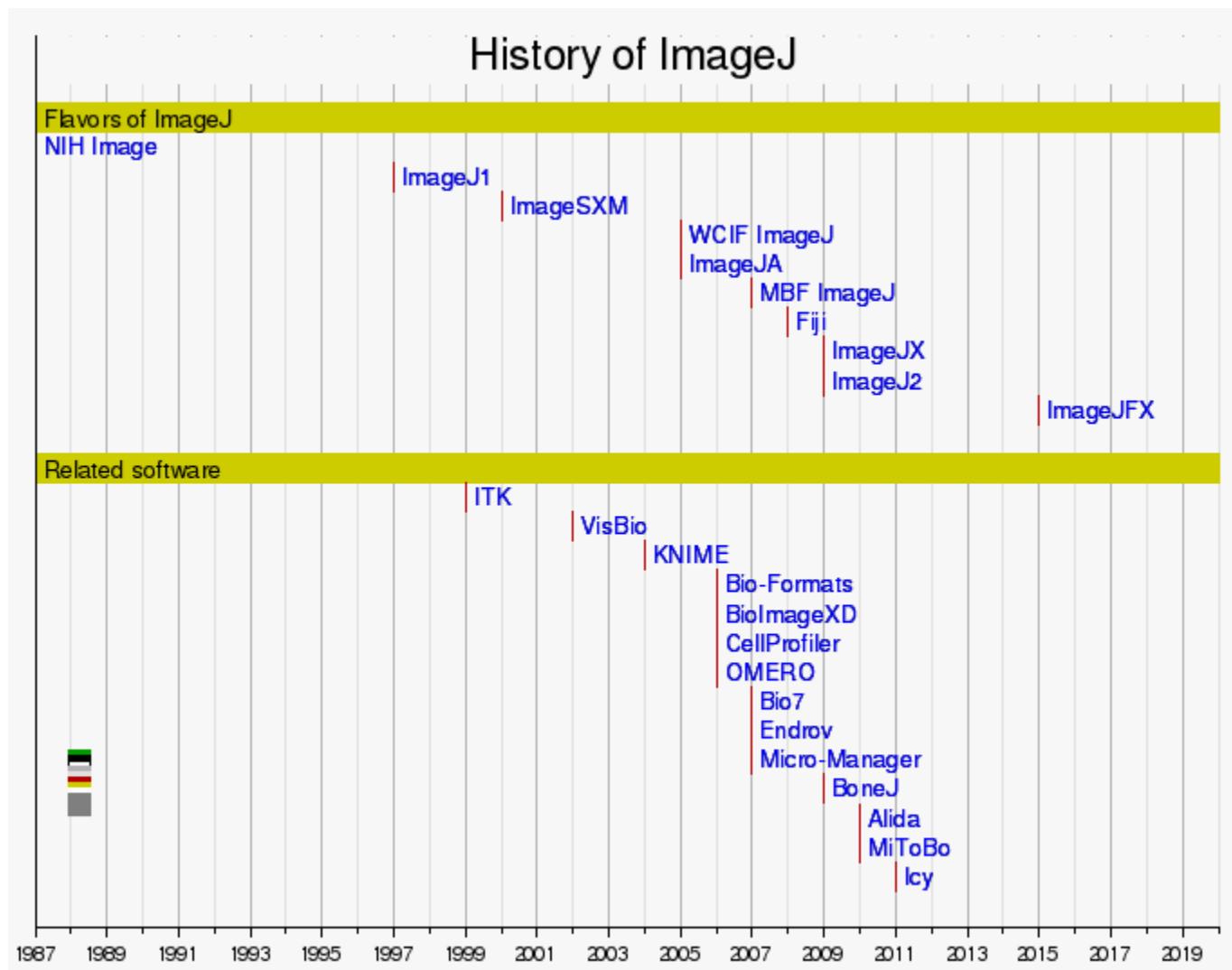
II/ Fiji: installing and updating, installing plugins

II/ Fiji: Flavors of ImageJ

There are many different derivatives of ImageJ with very similar names, and some confusion is inevitable. Below is a table which should help to clarify the purpose of each. For the historical context of these projects, see [History](#) below.

Name	Author/Maintainer(s)	Description	Initiated	Status
 ImageJ	Wayne Rasband	The original version of ImageJ which has been in development since 1997. It has a strong, established user base, with thousands of plugins and macros for performing a wide variety of tasks. Sometimes referred to as the original ImageJ or ImageJ 1.x for technical clarity, or informally with the shorthands ImageJ1 or IJ1 .	1997	Active
 ImageJ2	ImageJ2 developers	A new version of ImageJ targeting scientific multidimensional image data. It is a complete rewrite of the original ImageJ , but includes the original ImageJ with a compatibility layer, so that old-style plugins and macros can run the same as always. ImageJ2 provides several significant new features, such as an automatic updater , and improved scripting capabilities.	Dec. 2009	Active
 Fiji	Fiji contributors	Fiji is Just ImageJ, with extras. It is a distribution of ImageJ and ImageJ2 with many plugins useful for scientific image analysis in fields such as life sciences. It is actively maintained, with updates released often.	Dec. 2007	Active
 ImageJ.js	Wei Ouyang	ImageJ.js is a web version of ImageJ that runs in the browser without installation, compiled from Java to JavaScript using the Cheerpj compiler and integrated with the ImJoy plugin system. It's accessible from https://ij.imjoy.io and also supports mobile devices and tablets.	2020	Active

II/ Fiji: Flavors of ImageJ



II/ Fiji: Installation

<https://fiji.sc/>



Fiji

Fiji is an image processing package — a "batteries-included" distribution of [ImageJ](#), bundling many plugins which facilitate scientific image analysis.

[Download for Windows \(64-bit\)](#)

[More Downloads](#)

[Cite](#)

[Contribute](#)

Why Fiji?



Easy to Use

Fiji is easy to use and install - in one-click, Fiji installs all of its plugins, features an automatic updater, and offers comprehensive documentation.



Powerful

Fiji bundles together many popular and useful ImageJ plugins for image analysis into one installation, and automatically manages their dependencies and updating.



Free & Open Source

Like ImageJ itself, Fiji is an [open source](#) project hosted on [GitHub](#), developed and written by the community.

<https://imagej.net/software/fiji/downloads>



Fiji Downloads

Fiji is a distribution of ImageJ which includes many useful plugins [contributed by the community](#).

Operating system	Download links
Linux (64-bit)	imagej.net (USA) micron.ox.ac.uk (Europe)
macOS (Apple silicon)	imagej.net (USA)
macOS (Intel)	imagej.net (USA) micron.ox.ac.uk (Europe)
Windows (64-bit)	imagej.net (USA) micron.ox.ac.uk (Europe)
Windows (32-bit)	imagej.net (USA) micron.ox.ac.uk (Europe)
No JRE	imagej.net (USA) micron.ox.ac.uk (Europe)

- Download for your OS
- Unzip (/!\ Somewhere you have full permissions!)

<https://imagej.net/contribute/citing>

II/ Fiji: Installation

Explore your Fiji directory

To launch Fiji,
double click on the .exe
You can also create a
shortcut on your desktop.

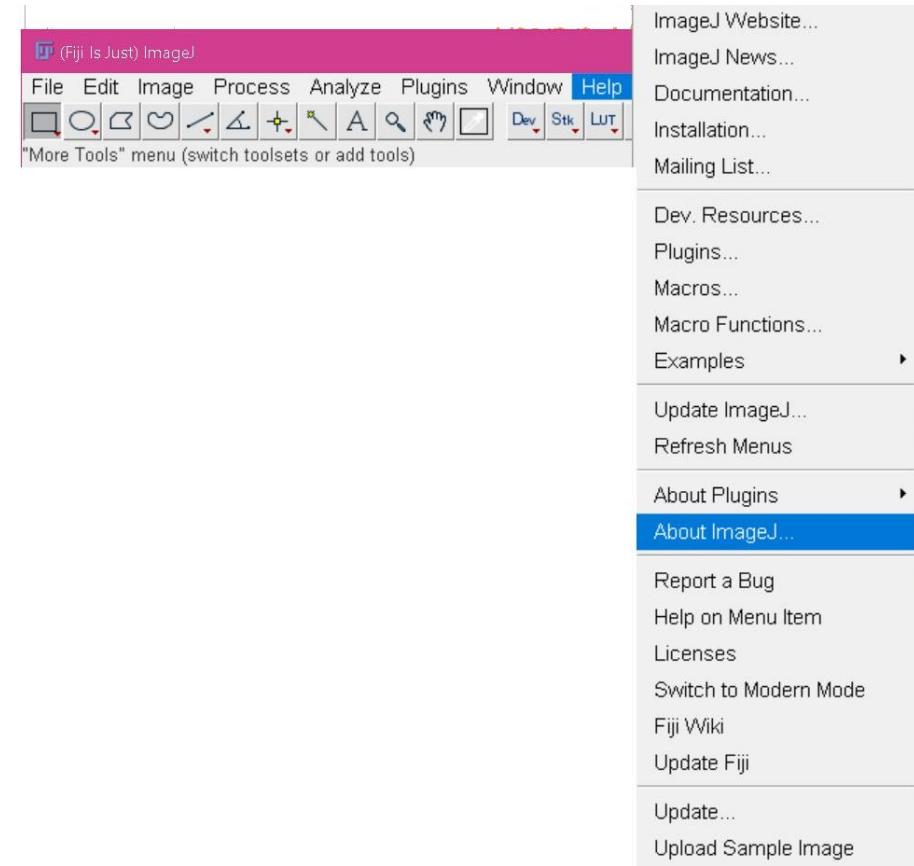
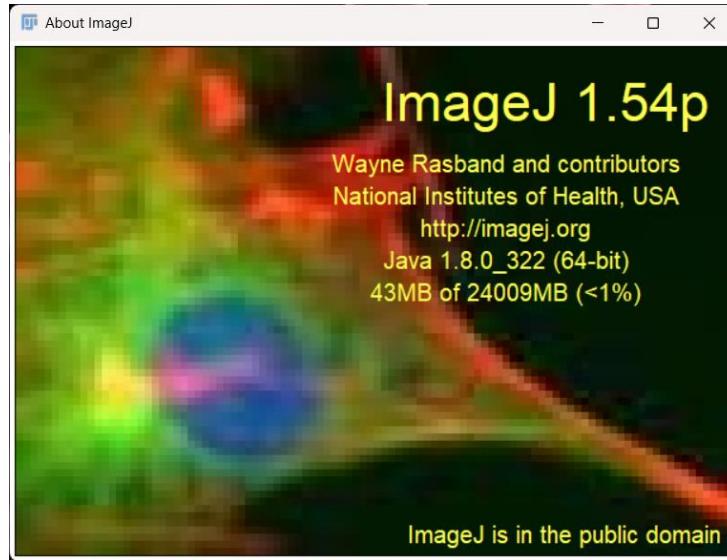
Nom	Type	Taille compres
config	Dossier de fichiers	
Contents	Dossier de fichiers	
Fiji.app	Dossier de fichiers	
images	Dossier de fichiers	
jars	Dossier de fichiers	
java	Dossier de fichiers	
lib	Dossier de fichiers	
licenses	Dossier de fichiers	
luts	Dossier de fichiers	
macros	Dossier de fichiers	
plugins	Dossier de fichiers	
retro	Dossier de fichiers	
scripts	Dossier de fichiers	
db.xml.gz	7-zip.gz	
fiji-windows-x64.exe	Application	
ImageJ-win64.exe	Application	
README.md	Fichier MD	
WELCOME.md	Fichier MD	

Fiji.app (18 éléments)

● Sélectionnez un seul fichier pour obtenir plus d'informations et partager votre contenu cloud.

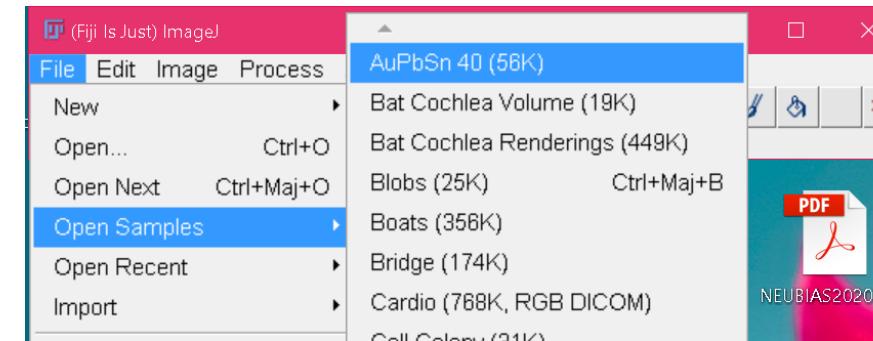
II/ Fiji: Installation

Check the version: *Help > About ImageJ*



II/ Fiji: Installation

Try to do *File > Open Samples > AuPnSn 40 (56k)*



Do you get



?

II/ Fiji: Installation: Plugins

Extending Fiji: plugins: <https://imagej.net/list-of-extensions>,
<https://imagej.net/list-of-update-sites/>

▼ Categories filter

All Any

<input checked="" type="checkbox"/> 3D	<input checked="" type="checkbox"/> Deep Learning	<input checked="" type="checkbox"/> ImgLib	<input checked="" type="checkbox"/> Noise	<input checked="" type="checkbox"/> Stacks
<input checked="" type="checkbox"/> Analysis	<input checked="" type="checkbox"/> Denoising	<input checked="" type="checkbox"/> Import-Export	<input checked="" type="checkbox"/> OME	<input checked="" type="checkbox"/> Stitching
<input checked="" type="checkbox"/> Annotation	<input checked="" type="checkbox"/> Development	<input checked="" type="checkbox"/> Integral Image	<input checked="" type="checkbox"/> Object Detection	<input checked="" type="checkbox"/> Super-resolution
<input checked="" type="checkbox"/> Automation	<input checked="" type="checkbox"/> Digital Volume Flattening	<input checked="" type="checkbox"/> Interactive	<input checked="" type="checkbox"/> Ops	<input checked="" type="checkbox"/> Time Signal Analysis
<input checked="" type="checkbox"/> Benchmark	<input checked="" type="checkbox"/> Digital Volume Unrolling	<input checked="" type="checkbox"/> Kymograph	<input checked="" type="checkbox"/> Optic Flow	<input checked="" type="checkbox"/> Tissue
<input checked="" type="checkbox"/> Binary	<input checked="" type="checkbox"/> Entropy	<input checked="" type="checkbox"/> Lifetime	<input checked="" type="checkbox"/> Particle Analysis	<input checked="" type="checkbox"/> Track analysis
<input checked="" type="checkbox"/> CellProfiler	<input checked="" type="checkbox"/> Example Data	<input checked="" type="checkbox"/> MATLAB	<input checked="" type="checkbox"/> Pattern Recognition	<input checked="" type="checkbox"/> Tracking
<input checked="" type="checkbox"/> Channel alignment	<input checked="" type="checkbox"/> Fast Fourier Transformation	<input checked="" type="checkbox"/> MRI	<input checked="" type="checkbox"/> Perfusion	<input checked="" type="checkbox"/> TrakEM2
<input checked="" type="checkbox"/> Cilia	<input checked="" type="checkbox"/> Feature Extraction	<input checked="" type="checkbox"/> Machine Learning	<input checked="" type="checkbox"/> Photogrammetry	<input checked="" type="checkbox"/> Transform
<input checked="" type="checkbox"/> Classification	<input checked="" type="checkbox"/> Filtering	<input checked="" type="checkbox"/> Macro	<input checked="" type="checkbox"/> Photography	<input checked="" type="checkbox"/> Tutorials
<input checked="" type="checkbox"/> Colocalization	<input checked="" type="checkbox"/> Fractal Dimension	<input checked="" type="checkbox"/> MacroExtensions	<input checked="" type="checkbox"/> Plotting	<input checked="" type="checkbox"/> Uncategorized
<input checked="" type="checkbox"/> Color Processing	<input checked="" type="checkbox"/> Frequency Analysis	<input checked="" type="checkbox"/> Mathematical Morphology	<input checked="" type="checkbox"/> Projection	<input checked="" type="checkbox"/> Unmaintained
<input checked="" type="checkbox"/> Complexity	<input checked="" type="checkbox"/> Gut	<input checked="" type="checkbox"/> Microscopy	<input checked="" type="checkbox"/> Registration	<input checked="" type="checkbox"/> User Interface
<input checked="" type="checkbox"/> Cookbook	<input checked="" type="checkbox"/> Help	<input checked="" type="checkbox"/> Microtubules	<input checked="" type="checkbox"/> SciJava	<input checked="" type="checkbox"/> Utilities
<input checked="" type="checkbox"/> Curvature	<input checked="" type="checkbox"/> Image Annotation	<input checked="" type="checkbox"/> Montage	<input checked="" type="checkbox"/> Scripting	<input checked="" type="checkbox"/> Visualization
<input checked="" type="checkbox"/> Curve Tracing	<input checked="" type="checkbox"/> ImageJ2	<input checked="" type="checkbox"/> Neuroanatomy	<input checked="" type="checkbox"/> Segmentation	
<input checked="" type="checkbox"/> Deconvolution	<input checked="" type="checkbox"/> ImageScience	<input checked="" type="checkbox"/> Neuron	<input checked="" type="checkbox"/> Skeleton	

Import-Export Uncategorized Scripting Filtering

Import-Export, SciJava, OME Uncategorized Uncategorized Uncategorized

Import-Export Uncategorized Tracking, Tutorials Color Processing

II/ Fiji: Installation: Plugins: *Hands-on!*

Example of plugin installation: Stardist



Step 1

Open Fiji Updater on Fiji

Help > Update...

Step 2

Click on Manage update sites and check
the following sites

Update sites : CSBDeep, StarDist and TensorFlow

Tip: use the Search bar!

Step 3

Close Manage update sites and Click on Apply changes

Step 4

Restart Fiji/ImageJ

Sometimes a simple Help> Refresh can do the job

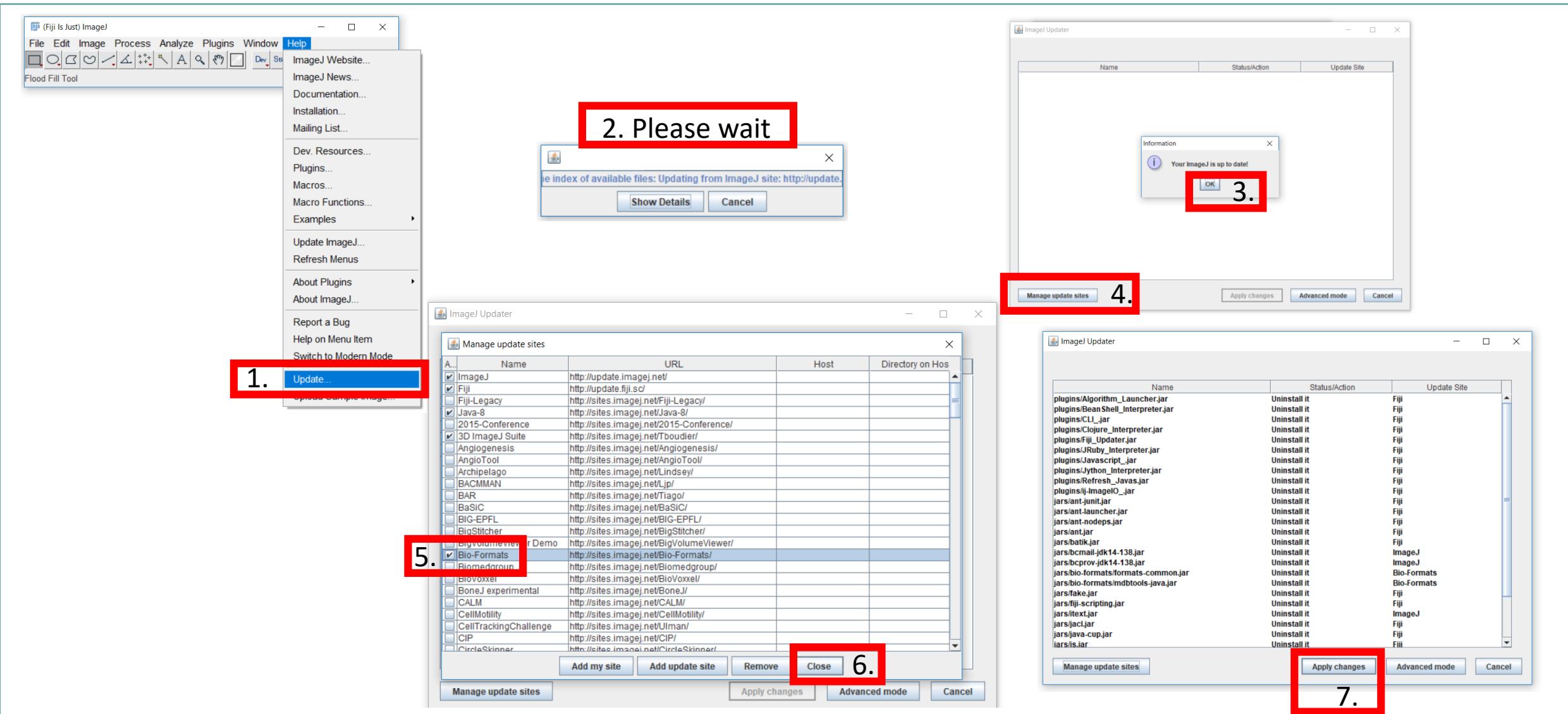
Step 5

The plugin is ready to use

It should appear under Plugins>StarDist

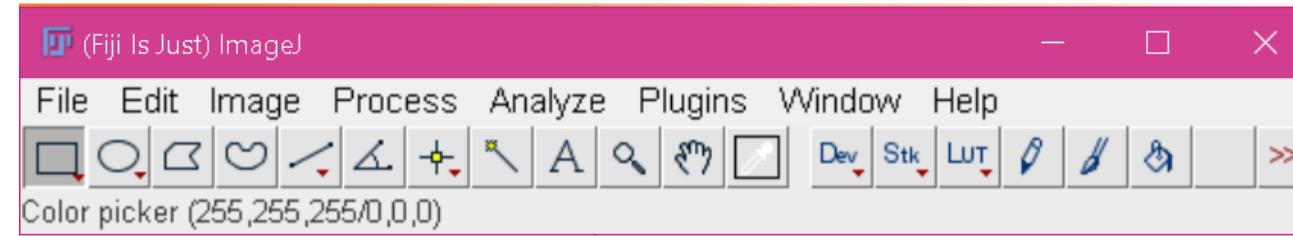
<https://imagej.net/plugins/stardist>

II/ Fiji: Installation: Plugins: Hands-on!



II/ Fiji: Exploring the menu

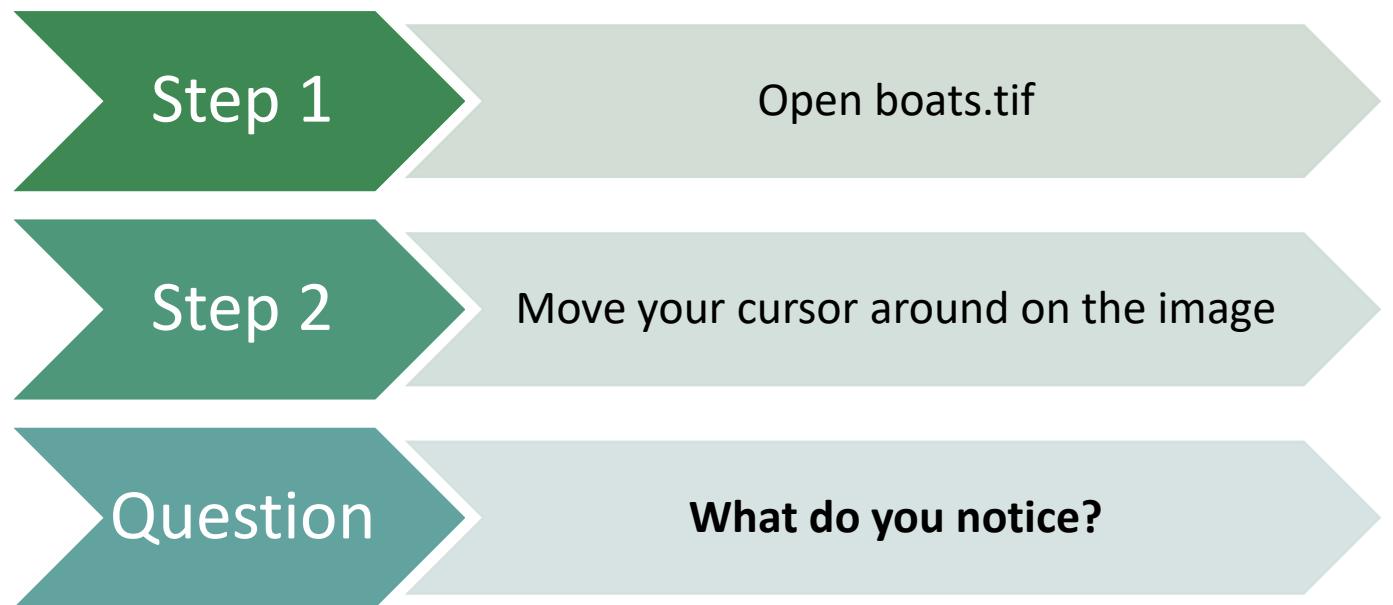
Going through the menu



II/ Fiji: Opening an image: *Hands-on!*

Two methods:

- File > open
- Drag and drop onto the Fiji bar



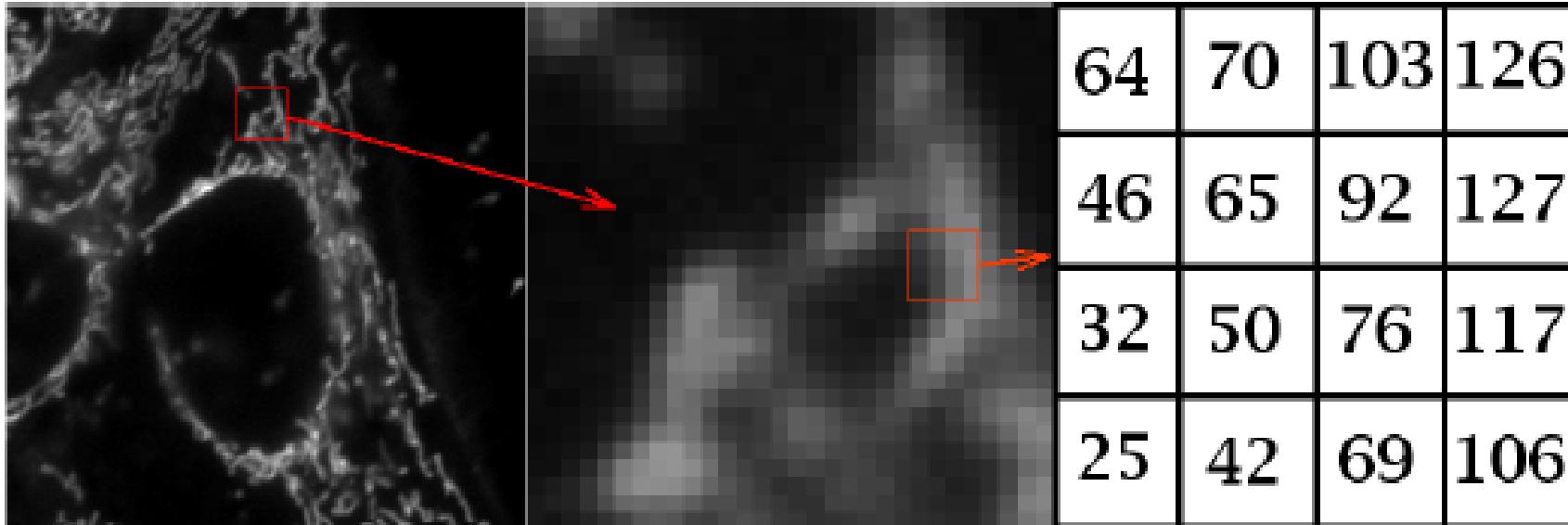
Rule number 1 in Fiji:

Anything you will do on an image will apply on the **active image**, i.e. the last image...

- which you have opened,
- or clicked on,
- or which was created by a process!

III/ Fundamentals: What are digital images?

III/ Fundamentals: What is a digital image?

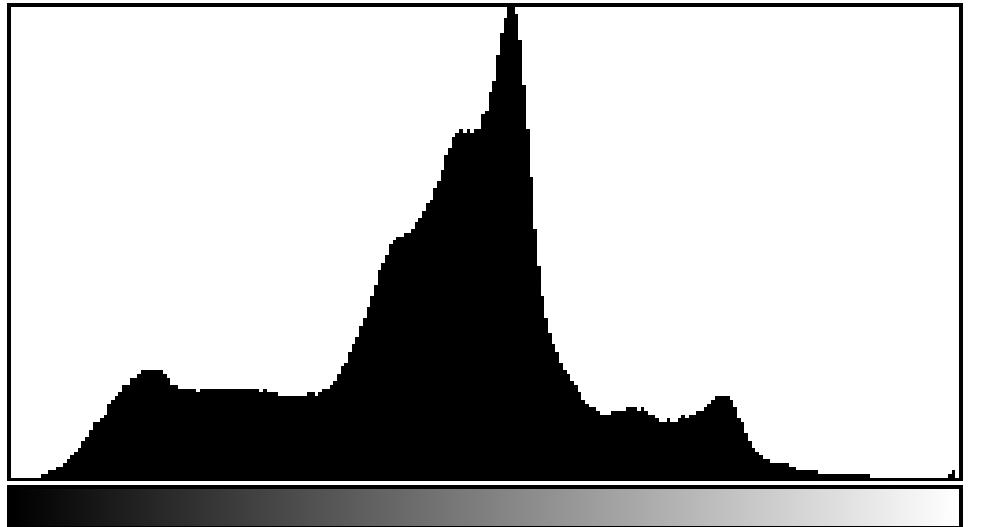


It is a matrix! There are only numbers, that we're going to name **intensity**

The cells are pixels (PICTure EElement)

In 3D = voxels (VOlumoc Element)

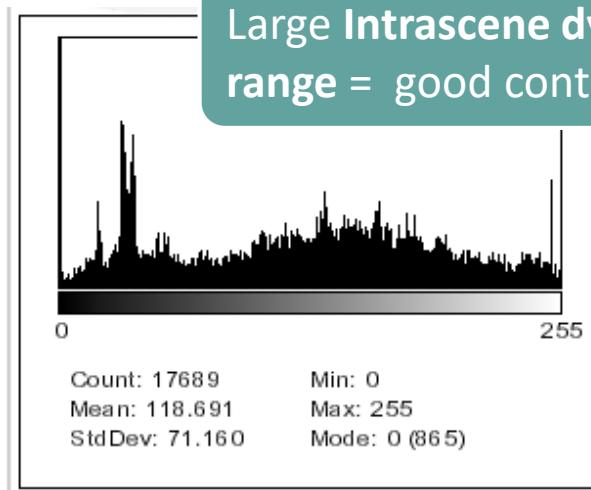
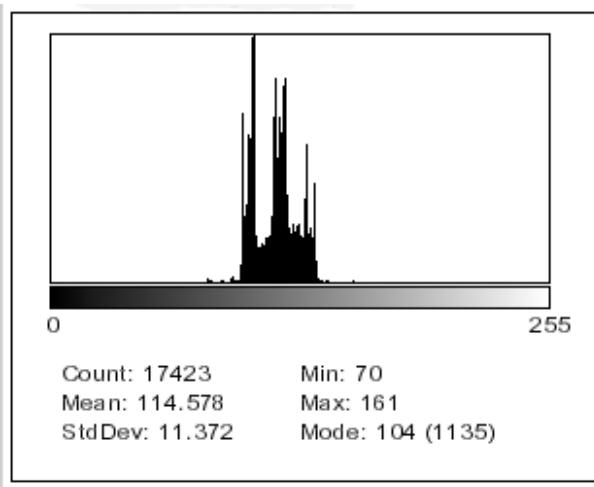
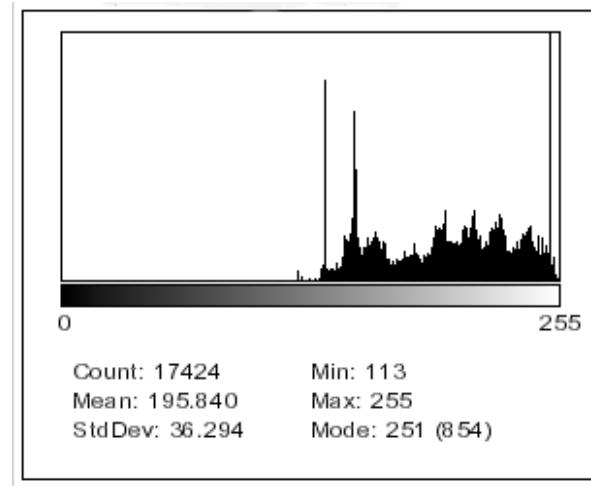
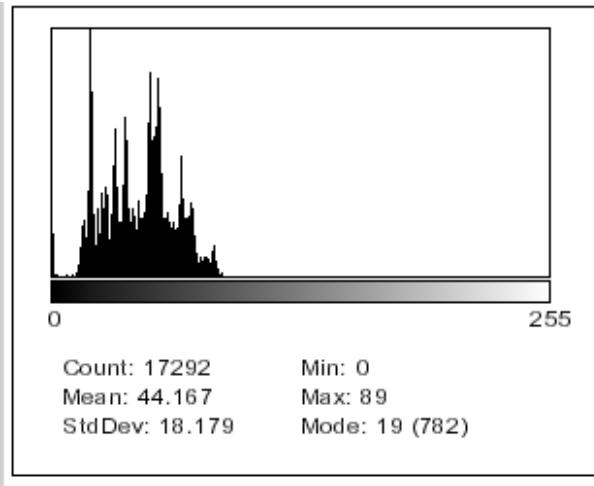
III/ Fundamentals: Histogram



Count: 1920000 Min: 0
Mean: 113.658 Max: 255
StdDev: 42.343 Mode: 135 (33747)
Bins: 256 Bin Width: 0.996

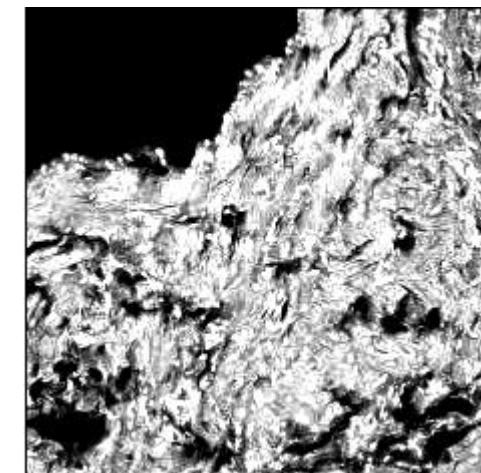
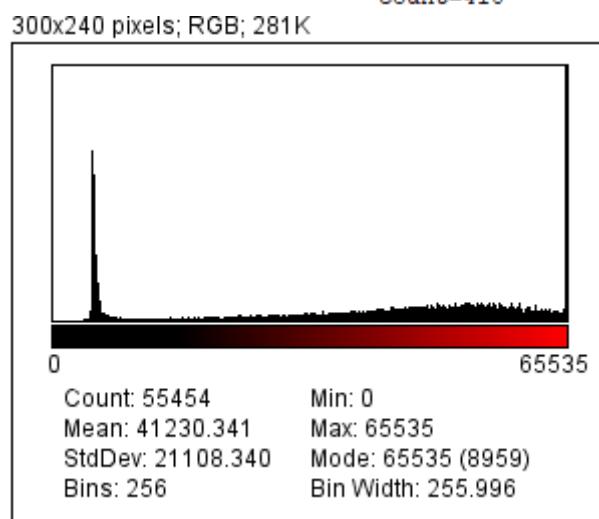
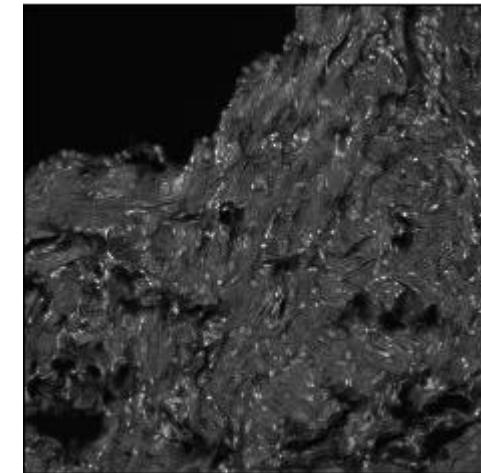
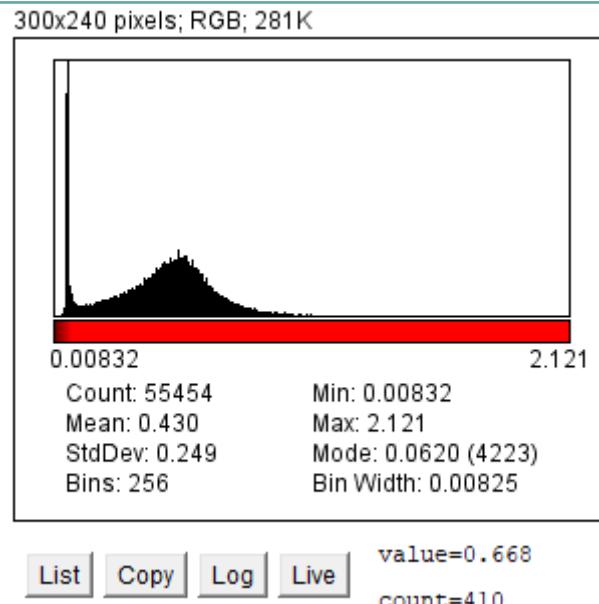
III/ Fundamentals: Using the full dynamic range of the camera

= ability to read very bright and very dark signals together

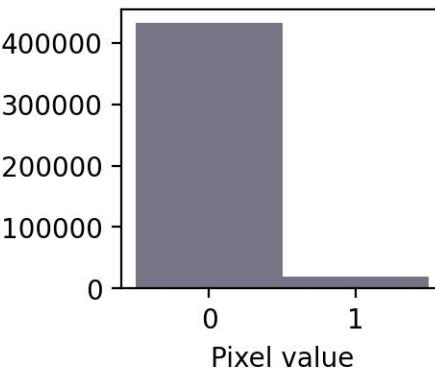
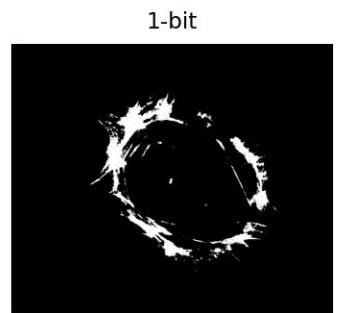


Large Intrascene dynamic range = good contrast!

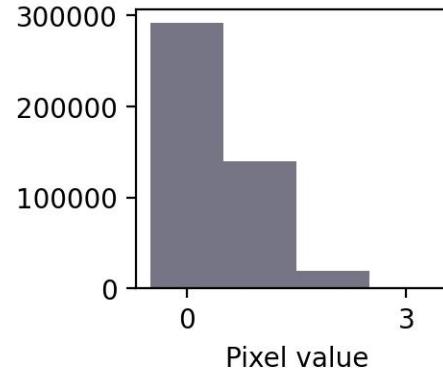
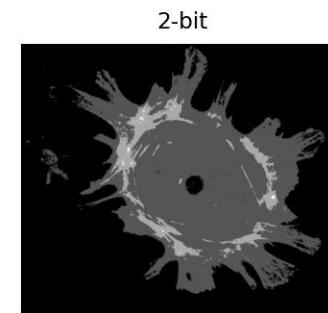
III/ Fundamentals: Saturation



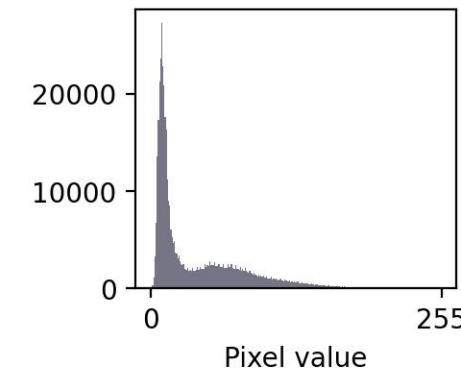
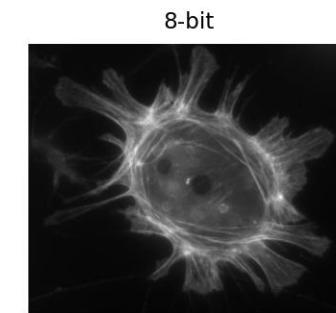
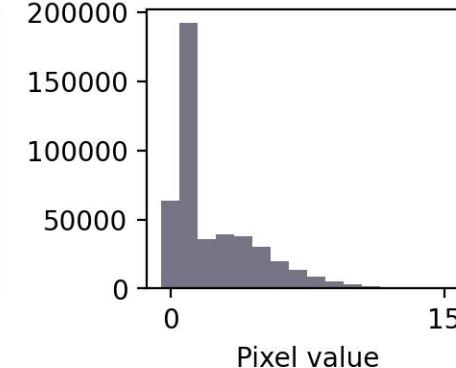
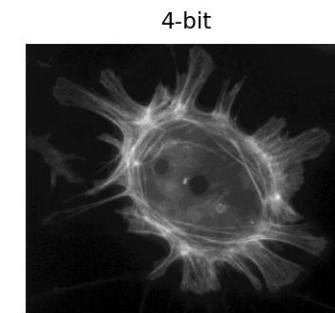
III/ Fundamentals: Bit depth



1-bit	
Decimal	Binary
0	0
1	1



2-bit	
Decimal	Binary
0	00
1	01
2	10
3	11



8-bit	
Decimal	Binary
0	00000000
1	00000001
2	00000010
3	00000011
4	00000100
5	00000101
6	00000110
7	00000111
...	...
255	11111111

- 8-bit: 0, 1, .., 255 ($2^8 - 1$)
 - 16-bit: 0, 1, .., 65 535 ($2^{16} - 1$)
 - 32-bit: 0, 1, .., 1 070 000 000 ($2^{32} - 1$)

https://bioimagebook.github.io/chapters/1-concepts/3-bit_depths/bit_depths.html#fig-bit-depths-demo

III/ Fundamentals: Detector type vs Image type

<u>Detector type</u>	<u>Range</u>	<u>Image type on the computer</u>	
8-bit	0 - 255	8-bit	<i>Much better dynamic range, good for intensity comparing</i>
12-bit (most PMTs & cameras)	0 - 4095		
14-bit (some EMCCD cameras)	0 - 16383	16-bit	<i>On the computer, everything is a multiple of 1 byte (=8 bits)</i>
16-bit (some PMTs)	0 - 65535		
∅	~ -Infinity - +Infinity (real numbers)	32-bit	<i>The only representation that permits negative numbers and floating point numbers</i>
Special			
Color cameras	RGB (color image)	3 layers of 8-bit each (red, green & blue)	

III/ Fundamentals: Display: Brightness Contrast

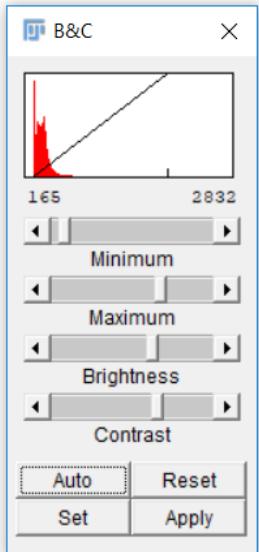


Image > Adjust > Brightness/Contrast...
Shift + C

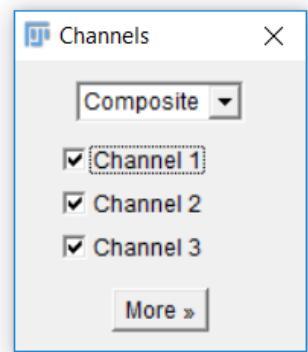
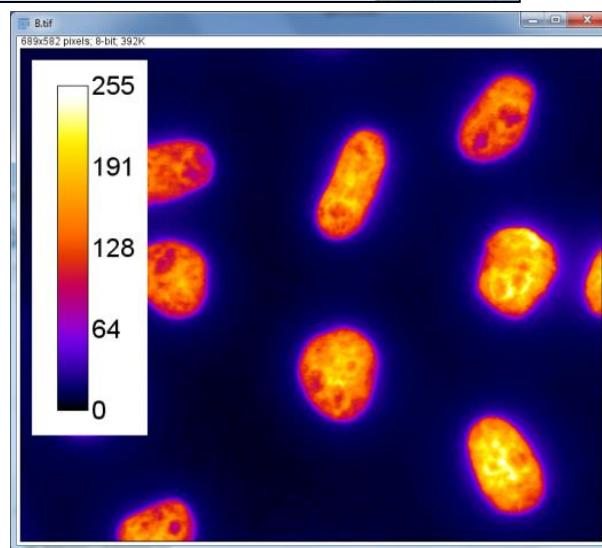
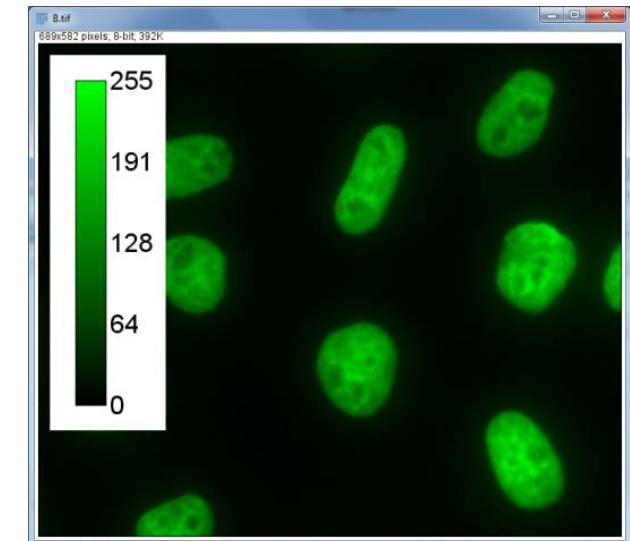
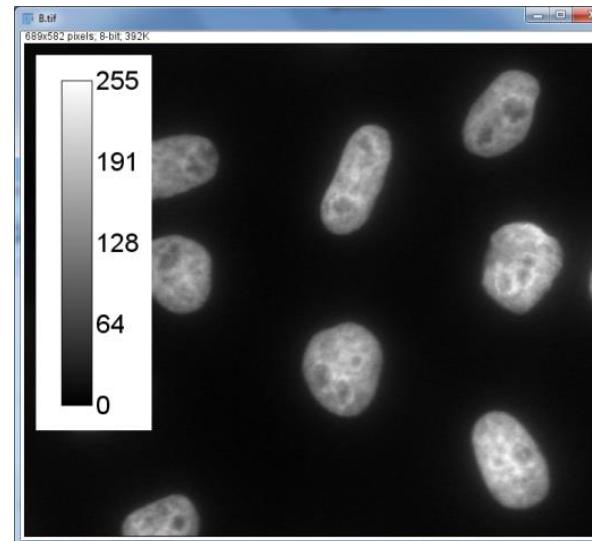
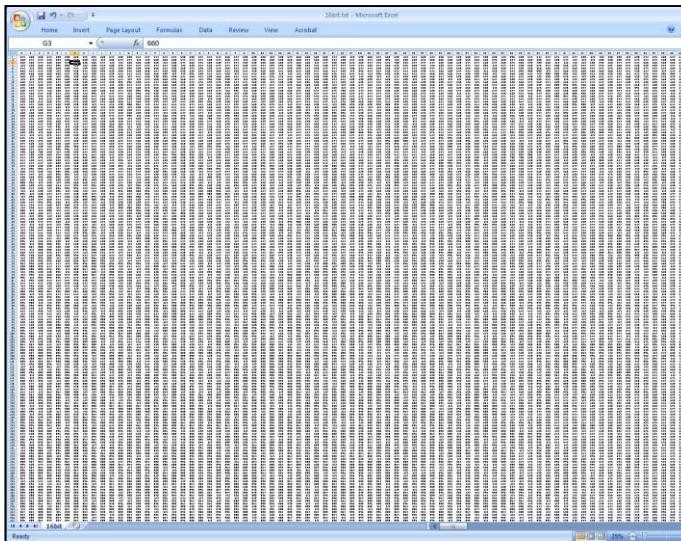


Image > Color > Channels Tool...
Shift + Z



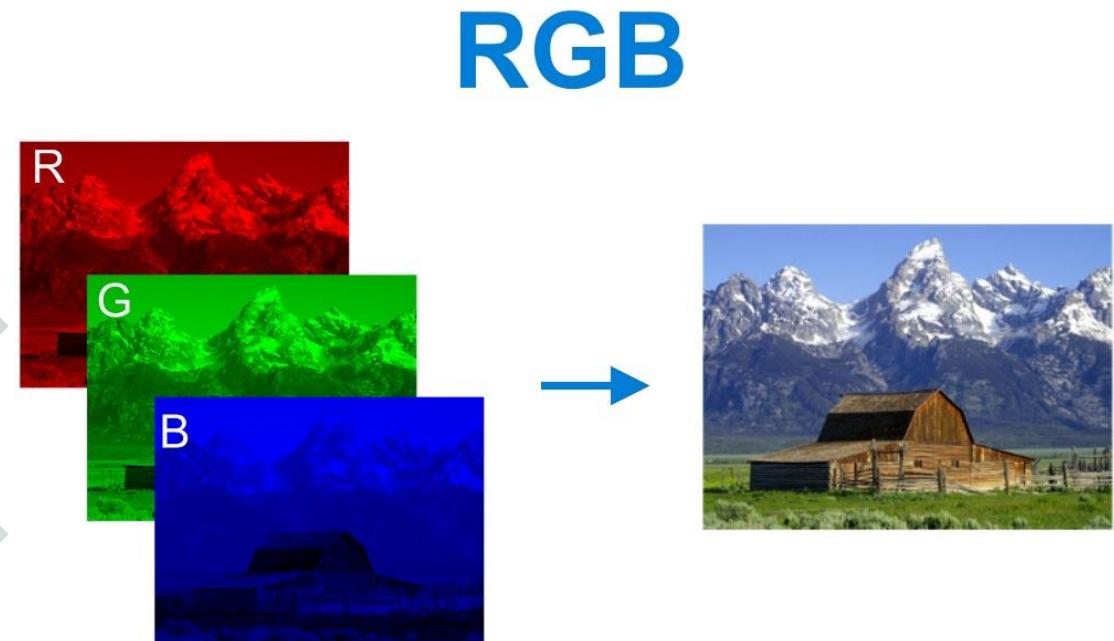
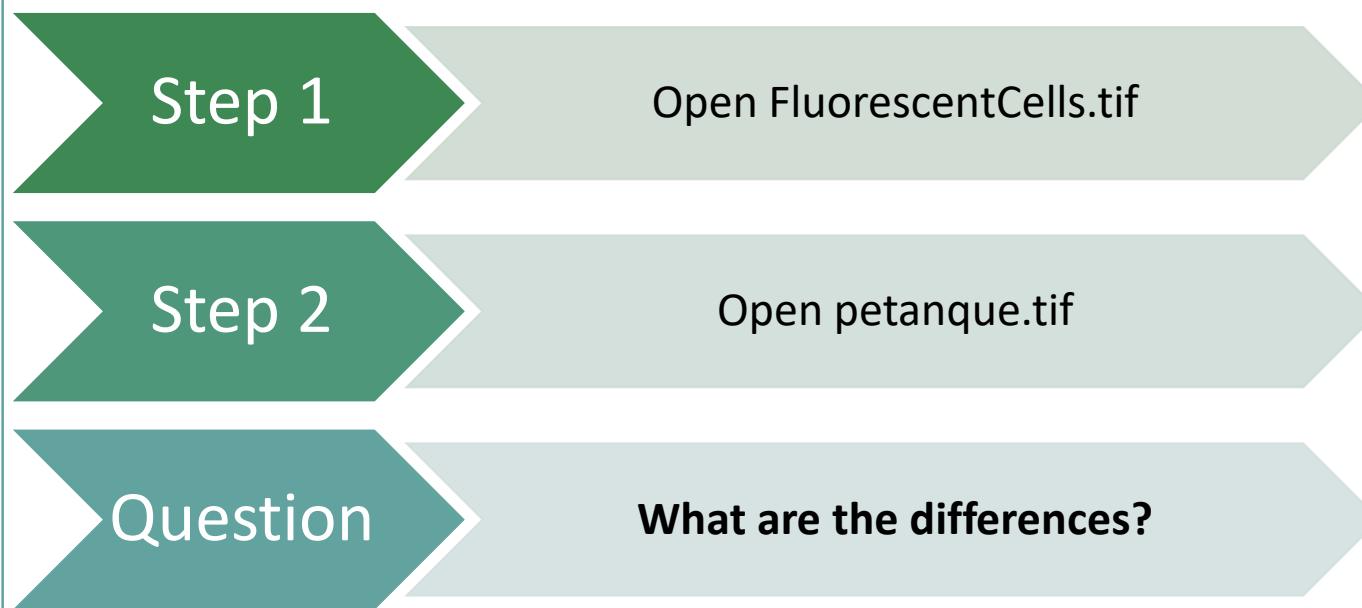
III/ Fundamentals: Display: Lookup Table (LUT) or colormap



Viridis LUT			
Red	Green	Blue	Color
68	1	84	
72	25	107	
70	47	124	
64	67	135	
56	86	139	
48	103	141	
41	120	142	
35	136	141	
30	152	138	
34	167	132	
53	183	120	
83	197	103	
121	209	81	
165	218	53	
210	225	27	
253	231	36	

LUT in RGB on Greyscale images:
the image is displayed thanks to
the mixing of 3 RED, GREEN and
BLUE values for one given
greyscale value.

III/ Fundamentals: RGB images



RGB images : each pixel has **three** values for RED, GREEN and BLUE

III/ Fundamentals: Read the information about an image



File>Open Samples> Mitosis

Aim: Get the dimensions, the sampling or pixel size (voxel in 3D),
the bit depth and the dynamic range of mitosis

Step 1

Get the dimensions (in X? Y? Z? C? T?)

*Observe the Information above the image.
Alternatively Image> Show Info*

Step 2

Get the bit depth

*Observe the Information above the image.
Alternatively Image> Show Info*

Step 3

Get the pixel size/ voxel size

Image>Properties

Step 4

Get the dynamic range

*Process>Histogram (min, max), Alternatively
Image>Show Info*

III/ Fundamentals: Read the information about an image



File>Open Samples> Mitosis

Aim: Get the dimensions, the sampling or pixel size (voxel in 3D),
the bit depth and the dynamic range of mitosis

Step 1

Get the dimensions (in X? Y? Z? C? T?)

X=171, Y=196, Z=5, C=2, T=51, 0,14286 seconds
between frames

Step 2

Get the bit depth

Bit-depth = 16 bits

Step 3

Get the pixel size/ voxel size

Pixel(Voxel in 3D) Size XYZ= 0.0885x0.0885x1 μm^3

Step 4

Get the dynamic range

c1 1582-6440; c2 1614-15787

Bonus: C1 and C2 content? Drosophila S2 cell
expressing GFP-Aurora B and mCherry-tubulin

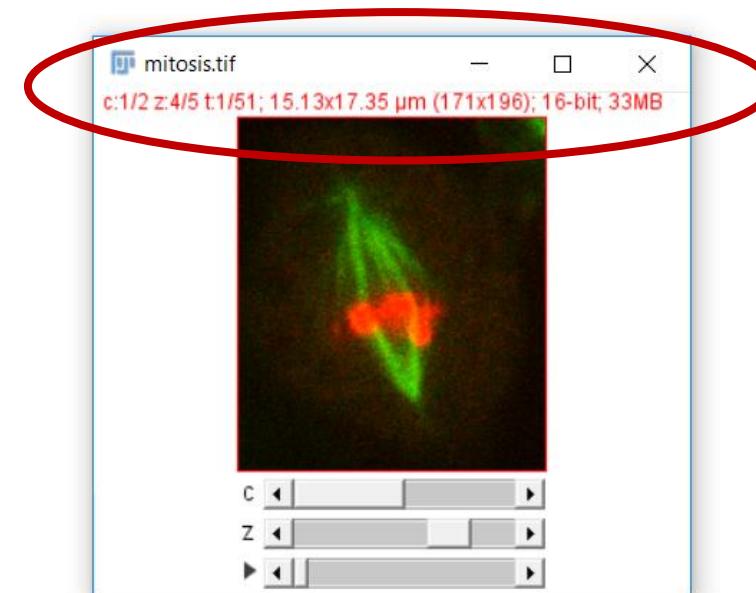
III/ Fundamentals: Show Info

Drosophila S2 cell expressing GFP-Aurora B and mCherry-tubulin fusion protein undergoing mitosis. Courtesy of Eric Griffis.

(Fiji Is Just) ImageJ 2.0.0-rc-69/1.52; Java 1.8.0_172 [64-bit]; Windows 10 10.0; 3145MB of 12141MB (25%)

Title: mitosis.tif
Width: 15.1335 μm (171)
Height: 17.3460 μm (196)
Depth: 5 μm (5)
Size: 33MB
Resolution: 11.2994 pixels per μm
Voxel size: 0.0885x0.0885x1 μm^3
ID: -26
Bits per pixel: 16 (unsigned)
Display ranges
1: 1582-6440
2: 1614-15787
Frame: 7/510
Frame interval: 0.14286 sec
Channel: 1/2
Slice: 4/5
Frame: 1/51
Composite mode: "composite"
No threshold
ScaleToFit: false
Uncalibrated
URL: <http://imagej.nih.gov/ij/images/Spindly-GFP.zip>
Screen location: 1171,328 (2048x1152)
Coordinate origin: 0,0,0
No overlay
No selection

Image > Show Info (i)



III/ Fundamentals: Hyperstacks

Image 2D

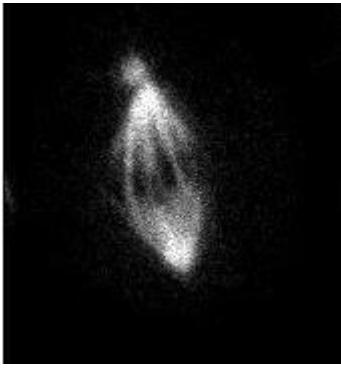


Image 3D:

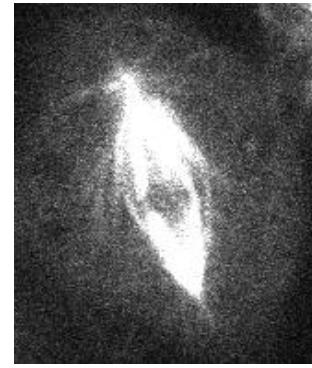
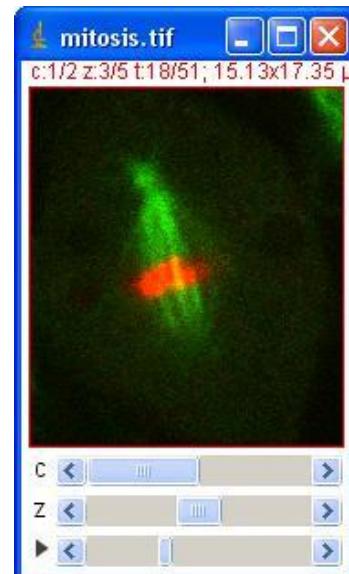
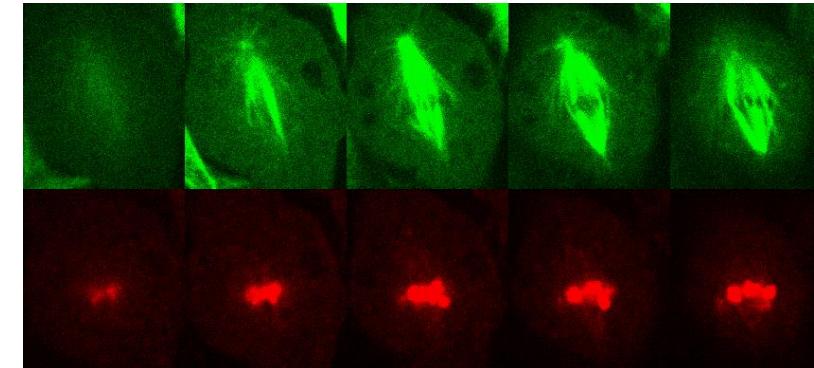


Image 4D+: Hyperstack

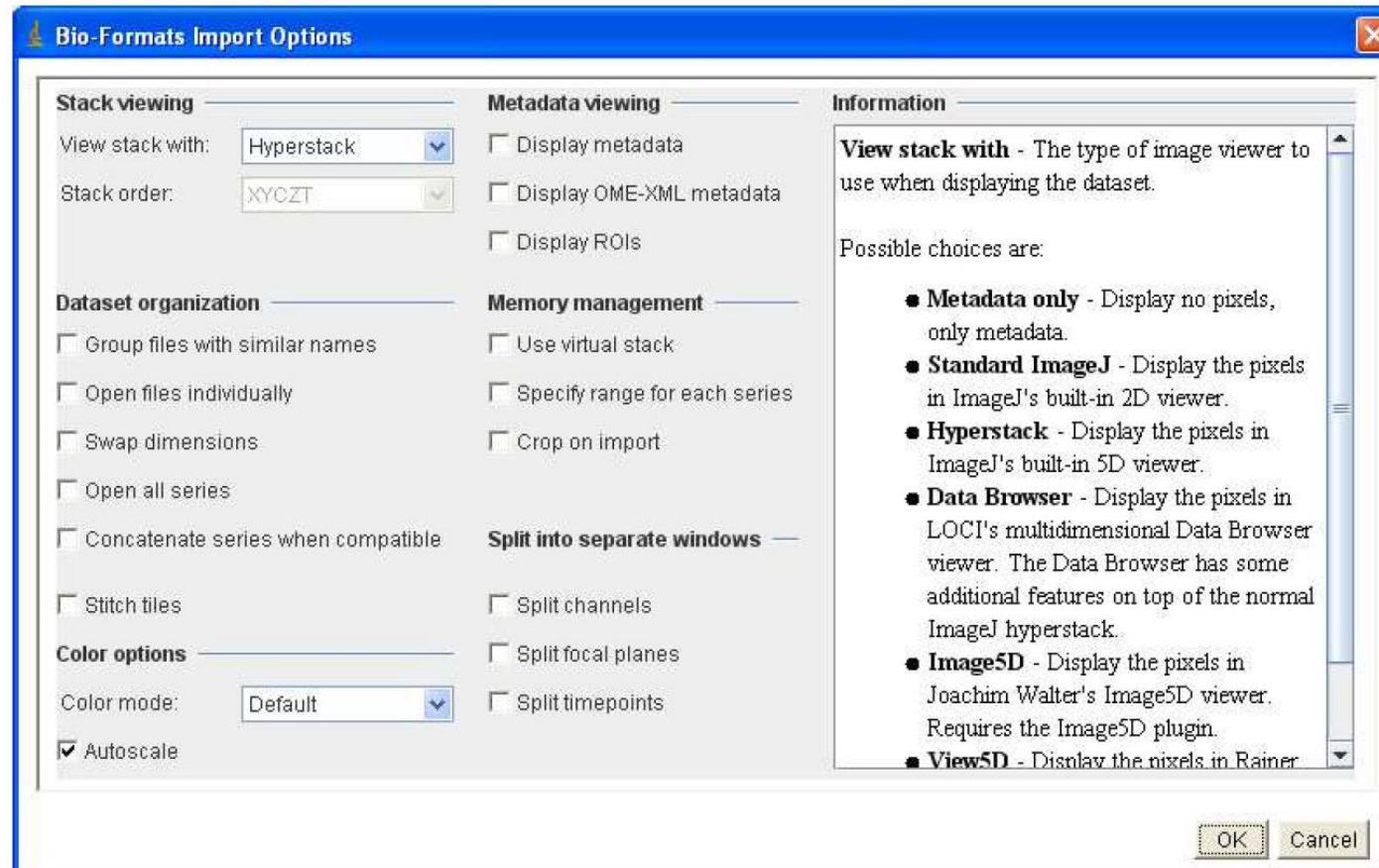


Hyperstack = 5 dimensions

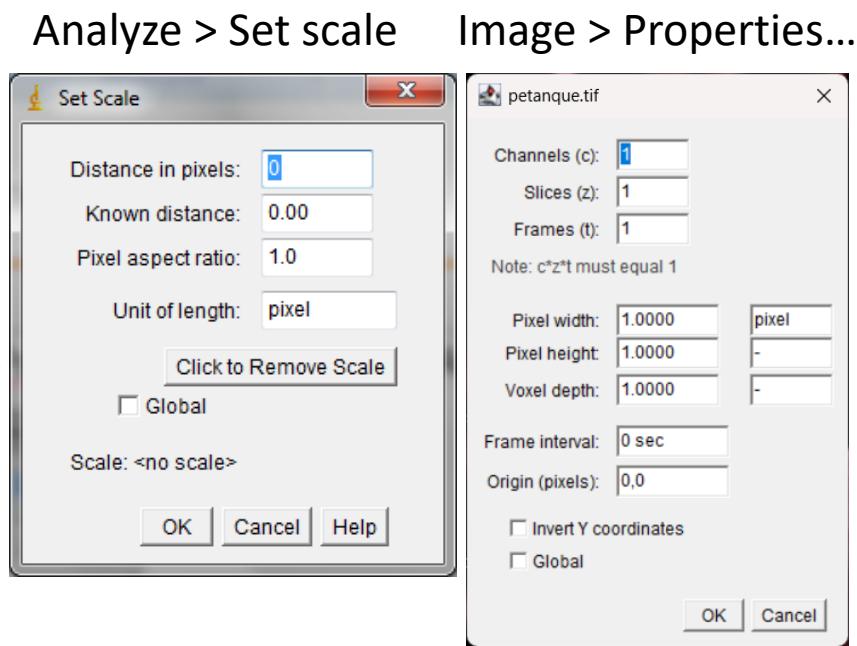
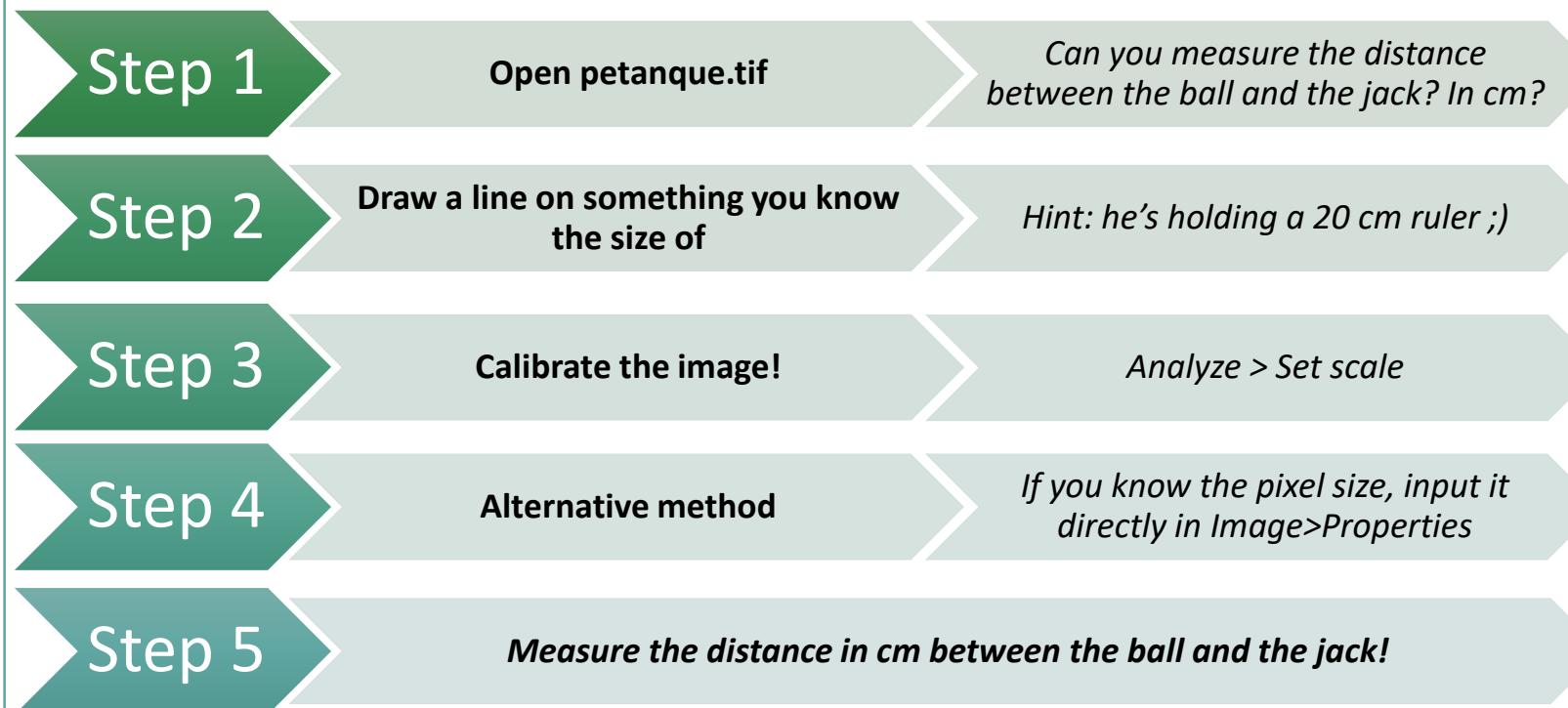
- x
- y
- z (slice)
- channel
- time (Frame)

III/ Fundamentals: Hyperstacks

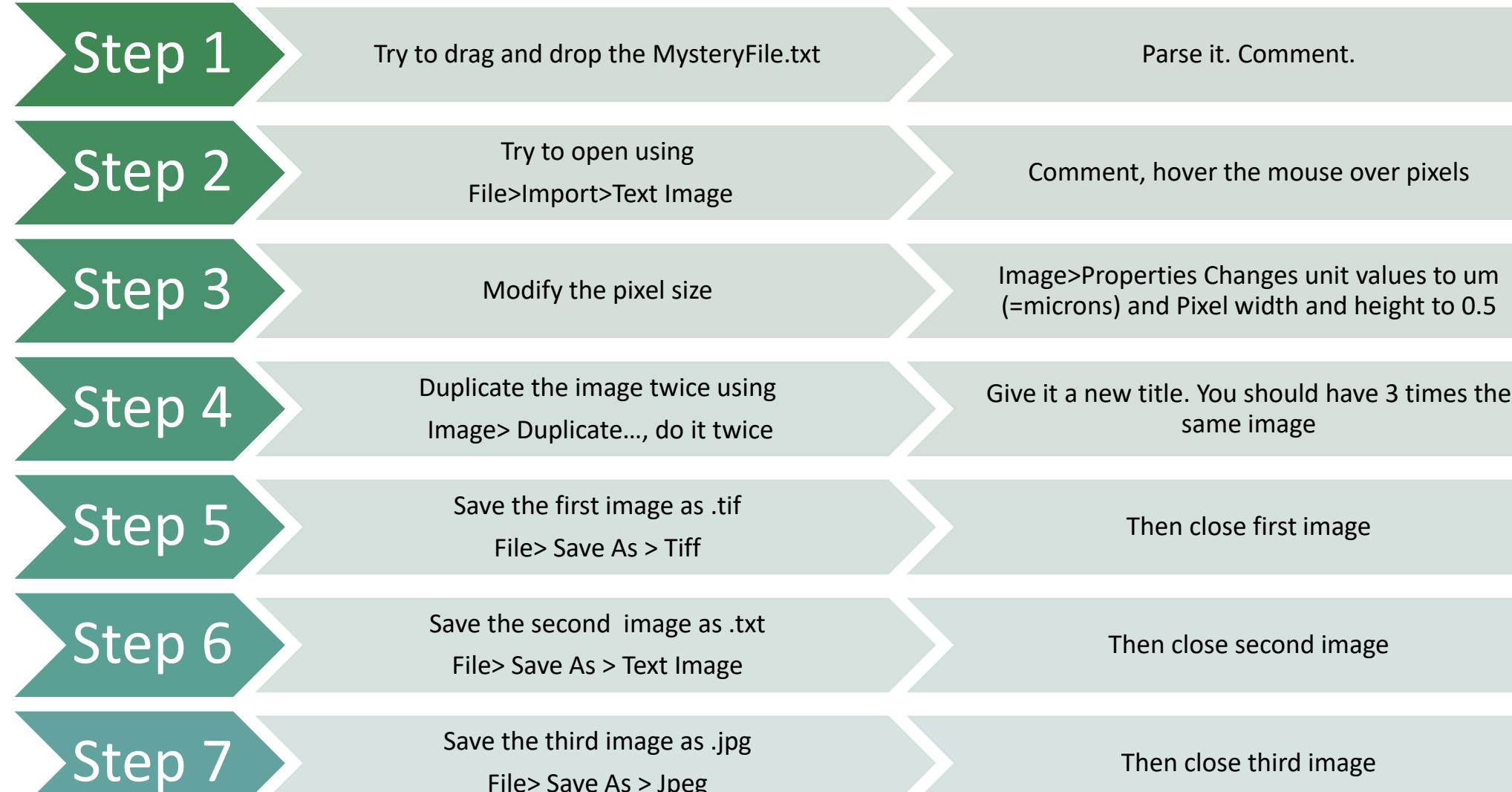
Can be opened with Bio-Formats



III/ Fundamentals: Calibration: *Hands-on!*



III/ Fundamentals: Image format and image storage: *Hands-on!*

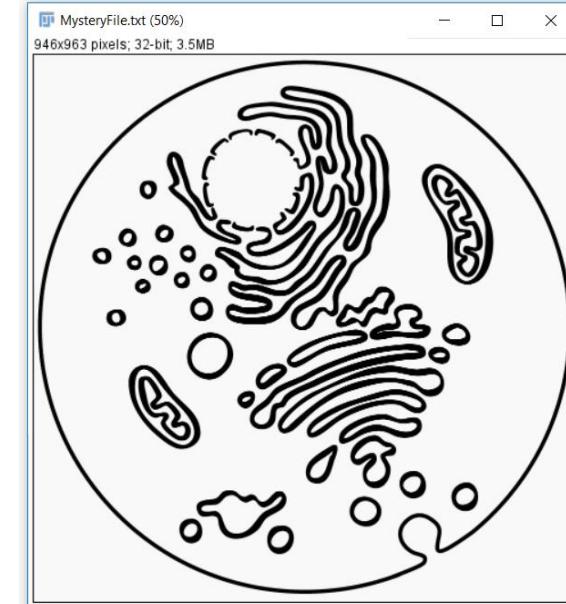
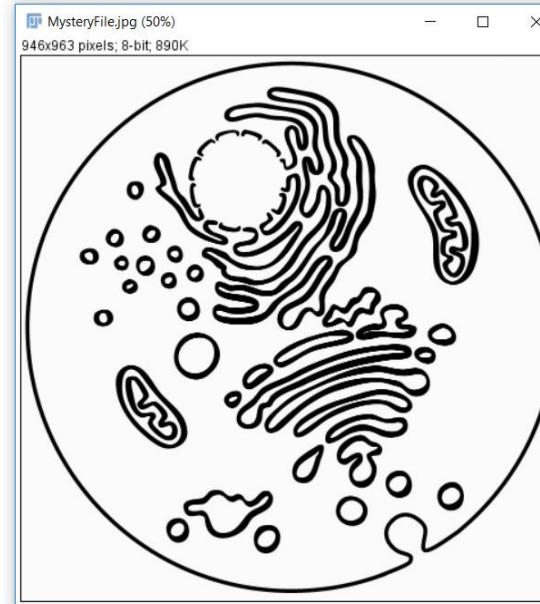
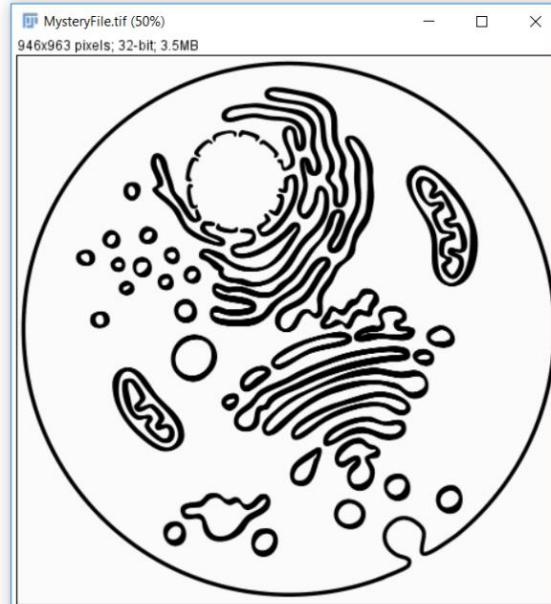


III/ Fundamentals: Image format and image storage: *Hands-on!*

Now look at the directory content. What do you notice?

 MysteryFile.tif	16/01/2019 18:39	Image TIFF	3 559 Ko
 MysteryFile.jpg	16/01/2019 18:39	Fichier JPG	141 Ko
 MysteryFile.txt	16/01/2019 18:39	Document texte	7 119 Ko

Open the 3 images again (Do not forget to use Import>Text Image for the .txt)



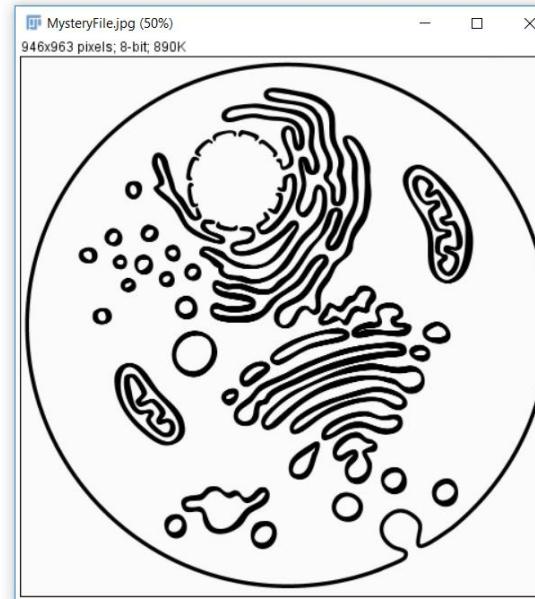
Different?

III/ Fundamentals: Image format and image storage: *Hands-on!*

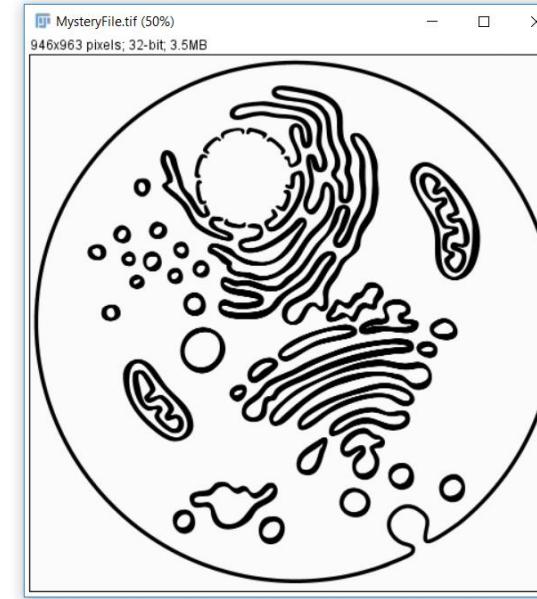
Conclusions:

Some formats store the metadata (here pixel size) in addition to the pixel data (ex: .tif)

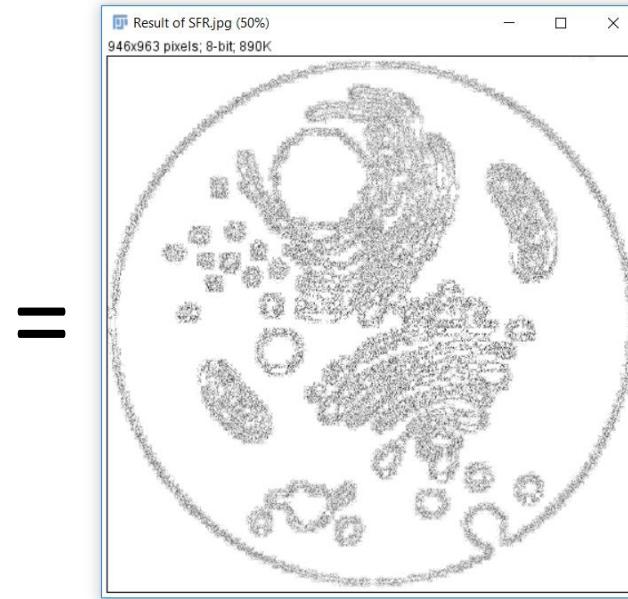
The way the data is stored is different -> compression



Jpeg image we created



Tiff image we created



Jpeg – tiff (pixel by pixel difference)

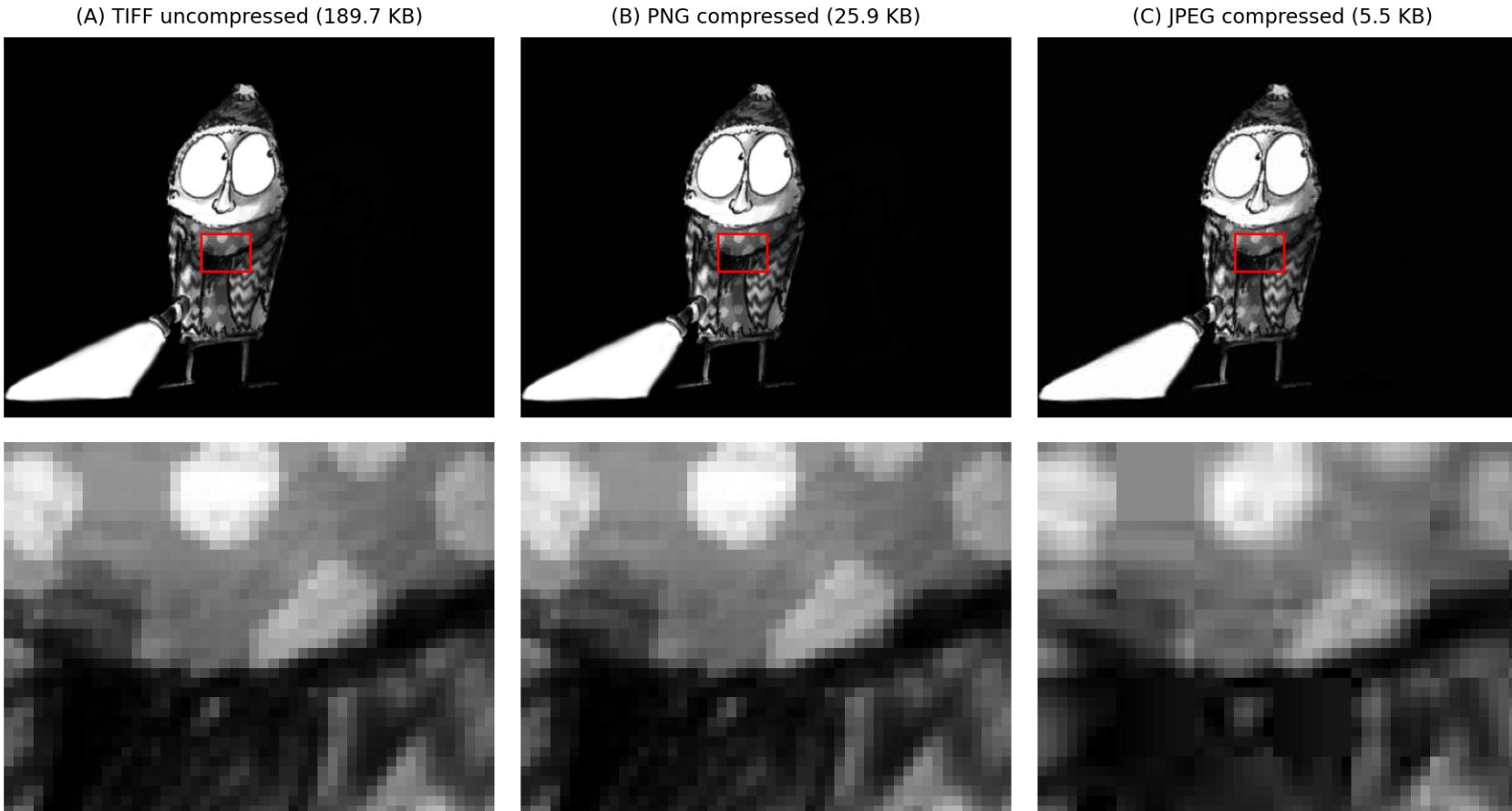
III/ Fundamentals: Image format and image storage

Images files= pixel values and metadata (dimension, image type, bit-depth, pixel size, microscope parameters...)

Format	Extensions	Main use	Compression	Comment
TIFF	.tif, .tiff	Analysis, display (print)	None, lossless, lossy	Very general image format
OME-TIFF	.ome.tif, .ome.tiff	Analysis, Display (print)	None, lossless, lossy	TIFF, with standardized metadata for microscopy
Zarr	.zarr	Analysis	None, lossless, lossy	Emerging format, great for big datasets – but limited support currently
PNG	.png	Display (web, print)	Lossless	Small(ish) file sizes without compression artefacts
JPEG	.jpg, .jpeg	Display (web)	Lossy (usually)	Small file sizes, but visible artefact

III/ Fundamentals: Image format and image storage

<https://bioimagebook.github.io/chapters/1-concepts/6-files/files.html>



III/ Fundamentals: Recap: concepts

Digital Images

They are matrices!

Histograms

An eagle-eye view of your data. Great to see what is wrong with it (e.g. saturation)

Dynamic range

The ability of the sensor to image bright and dark signal simultaneously: if low, bright signals can saturate the sensor while weak become lost in the noise. Adjust your settings so you make good use of it!

Bit depth

How finely chopped your signal is: the bigger it is, the more shades of grey, the more details can be replicated in an image. The

greater the bit depth, the better the full dynamic range can be resolved. Make sure you have the adequate one when sitting at the microscope

Display adjustments

- **Brightness & contrast:** they do not change your data, just the way it is represented on the screen.
- **LUT:** Same. Pick the best for a snapshot of your data.

Hyperstacks

3D = stack. 4D+=Hyperstack

Calibration

Demonstrated that the *metadata* is crucial in microscopy! Beware of lossy formats.

III/ Fundamentals: Recap: commands

Image > Adjust > Brightness/Contrast (**Shift + c**)

Image > Show info (**i**)

Image > Properties (**Shift + p**)

Image > Color > Channels Tool (**Shift + z**)

Analyze > Histogram (**h**)

Image > Lookup Tables

List of shortcuts: <https://imagej.net/ij/docs/shortcuts.html>

III/ Fundamentals: Other basic commands

Edit > Invert (**Shift + i**)

Image > Duplicate (**Shift + d**)

Image > Crop (**Shift + x**)

Image > Rename

Image > Transform > Rotate/Flip ...

Image > Adjust > Size

Image > Color > Split Channels/Merge Channels

Selection > Select All (**a**)

Selection > Select None (**Shift + a**)

Selection > Restore (**Shift + e**)

Selection > Add to Manager (**t**)

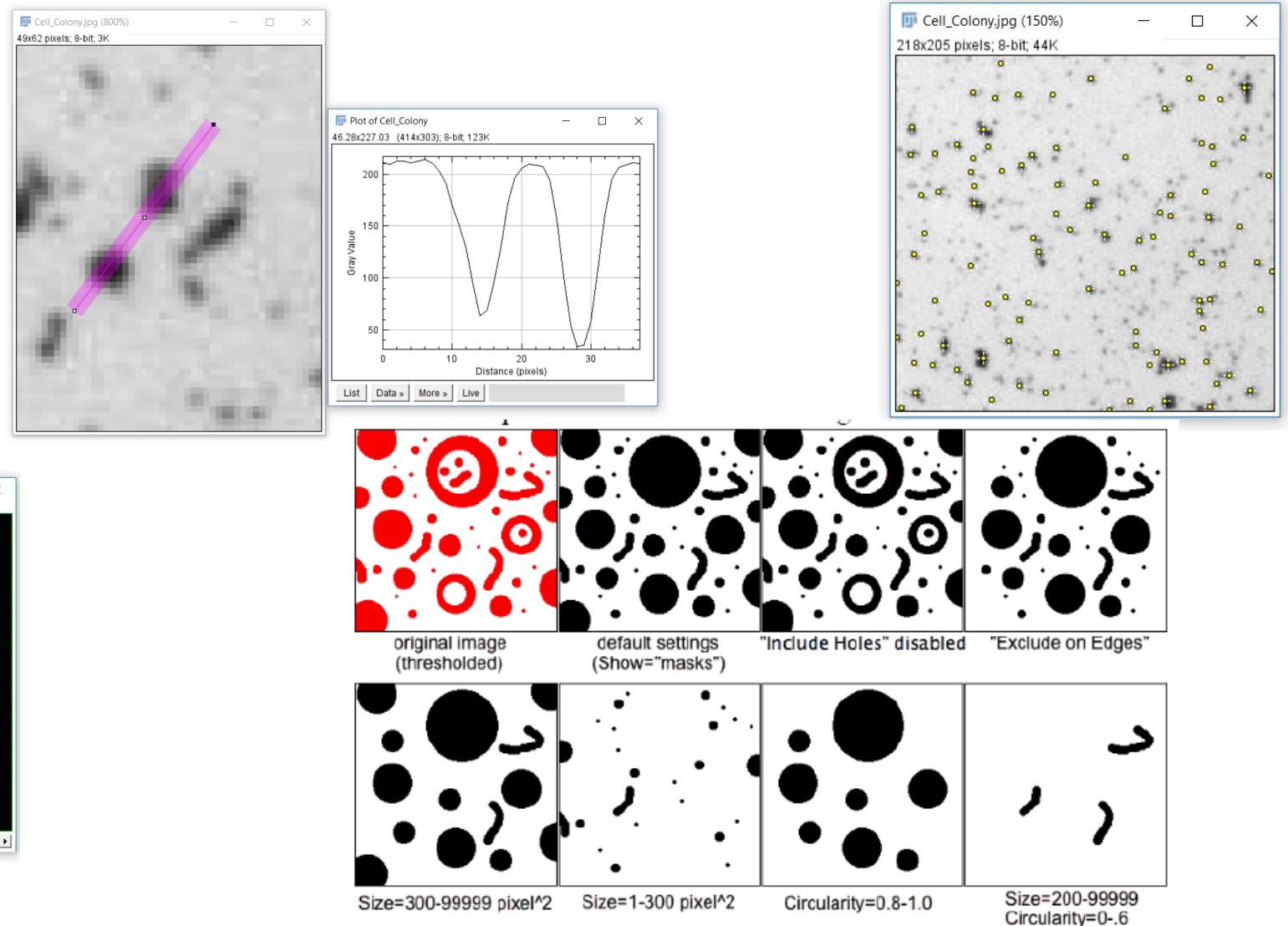
III/ Fundamentals: still a lot left to learn!

Image > Stacks > Z Project

Process > Find maxima

Analyze > Plot profile

Analyze > Analyze Particles



IV/ Image ethics

IV/ Image ethics: Ethical guidelines for the appropriate use and manipulation of scientific digital images (Cromey, 2010):

- 1/ Scientific digital images are **data that can be compromised** by inappropriate manipulations.
- 2/ Manipulation of digital images should only be performed on a **copy** of the unprocessed image data file (Always keep the original data file safe and unchanged!).
- 3/ **Simple adjustments** to the **entire** image are **usually acceptable**.
- 4/ **Cropping** an image is **usually acceptable**.
- 5/ Digital images that will be compared to one another should be **acquired under identical conditions**, and any **post-acquisition image processing** should also be **identical**.
- 6/ **[“Local”] manipulations** that are specific to one area of an image and are not performed on other areas are **questionable**.

Cromey, Douglas W. « Avoiding Twisted Pixels: Ethical Guidelines for the Appropriate Use and Manipulation of Scientific Digital Images ». *Science and Engineering Ethics* 16, n° 4 (2010): 639-67. <https://doi.org/10.1007/s11948-010-9201-y>.

IV/ Image ethics: Ethical guidelines for the appropriate use and manipulation of scientific digital images (Cromey, 2010):

- 7/ Use of **software filters to improve image quality** is usually **not recommended** for biological images.
- 8/ **Cloning or copying objects** into a digital image, from other parts of the same image or from a different image, is **very questionable**.
- 9/ **Intensity measurements** should be performed on **uniformly processed image data**, and the data should be **calibrated** to a known standard.
- 10/ **Avoid the use of lossy compression**.
- 11/ **Magnification** and **resolution** are important.
- 12/ Be **careful when changing the size** (in pixels) of a digital image.

Cromey, Douglas W. « Avoiding Twisted Pixels: Ethical Guidelines for the Appropriate Use and Manipulation of Scientific Digital Images ». *Science and Engineering Ethics* 16, n° 4 (2010): 639-67. <https://doi.org/10.1007/s11948-010-9201-y>.

IV/ Image ethics: Ressources for good practices

<https://focalplane.biologists.com/2021/05/25/preparing-your-manuscript-guidelines-for-writing-microscopy-methods-and-figures/>

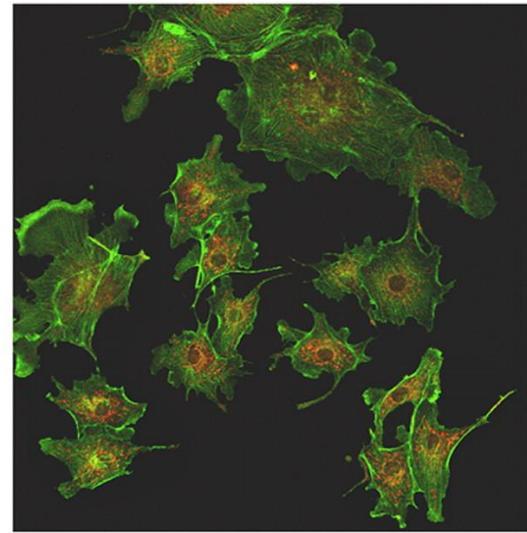
<https://www.nature.com/documents/nprot-guide-to-preparing-final-artwork.pdf>

1. Heddleston JM, Aaron JS, Khuon S, Chew TL. [‘A guide to accurate reporting in digital image acquisition – can anyone replicate your microscopy data?’](#) J Cell Sci. 2021 Mar 30;134(6):jcs254144. doi: 10.1242/jcs.254144.
2. Aaron J, Chew TL. [‘A guide to accurate reporting in digital image processing – can anyone reproduce your quantitative analysis?’](#) J Cell Sci. 2021 Mar 30;134(6):jcs254151. doi: 10.1242/jcs.254151.
3. Jambor H, Antonietti A, Alicea B, Audisio TL, Auer S, Bhardwaj V, Burgess SJ, Ferling I, Gazda MA, Hoeppner LH, Ilangoan V, Lo H, Olson M, Mohamed SY, Sarabipour S, Varma A, Walavalkar K, Wissink EM, Weissgerber TL. [‘Creating clear and informative image-based figures for scientific publications’](#) PLoS Biol. 2021 Mar 31;19(3):e3001161. doi: 10.1371/journal.pbio.3001161.
4. Schmied C and Jambor HK. [Effective image visualization for publications – a workflow using open access tools and concepts](#) [version 2; peer review: 2 approved]. F1000Research 2021, 9:1373 (<https://doi.org/10.12688/f1000research.27140.2>).
5. Schmied, Christopher, Michael S. Nelson, Sergiy Avilov, Gert-Jan Bakker, Cristina Bertocchi, Johanna Bischof, Ulrike Boehm, et al. « Community-Developed Checklists for Publishing Images and Image Analyses ». *Nature Methods*, 14 septembre 2023, 1-12. <https://doi.org/10.1038/s41592-023-01987-9>.
6. Miura, Kota, et Simon F Nørrelykke. « Reproducible image handling and analysis ». *The EMBO Journal* 40, n° 3 (février 2021). <https://doi.org/10.15252/embj.2020105889>.

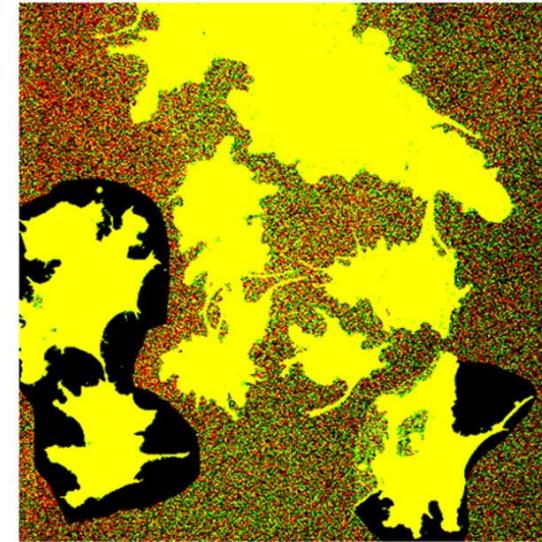
IV/ Image ethics: Mishandling can be intentional

Rossner & Yamada, JCB 2004

Manipulated image



Manipulation revealed by contrast adjustment

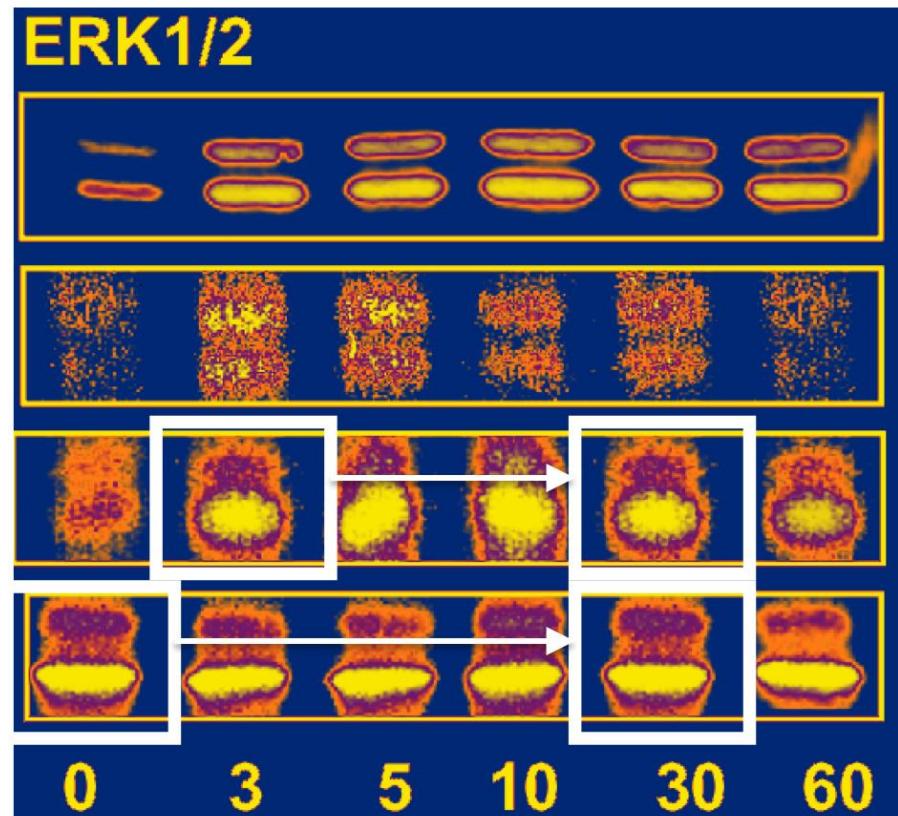


Copy / pasting cells into an image.

Slide: J-Y Tinevez

IV/ Image ethics: Mishandling can be intentional

Simple tools to detect image manipulation.



Copy/paste within images.
But you can detect them easily with false color LUT.

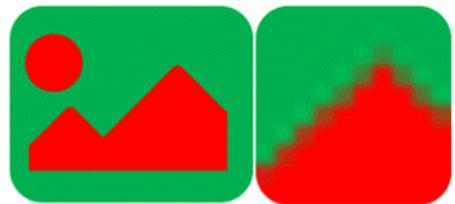
Solution to avoid mishandling? Yes,
REPORTING!

Document all you did to prepare the figures

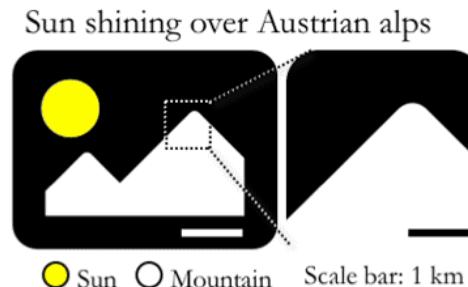
- Materials and Methods (supp info).
- Jupyter-like Notebooks or scripts to generate the figures.

IV/ Image ethics: Common mistakes

Schmied C and Jambor HK. [Effective image visualization for publications – a workflow using open access tools and concepts](#) [version 2; peer review: 2 approved]. F1000Research 2021, 9:1373 (<https://doi.org/10.12688/f1000research.27140.2>).

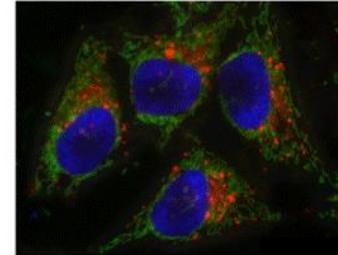


- Compression artifacts
- Inset not indicated
- Not color-blind safe
- Lack of scale information
- Lack of color annotation

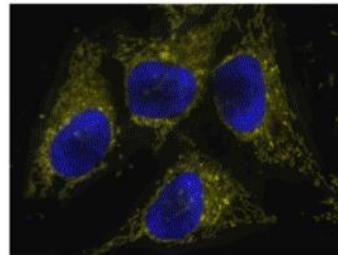


- Image quality good
- Inset origin clear
- Colors accessible
- Scale bar
- Color annotation

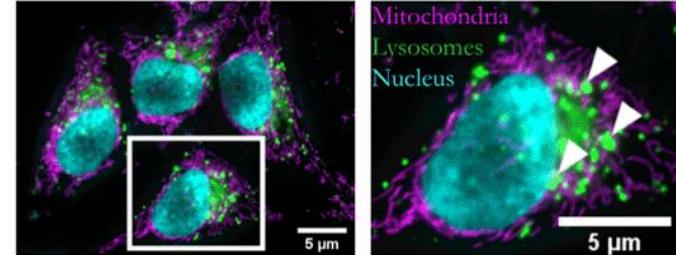
Poorly visualized image



- not color blind safe
- no annotation
- no scale



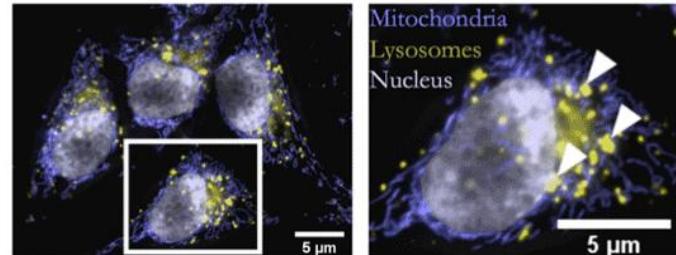
Workflow-based image processing for publication



HeLa cells

Test of Image quality:
Color blind (Deutanopia)
rendering of images

- colors accessible to color blind
- annotation, annotations color blind safe
- scale bar for image and inset



HeLa cells

IV/ Image ethics: Good practices

Schmied, Christopher, Michael S. Nelson, Sergiy Avilov, Gert-Jan Bakker, Cristina Bertocchi, Johanna Bischof, Ulrike Boehm, et al. "Community-Developed Checklists for Publishing Images and Image Analyses." *Nature Methods*, September 14, 2023, 1–12. <https://doi.org/10.1038/s41592-023-01987-9>.

Checklist for image publishing

Image format



Focus on relevant image content (e.g., crop, rotate, resize)

Minimal



Separate individual images



Show example image used for quantifications

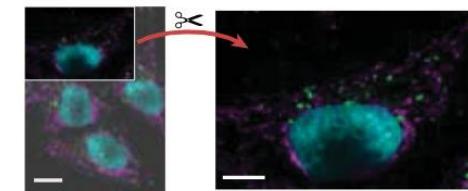
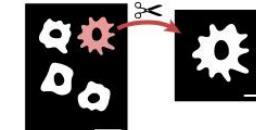


Indicate position of zoom view/inset in full-view/original image

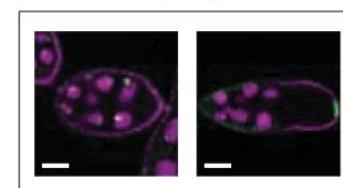


Show images of the range of the described phenotype

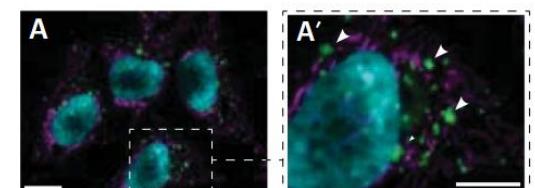
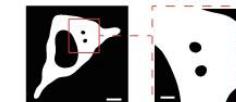
a Focus: crop, rotate, etc.



b Separate individual images clearly



c Show origin of inset Consider showing full-size image



IV/ Image ethics: Good practices

Image colors and channels

 Annotation of channels (staining, marker, etc.) visible

 Adjust brightness/contrast, report adjustments, use uniform color scales

 Image comparison: use the same adjustments

 Channel colors: high visibility on the background
Best visibility: grayscale

 Multicolors: provide grayscale for each color channel

 Multicolor: if channels are merged, make accessible to color-blind individuals

 Provide intensity scales (calibration bar) for grayscale, color, pseudocolor etc.

 Pseudocolored images: additionally provide grayscale version for comparison

 Gamma adjustments: additionally provide linear-adjusted image for comparison

Minimal

Recommended

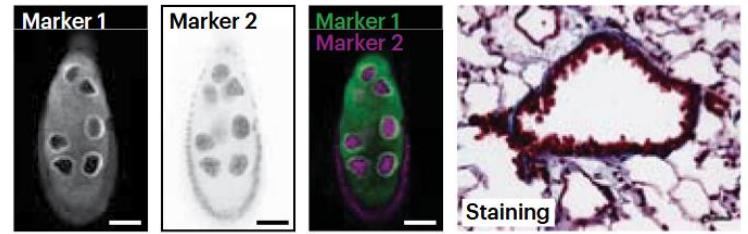
Ideal

e Channel annotations visible on background



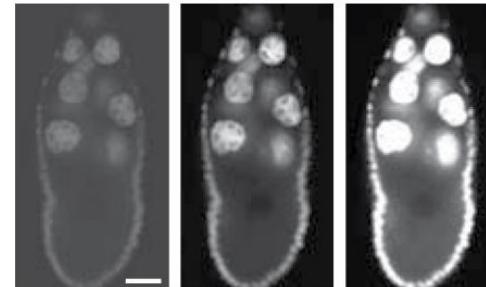






d Brightness/contrast adjustment





"This is also, why it is important to tell young researchers, that using the same method on all images does not mean that the same things happen on all of them and that they will stay comparable."

IV/ Image ethics: Good practices

Image colors and channels

 Annotation of channels (staining, marker, etc.) visible

Minimal

 Adjust brightness/contrast, report adjustments, use uniform color scales

 Image comparison: use the same adjustments

 Channel colors: high visibility on the background
Best visibility: grayscale

 Multicolors: provide grayscale for each color channel

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 Provide intensity scales (calibration bar) for grayscale, color, pseudocolor etc.

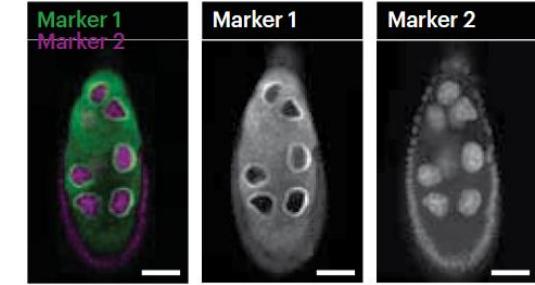
Recommended

 Pseudocolored images: additionally provide grayscale version for comparison

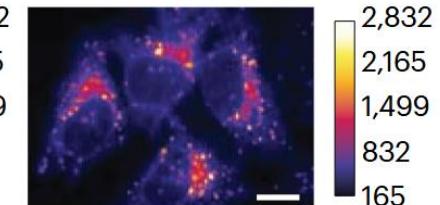
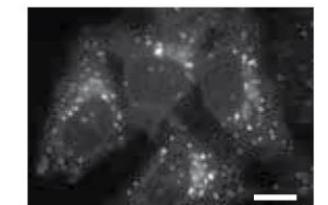
Ideal

 Gamma adjustments: additionally provide linear-adjusted image for comparison

f Grayscale channels improve visibility of image features



g Explain channel values with calibration scales/indicators



IV/ Image ethics: Good practices

Image annotation

 Add scale information (scale bar, image length in figure/figure legend)

Minimal

Aa= Explain all annotations (in figure/figure legend)

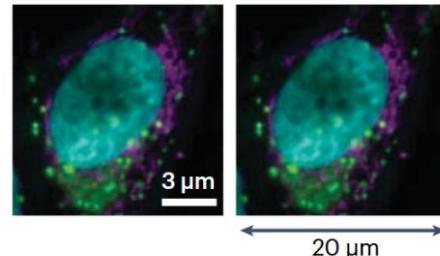
Aa Annotations should be legible (line width, size/point size, color)

A  Annotations should not obscure key data

 Annotate imaging details important for interpreting the figure (depending on the main message and imaging technique, this may be, e.g., image pixel size, imaging intervals (time-lapse in movies), exposure time or anatomical section)

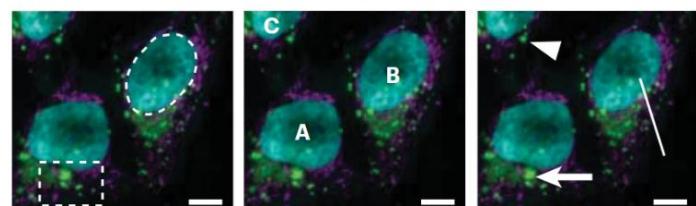
Recommended

a Scale information

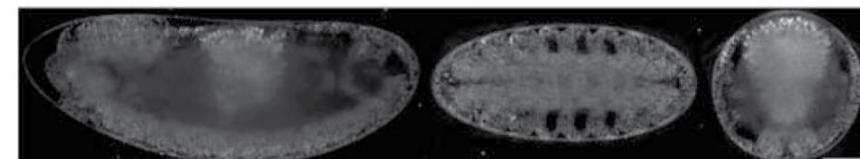


b Annotations: explained, visible, not obscuring content

Aa

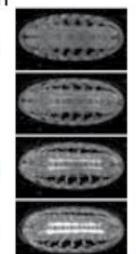


c Anatomical slices of a 3D volume



d Time lapse

hh:mm



04:30

05:15

06:00

06:45

IV/ Image ethics: Good practices

Image availability



Images are shared
(lossless compression/microscope images)



Image files are freely downloadable
(public database)



Image files are in dedicated image database
(added-value database or image archive)



Recommended



Ideal

e Share images



Local file server costs
Size limit, N/A

f Deposit images in public database



No costs/costs for >20 GB
Size limited, ~20–50 GB

g Images in dedicated image database



No costs
No/large size limit
Strict metadata/file type requirements

Provide lossless compressed or original microscope images