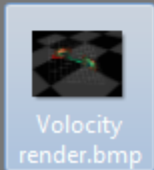
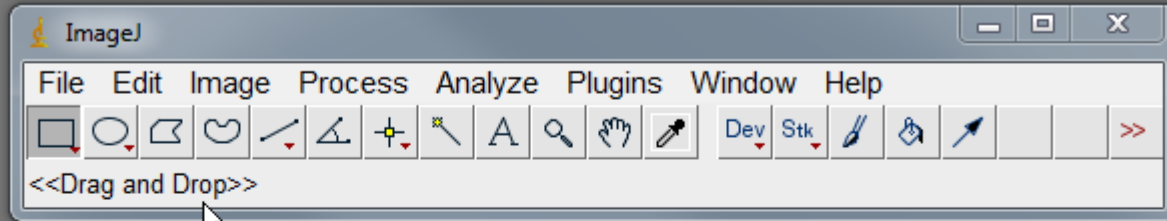


An introduction to image processing using ImageJ

Mark Willett, Imaging and Microscopy Centre,
Centre for Biological Sciences, University of Southampton.

Drag and drop onto the menu bar to open single or multiple images or stacks



[Shift] + Left click to select and open everything between two selections in a list

[Ctrl] + Left click to make multiple individual selections

A note on image formats

Images are comprised of pixels, each with an x,y coordinate and an intensity value that gives the pixel contrast in comparison to adjacent pixels.

The bit-depth of an image is defined by the number of intensity values available to each of the pixels in an image.

e.g.

Binary = 2 values, (black or white)

8-bit = 256 values. Black (0), white (255) and a scale of 254 grey steps in between

12-bit = 4,096 grey values

16-bit = 65,536 grey values

RGB formats are colour images comprised of 3 channels (red, green and blue) each channel has independent grey values, usually 8-bit

e.g. an RGB image that has 3 channels x 256 possible values per pixel is $3 \times 8\text{-bit} = 24\text{-bit}$

A note on image formats

The standard scientific image file format is the uncompressed Tagged Image File Format (.tiff, .tif) although some microscope manufacturers use their own similar proprietary lossless formats.

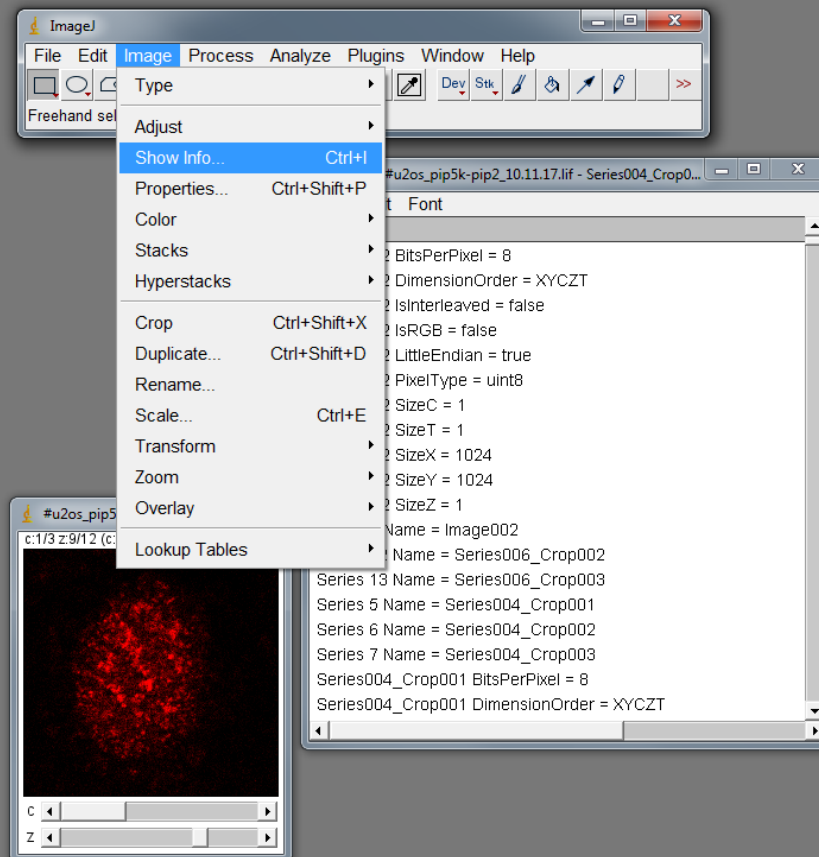
.Tiffs contain an embedded notepad file that contains essential information about the image such as scaling, laser wavelengths etc (metadata).

Non-scientific file formats such as .jpeg should be avoided as they are “lossy” (some of the image information is lost because they are compressed to reduce the file size, but are usually ok for presentations).

It’s usually also best to avoid Windows art packages and non-scientific image viewing and manipulation packages as they may apply compression to your image and display your image differently from scientific viewing and analysis packages.

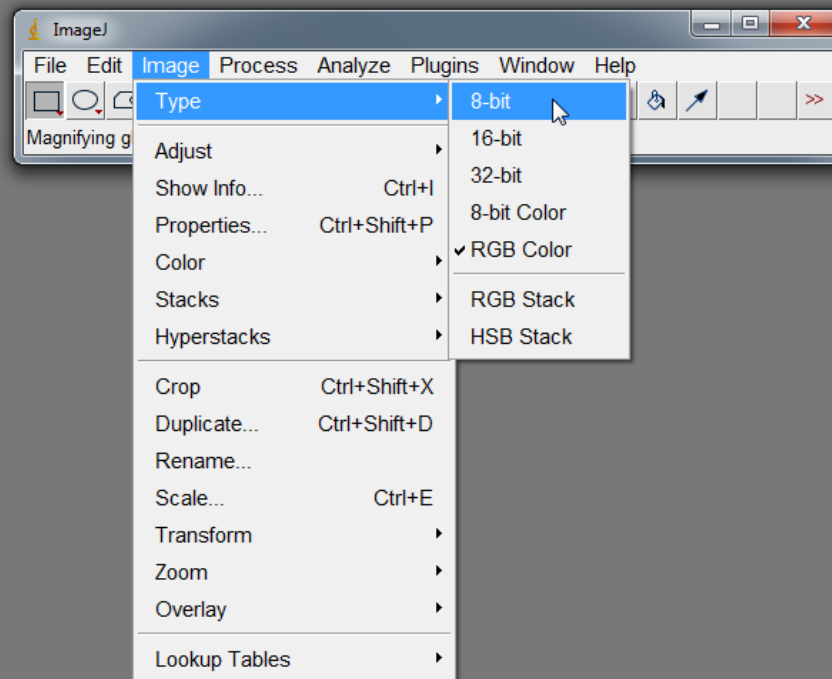
Viewing image metadata (.tiffs only)

View image metadata by selecting **Image>Show Info...** Metadata contains information such as scaling, pixel dimensions, bit depth, fluorescence wavelength etc.

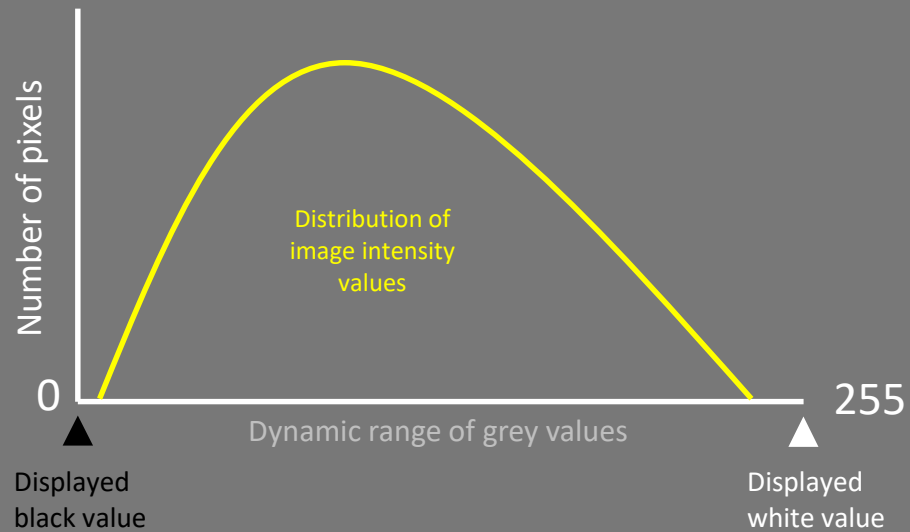


Some image analysis functions need to use 8-bit image formats (256 grey values)

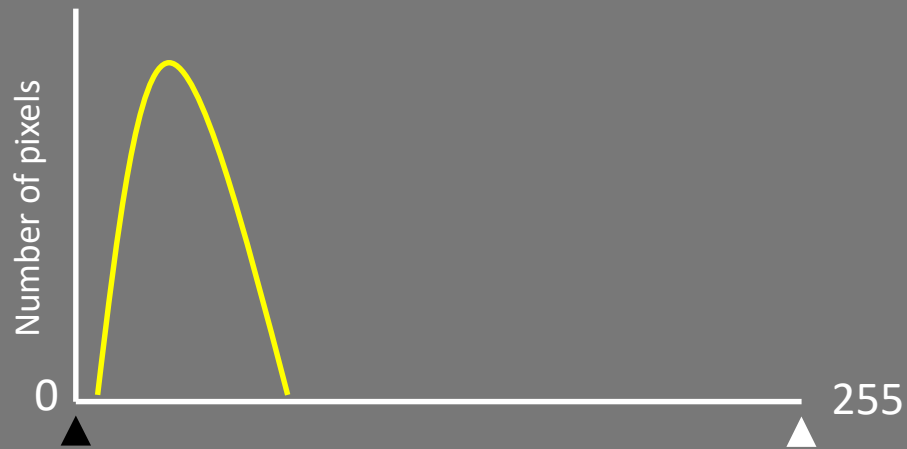
Change the bit depth using [Image>Type](#).



The image histogram (8-bit image example)



Maximum pixel value is 80 out of a possible 255. The image is too dark. This might have been necessary to retain a short exposure time or reduce photobleaching/phototoxicity of the specimen.



Intensity scaling

When the white value is moved all of the grey values between black and white are re-scaled so the image appears brighter, however actual pixel values remain unchanged.

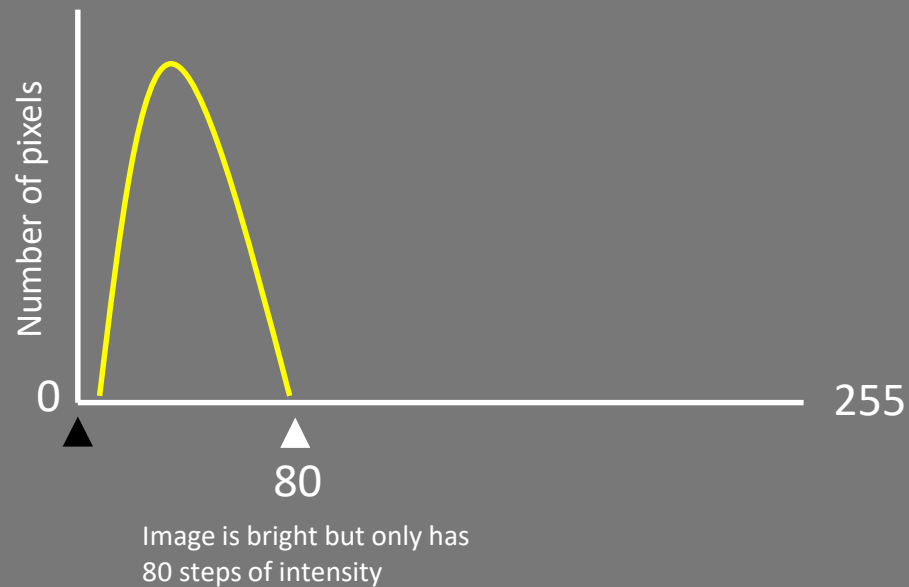


Image has been saturated during acquisition. Contrast of all pixels over 255 is permanently lost

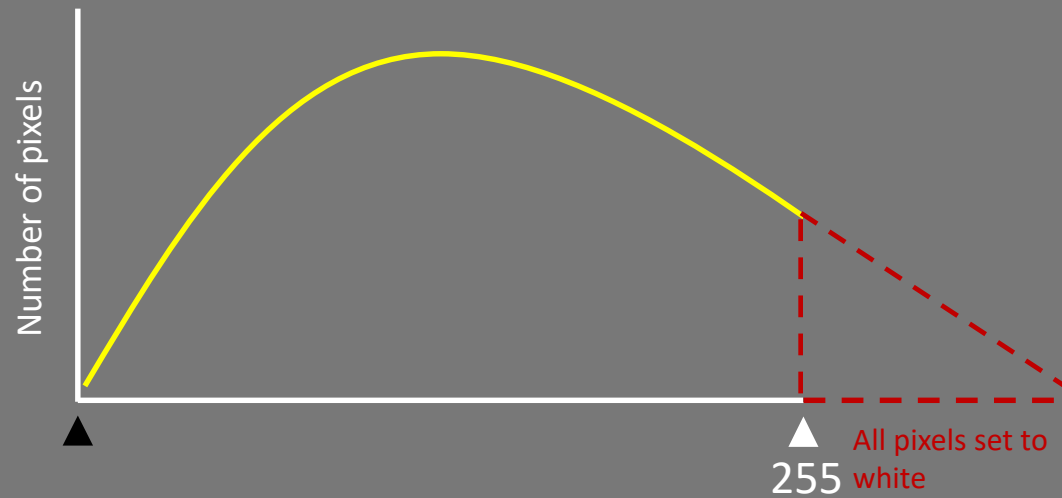
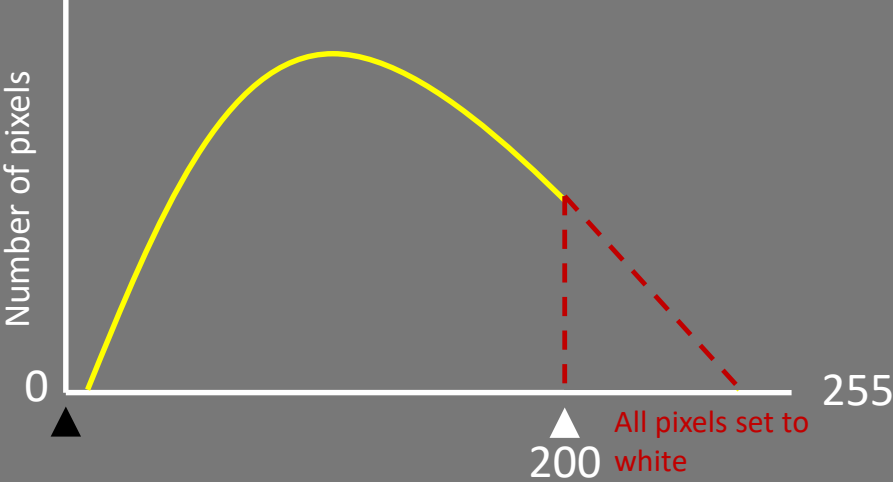
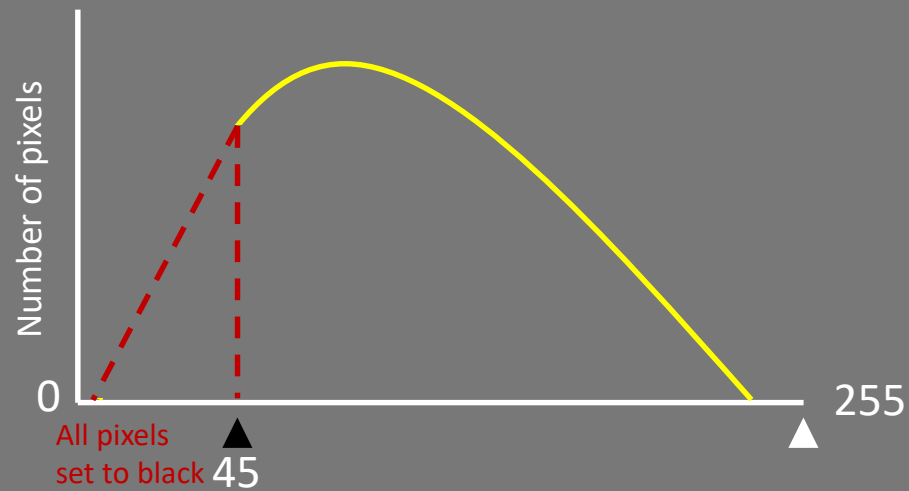


Image saturated during processing by incorrect placement of white value



Thresholding

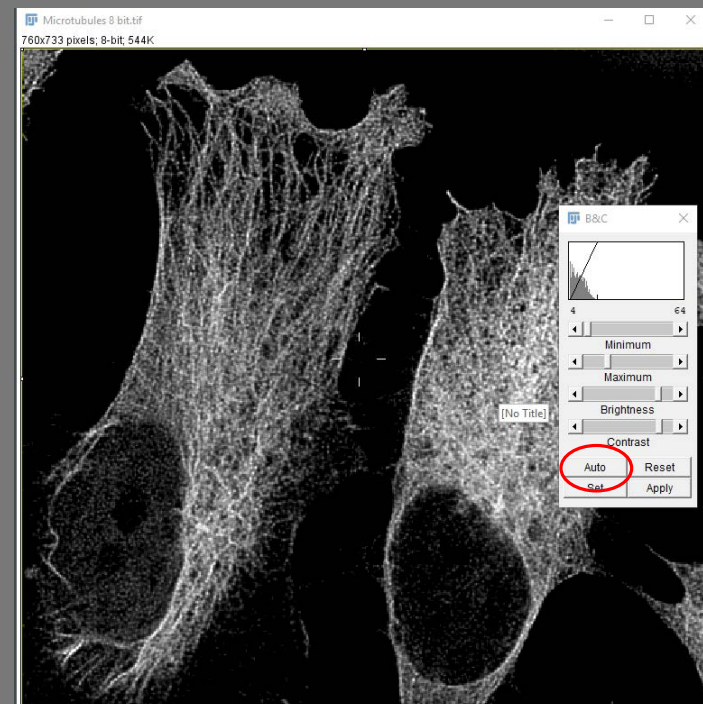
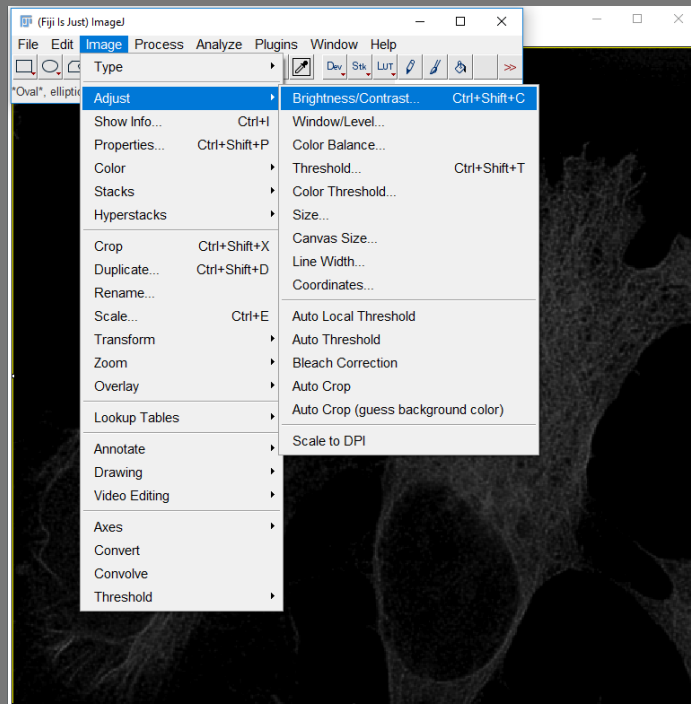
Removes background (and low intensity image information) noise or tell ImageJ which intensities to send to black and which to white when making a binary image.



Scaling image brightness automatically

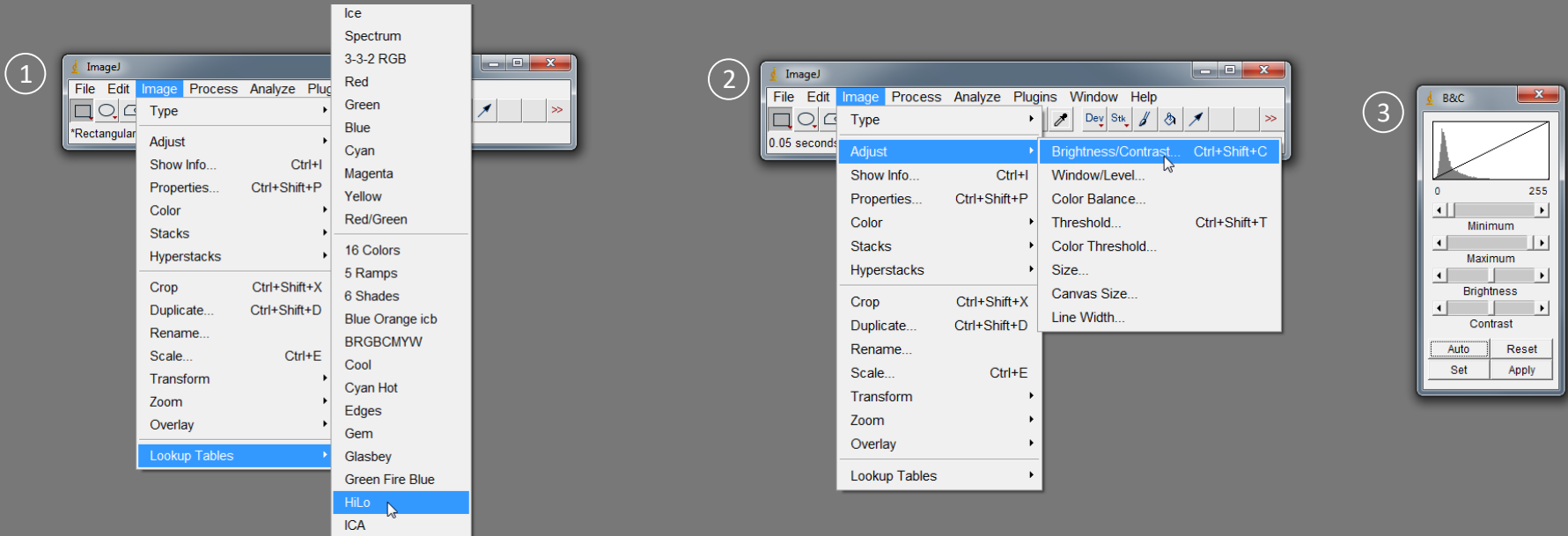
Open image “**Microtubules 8-bit**”. This image does not use the whole dynamic range. **Image>Adjust>Brightness/Contrast**, Select “**Auto**”. Don’t adjust sliders.

This function moves the displayed white value to the point where 0.4% of pixels are saturated. All the grey values are then re-scaled and the image appears brighter . As you can see from the histogram, actual pixel values remain unchanged and intensity can still be measured if desired.



Scaling image brightness manually

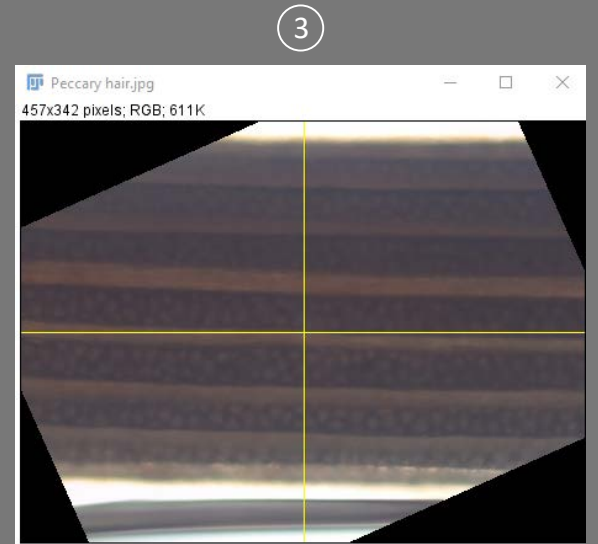
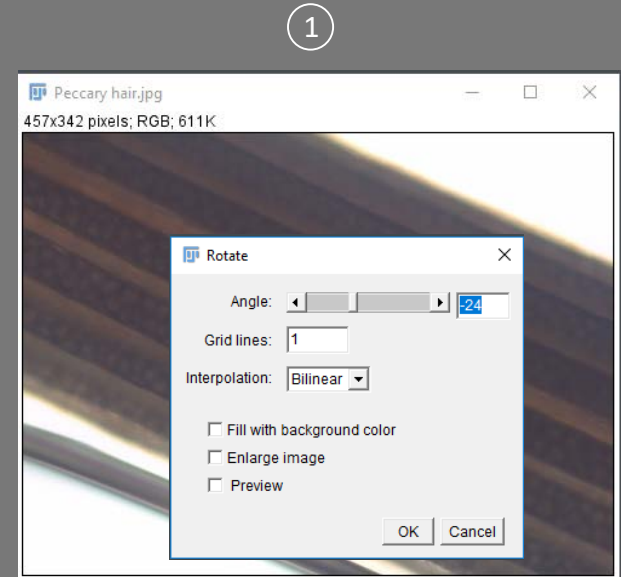
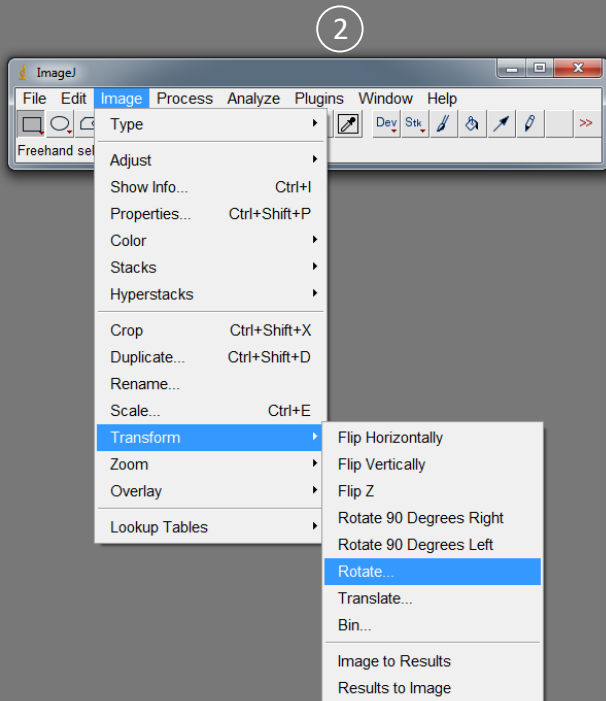
Open “**Microtubules 8-bit**” again. This method also does not change actual pixel values.



- 1 Select **Image>Lookup Tables>HiLo** and then **Image>Adjust>Brightness/Contrast**. Saturated pixels (value of 255) now appear red and pixels with a value of 0 appear blue.
- 2 Adjust the **Maximum** slider until you get few red pixels and then back it off until they just disappear.
- 3 Adjust the **“Minimum”** slider until the background turns blue. Click **“Apply”** then select **Image>Lookup Tables>Grays**.

Rotating images

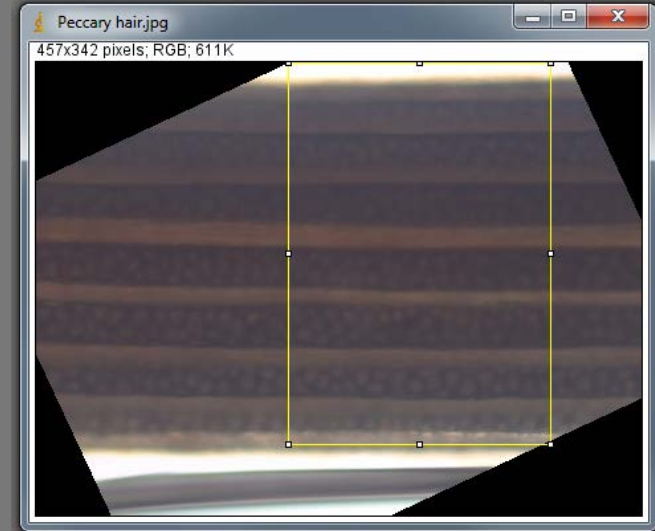
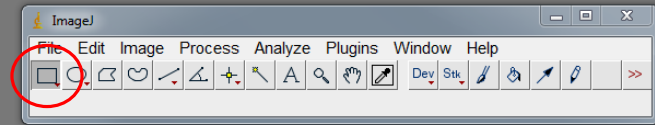
- 1 Open image “Peccary hair”
- 2 Select Image>Transform>Rotate....
- 3 Tick “Preview”. Adjust Angle slider until to achieve the desired rotation. Click OK.



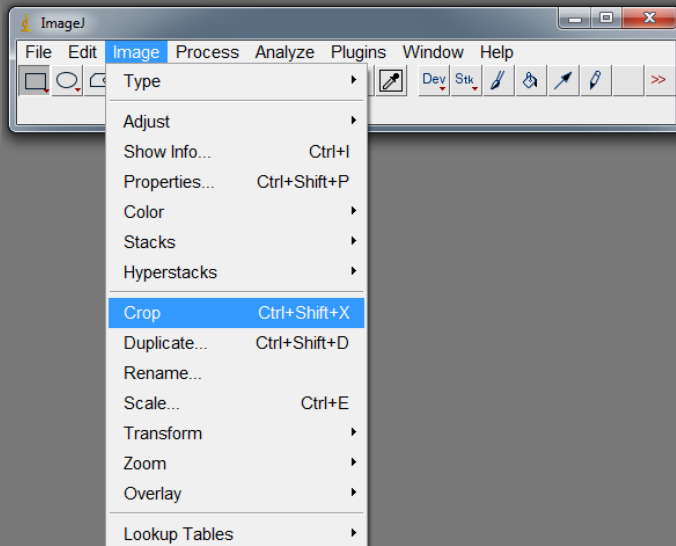
Cropping

- 1 With the image you just rotated, using the rectangular selection icon, drag a selection around the area that you want to keep.
- 2 Select **Image>Crop**
For consistency of size, crop regions can be stored in the ROI manager.
Analyze>Tools>ROI Manager...

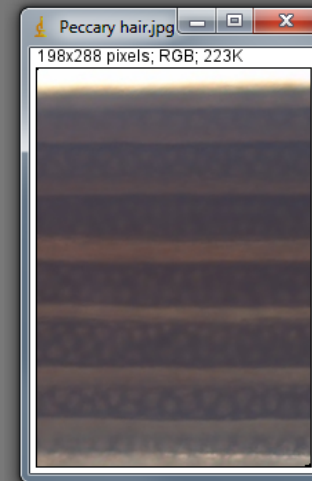
1



2



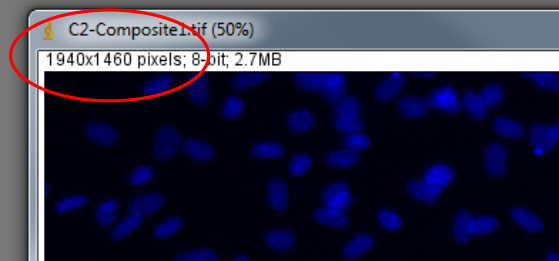
3



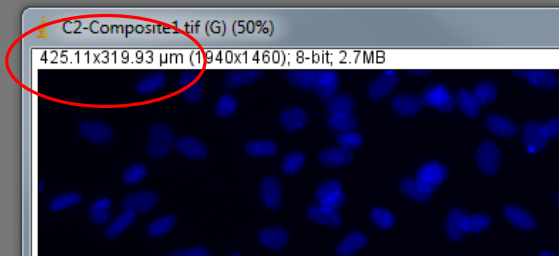
Calibrating images

Before you can add a scale bar or analyse images, the images have to be calibrated to the correct measurement units.

Many instruments automatically add spatial calibrations to the image metadata during acquisition. To check if your images are calibrated look in the top left hand corner of the image. If your image dimensions are given in pixel units the image has not been calibrated.



Uncalibrated

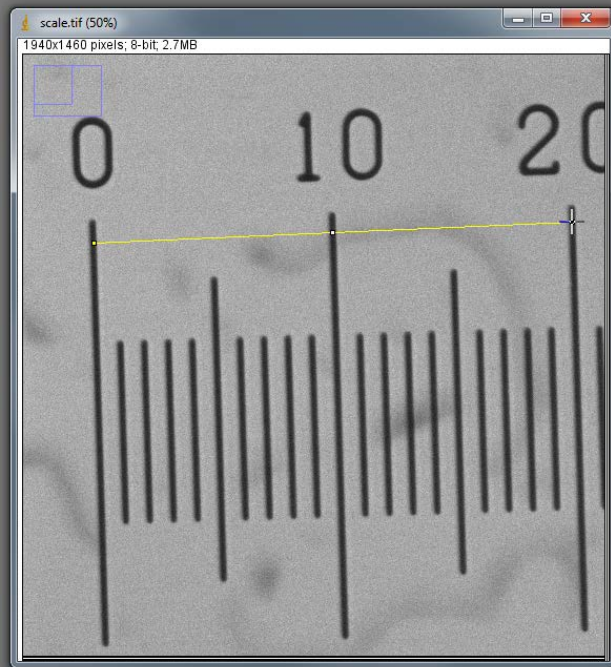
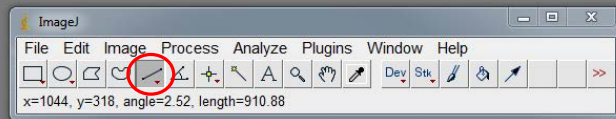


Calibrated

If your image is not automatically calibrated by the acquisition software, an image of a stage micrometer taken at the same magnification as your specimen can be used to calibrate your images.

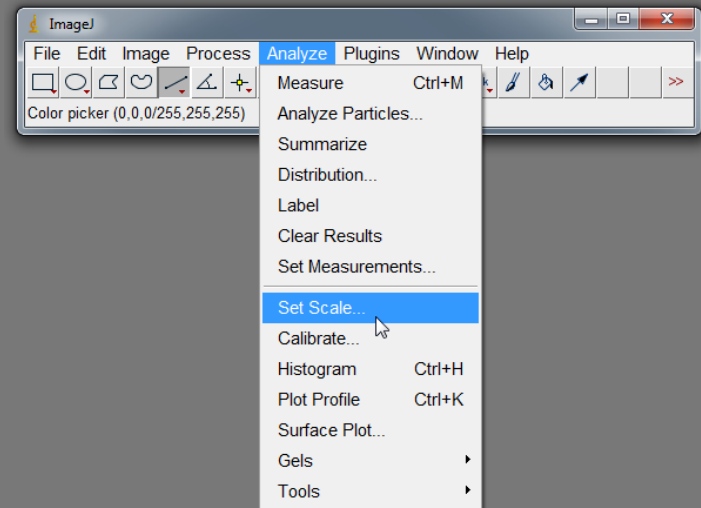
Manually calibrating images

- 1 Open images “calibration image” and “Uncalibrated image”. Using the line tool, draw a line of a known distance on the image of the stage micrometer.

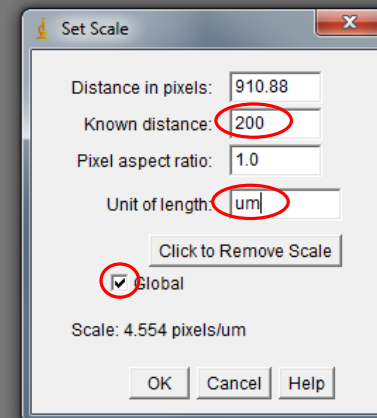


(1 small increment on the micrometer = 10 μm)

- 2 Select Analyze>Set Scale.



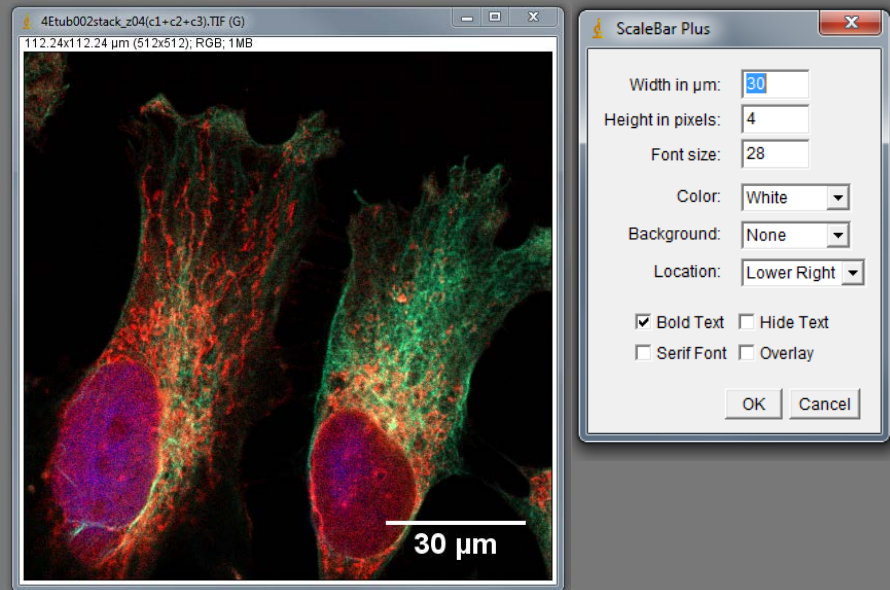
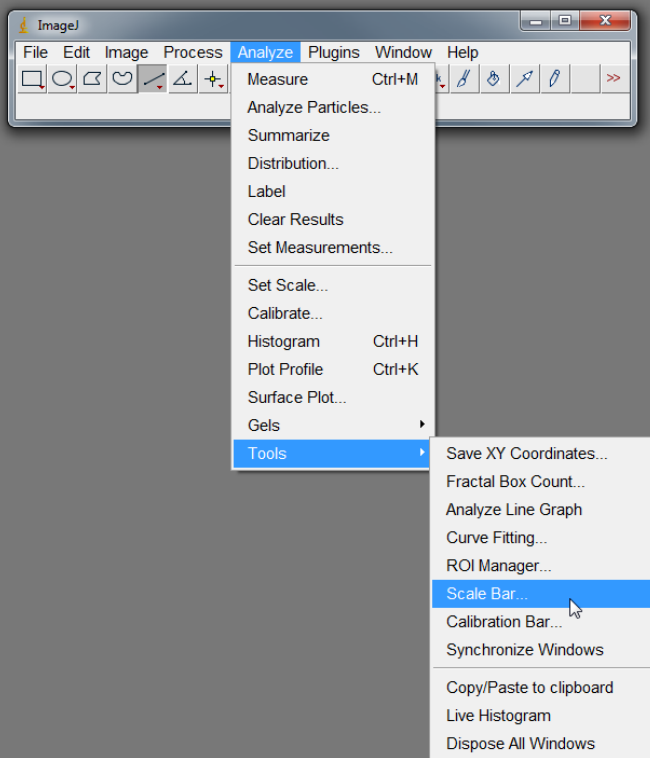
- 3 Enter the known distance and units in μm . Ticking “Global” applies the calibration to all images in the Imagej session.



Adding a scale bar

With your calibrated image open select **Analyze>Tools>Scale Bar**.

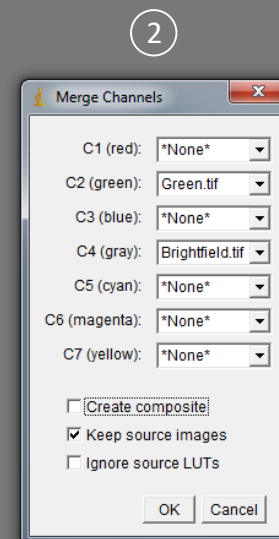
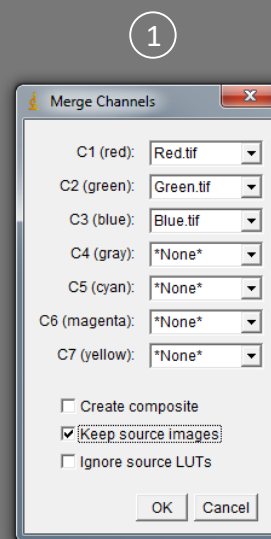
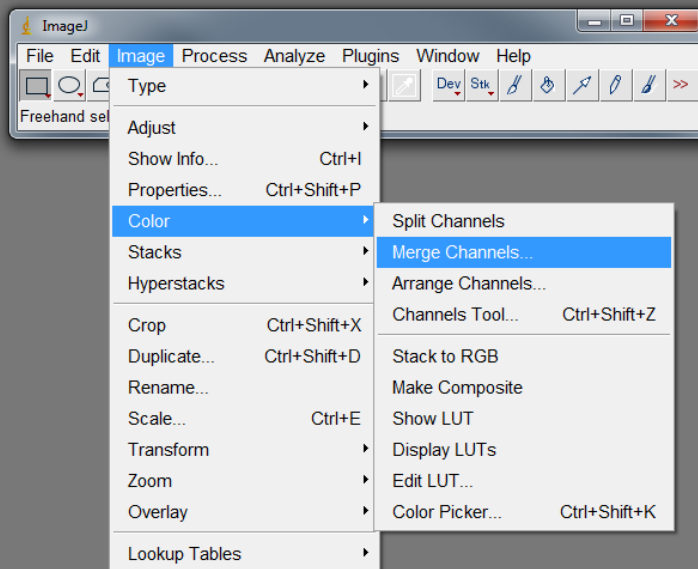
Select the required width and position and click “OK”.



Merging images into multichannel 24-bit RGB .tifs

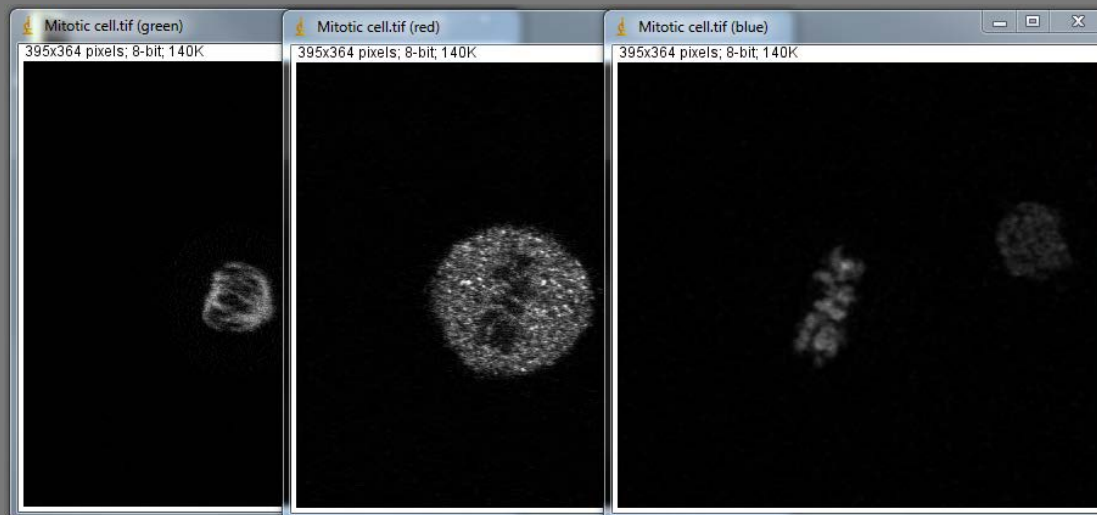
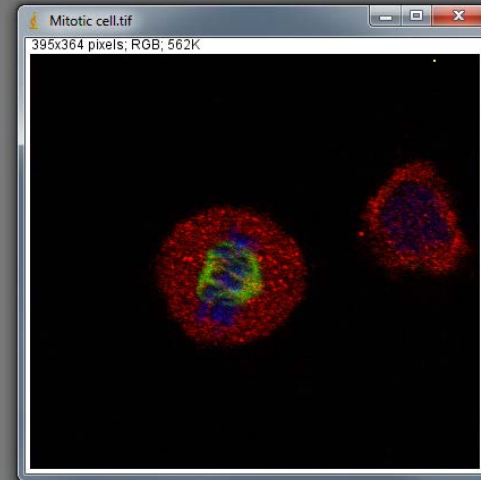
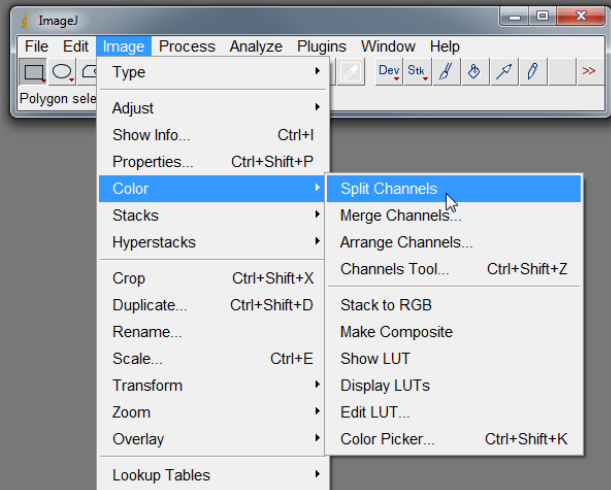
Open images “Blue.tif, Green.tif, Red.tif” and “Brightfield.tif”.

- 1 Select **Image>Colour>Merge Channels**
Select each image into their corresponding colour channels.
Untick “Create composite”, tick “Keep source images” and click “OK”.
- 2 Select **Image>Colour>Merge Channels**
Select **Green.tif** in the green channel and **Brightfield.tif** in the grey channel
Untick “Create composite”, tick “Keep source images” and click “OK”.



Splitting multichannel RGB images

Open image “Mitotic cell”.
Select Image>Colour>Split Channels.

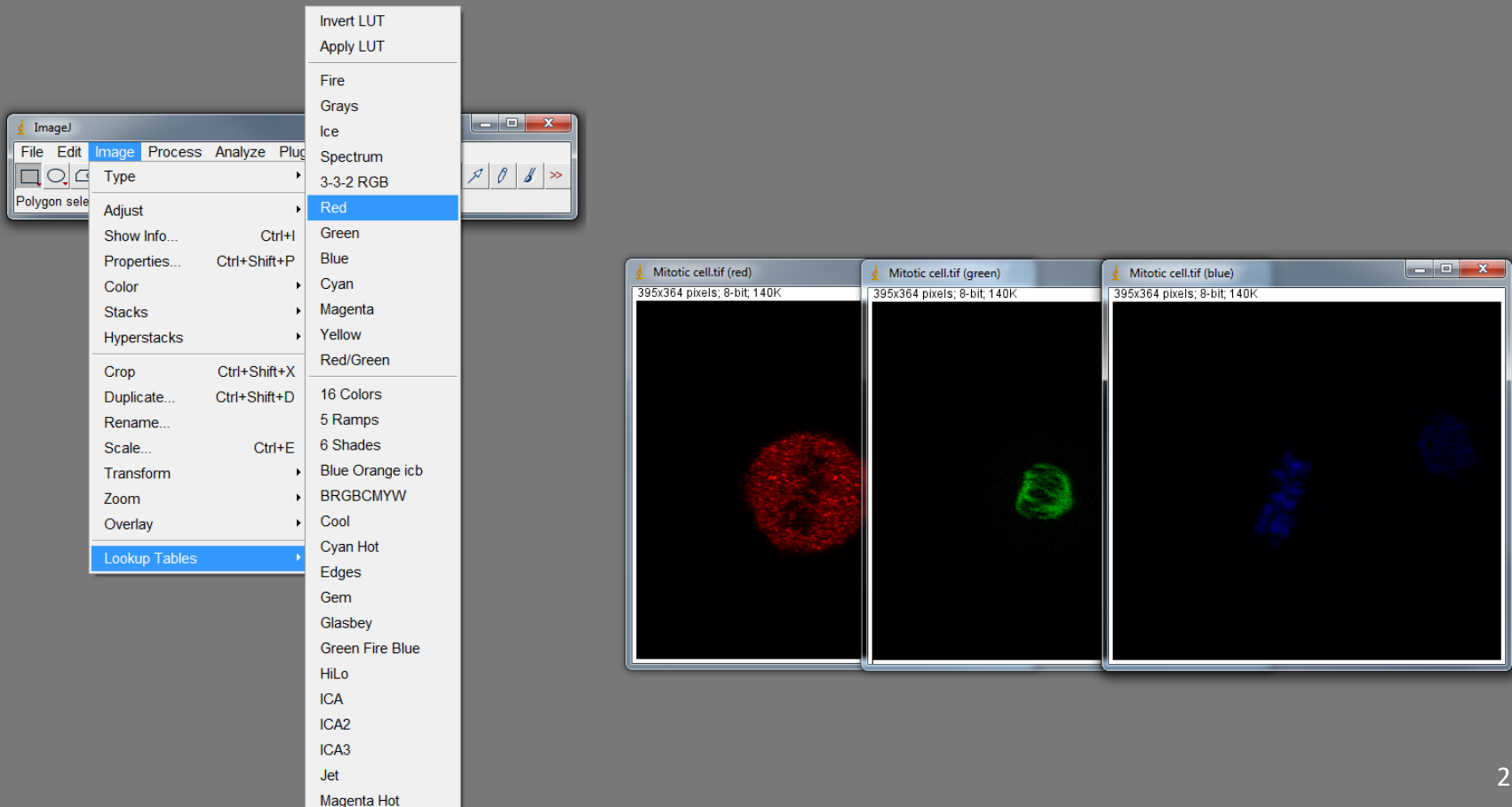


Applying pseudocolour to grey images

Using your open grey images split from “Mitotic cell.tif”, Select a channel image by clicking on it then select **Image>Lookup Tables**.

Select the correct corresponding colour from the drop down menu.

To return a channel to grey, select “Grays” from the drop down menu.



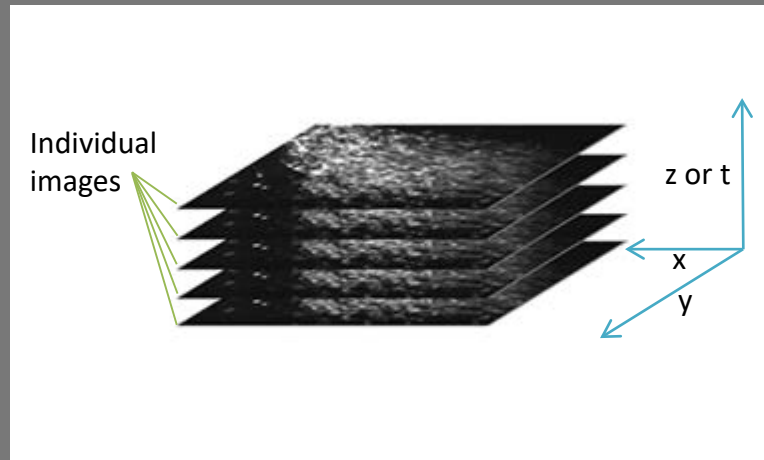
Stacks

Stacks are a method of handling multiple related images in one file.

They are often used to handle multiple slices through the vertical z-axis of a specimen (a z-stack), but are also used to handle sequential images in a time lapse experiment (a t-stack) or images acquired at different wavelengths (a λ -stack).

Stacks can have up to three dimensions e.g. $[x,y],[channels],[z \text{ or } t \text{ or } \lambda]$.

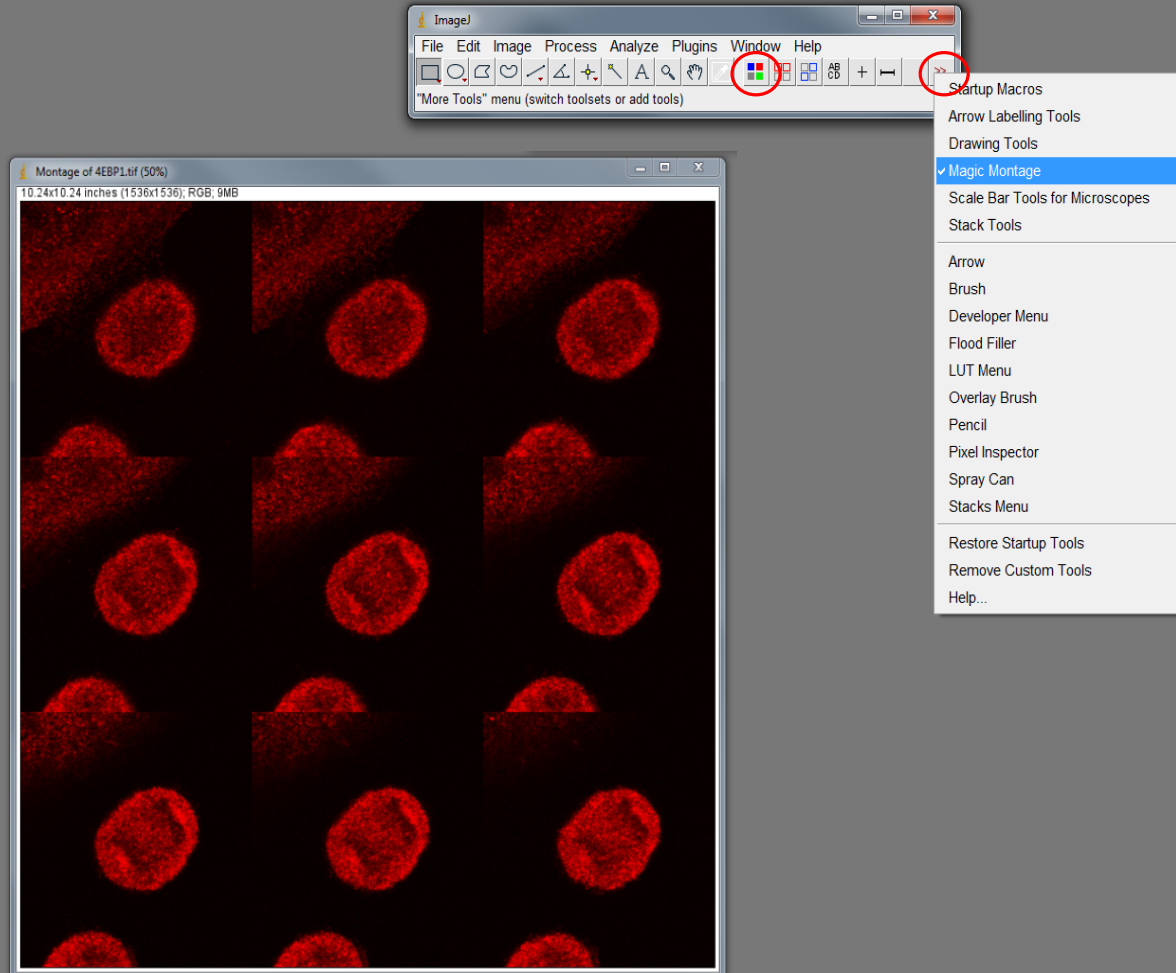
Stacks with more than three dimensions e.g. $\{[x,y],[channels],[z]],[t]\}$ are called hyperstacks.



View image stacks as a montage

Open z-stack 4EBP1.tif

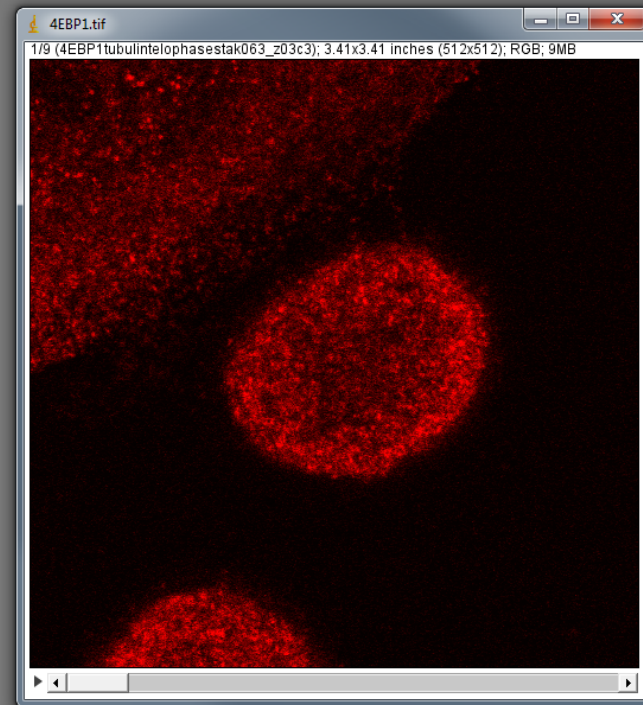
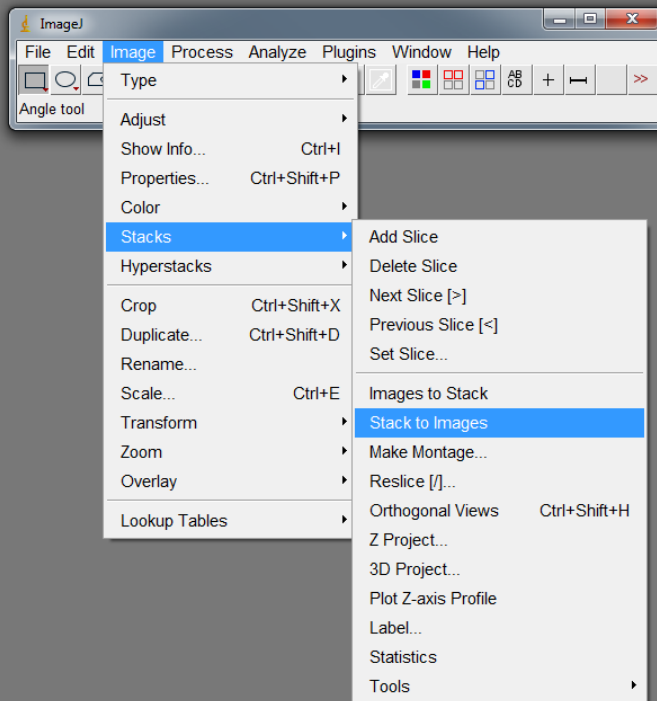
Select Image>Stacks>Make Montage



Convert stacks to single images

Open z-stack 4EBP1.tif

Select Image>Stacks>Stack to Images

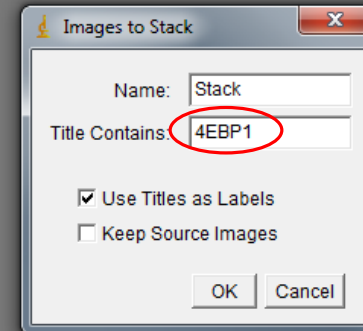
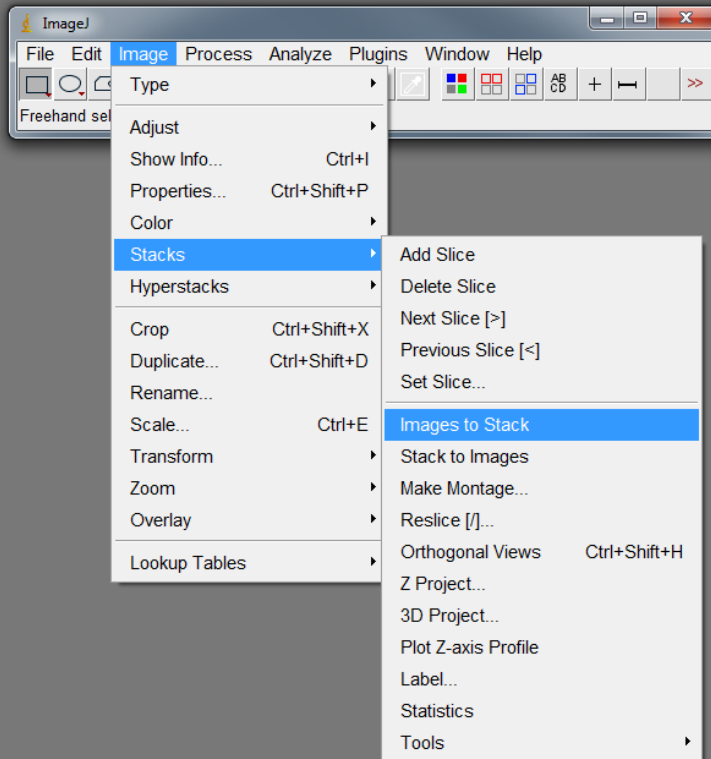


Convert single images to stacks

Using your single images split from z-stack 4EBP1.tif

Select Image>Stacks>Images to Stack

In the pop up window, type something that is in the title of all of the images.

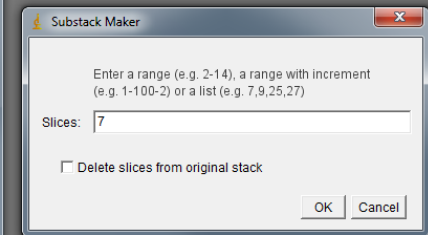
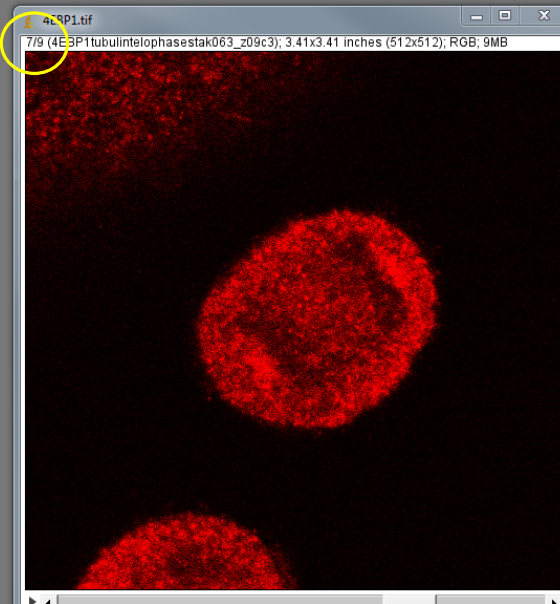
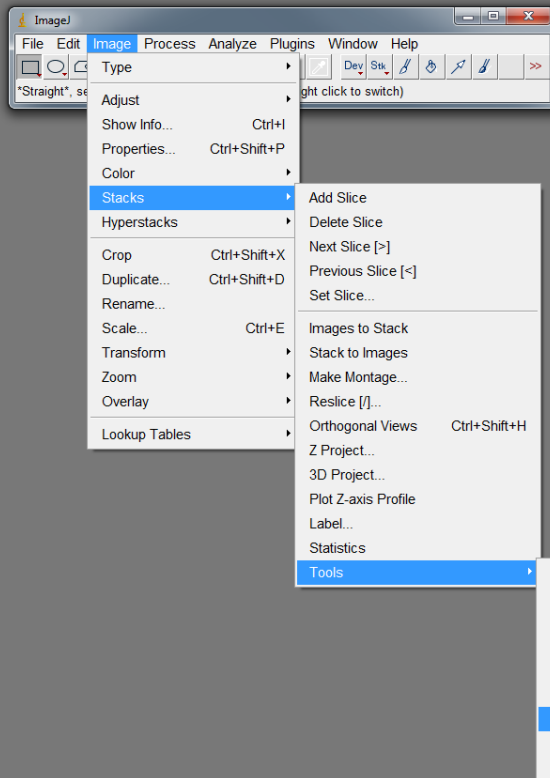


Make a substack or a single slice from a stack into a separate image

Displayed Slice number is shown in the top left hand corner of the stack

Select **Image>Stacks>Tools>Make Substack**

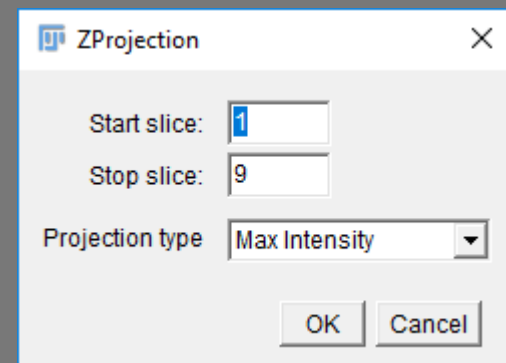
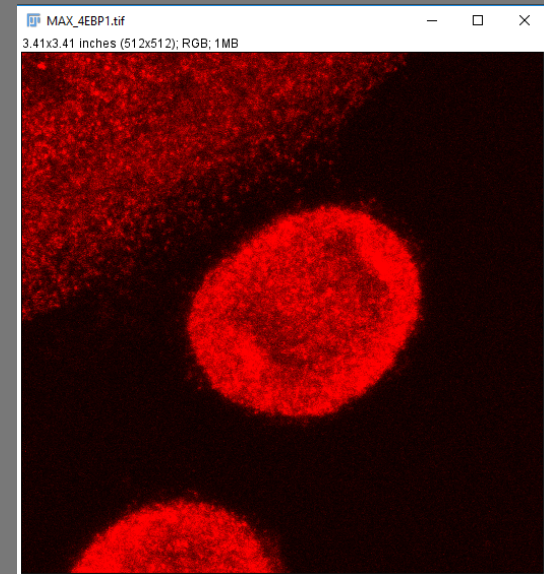
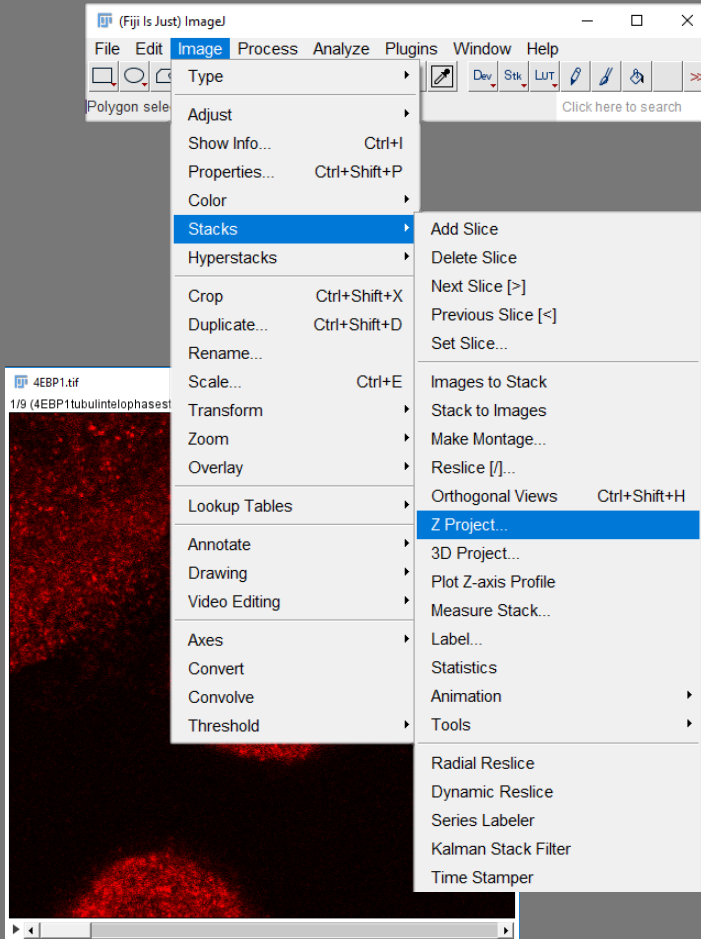
Enter slice number for a single image, or range if you want to make a substack.



Project a stack to a 2D image

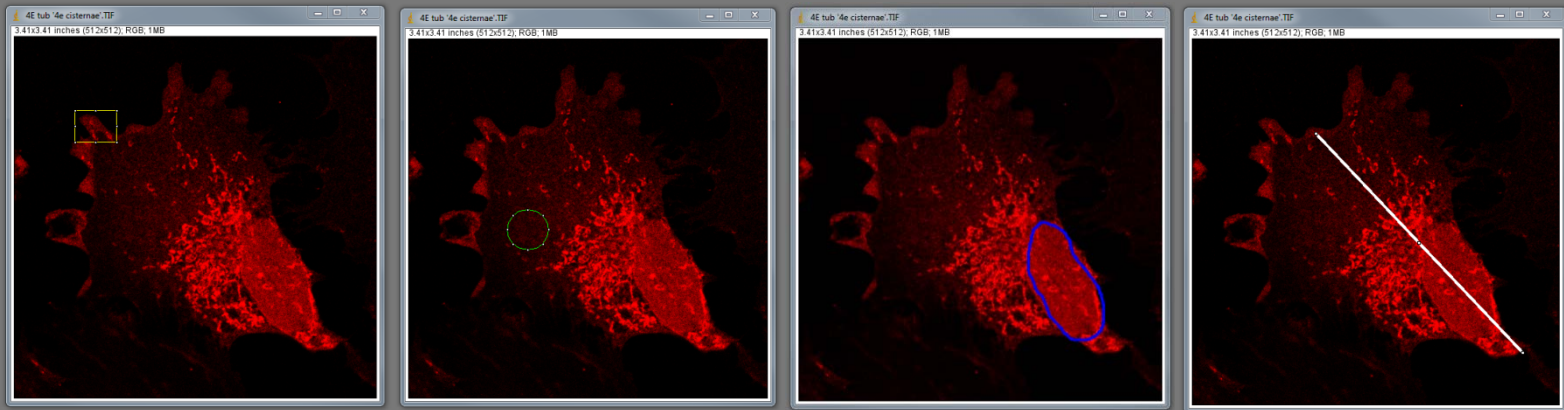
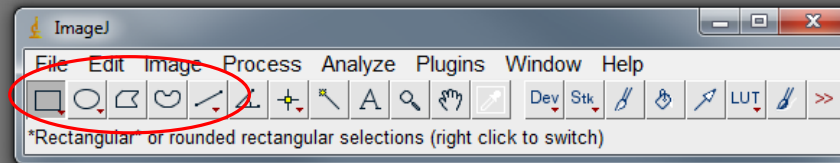
Select **Image>Stacks>Z Project....**

Select start and finish slices that you want to project and Projection type “Max intensity”



Basic selections and measurements

Open image 4E Cisternae.tif

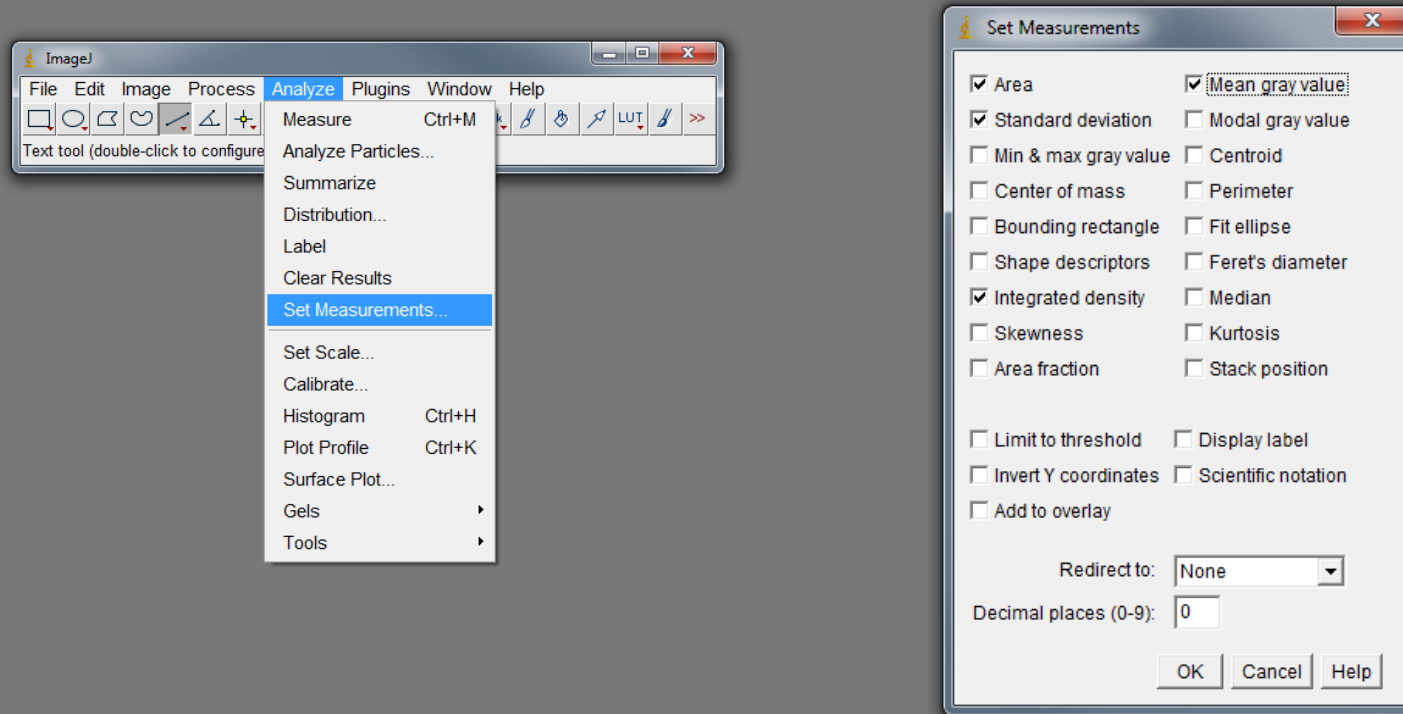


Select regions of interest (ROIs) for measurement using the Rectangular, Polygon, Oval, Freehand and Line selection tools.

When the cursor is a cross you can click and drag to make your ROI. Placing your cursor in the middle of your ROI (it changes to an arrow) allows you to move your ROI.

Basic selections and measurements

To set the type of measurement to be made with the ROI, select **Analyse>Set Measurements...** and tick the relevant boxes then click “OK”.



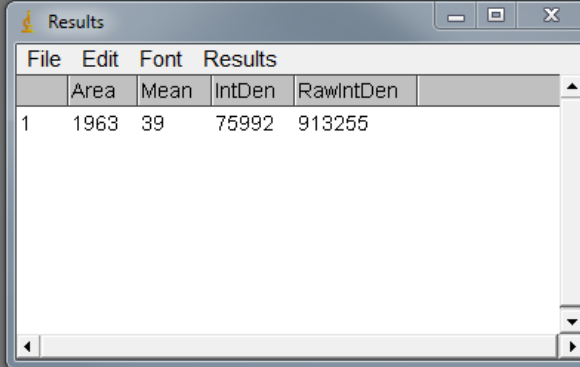
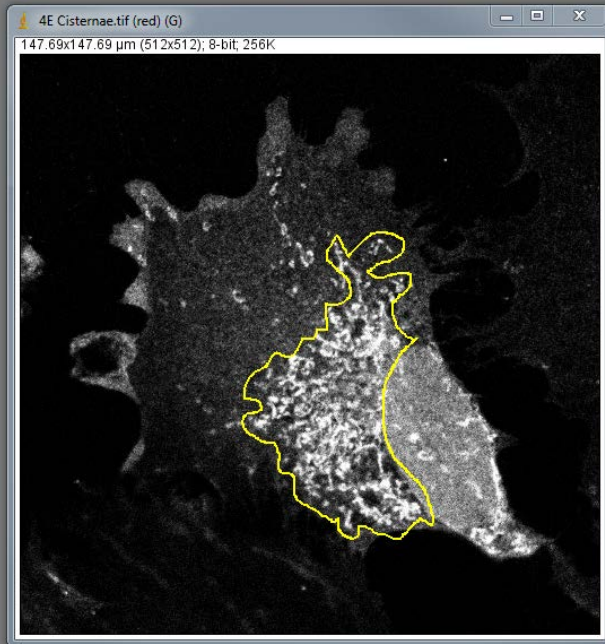
Area returns the area within the ROI in the calibrated units² (or pixels² if the image is uncalibrated).

Mean gray value returns the average grey value inside the ROI.

Integrated density is the average grey value * area inside the ROI.

Basic selections and measurements

Draw an ROI on the image using one of the selection tools.

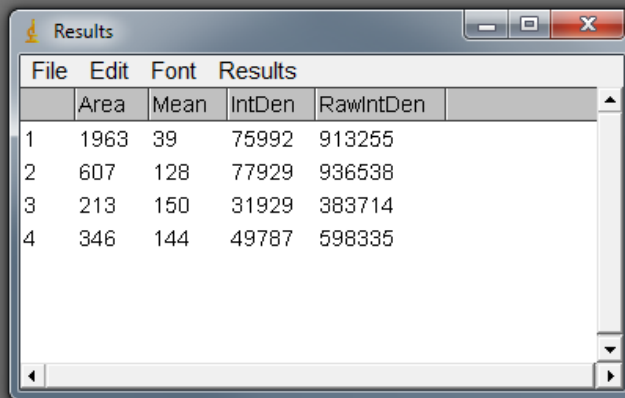


A "Results" window displaying measurement data for the ROI. The window has a menu bar with "File", "Edit", "Font", and "Results". The data is presented in a table with the following columns: "Area", "Mean", "IntDen", and "RawIntDen".

	Area	Mean	IntDen	RawIntDen
1	1963	39	75992	913255

Press [Ctrl+M] to make the measurement

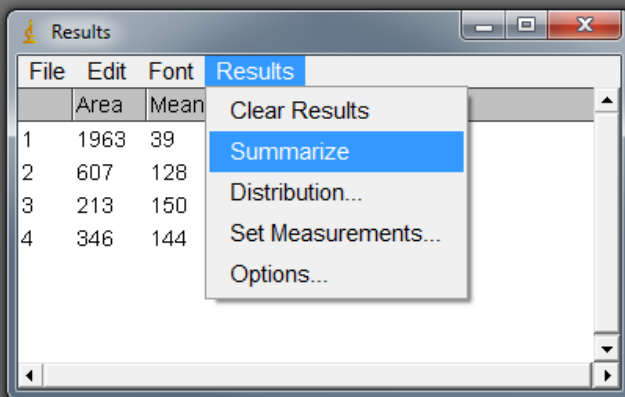
Basic selections and measurements



Results

	Area	Mean	IntDen	RawIntDen
1	1963	39	75992	913255
2	607	128	77929	936538
3	213	150	31929	383714
4	346	144	49787	598335

Multiple measurements can be made (select a new ROI and press **[Ctrl+M]** again).

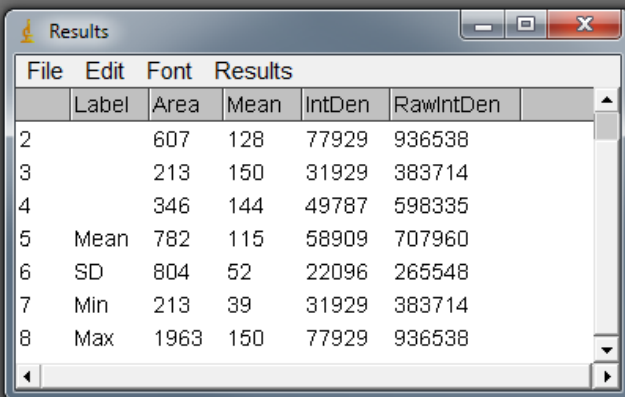


Results

	Area	Mean
1	1963	39
2	607	128
3	213	150
4	346	144

- Clear Results
- Summarize
- Distribution...
- Set Measurements...
- Options...

You can calculate the mean and standard deviation of the whole dataset by selecting **Results>Summarize** from the Results window.

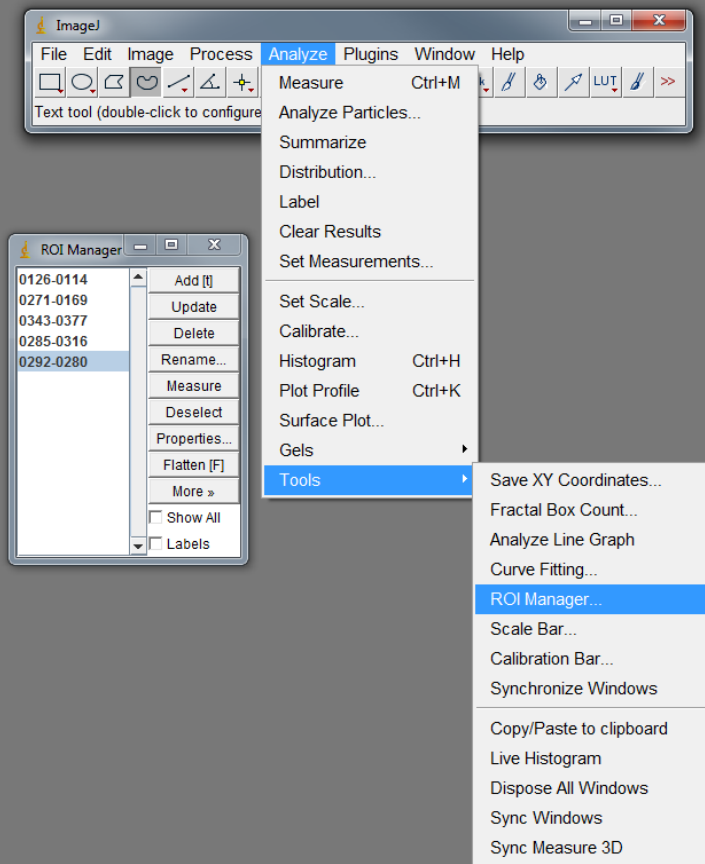


Results

	Label	Area	Mean	IntDen	RawIntDen
2		607	128	77929	936538
3		213	150	31929	383714
4		346	144	49787	598335
5	Mean	782	115	58909	707960
6	SD	804	52	22096	265548
7	Min	213	39	31929	383714
8	Max	1963	150	77929	936538

Results can also be copied and pasted into Excel for further analysis.

The ROI manager



Multiple ROIs can be stored and recalled using the ROI manager.

Select **Analyze>Tools>ROI Manager...**

To add the current ROI to the ROI manager click **"Add"**.

To recall the ROI into any image select it from the list on the left.

Select **More>>Save...** On the ROI manager window to save ROIs for later use.

To change the colour and weight of the ROI click **"Properties"** on the ROI manager window.

Background correction

Ideally the grey value for the image background should be as close to 0 as possible with similar values across the whole image background.

Poor quality or incorrectly configured illumination may cause an uneven background.

Some microscope systems allow grey value “offset” during acquisition, however this can also be achieved post-acquisition if necessary using several methods:

Rolling Ball – Corrects uneven illumination and preserves low intensity fine specimen detail.

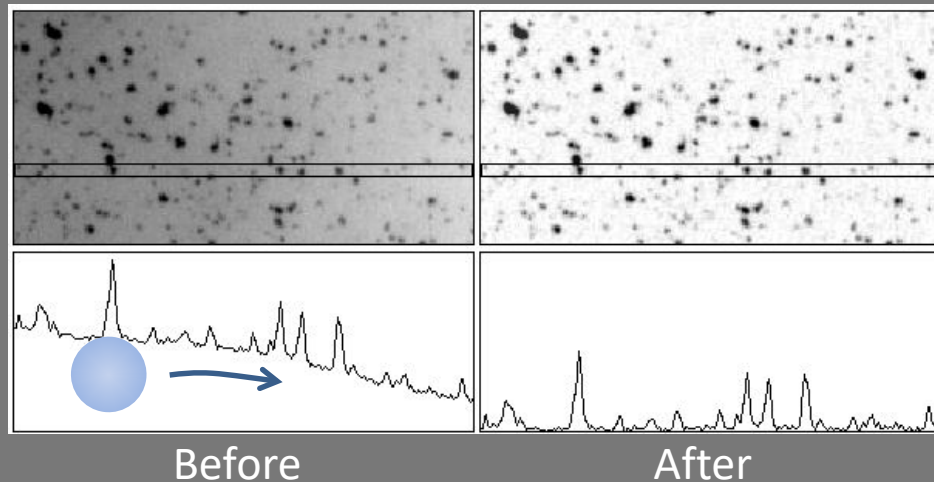
Thresholding – Subjective and removes low intensity image information as well as the background.

Background subtraction – Removes low intensity image information as well as the background. Based on average background intensity, better when quantification is required.

Background correction - Rolling Ball

This method imagines a ball or paraboloid rolled around the image. Areas of grey value large enough for the “ball” to touch to are removed.

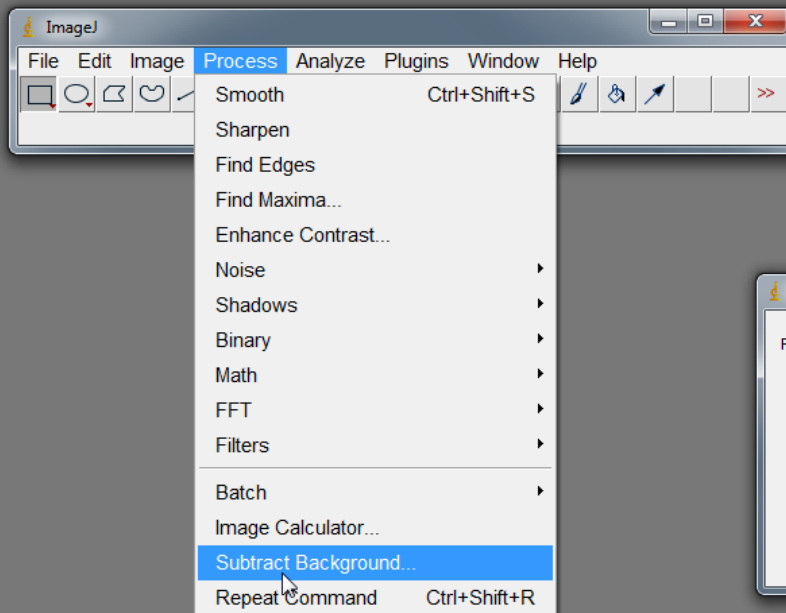
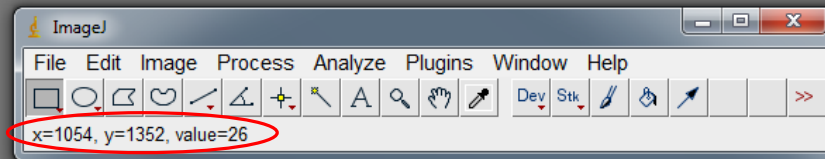
This has the advantage over simple background subtraction of preserving low intensity fine detail in the image and correcting uneven illumination.



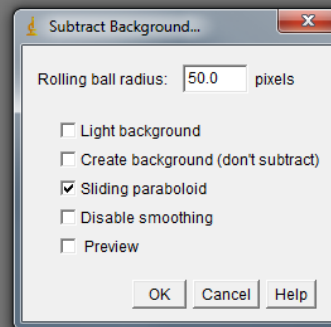
Background correction - Rolling Ball

Open image “**Background correction example**”. This image has a background problem as well as uneven illumination, and has a large circular artefact from the dish that the cells were imaged in.

If you hover the cursor over the image, the grey value of the pixel under the cursor is displayed on the menu bar.

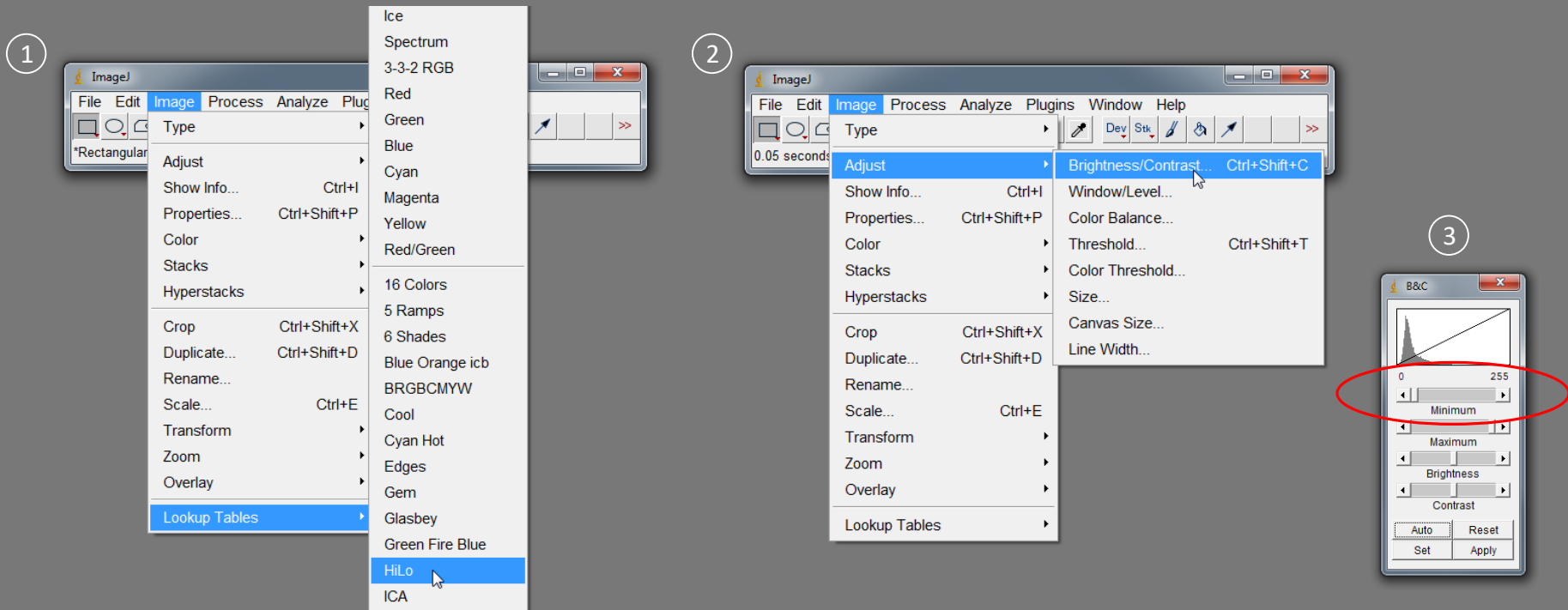


Select **Process>Subtract Background**. Tick “**Sliding paraboloid**” and choose a “**Rolling ball radius**” of 50 pixels and click “**OK**”.



Background correction - Thresholding

Close your background corrected image without saving and open image “Background correction example” again.

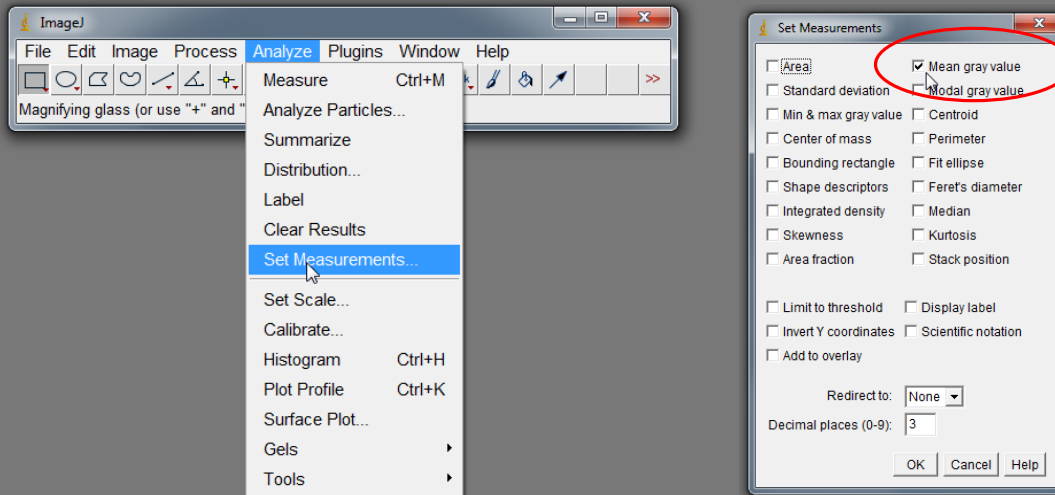


Select **Image>Lookup Tables>HiLo** and then **Image>Adjust>Brightness/Contrast**. Saturated pixels (value of 255) now appear red and pixels with a value of 0 appear blue. Adjust the “**Minimum**” slider until the entire background is blue. All pixels with a value below the selected threshold (including image information!) will be set to 0. Click “**Apply**” then Select **Image>Lookup Tables>Grays**.

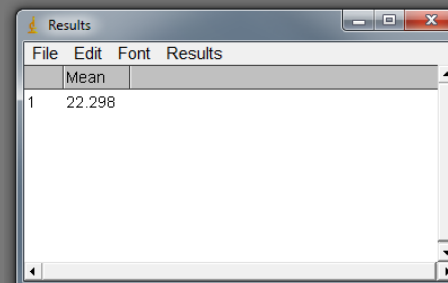
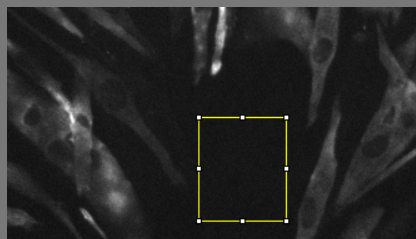
Background correction - Subtraction

Close your background corrected image without saving and open image “Background correction example” again.

Select **Analyze>Set Measurements** and then tick the “Mean gray value” box and click “OK”.

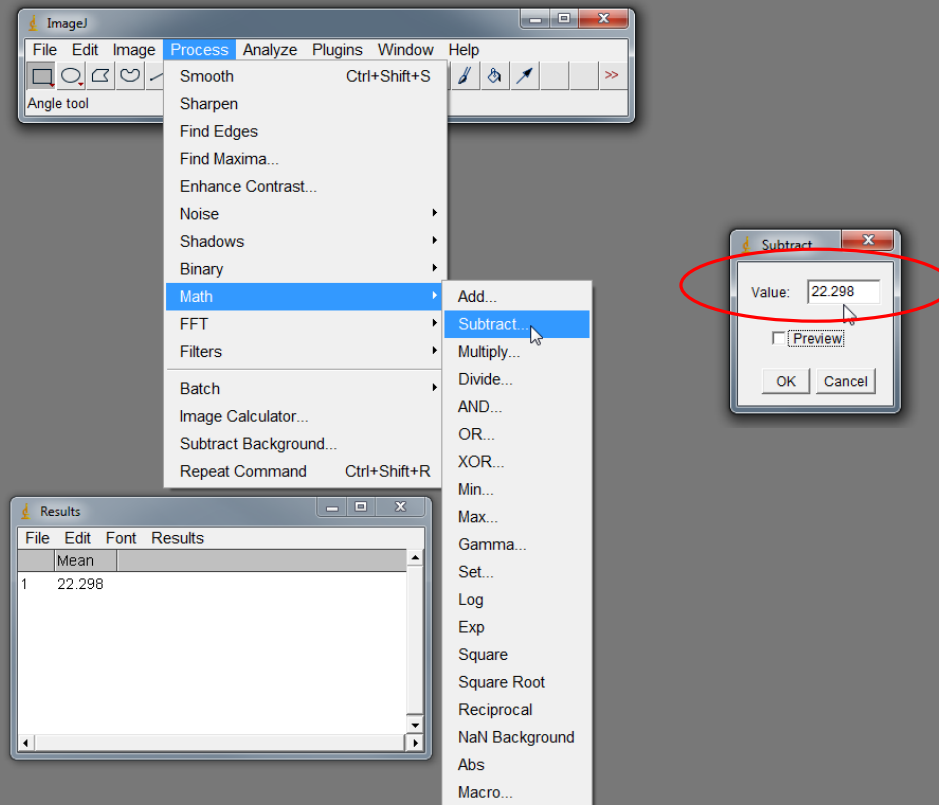


Select an ROI on an area of background and select **Analyze>Measure** or use [Ctrl+M]. A results window should appear with the mean grey value of the pixels within your ROI.



Background correction - Subtraction

Deselect your ROI by clicking on the image or the subtraction will only be applied within the ROI. Select **Process>Math>Subtract** and enter your mean background value into the value field on the window that appears. Click “OK” or select “Preview”.



Removing noise

Image noise tends to manifest as speckle and can be caused by:

- Electronic variations in imaging detectors.
- Analogue to digital conversion during image acquisition.
- Variations in photon detection, particularly from low signal specimens: “shot noise”.
- High detector gain: “dark noise”.

Speckle can also be caused by contamination such as dust in the sample auto-fluorescing and by precipitates from stains, so its good practice to wash or flame coverslips and slides and centrifuge stains prior to use.

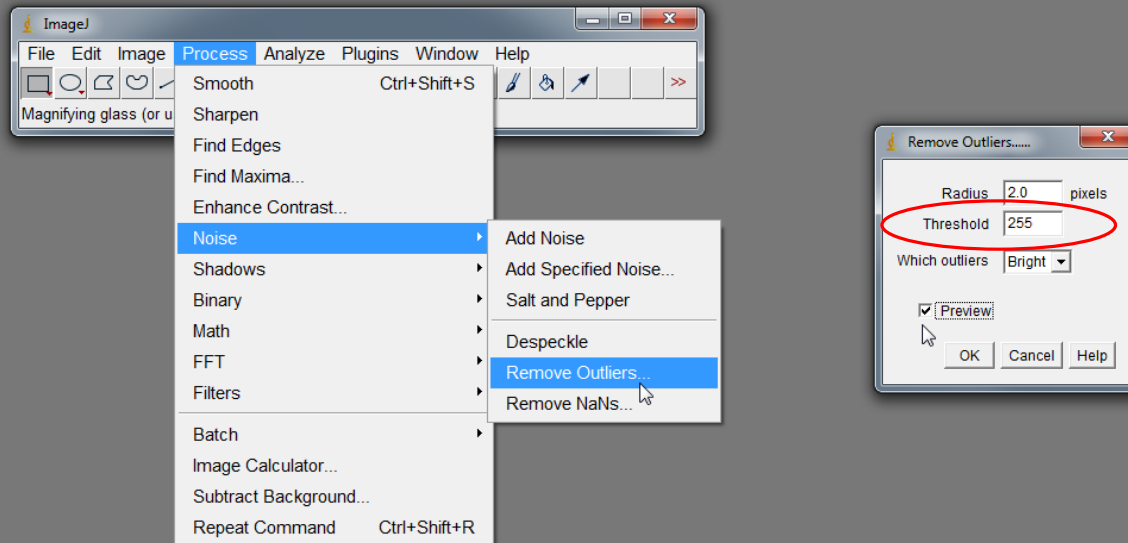
Care should be taken when doing noise corrections on images, which may not be suitable if quantitative measurements are to be made.

Removing noise - outliers

Open image “despeckle 3”. This image is suffering from speckle caused by high detector gain and also has some large speckles which are probably caused by stain precipitates reacting with the coverslip coating.

Remove the large bright speckles:

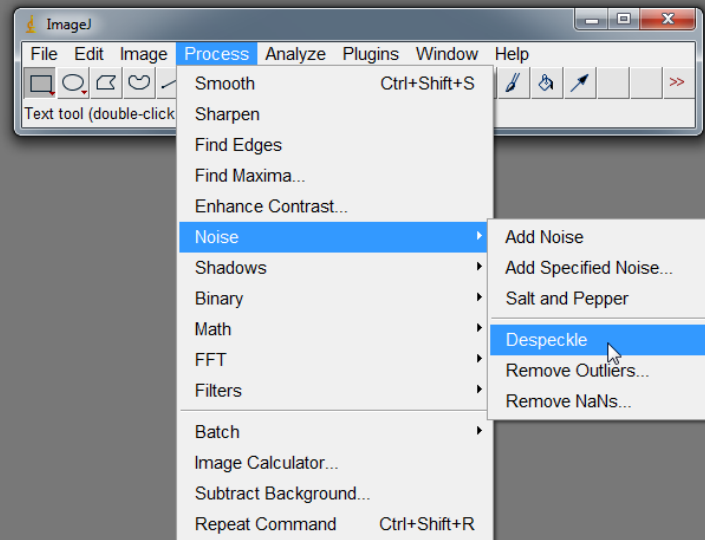
This function replaces a pixel with the median value of the surrounding pixel values if it deviates from the median by more than the thresholded value. Select “Preview” Reduce the threshold value until the bright outliers disappear (about 90 in this case). Click “OK” when finished



Removing noise - speckle

Remove the detector noise:

This function replaces each pixel with the median value of the surrounding 3x3 pixels. Select [Process>Noise>Despeckle](#).



Use [image>Adjust>Brightness/Contrast](#) and the minimum slider to background correct your image

Open the original image to compare the differences.

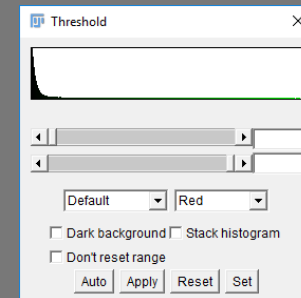
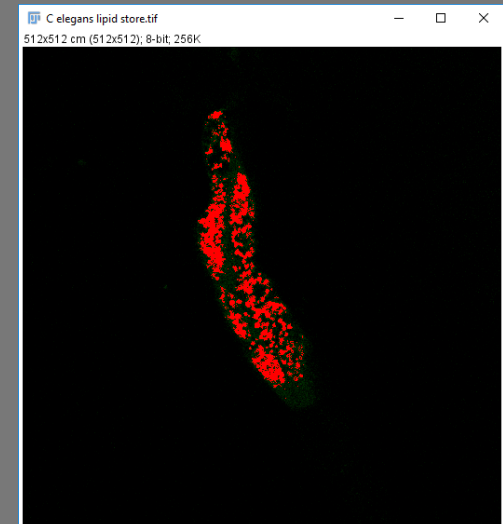
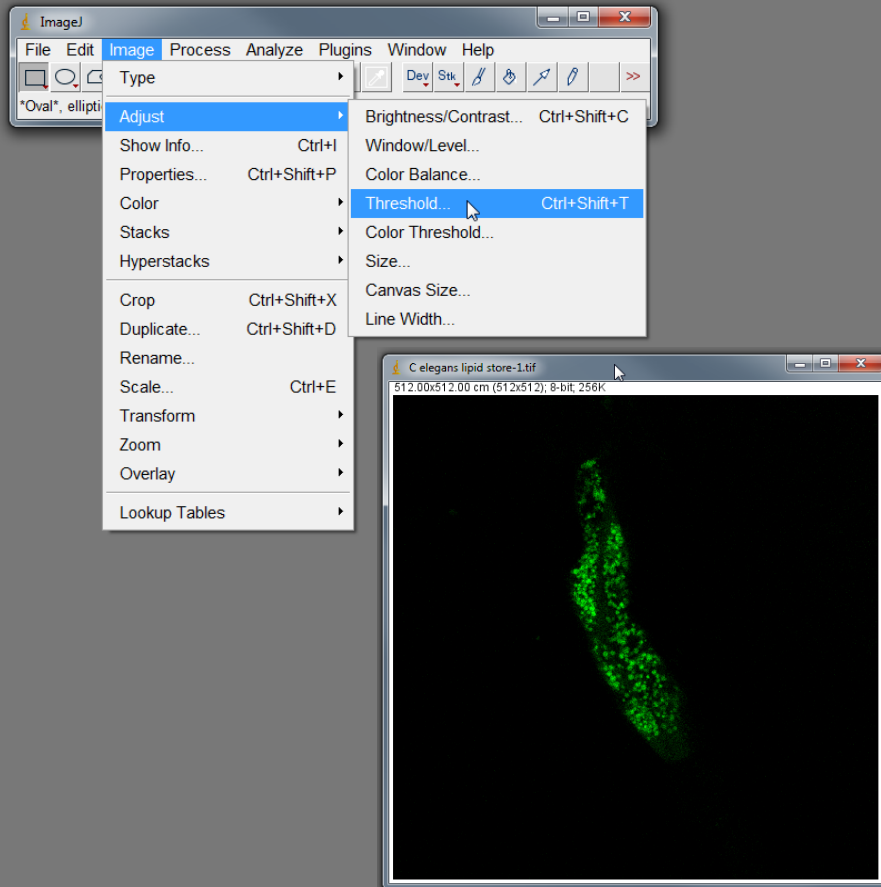
Making precise selections – binary selection method

Open image “C elegans lipid store”.

Set the second pulldown menu to “Red”.

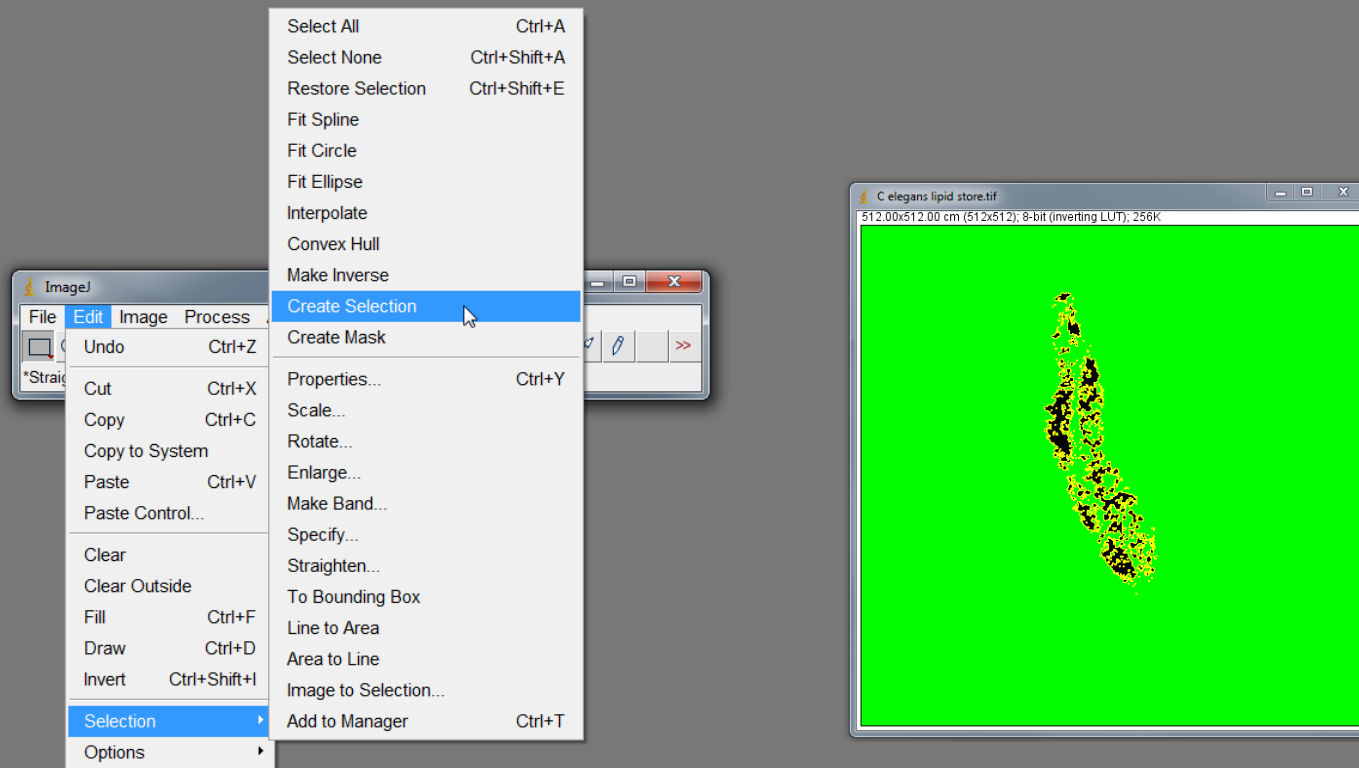
Select **Image>Adjust>Threshold** and threshold the image using the “minimum” slider.

Click “Apply” to make a binary image.



Making precise selections – binary selection method

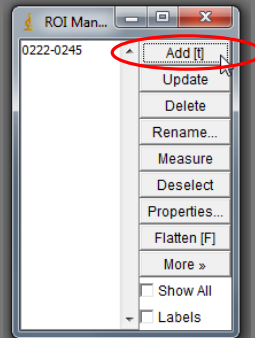
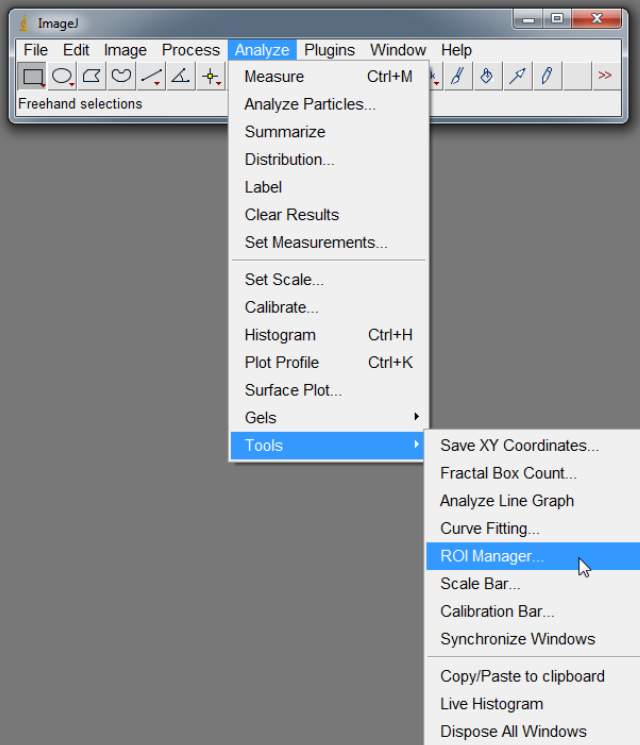
Select **Edit>Selection>Create Selection**. An ROI will appear over the binary image



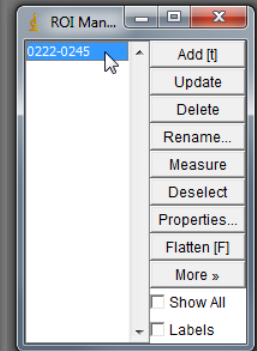
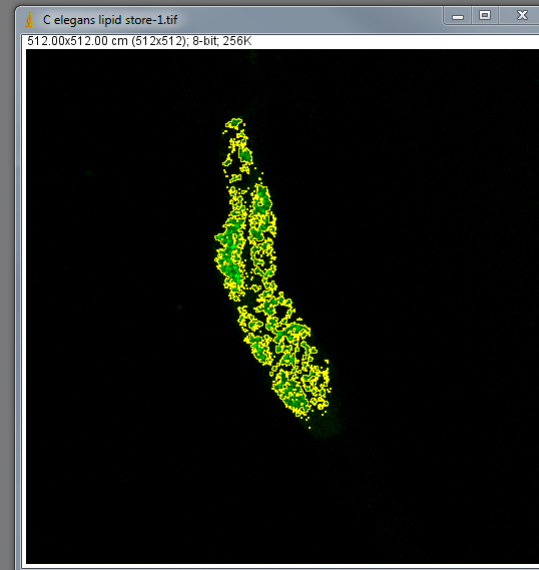
Making precise selections – binary selection method

- ① Select **Analyze>Tools>ROI Manager** and click “Add”. A new ROI appears in the ROI manager.
- ② Open the original image “**C elegans lipid store**”. Select the ROI in the ROI manager, the ROI will appear on the original image ready for analysis.

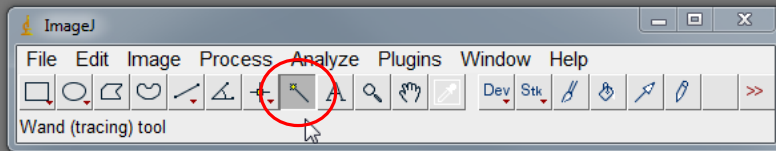
1



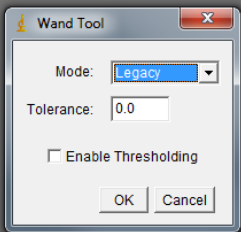
2



Making precise selections – Wand (tracing) tool



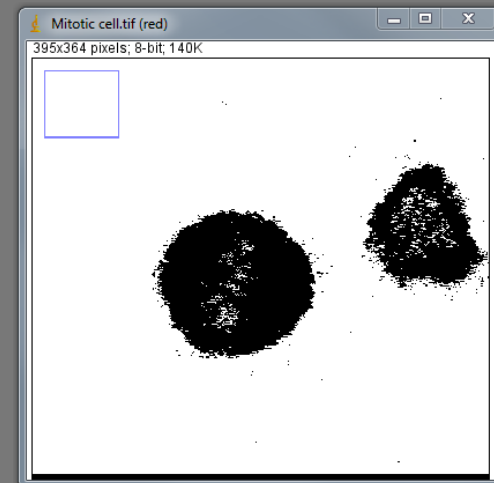
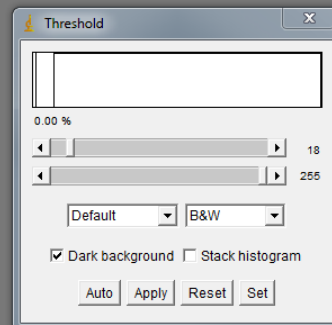
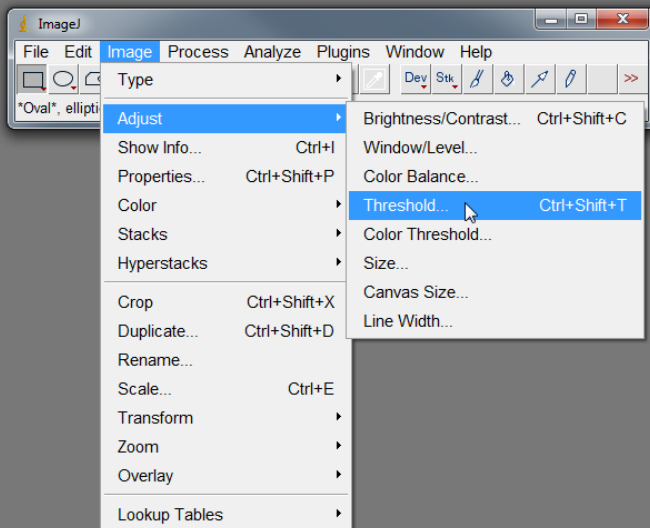
The wand (tracing) tool Makes selections based on grey values when clicked or dragged across an image.



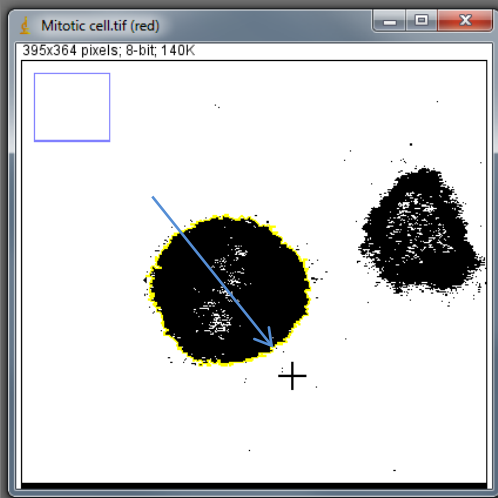
You can double click the icon to set the tolerance, but it works best with a thresholded or binary image.

Open “Mitotic cell.tif (red)”.

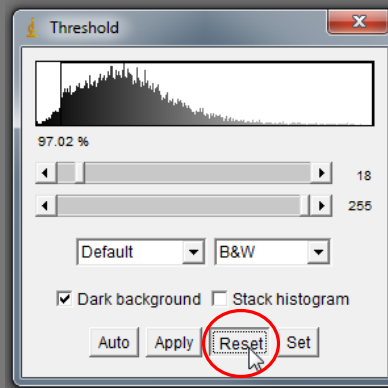
Select **Image>Adjust>Threshold** and threshold the image. Do not click “Apply”.



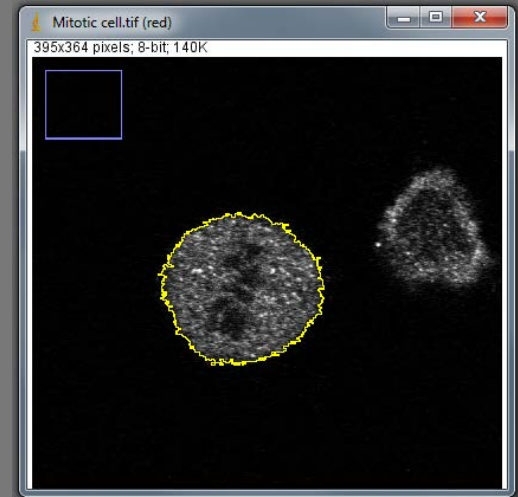
Making precise selections – Wand (tracing) tool



Select the **Wand (tracing) tool**. Click and drag to make the selection then add your selection to the ROI manager.



or reset the threshold



And make your measurements immediately