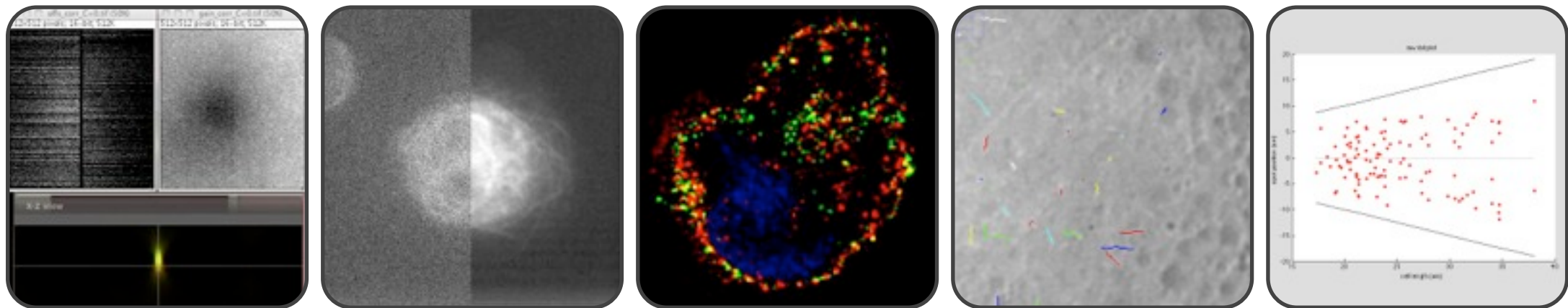


# Lecture 19 – Applied Image Analysis

**Graeme Ball**



# Three Questions

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1. Why should you be interested in image processing & analysis?
2. What are the key concepts & techniques?
3. How can you apply them most efficiently?

# 1. Science requires data

---

Hypothesis testing requires data analysis & statistics

Cell author guidelines:-

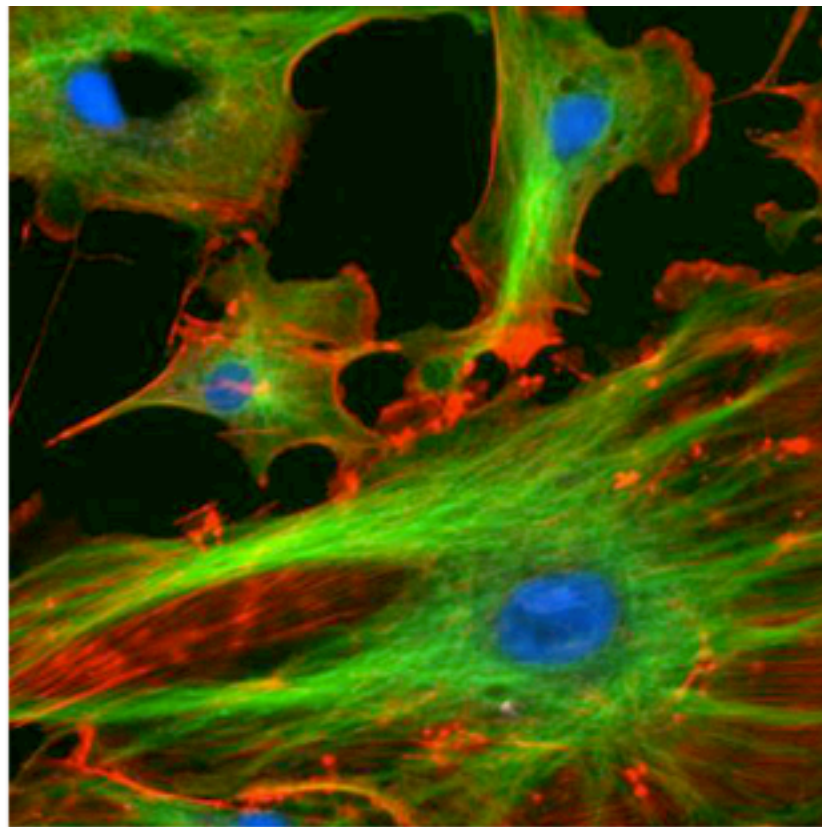
“processing may be unavoidable in certain instances and is permitted provided that the final data accurately reflect that of the original”

“any alterations must be clearly stated in the figure legend and in the methods section”

“ alterations must be applied to the entire image (e.g., brightness, contrast, color balance)”

“ authors will be required to make the original unprocessed data available to the editors”

# 1. Images are data

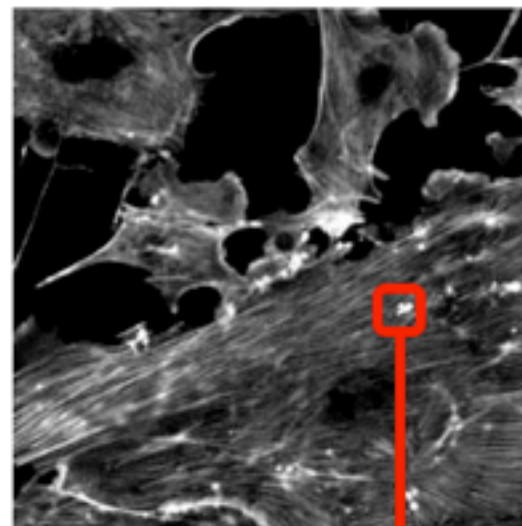


8-bit image (0-255), 3 channels

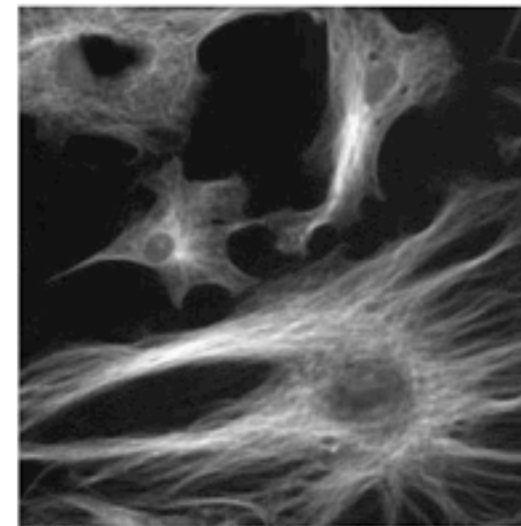
.. think "data", not "picture"

a digital image is:

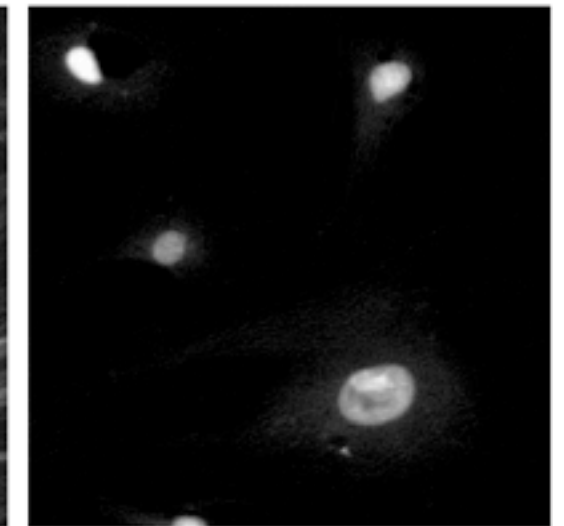
- data: arrays of carefully calibrated intensity measurements
- metadata: data about the data, which is essential for interpretation



"RED" (620nm?)

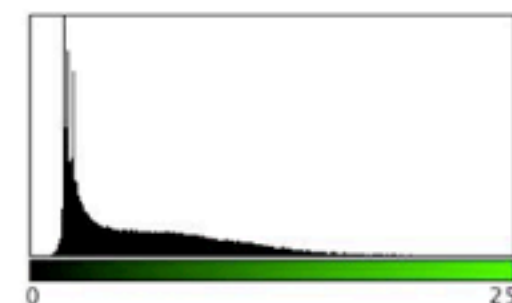


"GREEN" (520nm?)



"BLUE" (460nm)

Pixel Values
289 52 57 62 65 72 76 83 88 89 85 80 75 63 76 95 112 119 117 107 99 79
220 60 59 56 54 70 71 73 74 71 66 61 57 54 63 60 96 106 105 98 91 72
221 62 60 57 54 74 69 65 60 57 57 57 63 68 78 89 94 91 84 78 69
222 58 61 64 65 77 73 64 56 55 57 64 66 80 78 78 80 82 80 73 67 71
223 40 43 44 47 50 65 74 65 48 41 53 71 77 84 89 87 81 78 81 85 84
224 44 52 57 61 60 54 45 39 39 49 61 69 82 77 70 65 69 75 72 63 68
225 56 63 65 65 56 38 26 40 70 89 84 70 78 97 106 86 63 57 59 62 51
226 69 65 58 52 38 36 56 106 157 166 128 83 93 153 196 168 105 68 66 74 44
227 59 48 38 35 36 60 117 193 248 242 182 122 164 213 253 246 201 245 96 63 48
228 42 31 31 37 58 93 158 230 255 255 225 183 248 248 254 255 255 214 117 35 56
229 42 40 48 56 77 105 152 201 236 249 243 236 255 255 255 255 233 170 92 40 56
230 57 60 69 78 79 97 126 155 186 216 245 255 228 255 255 246 155 72 52 69 53
231 68 77 78 70 66 75 99 159 217 250 255 255 255 248 204 151 183 73 58 54 49
232 68 73 72 65 63 114 190 254 255 255 226 204 201 184 153 120 94 82 78 79 96
233 75 74 66 57 68 147 239 255 254 290 151 125 113 107 96 90 92 98 106 110 149
234 87 81 65 50 72 135 190 184 132 90 77 78 74 75 78 89 102 115 126 131 153
235 89 81 61 43 56 94 117 94 53 58 55 77 84 90 98 107 117 123 124 123 104
236 77 69 51 38 35 62 83 81 67 62 72 87 101 107 117 120 117 107 100 93 56
237 60 48 41 40 42 50 63 76 80 77 74 77 92 104 113 113 99 79 64 55 35
238 45 36 36 46 66 49 40 46 51 55 55 57 74 87 100 98 80 56 38 30 29
239 36 42 46 44 71 68 58 44 37 40 53 65 93 93 88 75 58 45 38 37 90

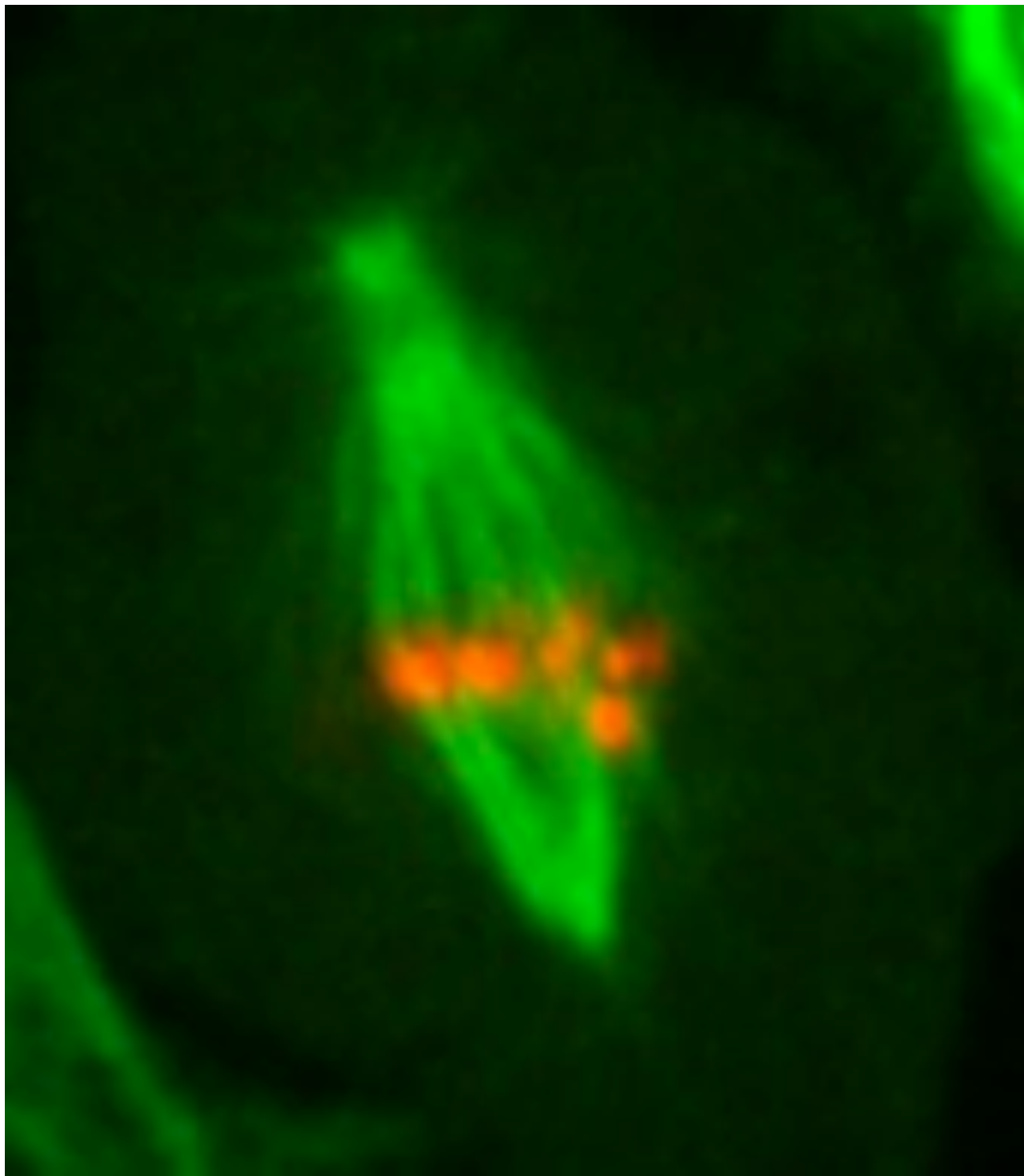


Count: 160000  
Mean: 60.209  
StdDev: 44.990  
Min: 0  
Max: 255  
Mode: 17 (9693)

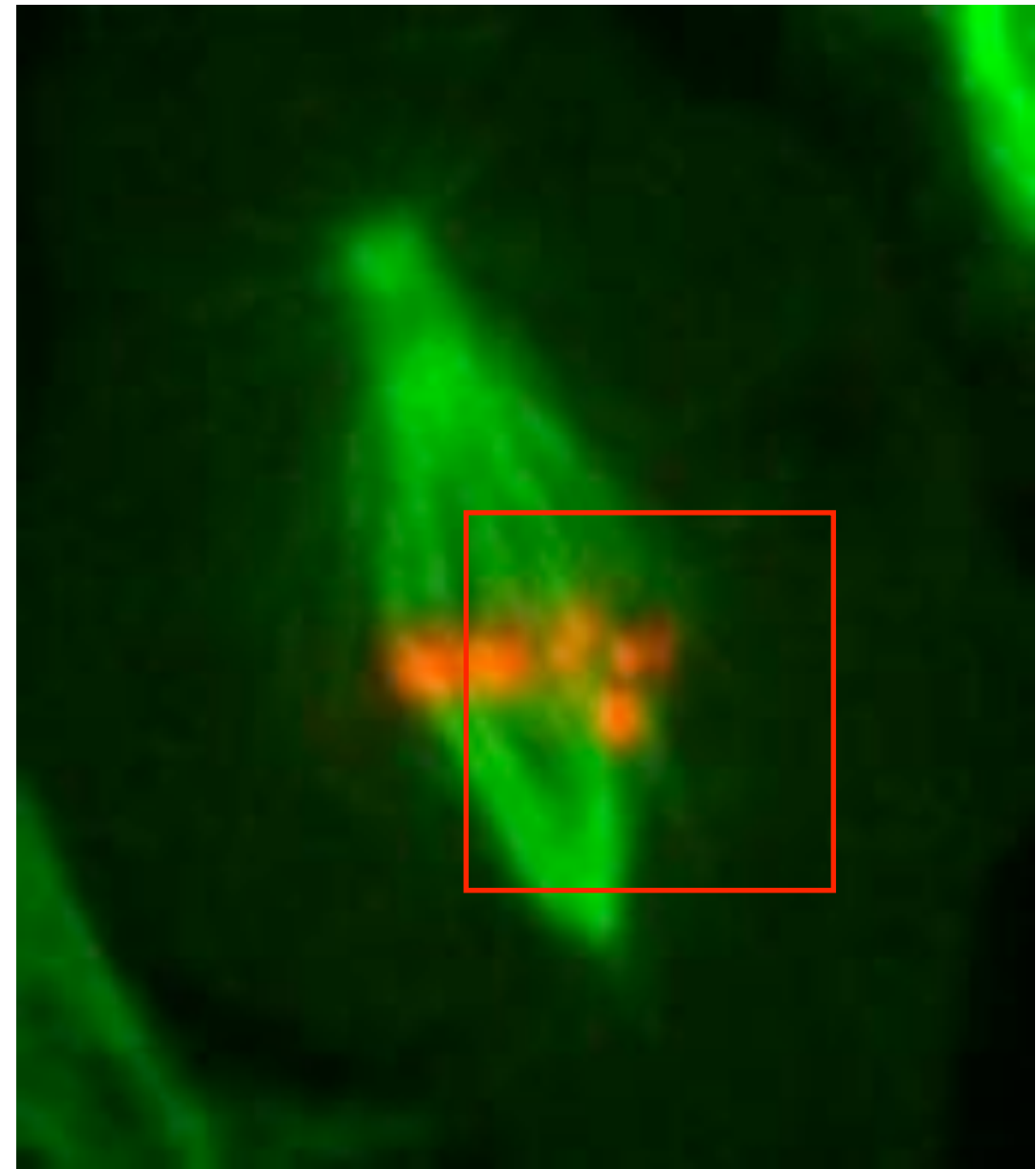


# 1. JPEGs are (much) less data

---



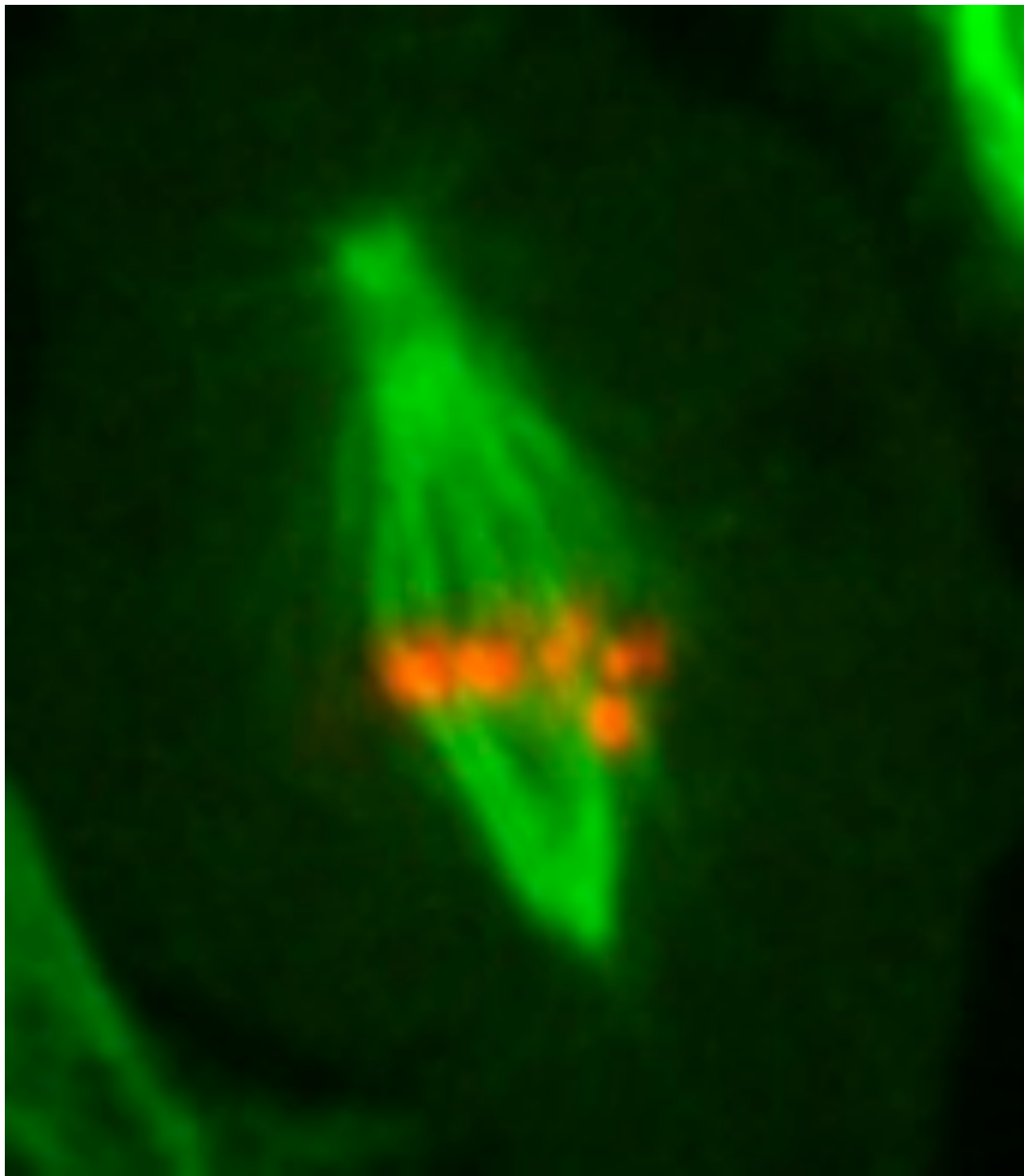
5D 16-bit data: 33 MB,  
17 million pixels (0-65535)  
134 KB per 2D plane



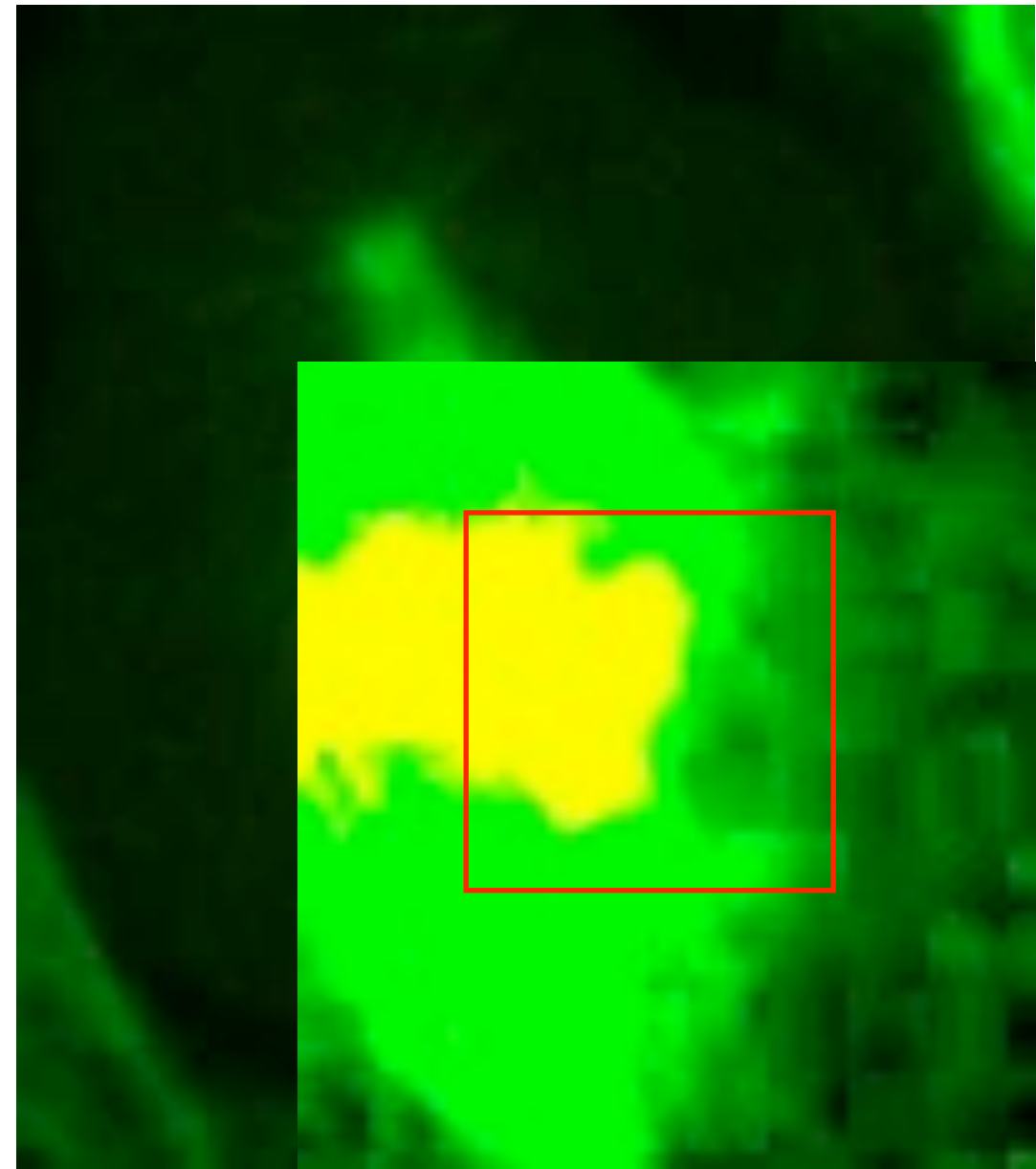
8-bit JPEG (0-255 per channel)  
3 KB per 2D plane

# 1. JPEGs are (much) less data

---



5D 16-bit data: 33 MB,  
17 million pixels (0-65535)  
134 KB per 2D plane



8-bit JPEG (0-255 per channel)  
3 KB per 2D plane

# 1. Collect the right data

---

What am I trying to measure?

# 1. Collect the right data

---

## What am I trying to measure?

Expression level?



accurate, calibrated intensities



# 1. Collect the right data

---

## What am I trying to measure?

Expression level?



accurate, calibrated intensities

Distance or colocalization?



contrast-to-noise, resolution,  
alignment

# 1. Collect the right data

---

## What am I trying to measure?

Expression level?



accurate, calibrated intensities

Distance or colocalization?



contrast-to-noise, resolution,  
alignment

Dynamics?



temporal resolution, photostability

# 1. Collect the right data

---

## What am I trying to measure?

Expression level?



accurate, calibrated intensities

Distance or colocalization?



contrast-to-noise, resolution,  
alignment

Dynamics?



temporal resolution, photostability



decide on the instrument (& technique)

# 1. Collect the right data

## What am I trying to measure?

Expression level?



accurate, calibrated intensities

Distance or colocalization?



contrast-to-noise, resolution,  
alignment

Dynamics?



temporal resolution, photostability



decide on the instrument (& technique)



choice of fluorophores



# 2. Image Processing & Analysis

---

## Image processing

Image restoration – correction, denoising, deblurring



Filtering to enhance features of interest

Segmentation – dividing an image into useful categories (e.g. feature, background)

## Image analysis

Identifying and tracking features

Summary statistics: intensity/amount, distribution, speed, colocalization

# 2. Image Restoration

---

## Image restoration aims to recover the true image

Flat-field correction – uneven illumination (also, pseudo-correction)

Denoising / noise filtering – smoothing, neighborhood filters, non-local

Deblurring – deconvolution with or without PSF, unsharp mask

Image registration – rigid/affine versus elastic – intensity-based or feature-based

Normalization – intensity of each time-point scaled to correct bleaching (& flicker)

# 2. Image Filtering

---

## Image filtering enhances some features / rejects others

Spatial filters for smoothing & sharpening

Frequency domain filters enhance/suppress features based on size

Histogram operations, e.g. histogram equalization for contrast enhancement

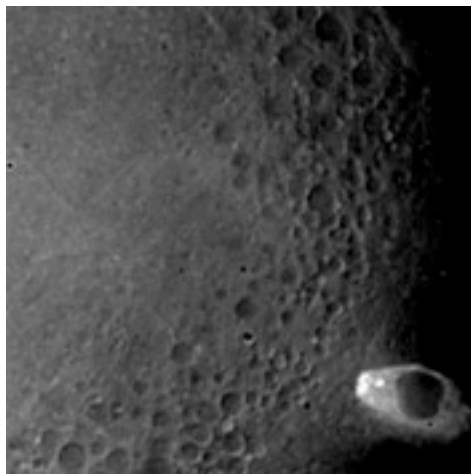
Adaptive filtering equivalent to global filtering after normalization

Time-domain filtering – see temporal median filter in the tracking example

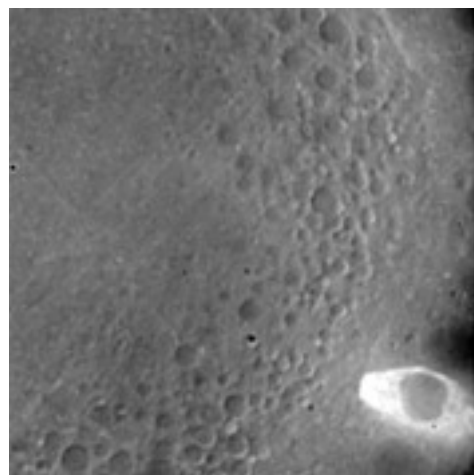
# 2. Image Processing Examples

pseudo-flat field

$$I_m = I_m - \text{mean} * I_m / \text{mean} - \text{filtered}$$



original

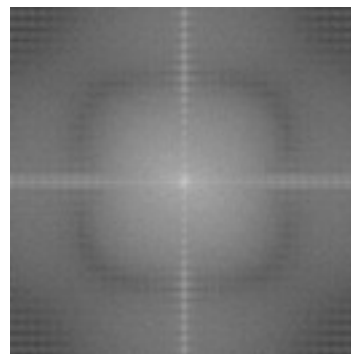


pseudo-corrected

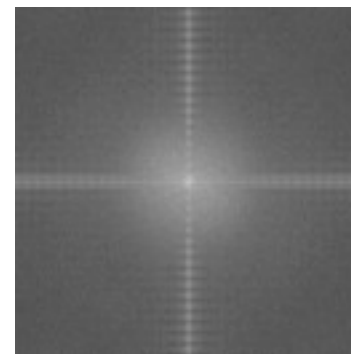


240x240 mean-filt

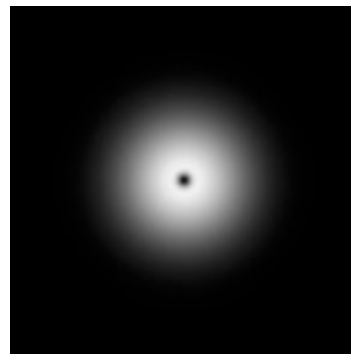
FFT reveals frequencies



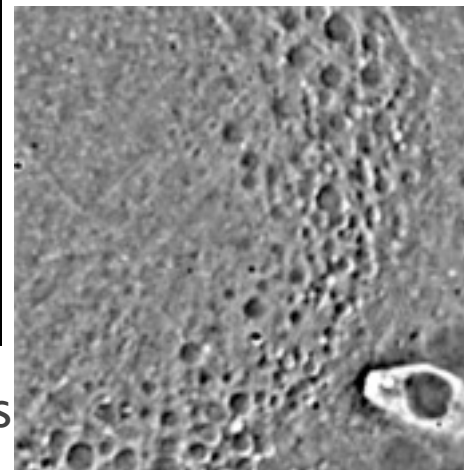
original



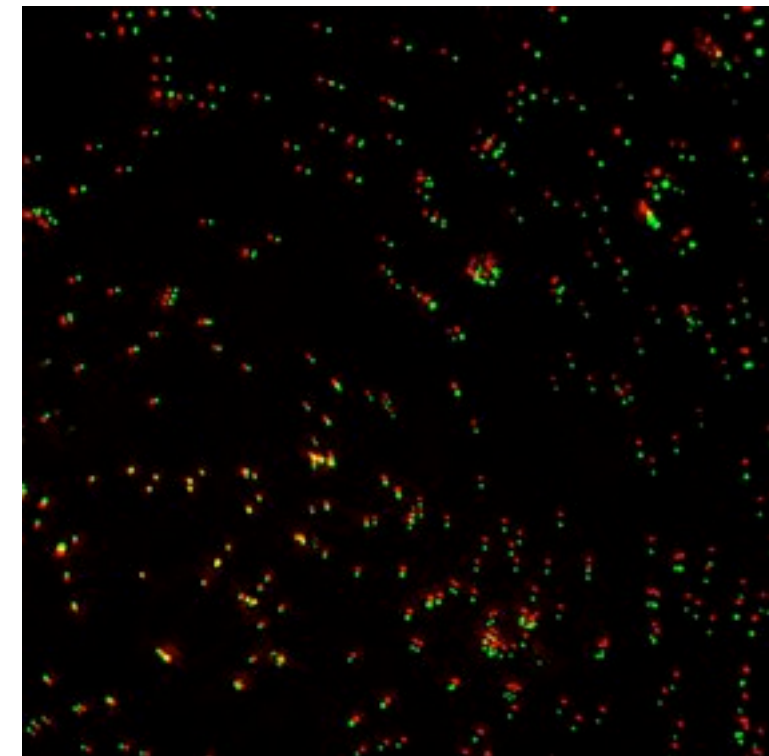
3x3 median



3-25 bandpass



registration can  
be critical!

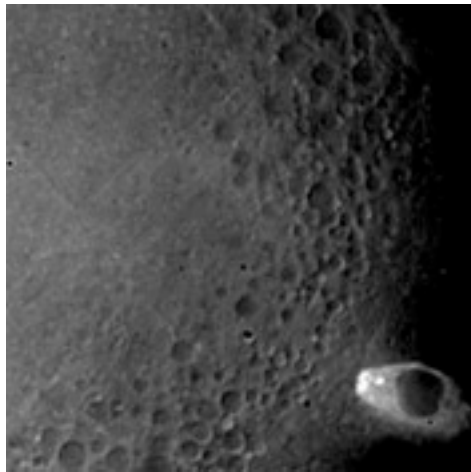




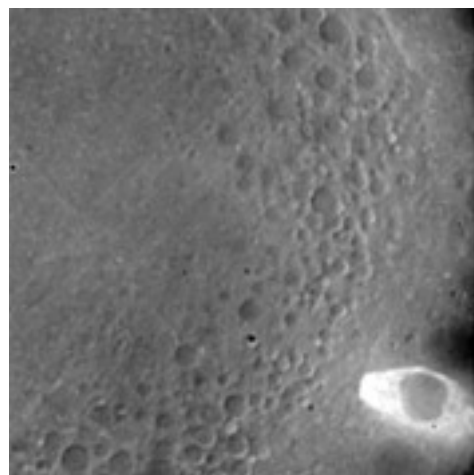
# 2. Image Processing Examples

pseudo-flat field

$$I_m = I_m - \text{mean} * I_m / \text{mean} - \text{filtered}$$



original

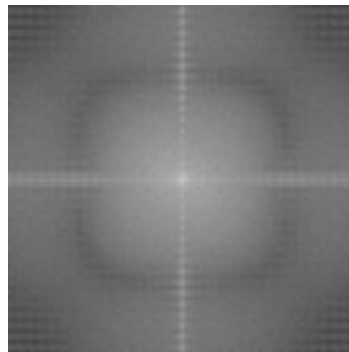


pseudo-corrected

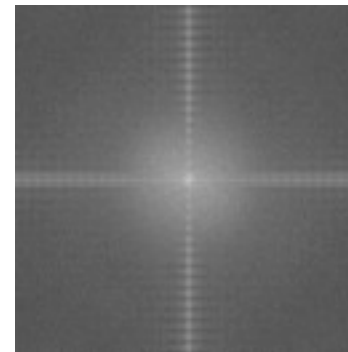


240x240 mean-filt

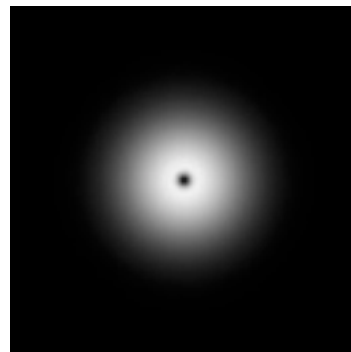
FFT reveals frequencies



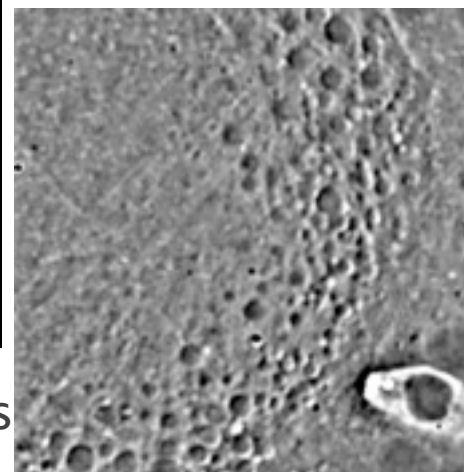
original



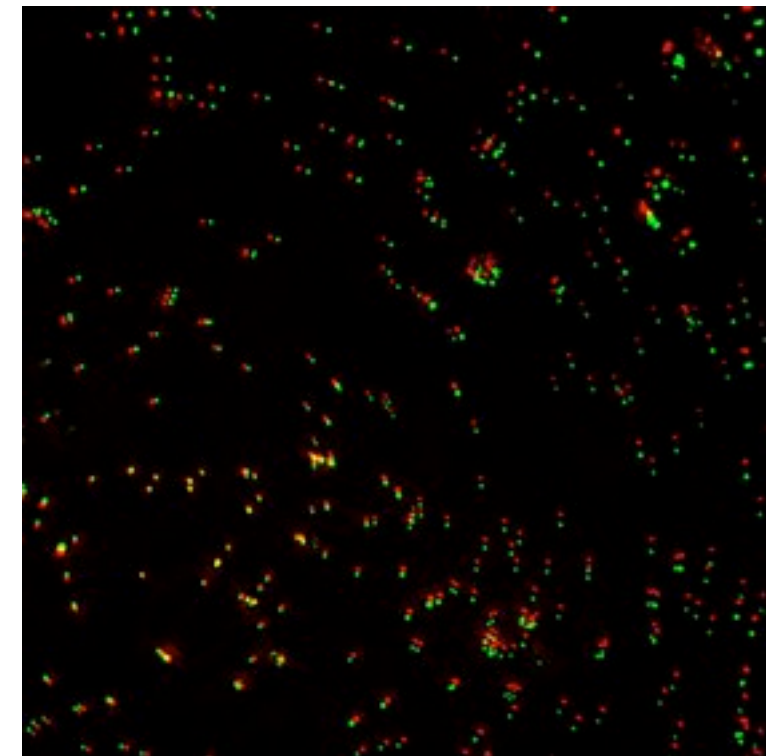
3x3 median



3-25 bandpass



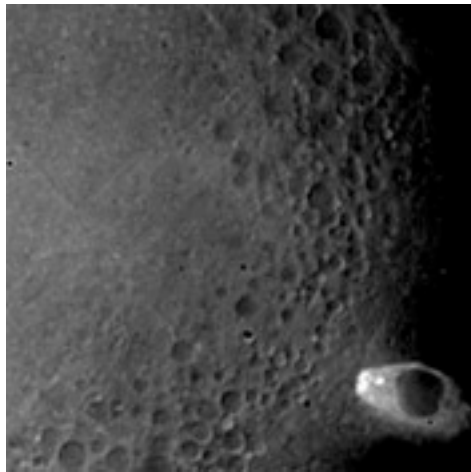
registration can be critical!



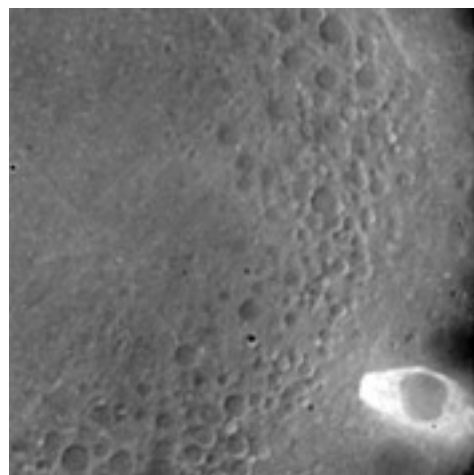
# 2. Image Processing Examples

pseudo-flat field

$$I_m = I_m - \text{mean} * I_m / \text{mean} - \text{filtered}$$



original

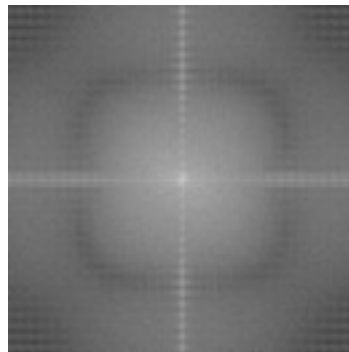


pseudo-corrected

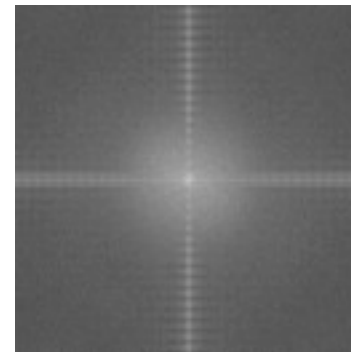


240x240 mean-filt

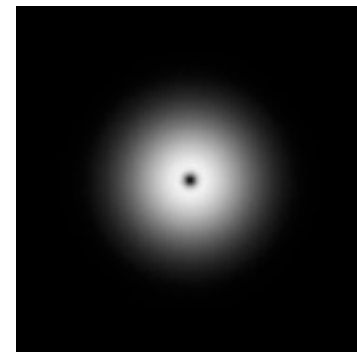
FFT reveals frequencies



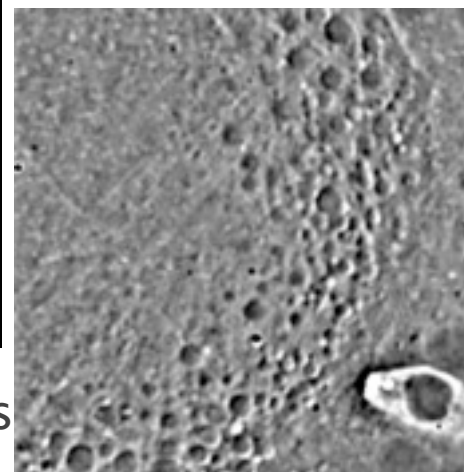
original



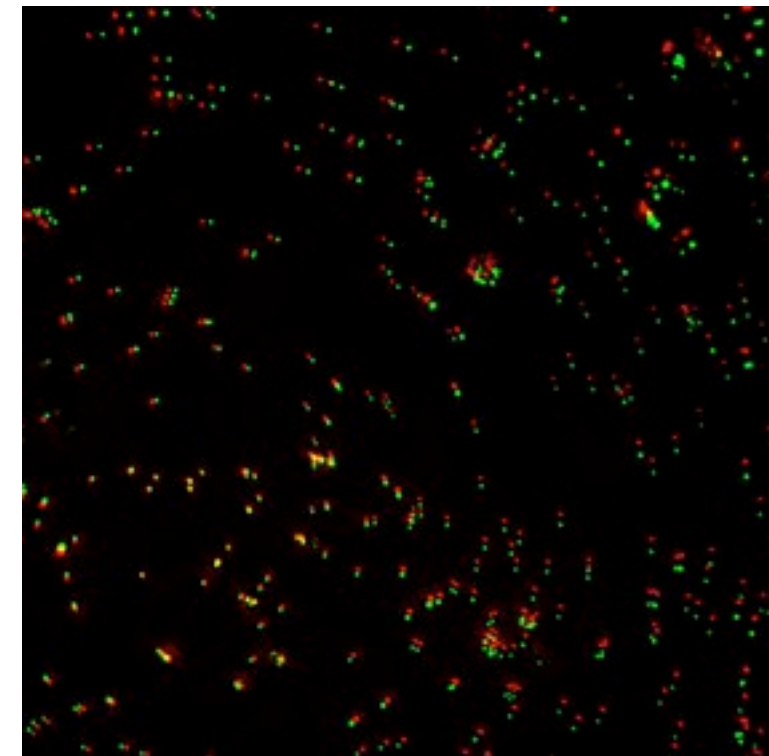
3x3 median



3-25 bandpass



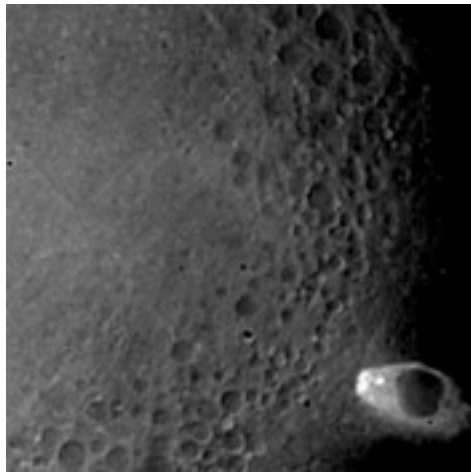
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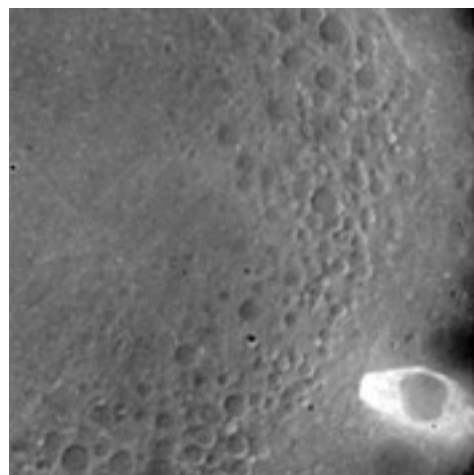
# 2. Image Processing Examples

## pseudo-flat field

$$Im = Im - mean * Im / mean - filtered$$



original

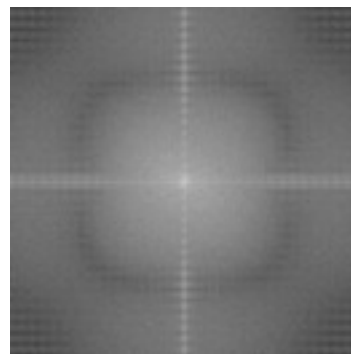


pseudo-corrected

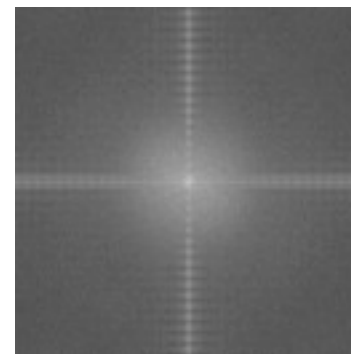


240x240 mean-filt

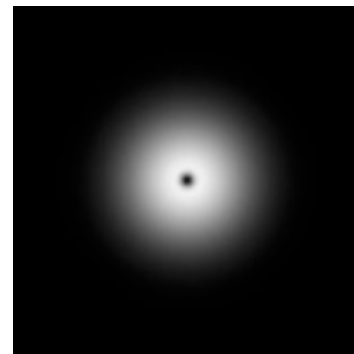
## FFT reveals frequencies



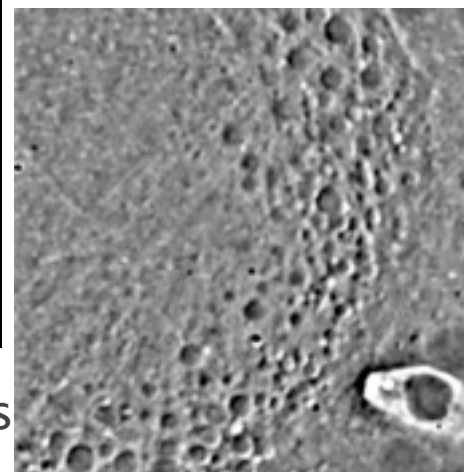
original



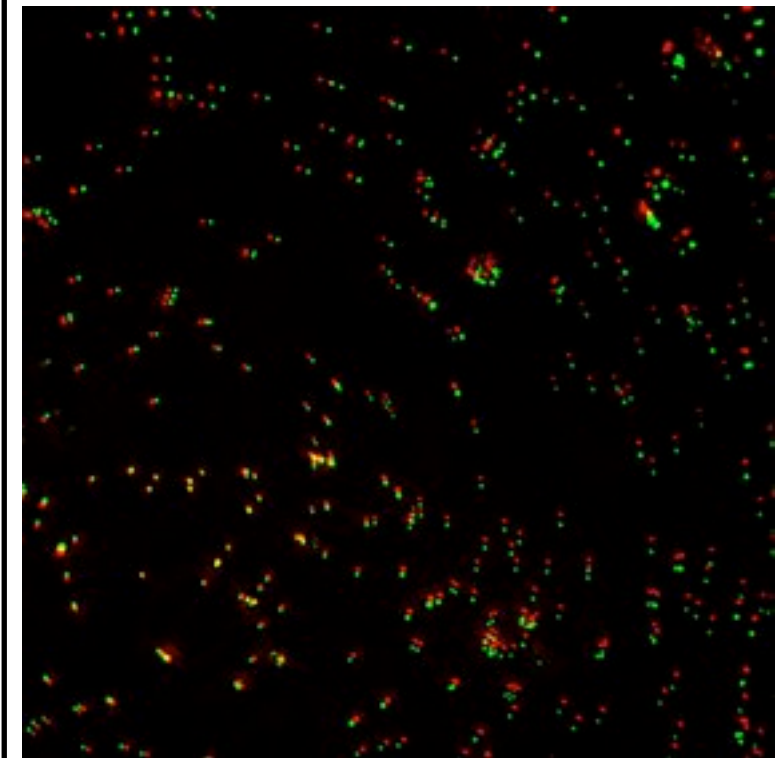
3x3 median



3-25 bandpass



registration can  
be critical!





# Patch-based denoising: 10-100 x less light?

8 ms exposure, 10% 488 Laser power

8 ms exposure, 0.1% 488 Laser power

Macrophage: Jupiter-GFP 7Z, 3stacks/s (Richard Parton)

Jerome Boulanger: SAFIR Denoising software

Integrated into Priism by the John Sedat Group UCSF

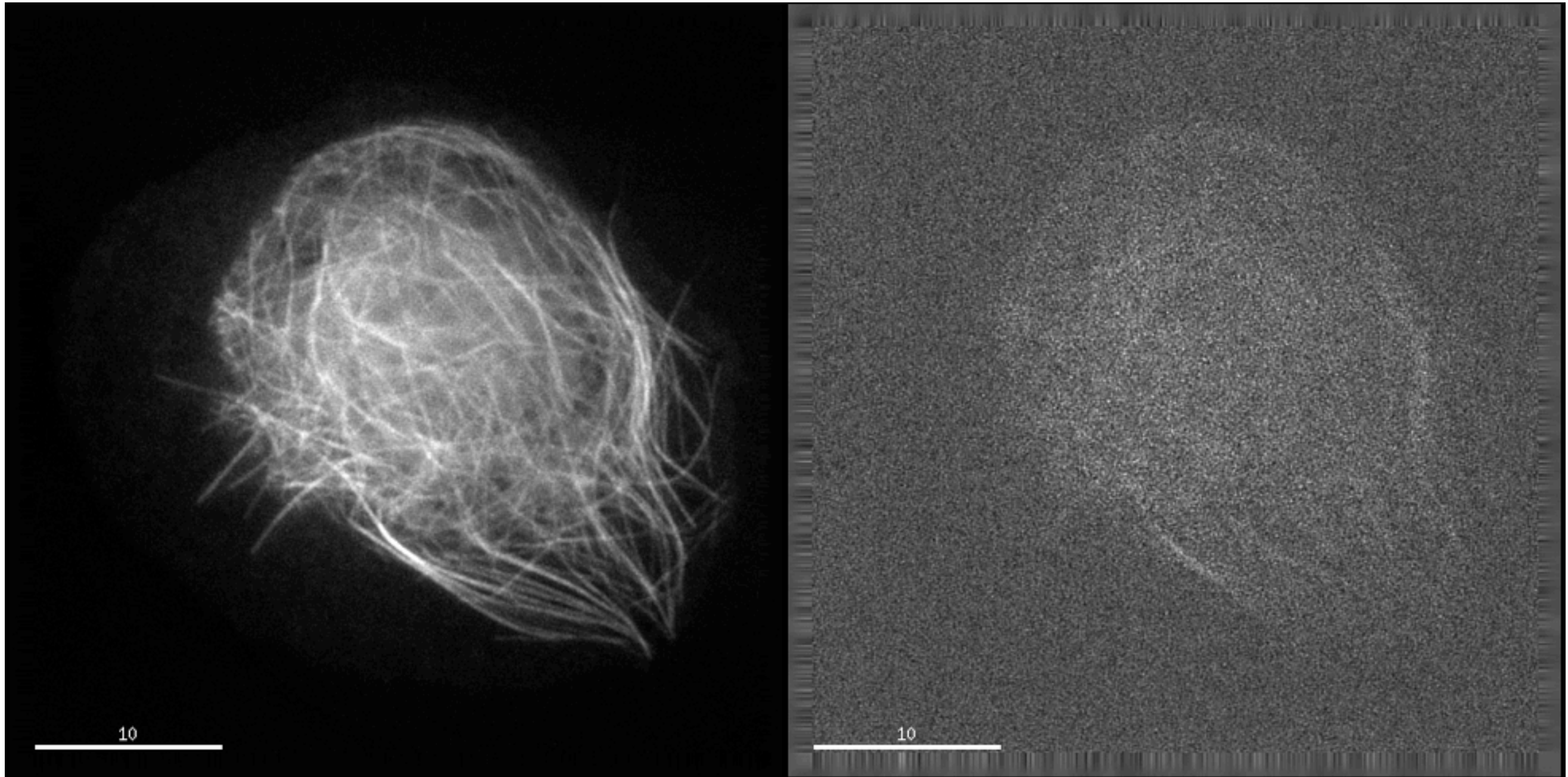
J. Boulanger, C. Kervrann, and P. Bouthemy, "Space-time adaptation for patch-based image sequence restoration," *IEEE Trans. on Pattern Analysis and Machine Intelligence*, vol. 29, no. 6, pp. 1096–1102, June 2007



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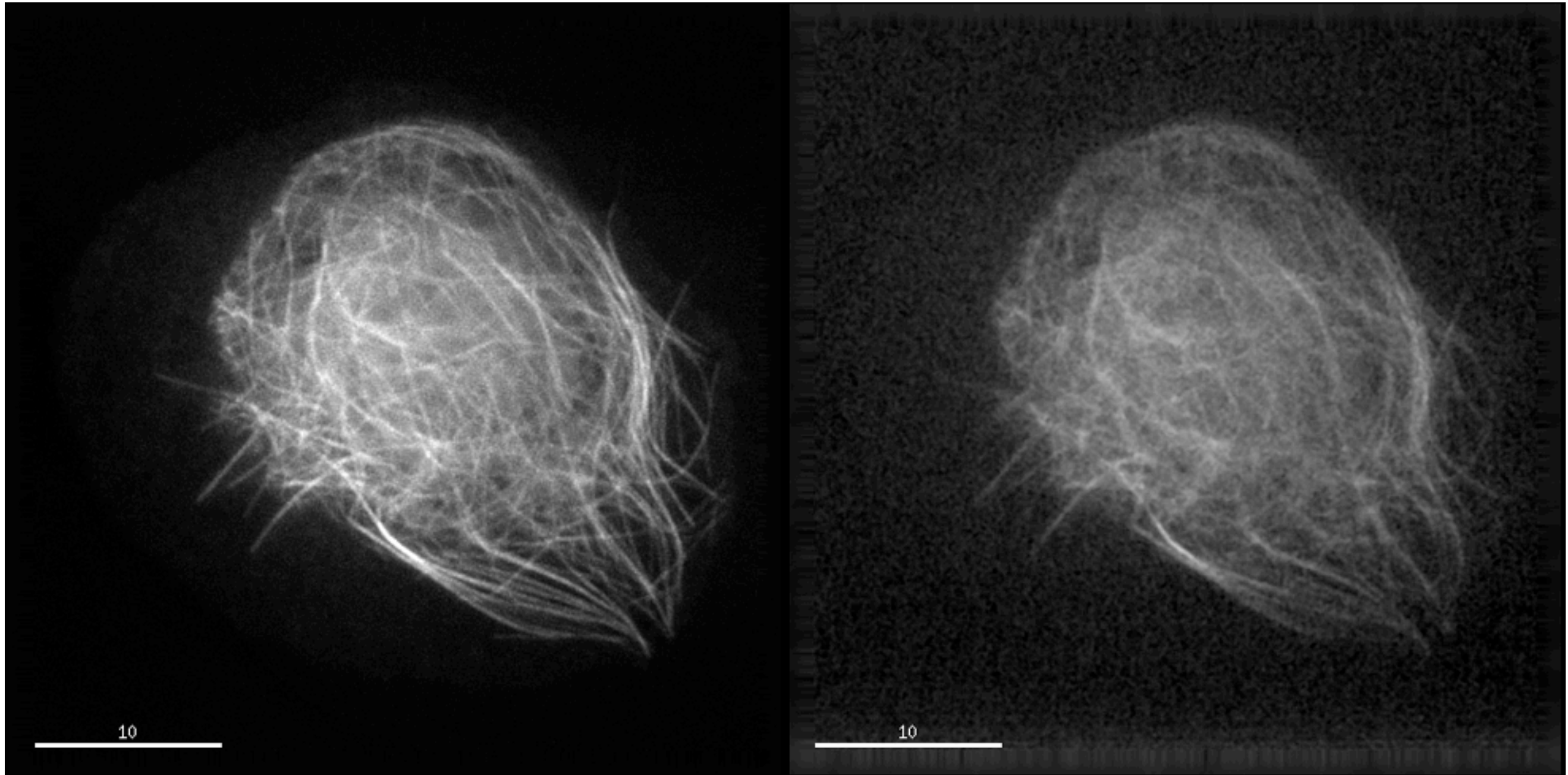
Integrated into Priism by the John Sedat Group UCSF

J. Boulanger, C. Kervrann, and P. Bouthemy, "Space-time adaptation for patch-based image sequence restoration," *IEEE Trans. on Pattern Analysis and Machine Intelligence*, vol. 29, no. 6, pp. 1096–1102, June 2007

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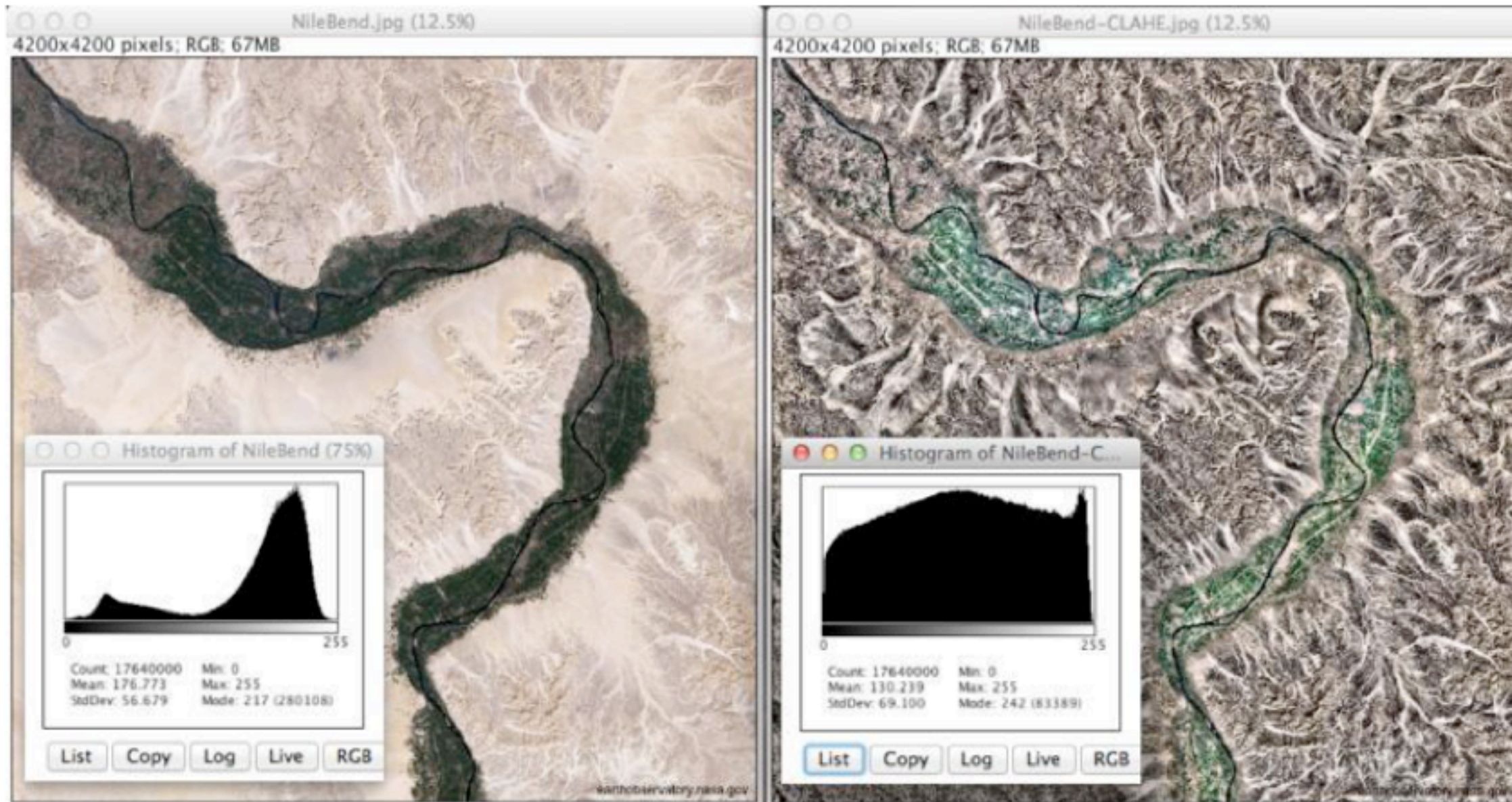
Integrated into Priism by the John Sedat Group UCSF

J. Boulanger, C. Kervrann, and P. Bouthemy, "Space-time adaptation for patch-based image sequence restoration," *IEEE Trans. on Pattern Analysis and Machine Intelligence*, vol. 29, no. 6, pp. 1096–1102, June 2007



# CLAHE

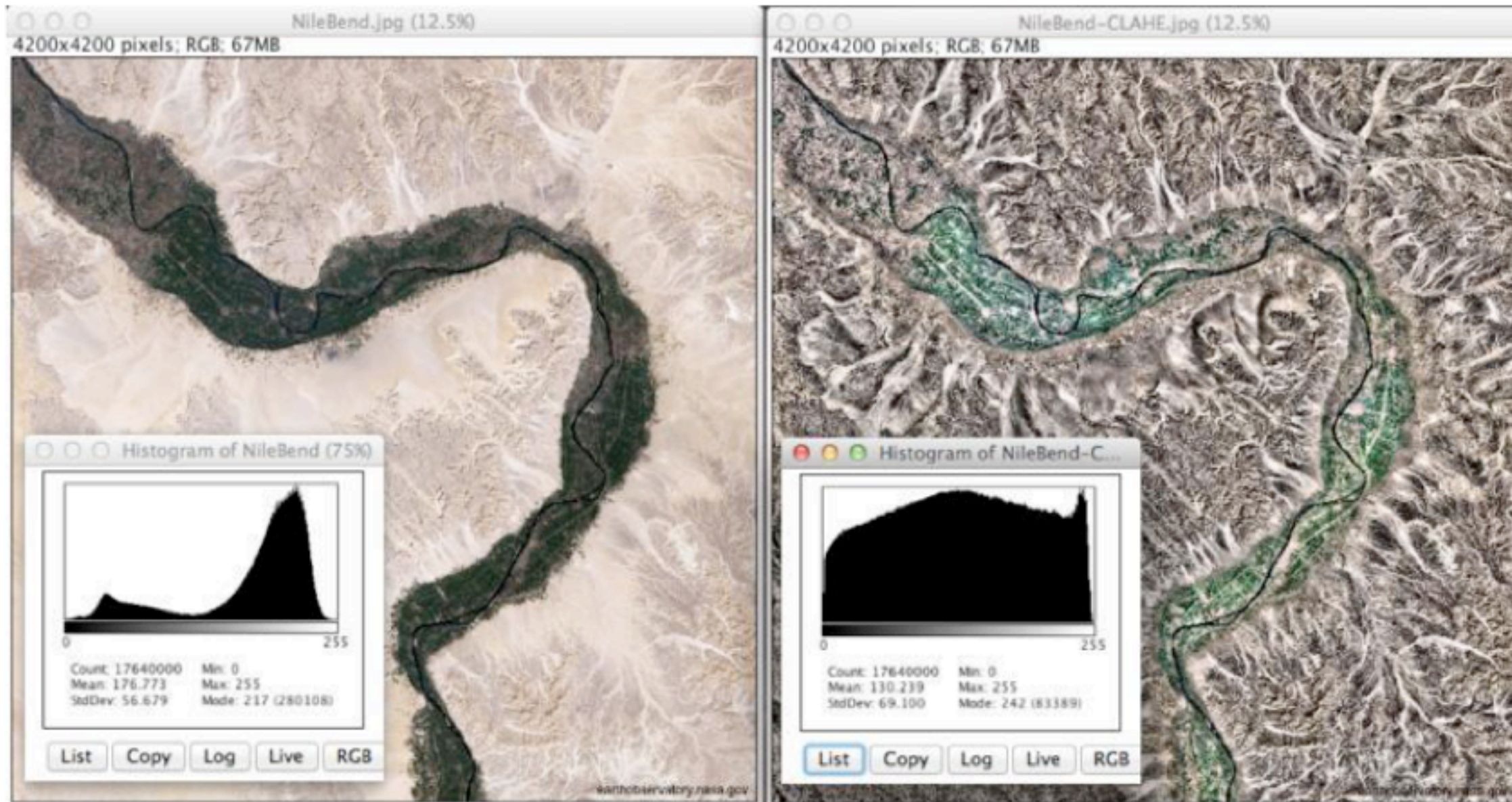
- Contrast Limited Adaptive Histogram Equalization





# CLAHE

- Contrast Limited Adaptive Histogram Equalization



# 2. Image Segmentation

---

## Segmentation separates an image into subregions

2 common representations: “binary mask” images, and Regions Of Interest (ROIs)

Simple intensity thresholding – several methods (e.g. Otsu) to estimate threshold

Spot/particle detection – intensity, size and shape

Edge detection (e.g. Sobel) & Morphological image processing\* (erosion, dilation)

Watershed calculation, Voronoi diagram, Ultimate eroded points

Machine learning and Manual options

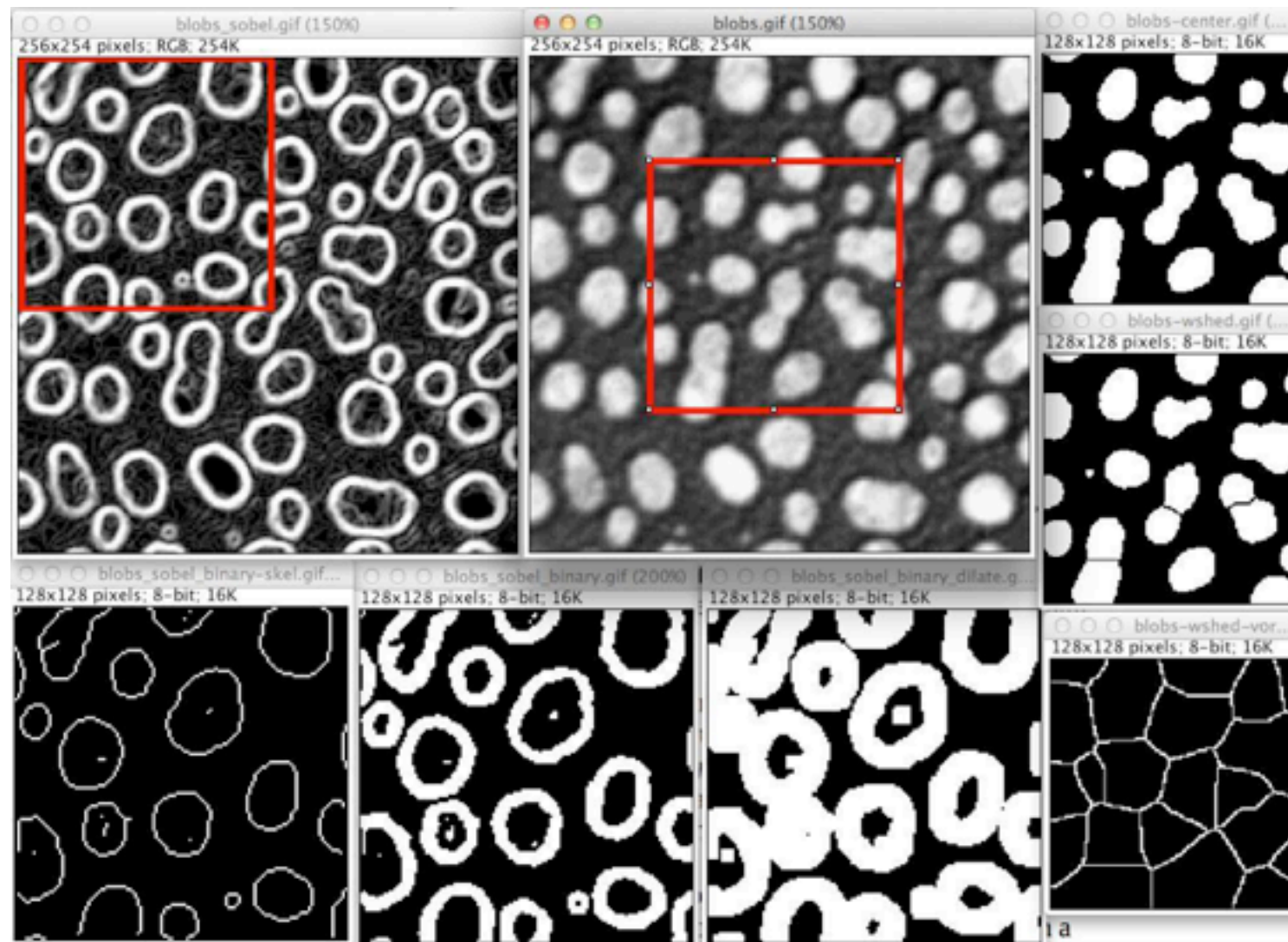


# 2. Binary masks & operations

Binary images usually created using a intensity threshold

ImageJ binary images are 8-bit gray, 255 and 0 (instead of 1 and 0)

Binary operations: erode, dilate, skeletonize, watershed, Voronoi



# Edge detection

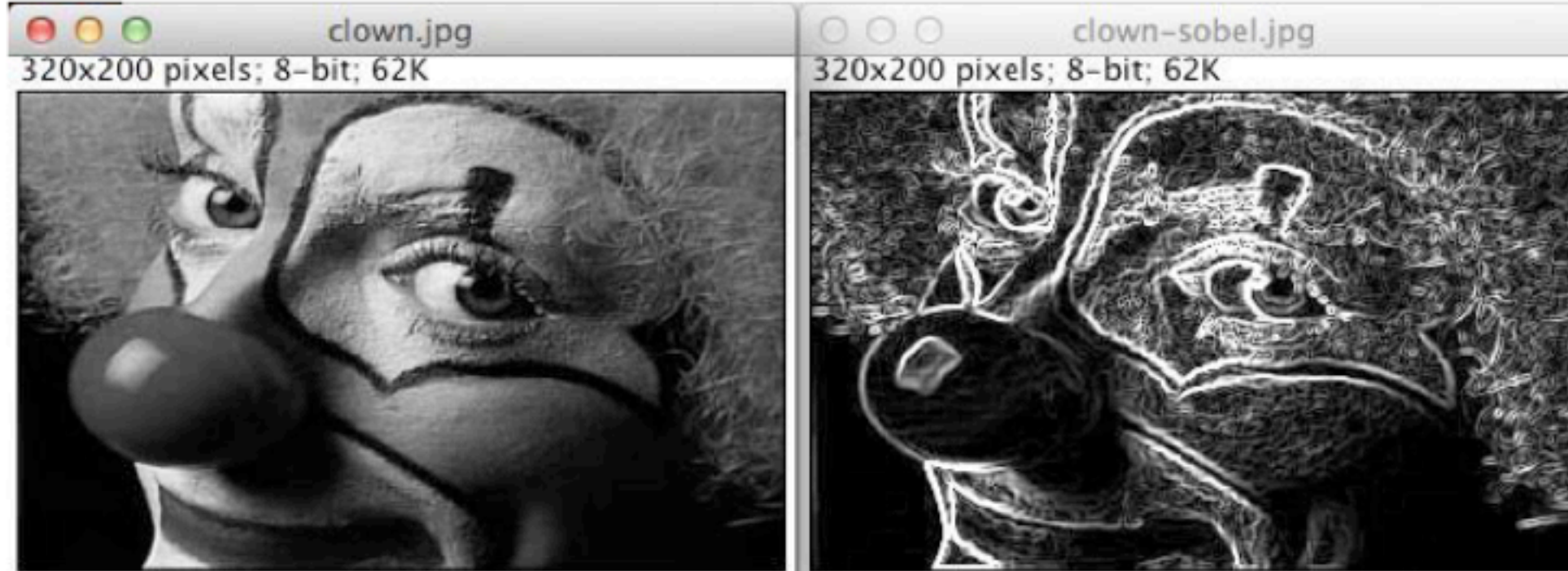
- "Find edges" is a Sobel operator: detects high gradient at edges

-1	0	+1
-2	0	+2
-1	0	+1

Gx

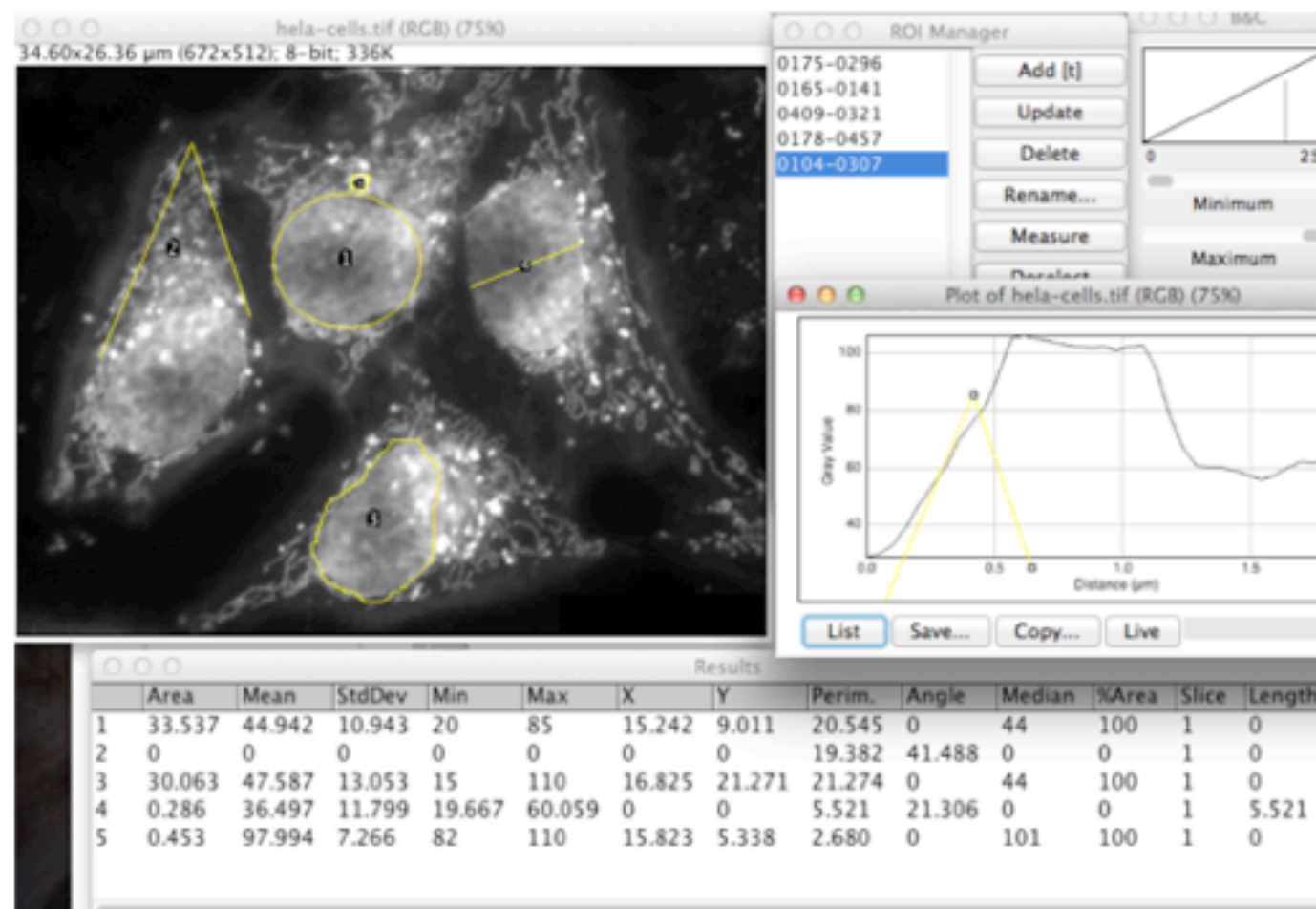
+1	+2	+1
0	0	0
-1	-2	-1

Gy



# Selections & measurements

- various selection tools: points, lines, enclosed regions, wand tool
- measure distances & angles
- add selection to ROI manager "t"
- *Measure* command (& *Set Measurements*)





# 2. Fiji/ImageJ tips

---

- useful tools that are easy to miss: the wand, ROI manager, brush selection
- understand how to manipulate stacks, hyperstacks and virtual stacks – e.g. how to convert, project, reduce, combine; channels tool
- make use of image histogram, “plot profile” and threshold tool
- learn how to “set measurements” and measure
- read the manual: <http://rsbweb.nih.gov/ij/docs/user-guide.pdf>

# 3. Free tools available



ImageJ / Fiji – versatile 2D+ image analysis tool with many plugins



Icy, Vaa3D, BioimageXD – 3D image visualization & analysis

VAA3D 挖三维

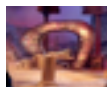


KNIME – data analysis workflow for large datasets / batch



CellProfiler – quantitatively measure cell phenotypes in large datasets (+ Worm Toolbox)

UCSF



Priism/IVE, Priithon & Editor – 2D image processing/analysis (DV/OMX)



OMERO – image repository & visualization

# 3. Commercial Analysis Tools



Imaris, Amira – 3D visualization & analysis packages



Volocity – 3D visualization & analysis package, spinning disk



SoftWoRx – API Deltavision, deconvolution, SI reconstruction



Metamorph – microscope control, image processing/analysis



Huygens, AutoQuant – Deconvolution software



# 3. 3D visualization & analysis

---

- ImageJ/Fiji has 3D visualization & analysis functionality:  
3D viewer, Volume Viewer, Image 5D, hyperstacks, orthogonal view,  
3D objects counter
- Imaris, Amira and Volocity are 3D image visualization & analysis packages  
designed for microscopy (and medical imaging)
- Choice between viewing fluorescence intensity (often MIP) versus  
generating surface representations of objects

# 3. Imaris movie

---

Image of bacterial septa (GFP) and DNA (DAPI), Christian Lesterlin

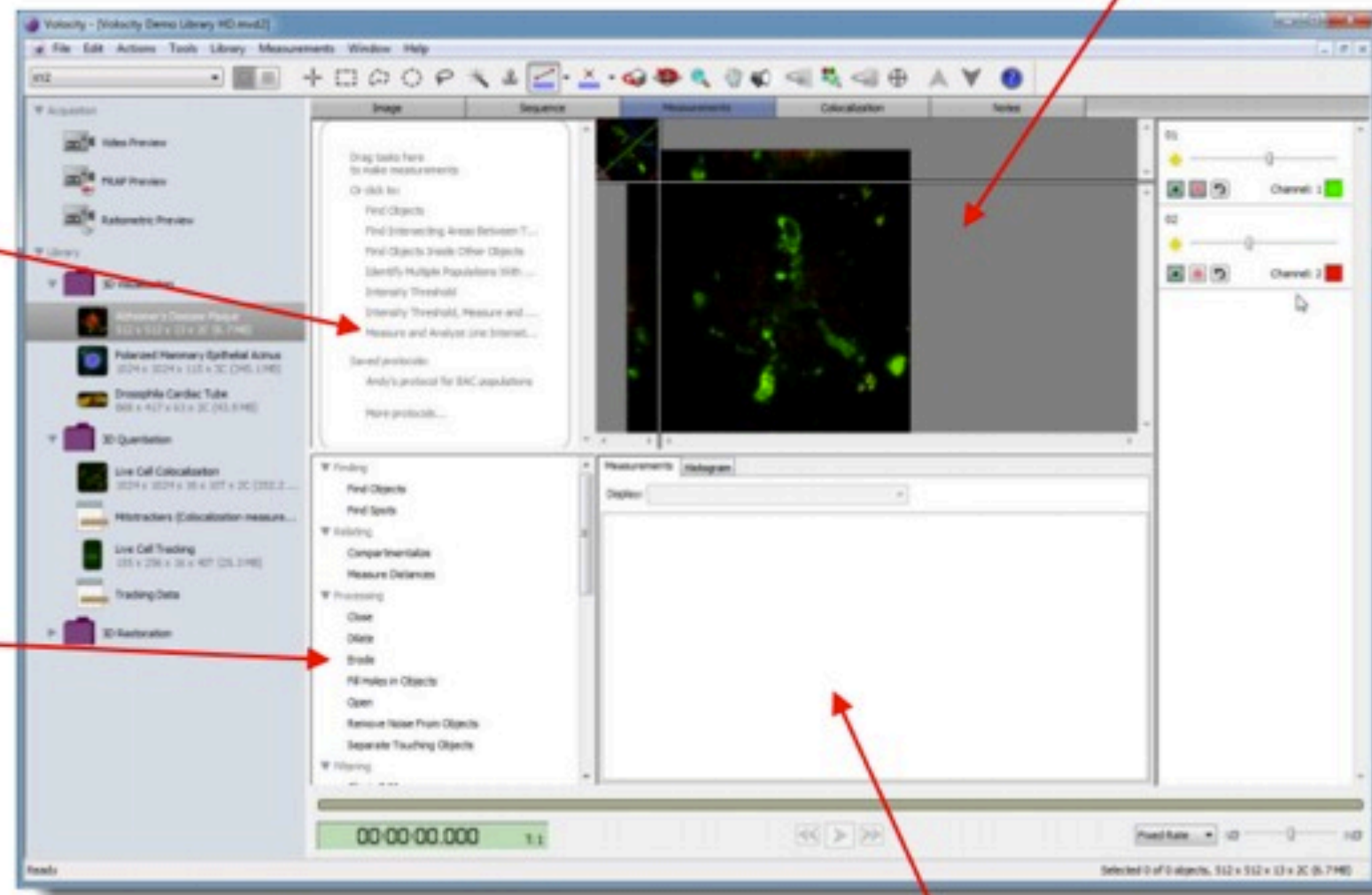
# 3. Volocity

View the Measurements tab. The Measurements View contains all the tools and information needed for selecting objects.

Image preview shows feedback as measurements are made

Drag measurement protocol tasks to this pane to make protocols

Measurement protocol tasks



Measurements are shown here as a table or histogram

<http://www.perkinelmer.com/PDFs/downloads/CreatingMeasurementProtocolVolocitySoftware.pdf>

# 3. Measuring Colocalization

Read Bolte & Cordelieres' 2006 review :-

<http://www.ncbi.nlm.nih.gov/pubmed/17210054>

- prerequisites: make **very** certain that:-
  1. you do not have bleed-through! (or crosstalk)
  2. your channel alignment is properly calibrated
  3. your images are as noise-free and deblurred as possible
- many colocalization statistics rely on segmenting both channels  
=> flat field & meticulous background correction
- use ROIs or masks to **analyze different compartments separately**
- 2 fundamental approaches:
  1. **intensity correlation** scatter plot, Pearson's (PCC)
    - 1b. Manders coefficients (M1, M2)
  2. **object-based analysis** (distances)

**JaCoP**  
(ImageJ plugin)

# 3. Scatter plots and ICC

Scatter plot of channel intensities

reveals presence/absence of colocalization:-

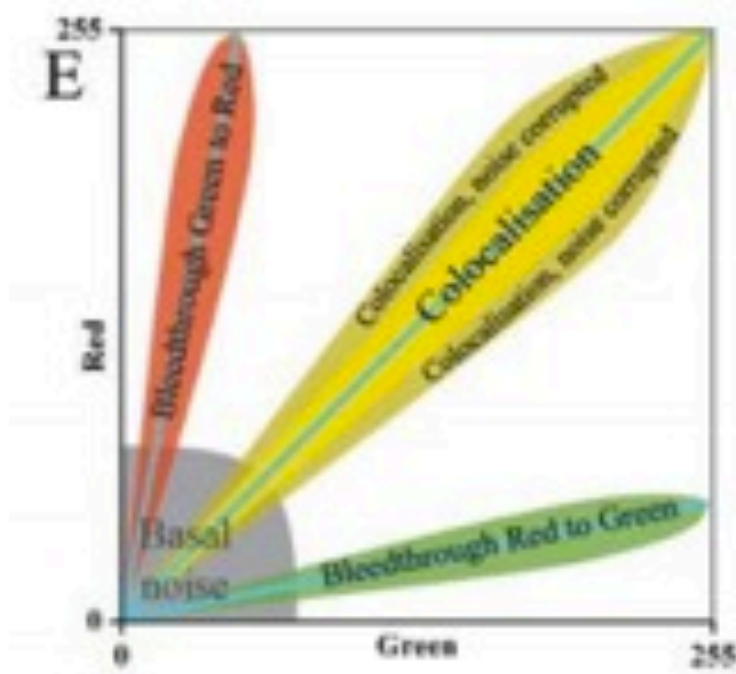


Fig. 5E

(Bolte & Cordelieres)

- Pearson's Correlation Coefficient (PCC) of 0.5–1 indicates colocalization
- large difference between M1 and M2 indicates more of one label – NB. if one label is everywhere, colocalization is meaningless!
- unlike PCC, negative values of Li's ICQ indicate mutual exclusion (0 means uncorrelated, 0.5 means colocalization)

Fiji & ImageJ have **Coloc\_2** and **JACoP** plugins (JACoP has object-based method)



# 3. Scripts & statistics

---

## Measurement, automation & statistics

Quantitative results: care, consistency, avoid systematic errors, avoid bias\*

Manual analysis vs. Macros vs. customized software tools for automation

In addition to Excel, other useful statistics software: R, MATLAB



\* recommend blind analysis to avoid bias



# 3. ImageJ Macros save time

---

- first, try out some processing/analysis options manually
- turn on the recorder ... “Plugins > Macros > Record”
- you will see a command equivalent to every task you carry out
- paste a sequence of commands into new Macro (Plugins > New > Macro)
- for a description of how Macros work and info about in-built functions, see –  
<http://rsbweb.nih.gov/ij/developer/macro/macros.html>  
<http://rsbweb.nih.gov/ij/developer/macro/functions.html>
- result: gbSumMaskedSignal.ijm; for NMJ screen, James Halstead (Davis Lab)

# 3. KNIME for data mining

The screenshot displays the KNIME software interface with a workflow titled 'CountChromosomes'. The workflow is organized into four main stages:

- Preprocessing:** Includes 'Image Reader' (Node 175) and two 'Image Cropper' nodes (Node 176 and Node 177).
- Segmentation:** Includes 'Segment Chromosomes' (Node 112) and 'Segment Nuclei' (Node 113).
- Labeling Combination:** Includes 'Labeling to Table' (Node 141), two 'Java Snippet' nodes (Node 172 and Node 173), and a 'Concatenate' node (Node 136).
- Analysis and Visualization:** Includes 'GroupBy' (Node 180), 'Analysis' (Node 160), 'Joiner' (Node 164), 'Histogram' (Node 159), 'Interactive Table' (Node 166), and 'Interactive Segmentation View' (Node 165).

Additional components include a 'RowID' node (Node 174) and a 'Views on data' section. The interface also features a 'KNIME Explorer' on the left, a 'Node Repository' on the bottom left, and a 'Console' at the bottom right showing various warning messages.

**GroupBy**

Groups the rows of a table by the unique values in the selected columns. A row is created for each unique value group of the selected column(s). The remaining rows are aggregated by the defined method. The output table therefore contains one row for each existing value combination of the selected group column(s).

To change the aggregation method of more than one column select all columns to change, open the context menu with a

# 3. MATLAB for image processing

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- MATLAB is not free, but many academic institutions have licenses
- Much quicker and easier to prototype new algorithms in MATLAB than e.g. java or C++
- MATLAB is interactive, can use Bioformats to open images, and has an extremely powerful image processing toolbox
- MIJ for ImageJ integration, plus DIPimage image processing library (Delft)

# 3. Tracking in MATLAB

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## Custom particle tracker

- Based on Single Particle Tracker from the MOSAIC group (ETH Zurich), which is available as ImageJ plugin and MATLAB code

I. F. Sbalzarini and P. Koumoutsakos. Feature Point Tracking and Trajectory Analysis for

Video Imaging in Cell Biology, Journal of Structural Biology 151(2):182–195, 2005.

- Used MATLAB to build up a custom processing and detection scheme
- See: <http://www.ncbi.nlm.nih.gov/pubmed/21746854>

# 3. Tracking in General

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## Tracking

Most common scheme: process, detect/refine, link, correct

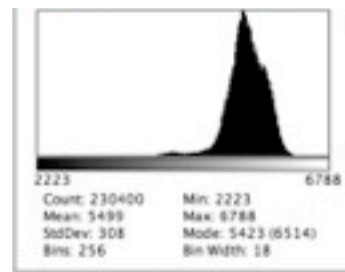
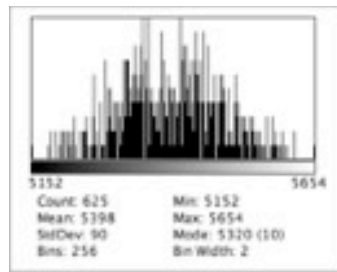
Reliable automatic detection is usually the hard part

Two essential prerequisites:-

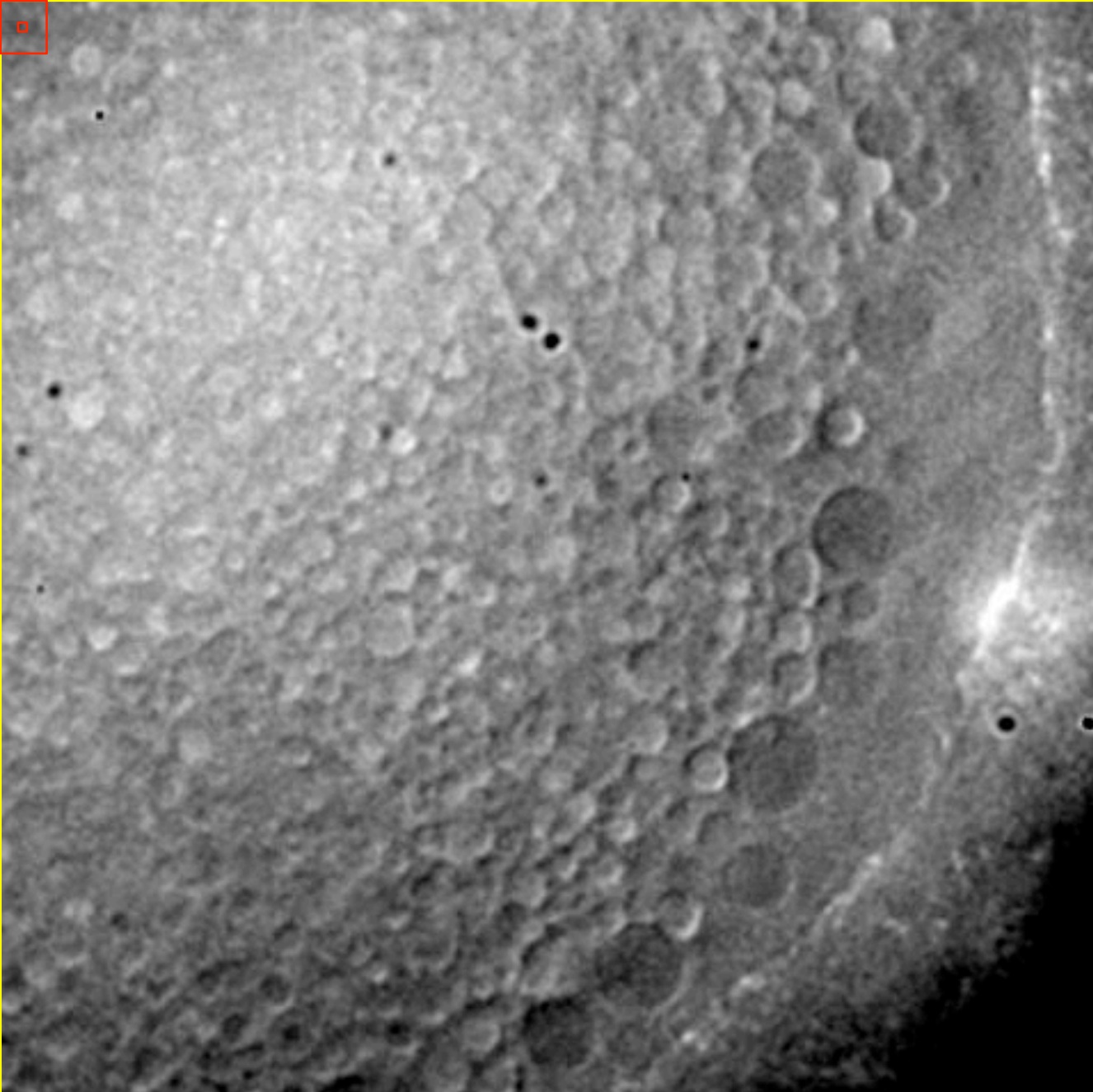
1. contrast-to-noise ratio of  $>4$
2. displacement per. frame less than inter-particle distance



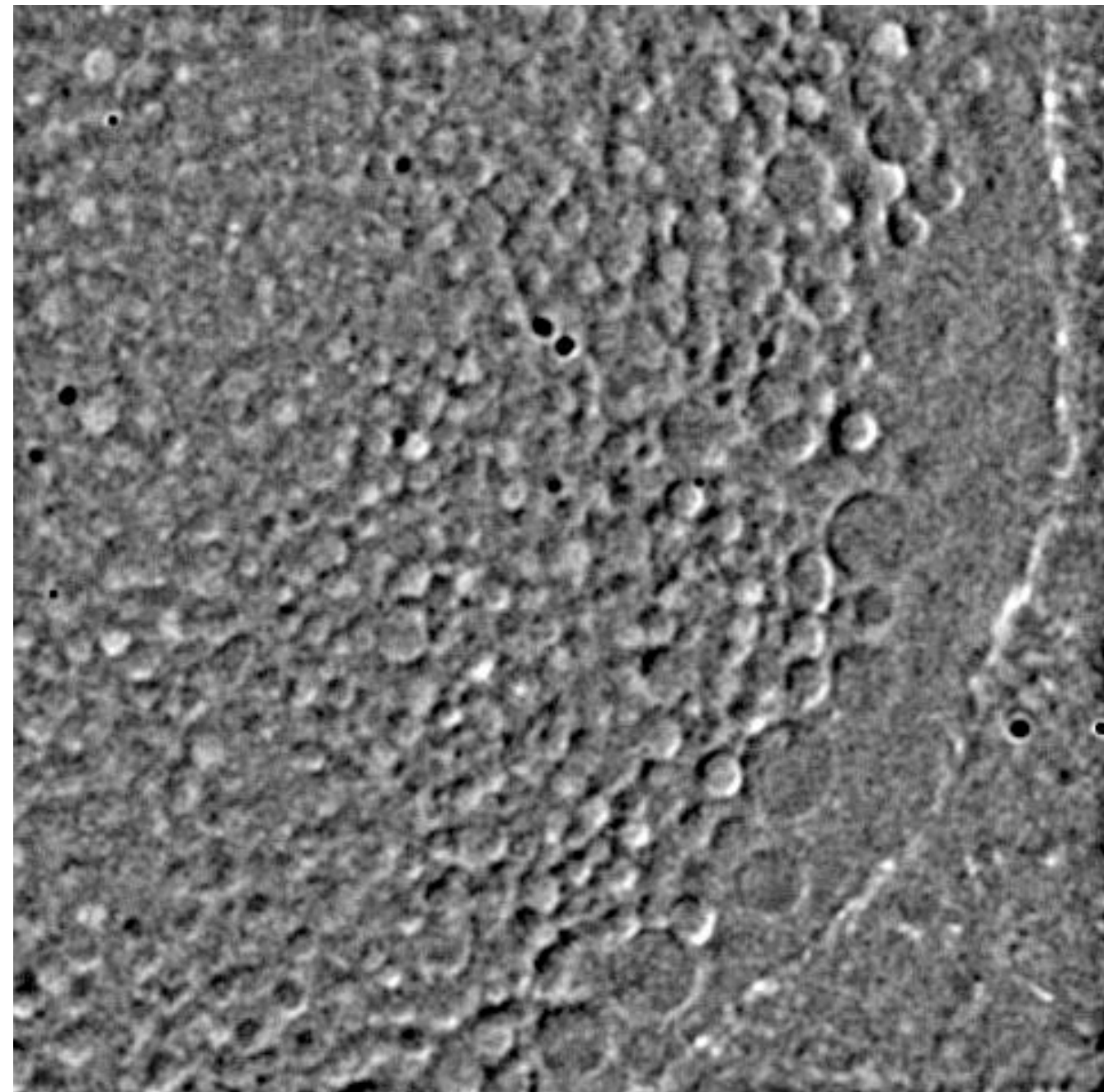
# 1. Image restoration / filtering



**example of a custom intensity transform: scale according to local median**



raw data showing uneven illumination



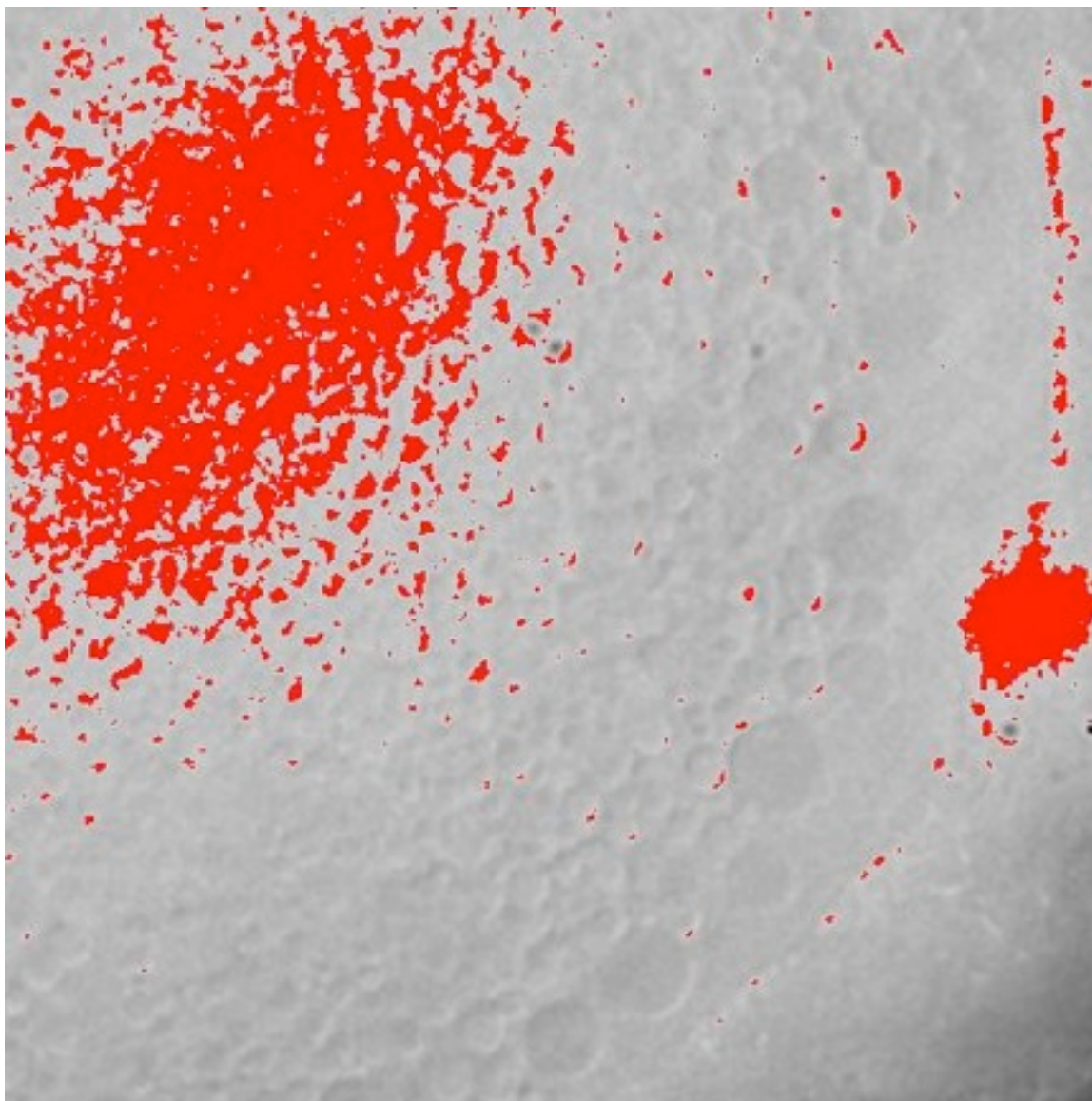
'normalized' image



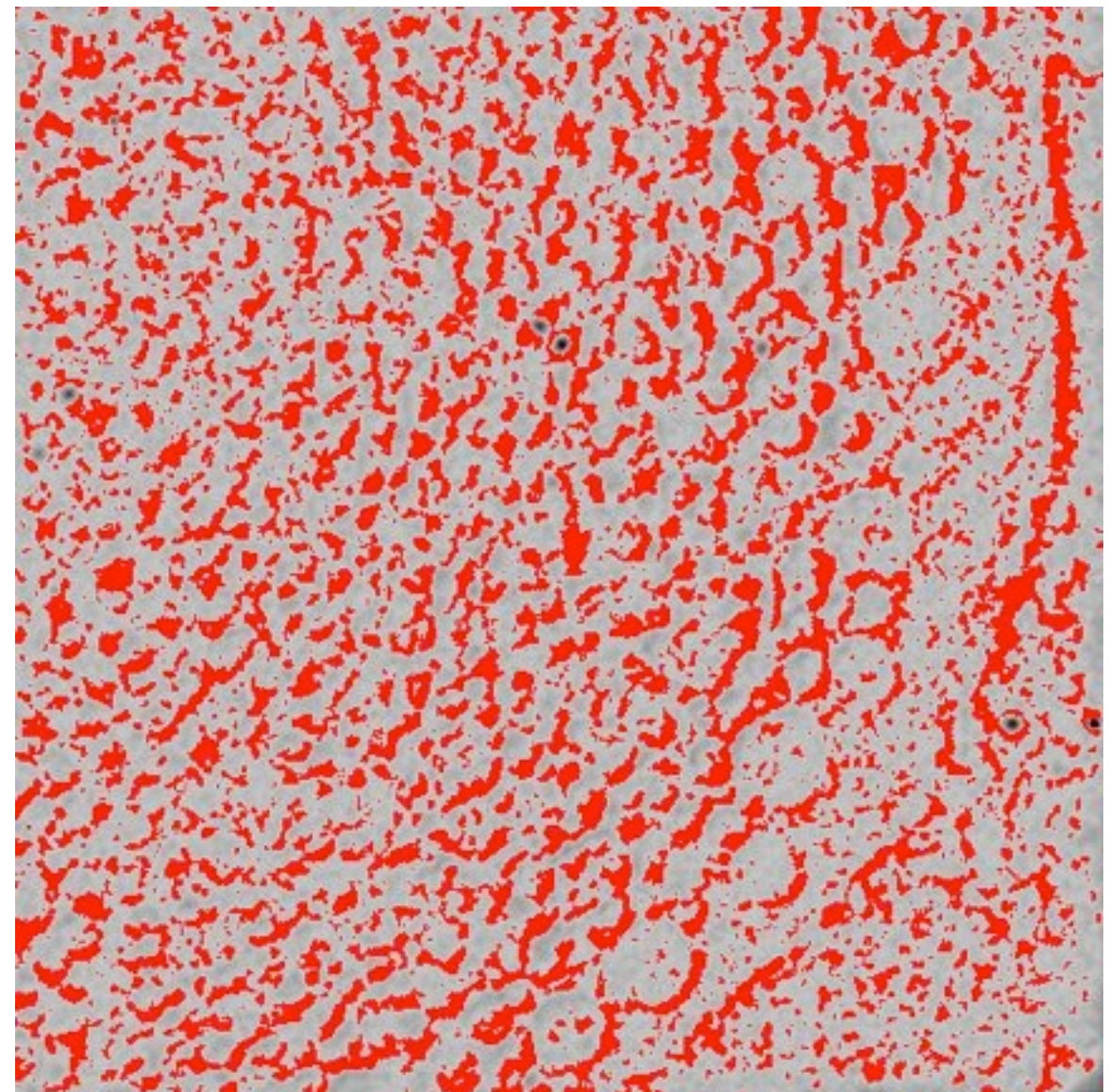
## 2. Feature extraction

### Image segmentation: thresholding

- a global threshold only works if the image is very ‘even’
- using raw data



using ‘normalized’ data

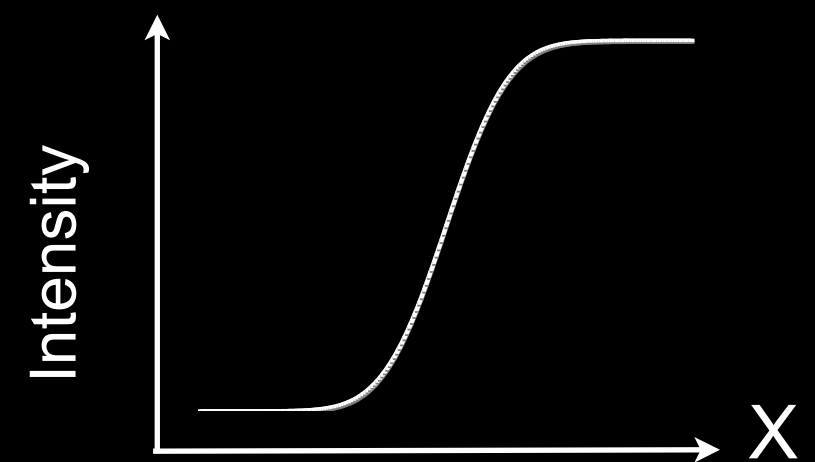
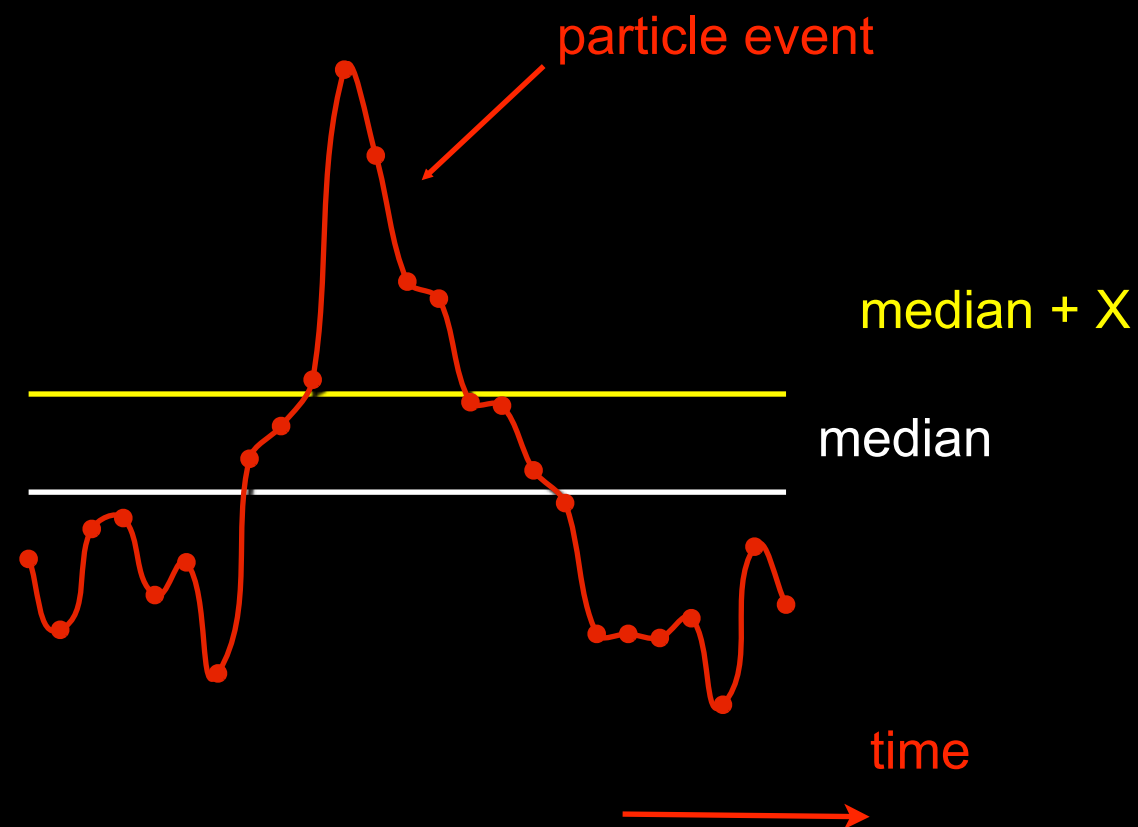


=> ‘adaptive thresholding’, or prior normalization

## 2. Feature extraction

### Image segmentation: identifying 'foreground' features

- easy to implement custom filters in MATLAB, like this **temporal median filter** to identify moving foreground



## 2. Feature extraction

### **Image segmentation: identifying 'foreground' features**

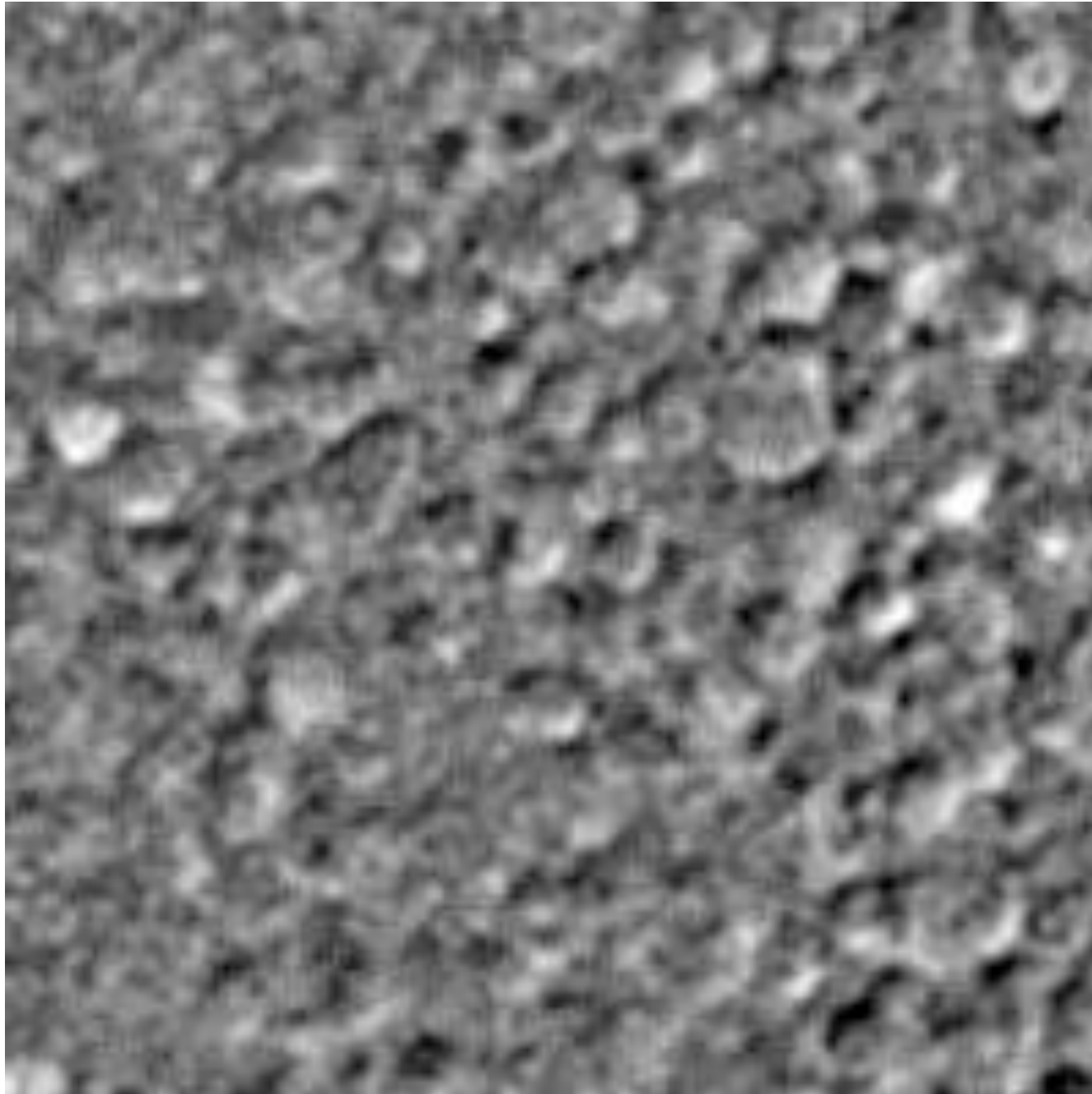
200x200 area, normalized

200x200 area, non-background ('foreground')

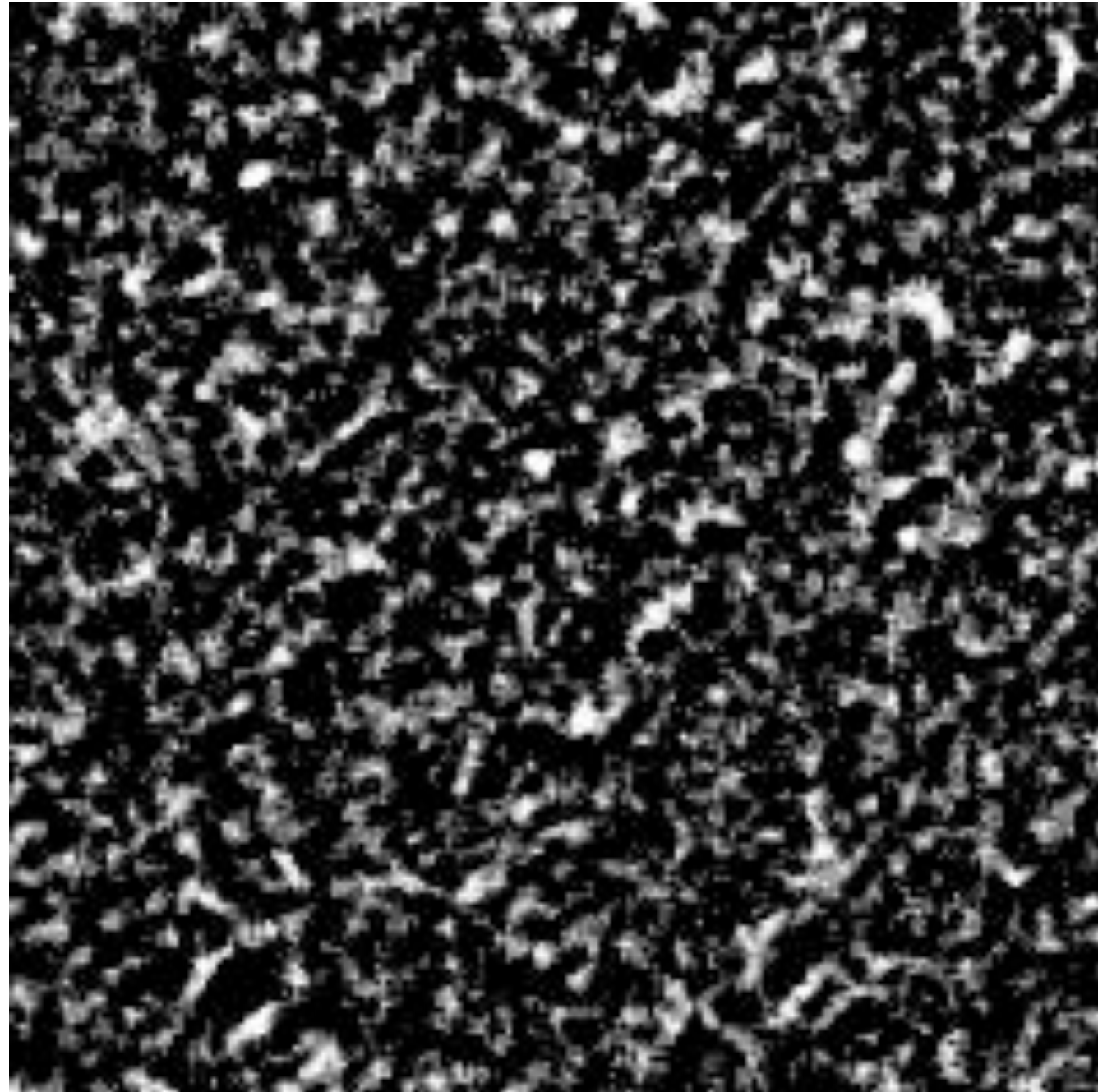


## 2. Feature extraction

### Image segmentation: identifying 'foreground' features



200x200 area, normalized



200x200 area, non-background ('foreground')



## 2. Feature extraction

### Object recognition

- many tools for point, line & edge detection in MATLAB
- generally work by either:
  - applying a mask to find maxima
  - or
  - calculating intensity gradient (steep gradient = edge)

e.g. detection of Haar-like features to find particles

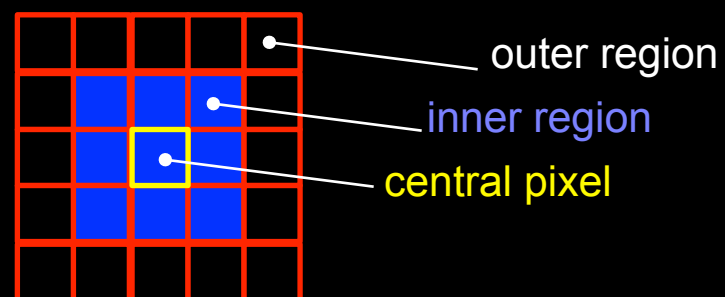
## 2. Feature extraction

### Object recognition

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e.g. detection of Haar-like features to find particles

square Haar-like feature



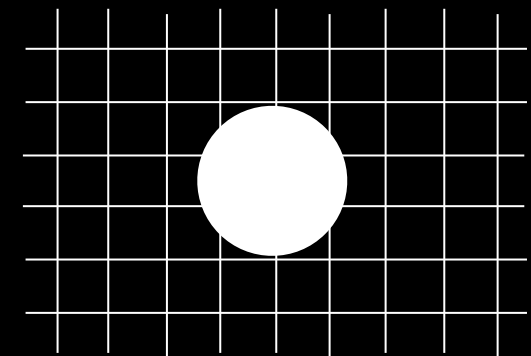
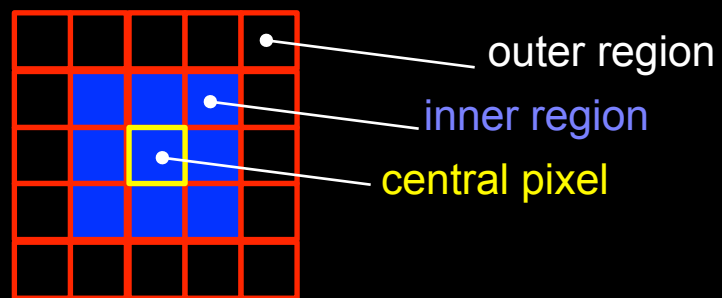
## 2. Feature extraction

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square Haar-like feature



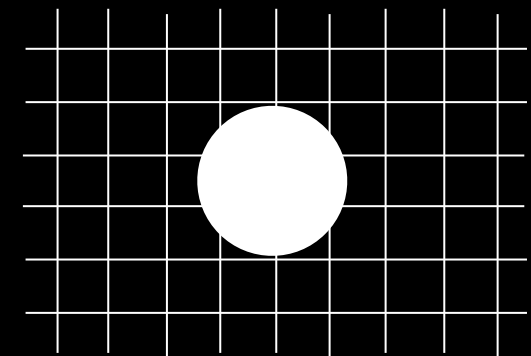
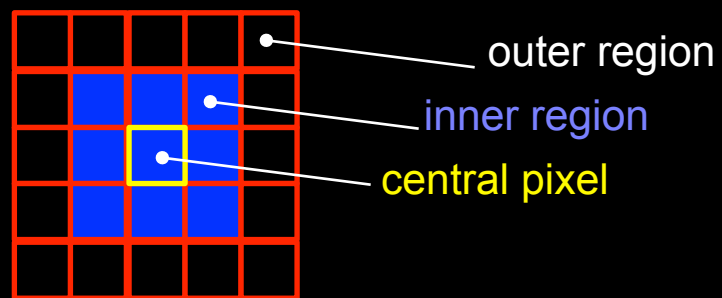
## 2. Feature extraction

### Object recognition

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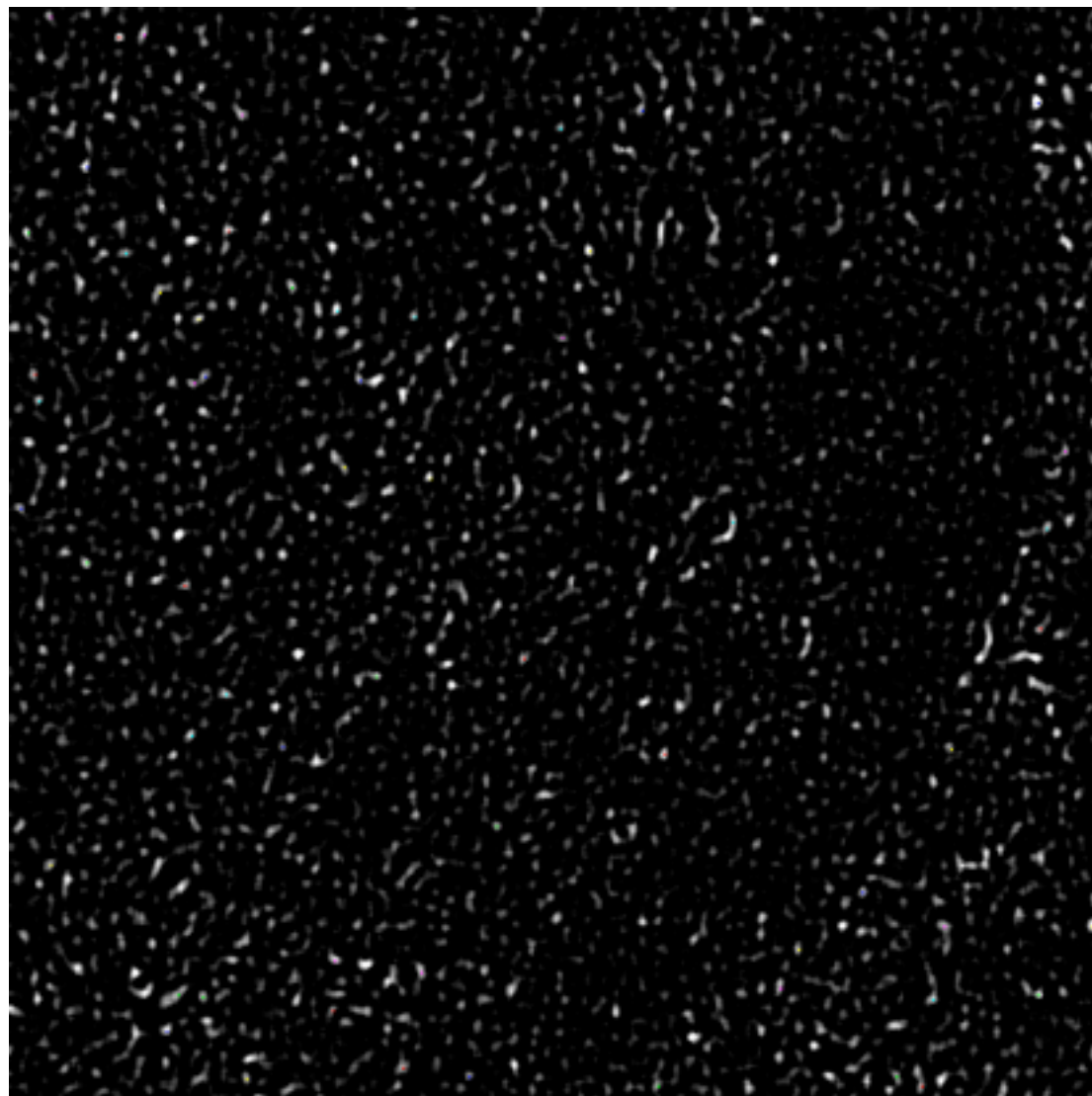
e.g. detection of Haar-like features to find particles

square Haar-like feature

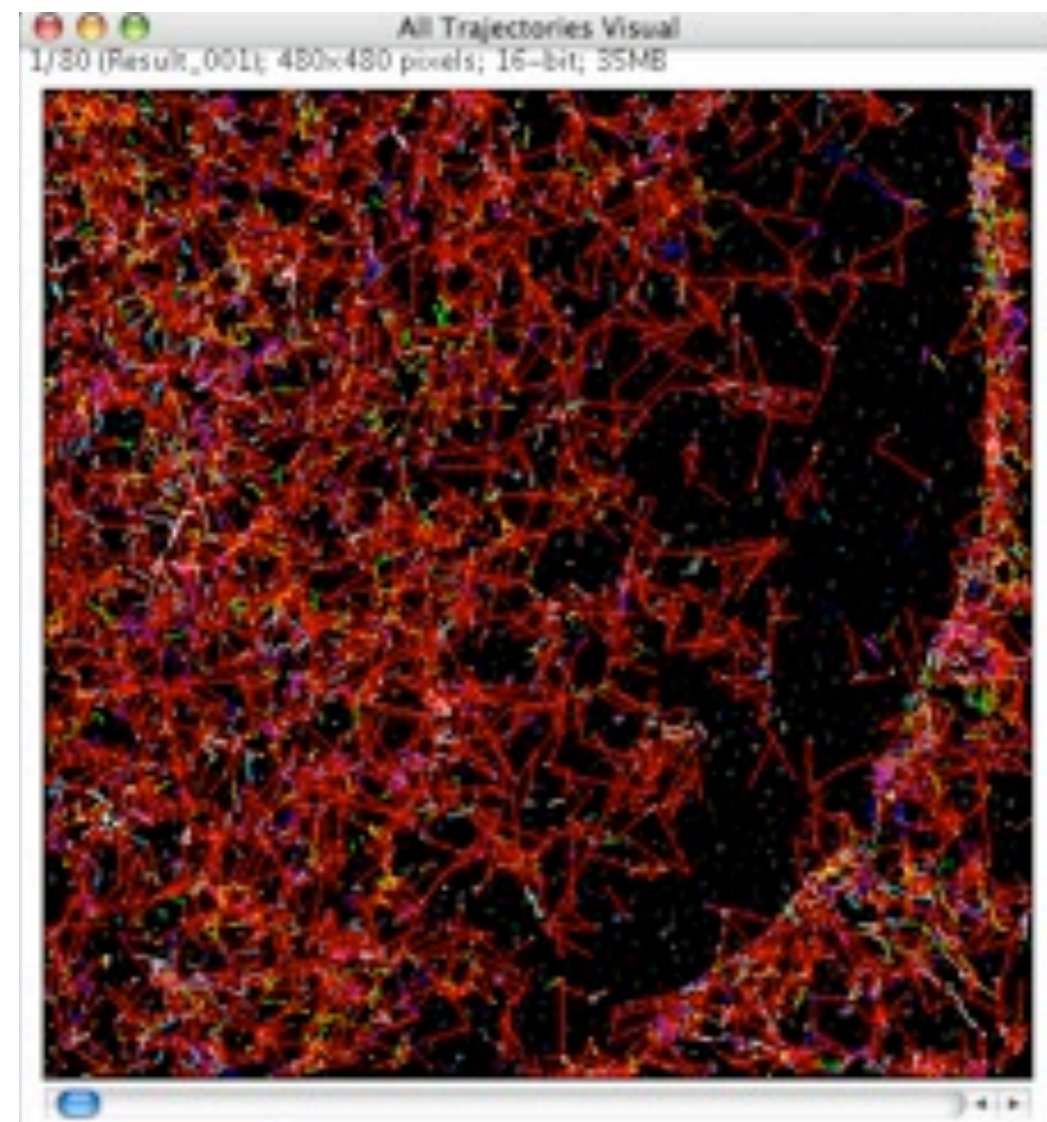


# 2.4. Tracking in MATLAB

## Custom particle tracker



final “particle image” with tracks



MOSAIC imageJ tracker results



# Summary

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- importance of quantitative image analysis
- importance of experimental design – the right data and trade-offs
- overview of processing / analysis concepts
- survey of available software – choosing the right tool (default to Fiji/ImageJ)
- automation: is it necessary? if so, don't be afraid to try / ask for help
- keep data secure, well-organized and annotated
- feedback – problems you are interested in that I haven't covered
- Demo tomorrow – please ask questions

# 3. Summary (7 slides)

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## Overview of image processing & analysis

- importance of experimental design & optimization (identify problems early)
- summary of software / choosing the right tool for the job
- processing / analysis tips
- automation: is it necessary? if so, ask / don't be afraid to try
- keep data secure, well-organized and annotated
- feedback – problems you are interested in that I haven't covered
- Demo