

# Lecture 15

# Localization Microscopy

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# localization microscopy – one technique, many acronyms

sptPALM      GSDIM      dSTORM      d<sup>4</sup>STORM

PALM      STORM      rapidSTORM

P-FPALM

FPALM      SPDM

PALMIRA      RPM

SALM      SOFI      DAOSTORM      CHIRON

LOBSTER      FIONA

PRILM      3B      uPAINT

# localization microscopy

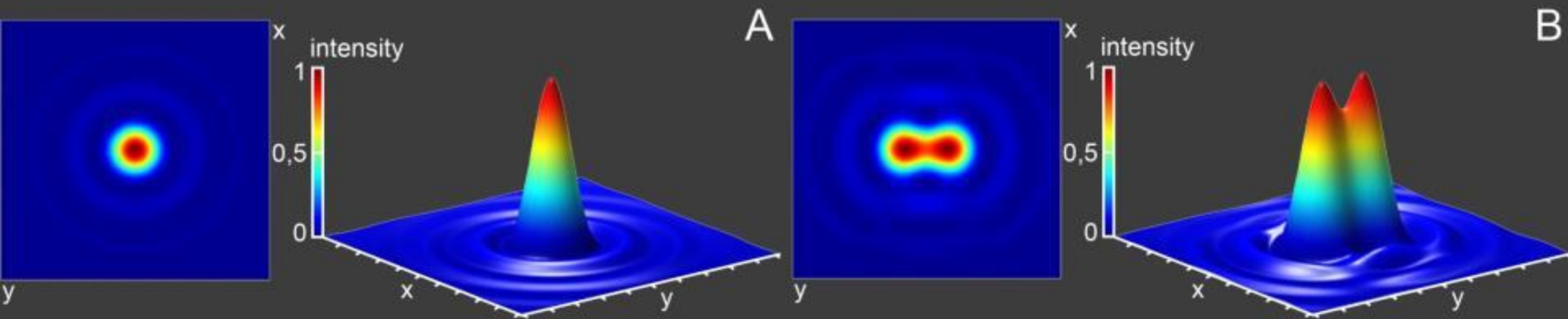
## Outline:

- introduction and general idea of localization microscopy
- first approaches: “original” (F)PALM and STORM
- dSTORM, SPDM, GSDIM – using standard fluorophores
- 3D
- live-cell (4D)
- statistical analysis using the additional single molecule information
- alternative approaches
- conclusion

# introduction to localization microscopy

problem in light microscopy: resolution limited by diffraction

$$\rightarrow I_{im}(x_2, y_2) = PSF \otimes P_{fl}(x_1, y_1)$$



Rayleigh criterion:  $D = 0.61 \frac{\lambda}{NA}$

# introduction to localization microscopy

general idea:

look at signals of single molecules individually instead of all fluorophores at the same time

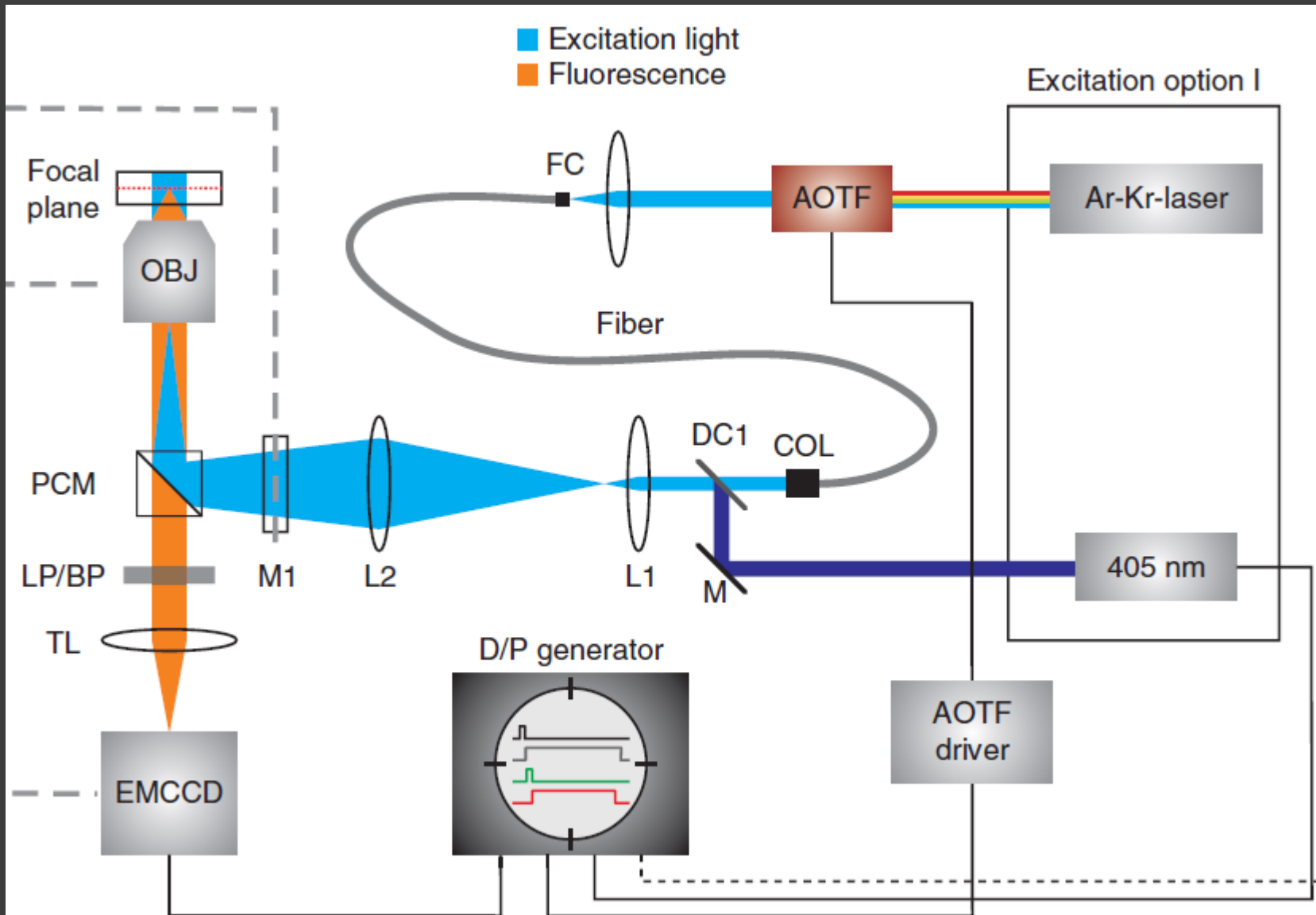
→ this allows a very precise determination of the molecule position

→ reconstruct super-resolution image from position data of the detected molecules



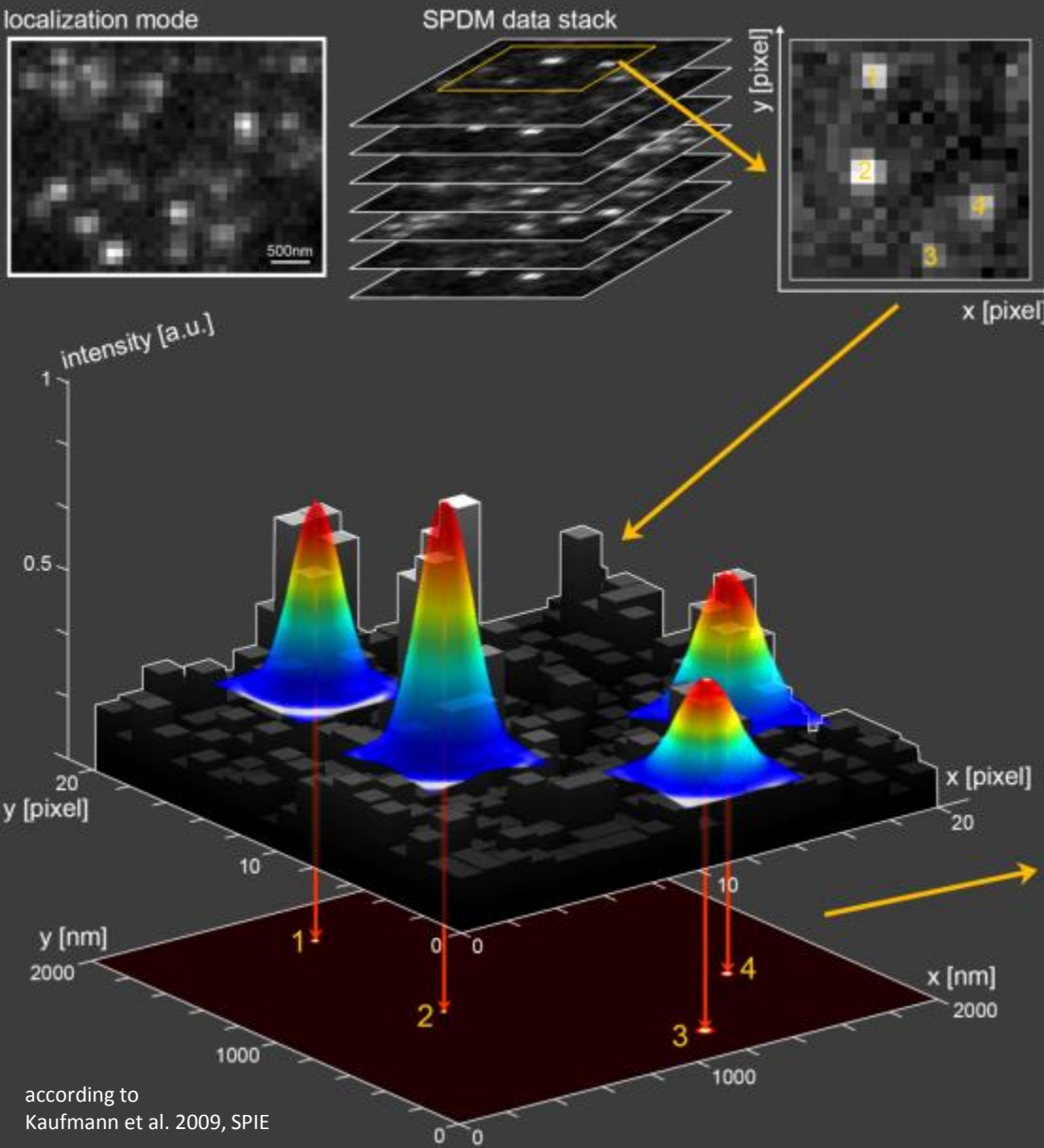
# introduction to localization microscopy

setup

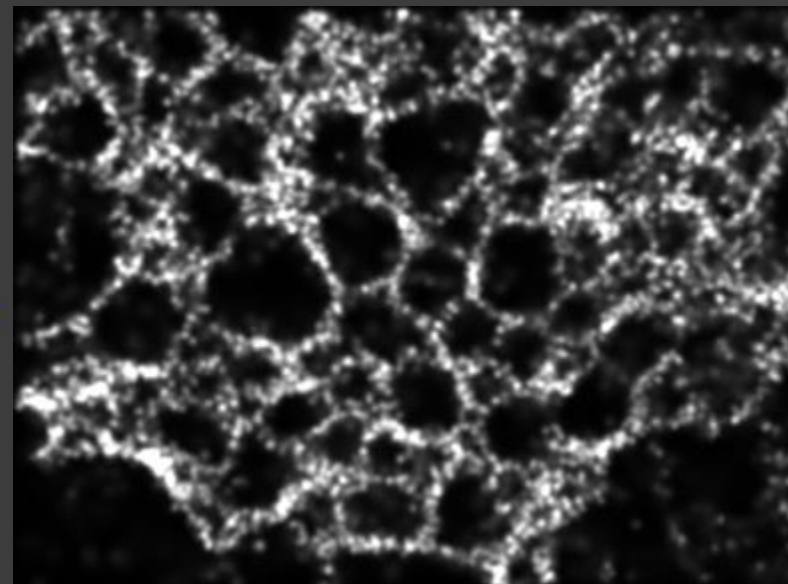
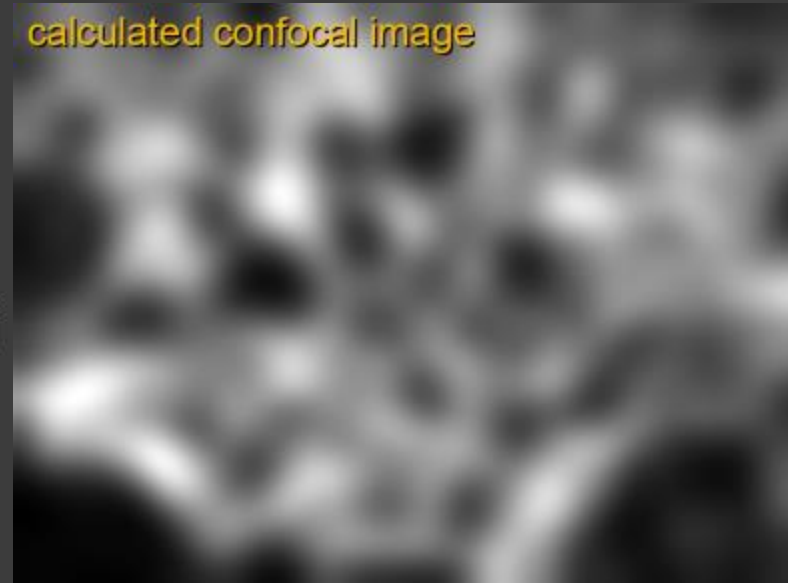


# principle of localization microscopy

image reconstruction



calculated confocal image



# principle of localization microscopy

image reconstruction

localisation accuracy  $\sigma$  of a single molecule is depended on

- width of the PSF  $s$
- number of detected photons  $N$
- background intensity  $b$
- size of the pixels on the camera  $a$

$$\sigma^2 = \frac{s^2 + a^2/12}{N} + \frac{8\pi s^4 b^2}{a^2 N^2}$$

typical model function: 2D Gaussian + linear background

$$I(x, y) = I_0 \exp \left[ -\frac{(x - x_0)^2 + (y - y_0)^2}{2s^2} \right] + b$$



# principle of localization microscopy

image reconstruction

structural resolution in localization microscopy is dependent on:

- the **localisation accuracy**  $\sigma_{xy}$  of the individual molecules
- **density** of detected molecules (sampling theorem)

$$\Rightarrow \text{structural resolution} = \sqrt{(2.35 \bar{\sigma}_{xy})^2 + (2 \bar{d}_{NN})^2}$$

$\bar{\sigma}_{xy}$  : mean localisation accuracy

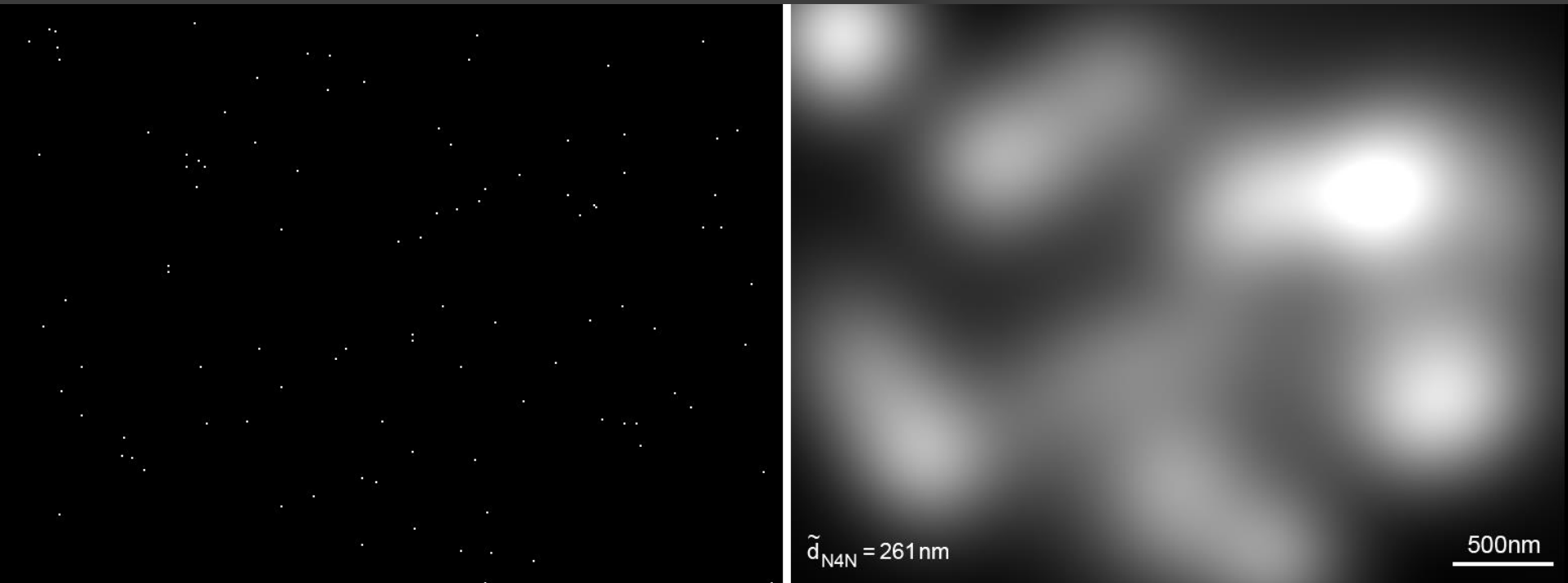
$\bar{d}_{NN}$  : mean distance to next neighbouring molecule

# principle of localization microscopy

image reconstruction

structural resolution in localization microscopy is dependent on:

- the **localisation accuracy**  $\sigma_{xy}$  of the individual molecules
- **density** of detected molecules (sampling theorem)

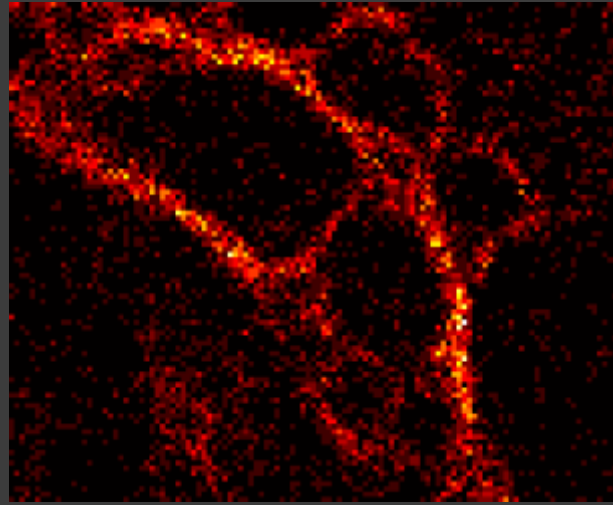


# principle of localization microscopy

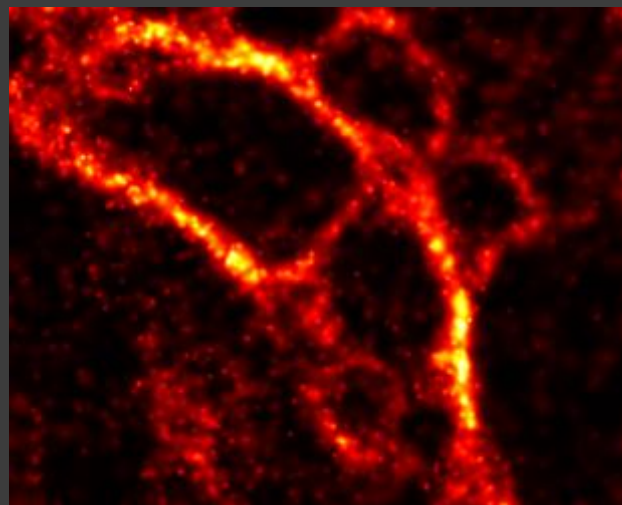
image reconstruction



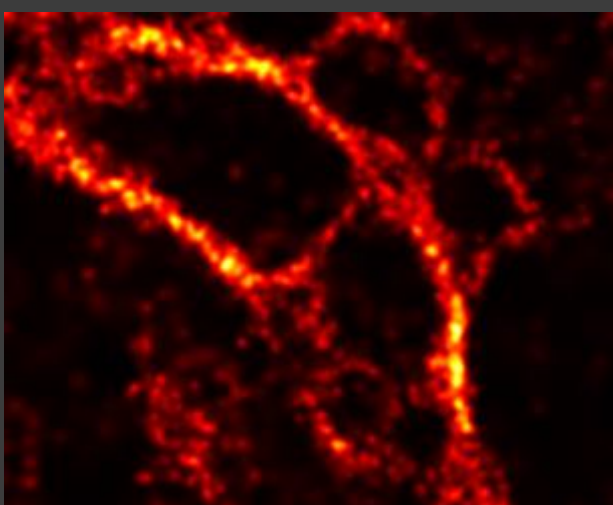
scatter plot



histogram with equal bins



visualisation of  $\sigma_{xy}$



visualisation of structural resolution

more about visualisation of  
localization microscopy data:

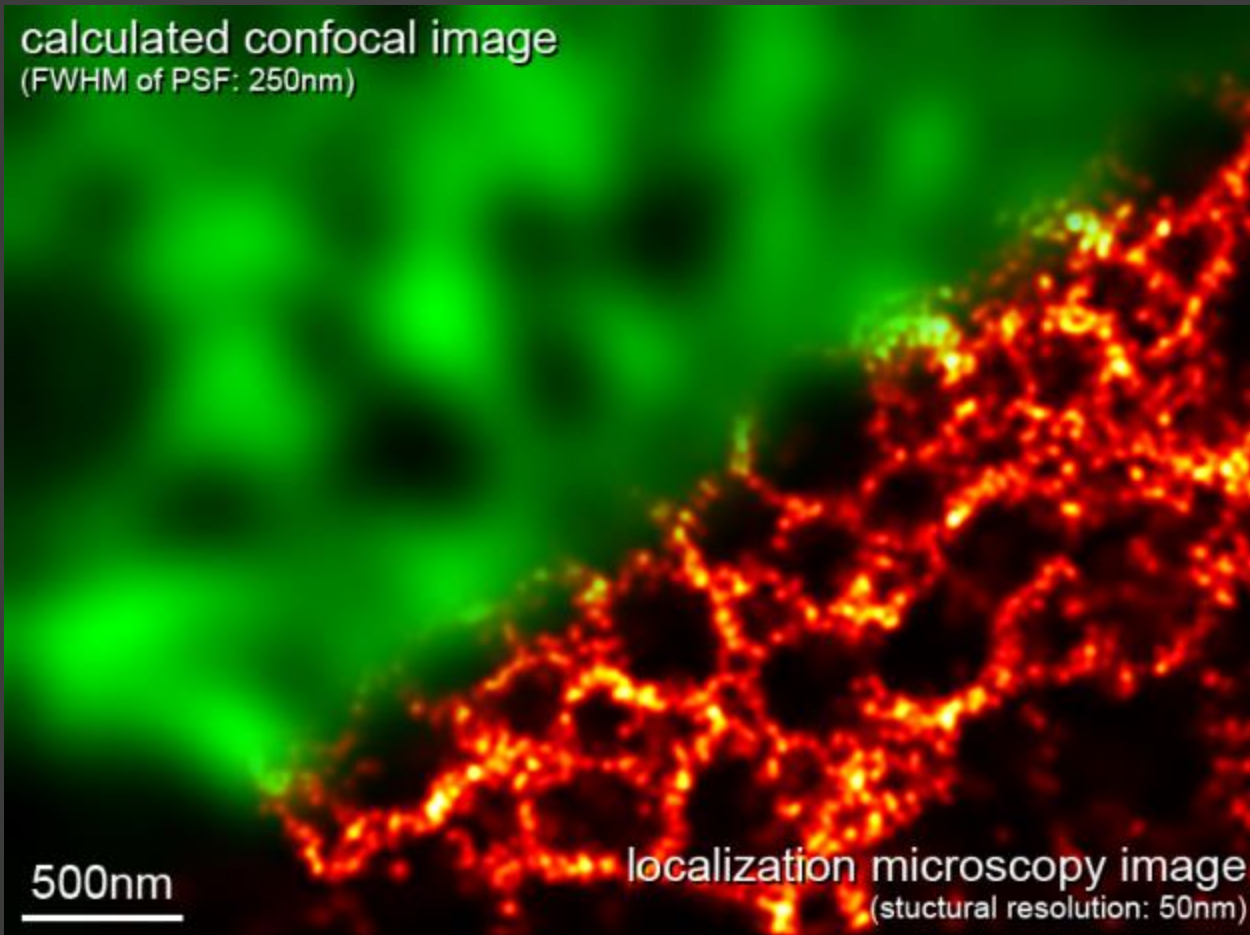
Baddeley et al., Microscopy  
and Microanalysis, 2010

# principle of localization microscopy

summary

enhanced structural resolution down the 20 nm range

calculated confocal image  
(FWHM of PSF: 250nm)



fluorophores are detected individually



single molecule information

- positions
- number of det. photons
- ...

(F)PALM and STORM

# (some) history of localization microscopy

## localisation of single molecules / point-like objects

- |                        |   |
|------------------------|---|
| Burns et al., 1985     | theoretical paper about super-resolution distance measurements using spectral characteristics       |
| Betzig, 1995           | first measurements with SNOM under cryo conditions  |
| Bornfleth et al., 1998 | CLSM measurements of 3D distances < 60 nm using fluorescent markers of different wavelengths (@ RT) |
| Heilemann et al., 2002 | using single molecule live time instead of colours to measure distances of 40 nm                    |

## localisation of many molecules to reconstruct structural information

2006: (PALM, FPALM, STORM) – photoswitchable / photoactivatable dyes

2008: (dSTORM, SPDM, GSDIM) – using standard fluorophores

# (F)PALM – (fluorescence) photo activated localization microscopy

uses **photoactivatable fluorophores** (e.g. PA-GFP, caged Fluorescein, ...)

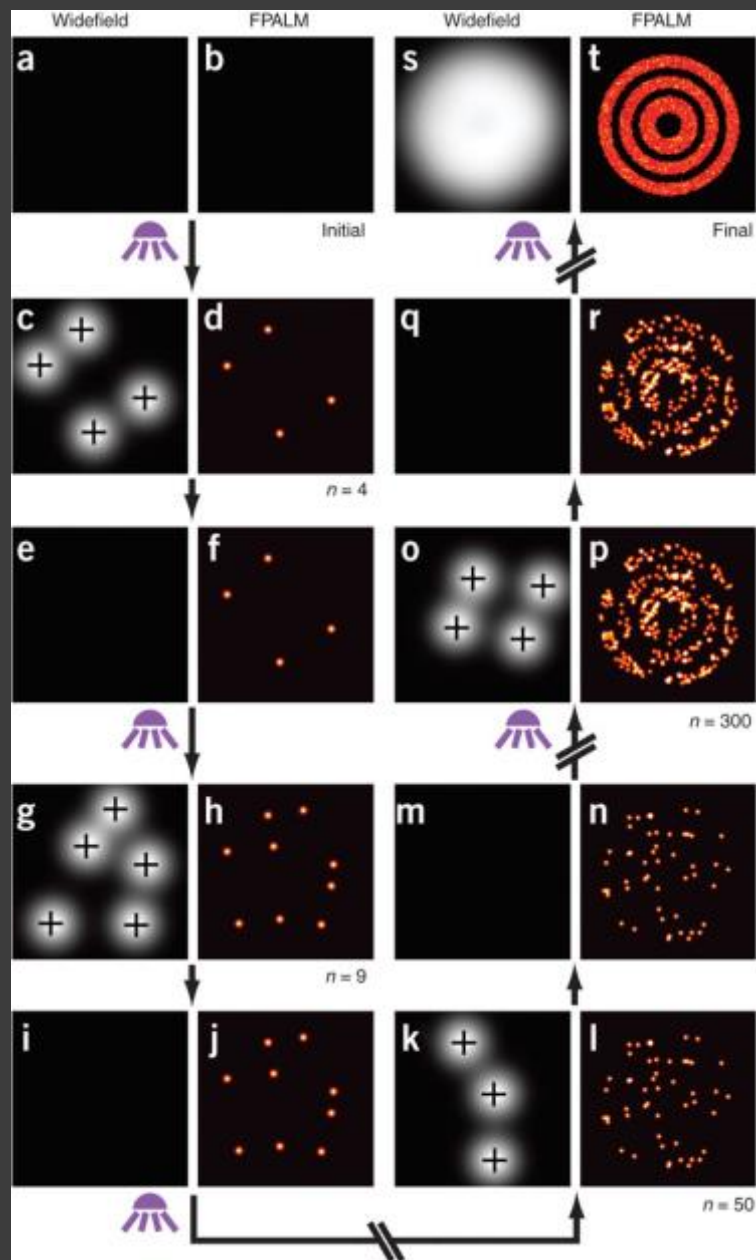
- at the beginning all fluorophores are “dark” (not fluorescent at their excitation wavelength)
- fluorophores can be “activated” to a “bright” state
- after bleaching the molecules they do not reappear

→ **irreversible process**

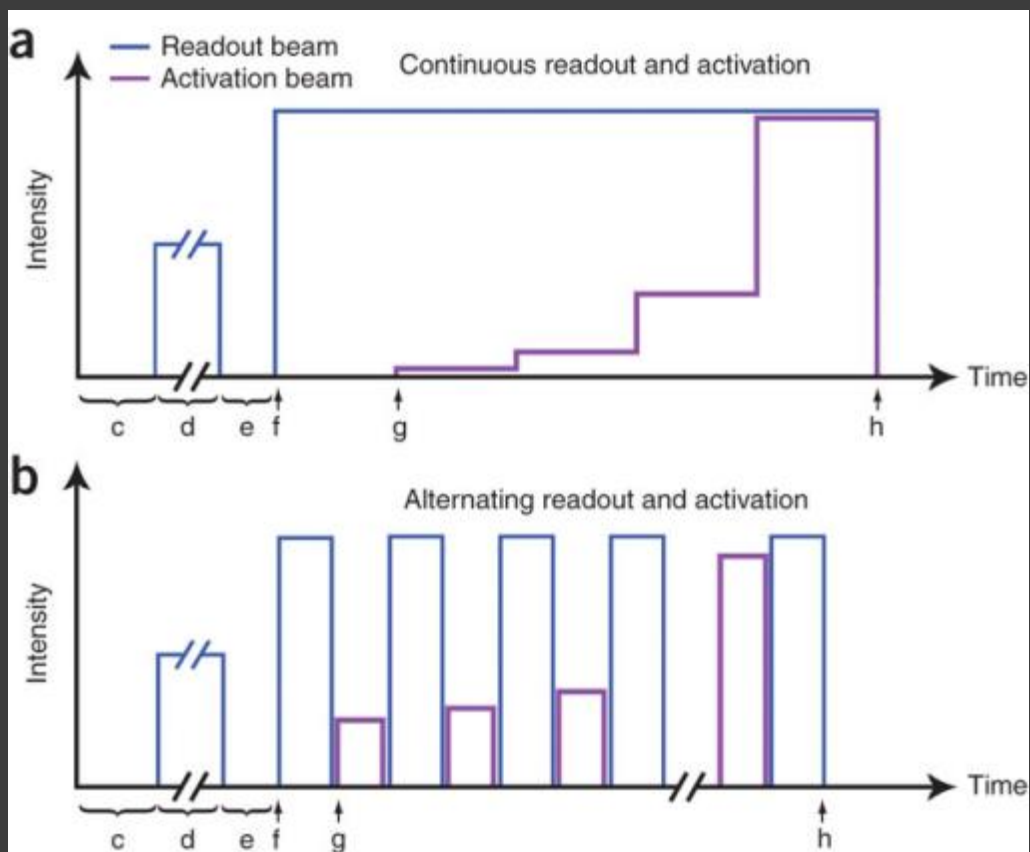
original publications:

- PALM: Betzig et al., Science, 2006
- FPALM: Hess et al., Biophysical Journal, 2006

# (F)PALM – (fluorescence) photo activated localization microscopy

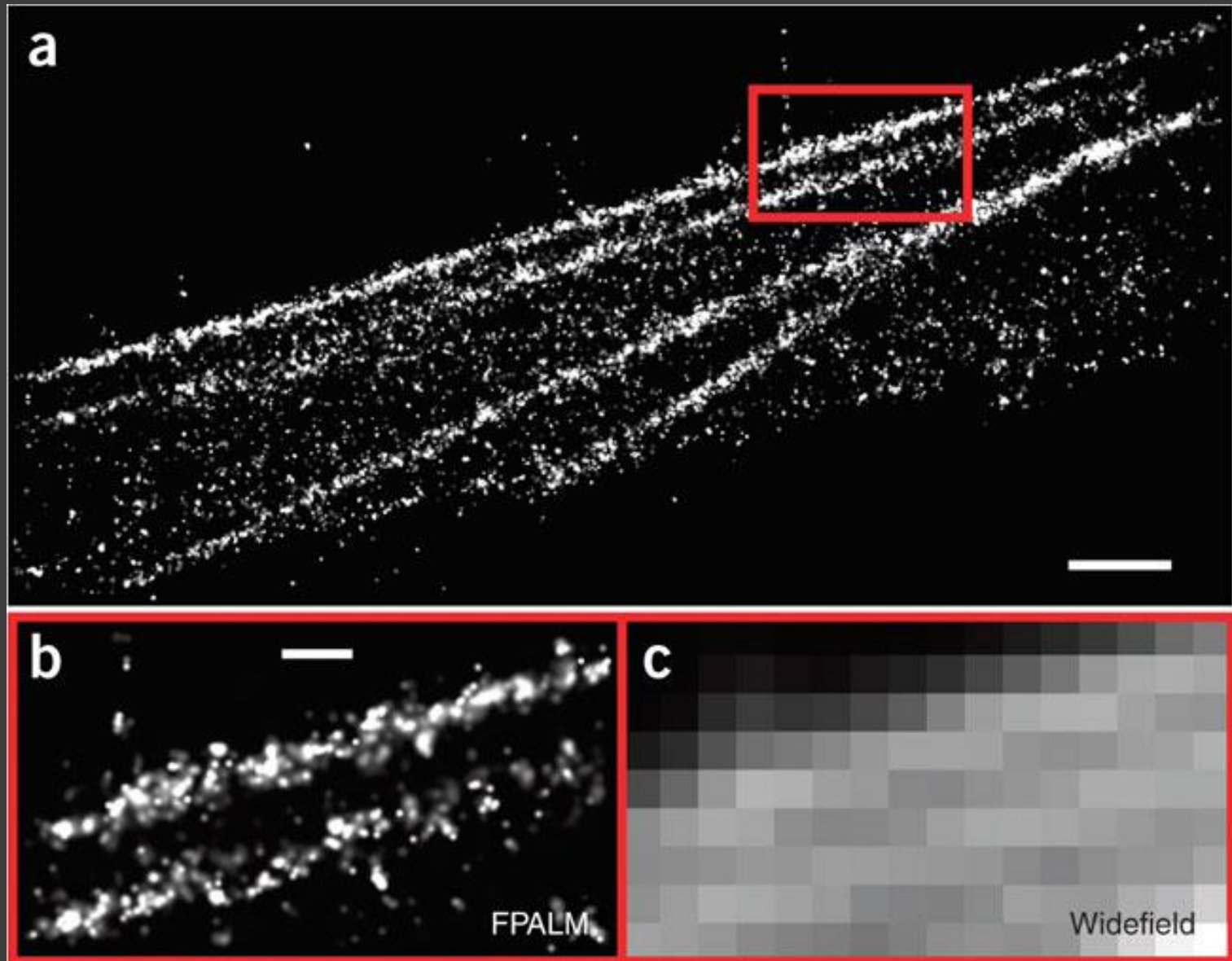


Gould et al., Nature Protocols, 2009





# (F)PALM – (fluorescence) photo activated localization microscopy



# STORM – stochastic optical reconstruction microscopy

uses **photoswitchable fluorophores** (dye pairs (e.g. Cy3-Cy5) or proteins like Dronpa)

- fluorophores can be switched many times between a “bright” and a “dark” state

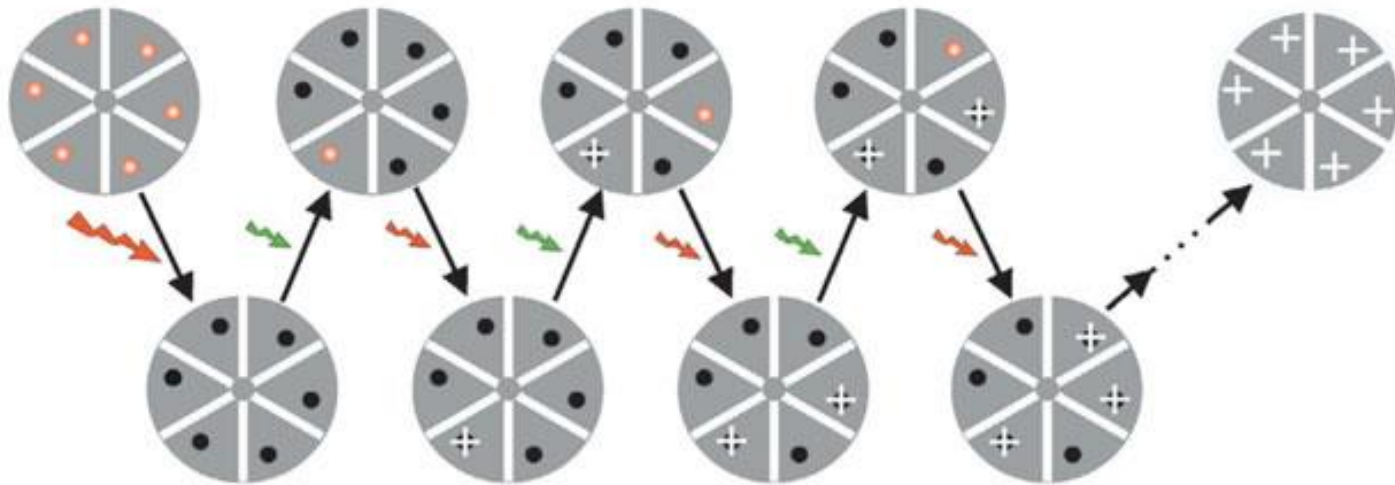
→ reversible process

original publication:

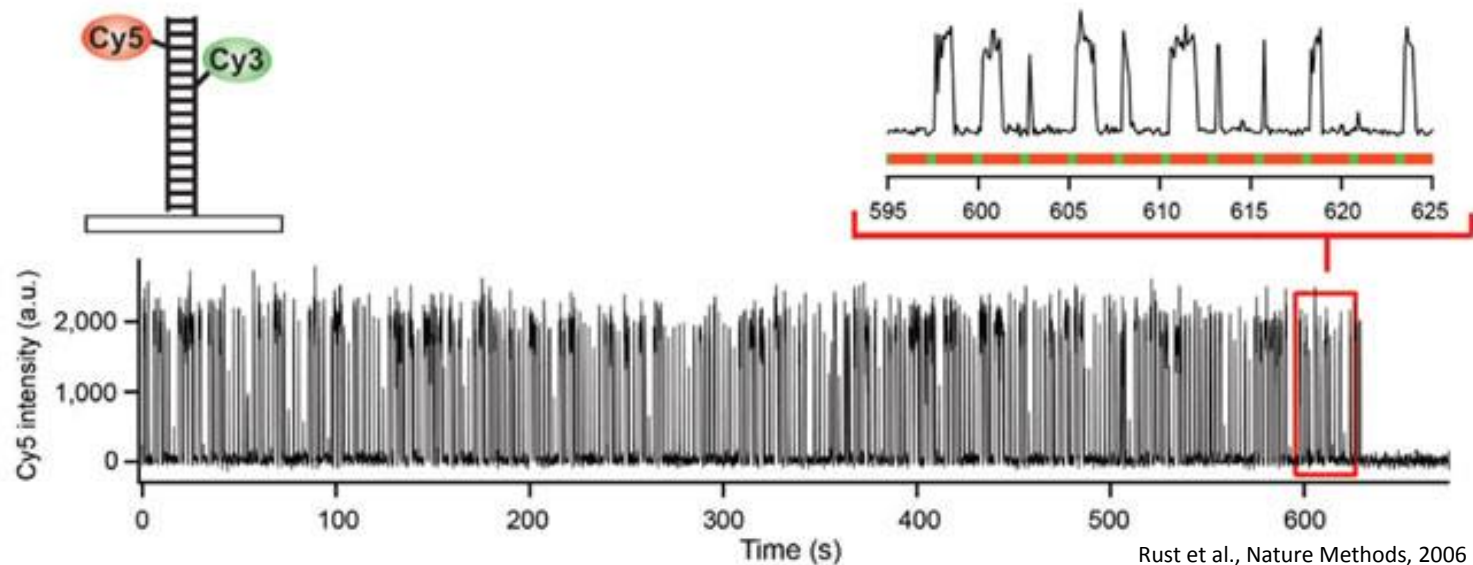
- Rust et al., Nature Methods, 2006

# STORM – stochastic optical reconstruction microscopy

**a**

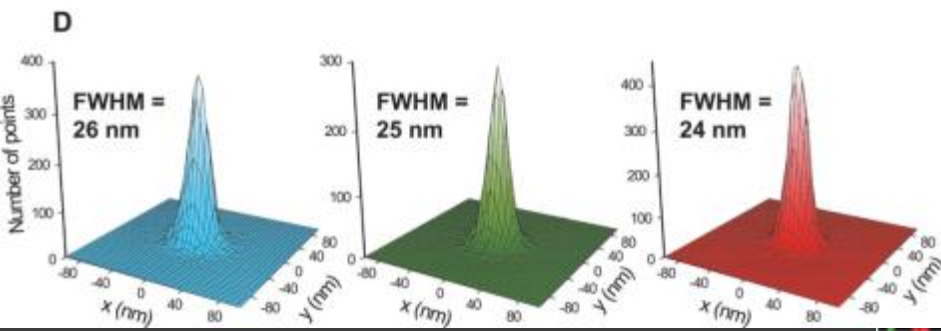
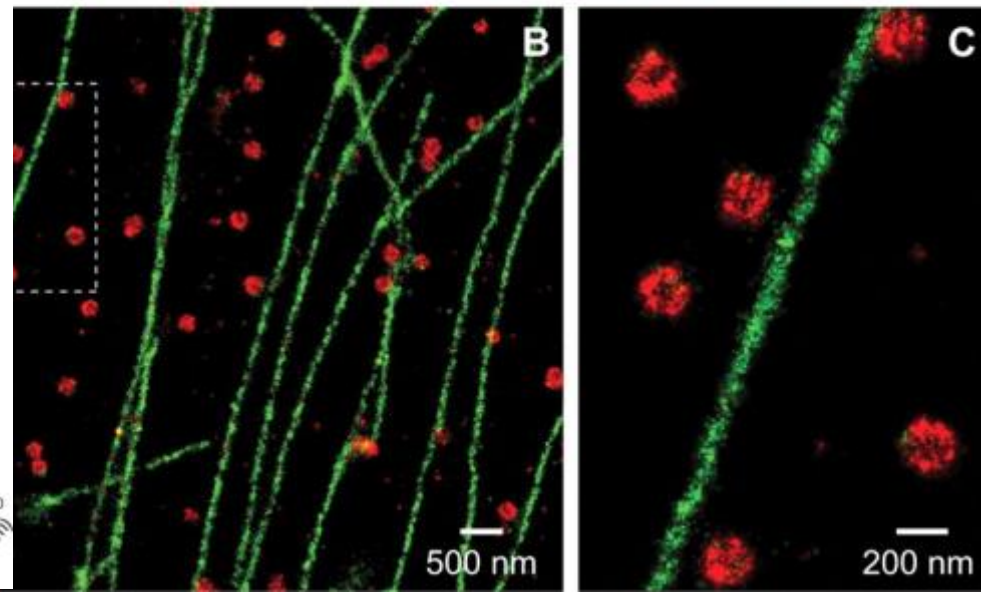
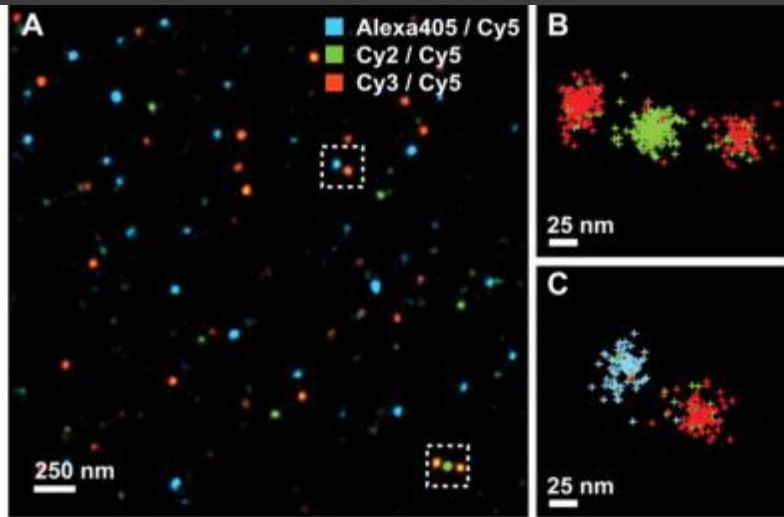
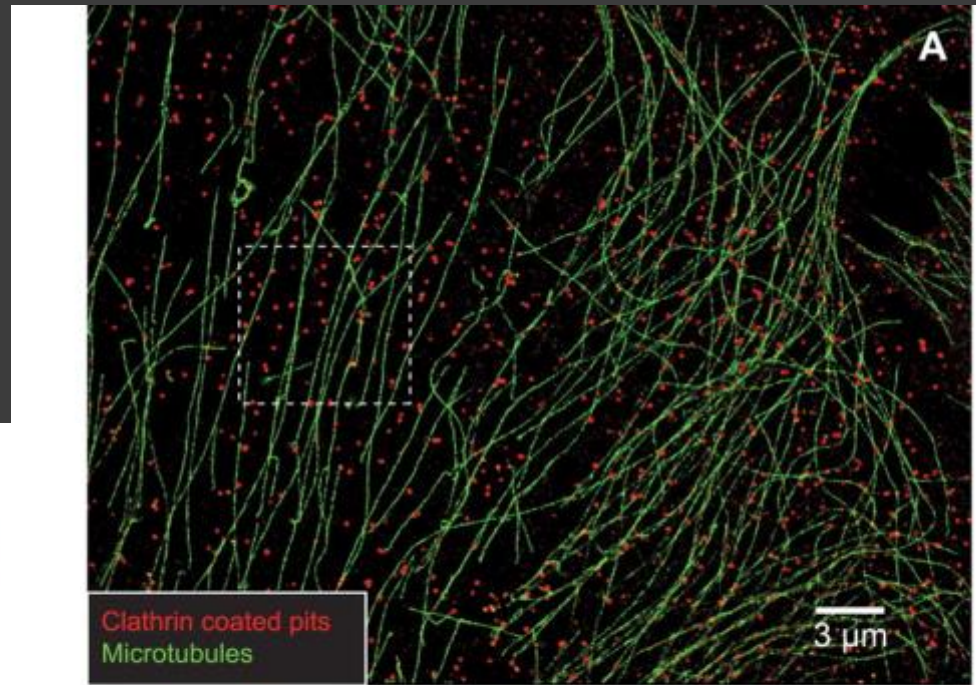


**b**



# STORM – stochastic optical reconstruction microscopy

Bates et al., Science, 2007



dSTORM, SPDM, GSDIM, ...

# dSTORM, SPDM, GSDIM, ...

direct STORM

spectral position determination microscopy

ground state depletion microscopy followed by individual molecule return

uses **standard fluorophores** (e.g. Alexa and Atto dyes, GFP, YFP, RFP, ...)

- switching mechanism based on **a light induced long-lived “dark” state**
- stochastic recovery to “bright” (fluorescent) state is used for optical isolation of the single molecule signals

original publication:

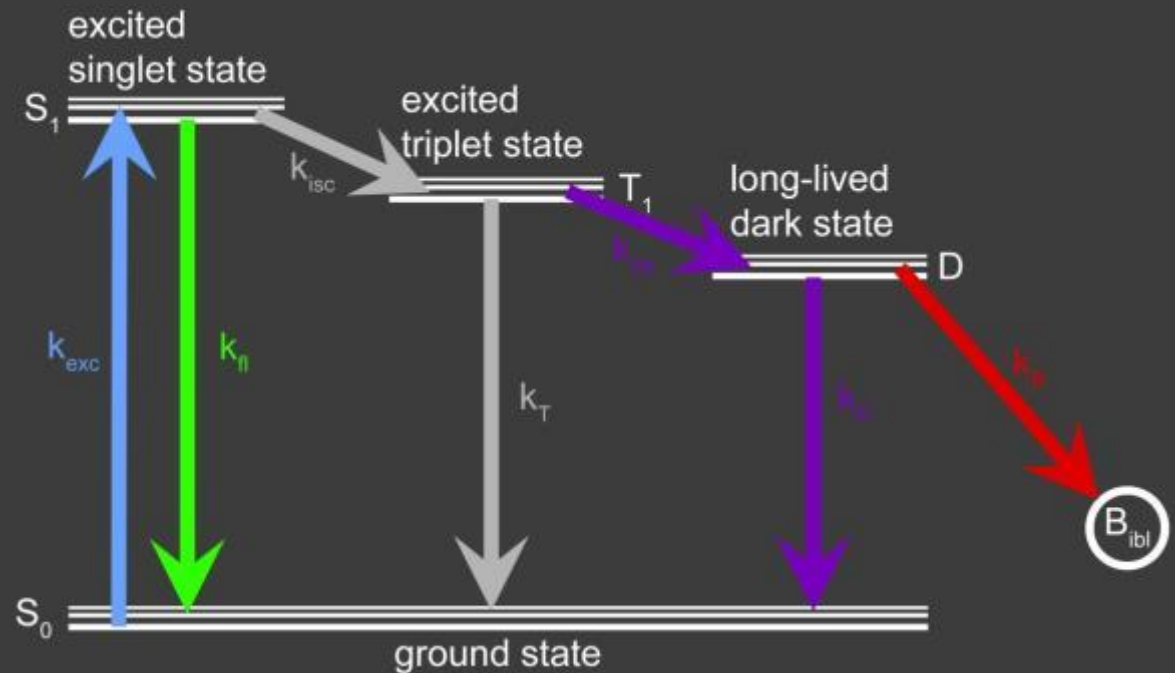
- dSTORM: Heilemann et. al., Angewandte Chemie International Edition, 2008
- SPDM: Lemmer et al., Applied Physics B, 2008
- GSDIM: Fölling et al., Nature Methods, 2008

# dSTORM, SPDM, GSDIM, ...

light induced long-lived (ms – 100 s) dark state

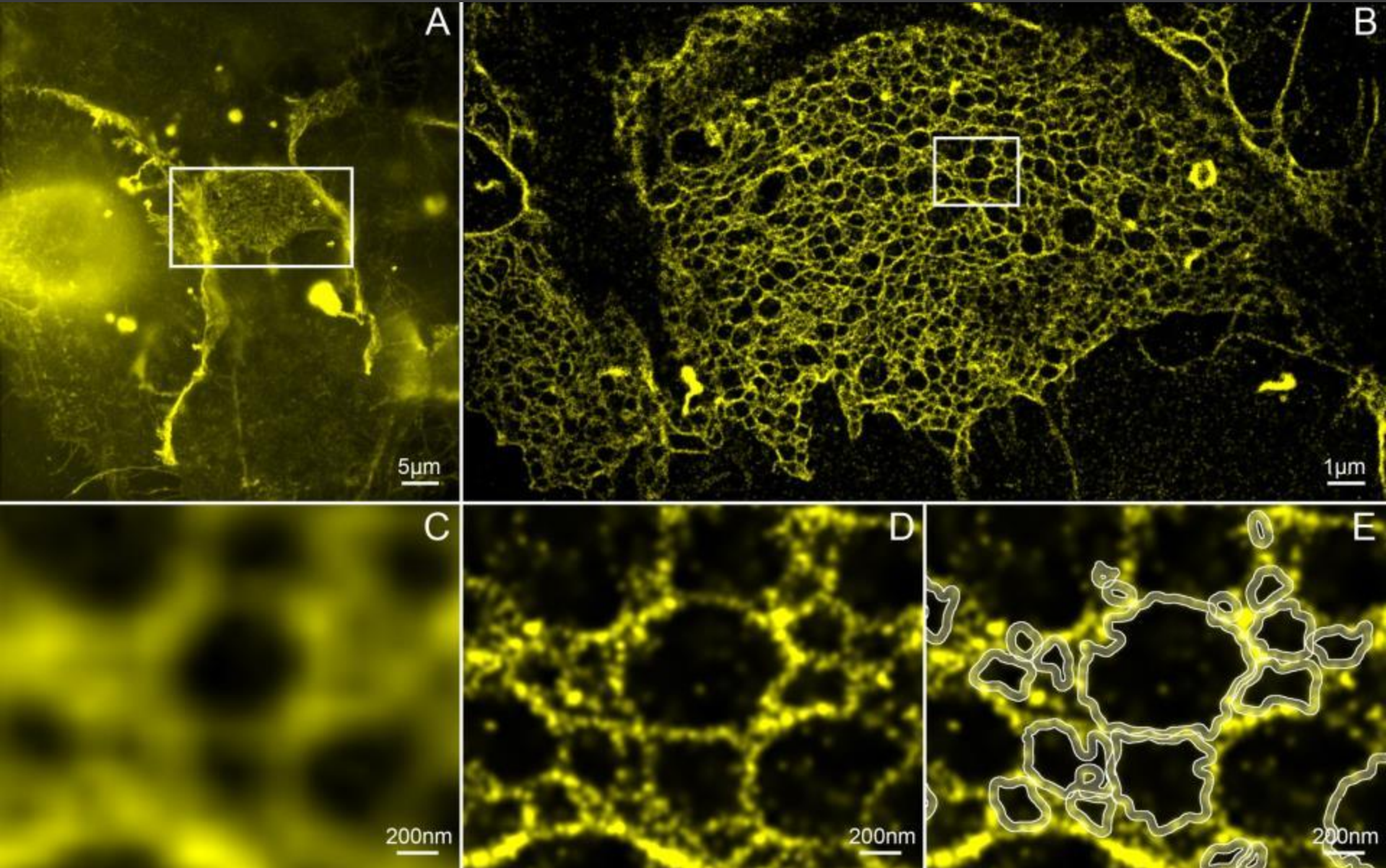
critical parameters for driving fluorophores into the long-lived dark state:

- illumination intensity
- wavelength
- embedding medium



**statistical recovery** of fluorophores from the light induced long-lived dark state can be used for **optical isolation** of single molecules

# dSTORM, SPDM, GSDIM, ...





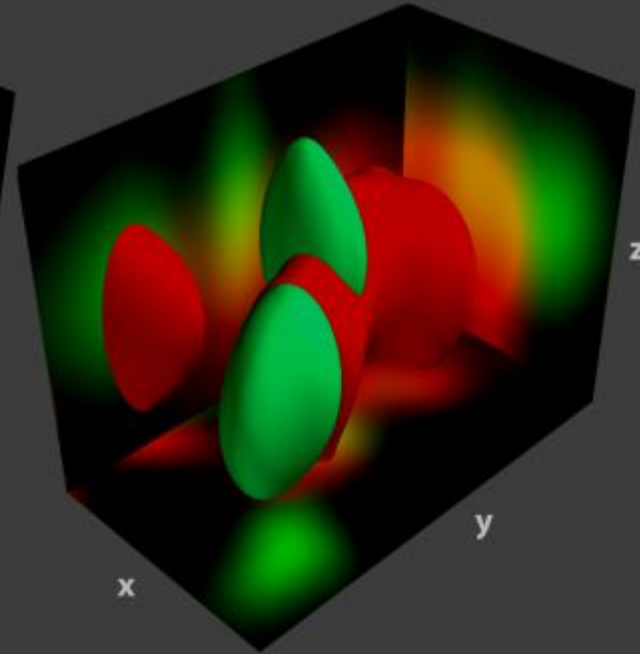
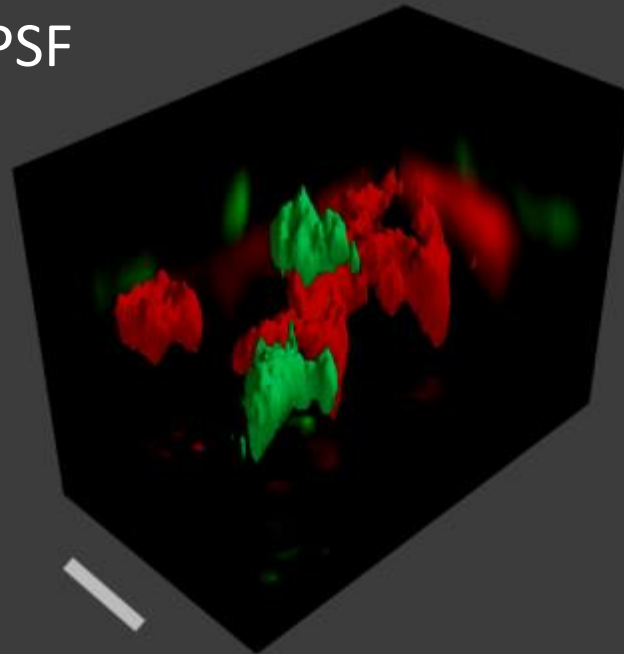
# 3D

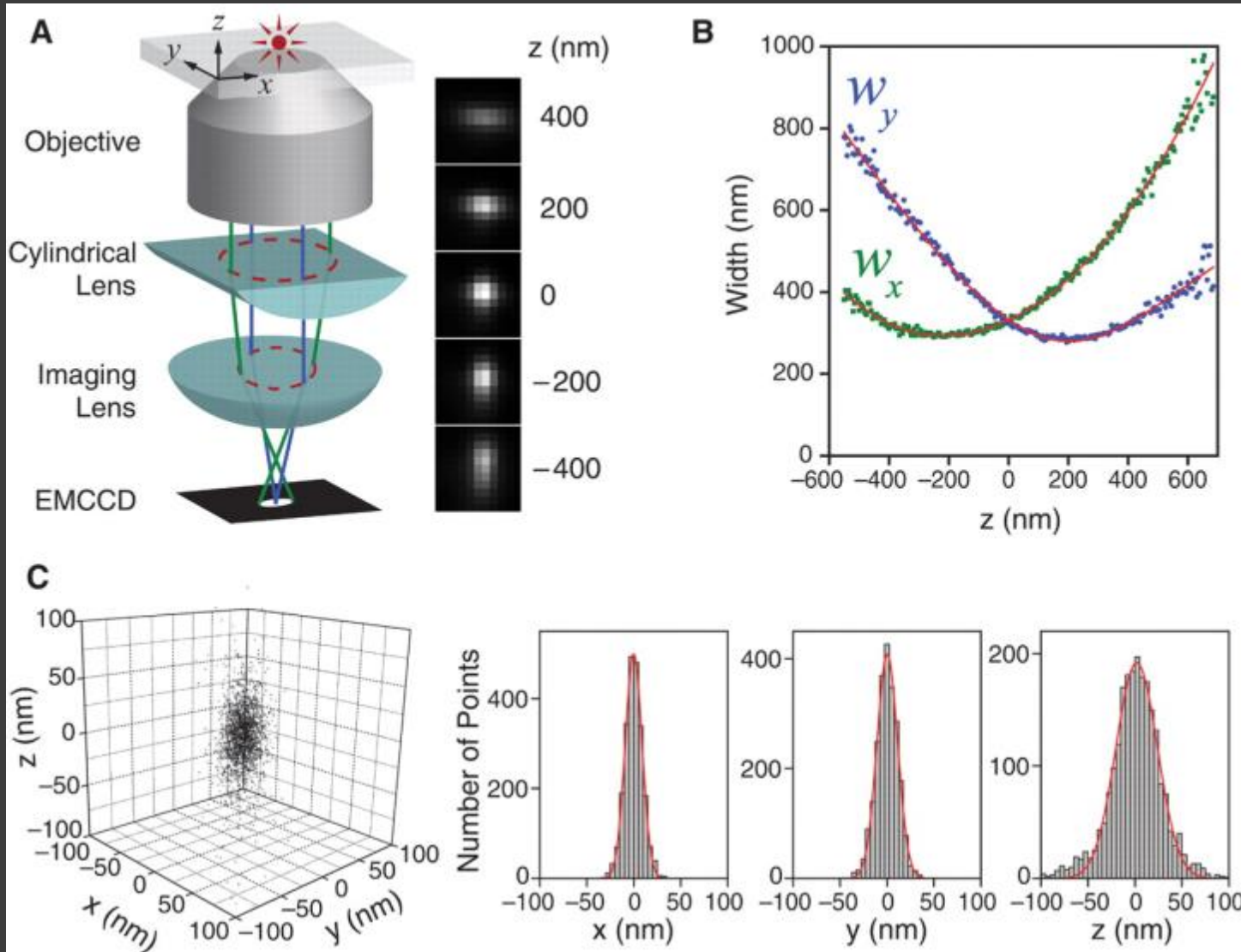
astigmatic (elliptical) PSF

biplane

double helical PSF

iPALM

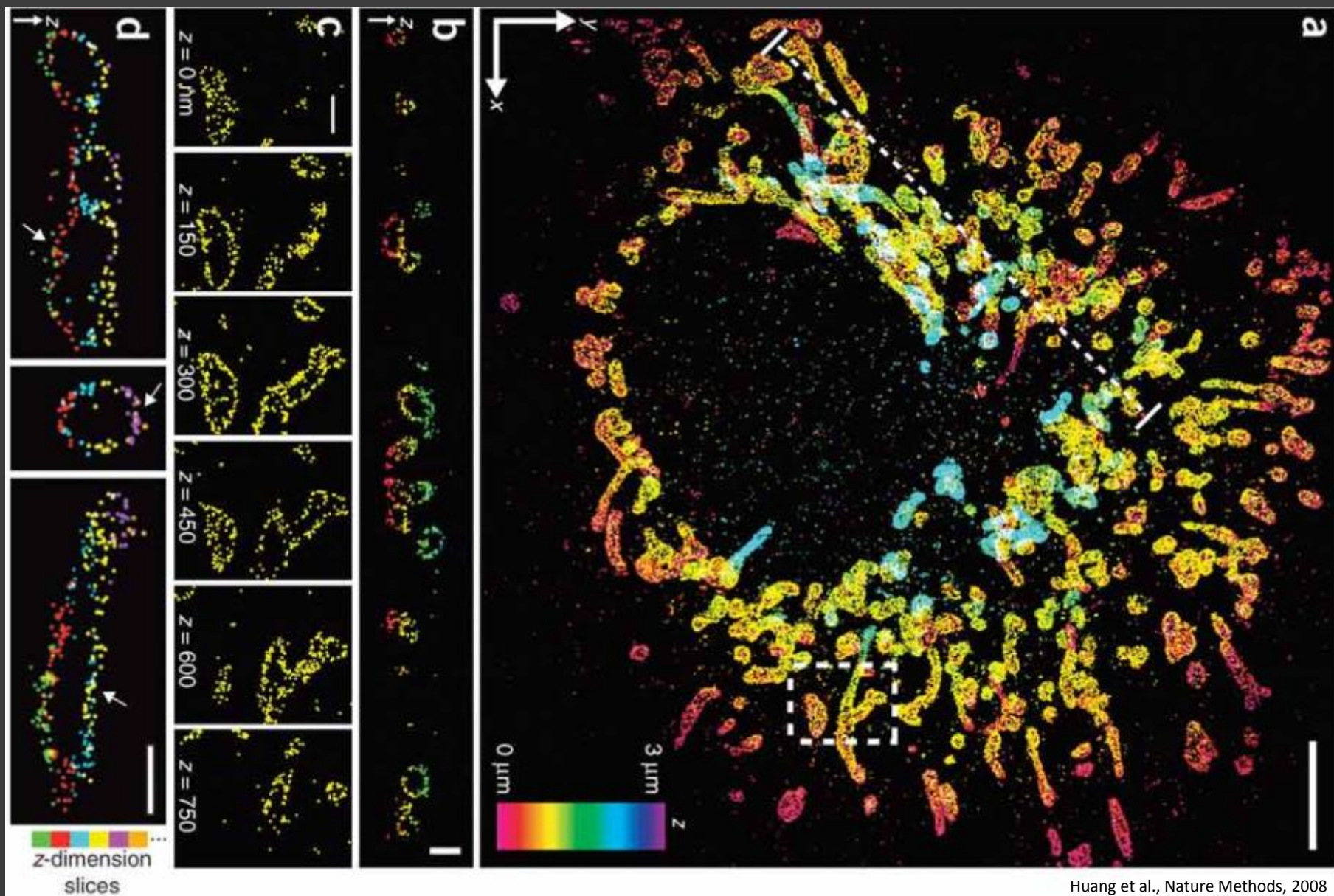




resolution  
lateral: 30 nm  
axial: 50 nm

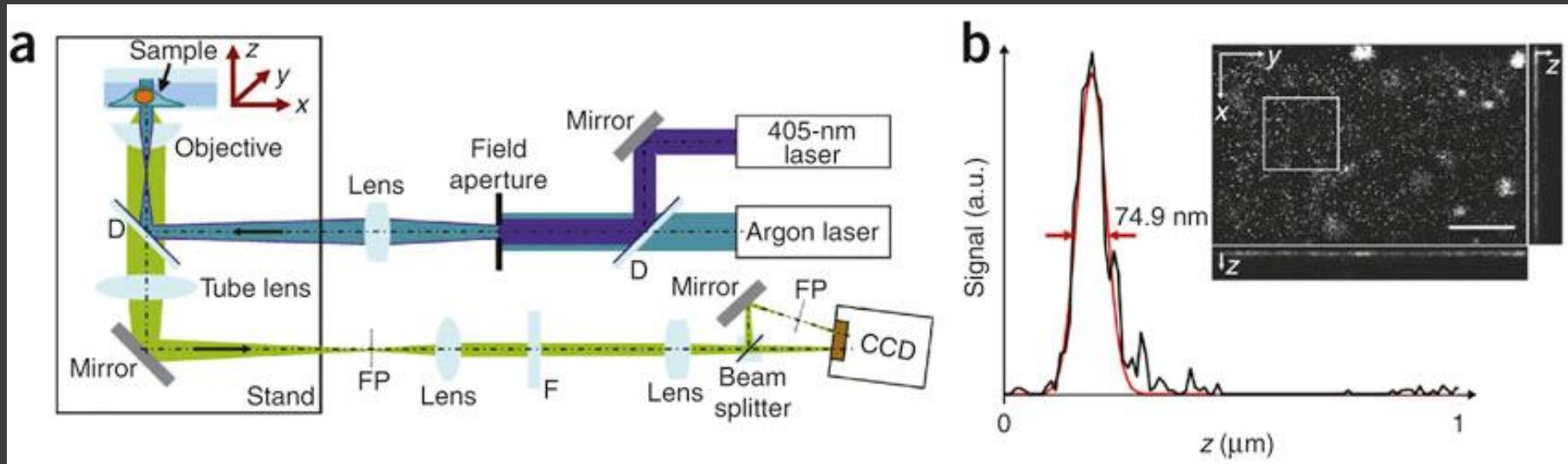
Alexa405-Cy5-mitochondria

astigmatic imaging system



imaging of two different axial plane simultaneously

→ fitting of 3D-PSF yields 3D position of the fluorophore



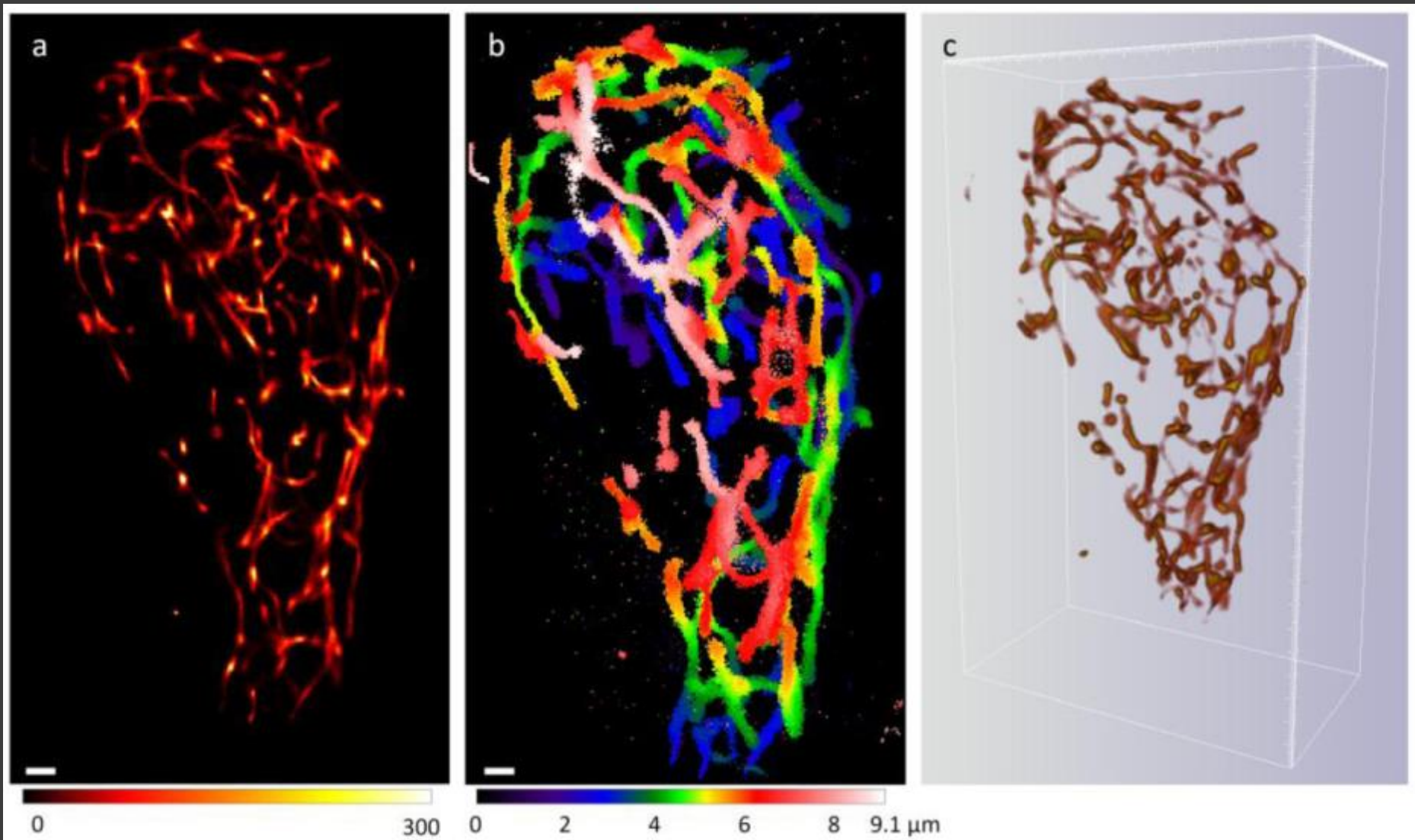
Juette et al., Nature Methods, 2008

resolution

lateral: 30 nm

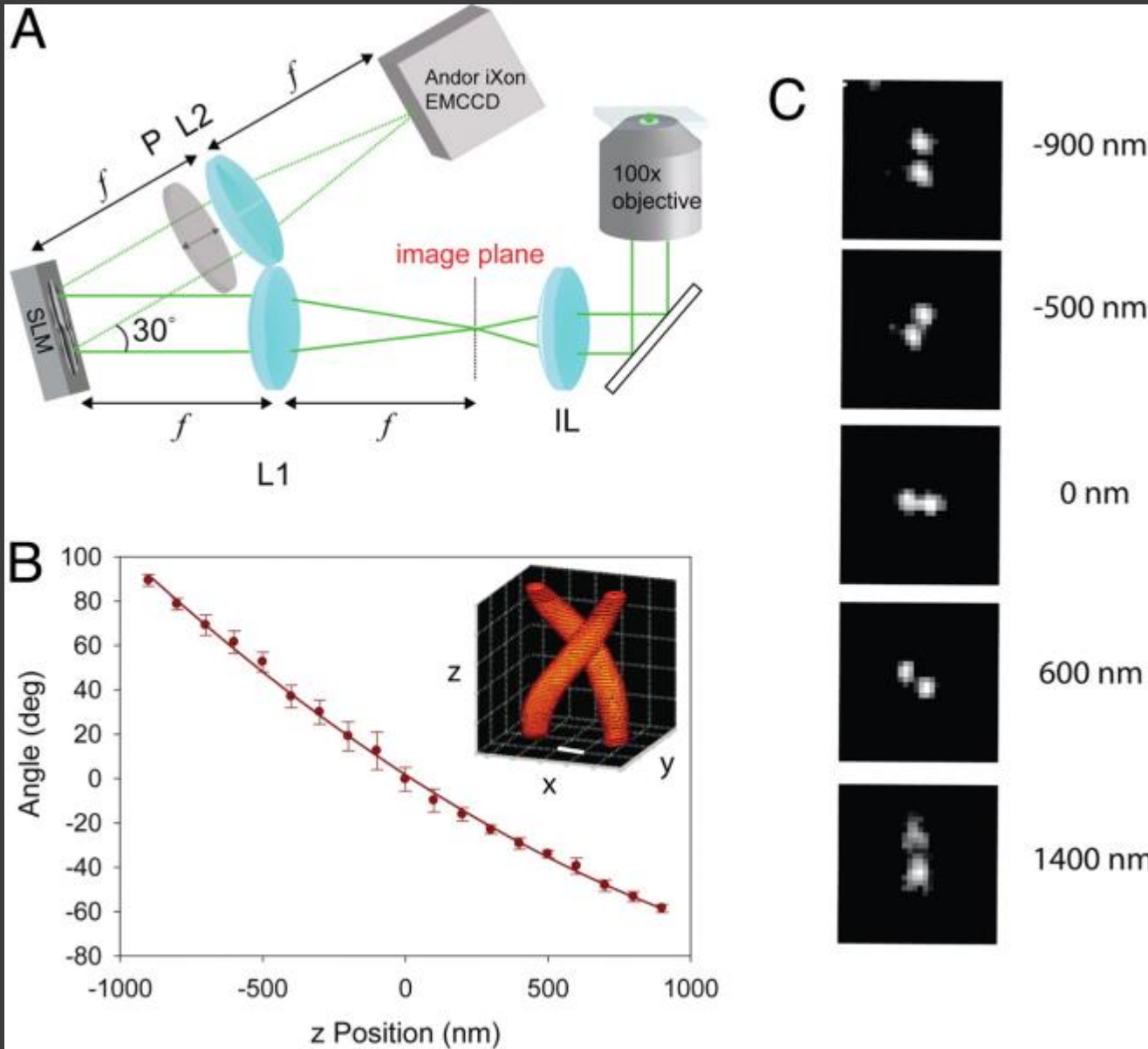
axial: 60 nm

mtEos2-mitochondria



# 3D

double helical PSF



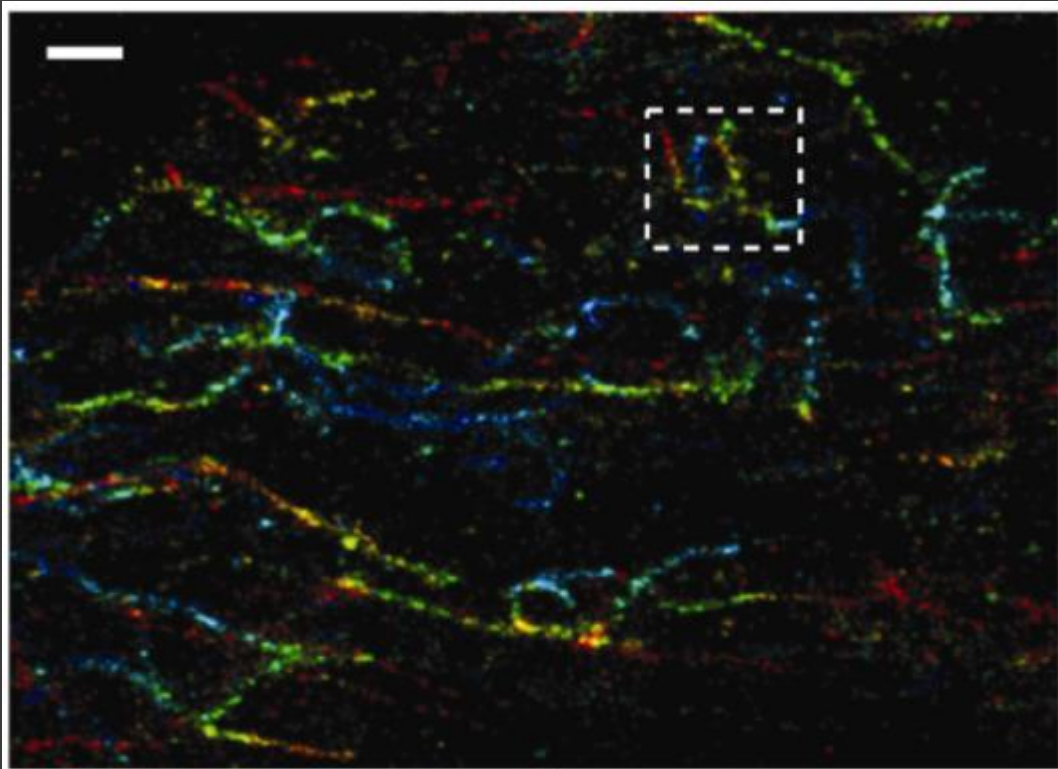
fitting of two 2D Gaussians

→ 3D position of the molecule

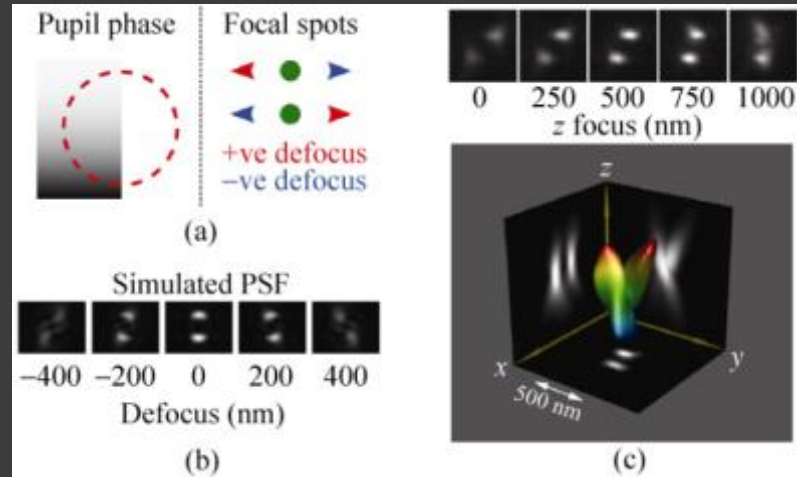
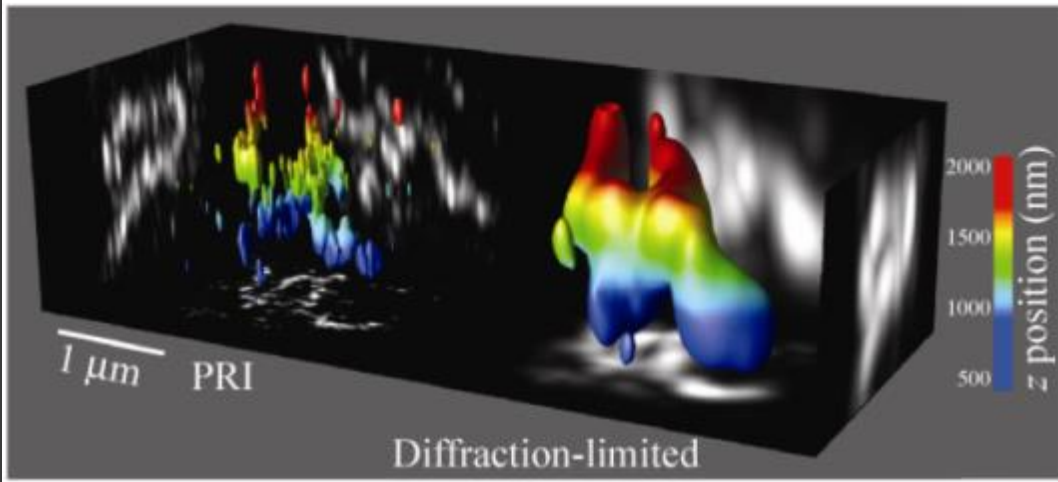
# 3D

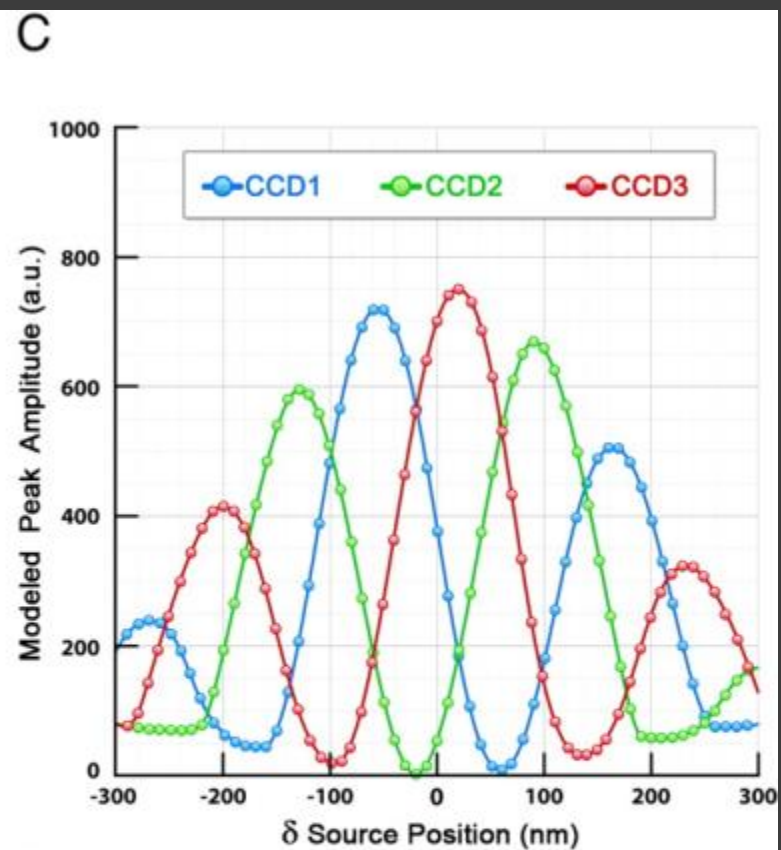
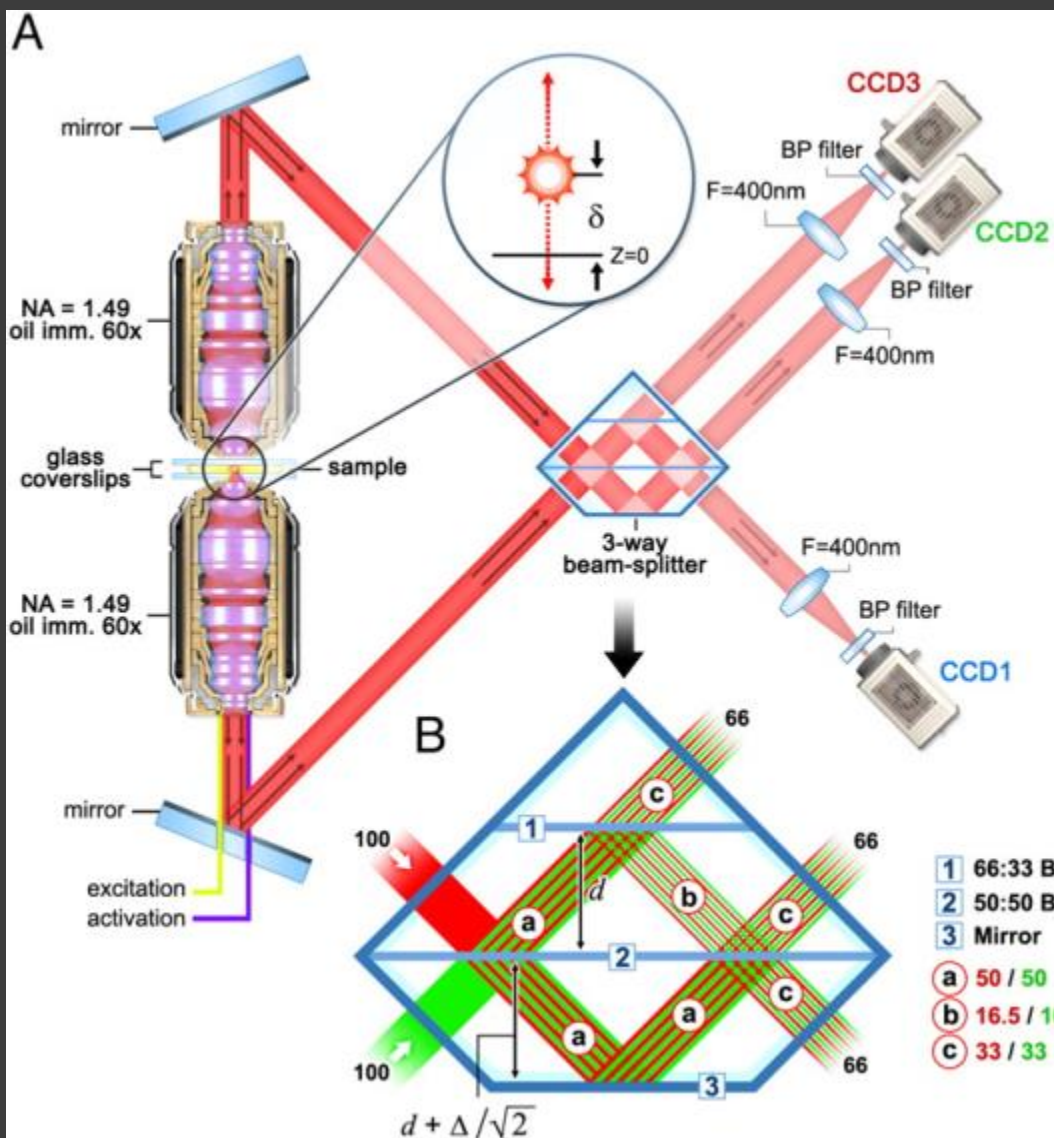
double helical PSF

resolution  
xy: 30 nm  
z: < 100 nm



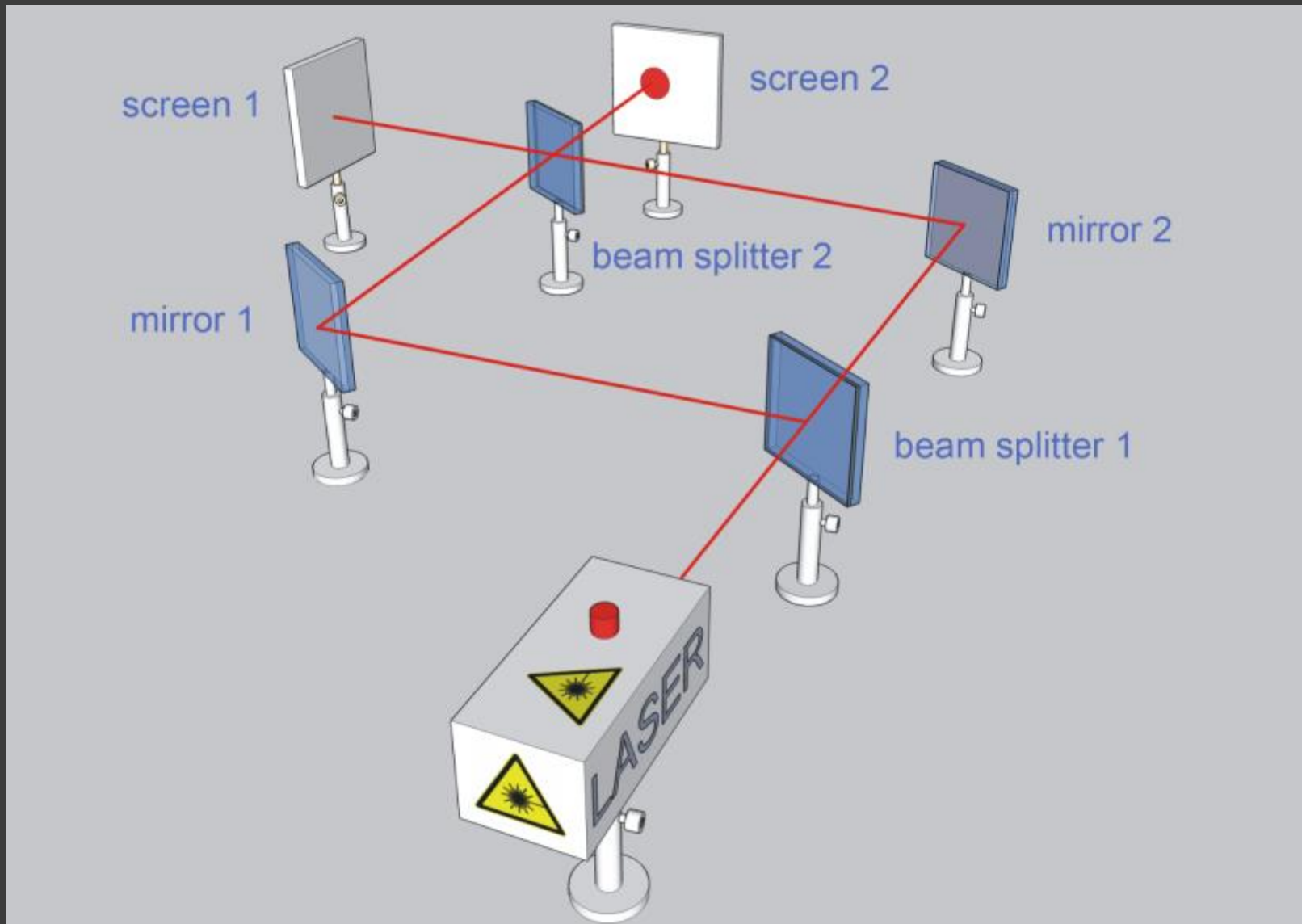
(a)

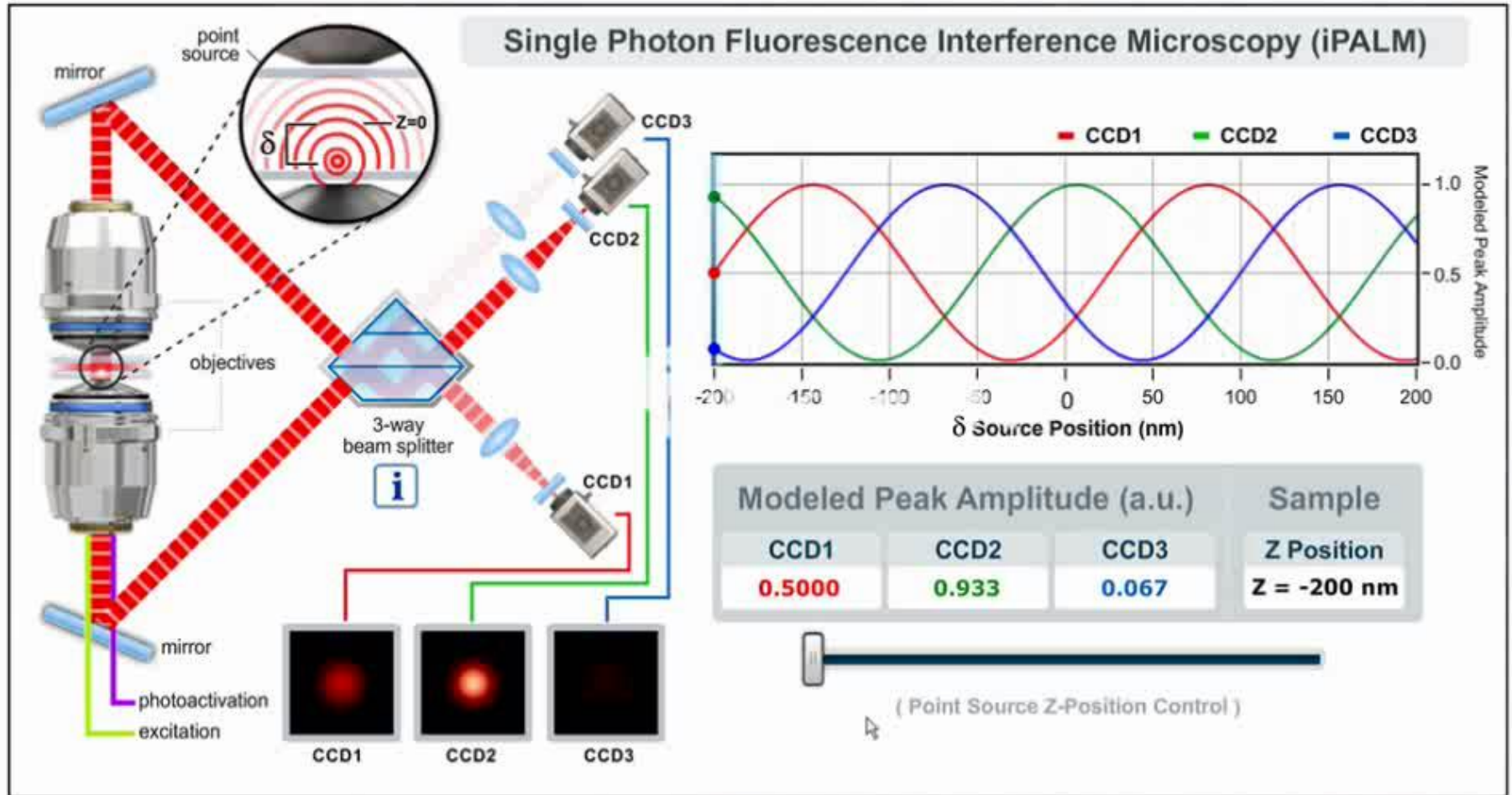


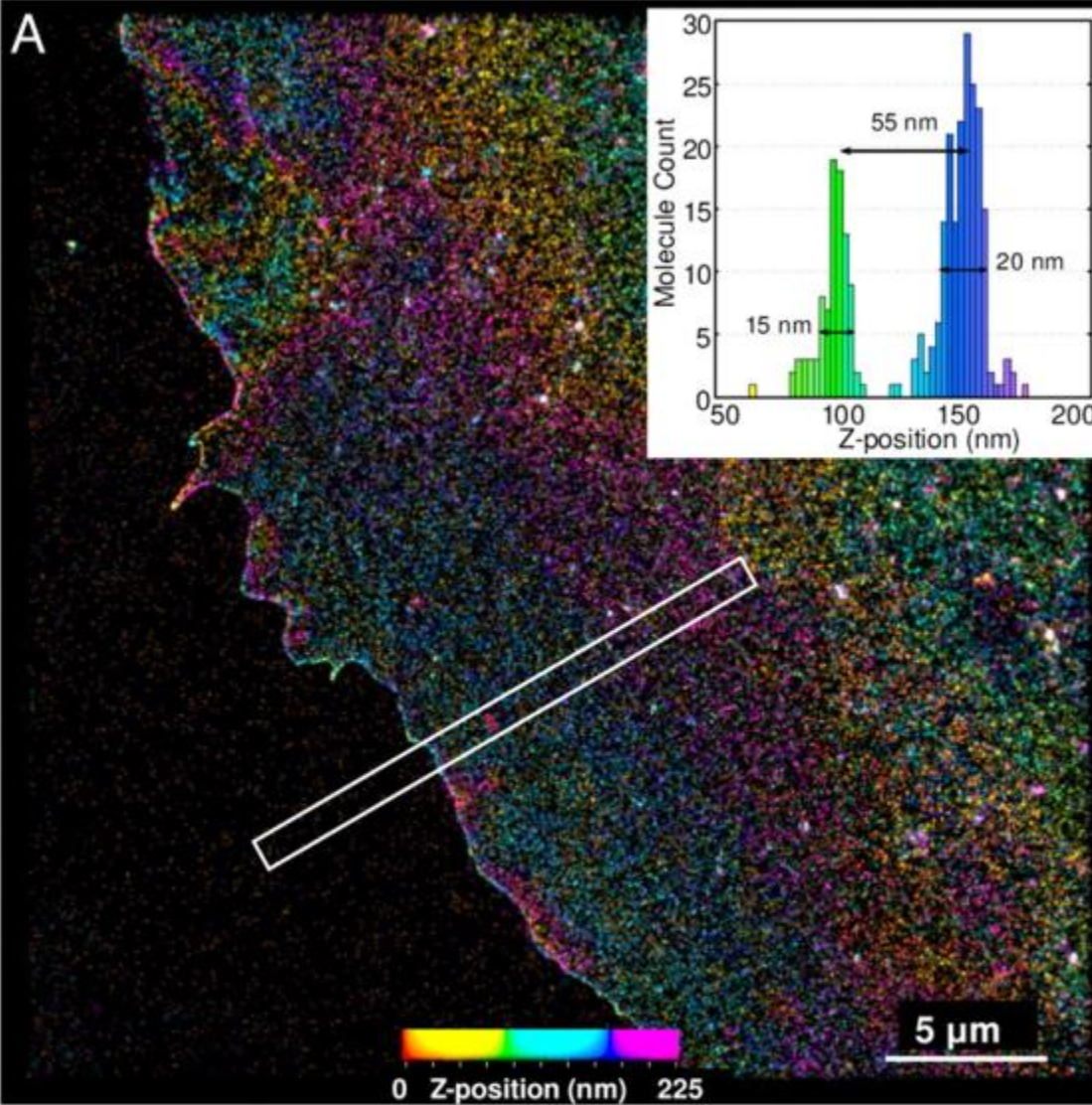




## Mach-Zehnder-Interferometer



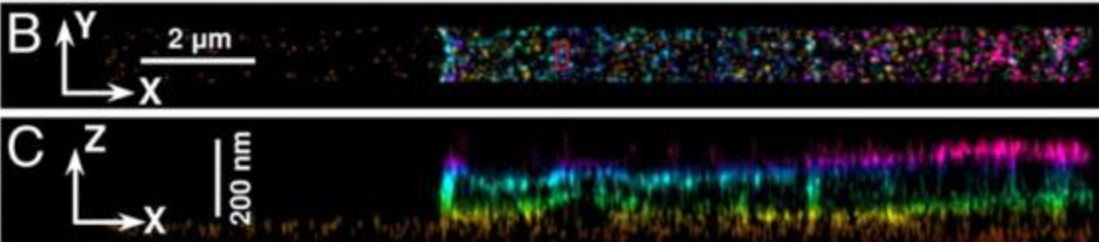




td-EosFP - VSVG

3D  
iPALM

resolution:  
50 nm in all 3 directions



two examples for “live-cell” applications

# live-cell STORM (dSTORM)

resolution

2D

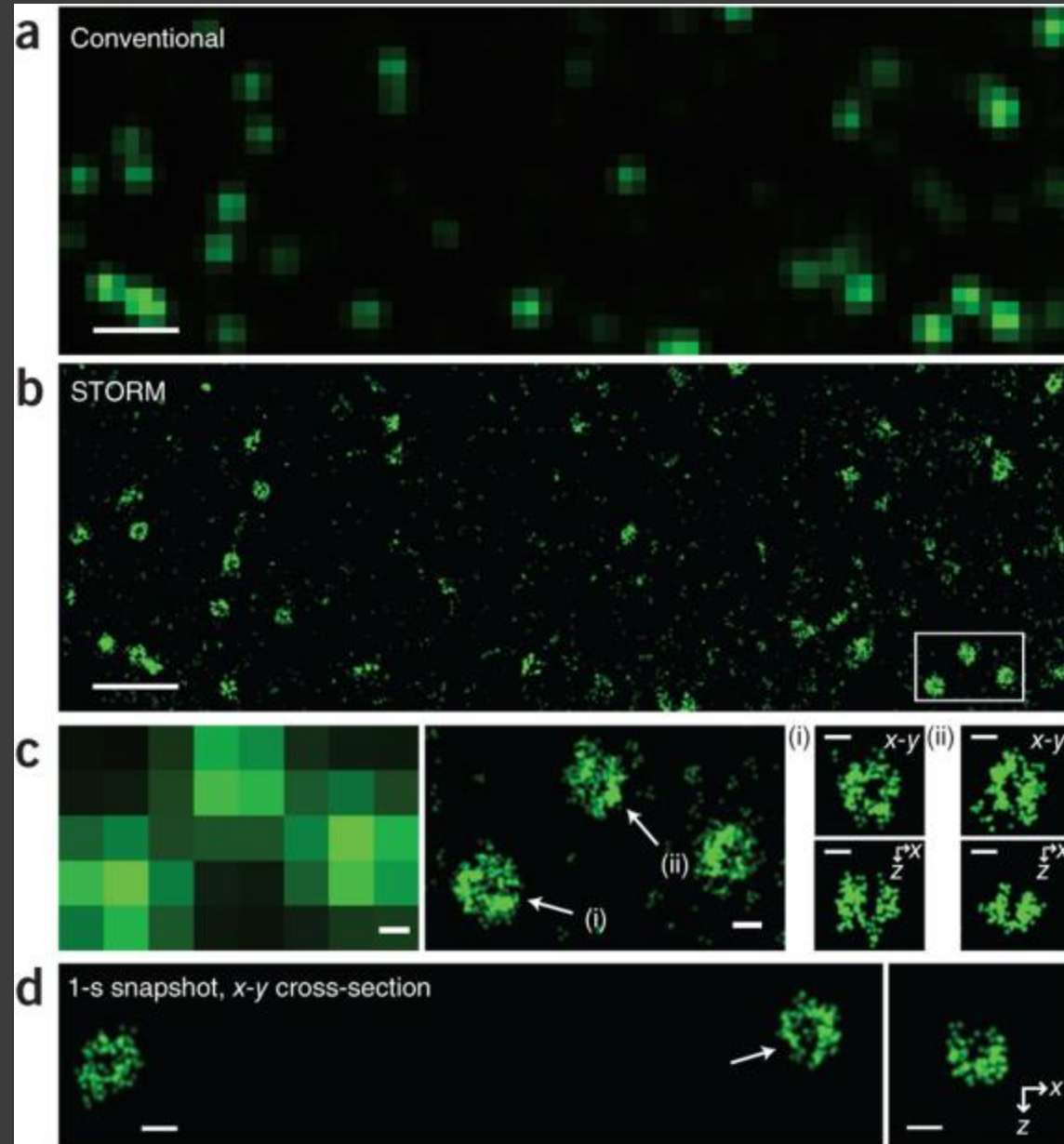
spatial: 25 nm

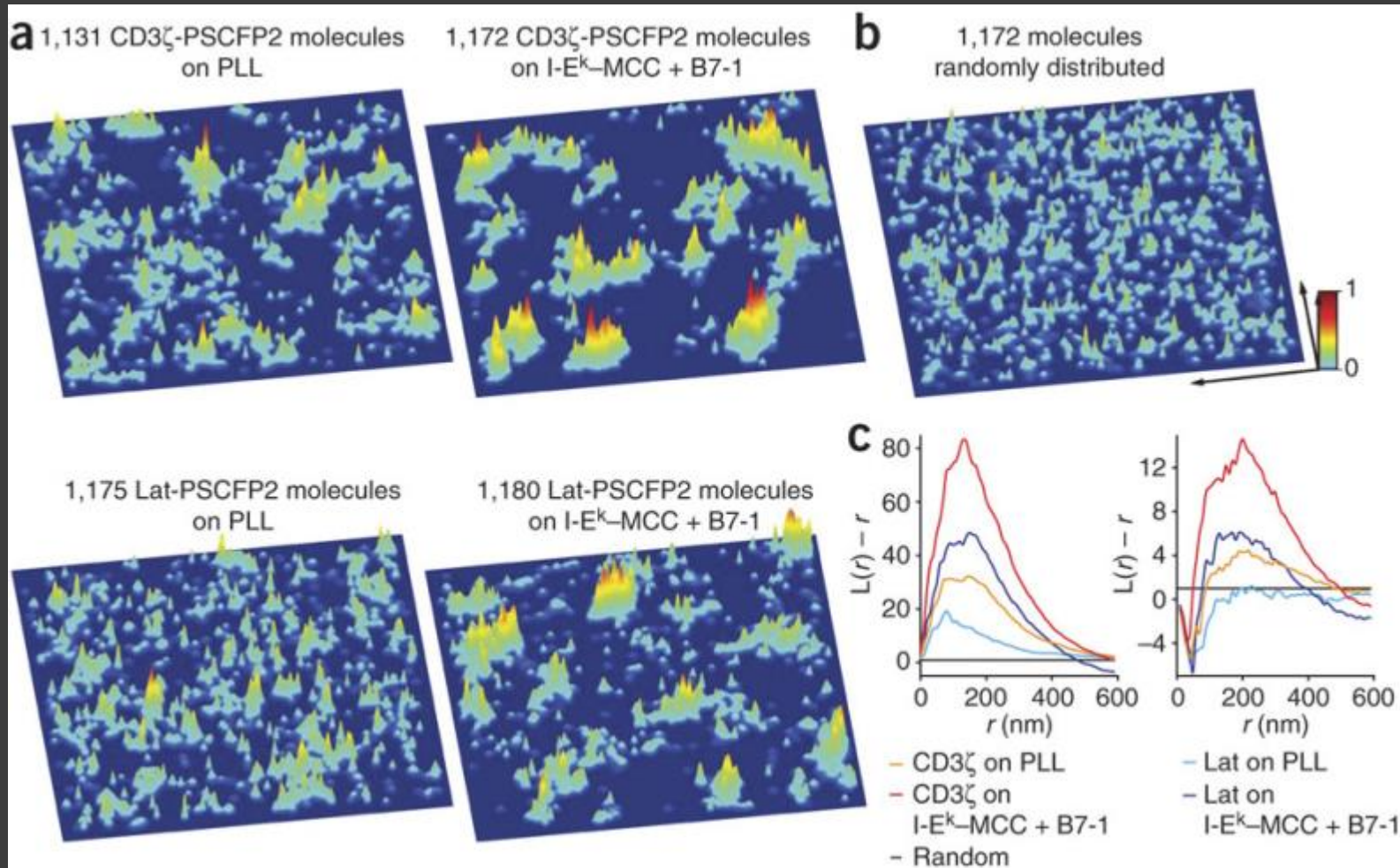
temporal: 500 ms

3D

spatial: xy: 30 nm, z: 50 nm

temporal: 1-2 s





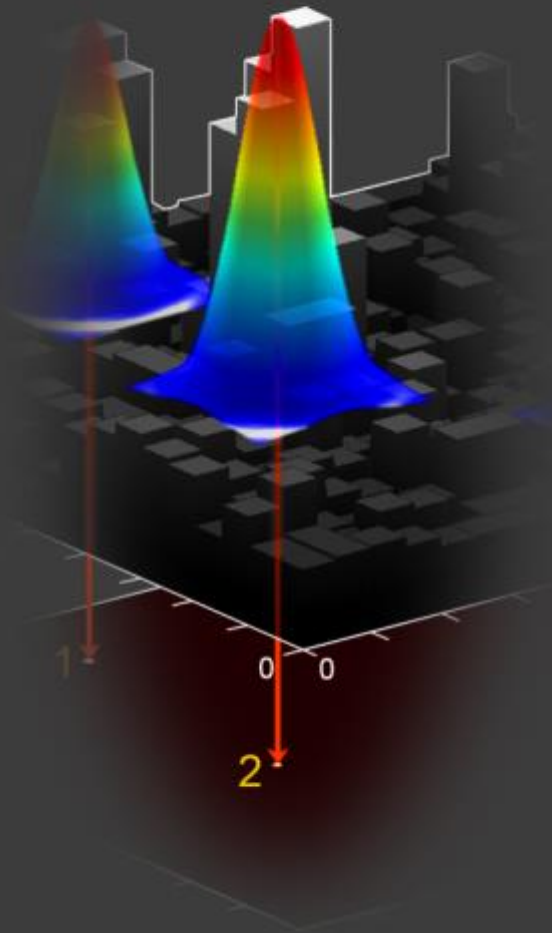
Lillemeier et al., Nature Immunology, 2009

2D, spatial resolution: 60 nm, temporal resolution: 4-10 s

how to get a lot more information from the data

# the additional single molecule information

remember?

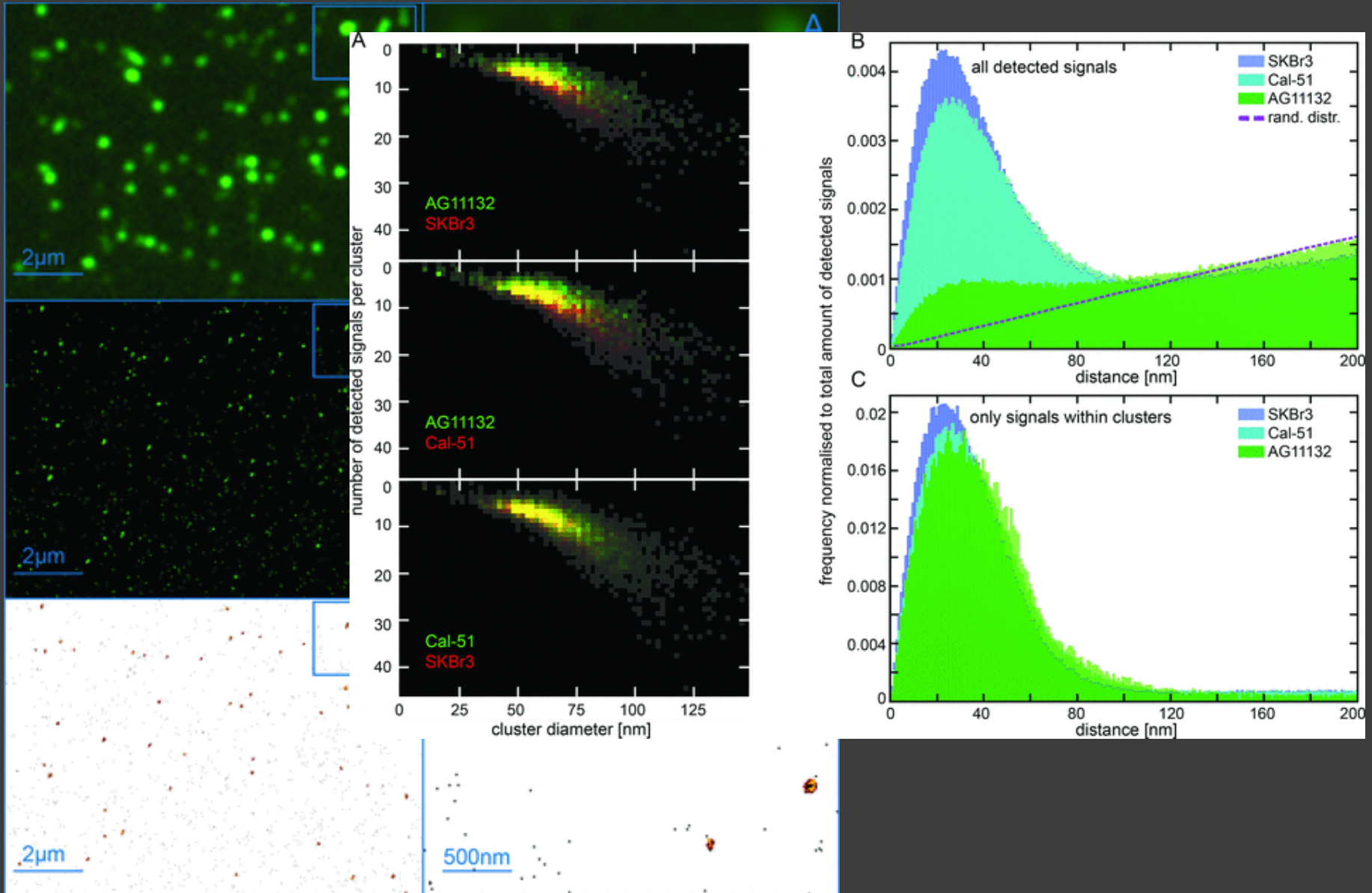


all the molecules in the image have been detected one by one

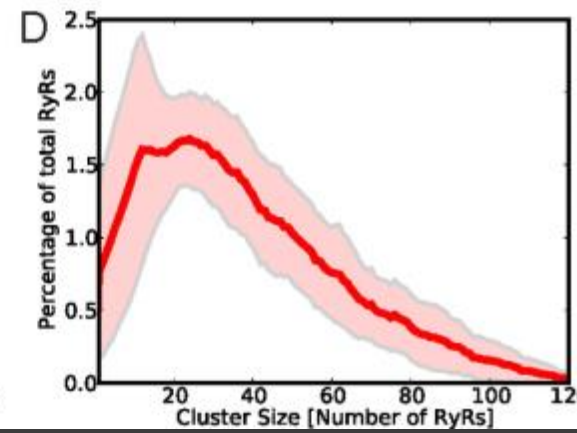
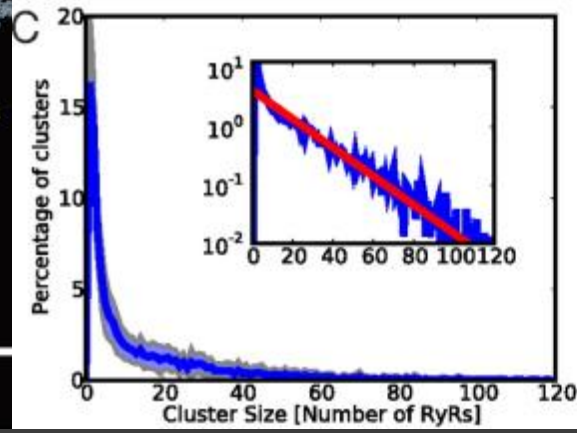
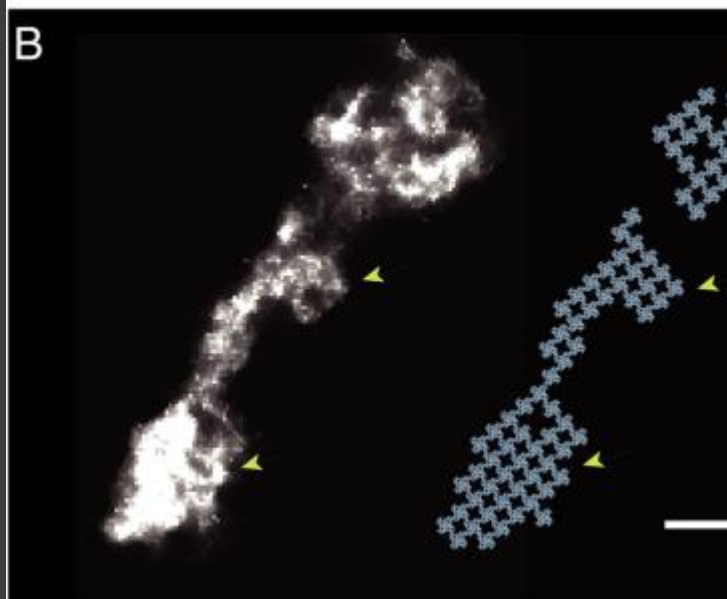
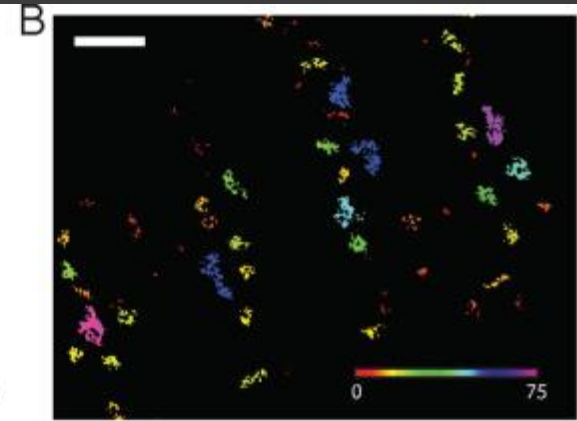
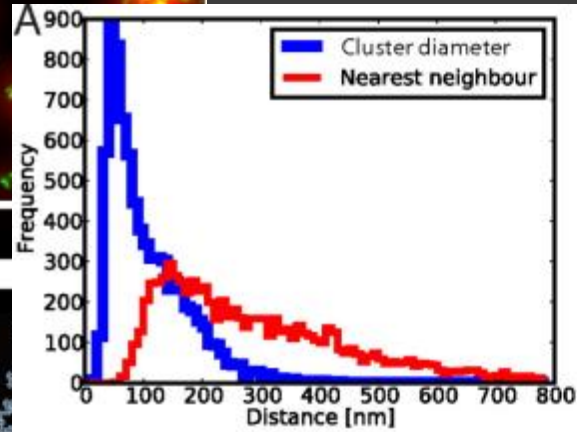
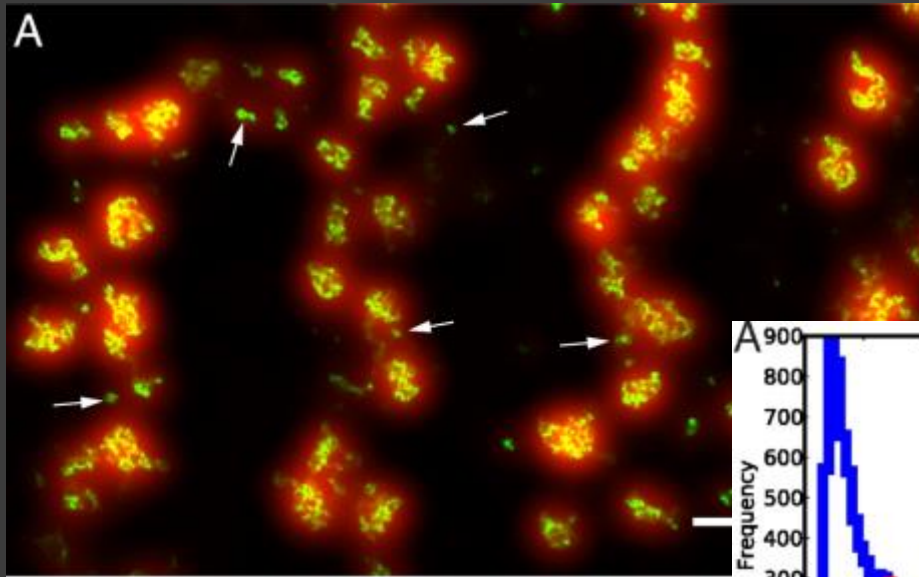
- position of each molecule
- number of detected photons
- shape of the PSF
- polarisation
- wavelength
- dynamics (in living cells)
- ...

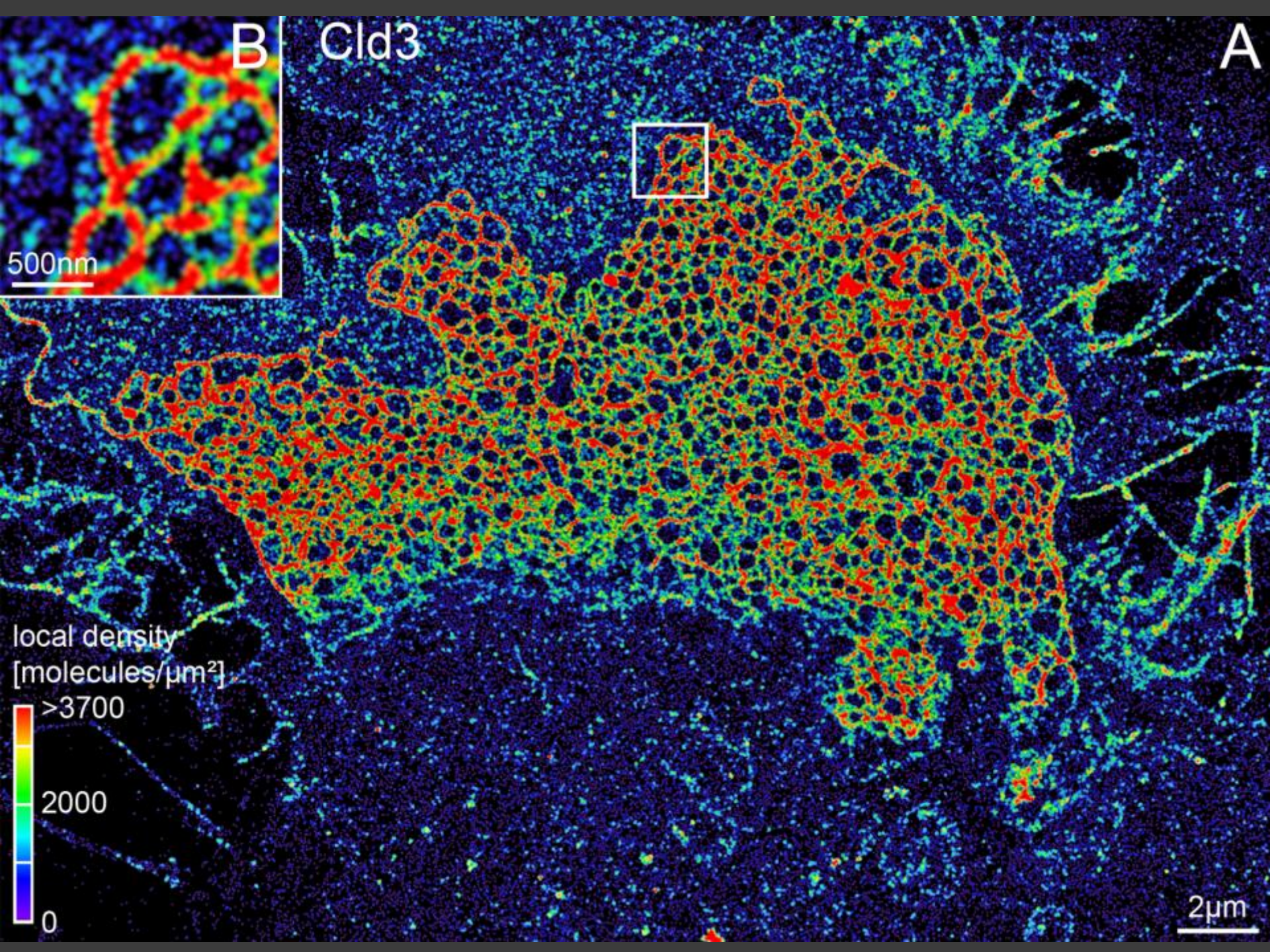


# statistical analysis of small protein clusters

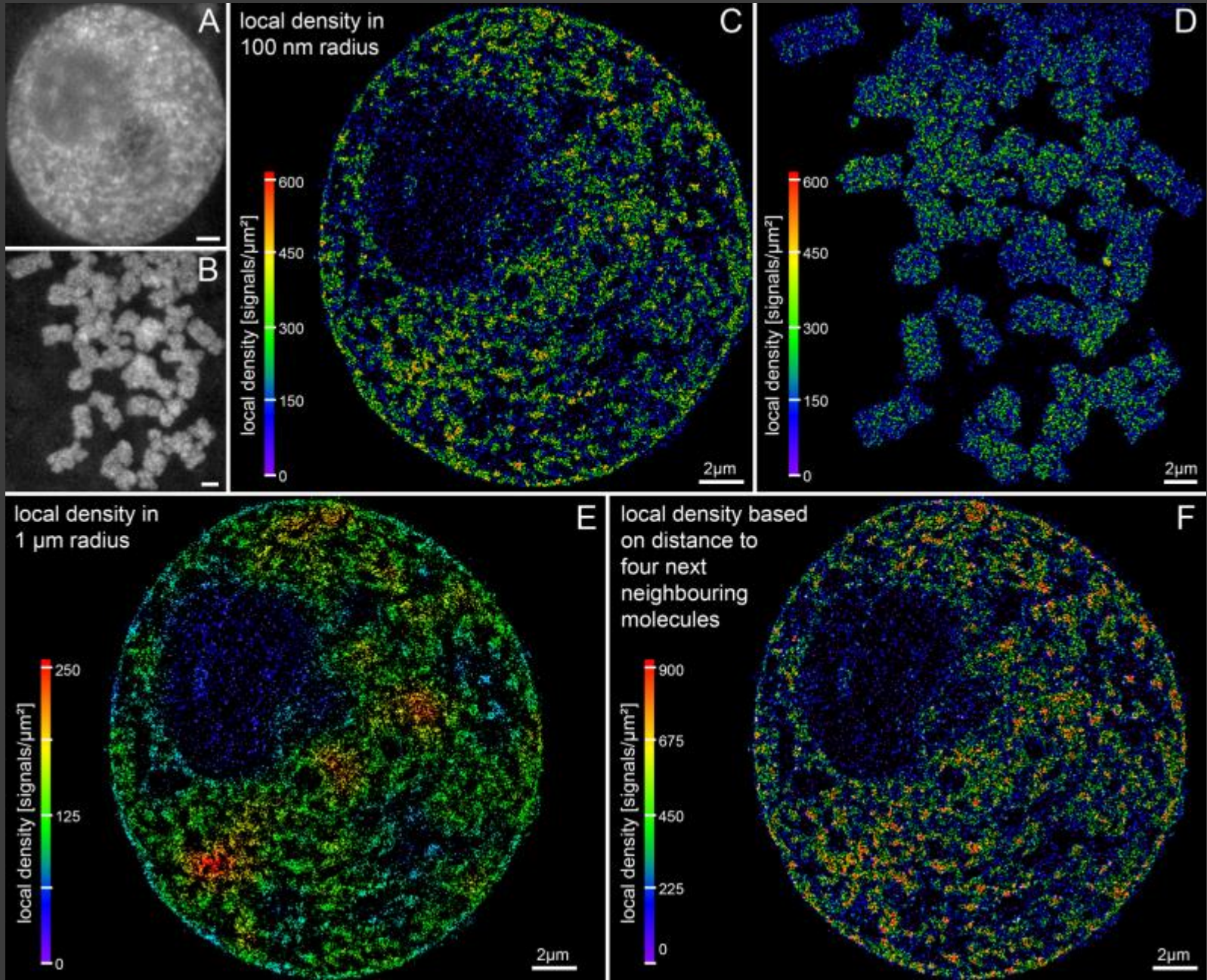


# statistical analysis of large protein clusters



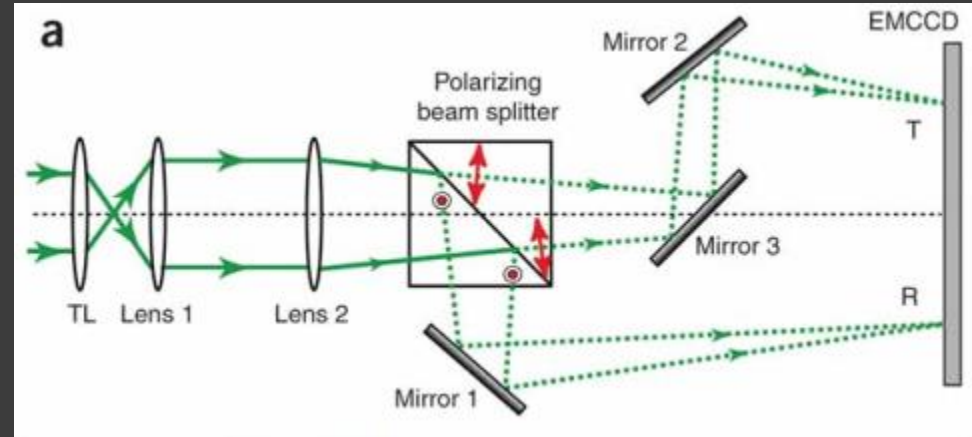
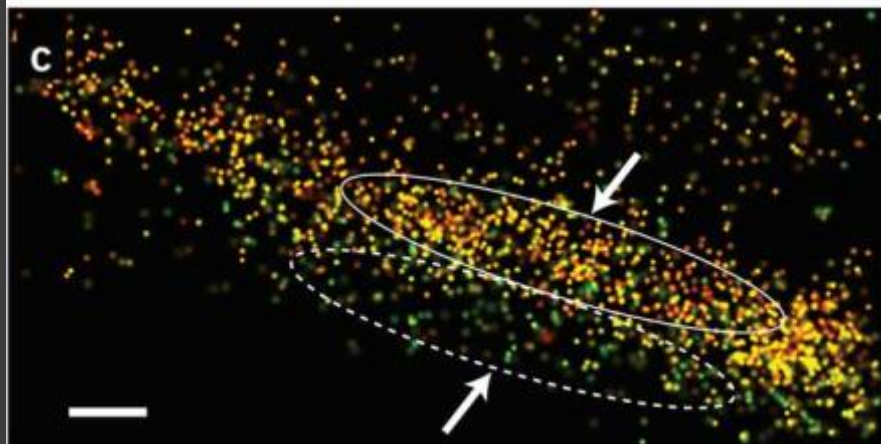
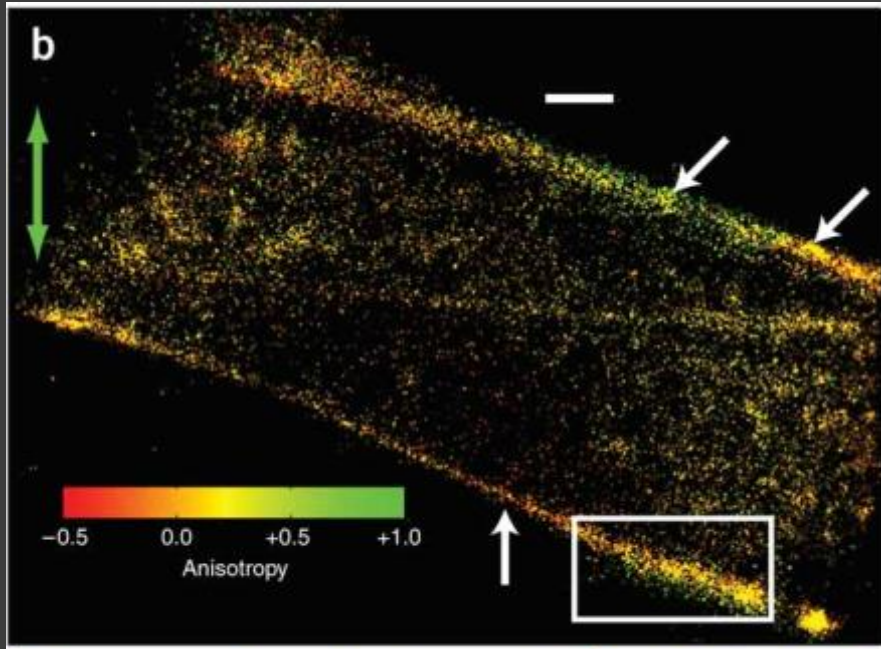


# visualisation of protein densities



# polarisation of the detected fluorophores

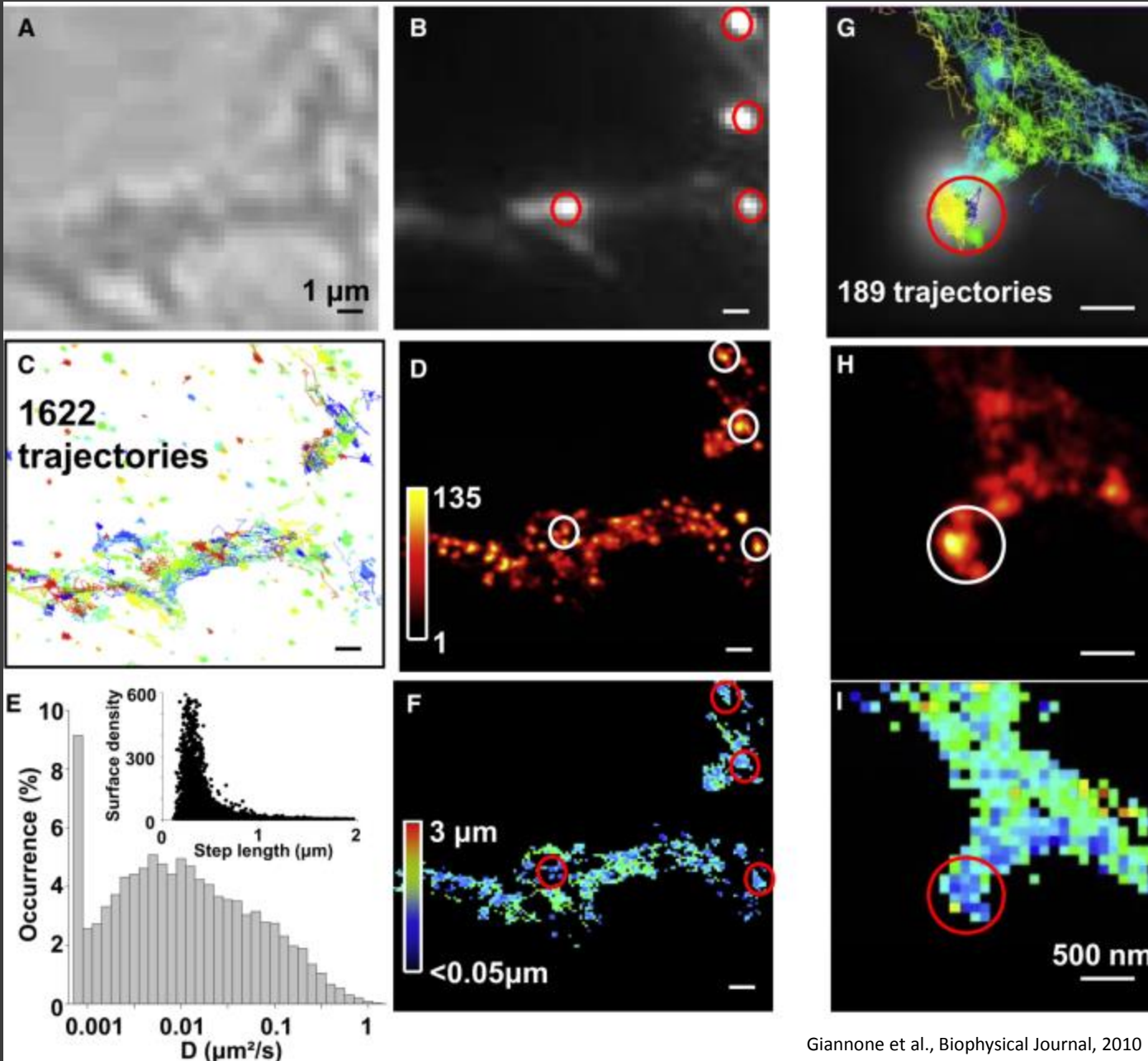
Dendra2-actin



Gould et al., Nature Methods, 2008

# high density particle tracking in living cells

sptPALM or uPAINT



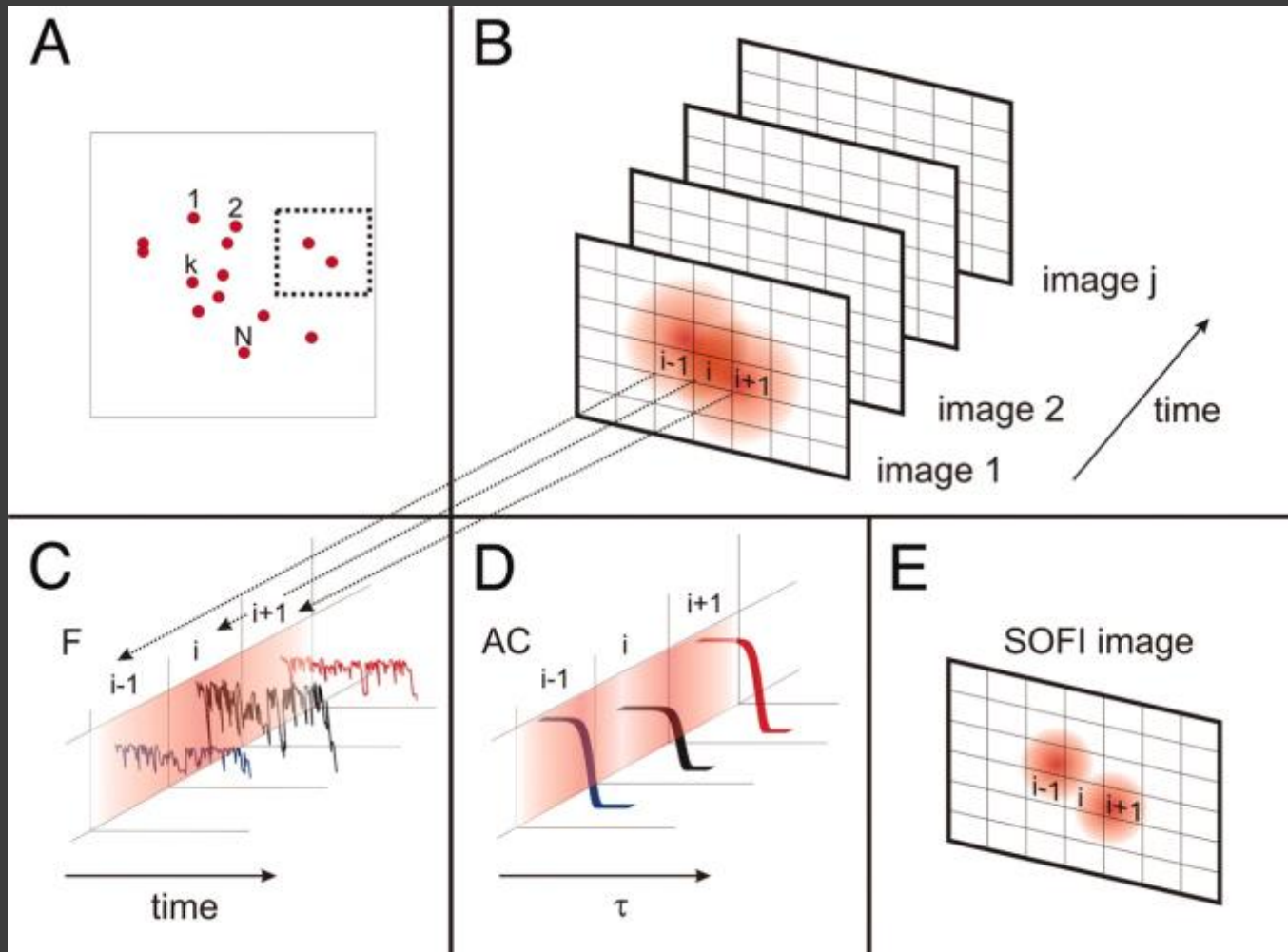
sptPALM:  
Manely et al., Nature  
Methods, 2008

antiGluR2-AT647N-AMPA

alternative approaches

# SOFI - making the setup even more simpler

localization microscopy using a lamp!



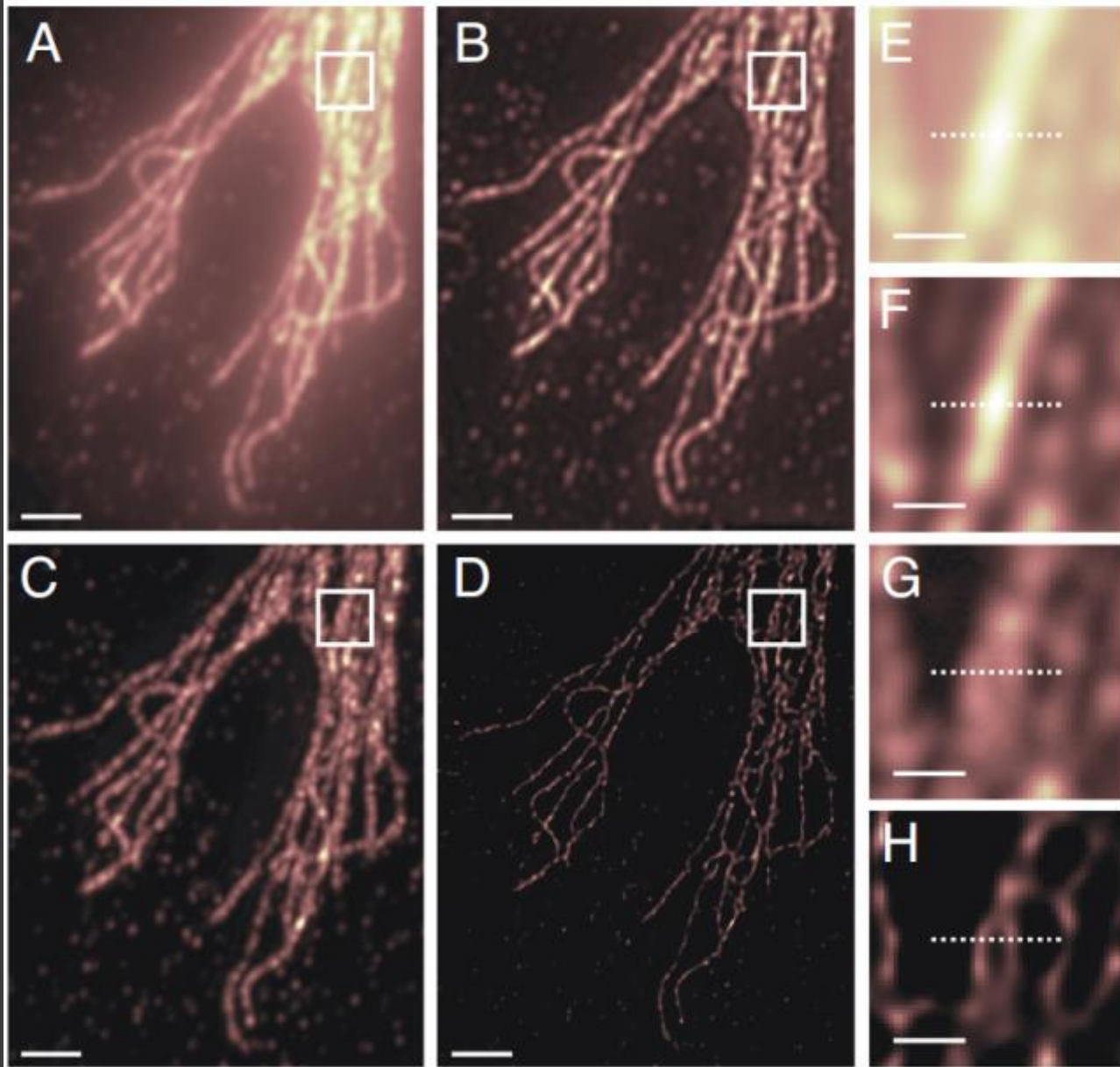
Dertinger et al., PNAS, 2009

$$\sum_{k=1}^N U(\mathbf{r} - \mathbf{r}_k) \cdot \boldsymbol{\varepsilon}_k \cdot s_k(t)$$

$$G_2(\mathbf{r}, \tau) = \sum_k U^2(\mathbf{r} - \mathbf{r}_k) \cdot \boldsymbol{\varepsilon}_k^2 \cdot \langle \delta s_k(t + \tau) s_k(t) \rangle$$



# SOFI - making the setup even more simpler



lateral resolution:  
70-100 nm

**BUT!**

no single molecule  
information

only resolution  
enhancement

# 3B analysis localization microscopy

similar approach as SOFI but some differences:

+ also based on very high molecule densities fluorescent in one frame

→ very fast: only several hundred frames needed for reconstruction of an image  
with a resolution of 50 nm → time resolution: 4 s

+ single molecule information is still accessible

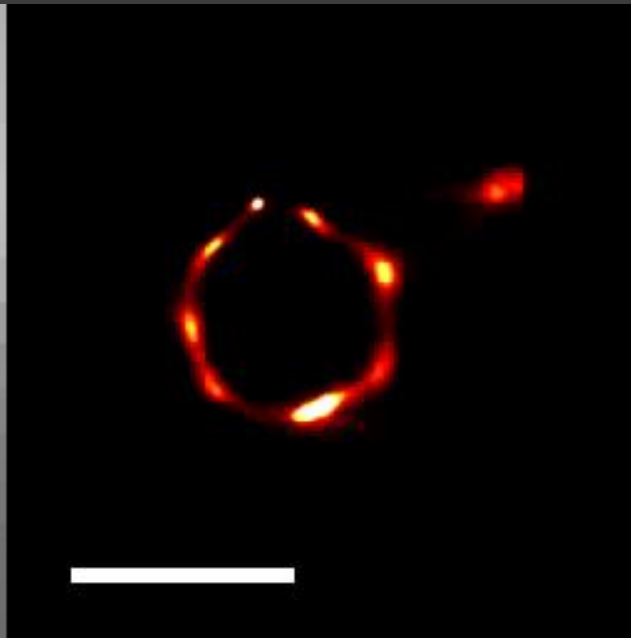
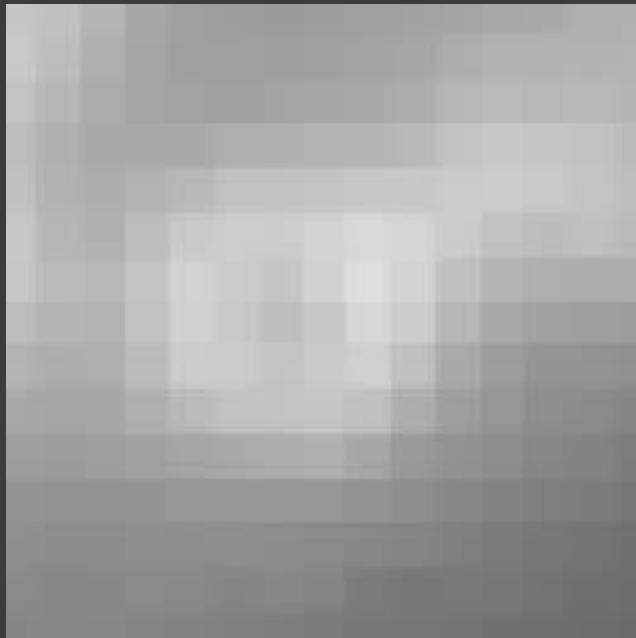
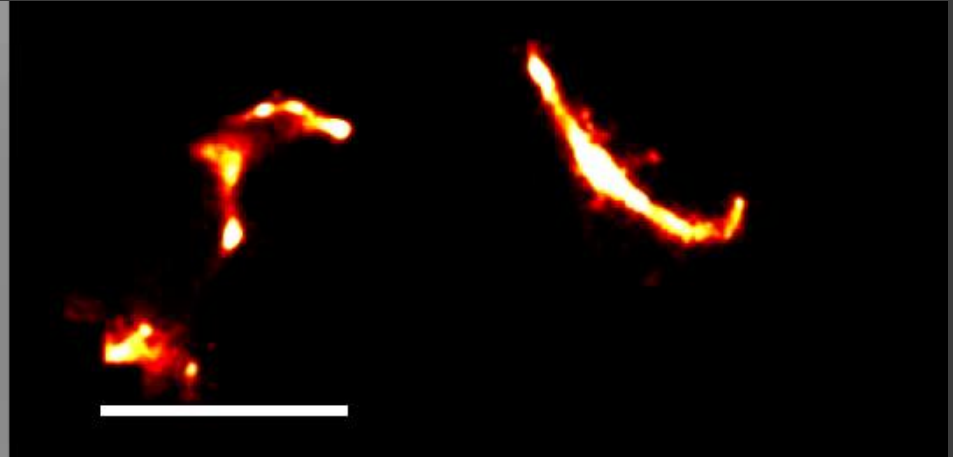
- extremely extensive computation effort

→ regions larger than  $2 \times 2 \mu\text{m}$  would need to be processed for days on a  
conventional (core i7) CPU

# 3B analysis localization microscopy

wide-field

reconstruction (resolution: 50 nm)



conclusion

PALM: irreversible photoactivation

- quantitative analyses, particle tracking, counting
- needs (in most cases) TIRF!

STORM, dSTORM, GSDIM, SPDM: reversible photoswitching

- resolution, fast
- also works without TIRF → imaging deep inside cells

SPDM with FPs: (almost) irreversible photoswitching

- quantitative analyses using conventional FPs
- also works without TIRF → imaging deep inside cells

# conclusion

resolution	quantitative and counting	particle tracking	speed	imaging in cells	use standard fluorophores	3D
(F)PALM	green	green	yellow	red	red	green
STORM	yellow	yellow	green	green	red	green
dSTORM	yellow	yellow	green	green	green	green
SPDM	green	green	yellow	green	green	green
GSDIM	yellow	yellow	green	green	green	green
SOFI	red	red	green	yellow	green	yellow
3B	yellow	red	green	yellow	green	red

referring to the original ideas of the methods

If you have a wide-field microscope with a laser for excitation of the fluorophores and one for switching/activating you can do ALL of these methods!

resolution	quantitative and counting	particle tracking	speed	imaging in cells	use standard fluorophores	3D
<h2>Localization Microscopy</h2>						
SOFI						
3B						

referring to the original ideas of the methods

If you have a wide-field microscope with a laser for excitation of the fluorophores and one for switching/activating you can do ALL of these methods!

original (F)PALM and STORM:

<http://www.sciencemag.org/content/313/5793/1642.short>

<http://www.nature.com/nmeth/journal/v3/n10/full/nmeth929.html>

<http://www.sciencedirect.com/science/article/pii/S0006349506721403>

dSTORM, SPDM and GSDIM (with standard fluorophores):

<http://onlinelibrary.wiley.com/doi/10.1002/anie.200802376/full>

<http://www.springerlink.com/content/vx05p35kr3424228/>

<http://www.nature.com/nmeth/journal/v5/n11/full/nmeth.1257.html>

3D:

[http://apl.aip.org/resource/1/applab/v97/i16/p161103\\_s1?view=fulltext](http://apl.aip.org/resource/1/applab/v97/i16/p161103_s1?view=fulltext)

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<http://www.super-resolution.biozentrum.uni-wuerzburg.de/home/rapidstorm/>

<http://code.google.com/p/quickpalm/>

the end