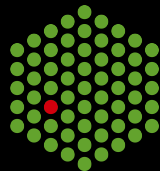


Image Analysis in Cell and Developmental Biology

EMBL



Centre for Molecular and Cellular Imaging (CMCI)

EMBL Heidelberg, Germany

20120807 Biocity, Turku

Kota Miura

Experiment → Image Acquisition → Processing → Analysis

Aug 6, 2012

9:00	Lecture Image Analysis: an overview	
10:30	Lecture: BioimageXD, Image Analysis Software	
13:00	Lecture: Microscopy	

Aug. 7, 2012

9:00 - 12:00	Practical: Microscopy / ImageJ (sub-groups)	1.5 hrs
13:00 – 14:30	Practical : BioimageXD	
14:30 – 17:00	Practical: ImageJ	2.5 hrs

Aug. 8, 2012

9:00 – 11:30	Lecture: Microscopy	
13:00 – 15:00	Practical: Microscopy / BioImageXD (sub-groups)	
15:00 – 17:00	Practical: ImageJ	2 hrs

Aug. 9, 2012

9:00 – 12:00	Practical: ImageJ Macro programming	3 hrs
13:00 – 14:30	Practical: BioImageXD	
14:30 – 17:00	Practical: Microscopy	

EMBL Heidelberg



Rainer Pepperkok (CBB, ALMF)

Heiko Runz (Uni. Heidelberg)

Fatima Verissimo (CBB)

Francois Pouthas (became an Artist)

Emmanuel Reynaud (now in Dublin)

Cihan Cetin (now in Pasadena)

Clemens Grabher (now in Karlsruhe)

Adam Cliffe (now in Singapore)

Tina Diamantara (DB)

Stefano De Lenzis (DB)

Darren Gilmour (DB)

Gulcin Cakan (now in Boston)

Petra Haas (Heidelberg)

Asifa Akhtar (now in Freiburg)

Ritsuko Suyama (now in Kyoto)

Janina Karres

Steve Cohen (Singapore)

EMBL ALMF

Jens Rietdorf (now in Munich)

Timo Zimmermann (now in Barcelona)

Stefan Terjung (ALMF)

Univ. Munich (LMU)

Florian Siegert

Charles David

Osaka Univ

Satoshi Ogiwara

cmci.embl.de



Daily Life

Images (pictures, movies)

→ interpretation (“Fantastic!”)

Daily Life

Images (pictures, movies)

→ interpretation (“Fantastic!”)

Scientific Research

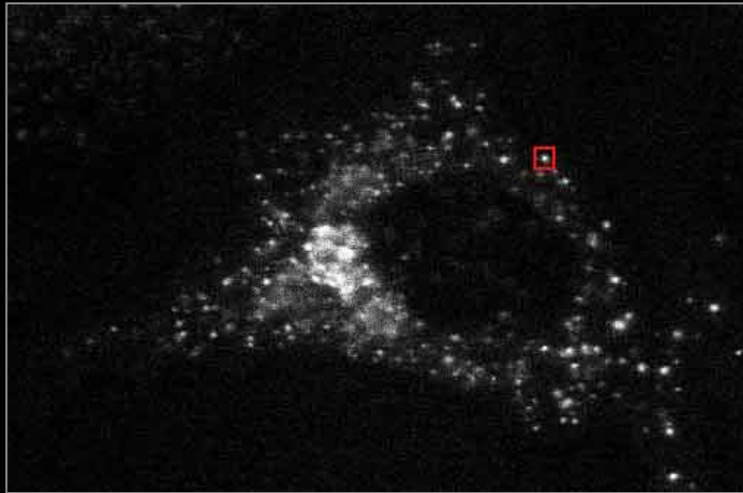
Images (static, dynamic)

→ extract numerical parameters (Results)

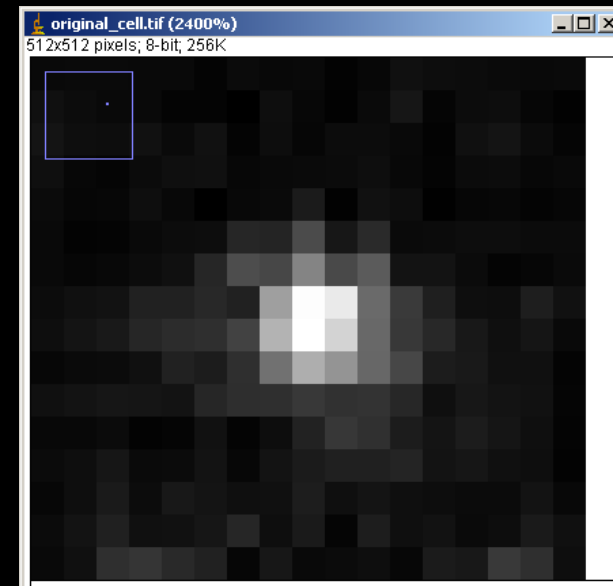
→ interpretation (Discussion)

See Rossner and Yamada (2004; pdf) for more about ‘**manipulation**’

Numerical Nature of Digital Images



Fatima Verissimo



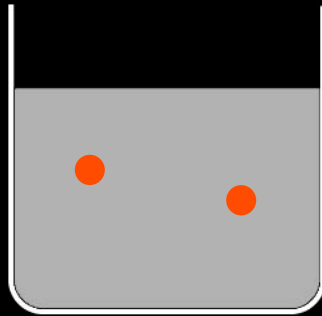
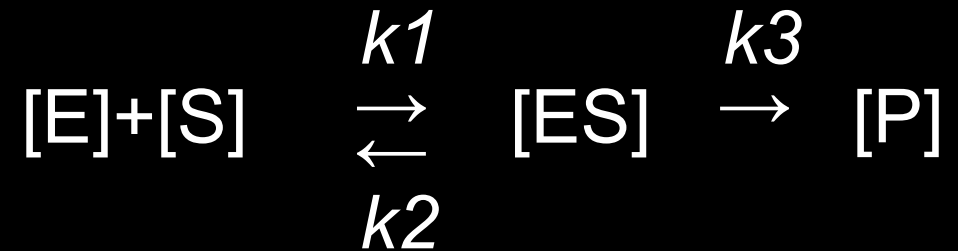
Pixel intensity



Protein density

19	22	25	19	0	0	10	37	0	15	20
29	16	8	11	16	16	13	12	0	13	17
14	5	1	11	29	36	23	5	5	11	13
0	5	19	33	54	70	62	43	19	12	13
12	29	46	70	115	152	140	103	38	19	18
58	56	57	92	176	238	210	140	61	30	28
63	58	50	82	175	249	217	135	82	42	38
34	42	40	61	138	208	186	115	95	50	45
17	38	51	51	60	78	78	64	42	29	29
8	21	22	15	26	53	67	63	49	37	37
7	18	17	6	14	37	49	43	42	29	29
13	31	38	30	31	40	34	17	25	12	12

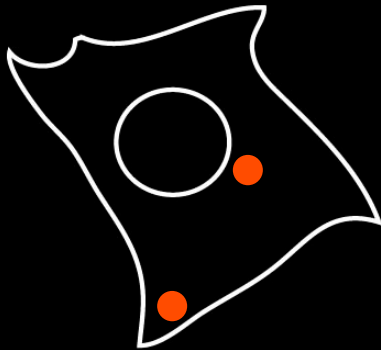
Biochemistry (solution)



$$B([E], [S], [ES], [P], t)$$

Topo-biochemistry

(in vivo)



$$B([E], [S], [ES], [P], t, x, y, z)$$

Extracting Parameters out of Images: Overview

Experiment → Image Acquisition → Processing → Analysis

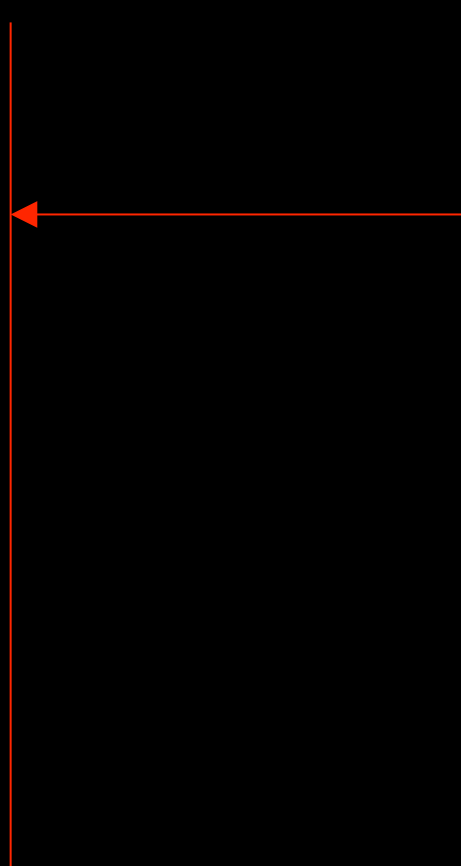
Extract numbers and interpret

Static images

- Intensity Measurement
- Morphometry
- Distribution
- Colocalization

Image sequences (dynamics)

- Intensity changes
- Shape changes
- Position changes:
 - kymograph
 - tracking
- Distribution changes
- Optical flow estimation



Static Images: Intensity Measurement

Static images

Intensity Measurement →

Morphometry

Distribution

Colocalization

Intensity Difference

Intensity Profile

Image sequences

Intensity dynamics

Shape dynamics

Position dynamics :

kymograph

tracking

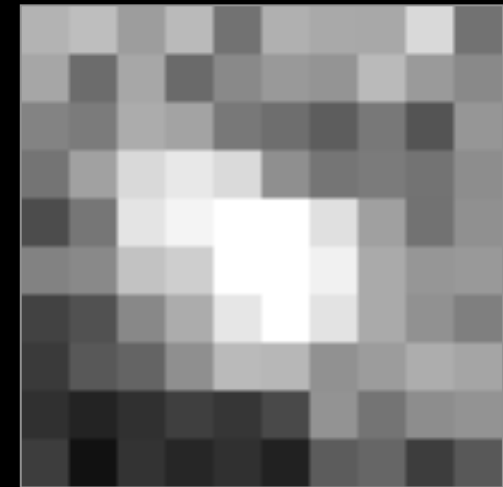
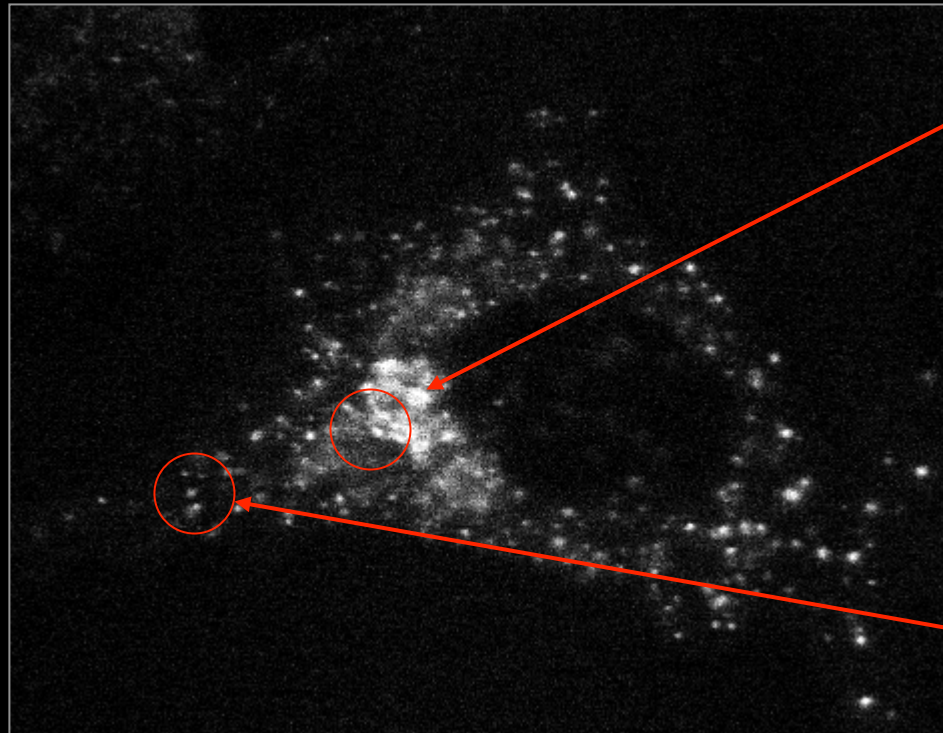
Distribution dynamics

Optical Flow Estimation

Static Images: Intensity Measurement

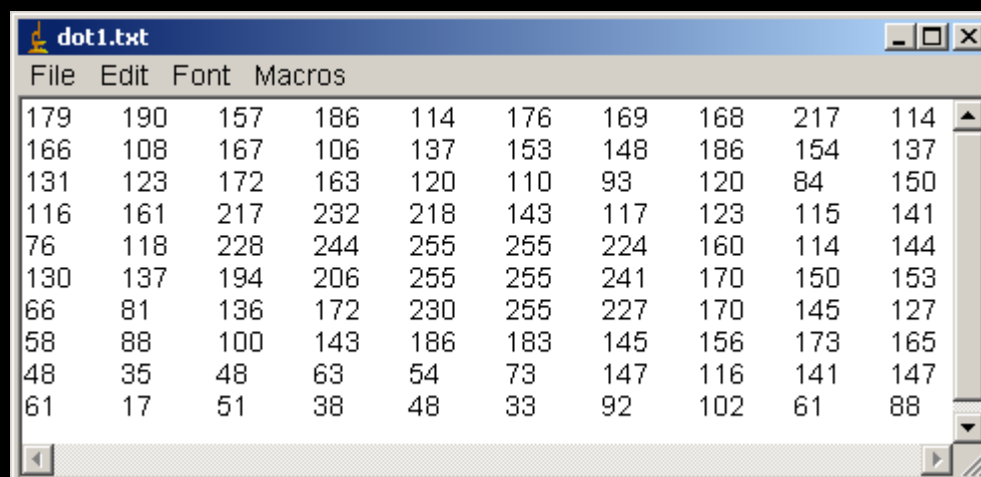
-Intensity Difference

-Intensity Profile

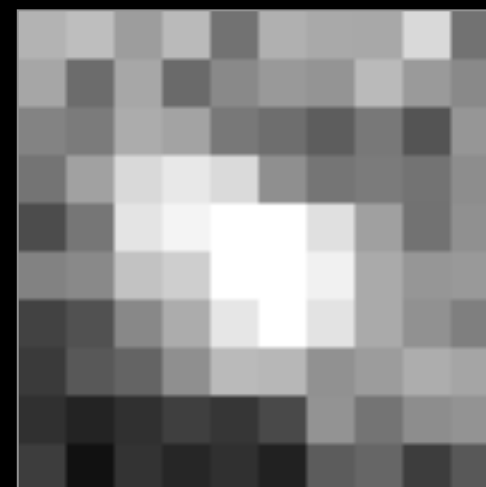


Fatima Verissimo

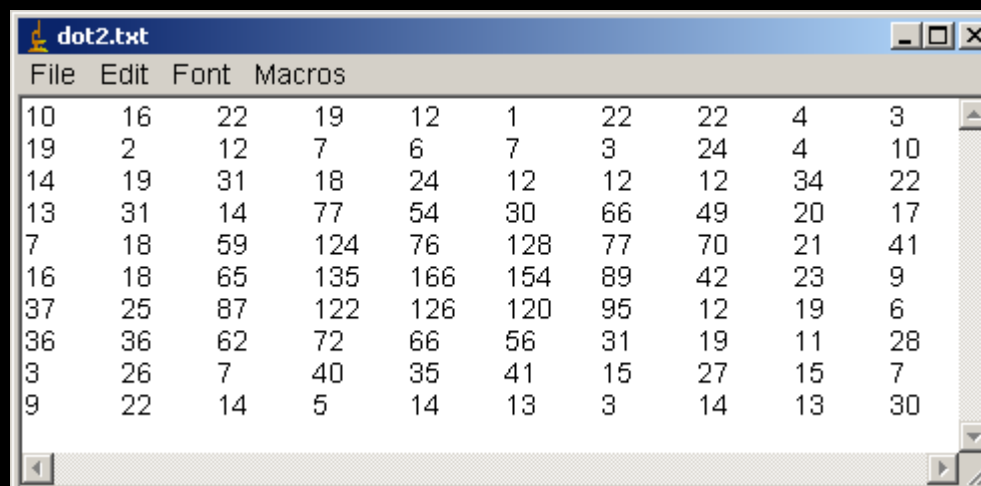
Static Images: Intensity Measurement



File	Edit	Font	Macros						
179	190	157	186	114	176	169	168	217	114
166	108	167	106	137	153	148	186	154	137
131	123	172	163	120	110	93	120	84	150
116	161	217	232	218	143	117	123	115	141
76	118	228	244	255	255	224	160	114	144
130	137	194	206	255	255	241	170	150	153
66	81	136	172	230	255	227	170	145	127
58	88	100	143	186	183	145	156	173	165
48	35	48	63	54	73	147	116	141	147
61	17	51	38	48	33	92	102	61	88



Mean Pixel Intensity: 141.6 ± 58.0 (saturated)



File	Edit	Font	Macros						
10	16	22	19	12	1	22	22	4	3
19	2	12	7	6	7	3	24	4	10
14	19	31	18	24	12	12	12	34	22
13	31	14	77	54	30	66	49	20	17
7	18	59	124	76	128	77	70	21	41
16	18	65	135	166	154	89	42	23	9
37	25	87	122	126	120	95	12	19	6
36	36	62	72	66	56	31	19	11	28
3	26	7	40	35	41	15	27	15	7
9	22	14	5	14	13	3	14	13	30

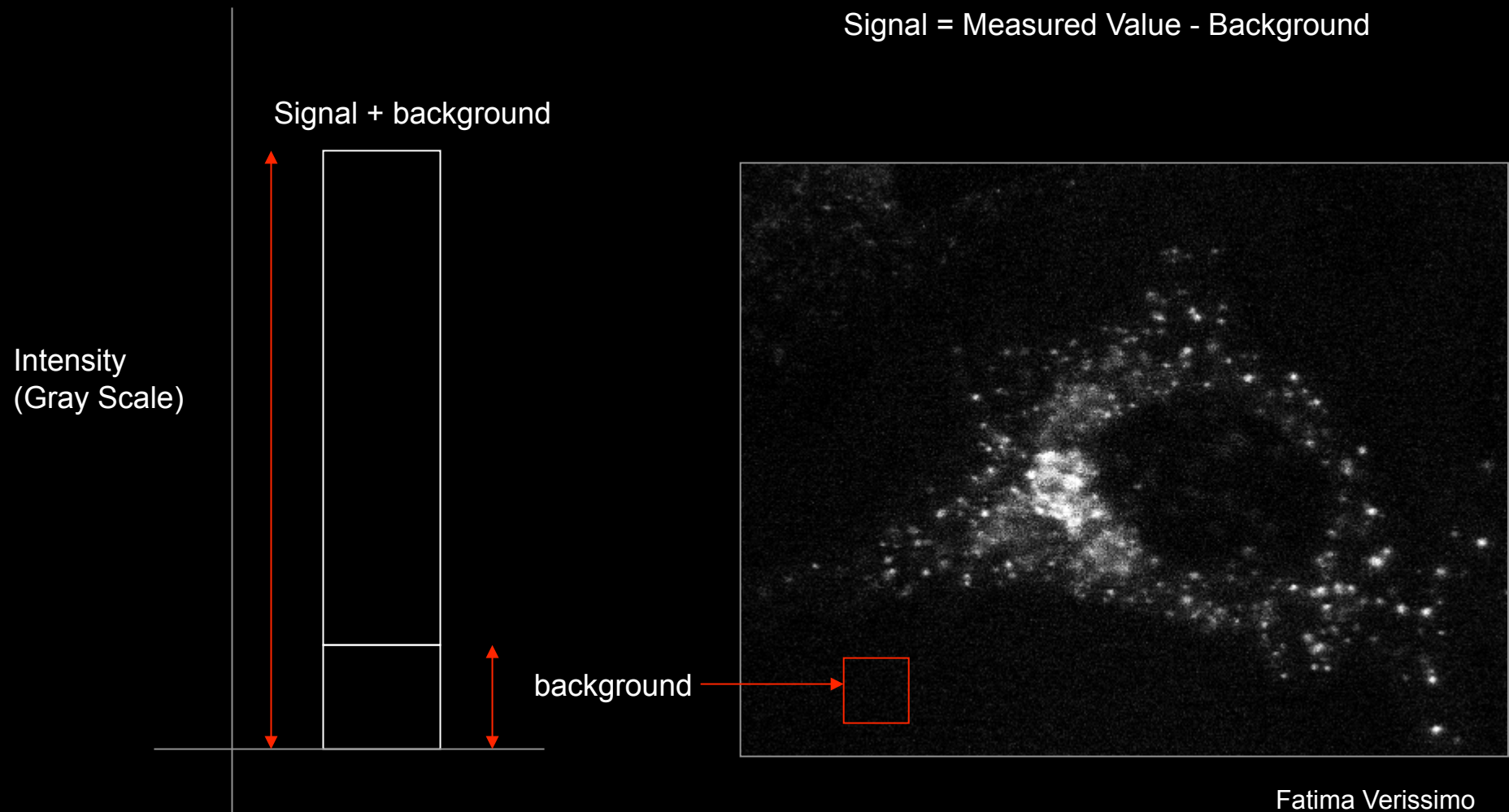


Mean Pixel Intensity: 35.5 ± 36.8

Static Images: Intensity Measurement

Background Subtraction

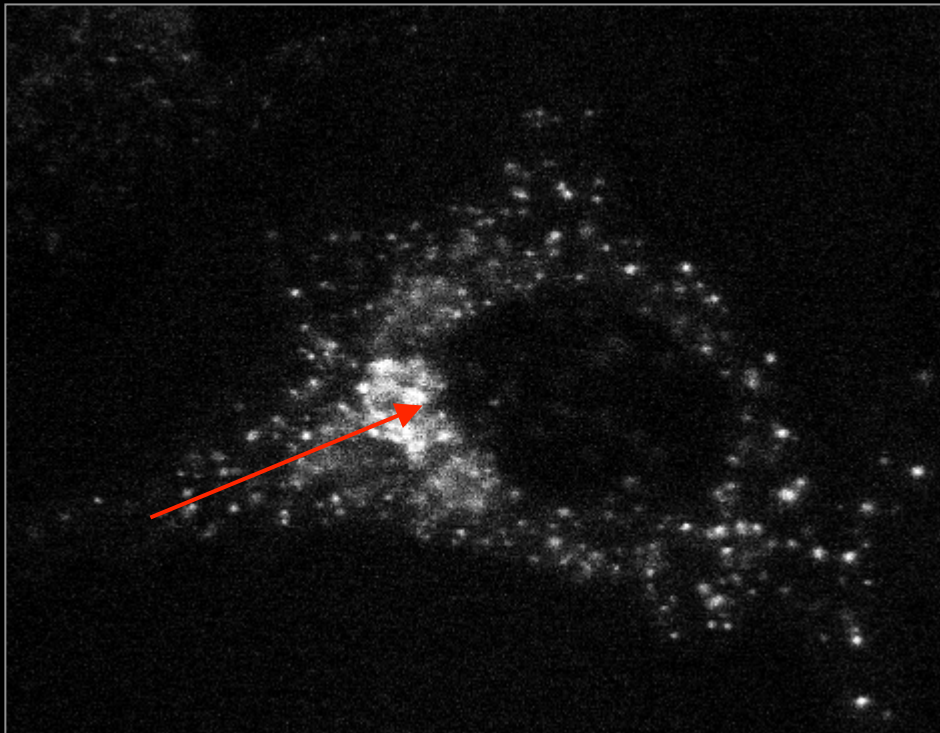
$$\text{Signal} = \text{Measured Value} - \text{Background}$$



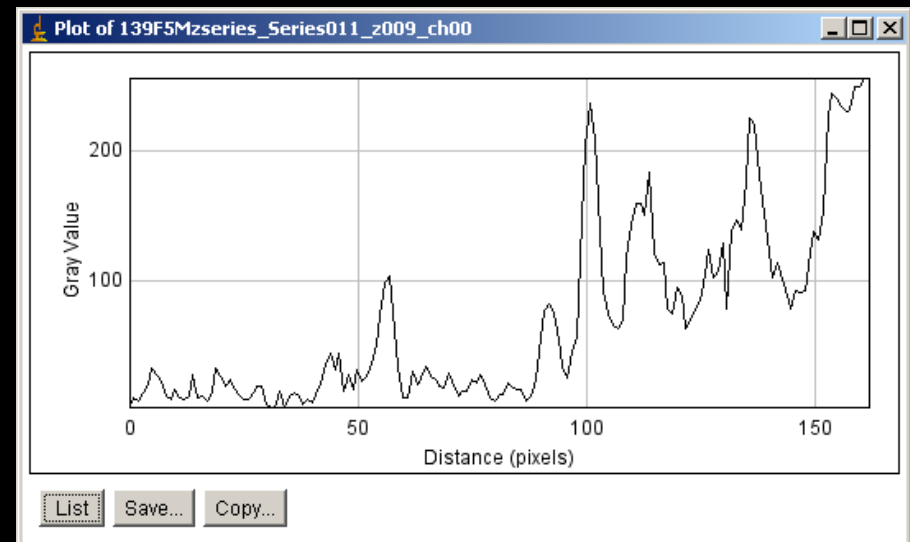
Static Images: Intensity Measurement

-Intensity Difference

-Intensity Profile



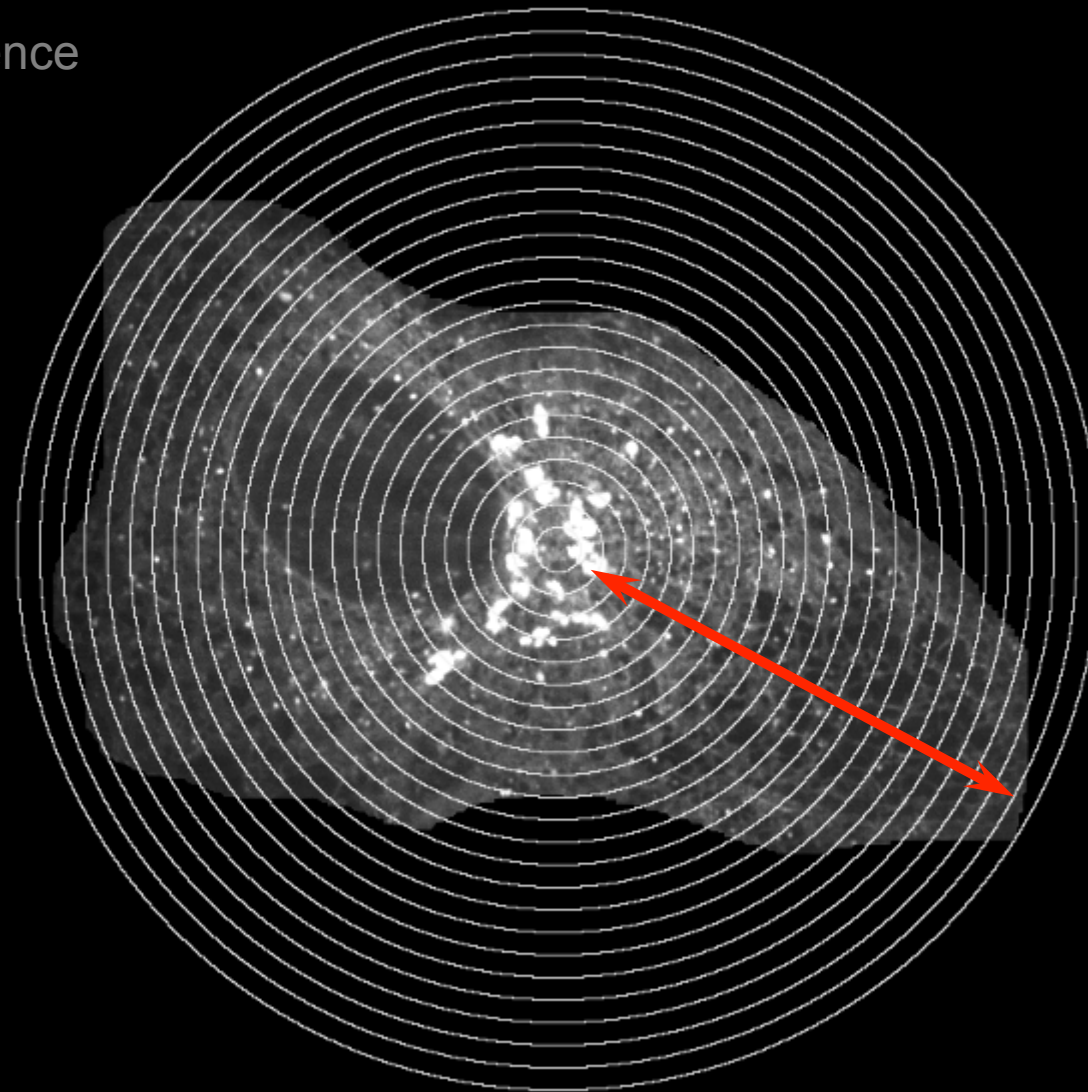
Fatima Verissimo



Static Images: Intensity Measurement An Example

-Intensity Difference

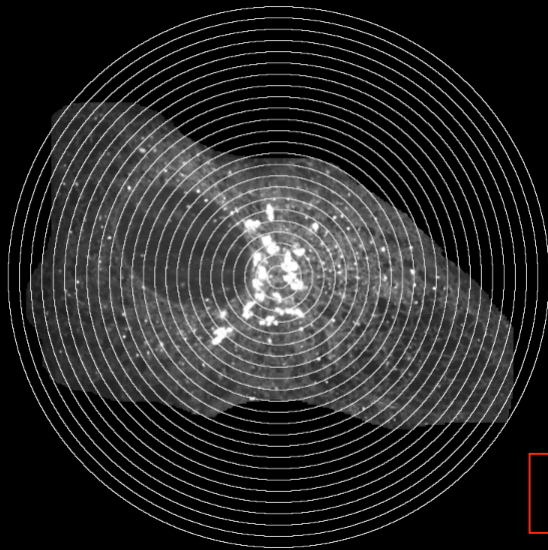
-Intensity Profile



Static Images: Intensity Measurement An Example

-Intensity Difference

-Intensity Profile



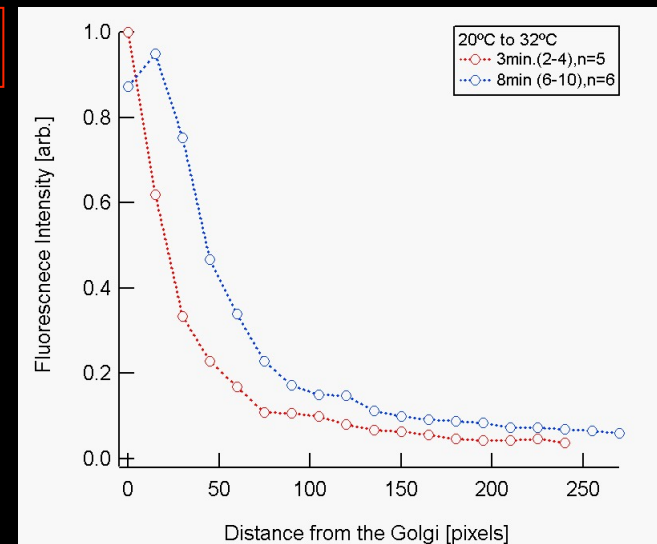
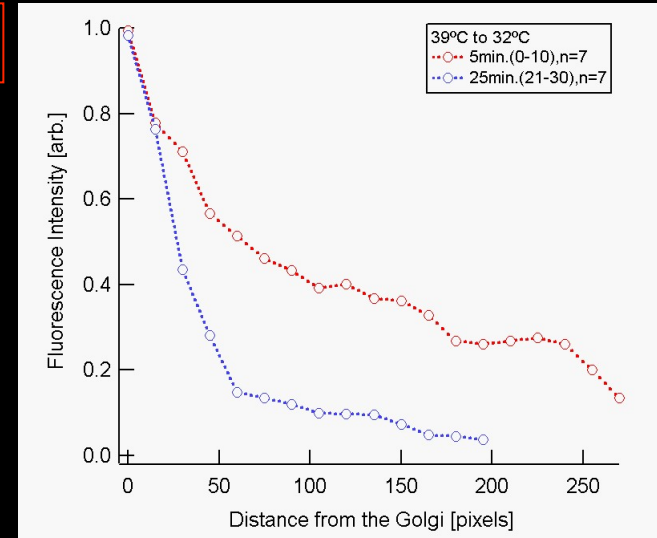
ER

Golgi

Golgi

PM

Radial Intensity Distribution



Static Images: Morphometry

Static images

Intensity Measurement
Morphometry
Distribution
Colocalization

Length
Area
Perimeter
Circularity

Image sequences

Intensity dynamics
Shape dynamics
Position dynamics :
 kymograph
 tracking
Distribution dynamics
Optical Flow Estimation

Static Images: Morphometry

Length

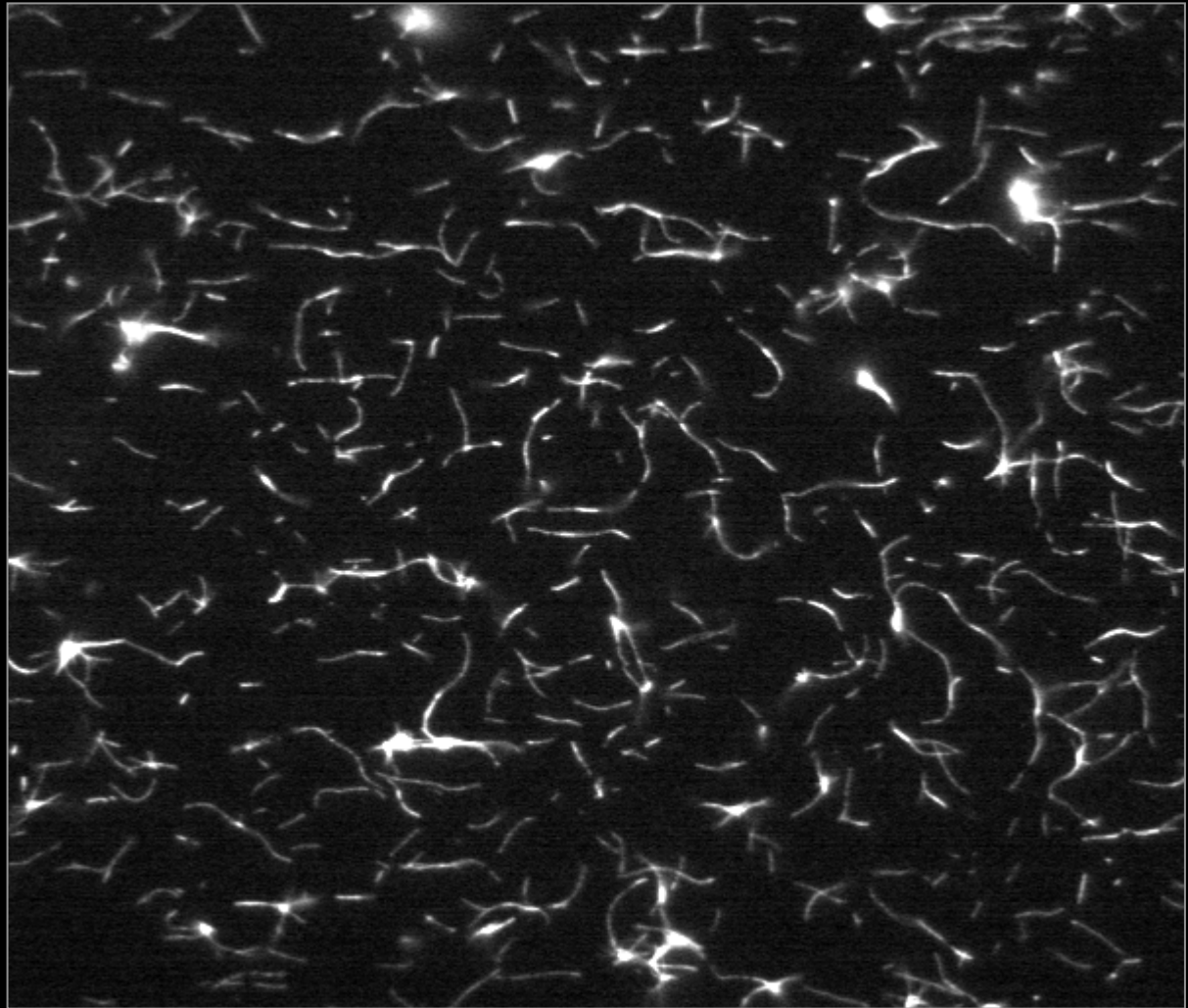
straight: (x_1, y_1) to (x_2, y_2)

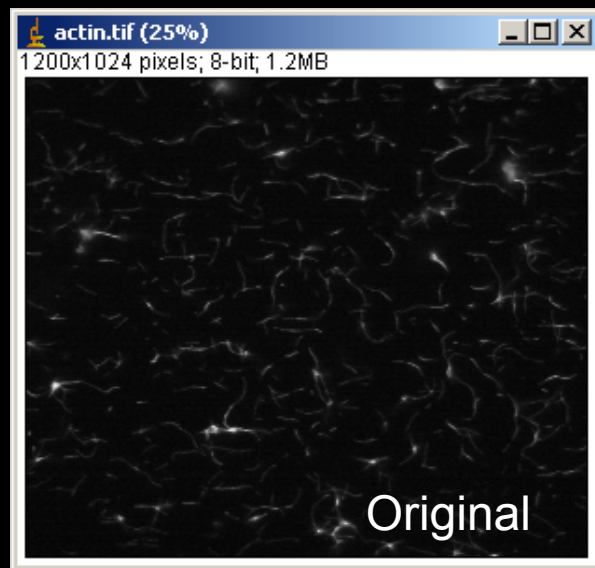
curves

Area

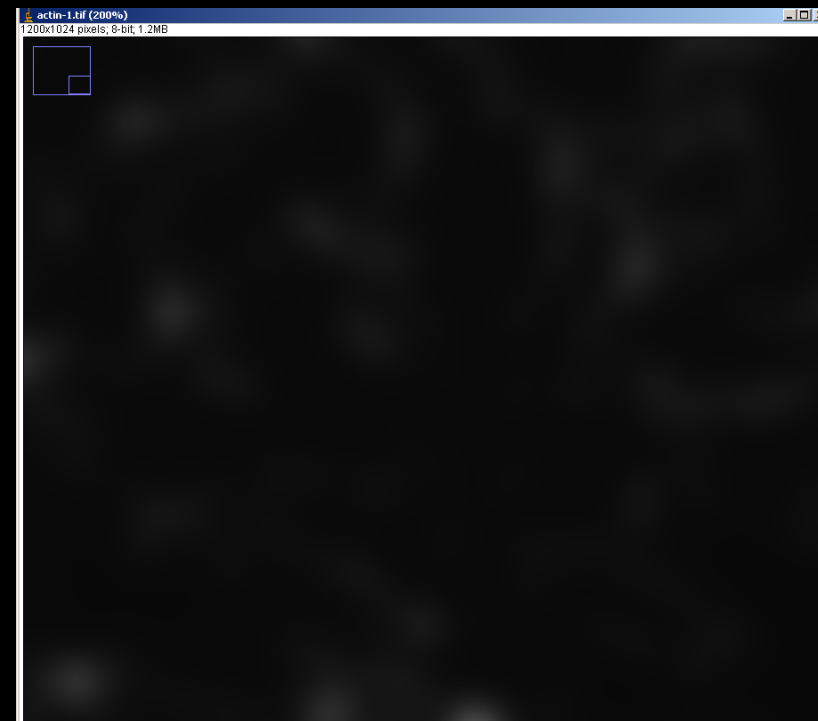
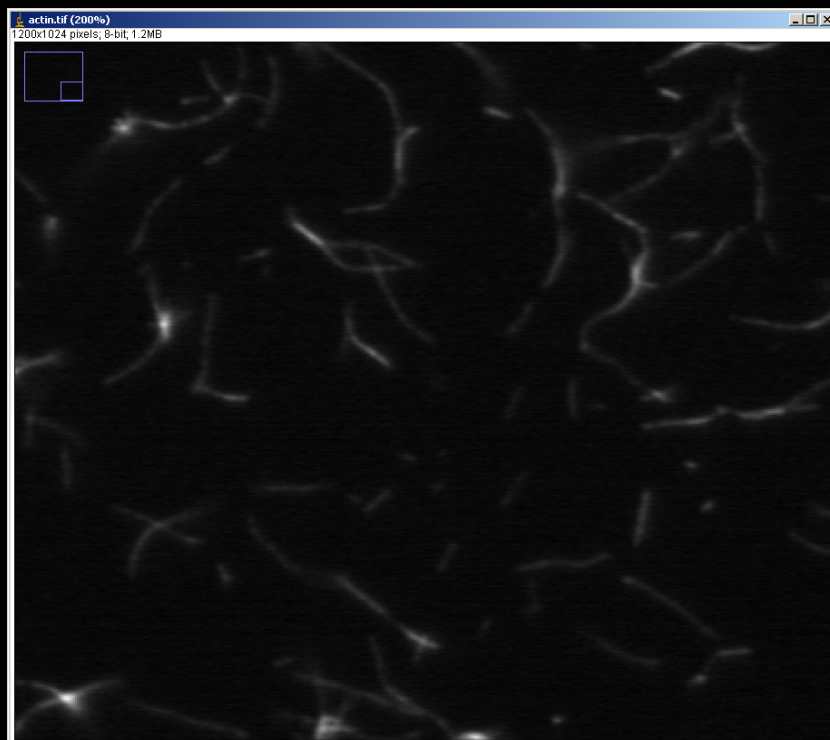
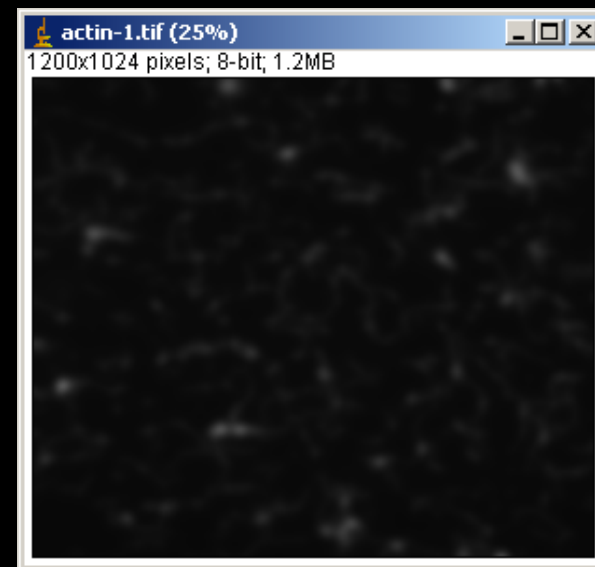
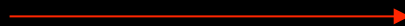
Perimeter

Circularity

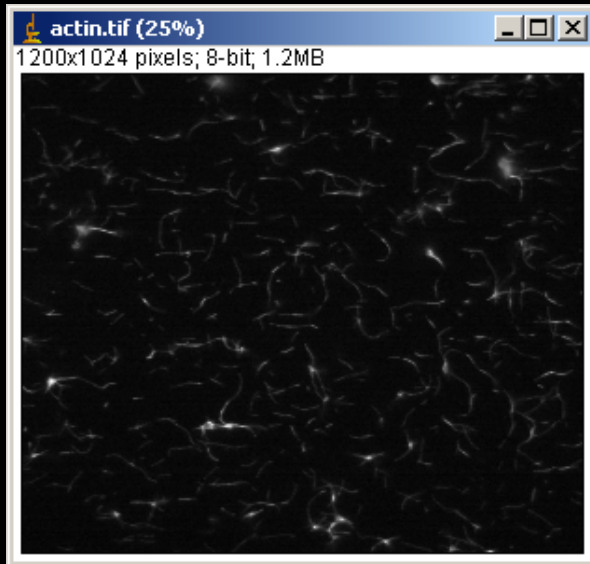




Gaussian Blur (10)



Remove local difference

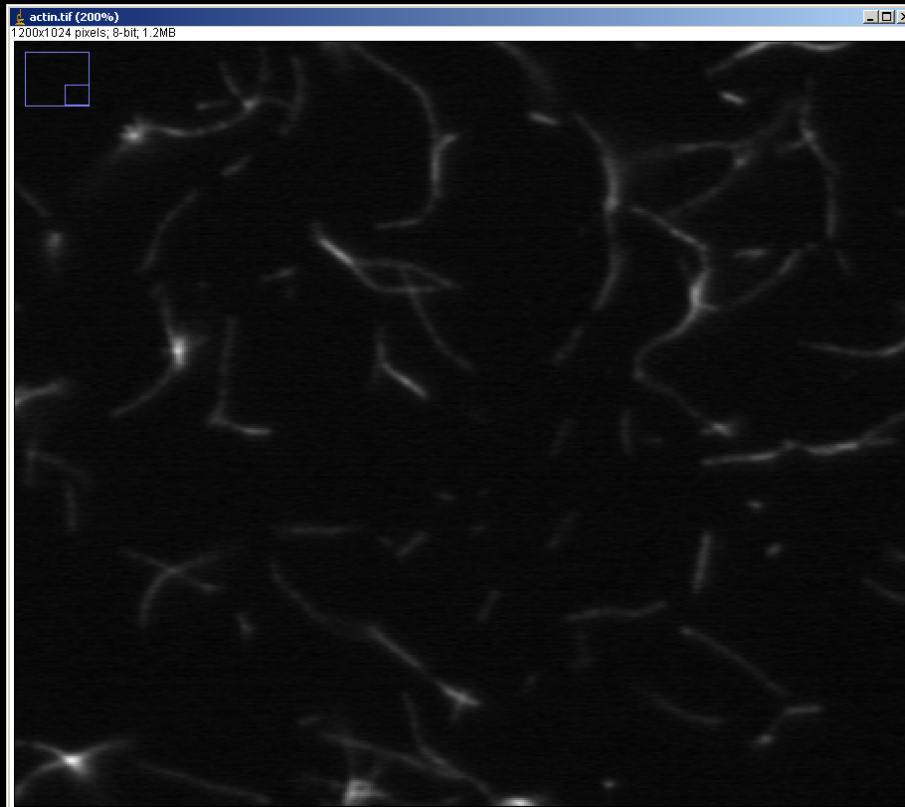
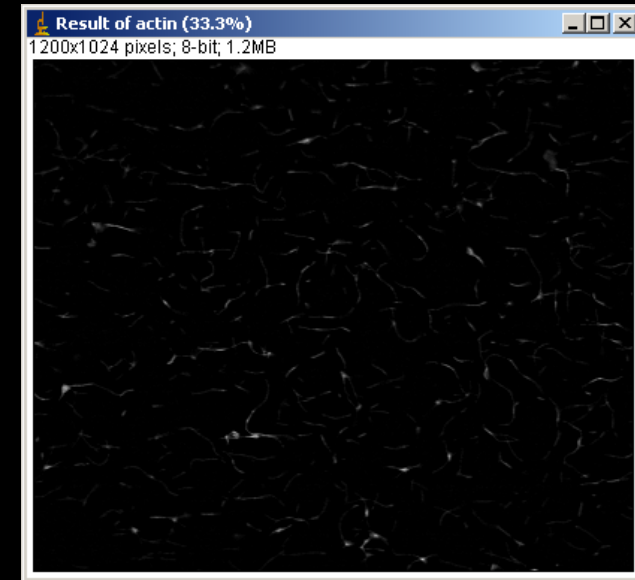


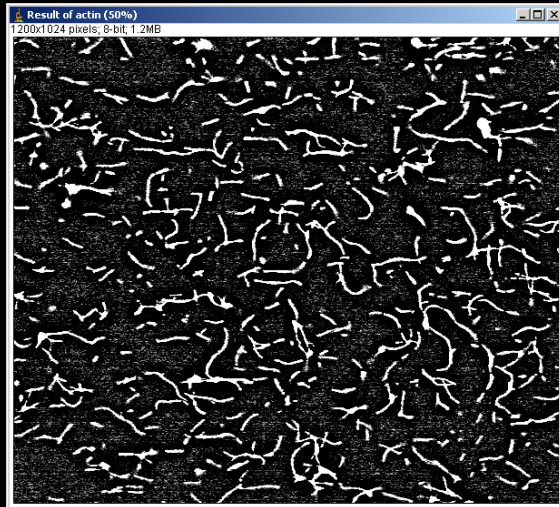
Original

Subtract Blurred

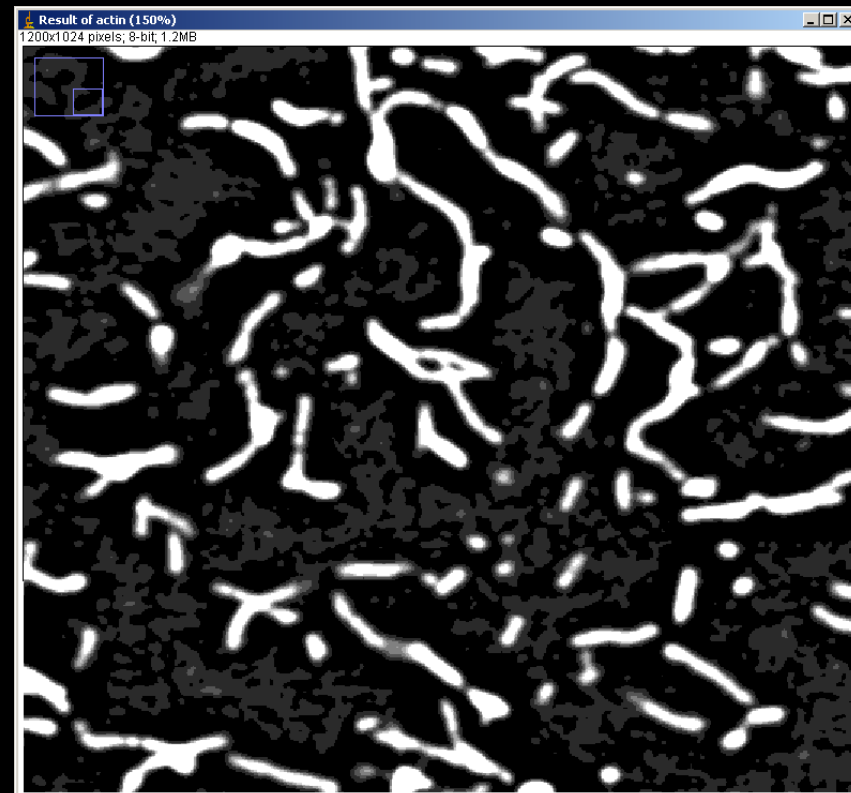
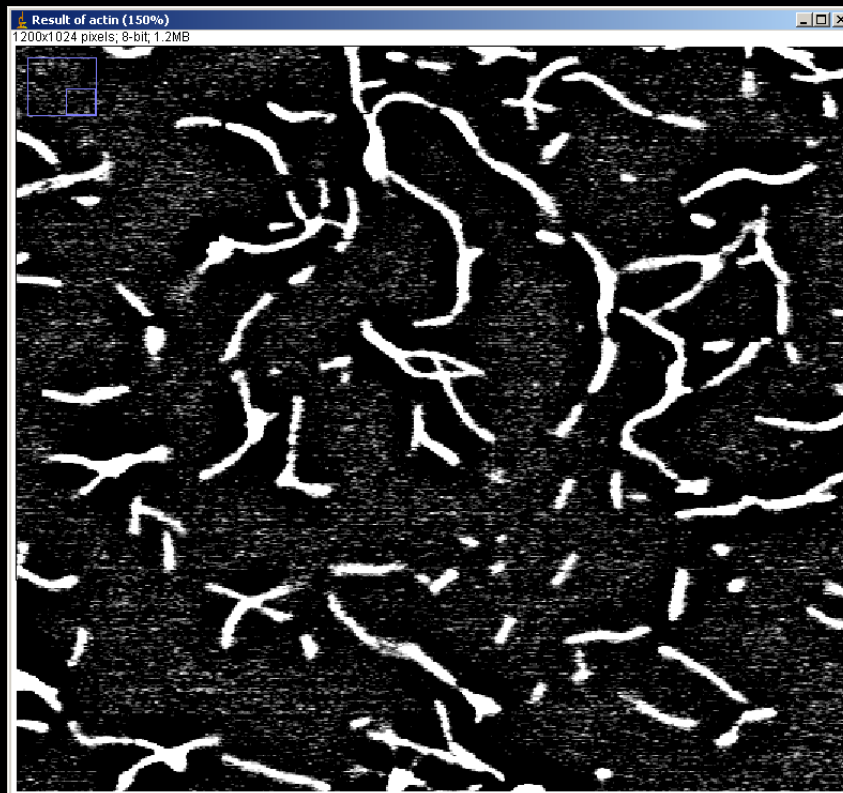
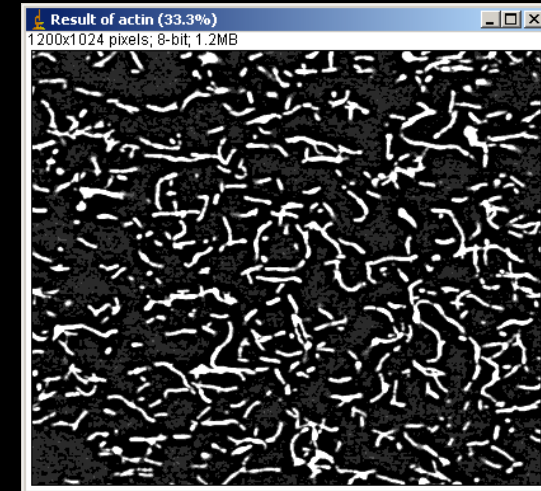



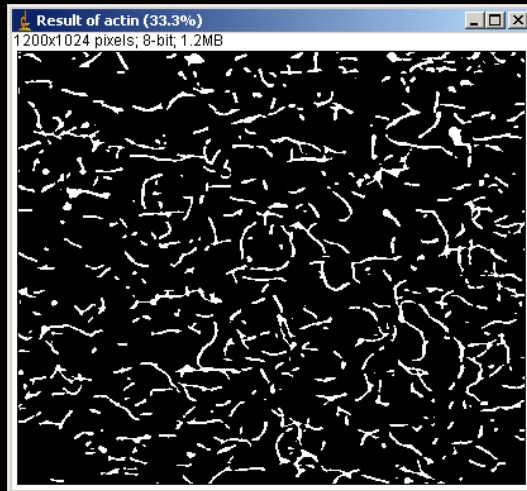
Local Difference
Suppressed





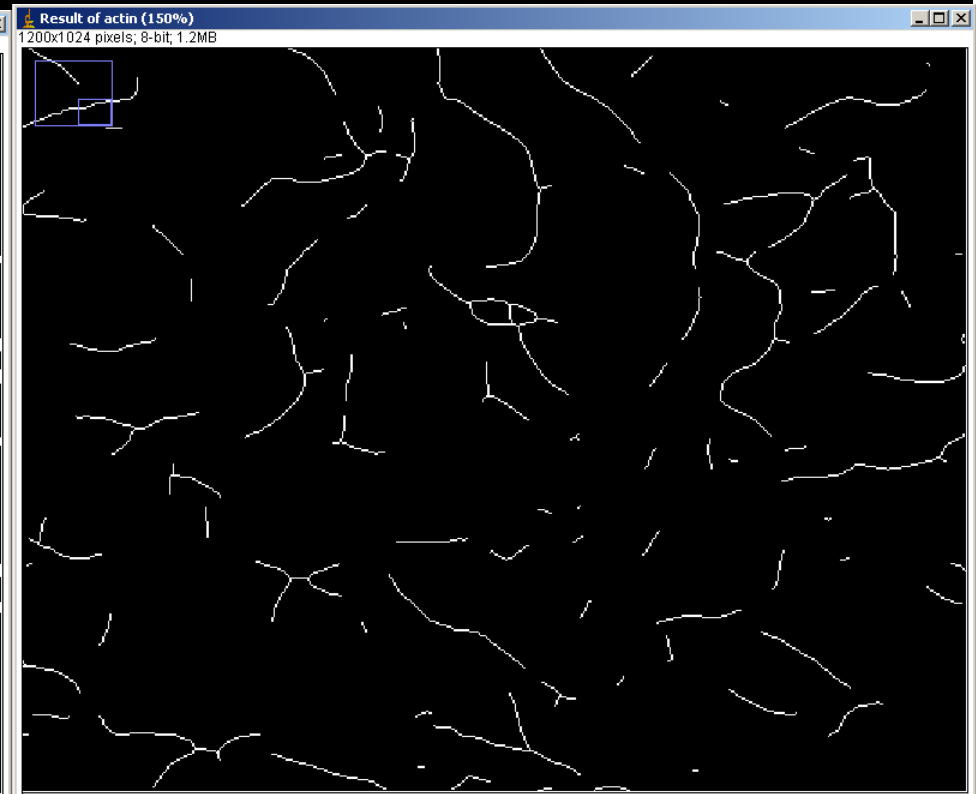
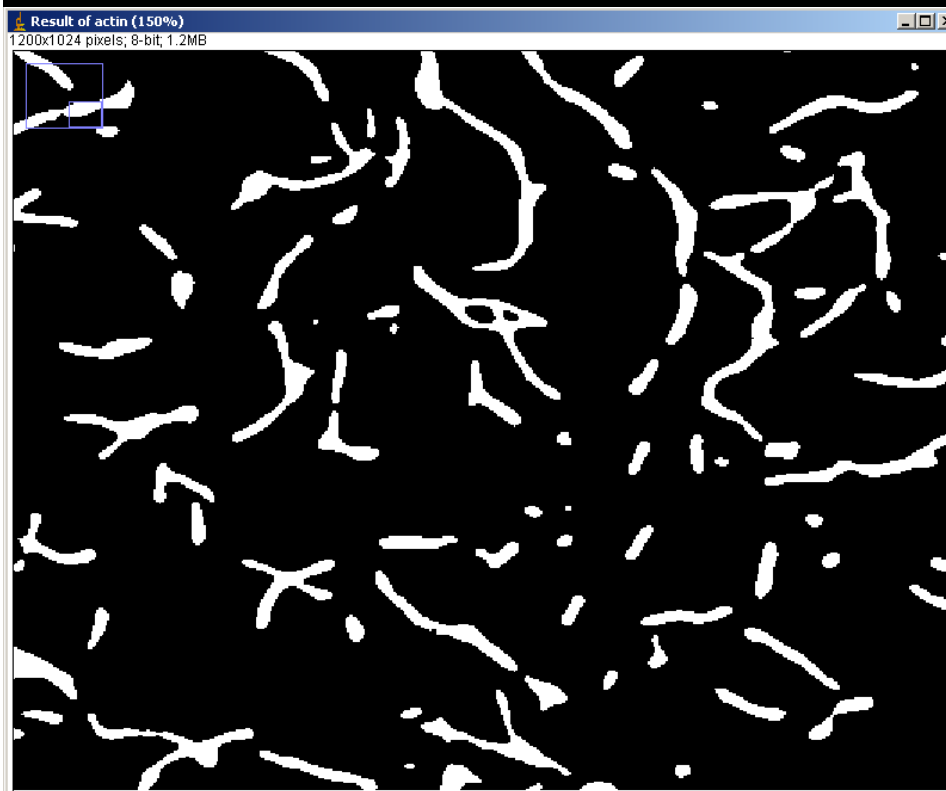
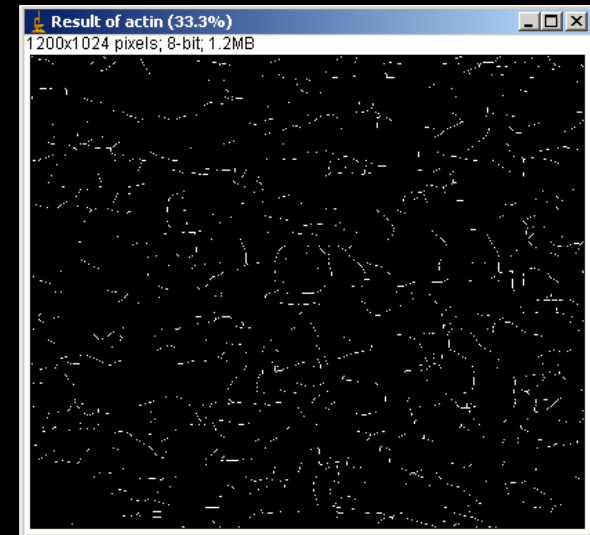
Gaussian Blur (2)
Remove noise



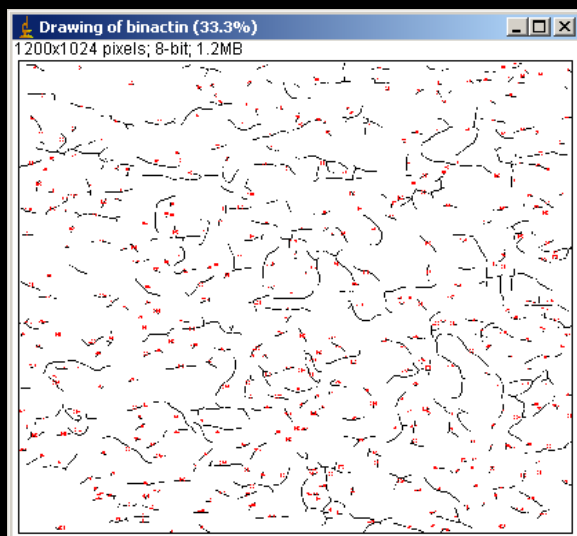


Binarized
(thresholding)

Skeltonize
(erosion)

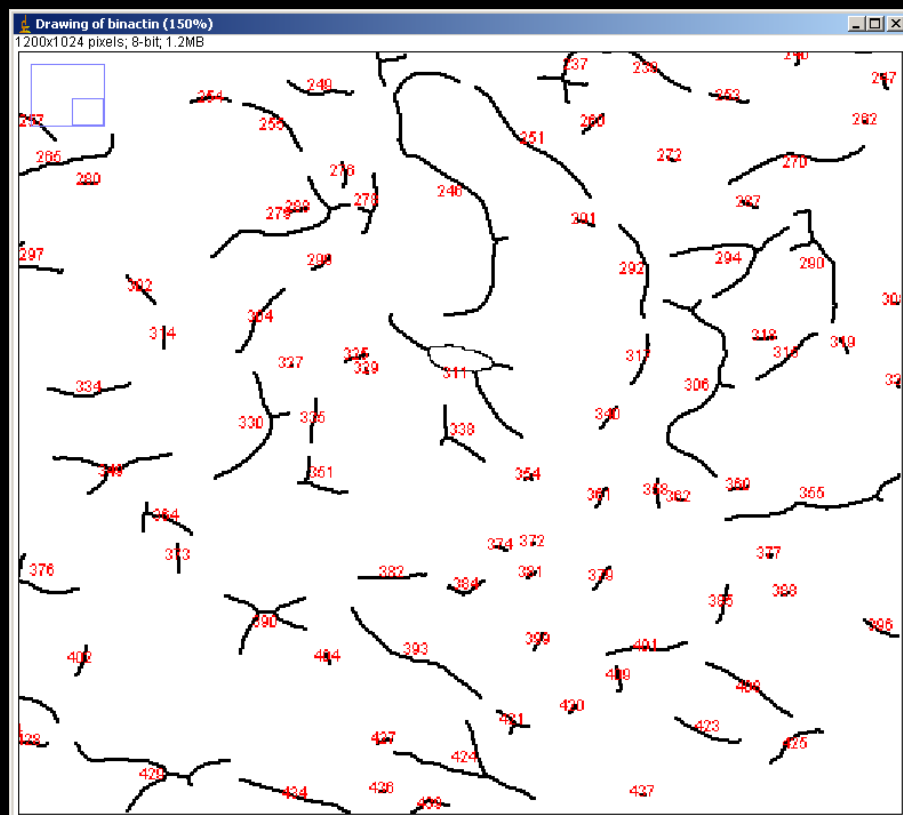


Threshold and Particle Count...



Summary				
File Edit Font				
Slice	Count	Total Area	Average Size	Area Fraction
binactin	440	17157.00	38.99	1.4

Results				
File Edit Font				
	Cut	Ctrl+X	IntDen	
408	Copy	Ctrl+C	9.48	12750
409	Clear		4.94	4080
410	Select All	Ctrl+A	3.86	6375
411			1.07	3570
412	Clear Results		2.50	2805
413	Summarize		6.50	1785
414	Distribution...		3.79	32385
415	Set Measurements...		1.50	6630
16	Duplicate...		2.75	2040
17			5.70	3825
18	17	255	503.74	966.50 4335
19	46	255	231.57	964.74 11730



Distribution

Parameter: Area

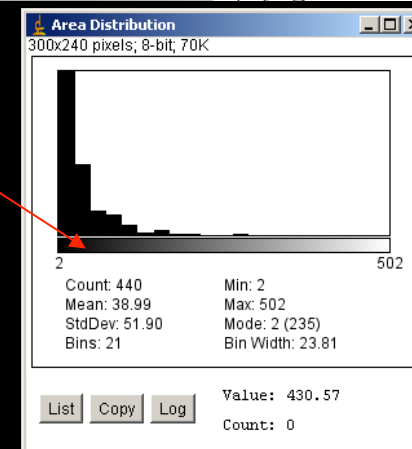
Data points: 440

☒ Automatic binning

or specify bins: 10

and range: 0-0

OK Cancel



Static Images: Morphometry

Length

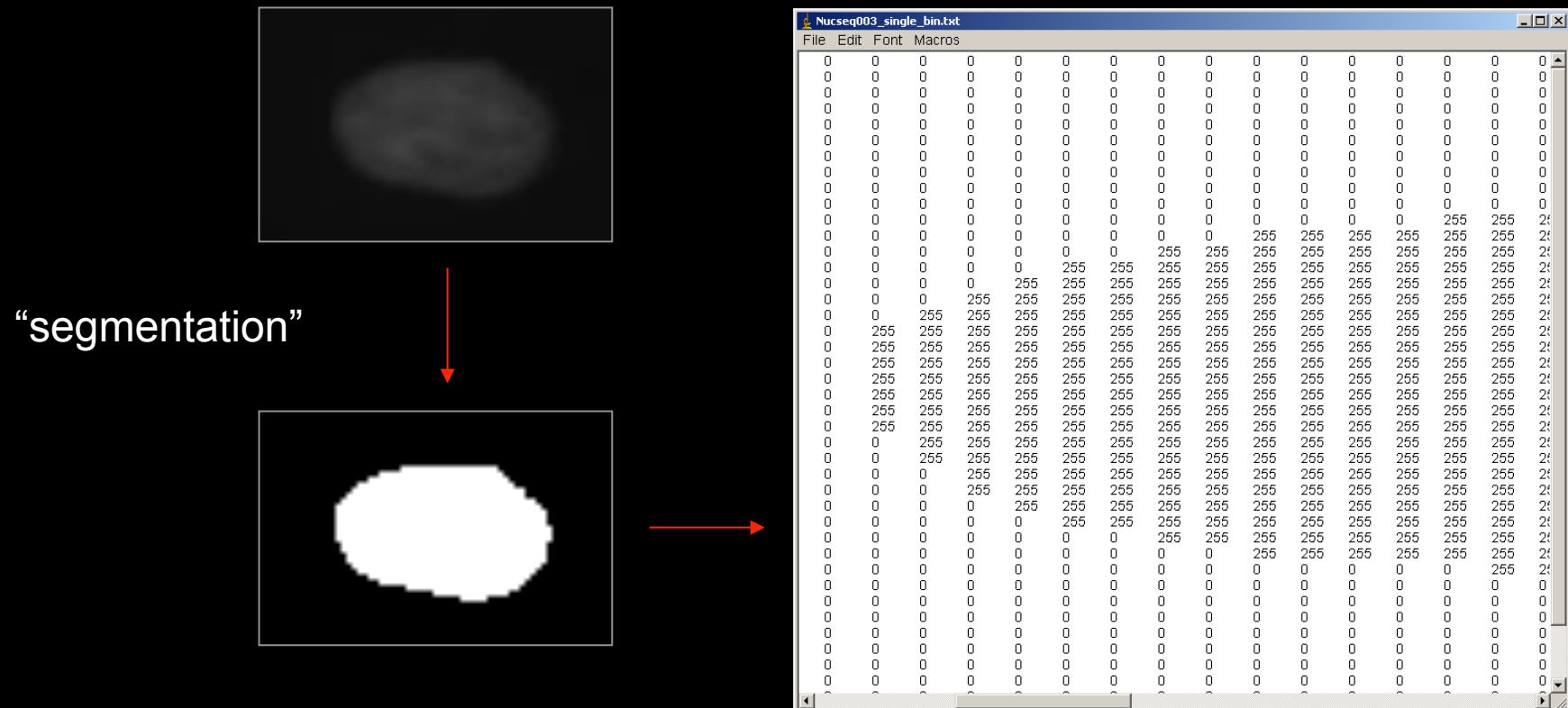
straight: (x_1, y_1) to (x_2, y_2)
curves

Area

Perimeter

Circularity

number of pixels with value 255
= Area



Static Images: Morphometry

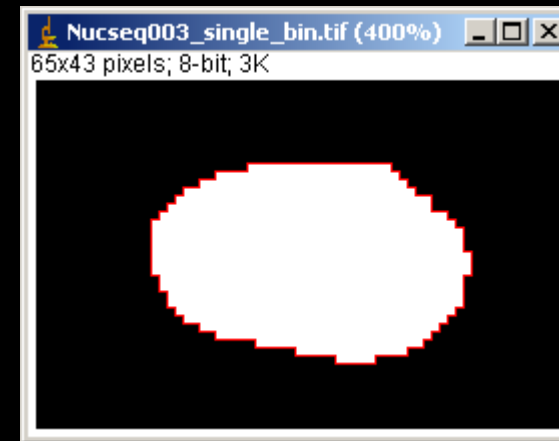
Length

straight: $(x1, y1)$ to $(x2, y2)$
curves

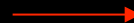
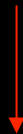
Area

Perimeter

Circularity



“segmentation”



Static Images: Morphometry

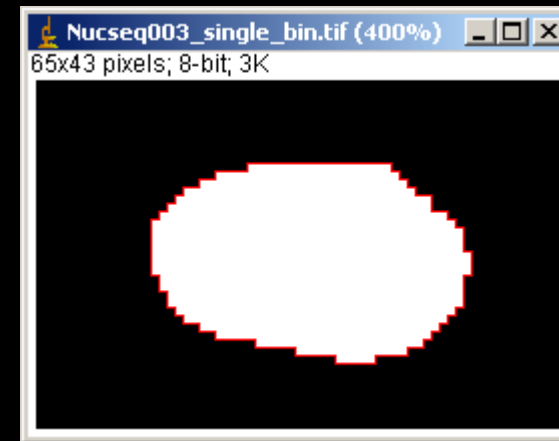
Length

straight: $(x1, y1)$ to $(x2, y2)$
curves

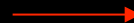
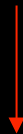
Area

Perimeter

Circularity



“segmentation”



Static Images: Morphometry

Length

straight: (x1, y1) to (x2, x2)

curves

Area

Perimeter

Circularity

$$\text{Circularity} = 4\pi(\text{area} / \text{perimeter}^2)$$

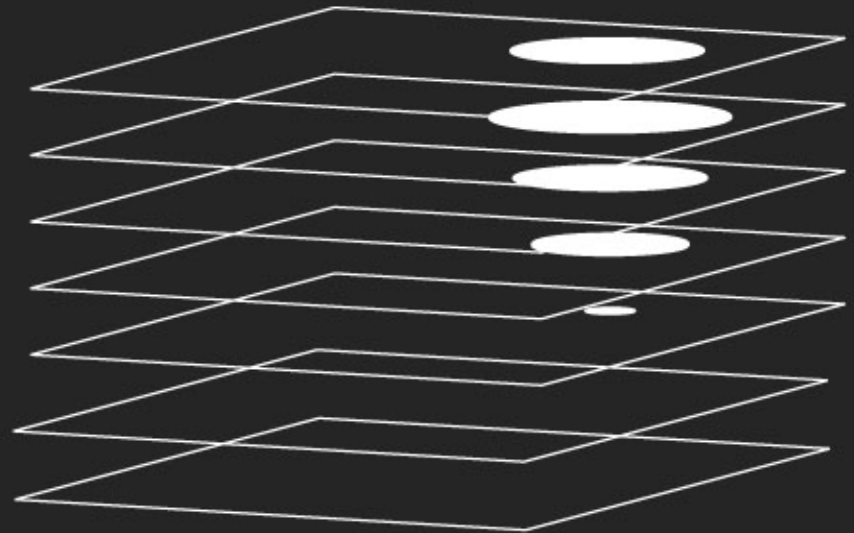
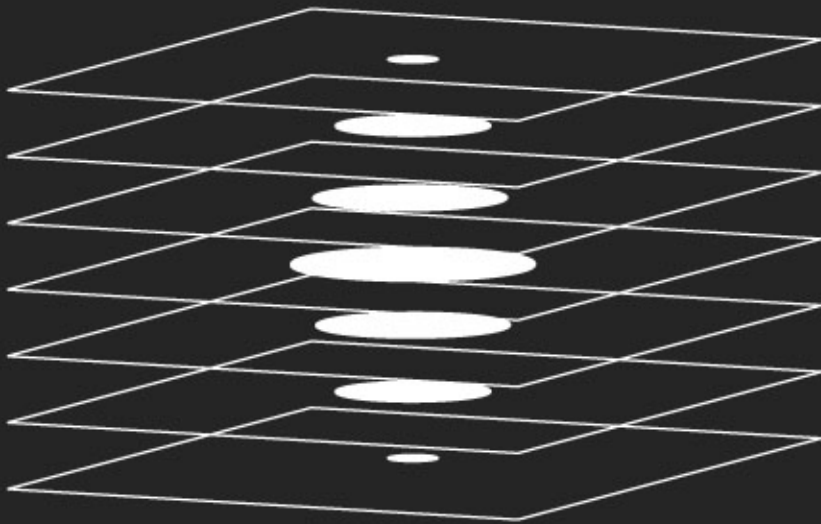


Circularity 0.906

0.843

0.348

Static Images: Morphometry of three dimensional objects



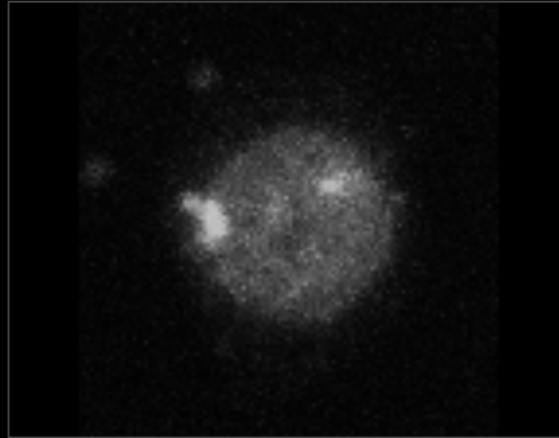
3D data sets (x-y-z): stack of optical sections

Static Images: Morphometry of three dimensional objects

XY stack



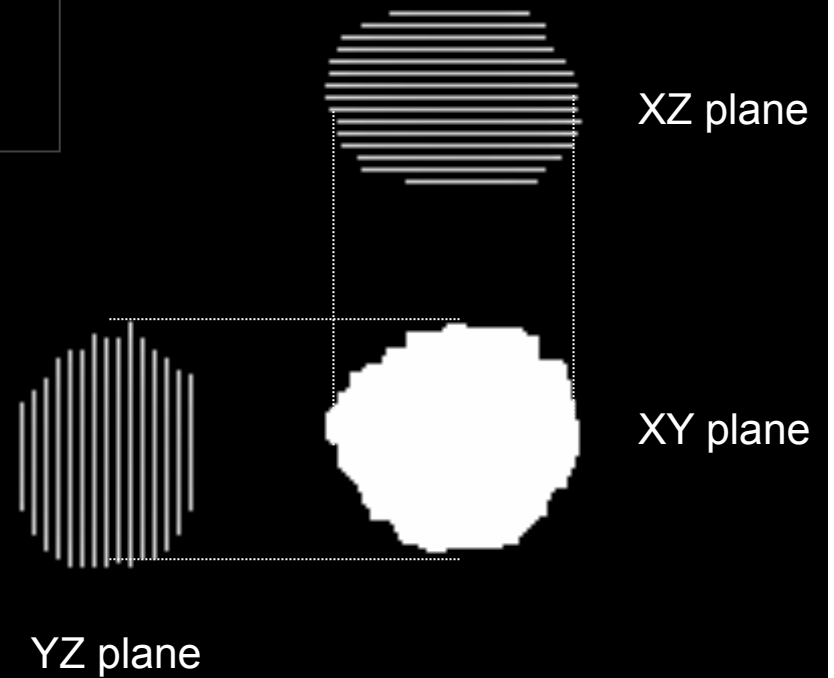
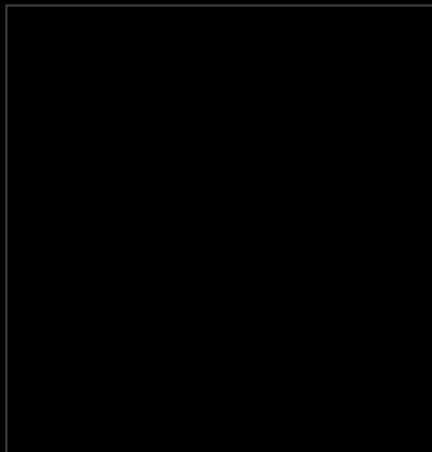
3D projection



Volume =

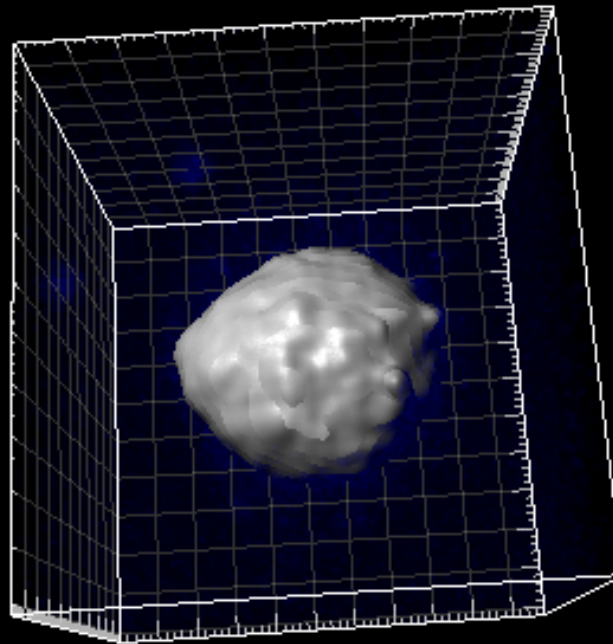
$\text{Sigma}(\text{Area}(\text{slice}) * \text{thickness})$

Segmented Image



Static Images: Morphometry of three dimensional objects

Software:
Imaris (Bitplane)



Visualization of
3D object

50 μm

Static Images: Distribution

Static images

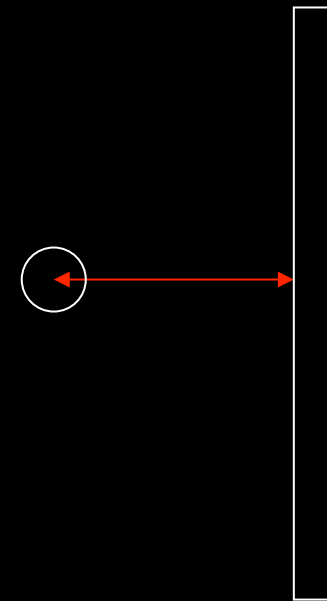
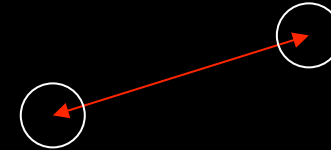
Intensity Measurement
Morphometry
Distribution
Colocalization

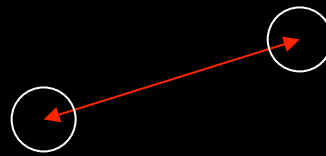
Image sequences

Intensity dynamics
Shape dynamics
Position dynamics :
 kymograph
 tracking
Distribution dynamics
Optical Flow Estimation

Distance Between Points

Distance from Specific Structure

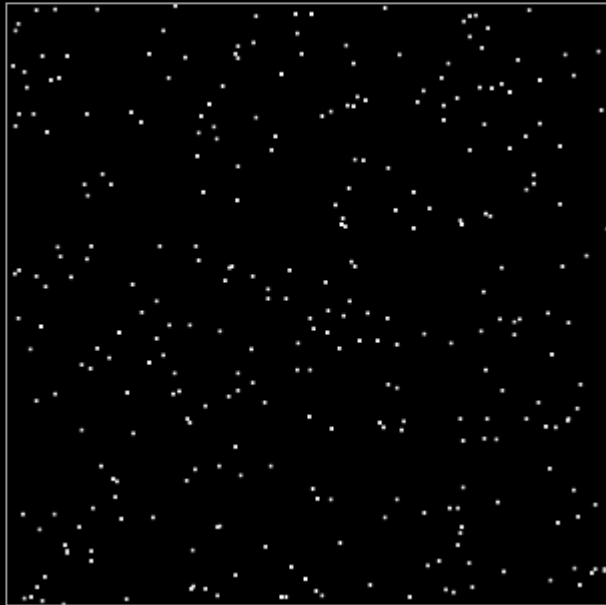




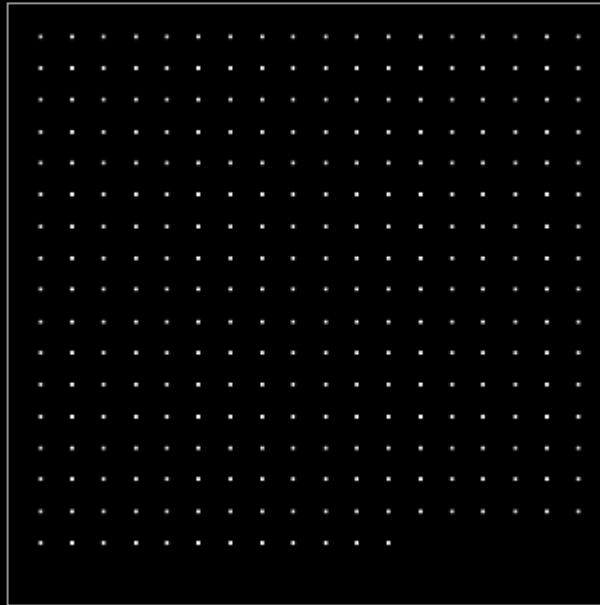
Distance Between Points

Static Images: Distance Between Points

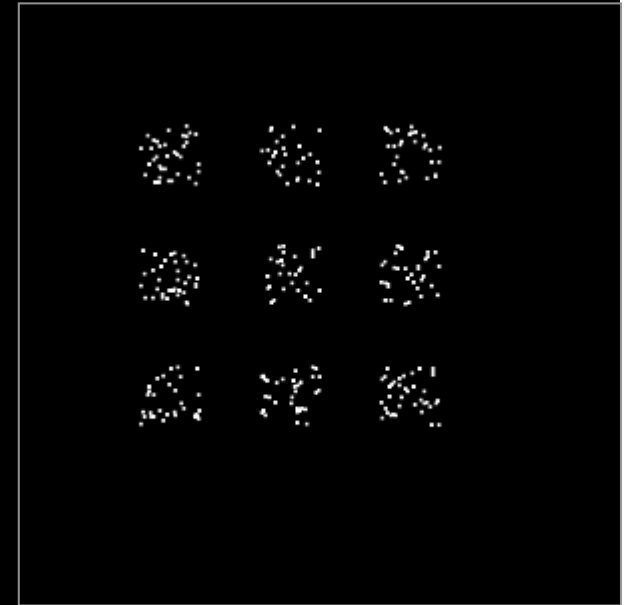
Quantification of Particle Distribution



random

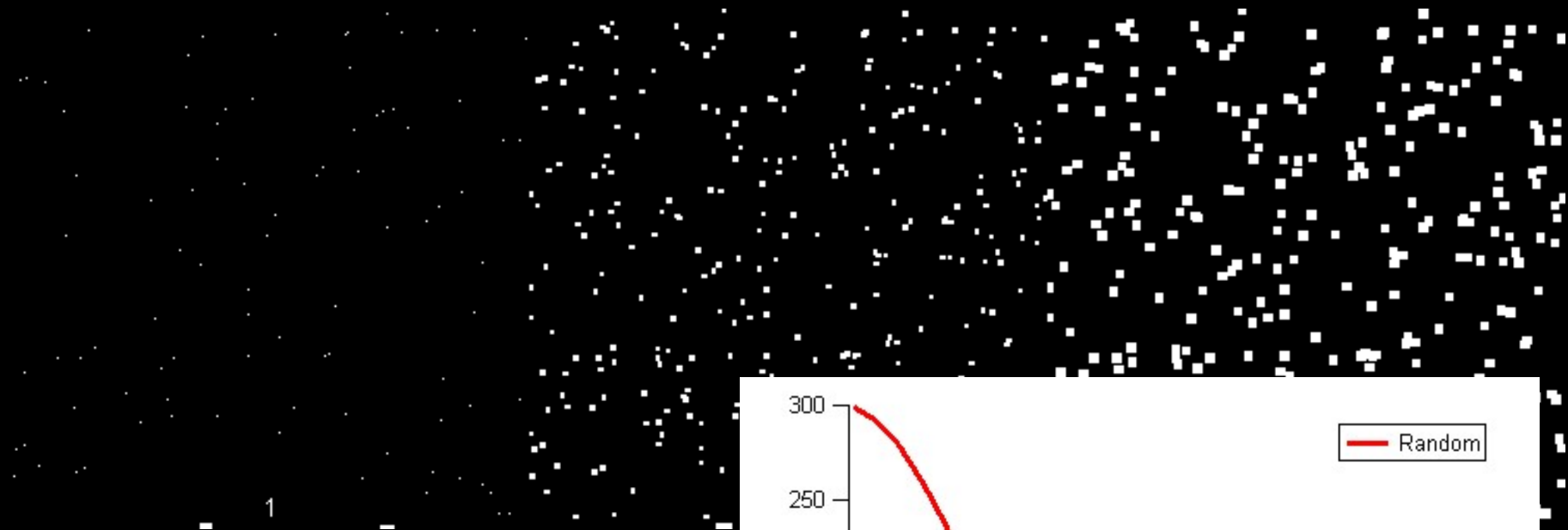


grid

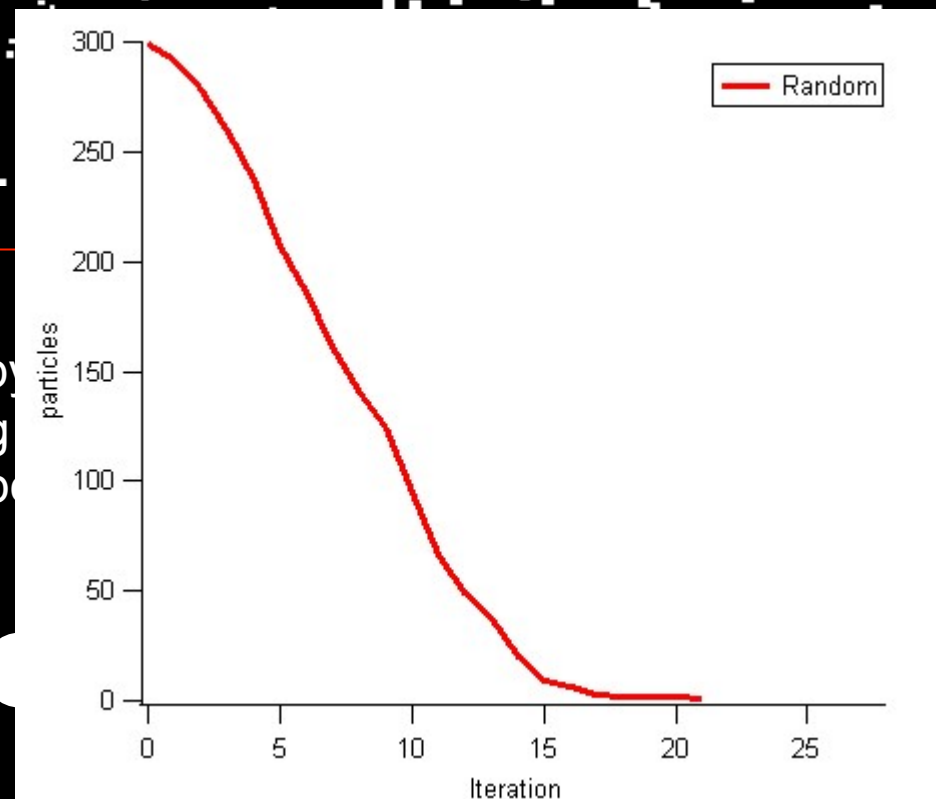


clusters

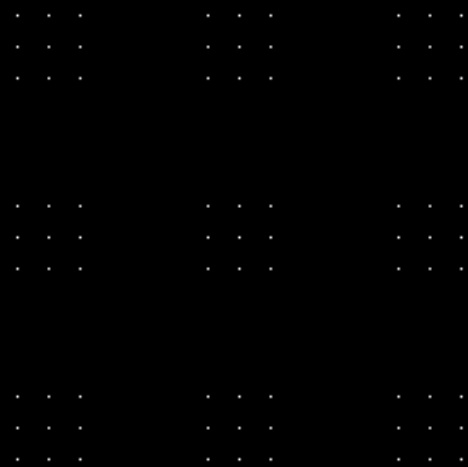
Static Images: Distance Between Points



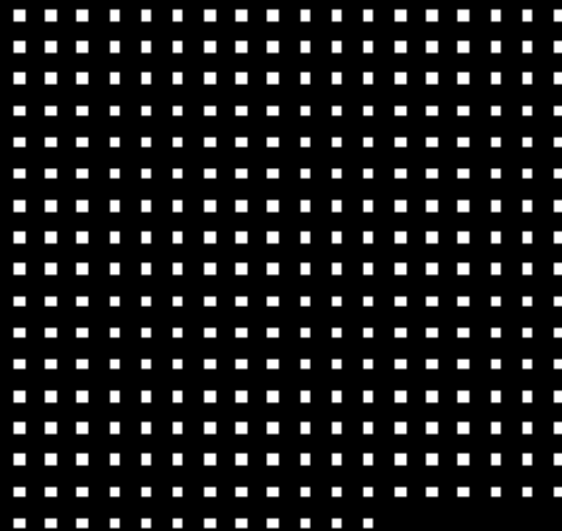
Dilating signals step-by-step
→ increasing
→ less number



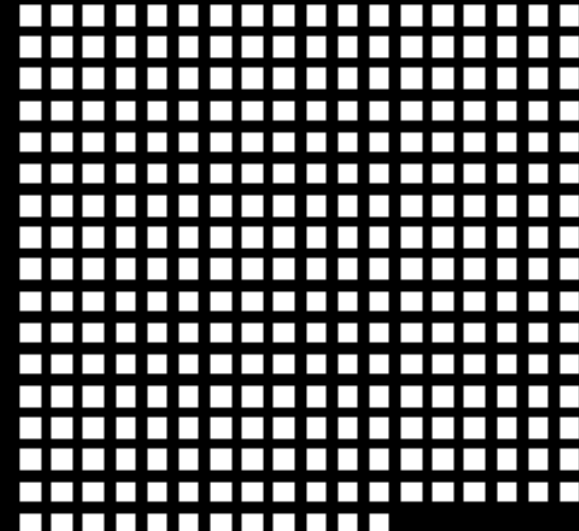
Static Images: Distance Between Points



1



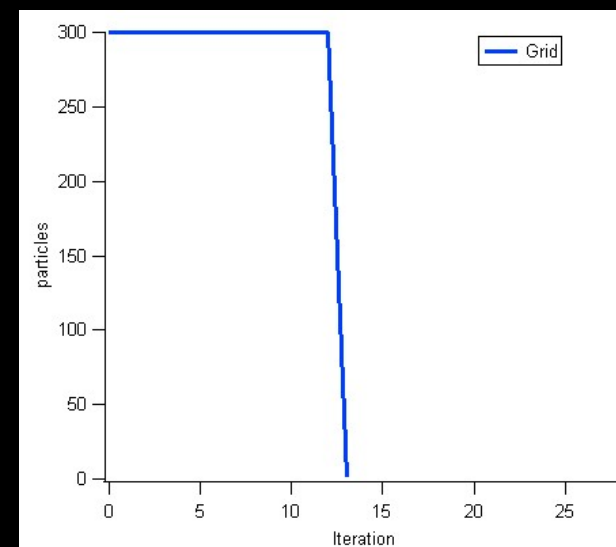
5



9

Dilating the signal

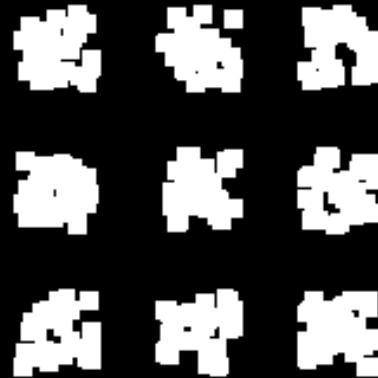
- increasing connections between particle
- less number of “particles”



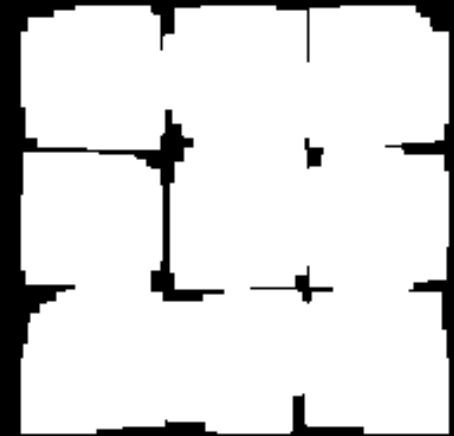
Static Images: Distance Between Points



Many



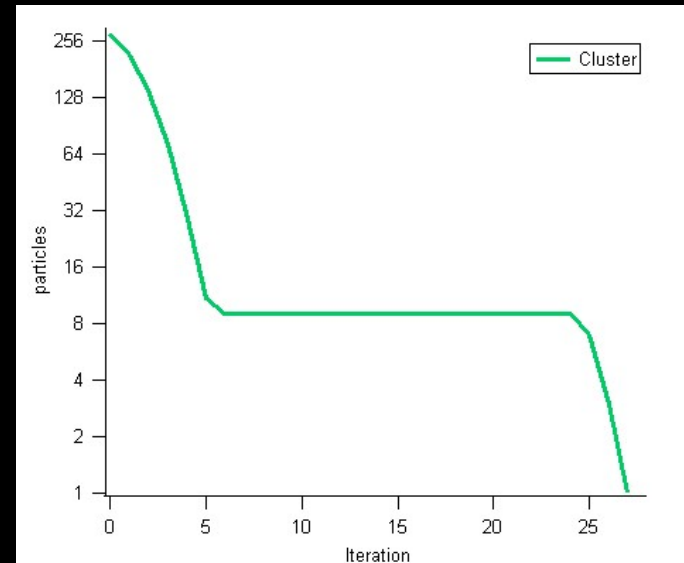
9 clusters



Single Cluster

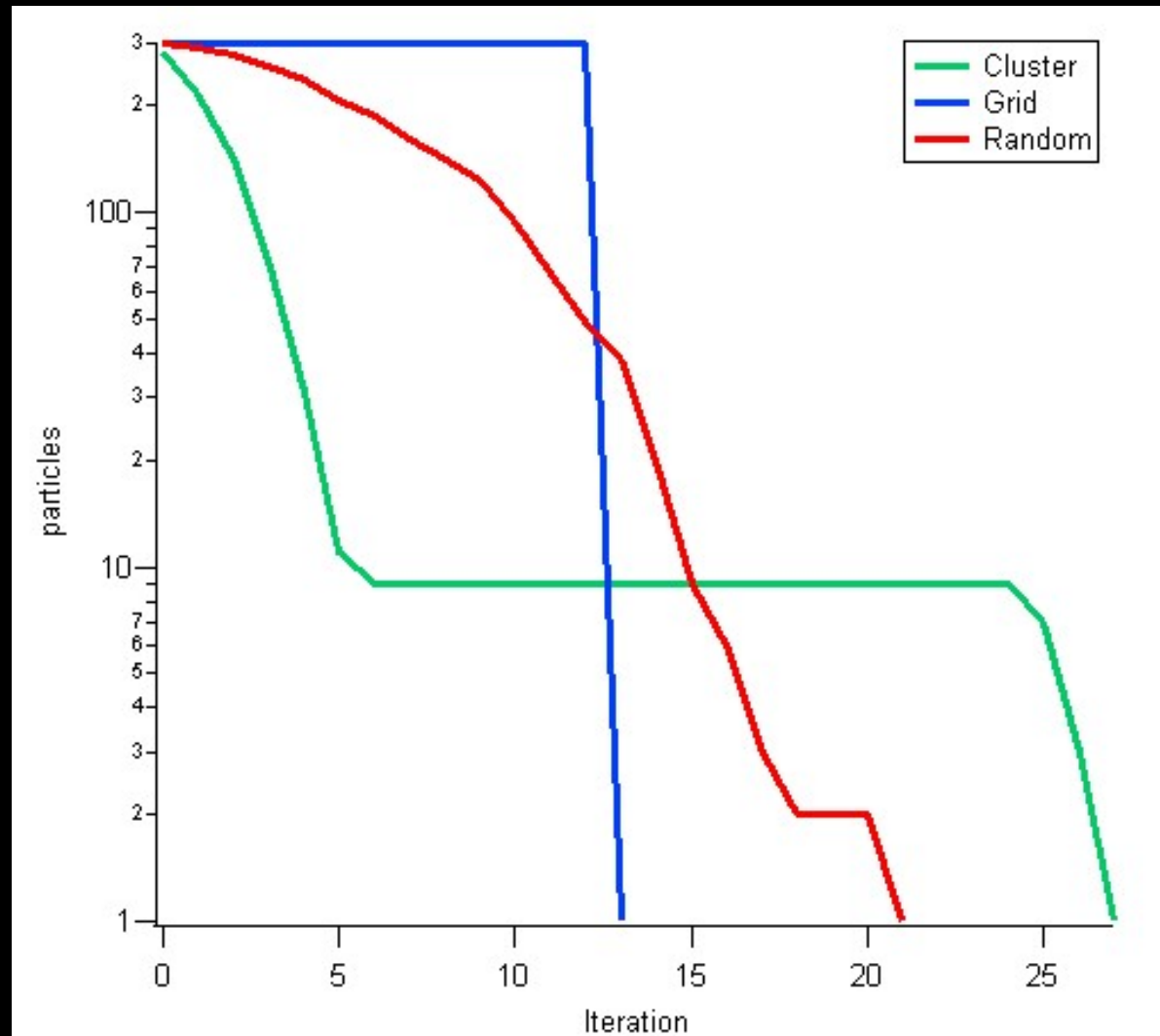
Dilating the signal

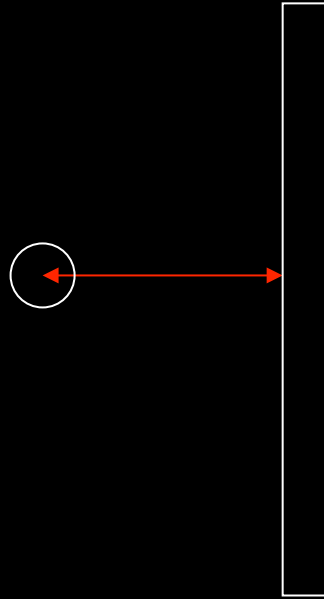
- increasing connections between particle
- less number of “particles”



Static Images: Distance Between Points

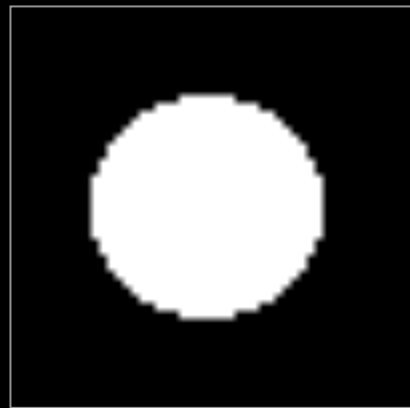
Indicator of particle distribution → Connected Region Statistics



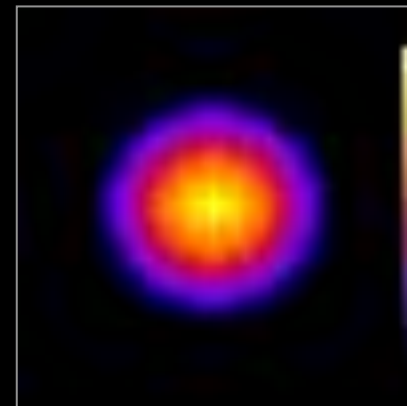
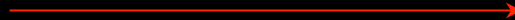


Distance from Specific Structure

2. Measurement of Distance



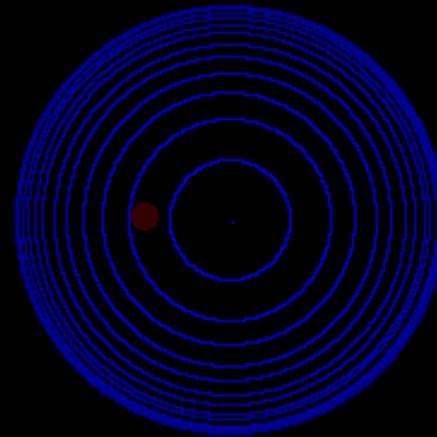
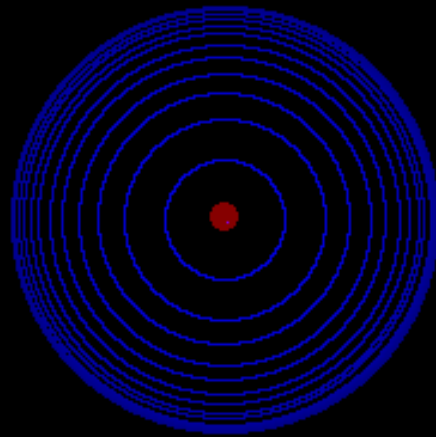
Distance Map (2D)



[illegible]

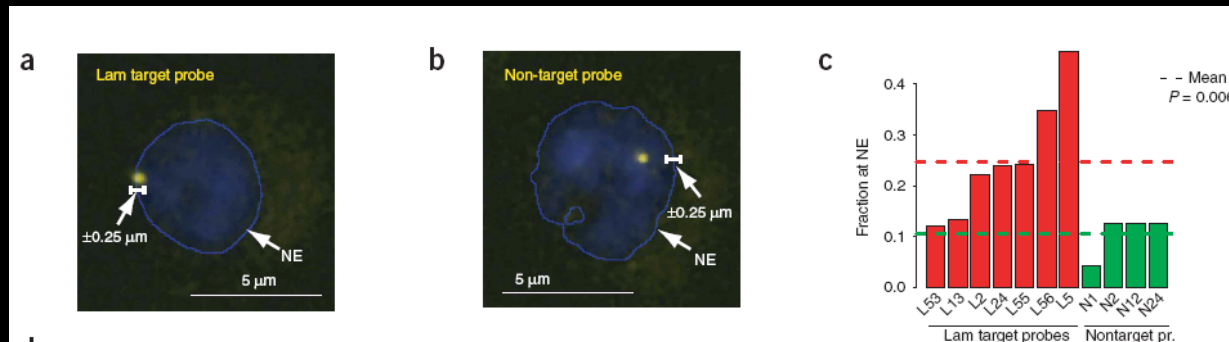
[illegible]

Static Images: Distance from Specific Structure in 3D

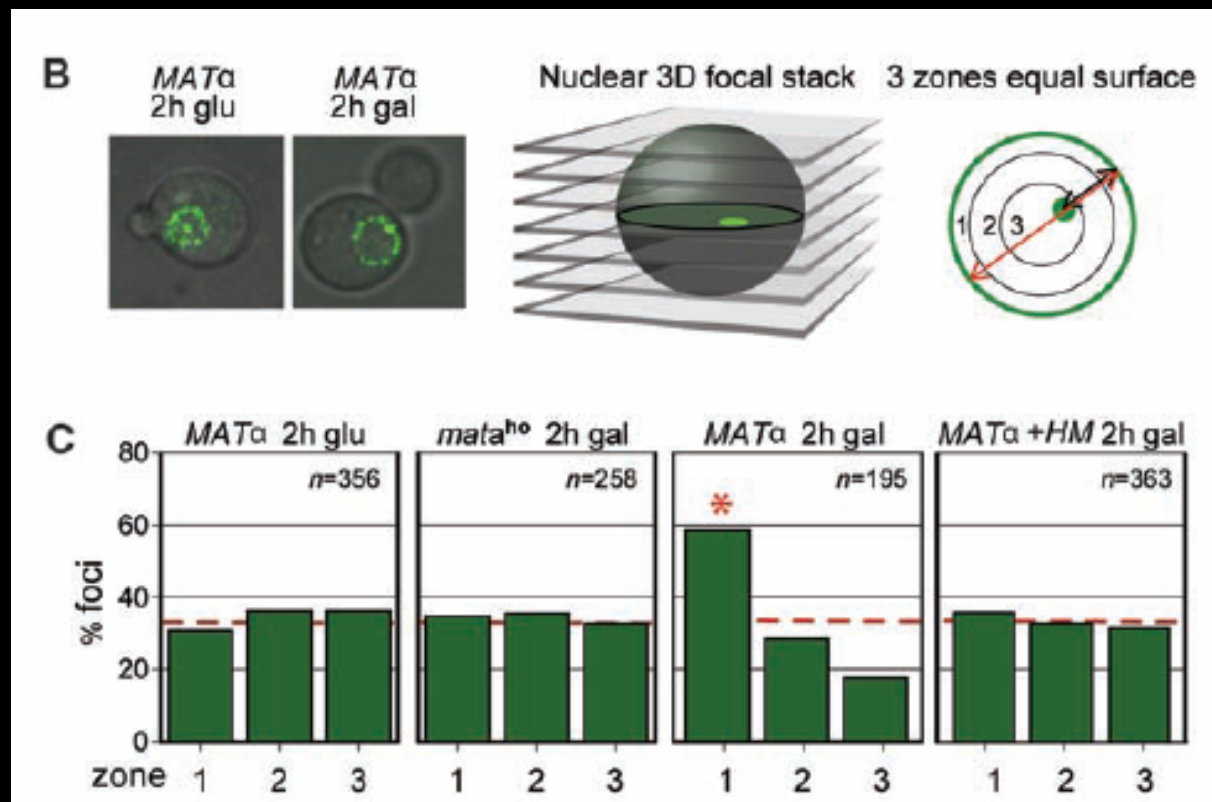


3D Localization of Chromosome signal: Conventional Method

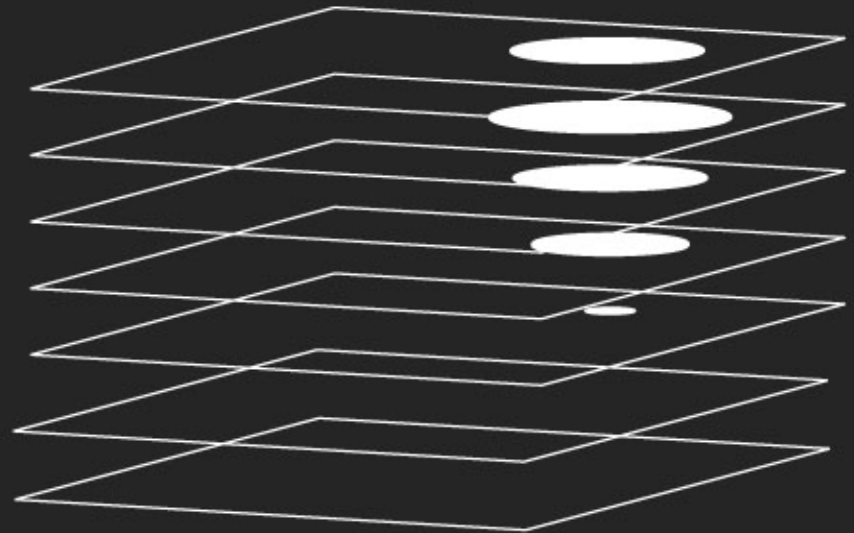
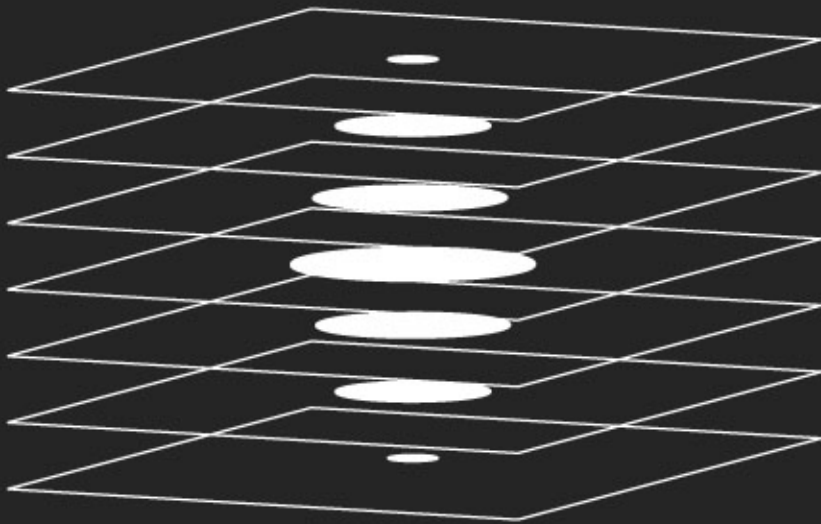
Pickersgill et al. (2006)
Nature Genetics 38 p1005



Nagai et al. (2008)
Science 332 p597



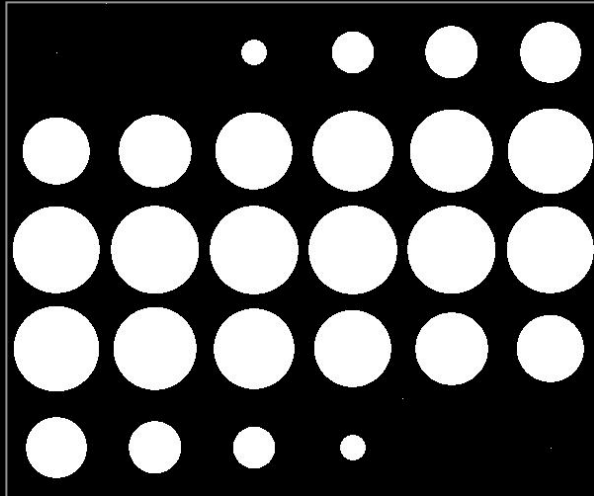
Static Images: Three dimensional data



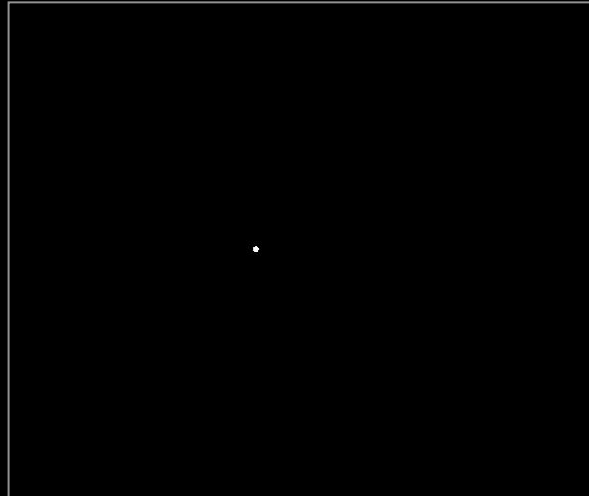
3D data sets (x-y-z)

Static Images: Distance from Specific Structure

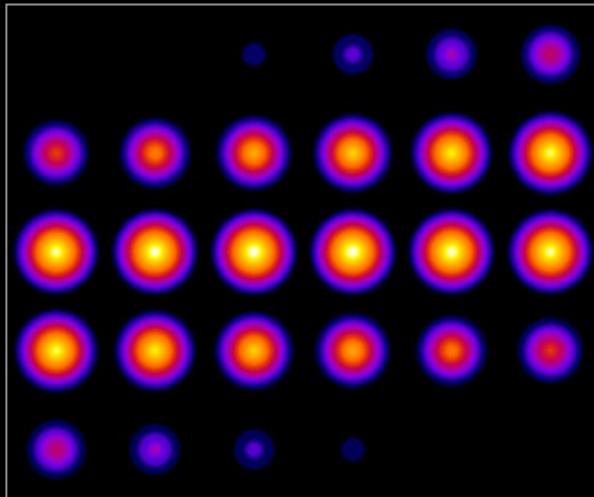
Nucleus



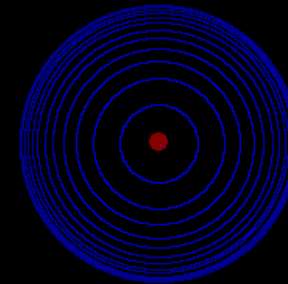
chromosome



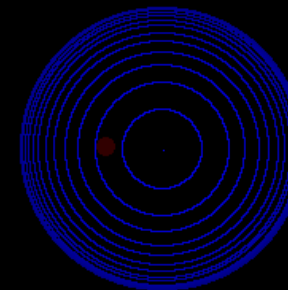
Nucleus 3D distance map



Center

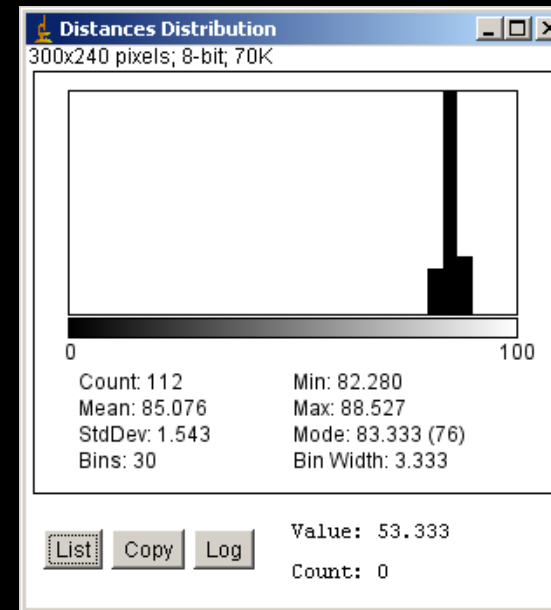
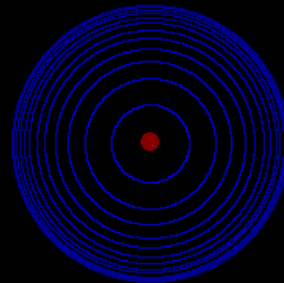


Periphery

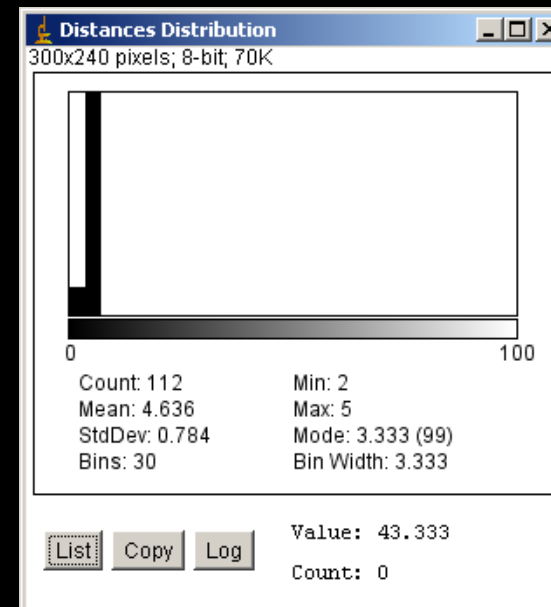
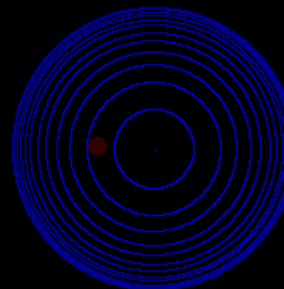


Static Images: Distance from Specific Structure

Center



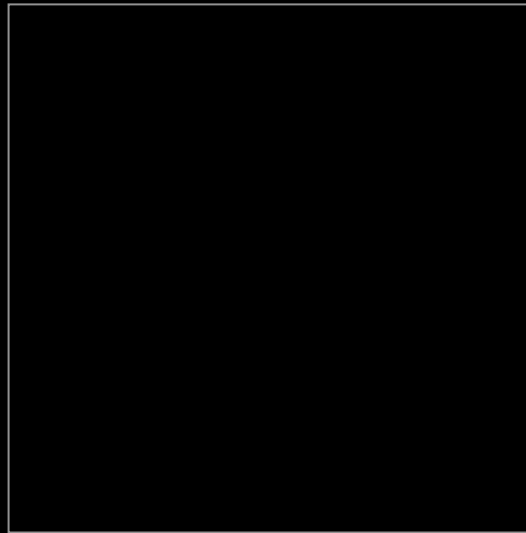
Periphery



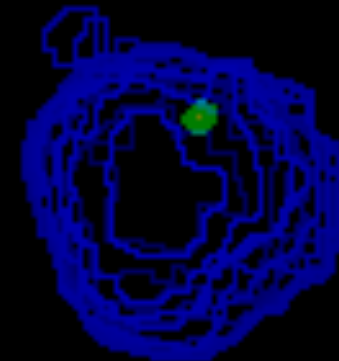
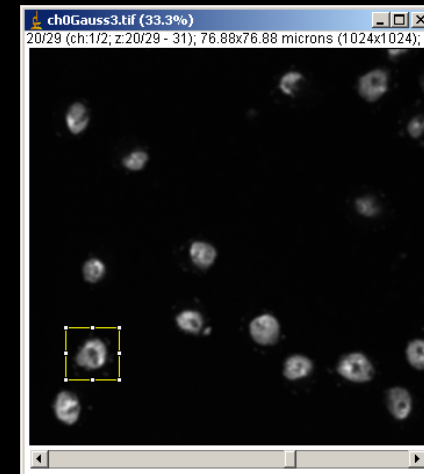
1. Segmentation of DAPI and FISH



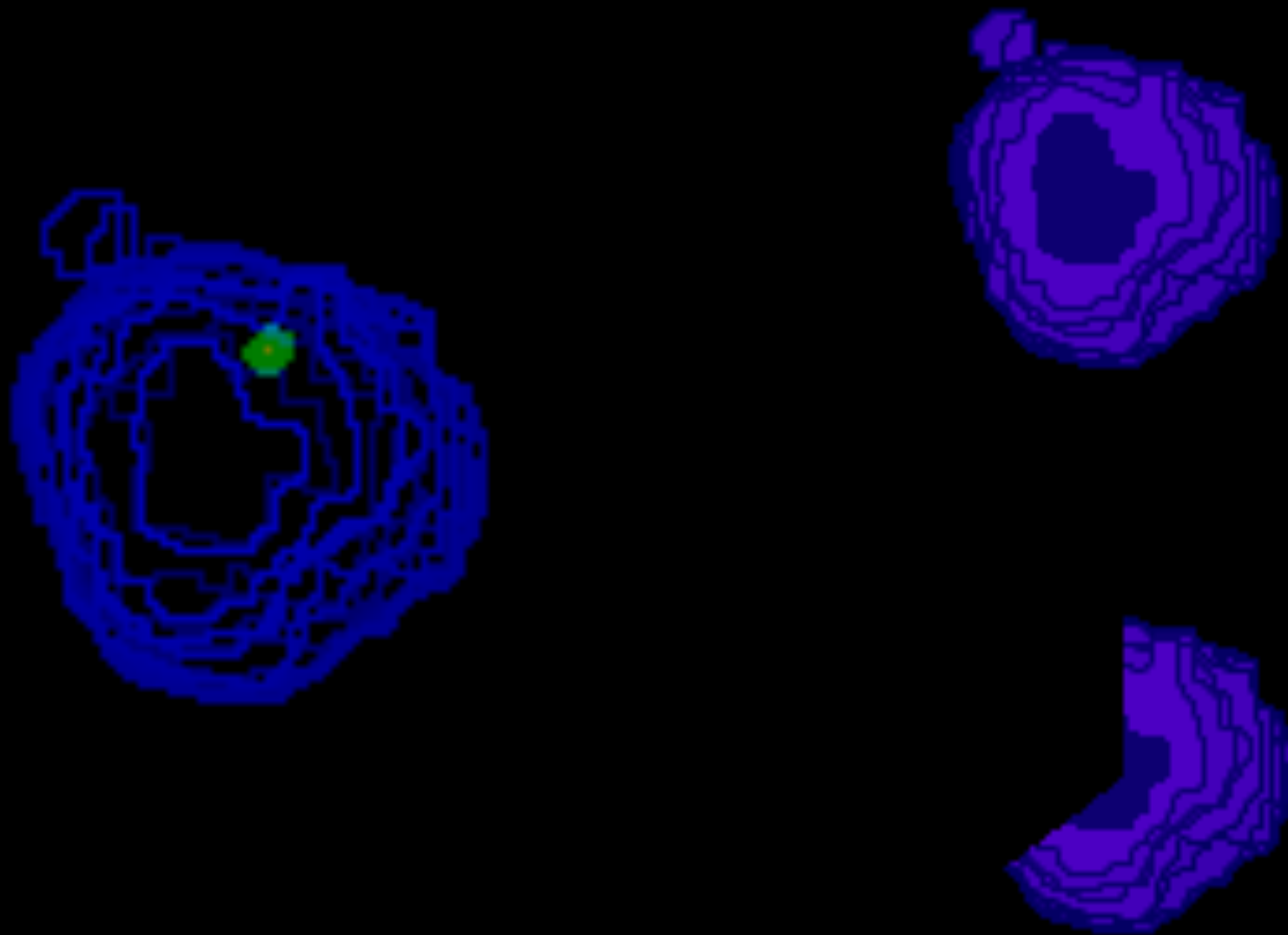
DAPI image



FISH image



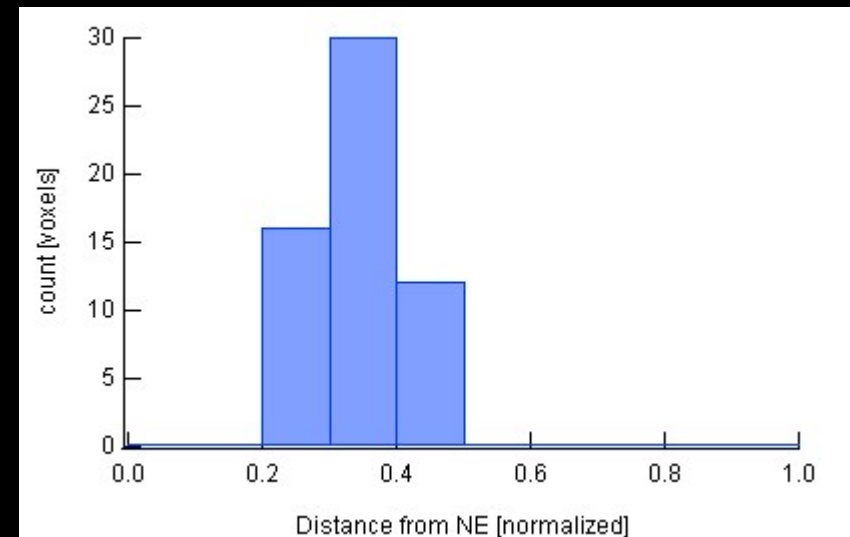
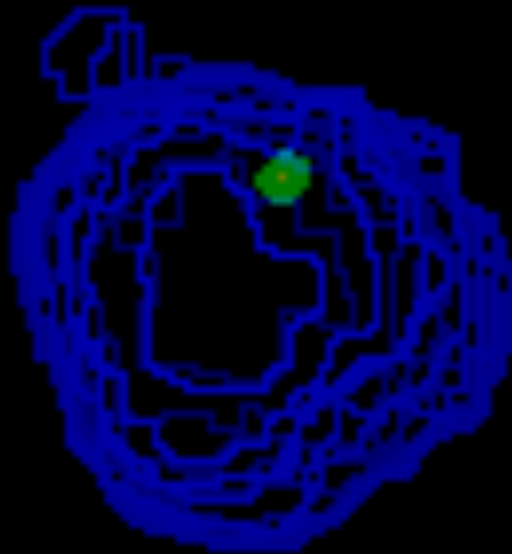
2. Measurement of Distance: Distance Map 3D



2. Measurement of Distance: Distance Map 3D

§

-> Leaves only FISH voxels with distances



1. Vaquerizas et al. Nuclear pore proteins nup153 and megator define transcriptionally active regions in the Drosophila genome. PLoS genetics. 2010;6(2):e1000846

Static Images: Colocalization

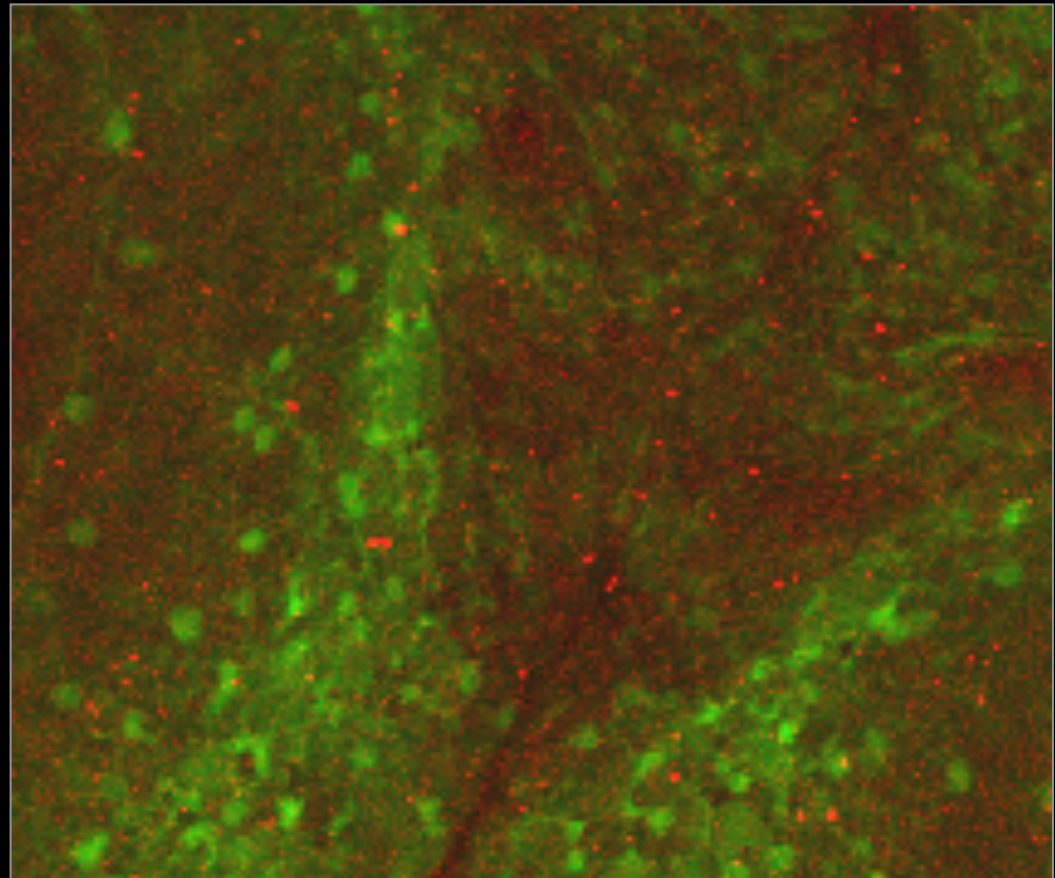
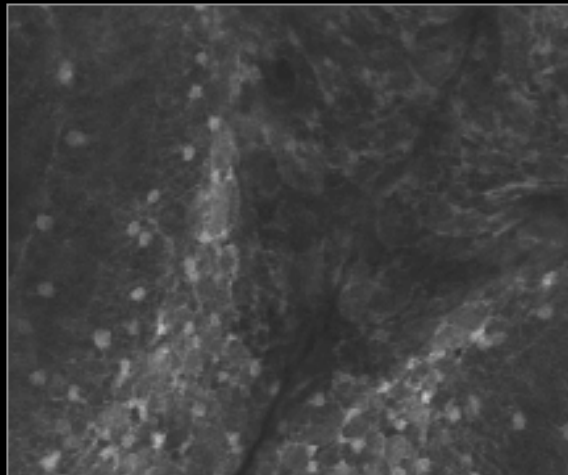
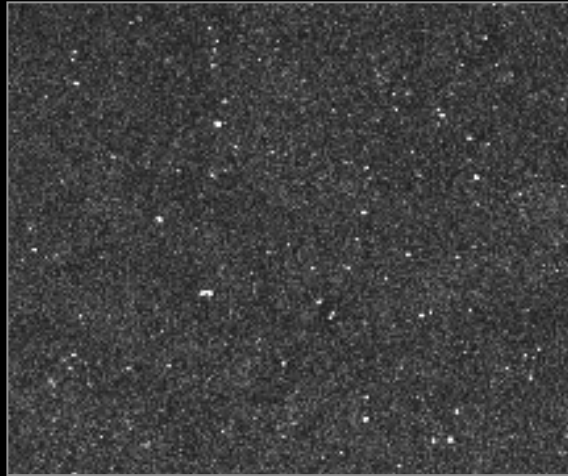
Static images

Intensity Measurement
Morphometry
Distribution
Colocalization

Qualitative
Quantitative

Image sequences

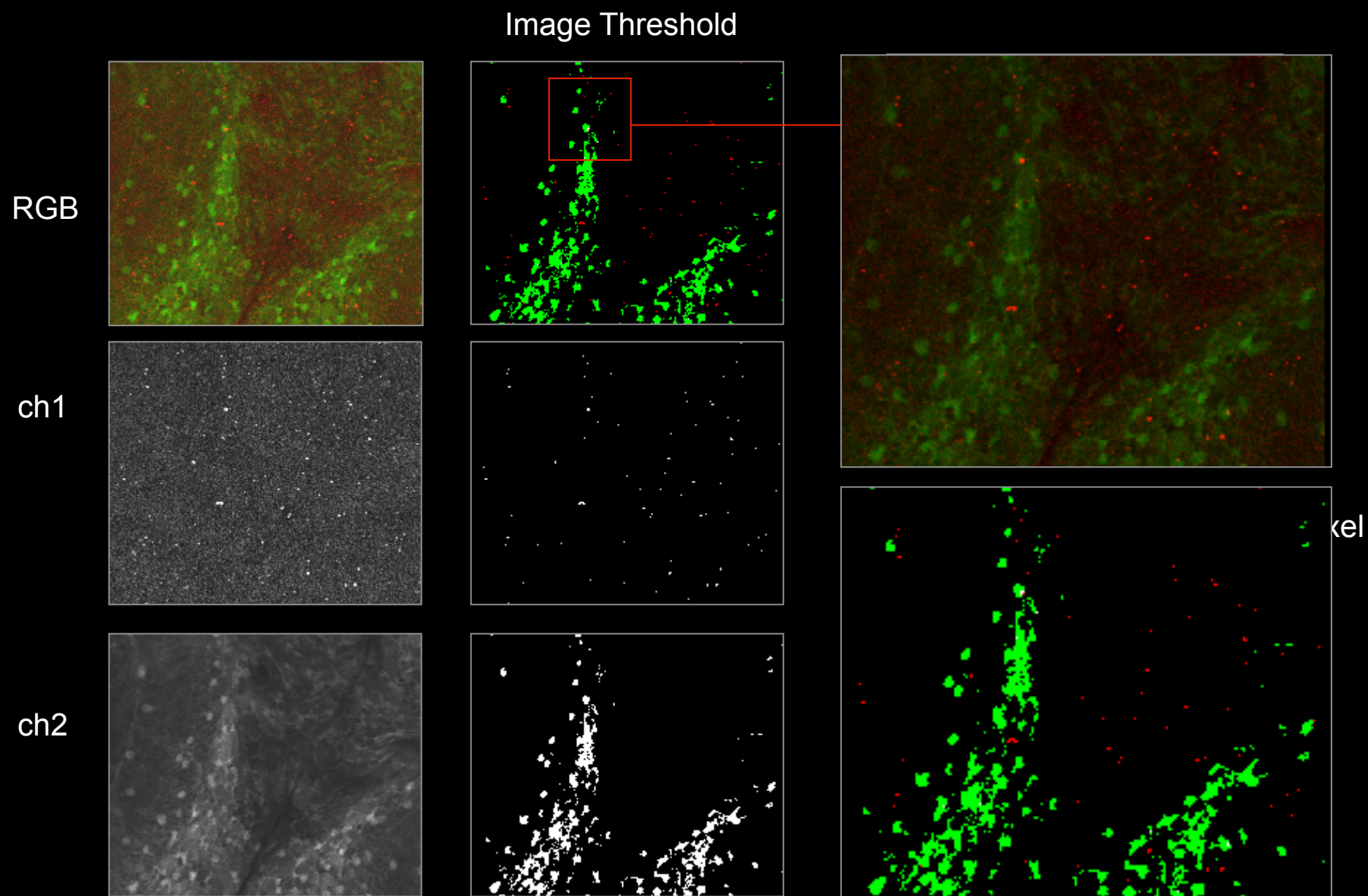
Intensity dynamics
Shape dynamics
Position dynamics :
 kymograph
 tracking
Distribution dynamics
Optical Flow Estimation



“see, they are colocalized!”

Static Images: Co-localization

Qualitative, a bit more quantitative



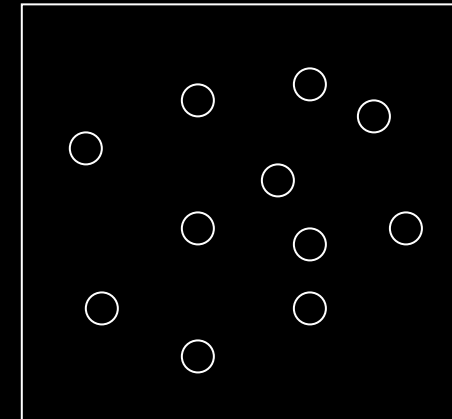
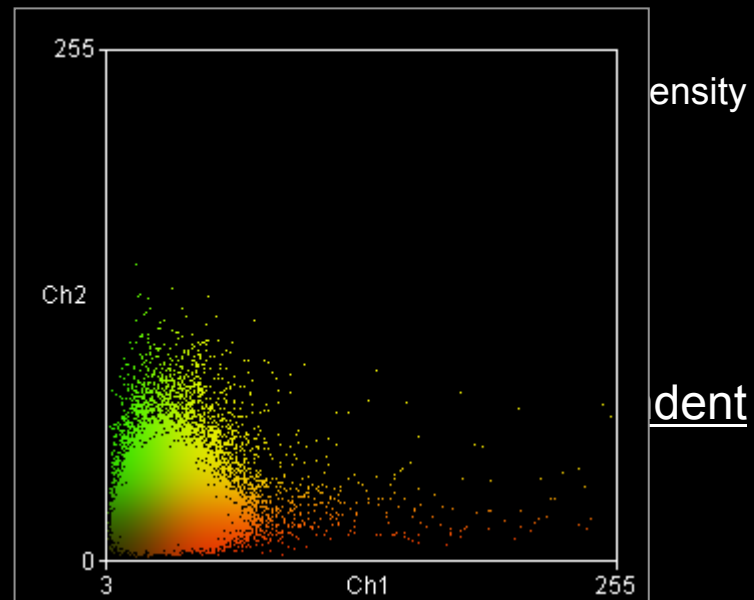
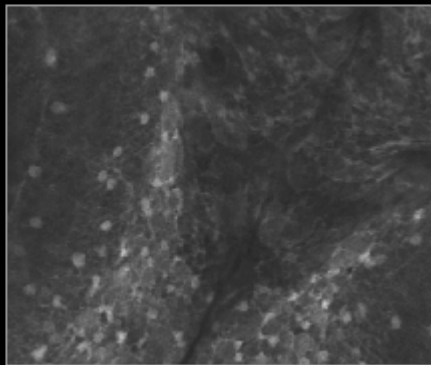
Static Images: Co-localization

Quantitative

ch1



ch2

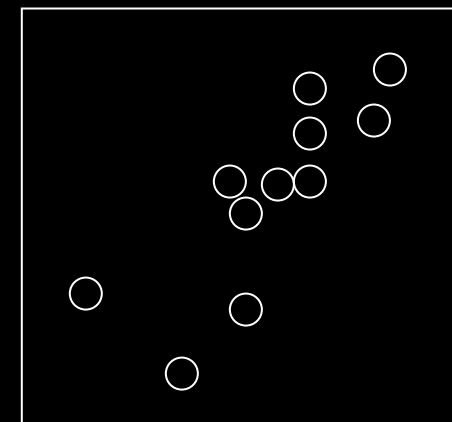


Ch1 intensity

Ch2 intensity

- Plot a point at (ch1 intensity, ch2 intensity)
- Color code the number of pixels for each position.

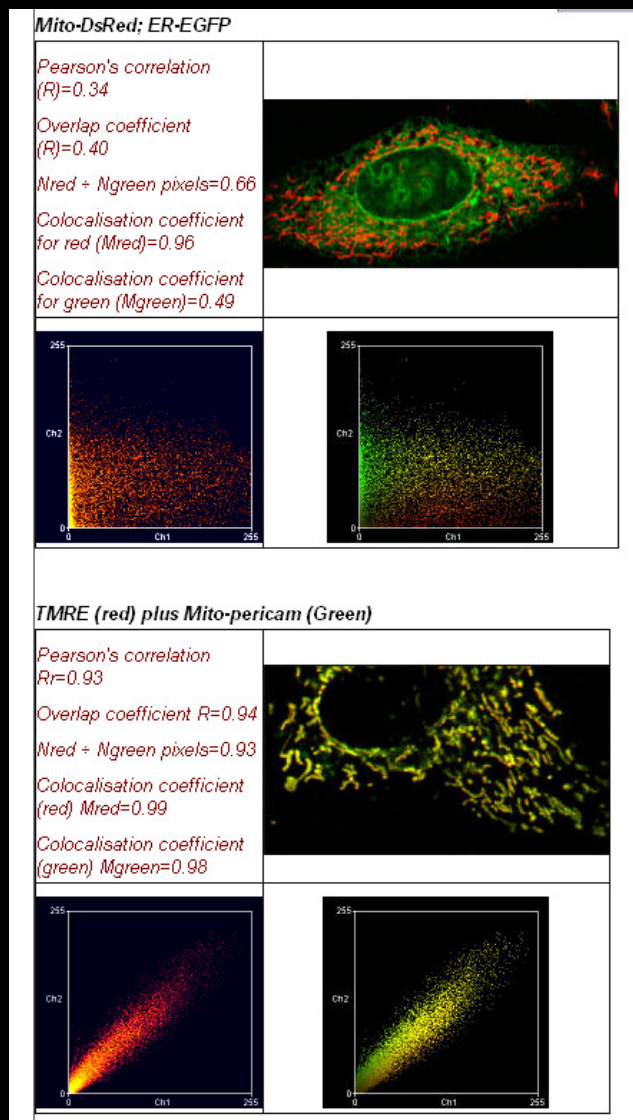
colocalized



Ch1 intensity

Static Images: Co-localization

Quantitative



Ch2 intensity

independent

Ch1 intensity

Ch2 intensity

colocalized

Ch1 intensity

Statistics:

van Steensels test
(J.Cell Sci. 109:787)

Costes test
(Costes et al. 2004. Biophys. J 86:3993)

Fay's test
(Fay et al. 1997 Exp. Cell Res. 231: 27)

Manders' coefficients test
(Manders et al. 1993. J. Microscopy. 169:375)

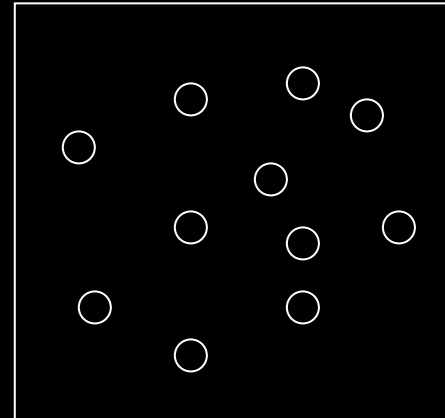
Review

Bolte, S. & Cordelieres, F.P. (2006) J. Microsc. 224, 213.
A guided tour into subcellular colocalization analysis in
light microscopy.

Ch2 intensity

independent

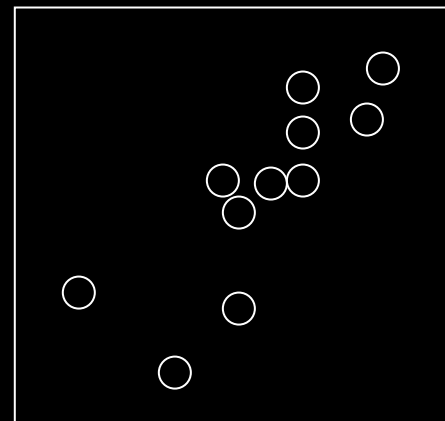
Ch1 intensity



Ch2 intensity

colocalized

Ch1 intensity



Dynamics: Image Series

Static images

- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

Image sequences

- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
 - optical flow
- Distribution dynamics

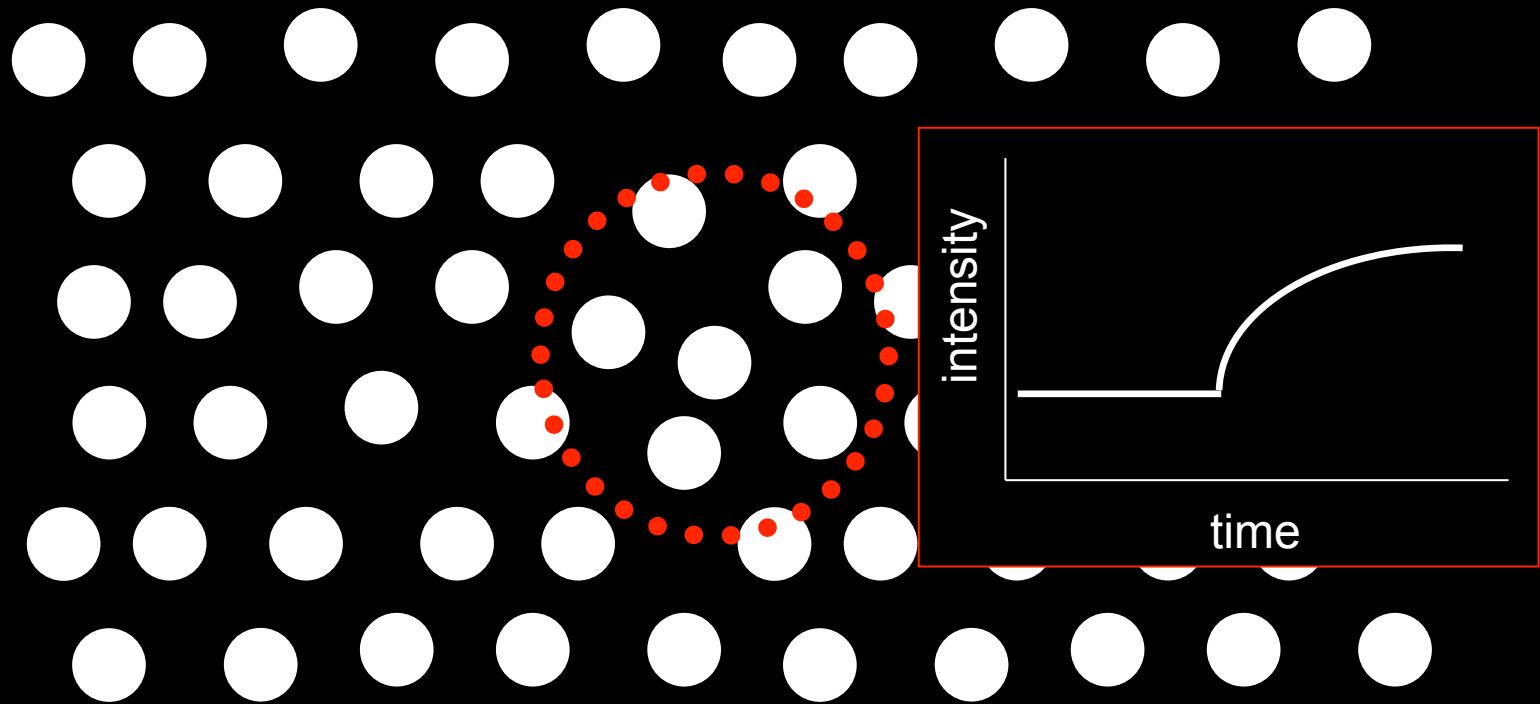
Static images

- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

Image sequences

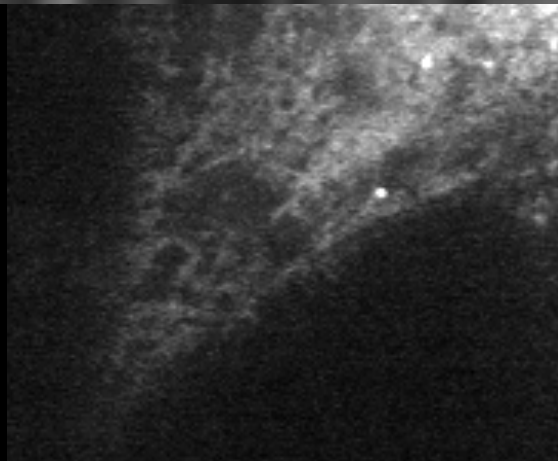
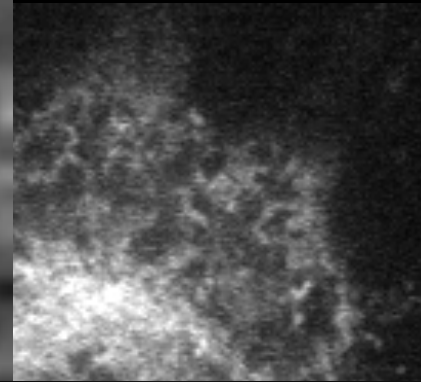
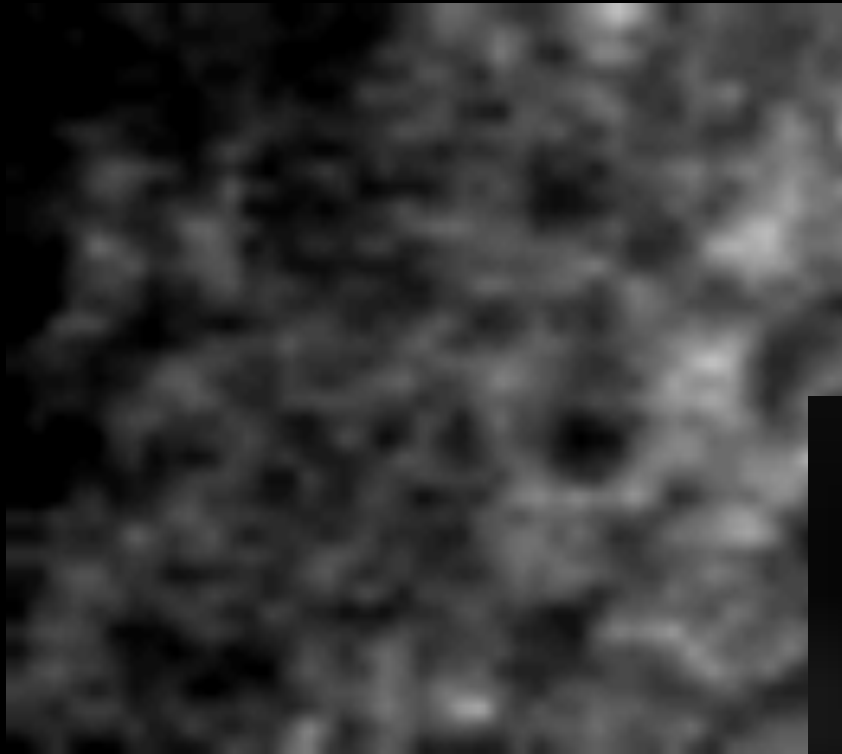
- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation

Kinetics, FRAP, FLIP ...



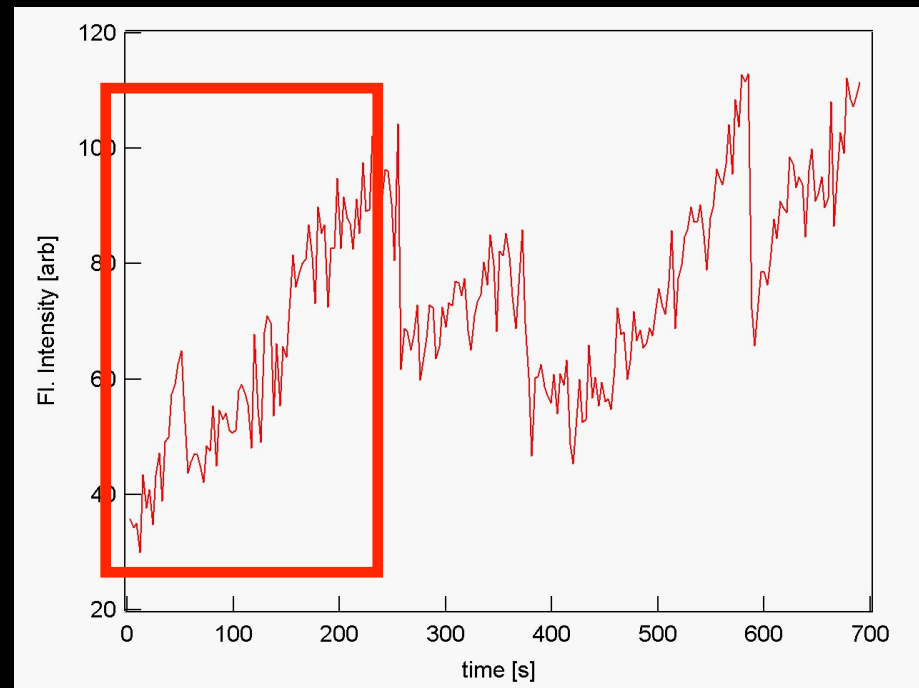
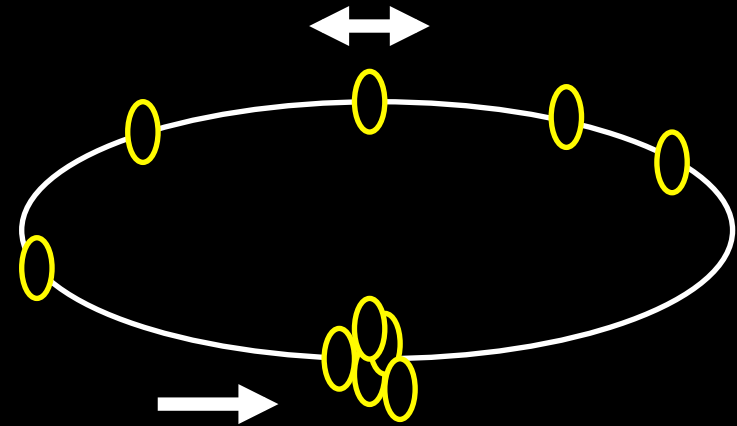
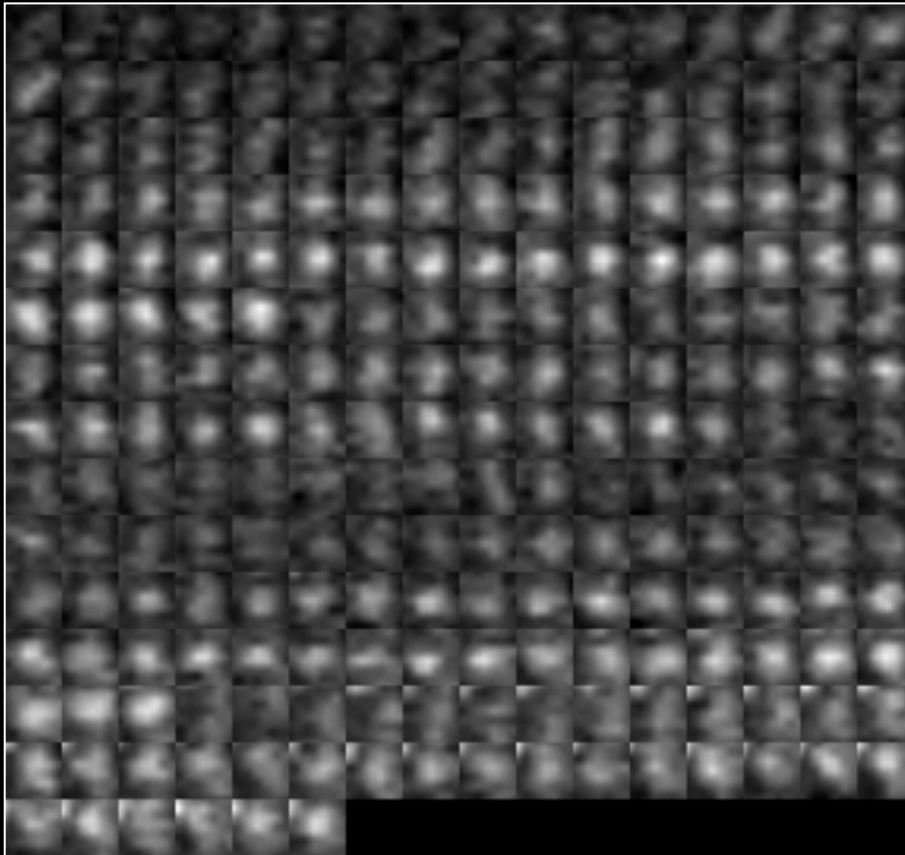
Intensity Dynamics

Accumulation of VSVG protein at the exit sites



Intensity Dynamics

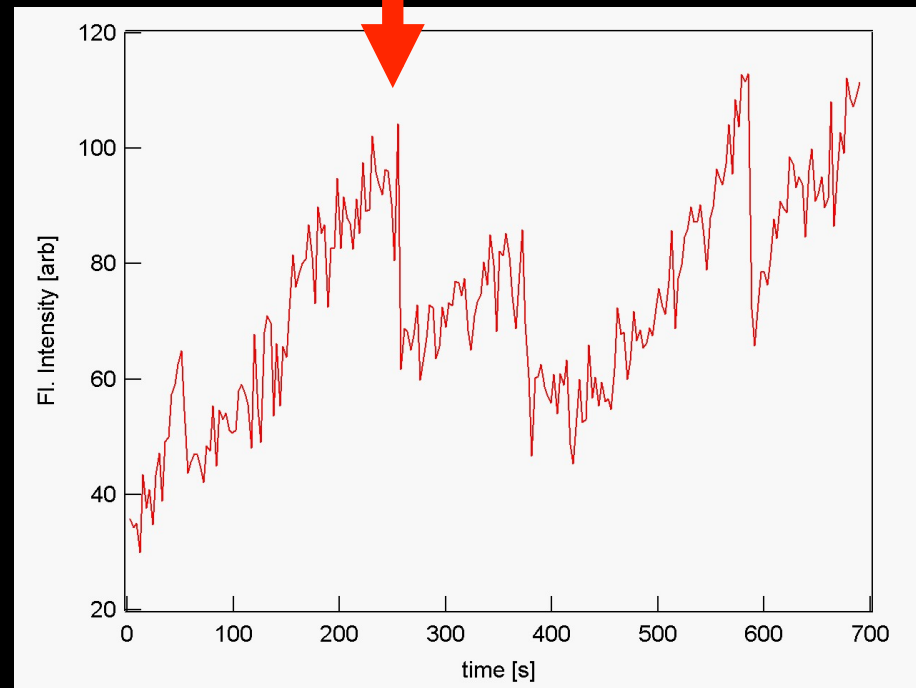
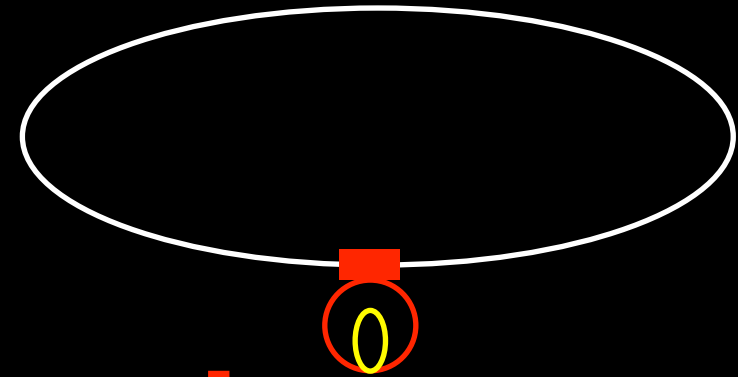
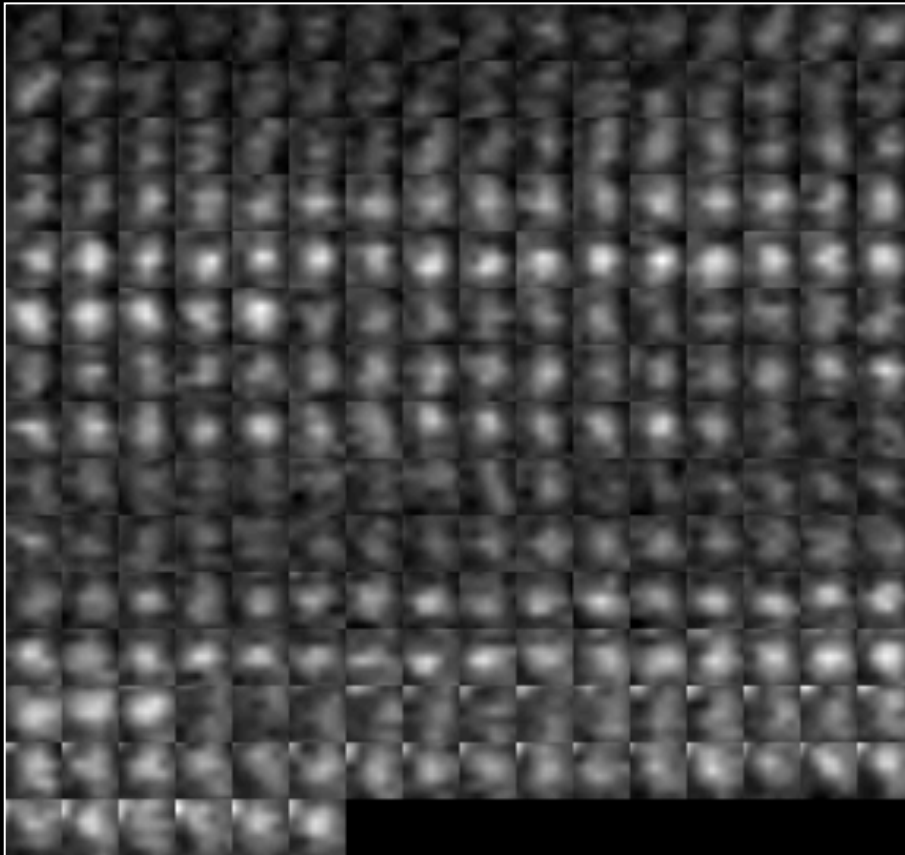
Accumulation of VSVG protein at the exit sites



→ Binding constants of VSVG protein to the ER exit site can be measured directly.

Intensity Dynamics

Accumulation of VSVG protein at the exit sites

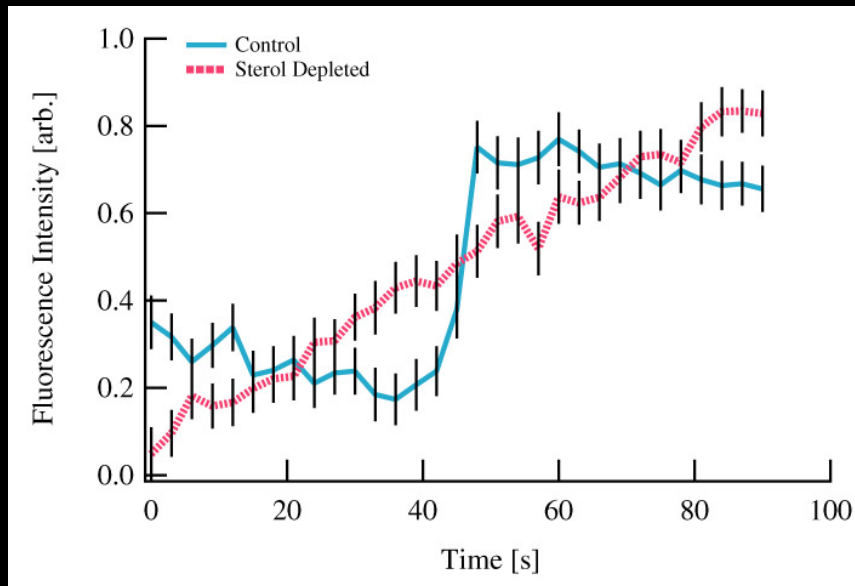


→ Binding constants of VSVG protein to the ER exit site can be measured directly.

Intensity Dynamics

Accumulation of VSVG protein at the exit sites

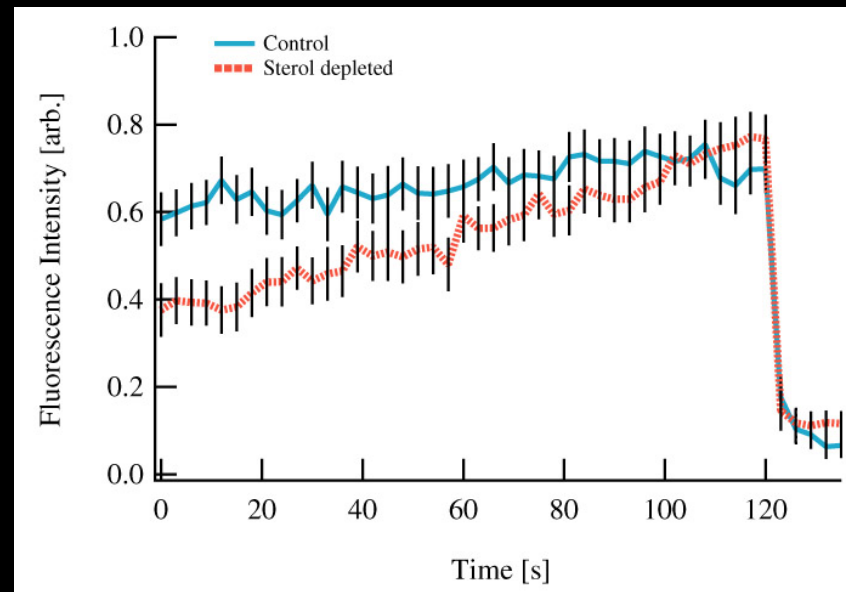
Loading



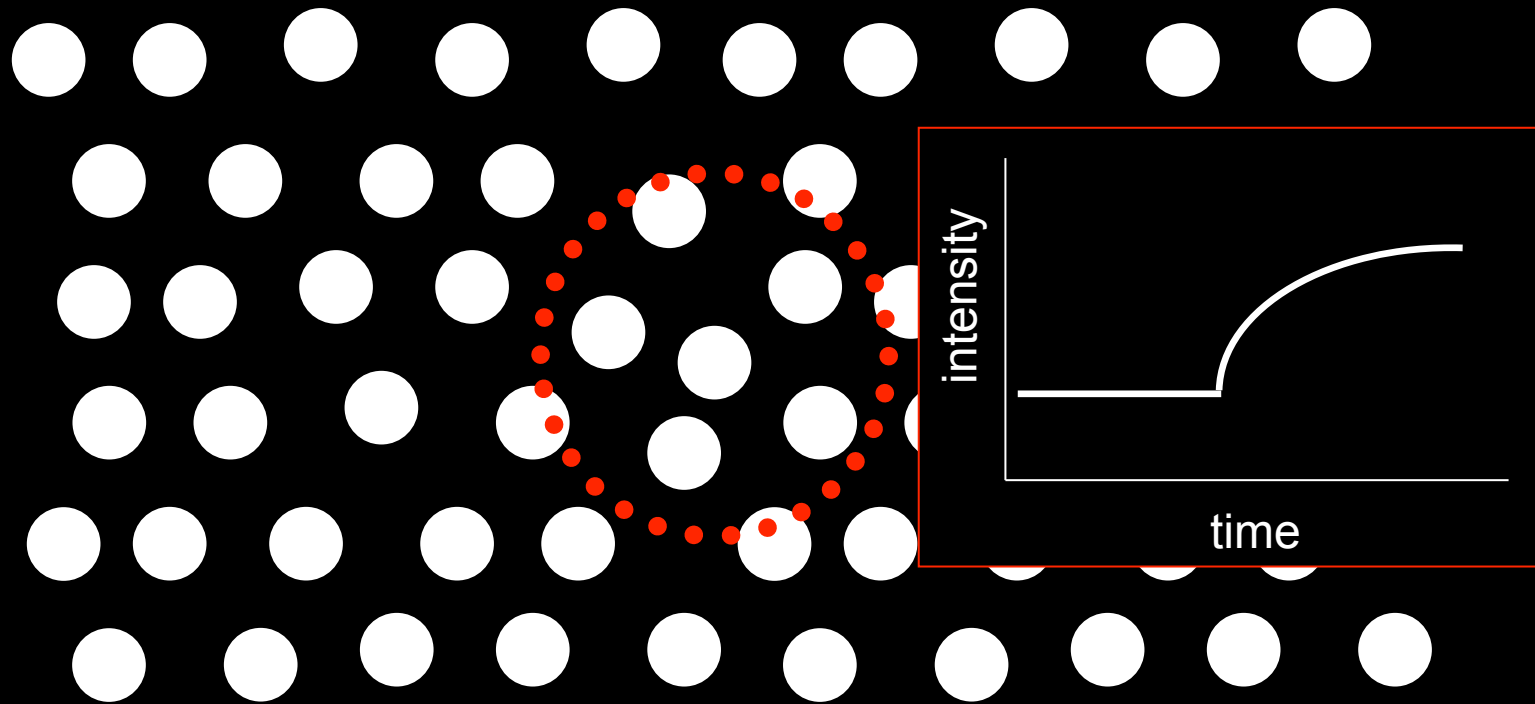
$$\frac{d[tsO45G_{ERES}]}{dt} = K_{on}[tsO45G_{free}] - K_{off}[tsO45G_{ERES}]$$

$$\left. \frac{d[tsO45G_{ERES}]}{dt} \right|_{initial} = K_{on}[tsO45G_{free}]$$

Budding



→ This strategy does not work under steady state...
→ FRAP



Static images

- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

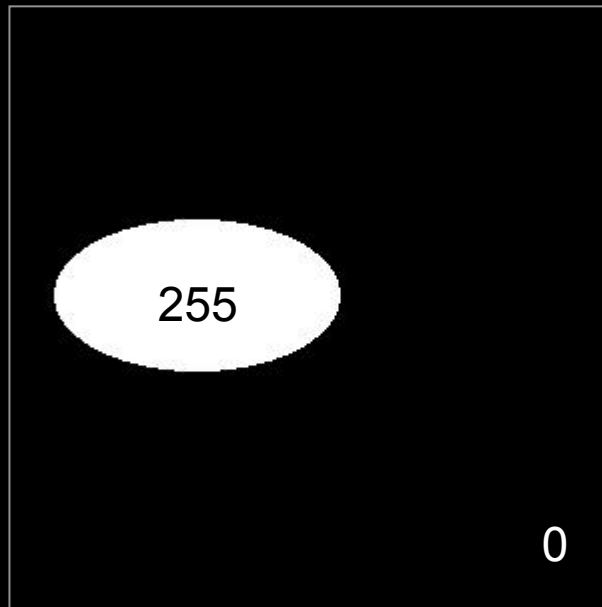
Image sequences

- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation

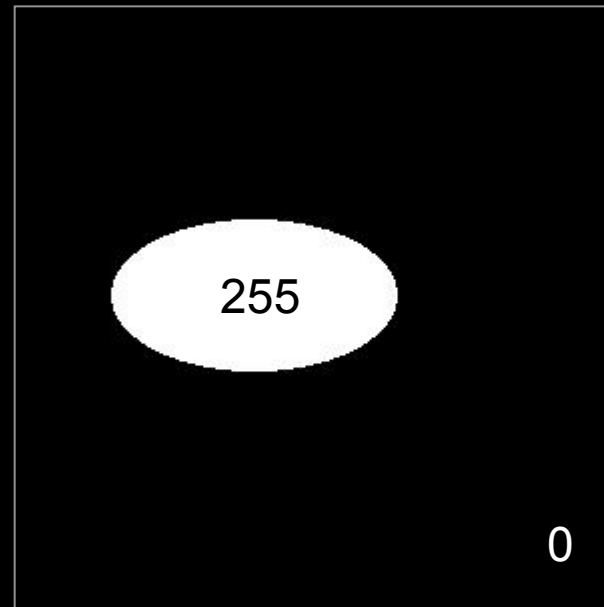
Length, Area dynamics

Measurement of Shape Changes: Area Dynamics

$$D_j(x,y) = I_{t+1}(x,y) - I_t(x,y)$$

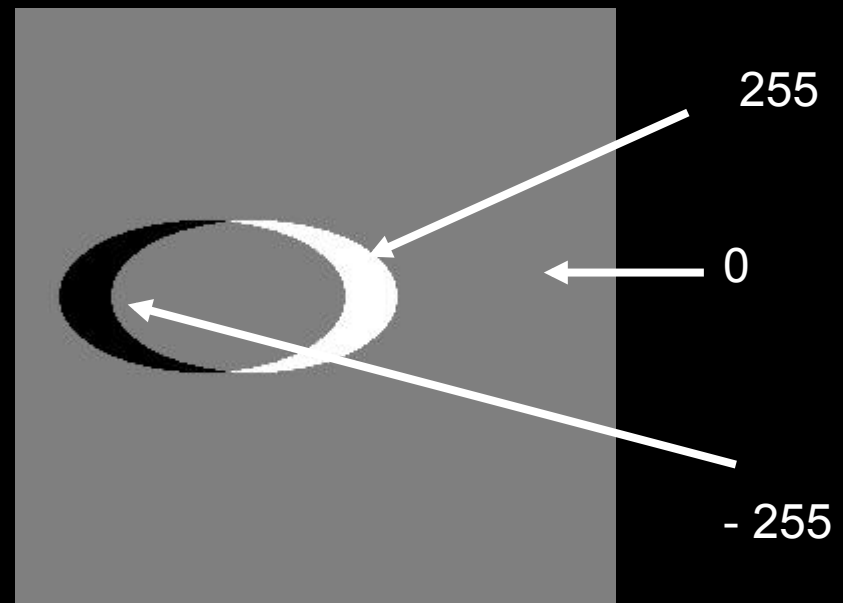


t



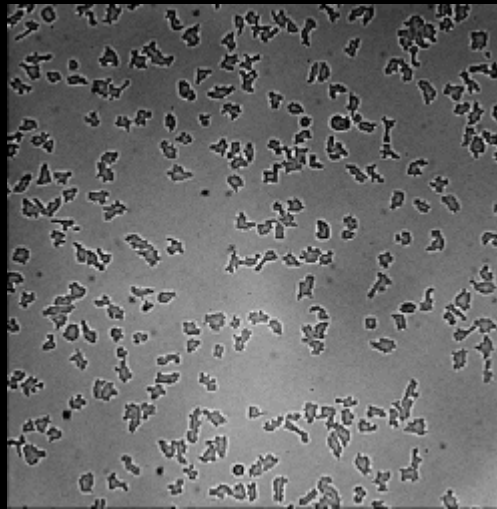
$t+1$

Image $[t+1] - \text{image}[t] =$



Measurement of Shape Changes: Area Dynamics

Light increases signaling waves

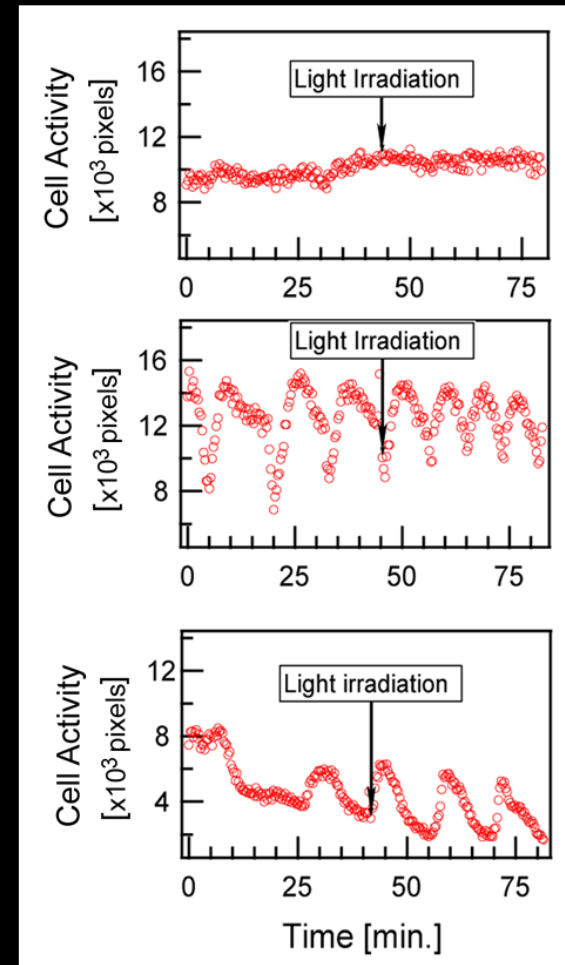


Starvation:

2hr

4hr

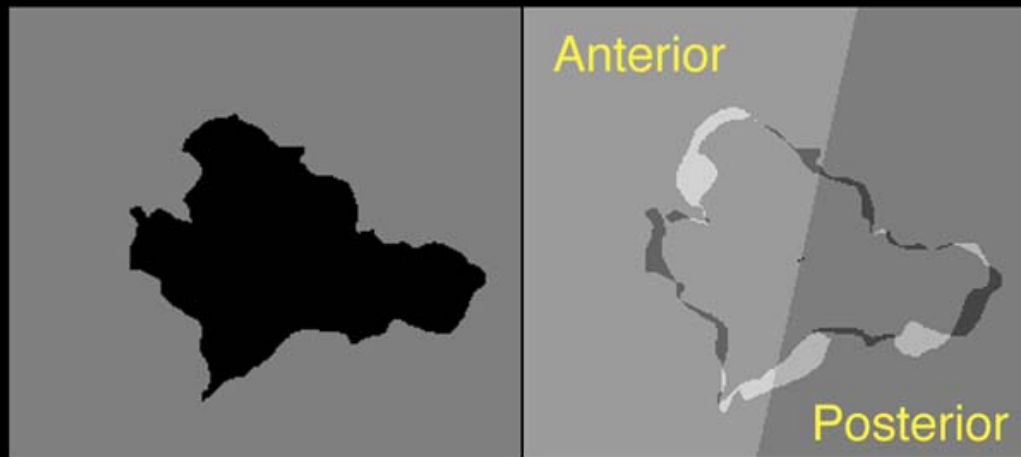
6.5hr





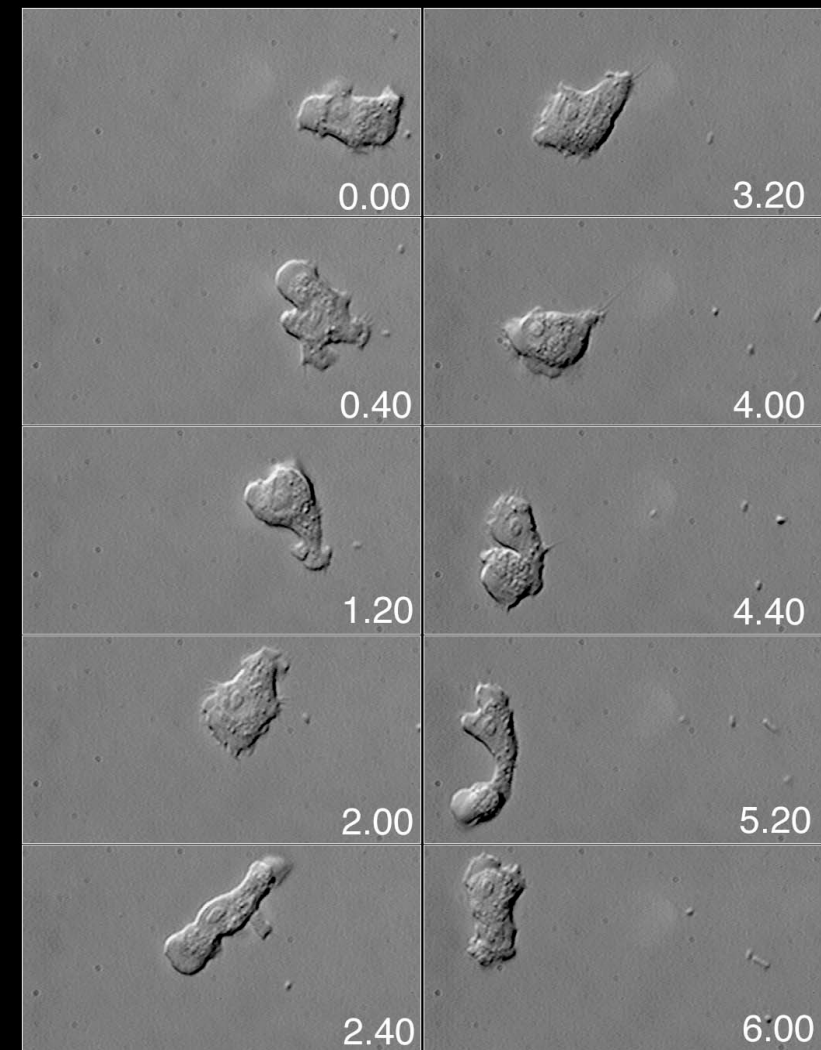
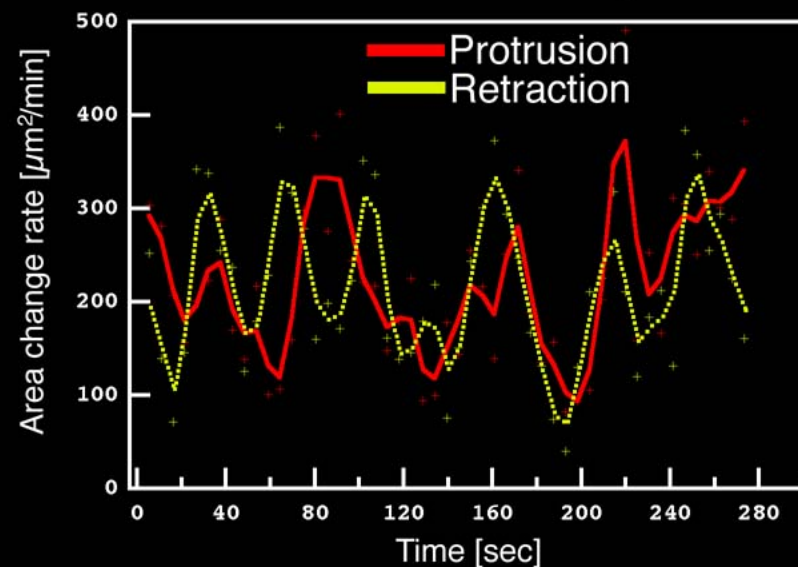
Measurement of Shape Changes: Area Dynamics

Single Cell Movement



Area changes in anterior half and posterior half.

Cell movement on a plane



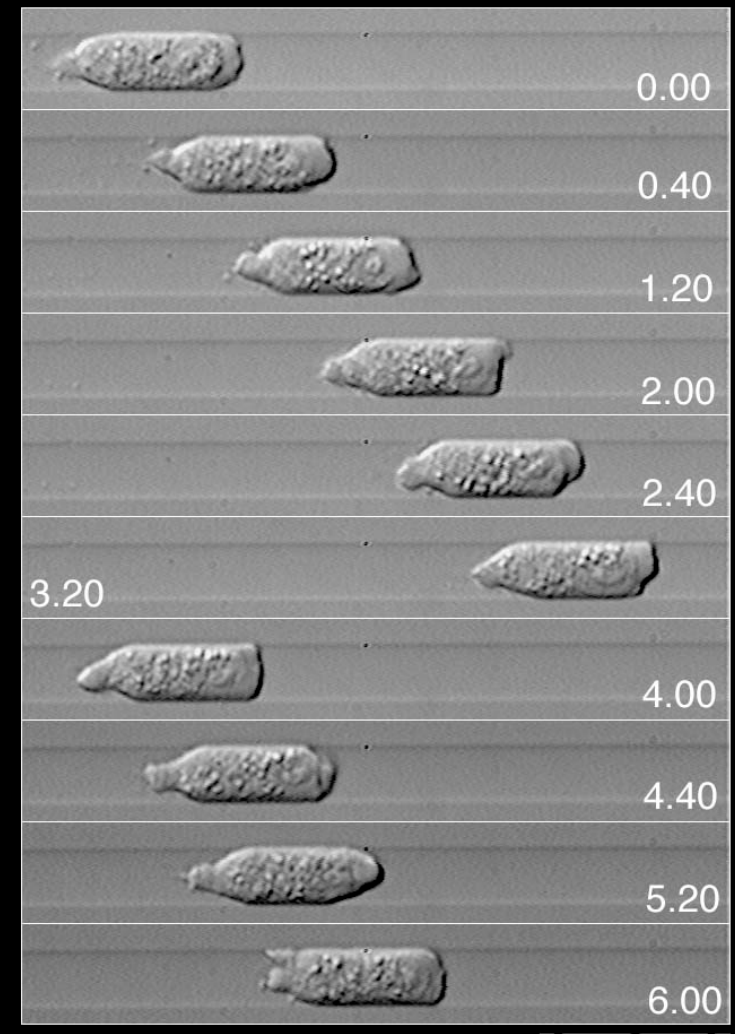
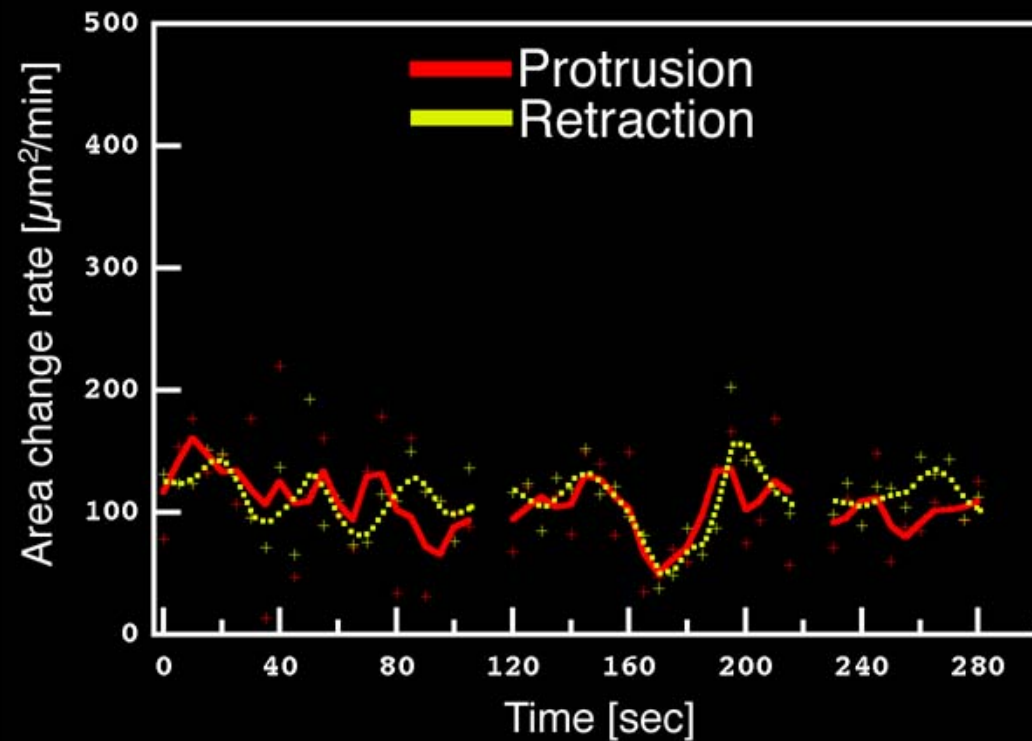
Measurement of Shape Changes: Area Dynamics

Cell Migration in agar capillary

02-27-96 19:20:55

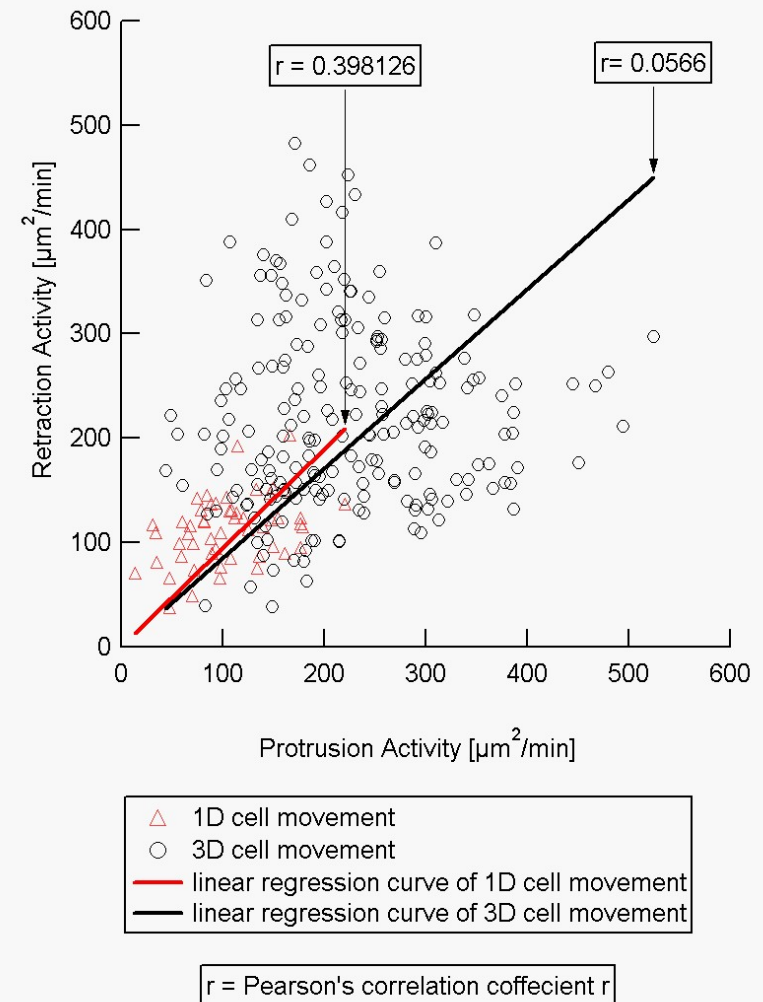
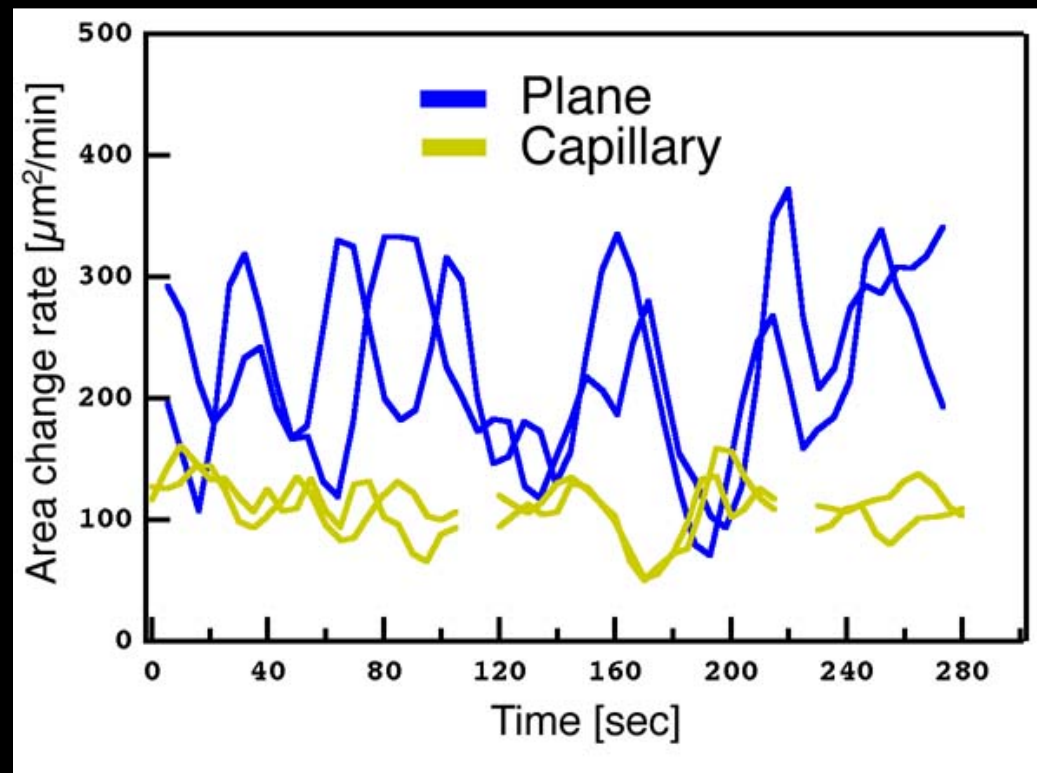


Cell movement in a capillary



Measurement of Shape Changes: Area Dynamics

Protrusion / retraction coupling

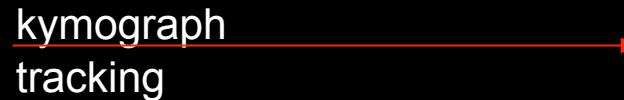


Static images

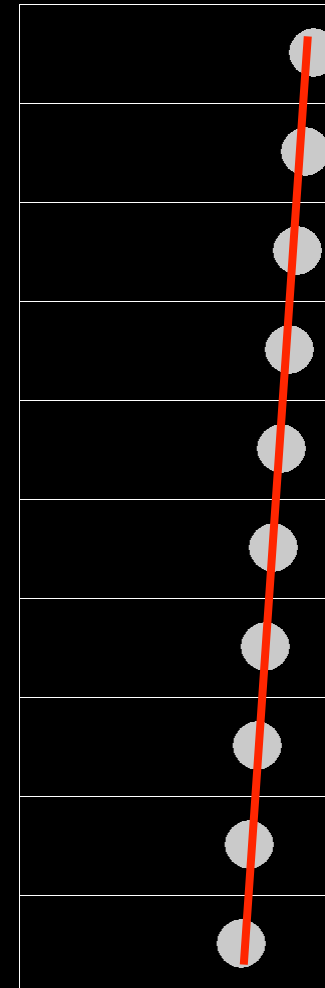
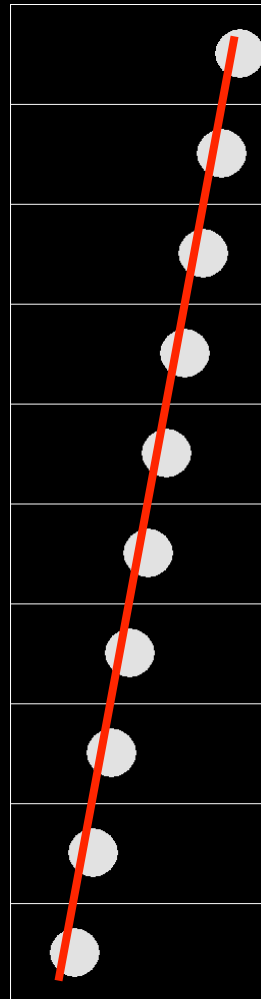
- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

Image sequences

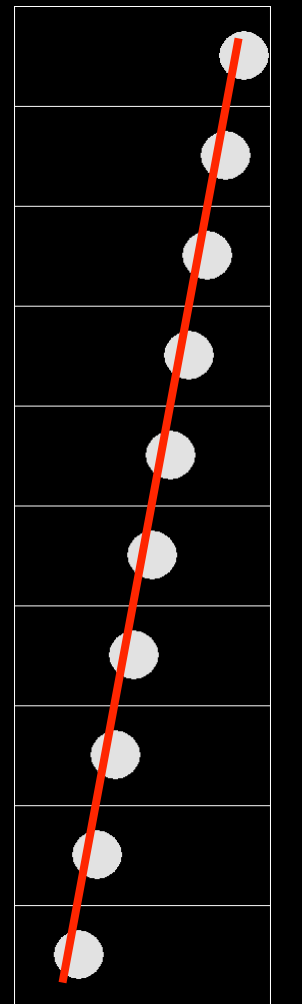
- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation



Measurement of Movement: kymograph



Measurement of Movement: kymograph

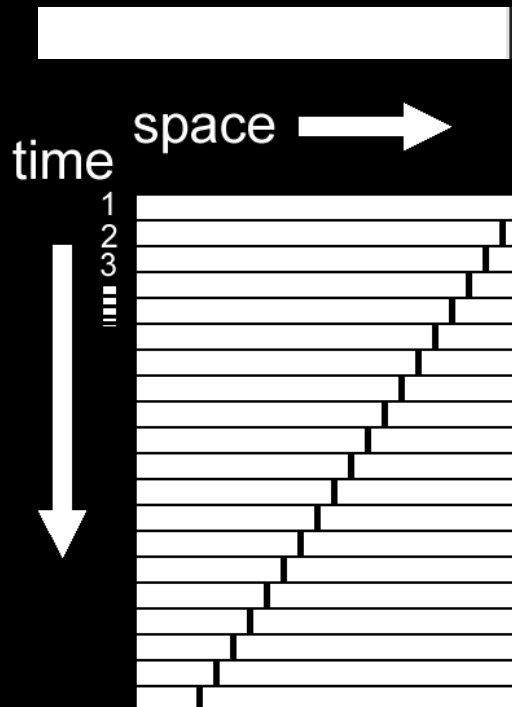
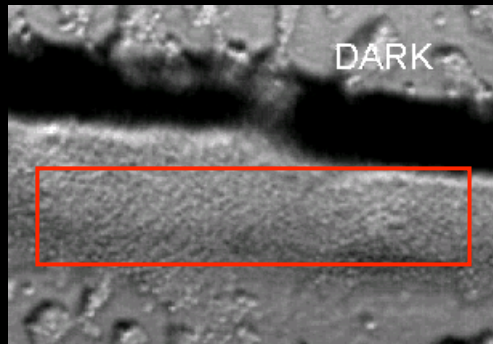


time

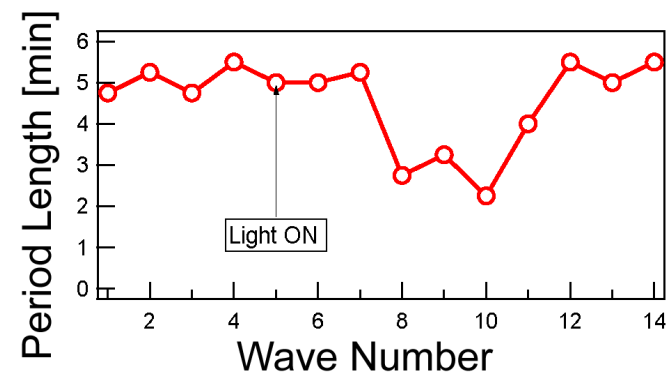
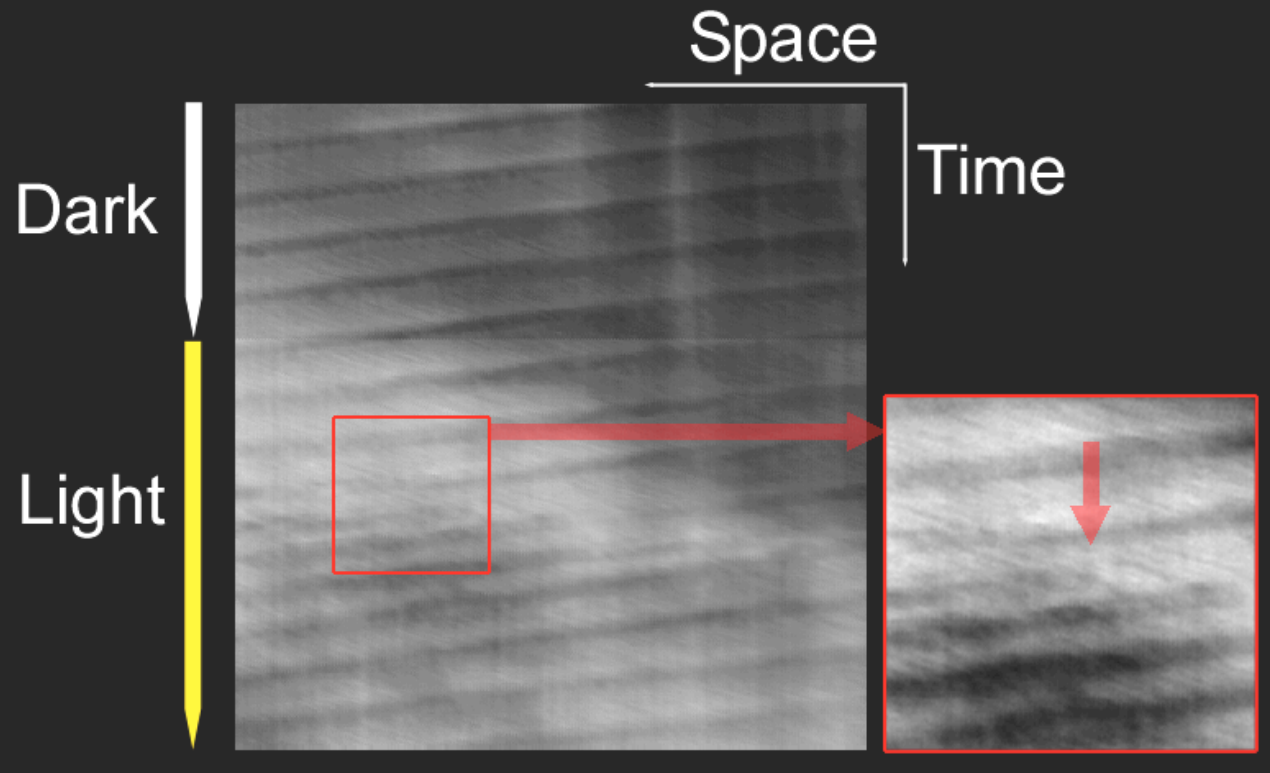
$$\text{Speed} = \text{Displacement} / \text{Time}$$

displacement

kymograph



Light Induces New Wave



Measurement of Movement

Static images

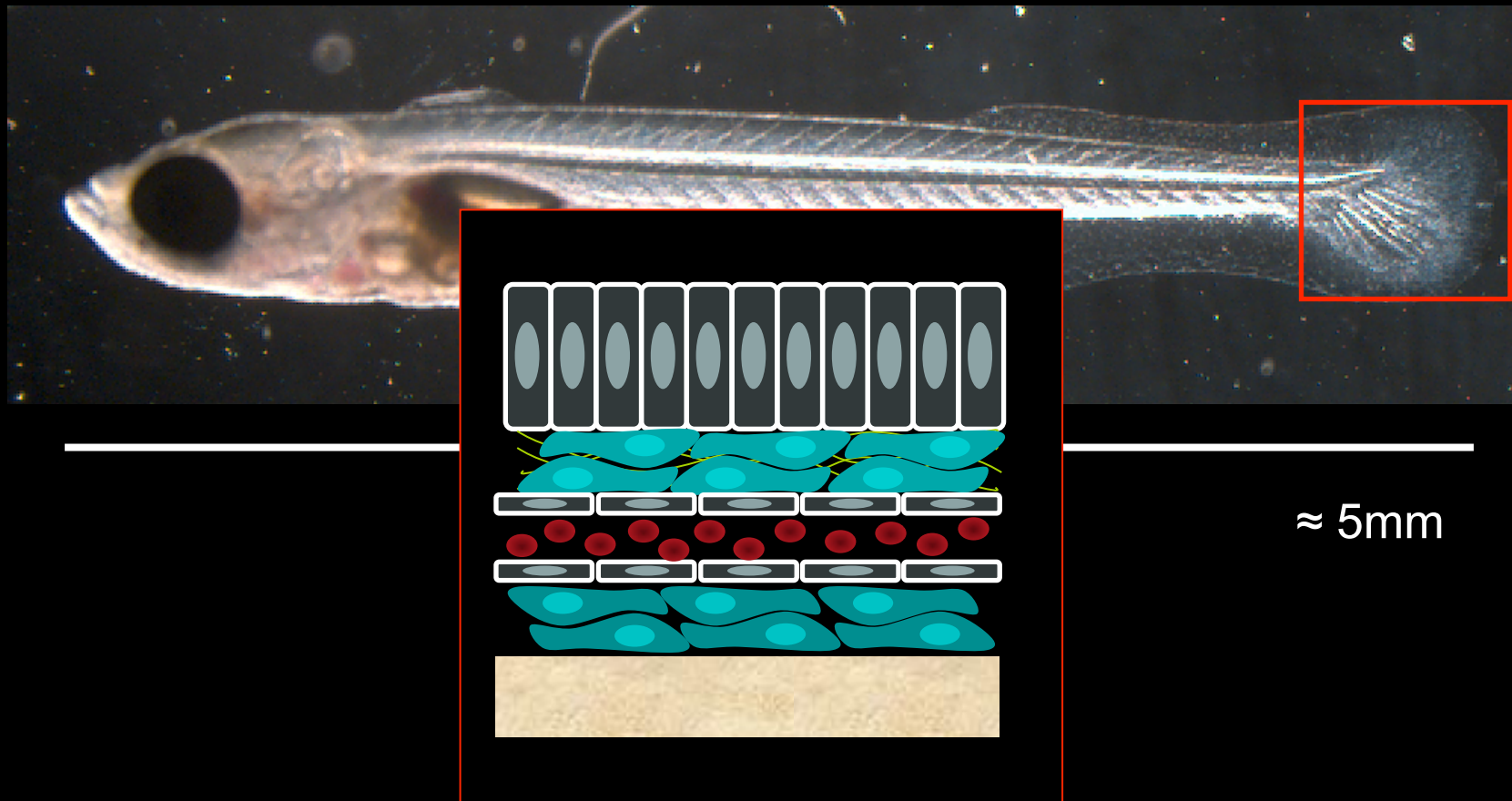
- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

Image sequences

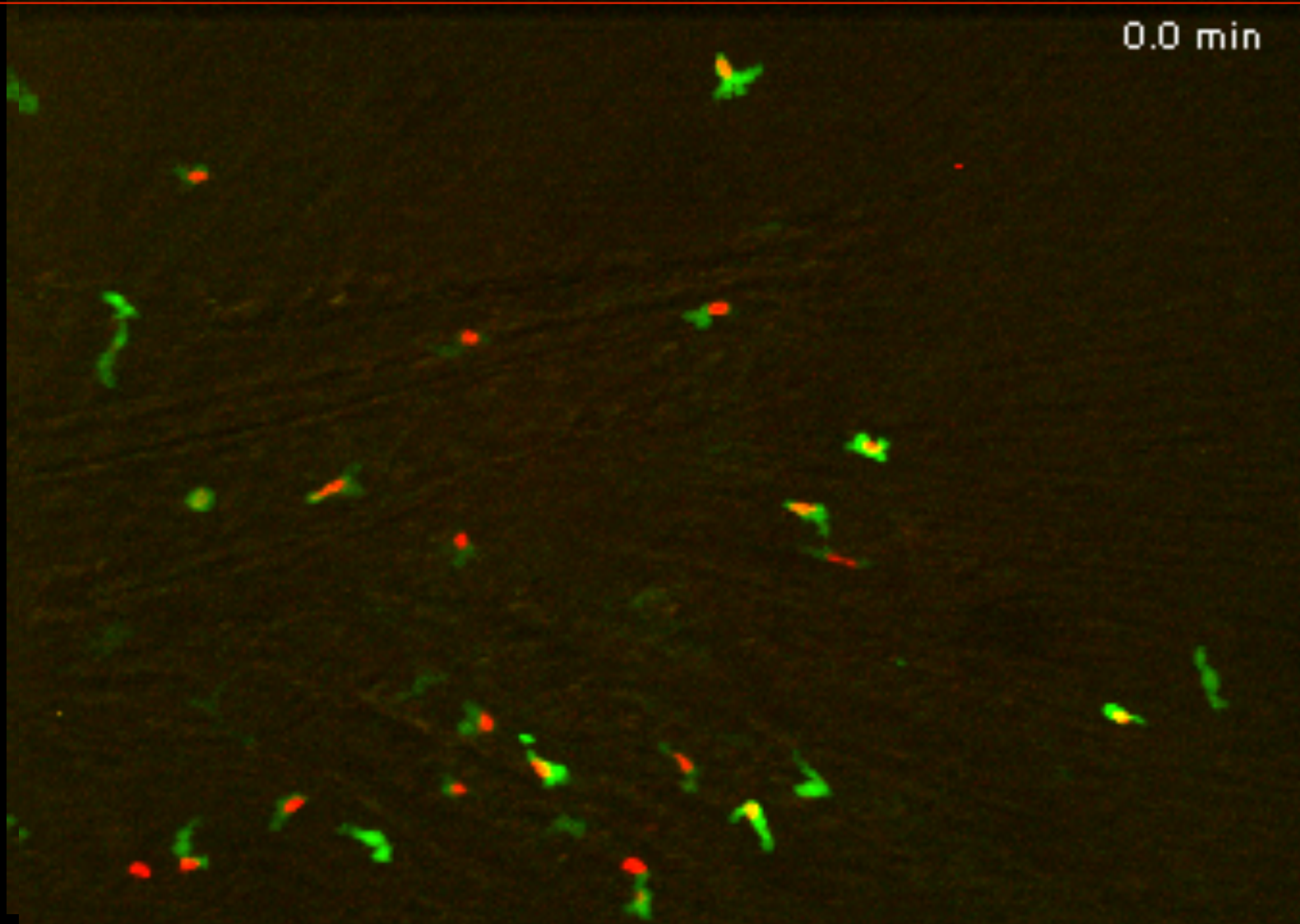
- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation



Measurement of Movement: cell tracking



Cell migration within *Medaka* tail-fin



1. TBL (unknown protein, random insertion & screening)
2. FMPO (myeloperoxidase promoter, Histone-mRFP, macrophage-marker)

Green GFP
Red GFP

Measurement of Movement: cell tracking

Pattern Matching Method

- Cross-Correlation

$$C(x, y) = \sum_{i=0}^{n-1} \sum_{j=0}^{m-1} I(x+i, y+j) \{K(i, j) - \bar{K}\}$$

- Sum of Absolute Difference

$$SAD(x, y) = \sum_{i=0}^{n-1} \sum_{j=0}^{m-1} |I(x+i, y+j) - K(i, j)|$$

$$x_c = \frac{\sum x \{C(x, y) - T\}}{\sum \{C(x, y) - T\}}$$

$$y_c = \frac{\sum y \{C(x, y) - T\}}{\sum \{C(x, y) - T\}}$$

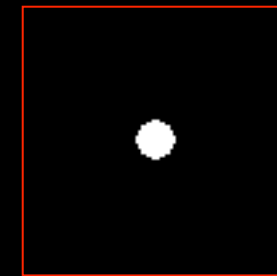
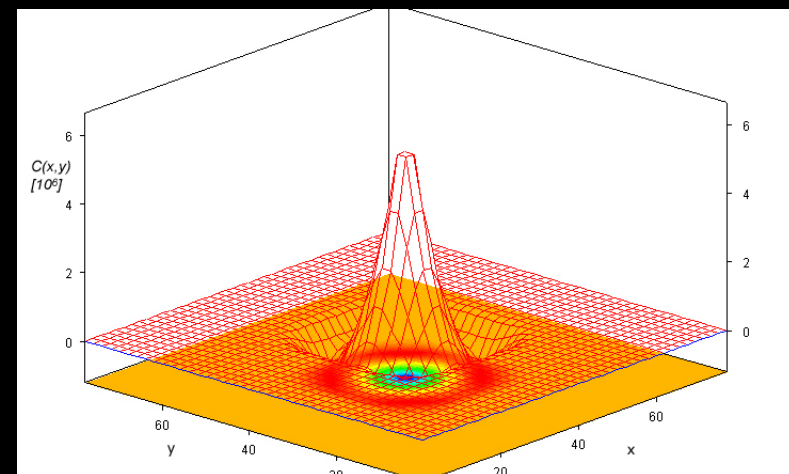
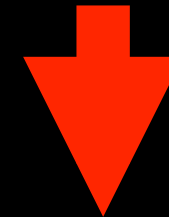


Image Frame

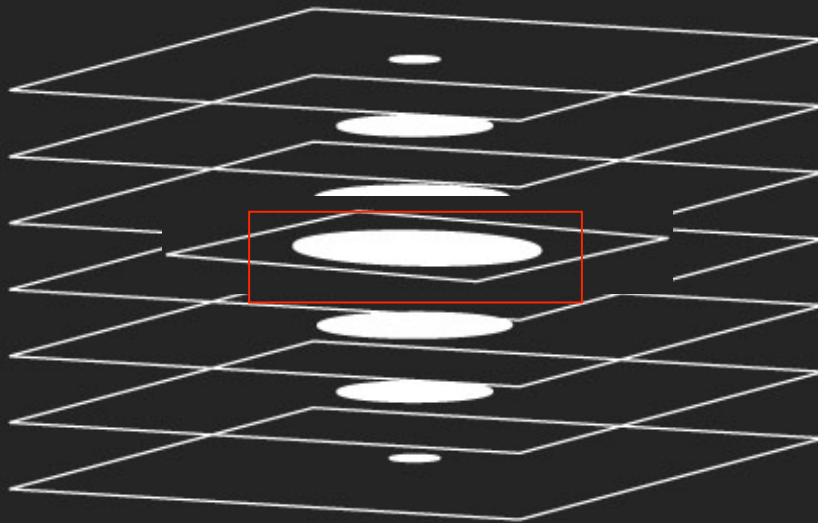


Kernel
(Template)

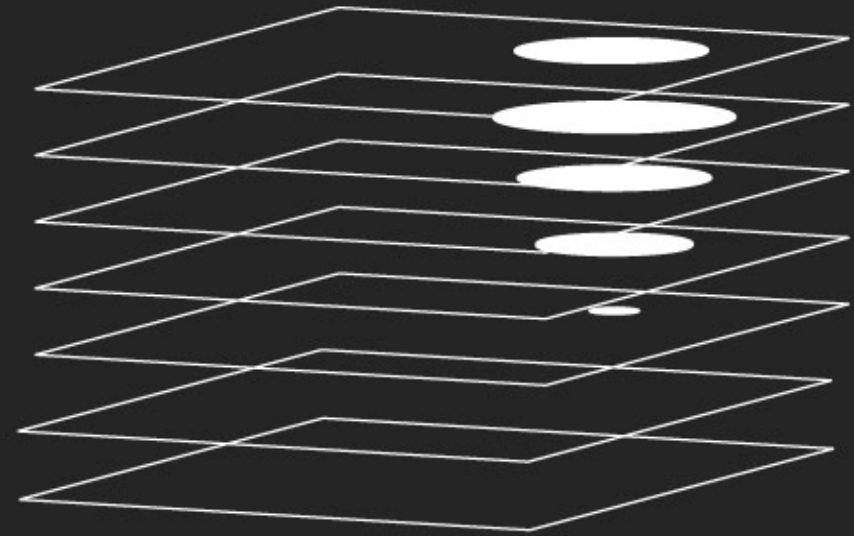


Measurement of Movement: cell tracking

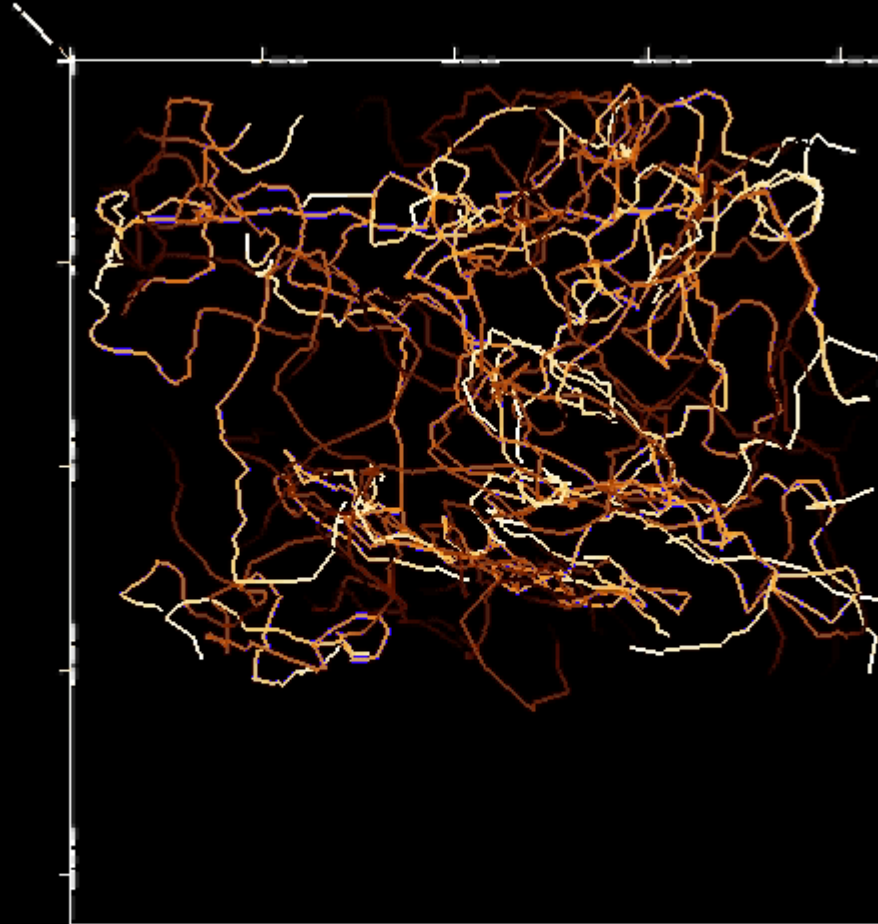
Pattern matching in 3D



t

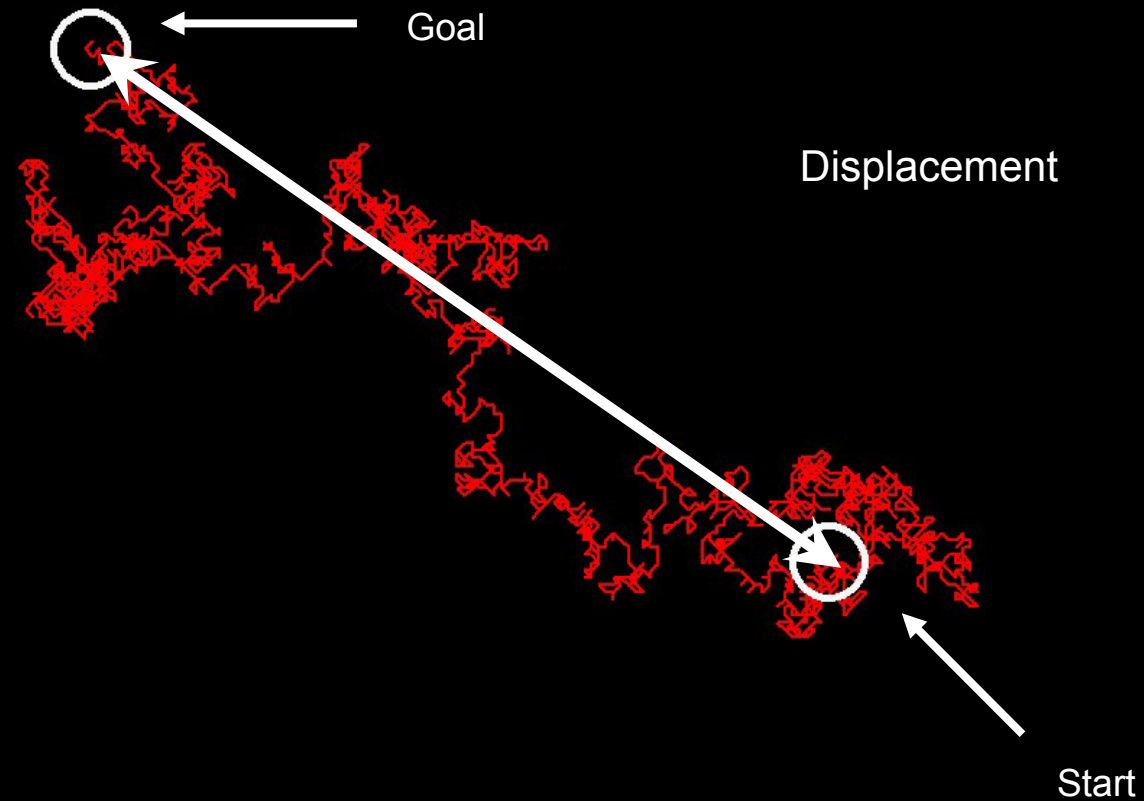


$t+1$



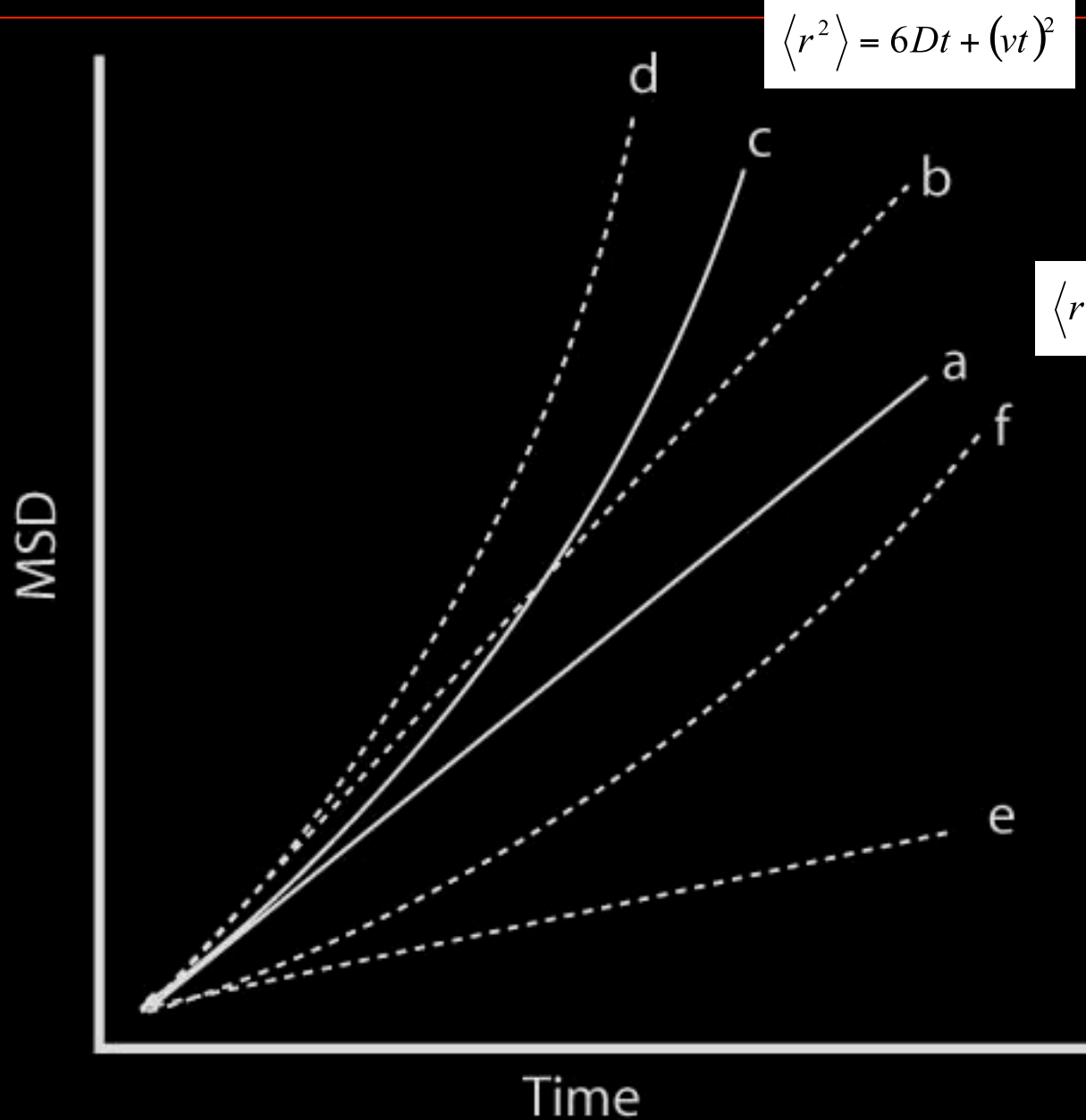
3D plotting

Measurement of Movement: cell tracking



$$\text{Mean Square Displacement (MSD)} = \langle (\text{Displacement})^2 \rangle$$

Measurement of Movement: cell tracking



$$\langle r^2 \rangle = 6Dt + (vt)^2$$

$$\langle r^2 \rangle = 6Dt$$

$$\langle d^2(t) \rangle = 6\mu \left\{ -P \left[1 - e^{-t/P} \right] \right\}$$

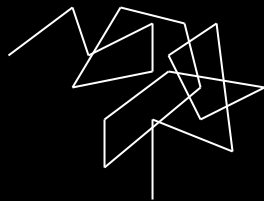
Fuerth's Formula

Measurement of Movement: cell tracking

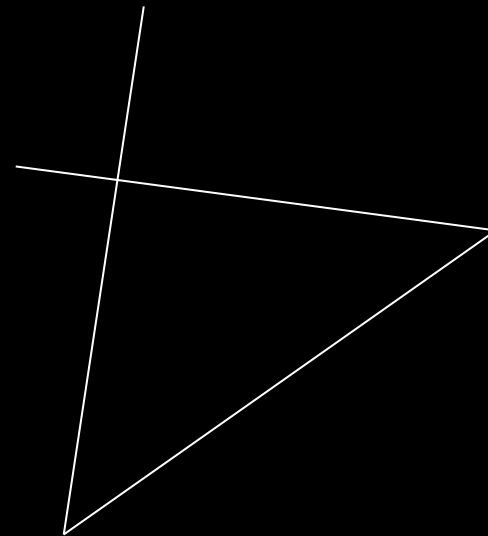
Fuerth's formula: Persistence Time

“characteristic amount of time between directional changes in cell movement” *1

$$\langle d^2(t) \rangle = 6\mu \left\{ -P \left[1 - e^{-t/P} \right] \right\}$$

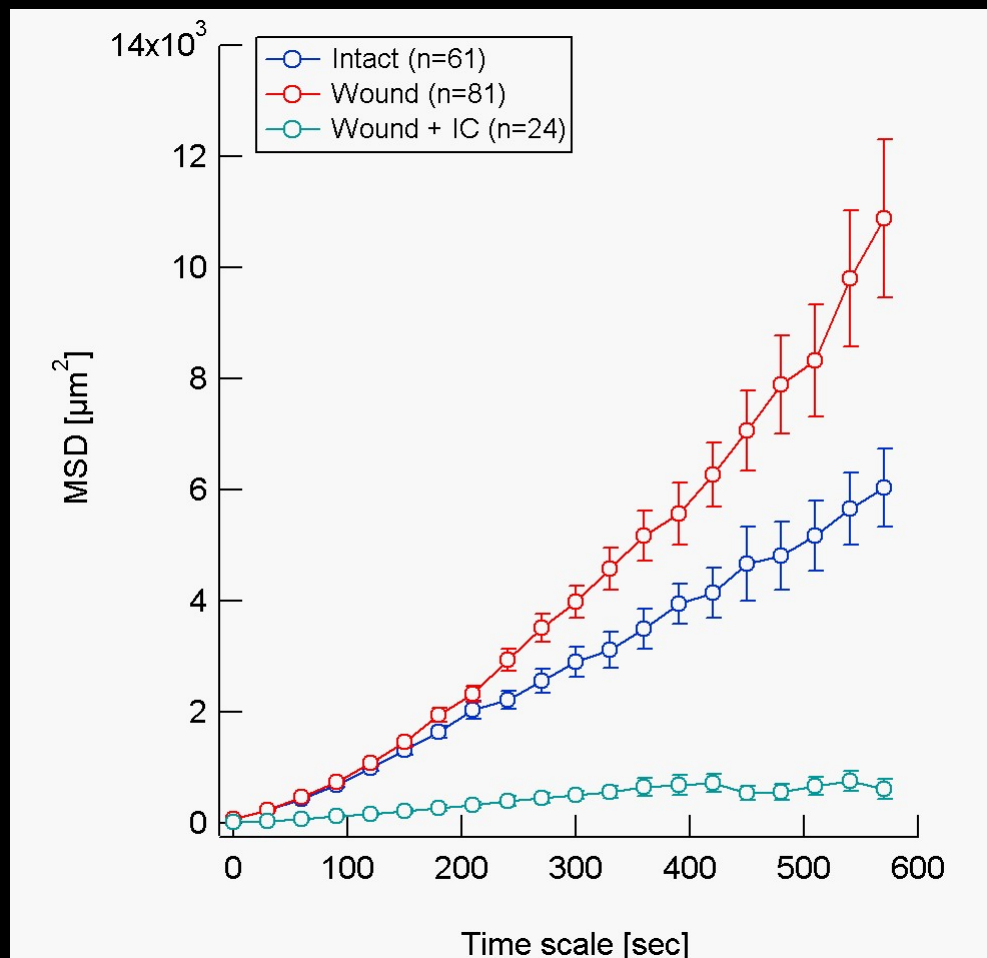


Short Persistence Time



Long Persistence Time

1. Bergman & Zygorakis (1999) Biomaterials
2. Selmeczi et. al. (2005) Biophys. J.
3. Fuerth (1920) Z. Physik. 2:244-256



$\mu = 17.4 \pm 1.8 \mu\text{m}^2/\text{min}$
 $P = 44 \pm 28 \text{ sec}$

Lower Cell Activity and
 Suppressed Chemotaxis
 (PI3 kinase inhibitor)

μ : Random Migration Coefficient
 P : Persistence Time

← Random Walk + Directed Motion

$\mu = 291 \pm 23.4 \mu\text{m}^2/\text{min}$
 $P = 282 \pm 40 \text{ sec}$

← Random Walk

$\mu = 114 \pm 1.8 \mu\text{m}^2/\text{min}$
 $P = 76 \pm 6 \text{ sec}$

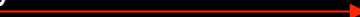
	Average Velocity [$\mu\text{m}/\text{min}$]	Persistence
Intact (61 tracks)	14.1 ± 10.1	0.48 ± 0.21
Wounded (81 tracks)	14.3 ± 8.7	0.67 ± 0.20
Wounded +Inhibitor (24 tracks)	5.9 ± 5.4	0.40 ± 0.22

Static images

- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

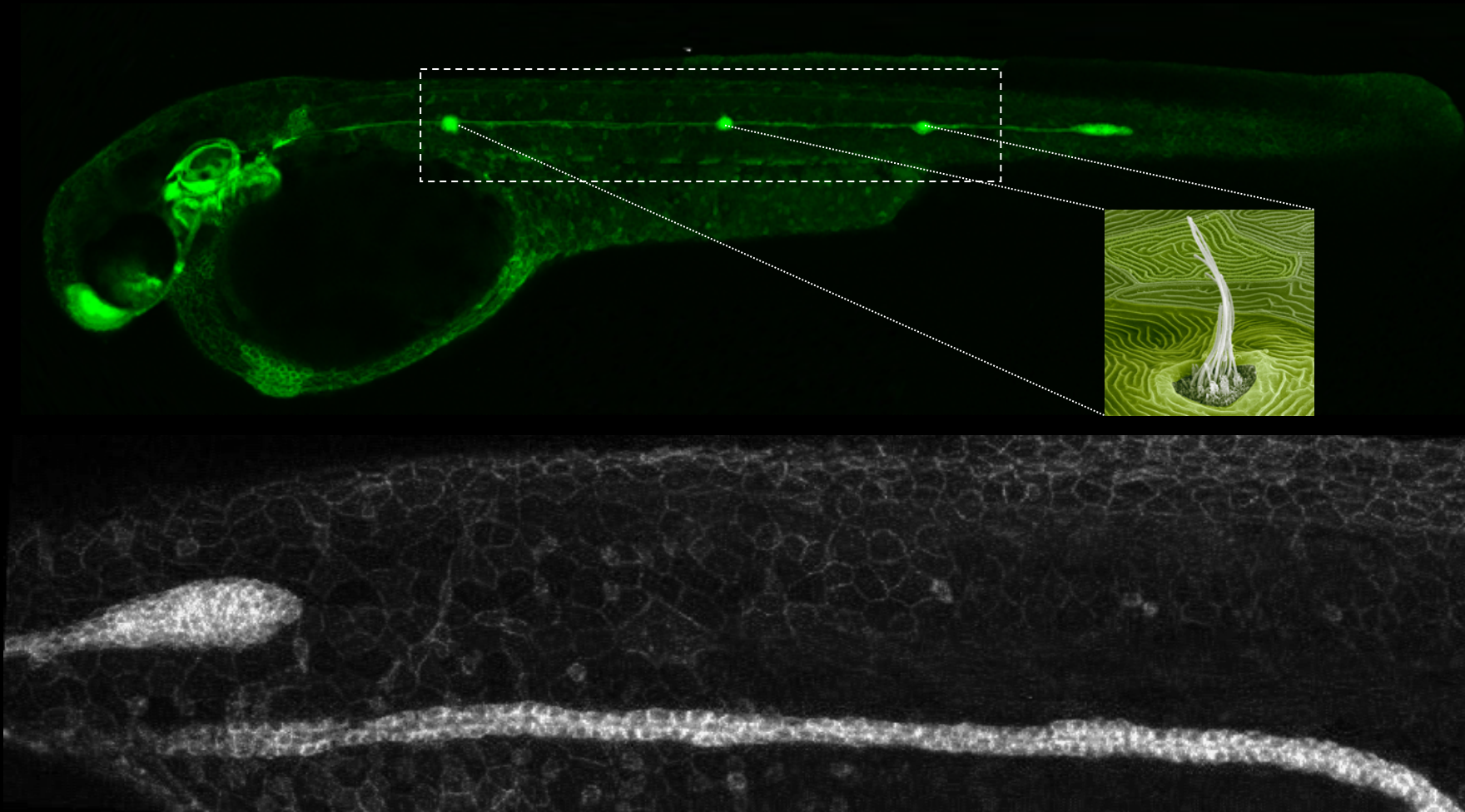
Image sequences

- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation



The zebrafish lateral line: A model system for in vivo collective cell migration

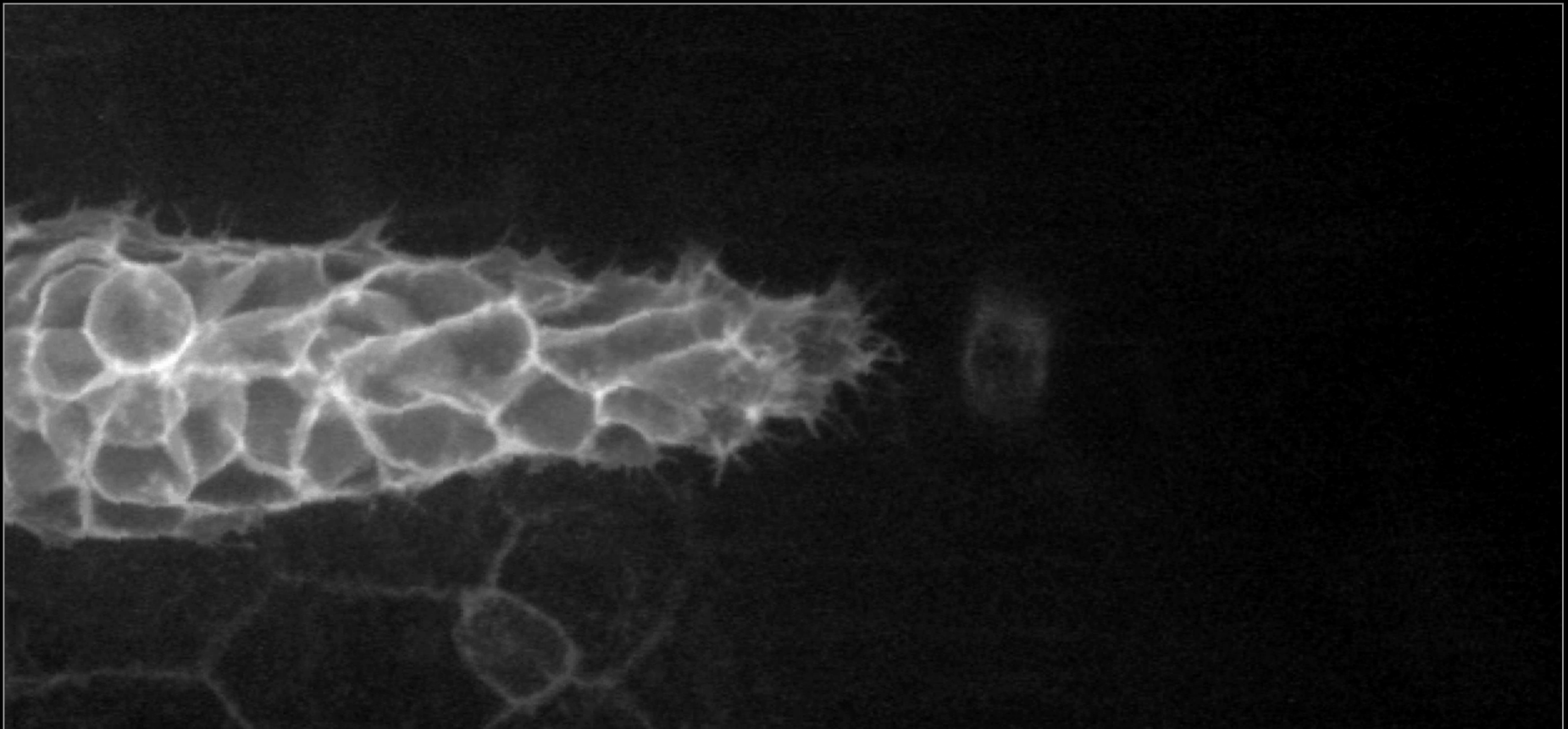
Darren Gilmour lab@EMBL



- The cells migrate efficiently as a highly coordinated unit along a predefined path.
- The forming organs repeatedly appear *de novo* within the moving tissue.
- Cells fated to become organs decelerate at the trailing edge and arrest migration.

How does this group move in a coordinated way?

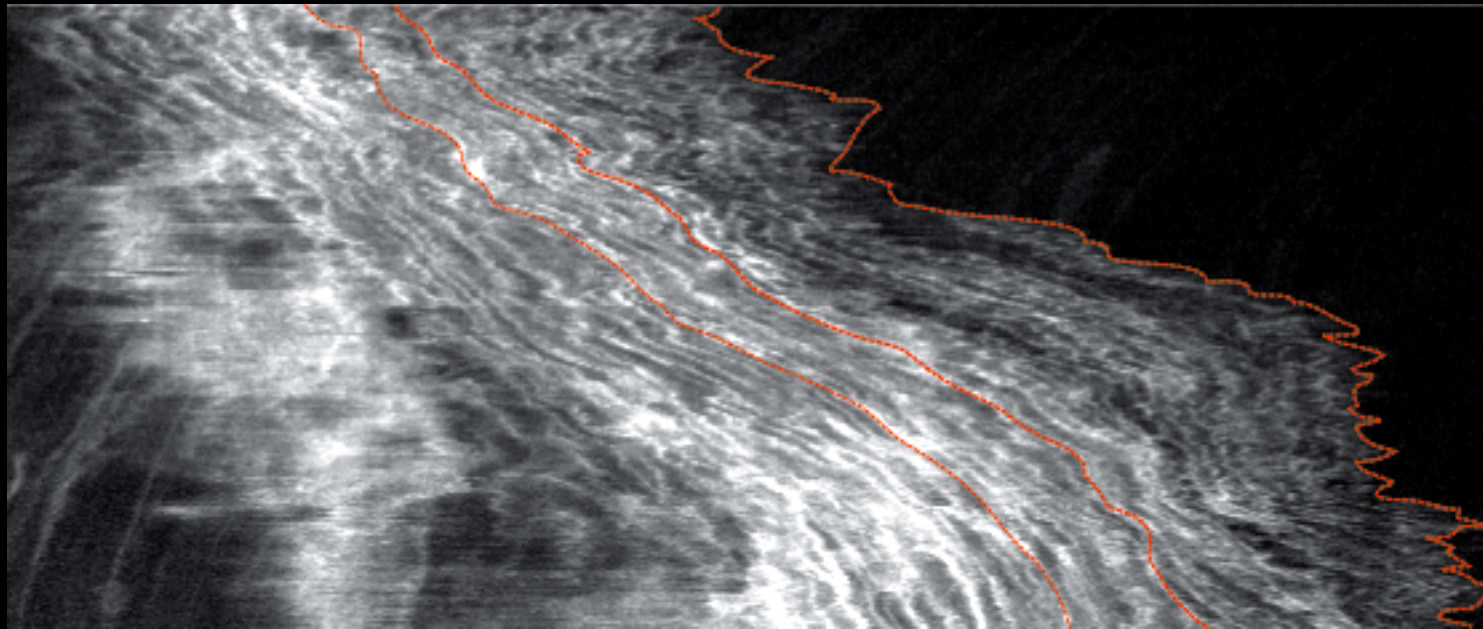
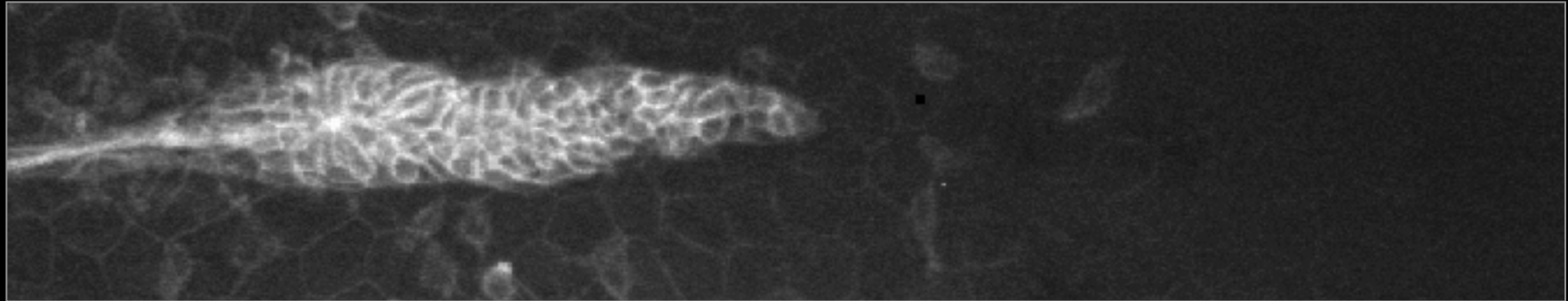
Darren Gilmour lab@EMBL

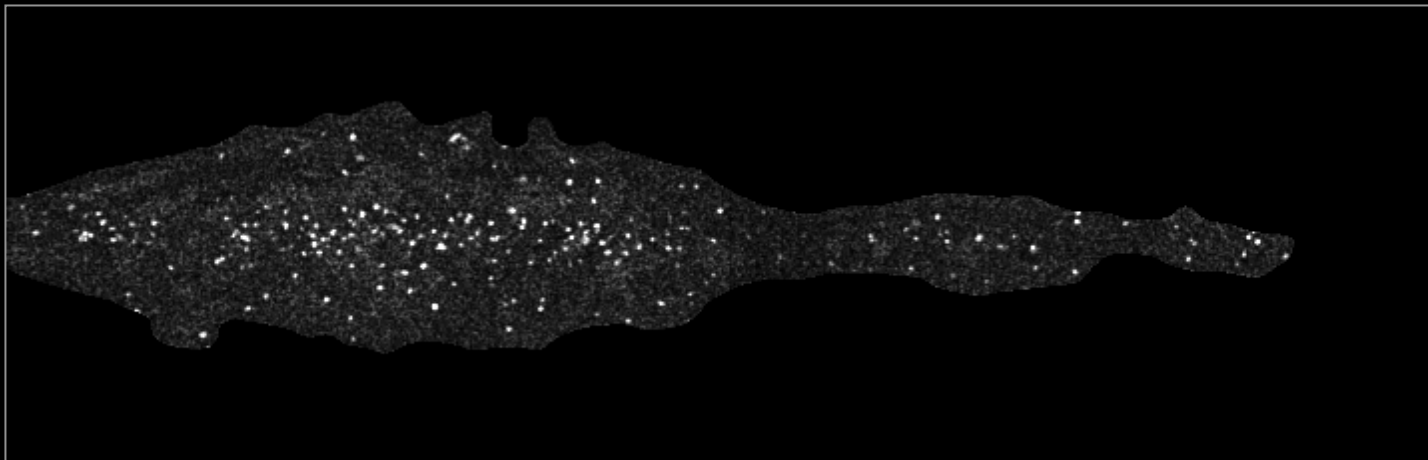
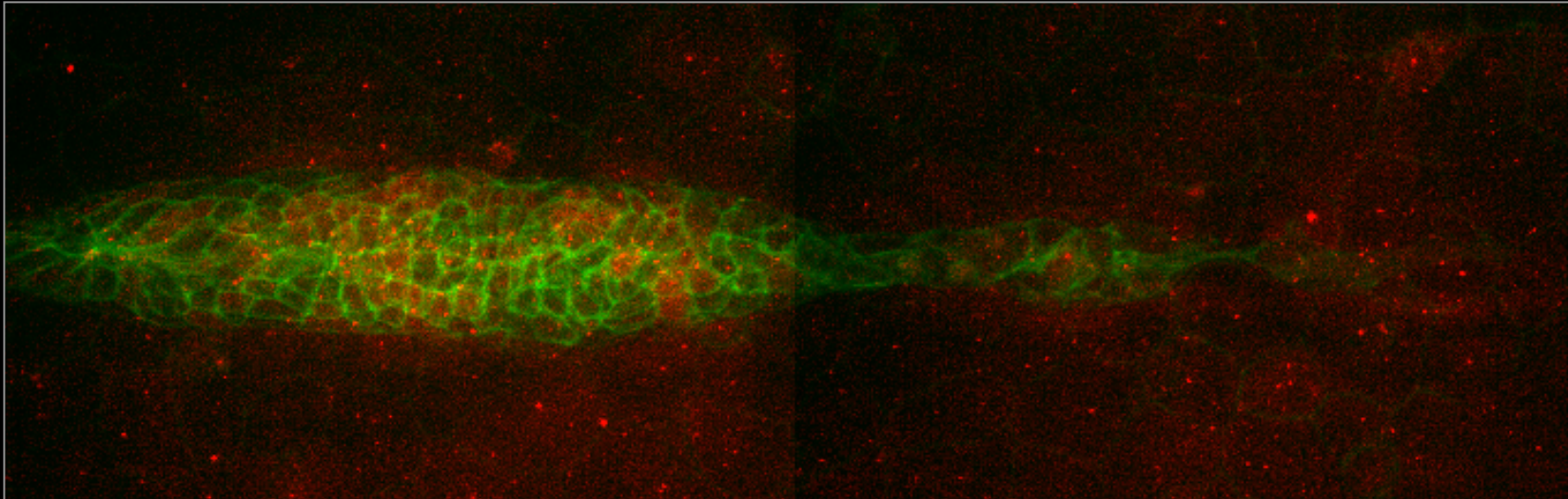


Many cells of the pLLP appear to respond directly to extrinsic guidance cues.

Loss of FGF signaling *uncouples* the migration of leaders and followers

Darren Gilmour lab@EMBL





Task: Gulcin's Project
quantitative analysis of the formation of hair cell "garlic" aggregates

Tool: Centrin labeling and image analysis
FGF inhibitor, two hours after the wash-out FGF inhibitor

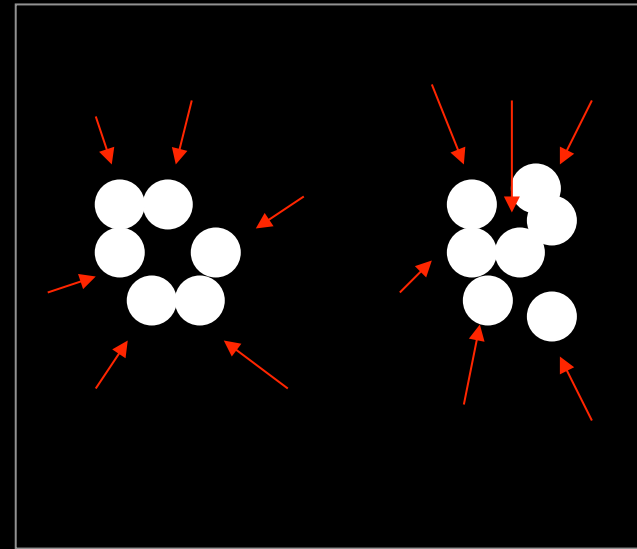
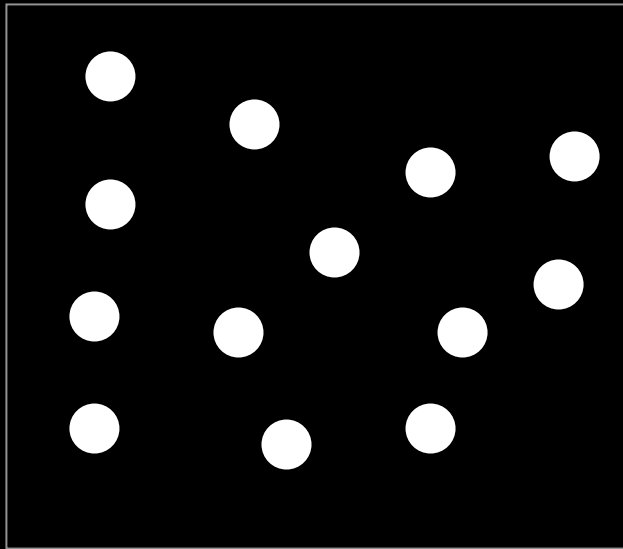
Analysis of Centrin Distribution Dynamics

General Parameters

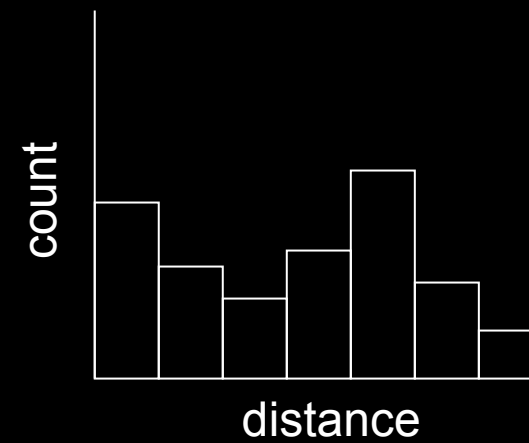
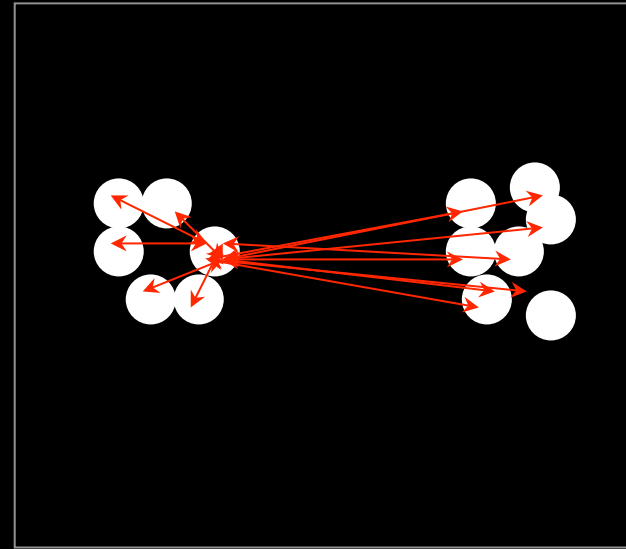
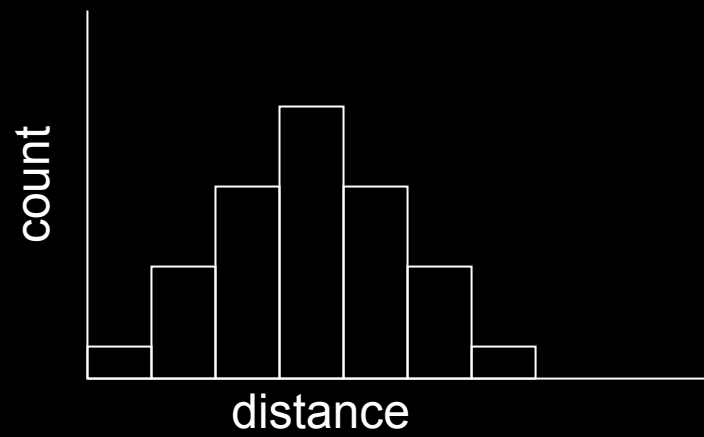
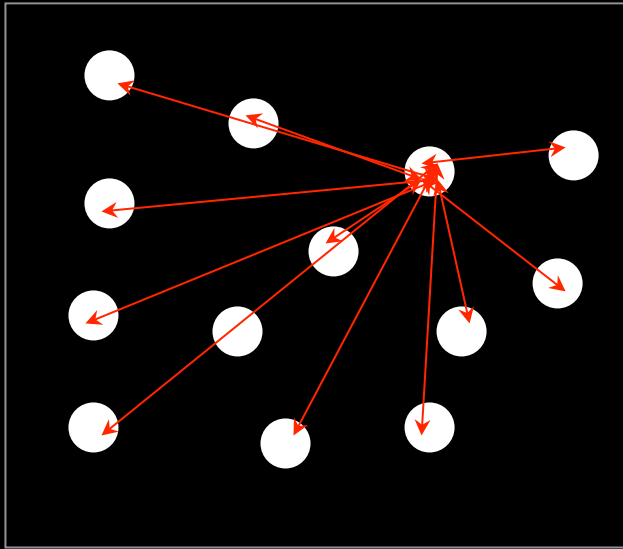
- Velocity
- Directionality

Distribution Function

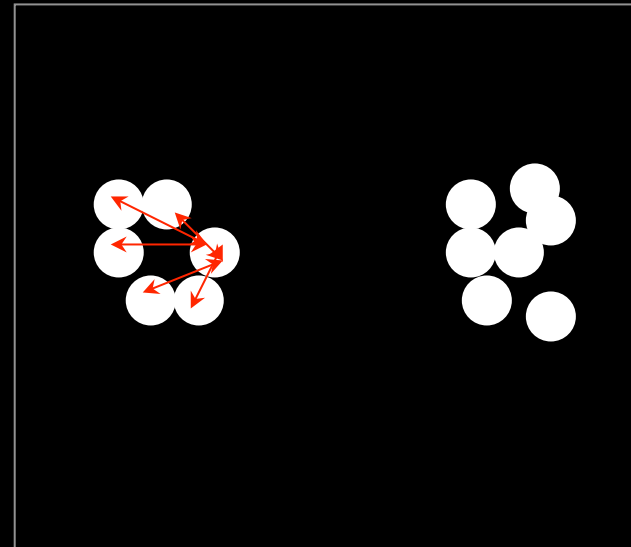
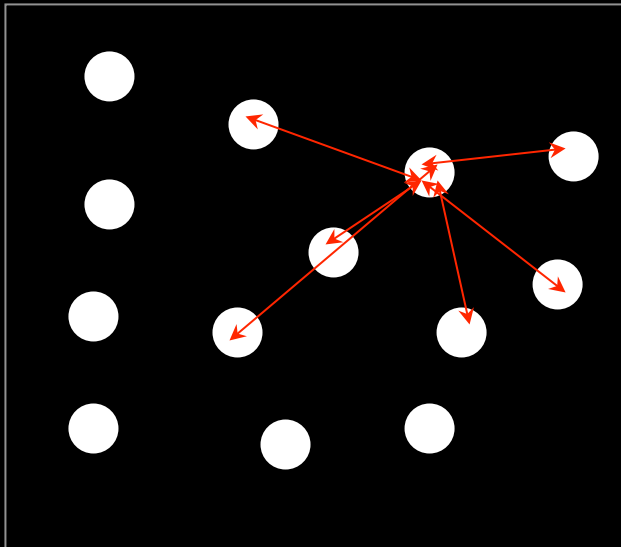
- how close are the centrin?
→ **Sparse to Dense**



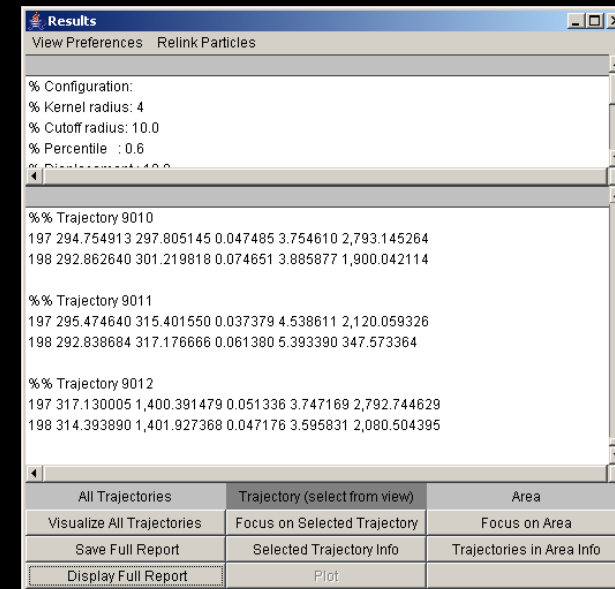
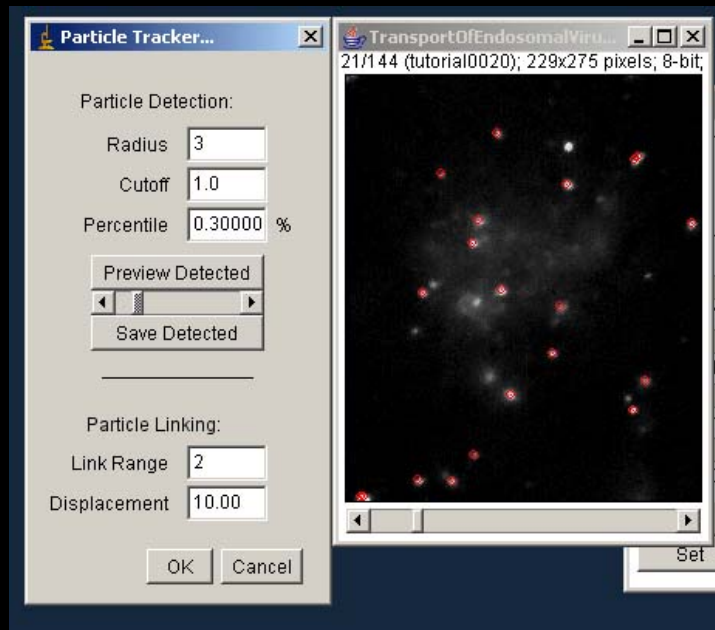
Distribution Function



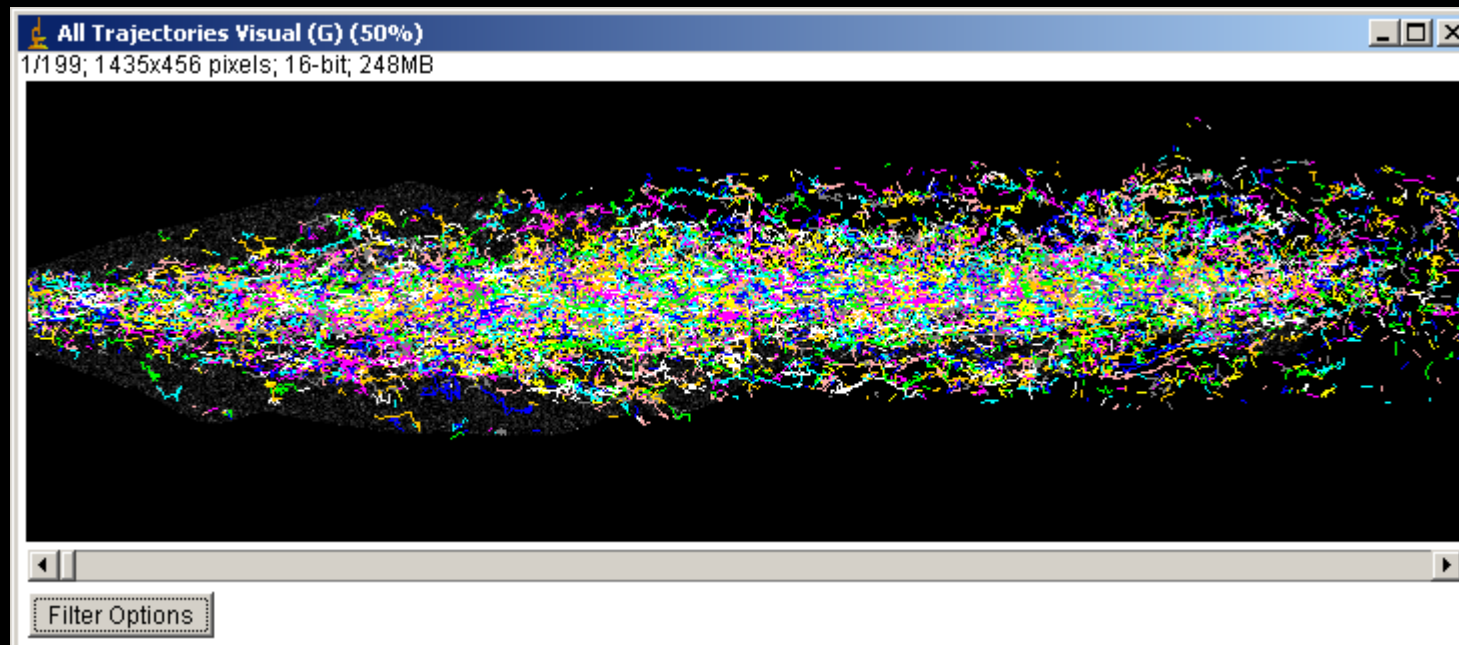
Average distance to nearest 5 centrans



Particle Tracking: using ImageJ Particle Tracker

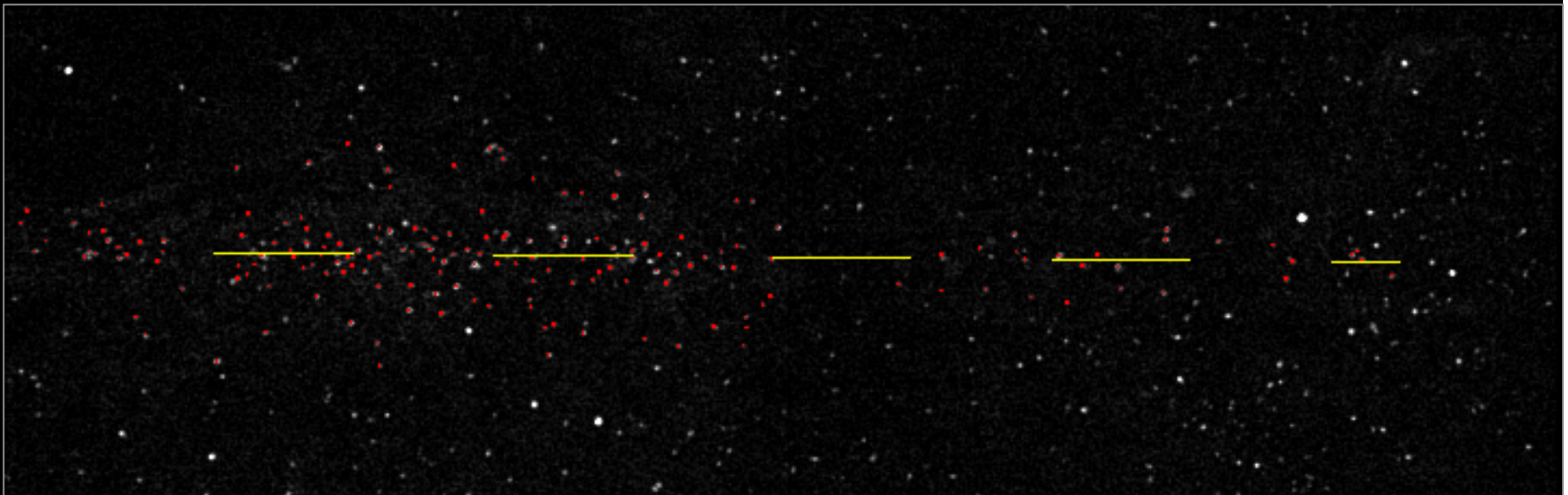
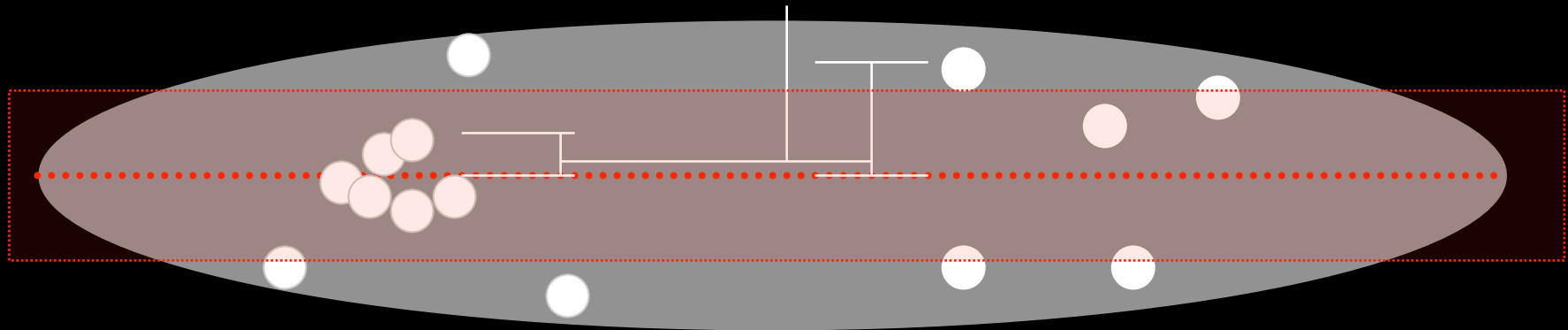


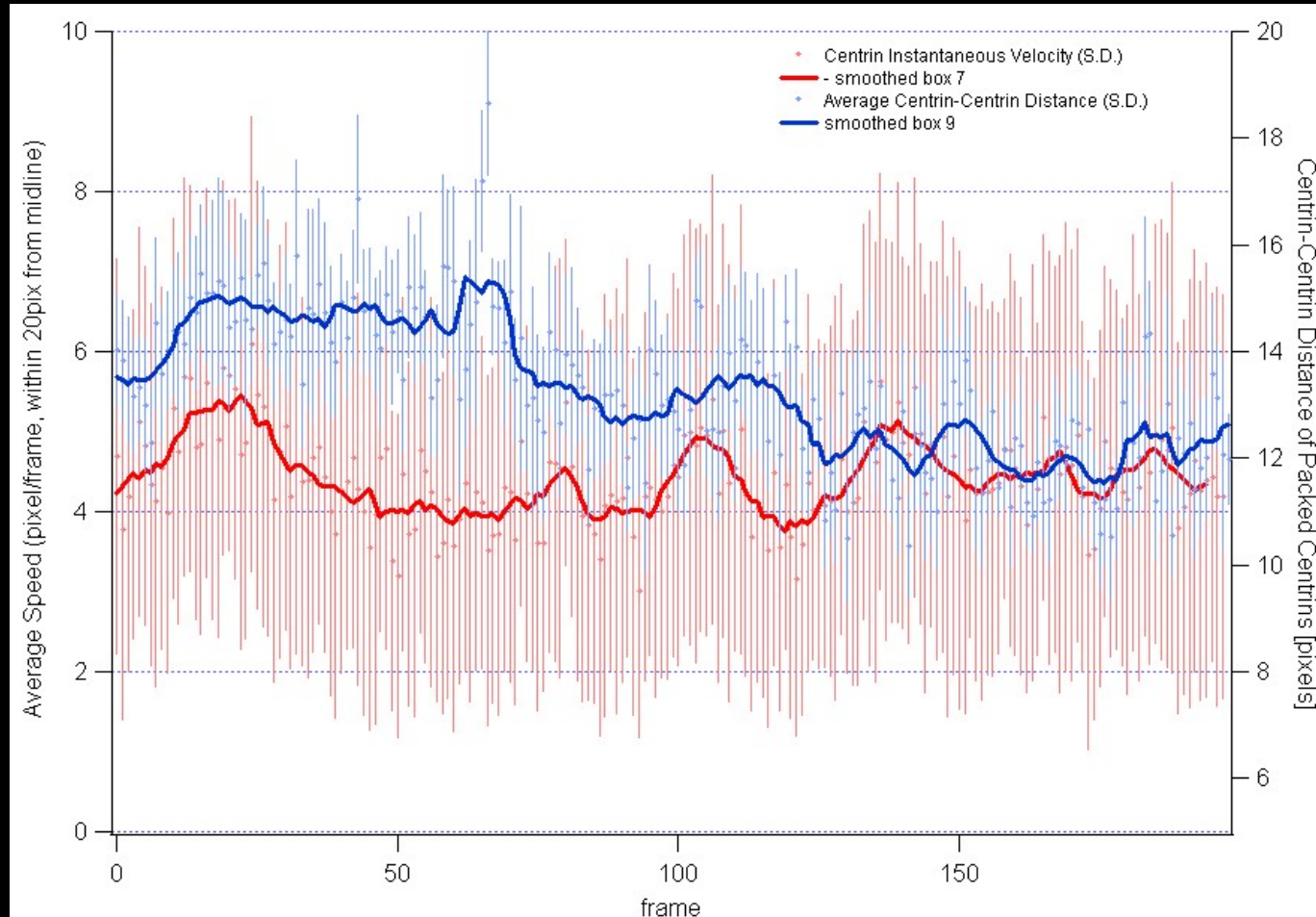
9012 trajectories



Export data
to IgorPro

Distance to the Midline

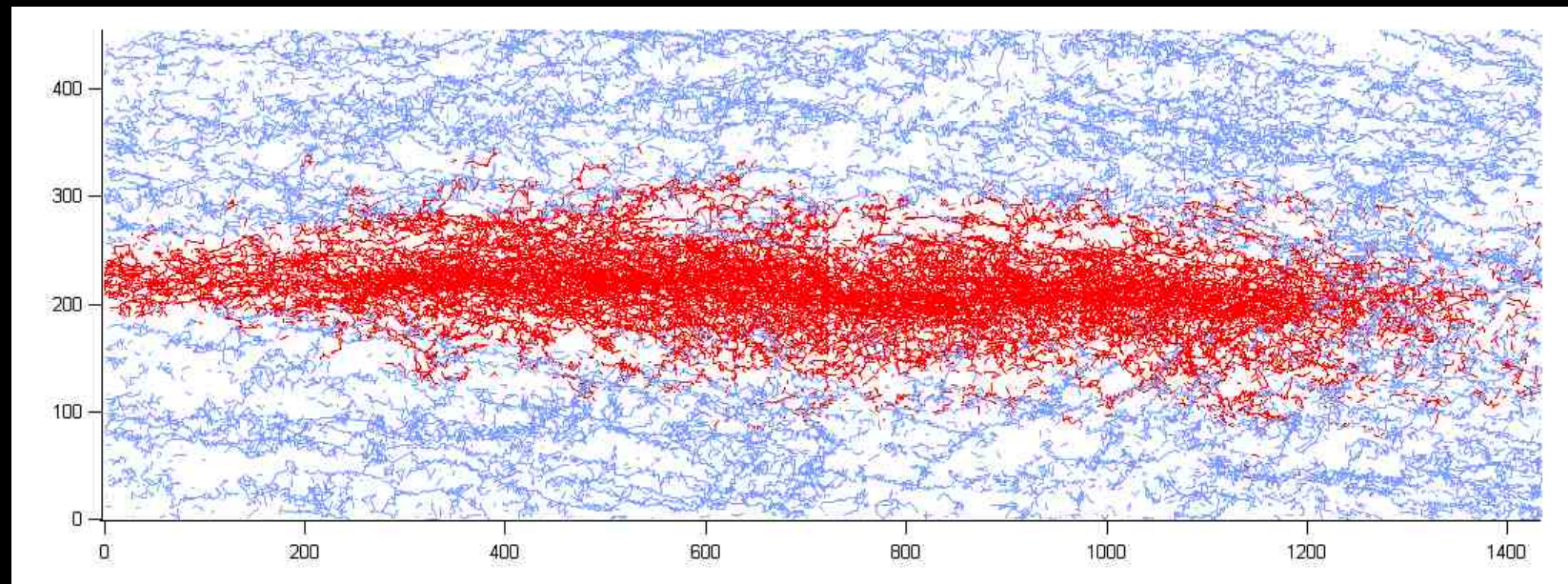
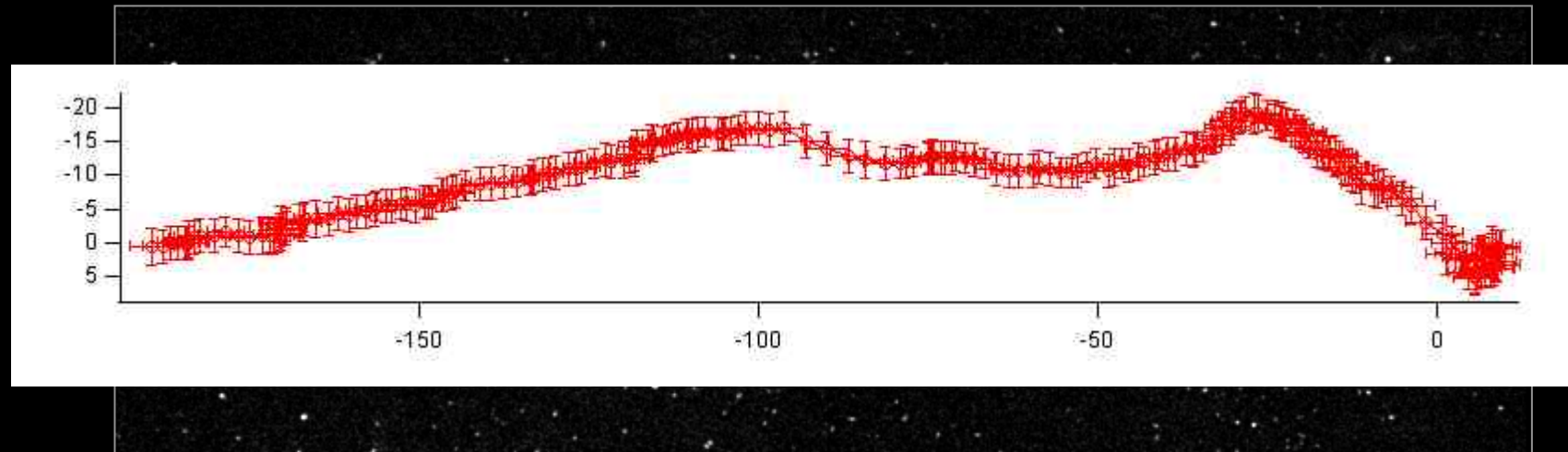




Velocity Decrease \rightarrow (delay) \rightarrow Clustering 1
 \rightarrow cells resume movement \rightarrow Clustering 2

Particle Tracking: Registration – Correction for the background movement

Background movement



Static images

- Intensity Comparison
- Morphometry
- Distribution
- Colocalization

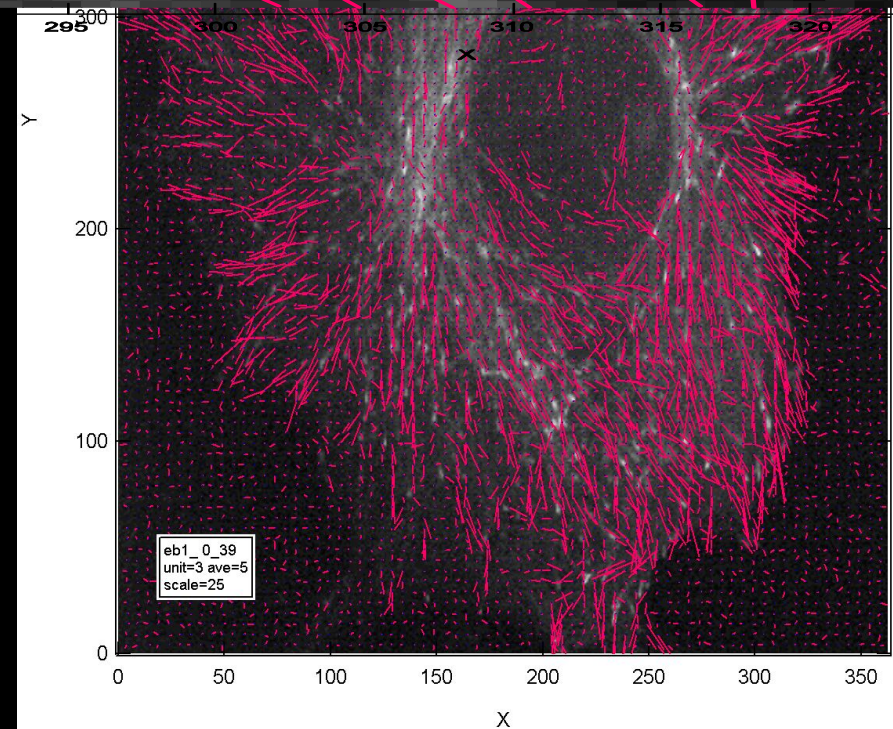
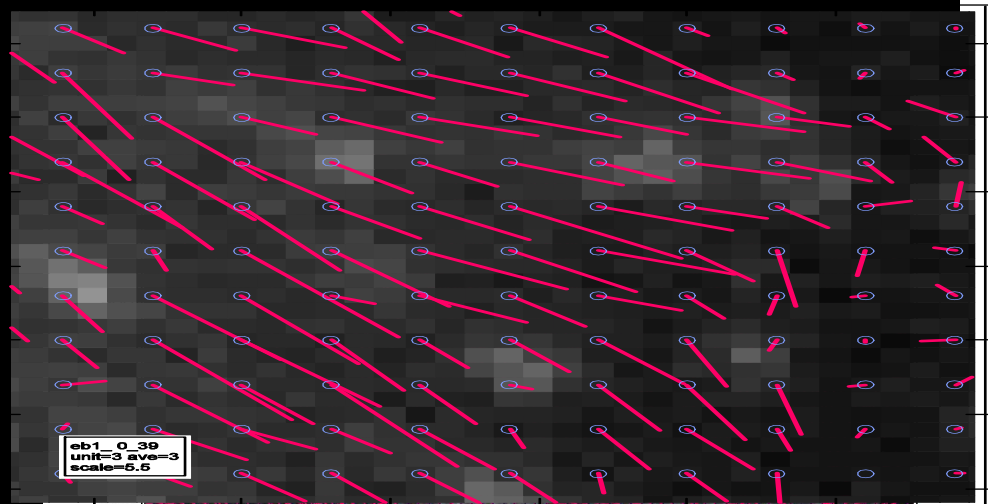
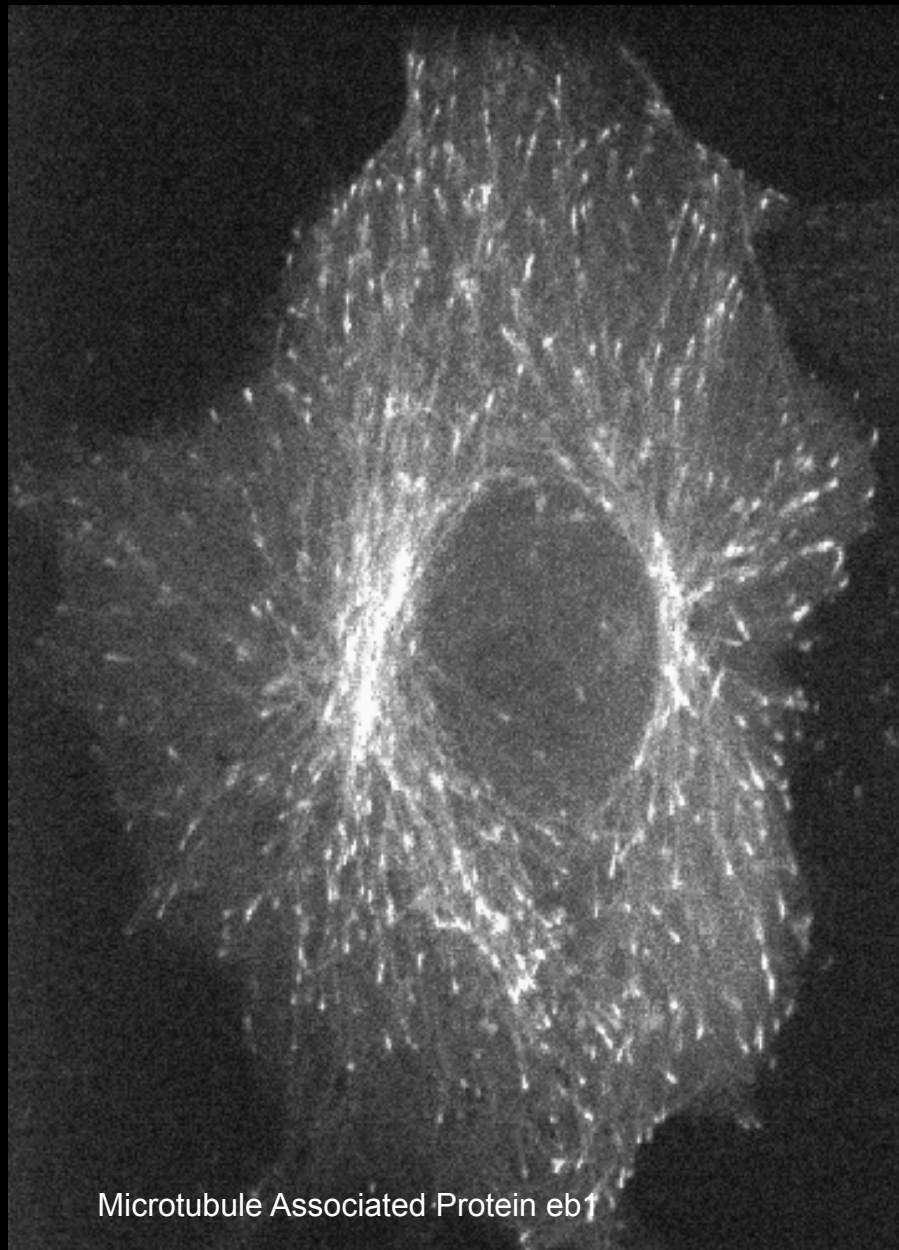
Image sequences

- Intensity dynamics
- Shape dynamics
- Position dynamics :
 - kymograph
 - tracking
- Distribution dynamics
- Optical Flow Estimation →

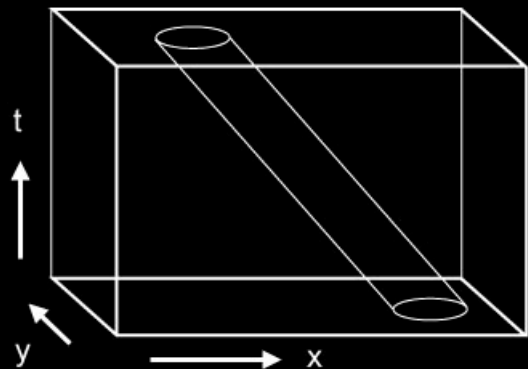
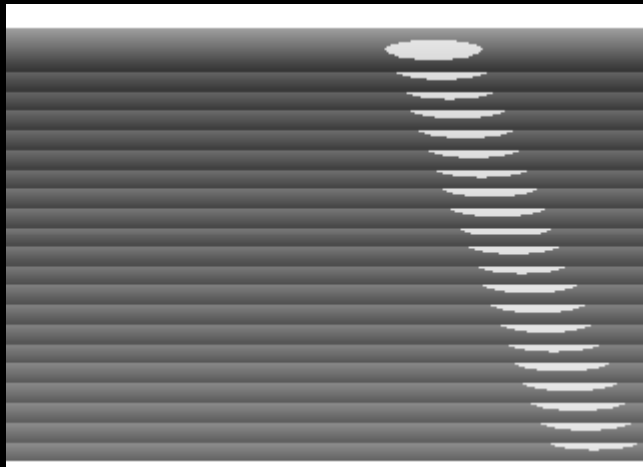
Measurement of Movement

Igor Pro, Wavemetrics

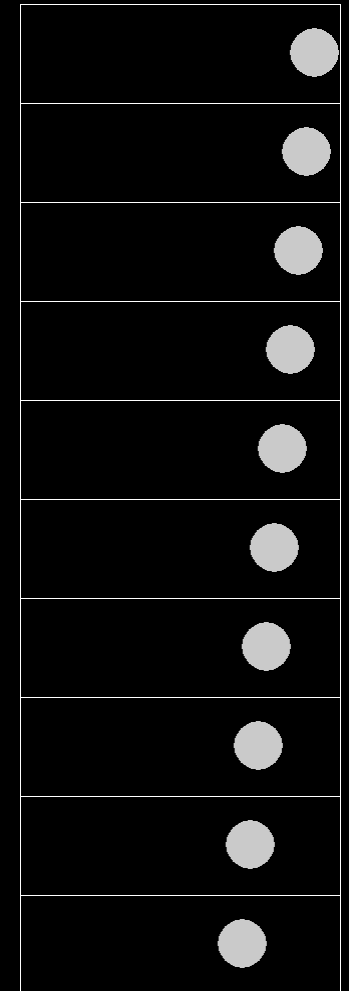
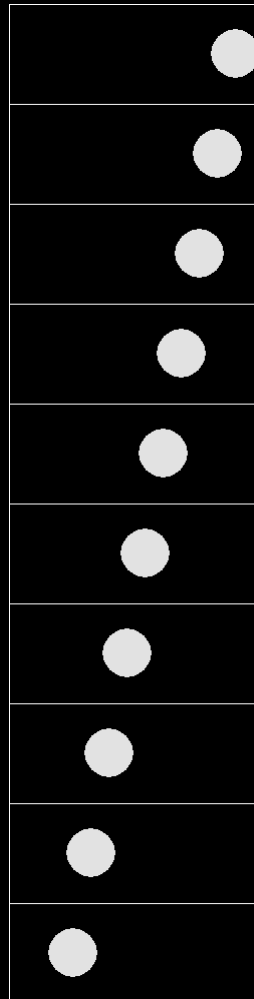
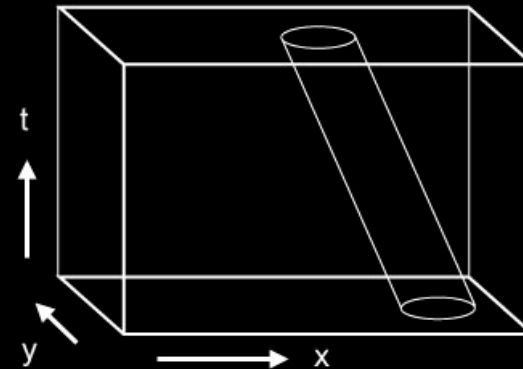
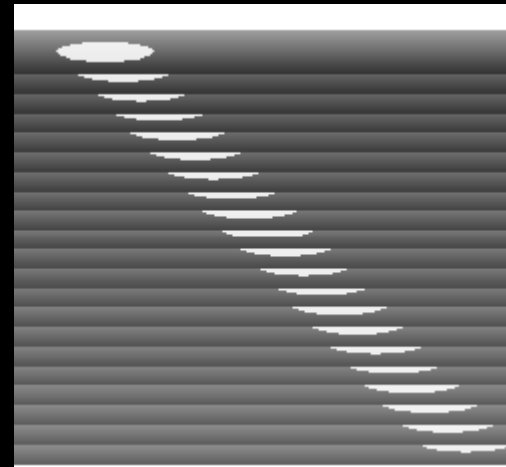
Optical Flow Estimation



Measurement of Movement

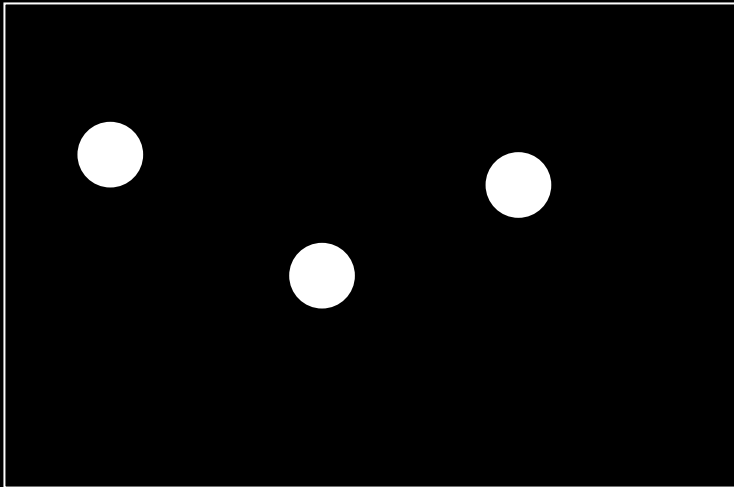


Optical Flow Estimation

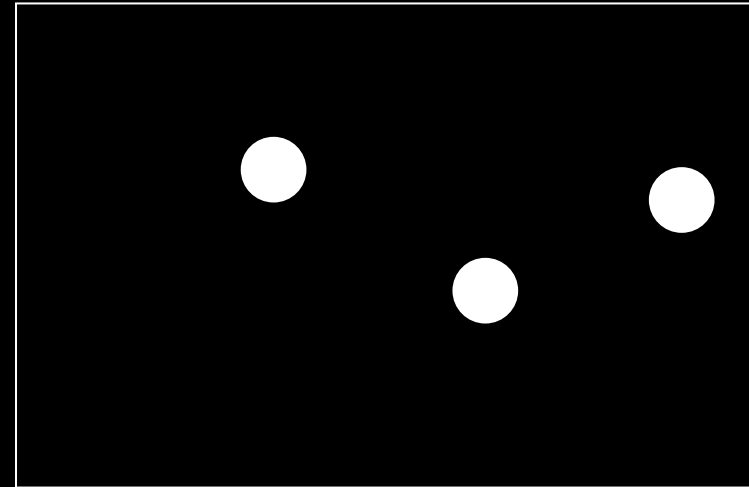


$$\frac{\partial f}{\partial t} + \frac{\partial f}{\partial x} v_x + \frac{\partial f}{\partial y} v_y = 0$$

Optical Flow Equation



t



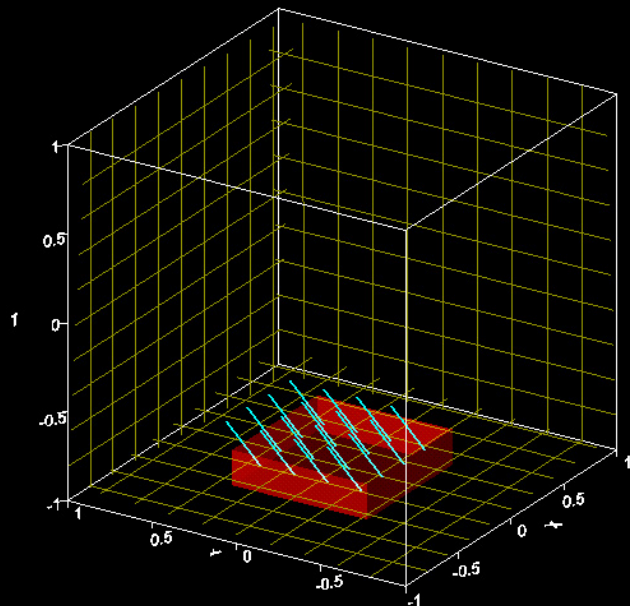
t+1

Under-determined → Need another constraint...

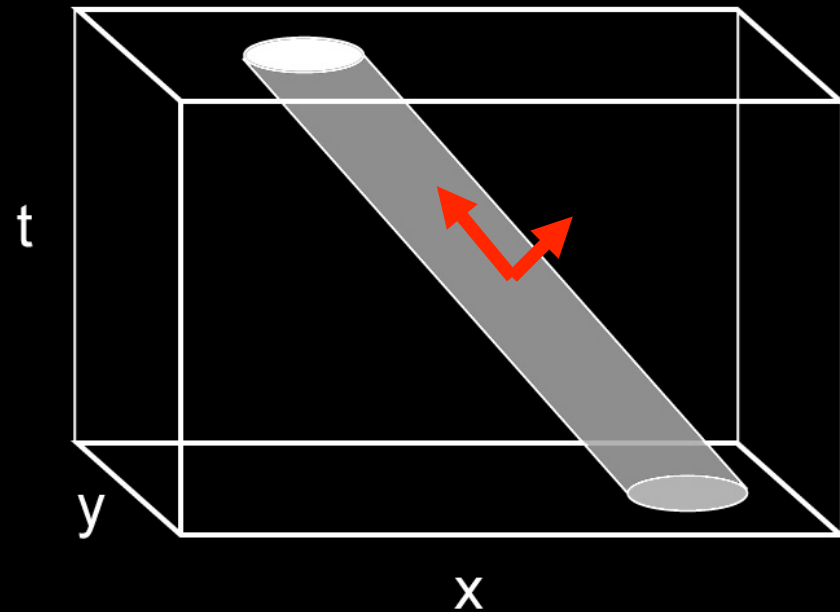
Second Constraint - I

Spatial Local Smoothness

$$\frac{\partial \mathbf{v}}{\partial x} = \frac{\partial \mathbf{v}}{\partial y} = 0$$



Structure Tensor Approach



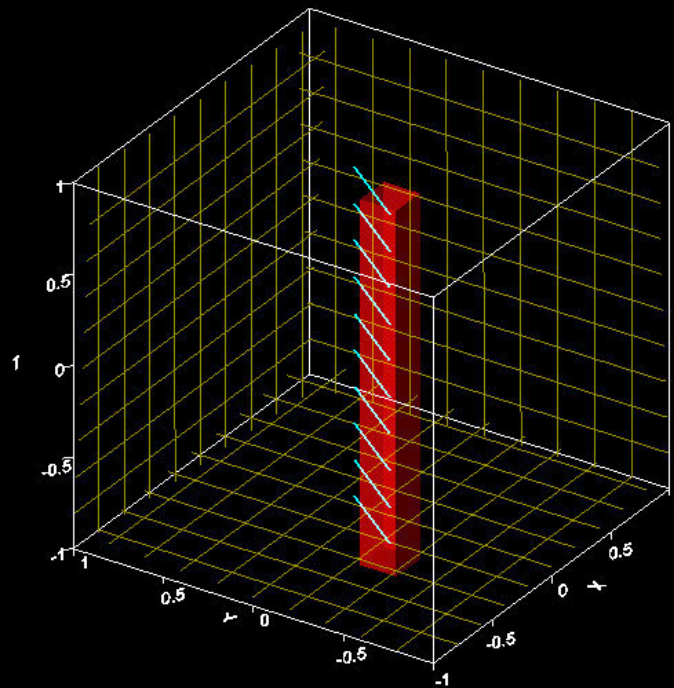
$$J = \begin{bmatrix} \frac{\partial I}{\partial x} \cdot \frac{\partial I}{\partial x} & \frac{\partial I}{\partial x} \cdot \frac{\partial I}{\partial y} & \frac{\partial I}{\partial x} \cdot \frac{\partial I}{\partial t} \\ \frac{\partial I}{\partial x} \cdot \frac{\partial I}{\partial y} & \frac{\partial I}{\partial y} \cdot \frac{\partial I}{\partial y} & \frac{\partial I}{\partial y} \cdot \frac{\partial I}{\partial t} \\ \frac{\partial I}{\partial x} \cdot \frac{\partial I}{\partial t} & \frac{\partial I}{\partial y} \cdot \frac{\partial I}{\partial t} & \frac{\partial I}{\partial t} \cdot \frac{\partial I}{\partial t} \end{bmatrix}$$

→
eigenvector

Second Constraint - II

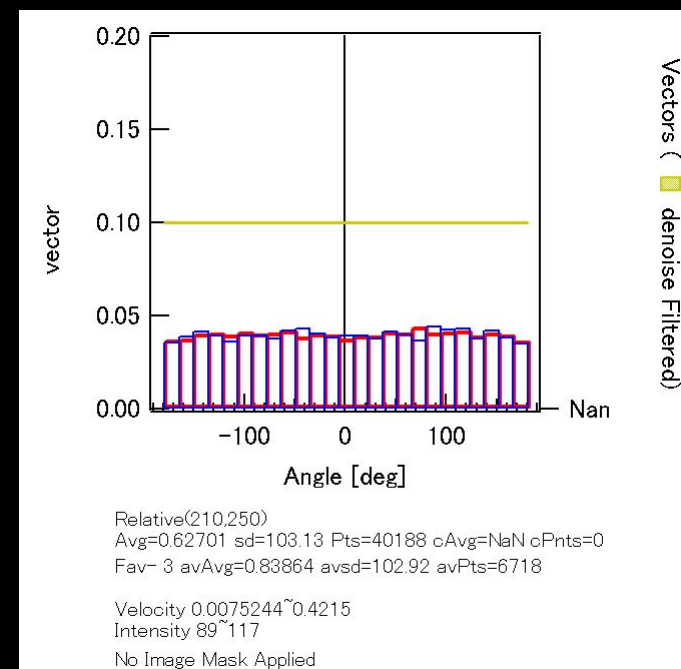
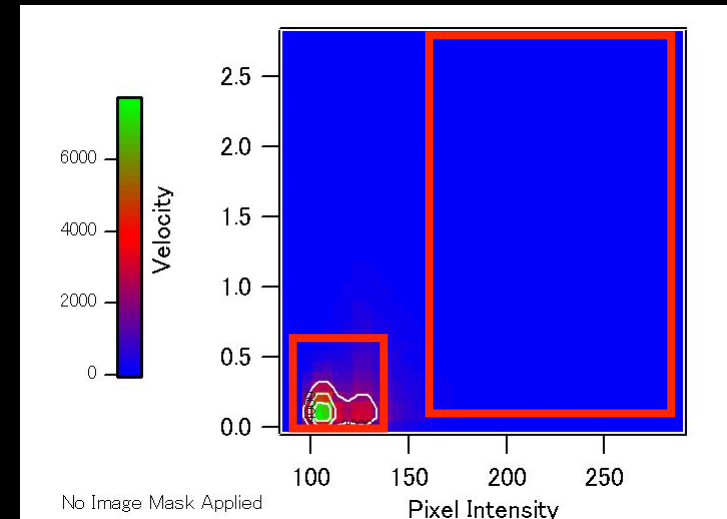
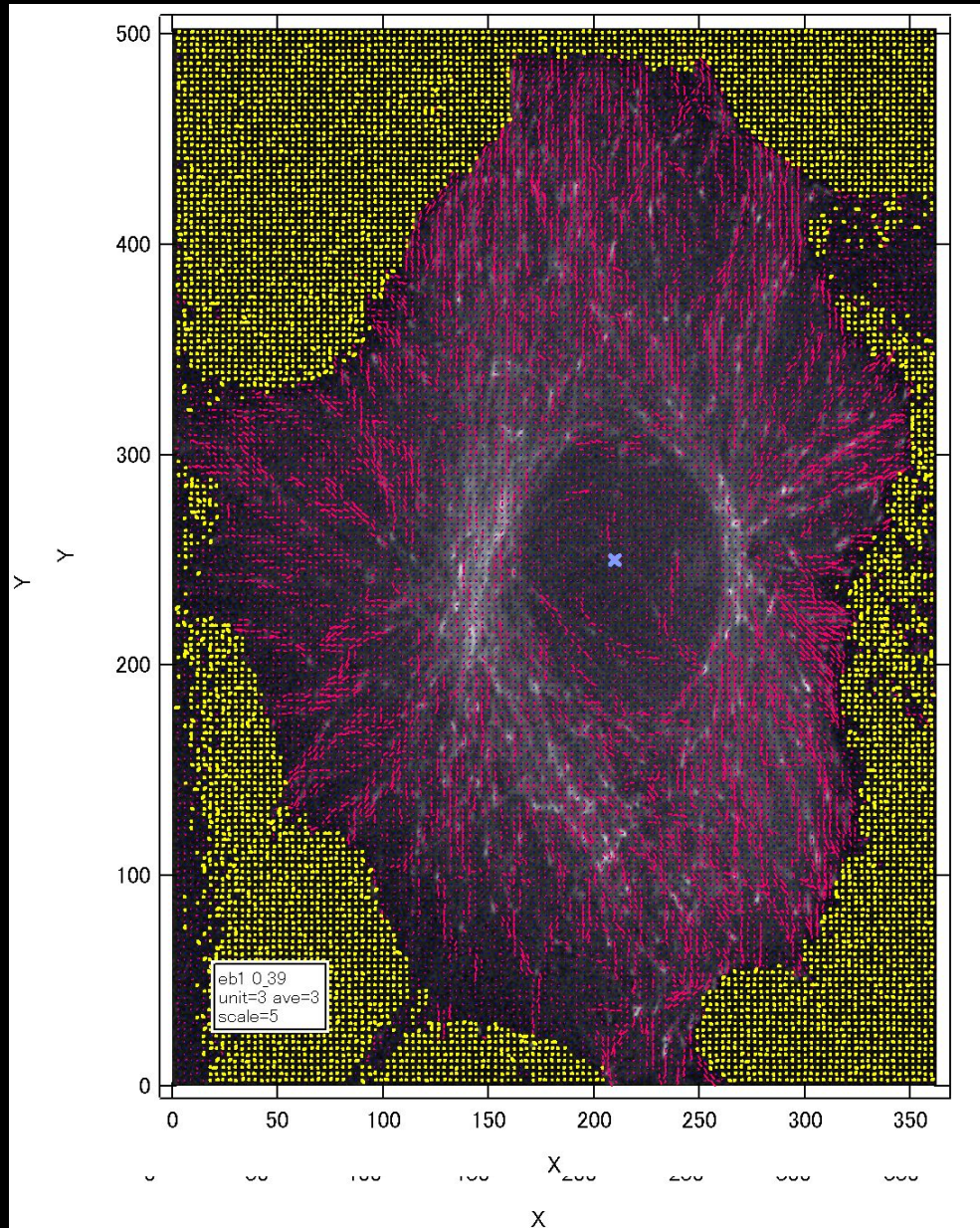
Temporal Local Smoothness

$$\frac{\partial v}{\partial t} = 0$$



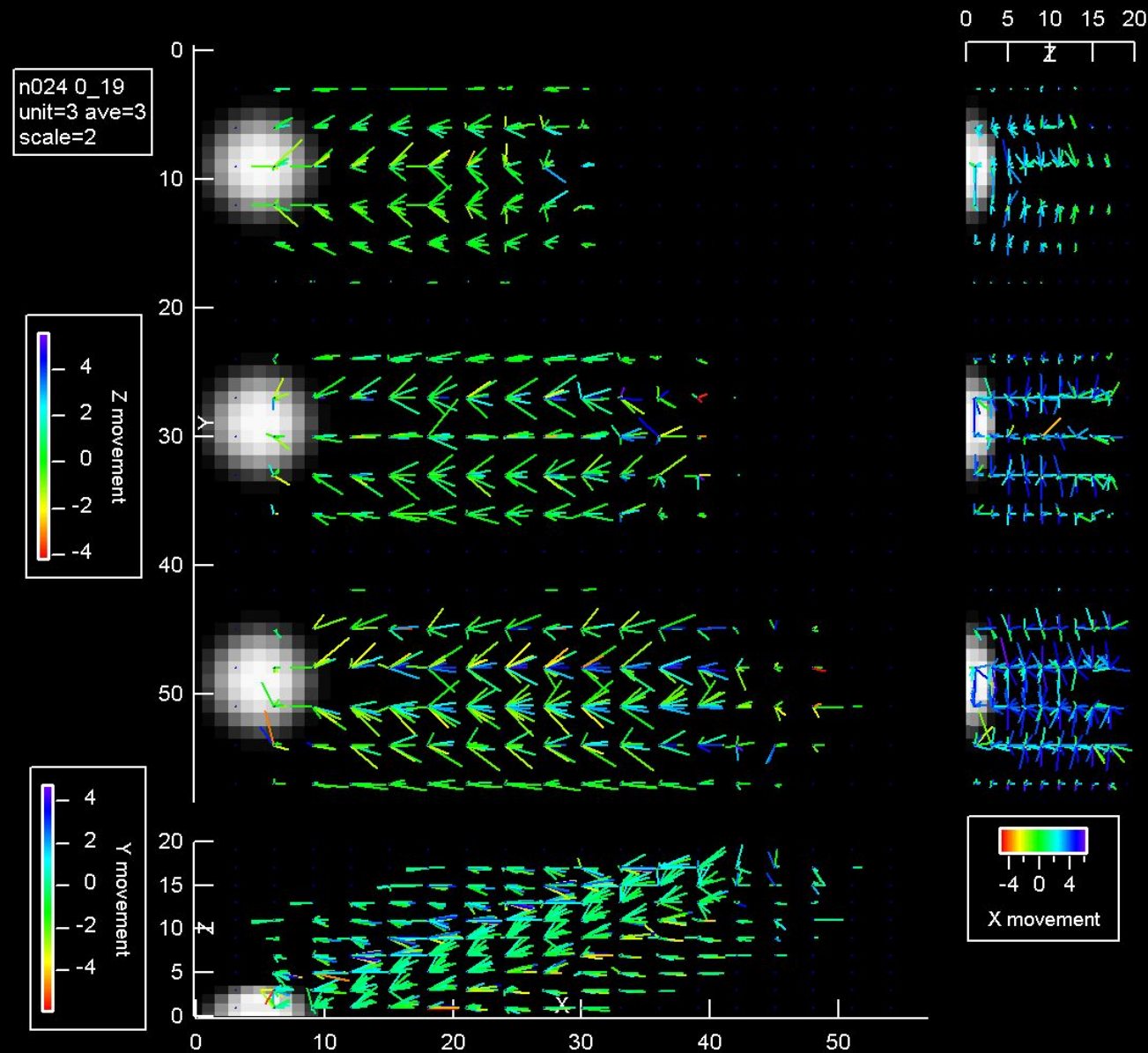
Measurement of Movement

Filtering of Vector Field by Speed & Pixel Intensity



Measurement of Movement: Optical Flow Estimation

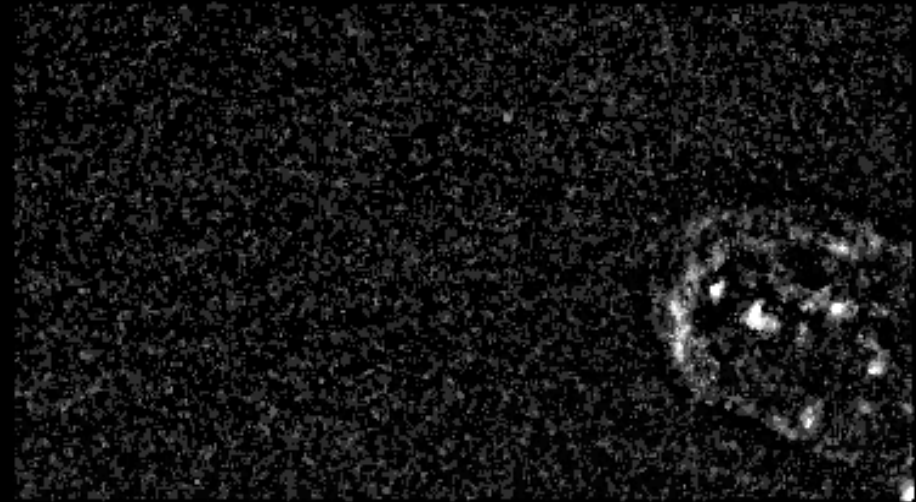
virtual 3D beads



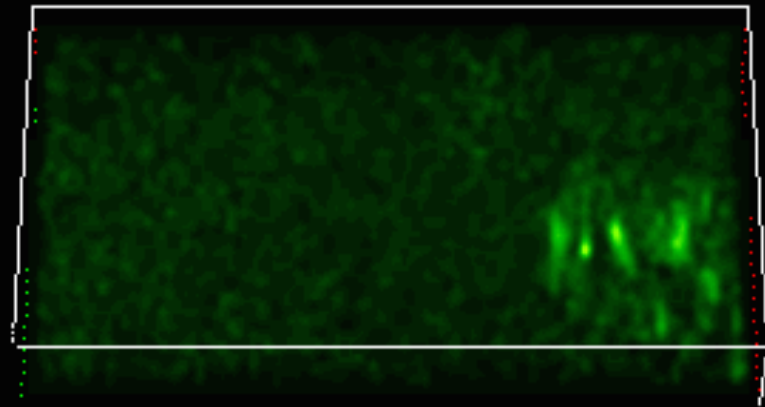
Measurement of Movement: Optical Flow Estimation 3D

3-D *Dictyostelium* Slug Cell Movement

Z-projection



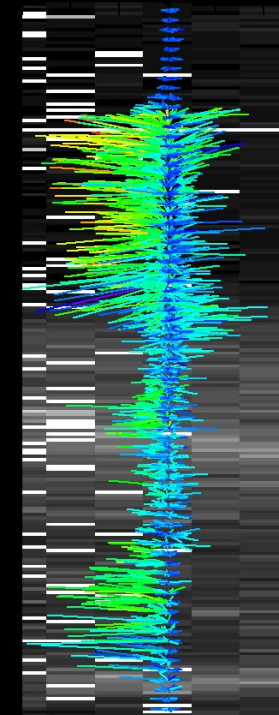
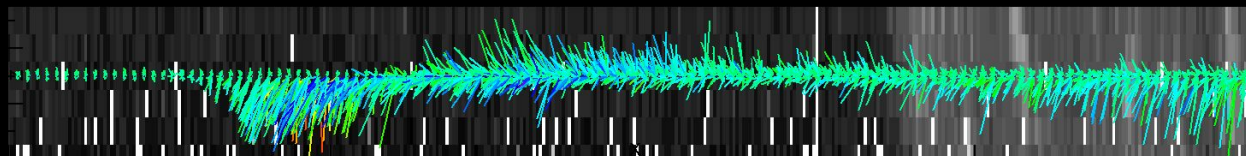
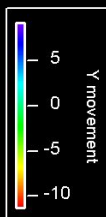
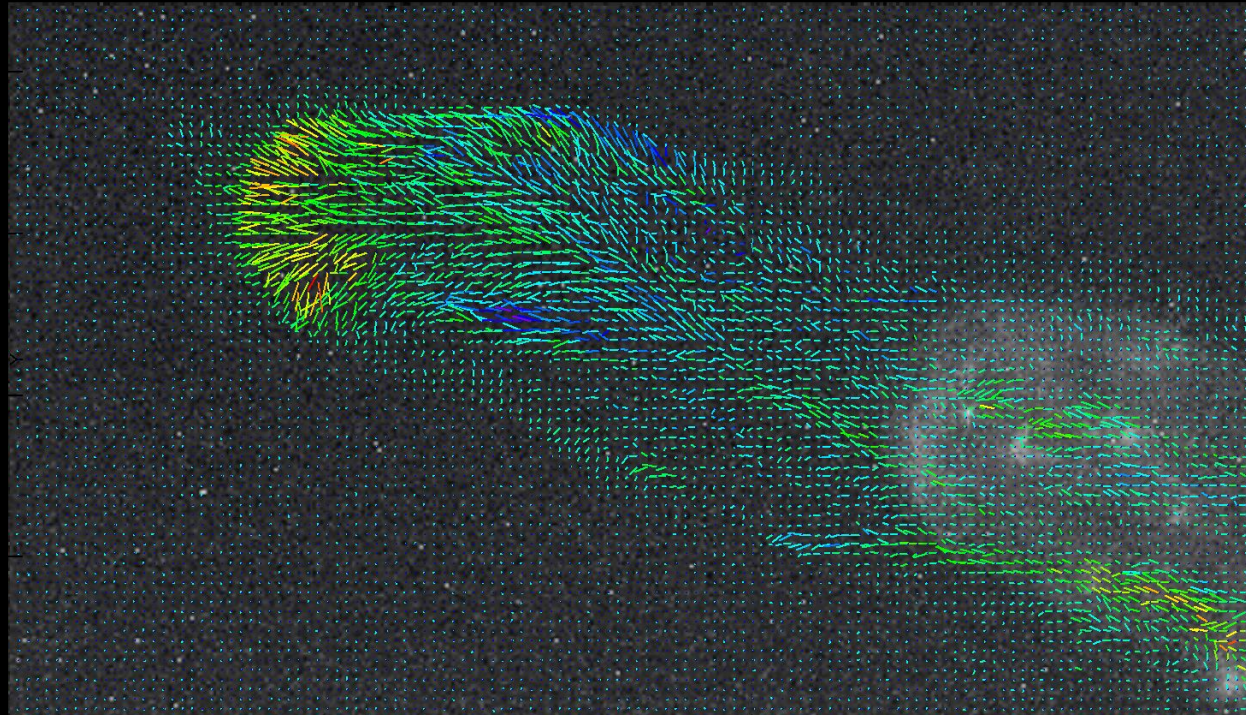
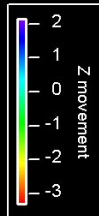
Max.point-projection
3D rendering

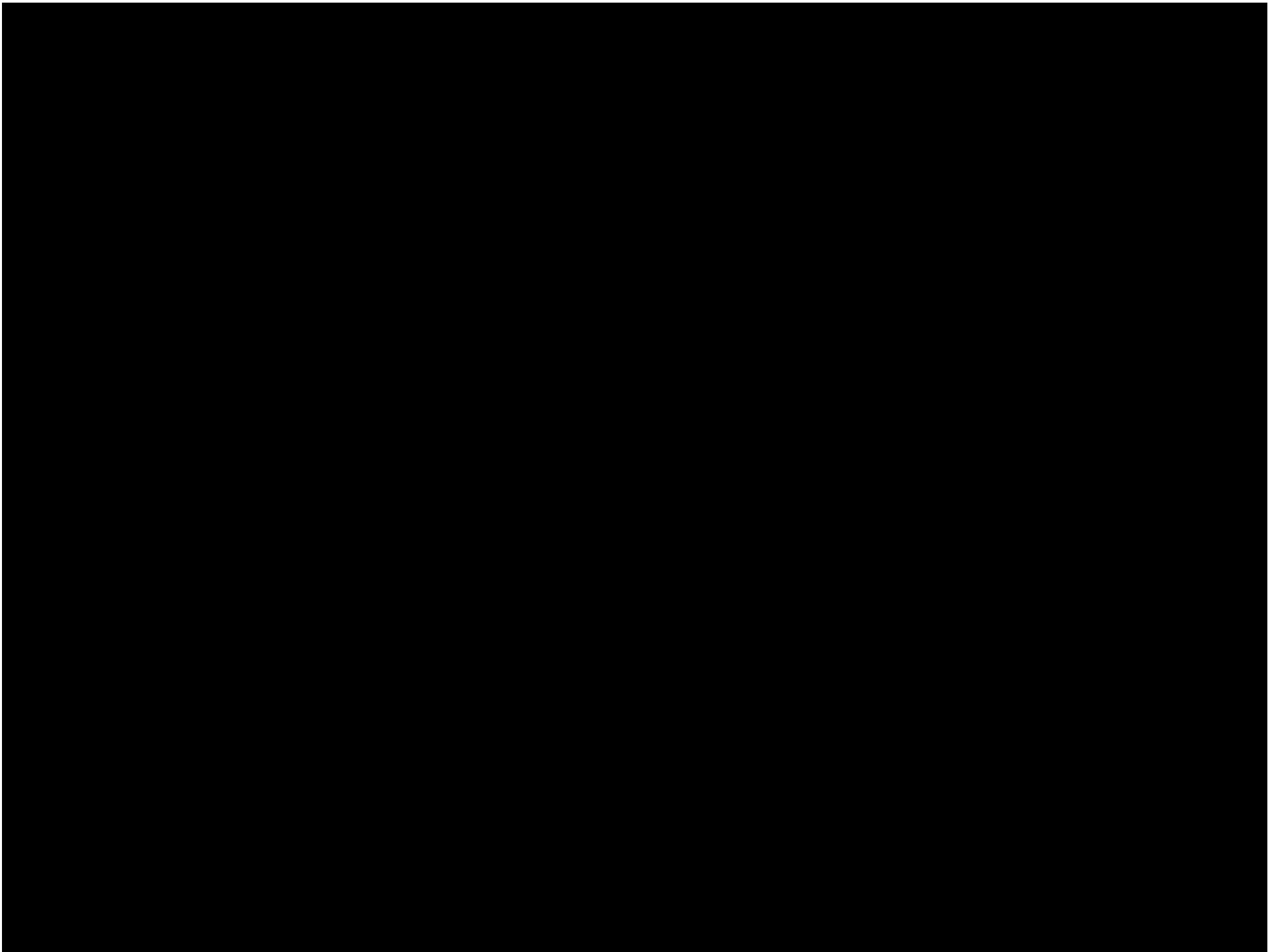


Measurement of Movement: Optical Flow Estimation 3D

3-D *Dictyostelium* Slug Cell Movement

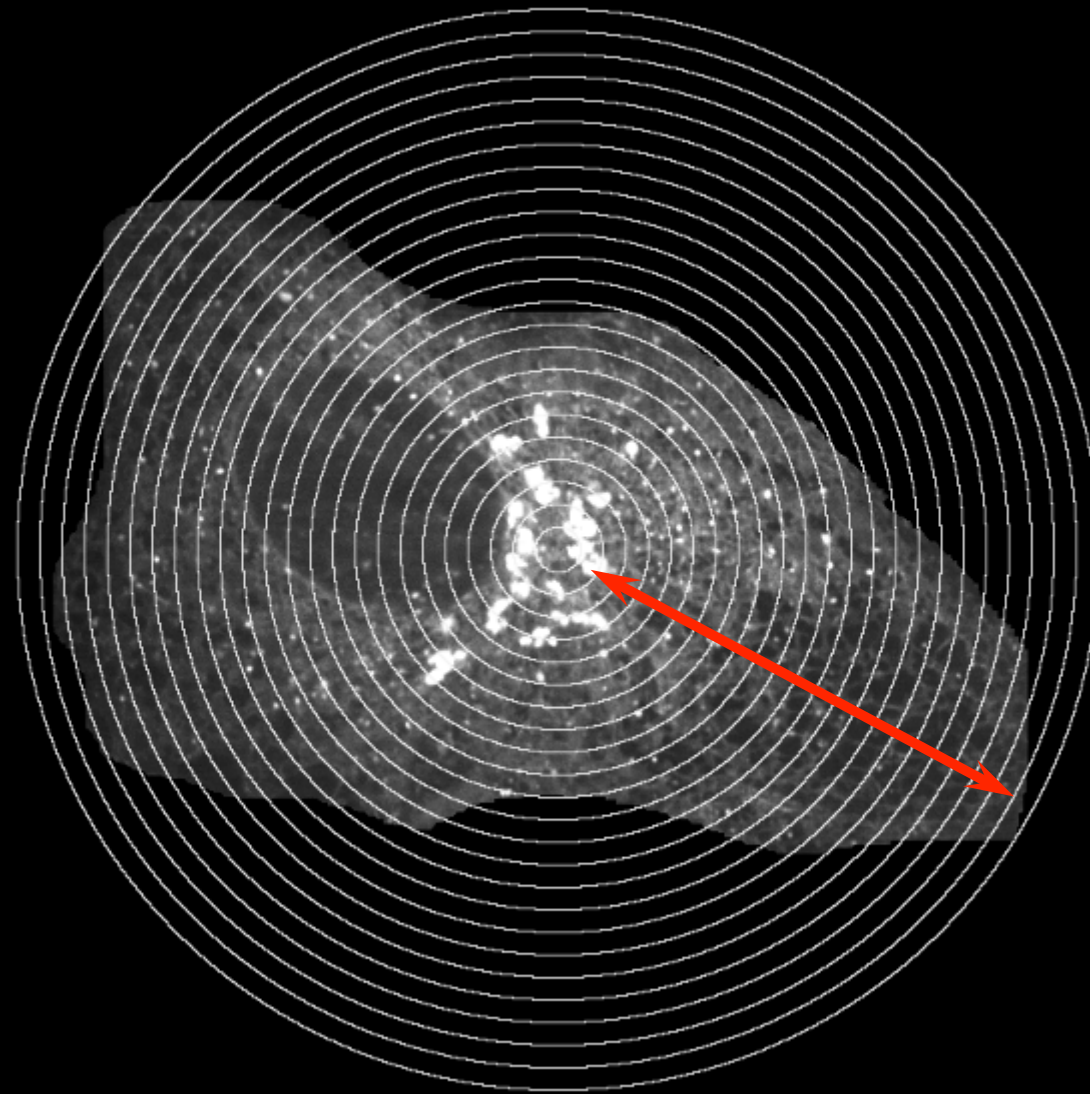
s7a 0_39
unit=3 ave=3
scale=5



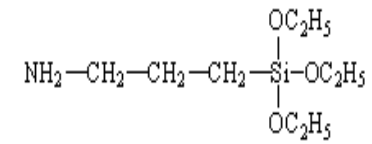


Vesicle Transport

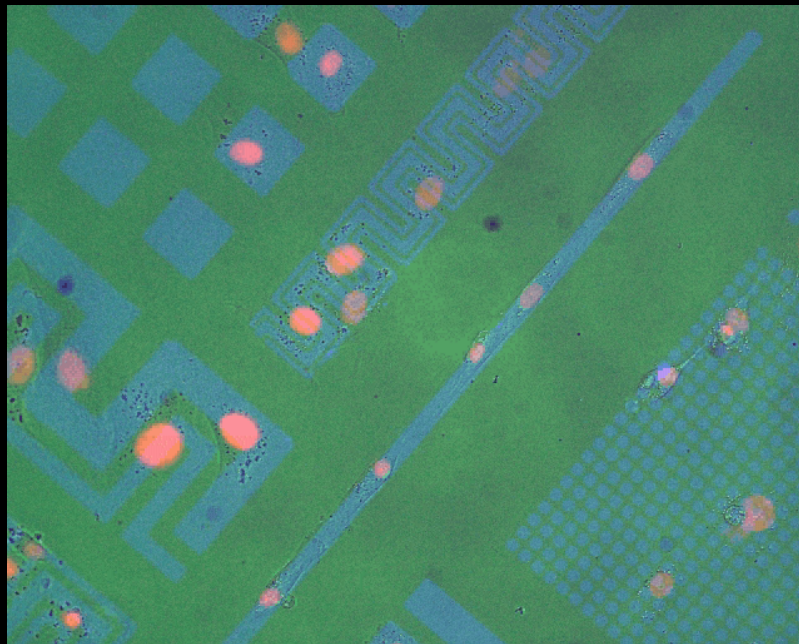
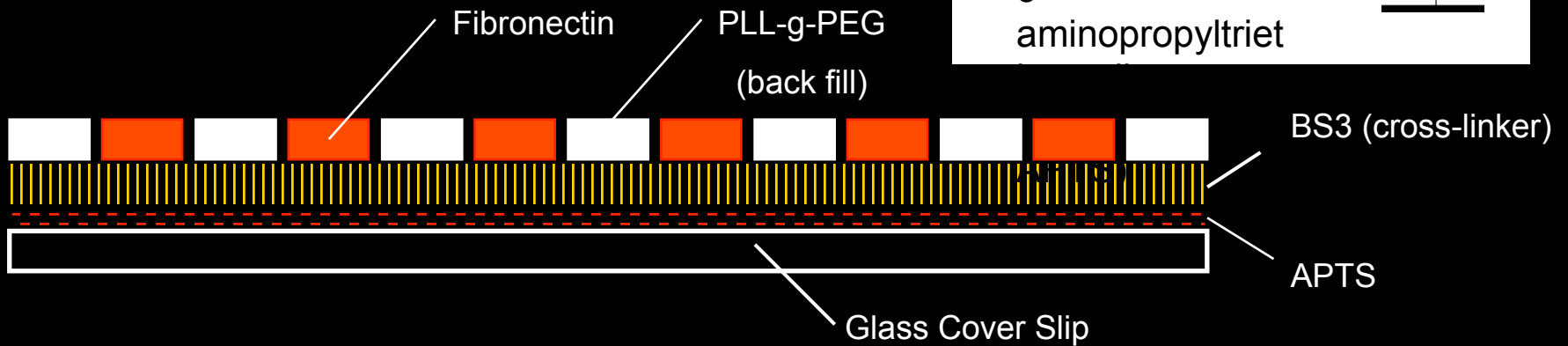
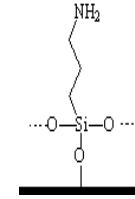
- Simulation: Initial Protein Distribution -



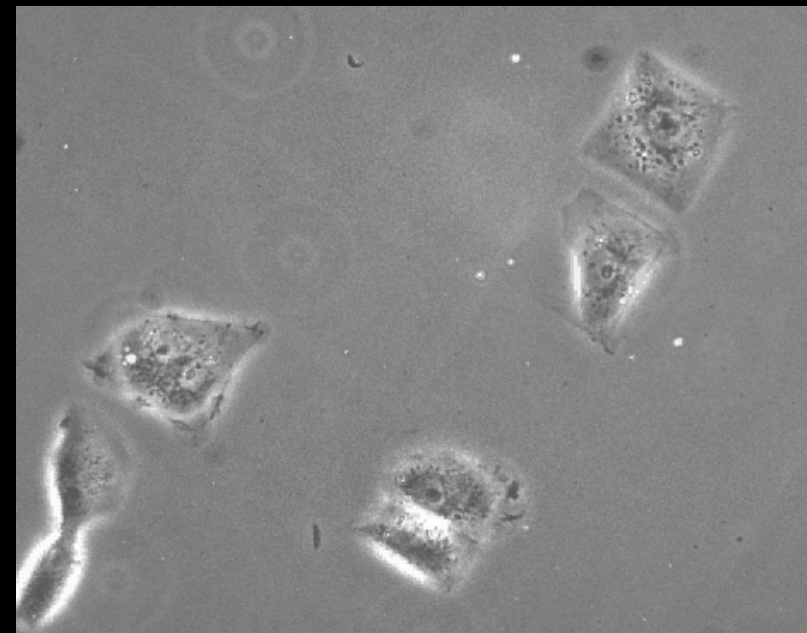
Patterned Substrate
→ constant boundary condition



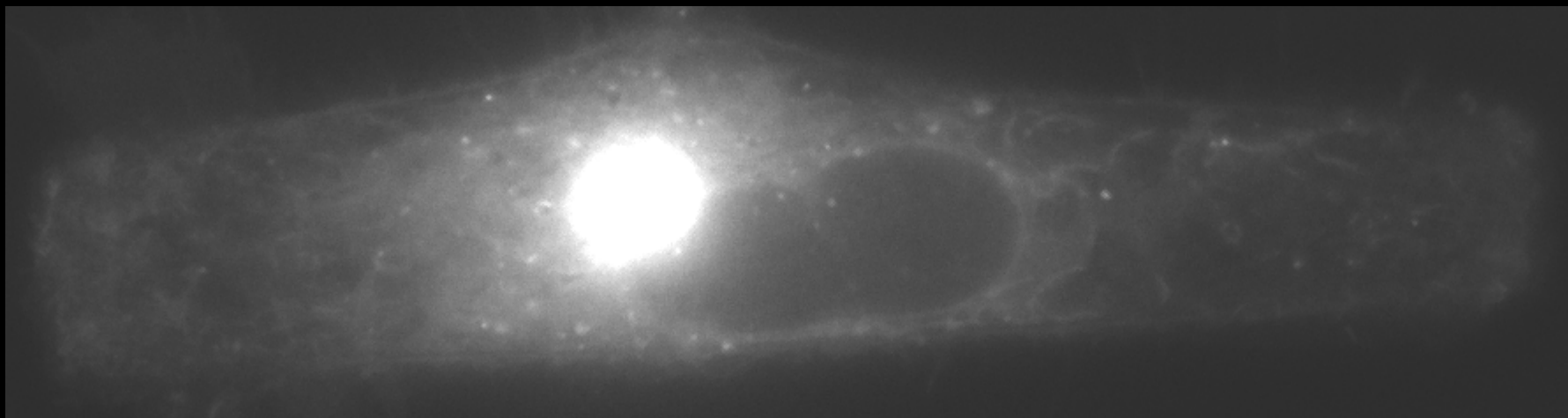
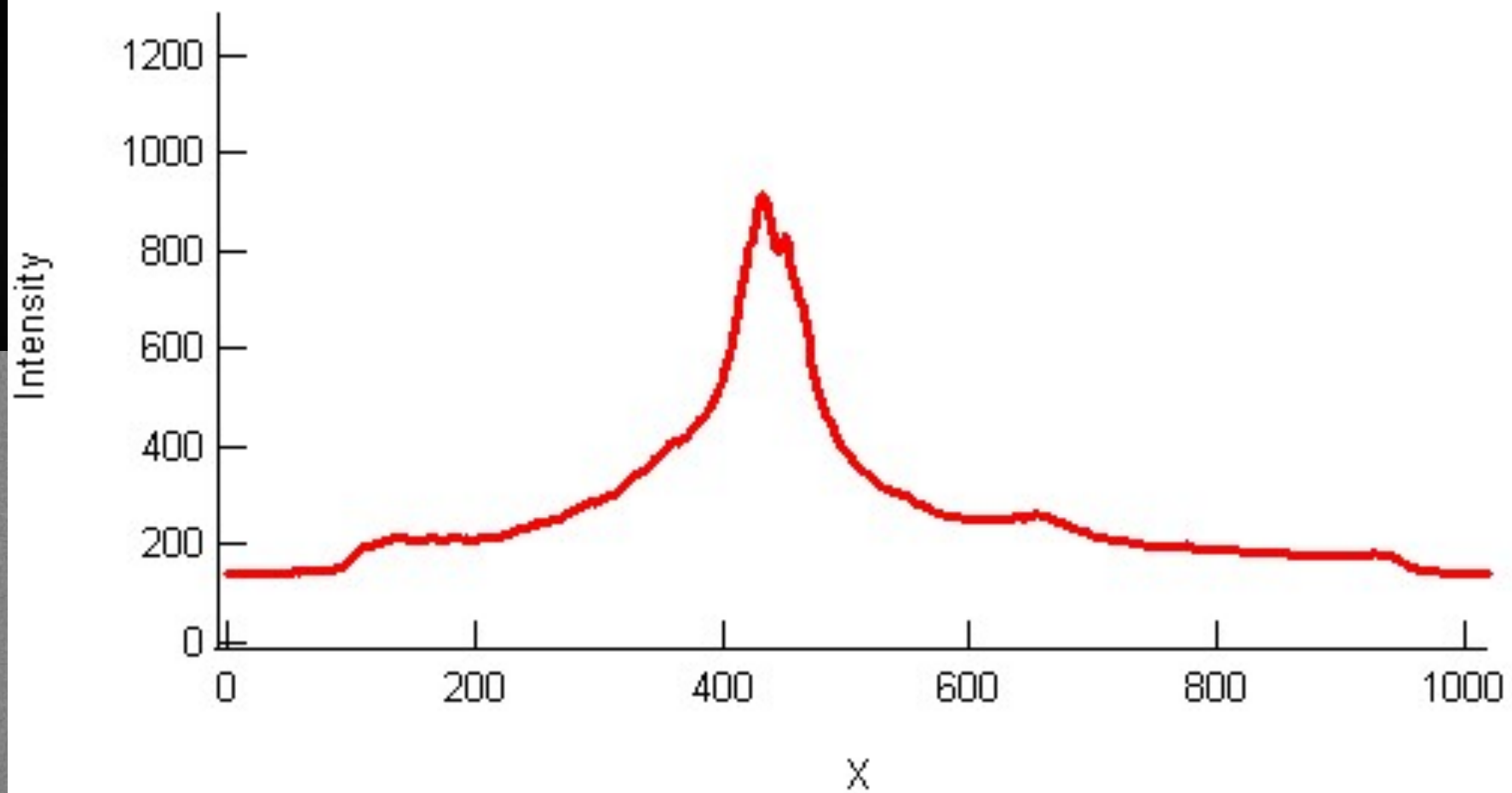
3-aminopropyltriethoxysilane



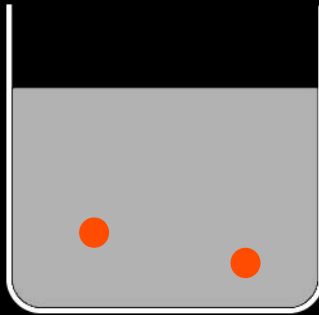
Line width: 10 μm



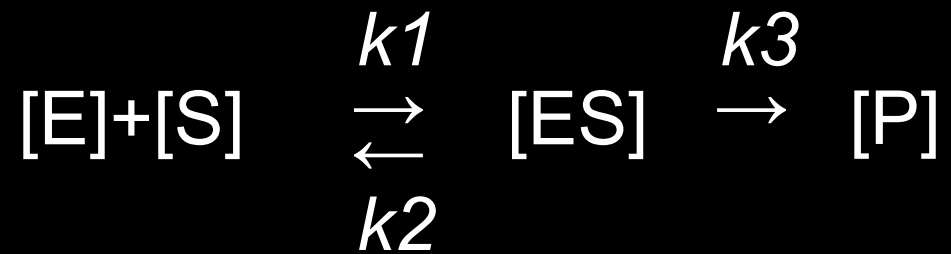
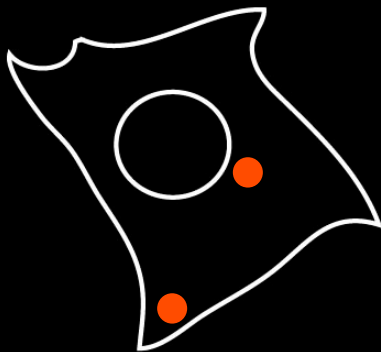
Squares: 50 x 50 μm



Biochemistry
(solution)



Topo-biochemistry
(in vivo)

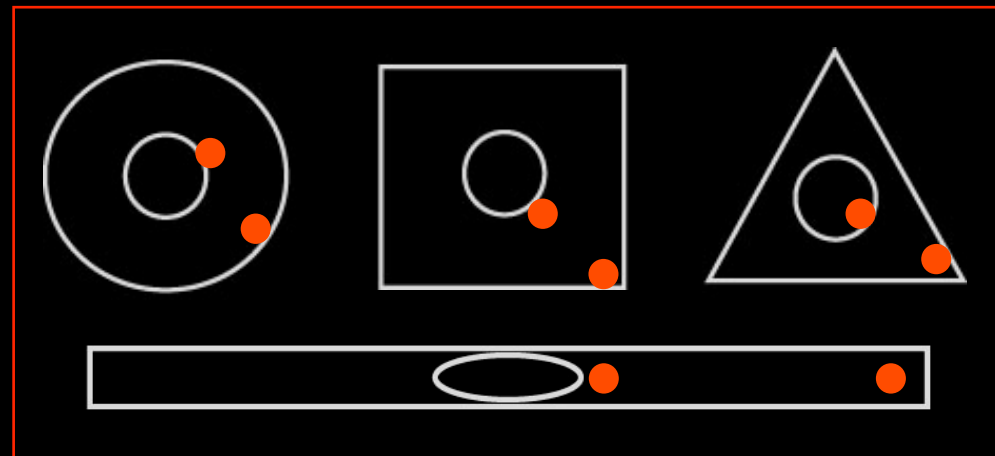


$$B([E], [S], [ES], [P], t)$$

$$B([E], [S], [ES], [P], t, x)$$

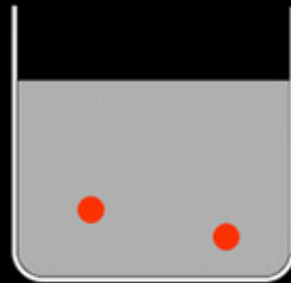
$$B([E], [S], [ES], [P], t, x, y)$$

$$B([E], [S], [ES], [P], t, x, y, z)$$



Topobiochemistry

A.



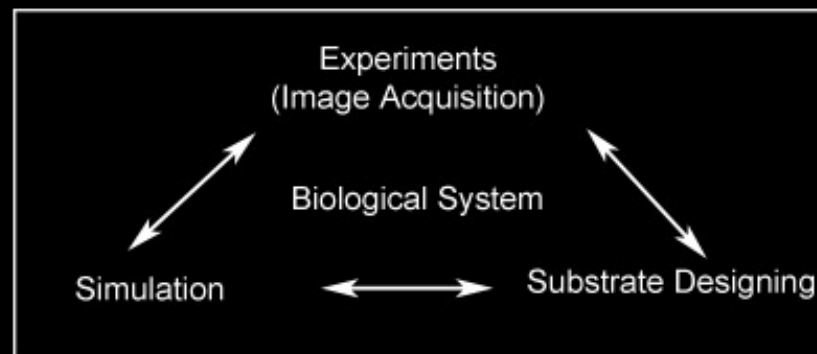
B.



C.



D.



Endoplasmic Reticulum
The Golgi apparatus

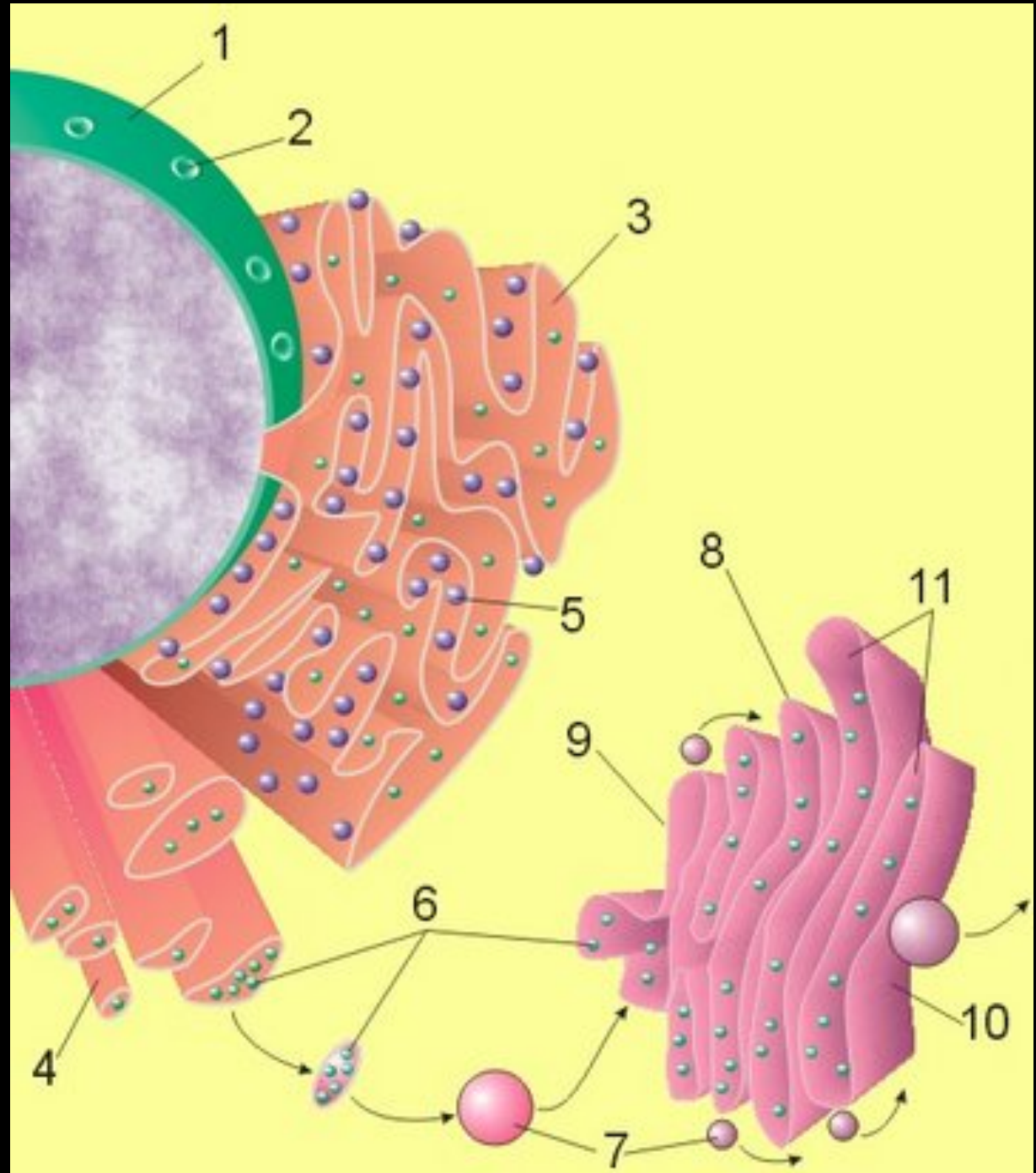
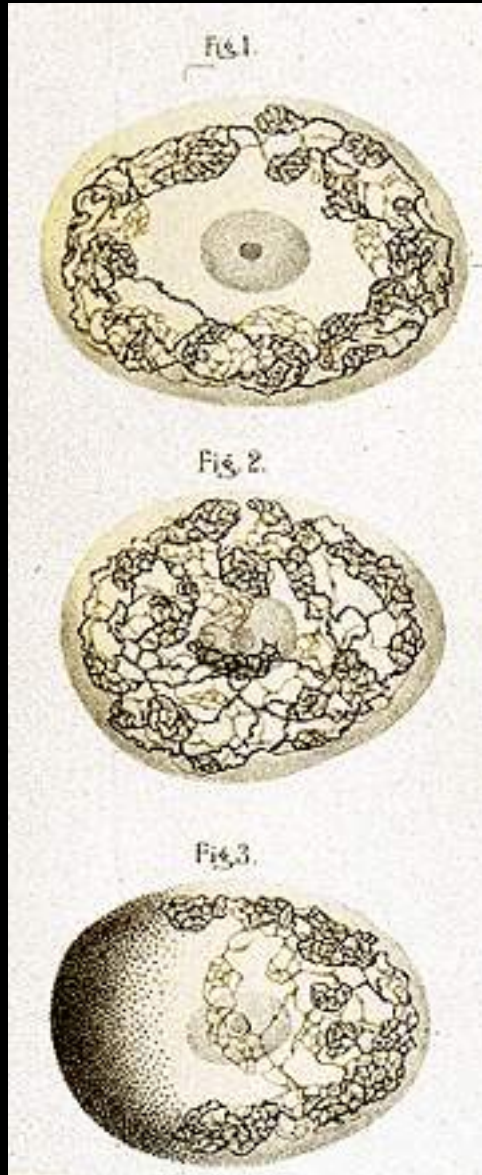


Image taken from Wikipedia

The Golgi Apparatus

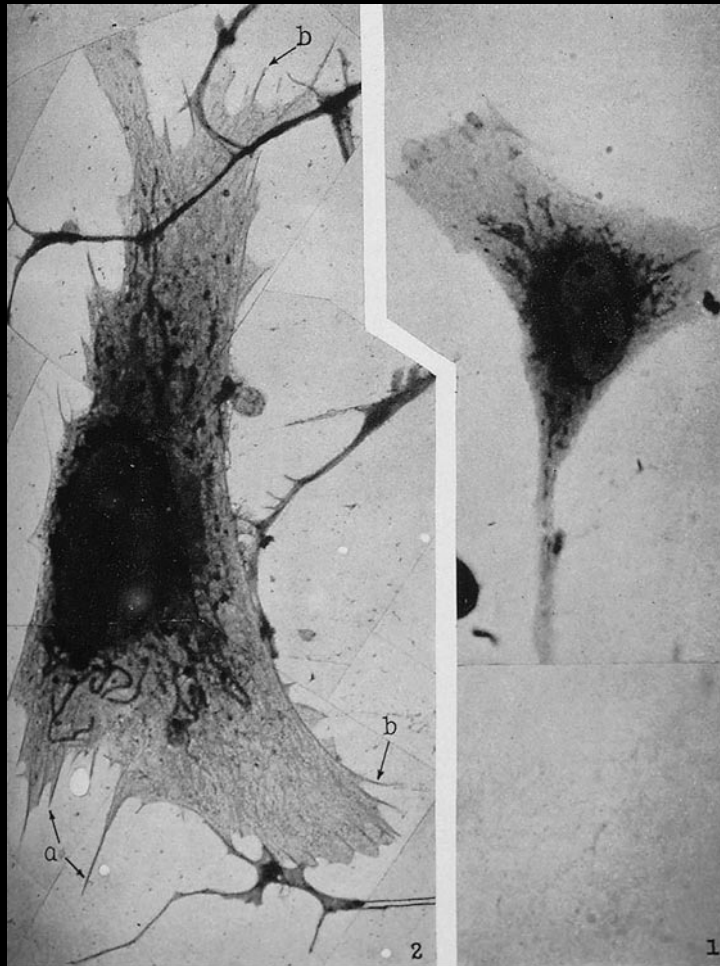


In 1897, studying the nervous system with his black reaction, Golgi noticed in neurons an intracellular structure, whose existence he officially reported in April 1898. This structure was designated by Golgi "internal reticular apparatus" and was soon named after him as Golgi apparatus (or much later as the Golgi complex and is frequently referred to nowadays only as "the Golgi").

Golgi's drawings of the "internal reticular apparatus" that he observed in spinal ganglia (the different drawings illustrate the variety of features Golgi observed with his metal impregnation, from *Opera Omnia*). This intracellular structure is universally known nowadays as "Golgi apparatus".

<http://nobelprize.org/medicine/articles/golgi/>

Endoplasmic Reticulum



This famous first electron micrograph of an intact cell was published in *The Journal of Experimental Medicine* in March 1945, in "A Study of Tissue Culture Cells by Electron Microscopy," by Keith R. Porter, Albert Claude, and Ernest F. Fullam. The cell is a cultured fibroblast originating from a chick embryo, which was grown by Porter on polyvinyl film, then peeled off and transferred to a wire specimen grid. The cell was fixed with osmium tetroxide, washed and then dried in order to prevent evaporation in the electron microscope's vacuum chamber. Magnified 1600 times, this first electron micrograph of a cell reveals mitochondria, the Golgi apparatus and a "lace-like reticulum" which Porter later named the "endoplasmic reticulum". The electron microscope used for this historic image was an RCA EMB model, operated by Fullam at the Interchemical Corporation in New York City.

<http://www.rockefeller.edu/rucal/journey/journey.html>

http://www.nature.com/ncb/journal/v1/n4/fig_tab/ncb0899_E99_F1.html

Intracellular Organelle → Protein Dynamics
(Static View)



Protein Dynamics → Intracellular Organelle
(Dynamic View)

