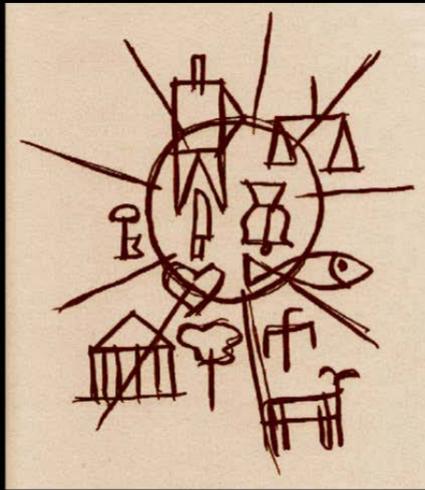


# Welcome





## Advanced Course

# Procesamiento de Imágenes y Bioseñales I & II

Primavera, 2012

Escuela de Postgrado, Facultad de Medicina, U-Chile



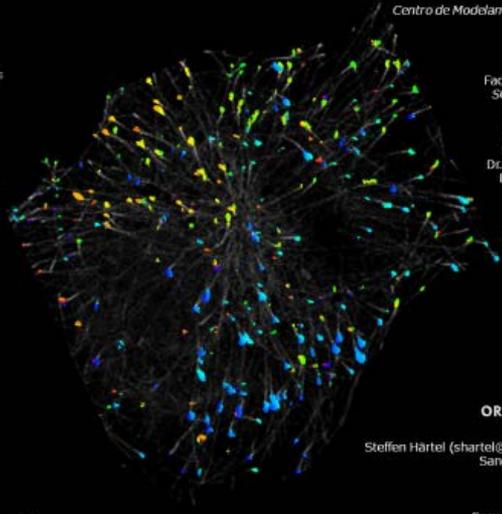
### OVERVIEW

#### Part I

- Biological and biomedical imaging acquisition
- Methods and techniques of signal and image processing
- Analysis of biological structures in digital images

#### Part II

- Interpretation of morphological, topological and dynamical information in biological and biomedical images
- High throughput microscopy
- Super-resolution microscopy



### ACTIVITIES

- Lectures
- Bibliographic Seminars
- Hands-on Microscopy
- Hands-on Computational Analysis

### PROFESSORS

Universidad de Chile  
 FCFM  
 Centro de Modelamiento Matemático  
 Takeshi Asahi

Facultad de Medicina  
 SCIAN-Lab / CEDAI

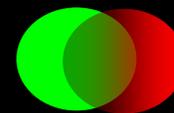
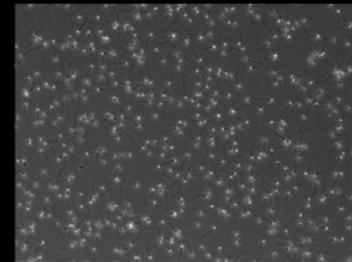
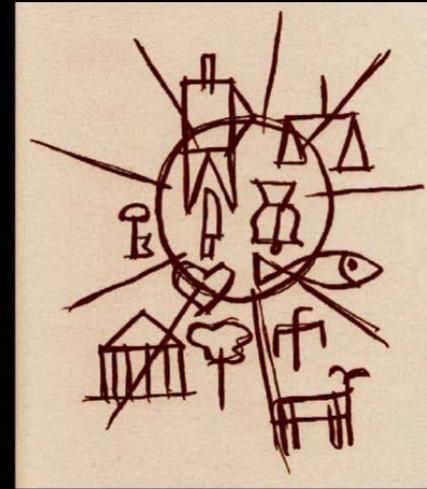
Dr. Steffen Härtel  
 Dr. Enzo Brunetti  
 Dr. Omar Ramírez  
 Dr. Victor Castañeda  
 Dr. Mauricio Cerda  
 Dr. Jarno Ralli  
 Jorge Jara  
 Luis Briones  
 Felipe Santibañez

### ORGANIZATION

Steffen Härtel (shartel@med.uchile.cl)  
 Sandra de la Fuente  
 Pamela Weber

Register @  
 Escuela de Postgrado  
 Facultad de Medicina  
 Universidad de Chile

More Information @  
[www.scian.cl/mim](http://www.scian.cl/mim)





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2012 Curso PIB

> MSc Informatica Medica

Programa

> U-Chile

- 2006 Curso 3D-Microscopy I
- 2007 Curso 3D-Microscopy II
- 2007 Curso Colocalization
- 2007 Curso Microscopía
- 2008 Curso Fotografía
- 2008 Curso Colocalization
- 2008 Curso Microscopía
- 2009 Curso Colocalization
- 2011 Curso MMIMB
- 2011 Curso Bioinformatica
- 2012 Curso PIB
- 2012 Curso Bioinformática
- 2012 Curso Seminario MIM

> U-Chile / International

2005 Microscopy Uruguay

Procesamiento de Imágenes y Bioseñales I & II

| Santiago de Chile 18 de Agosto 2012 - 12 de Diciembre 2012, Facultad de Medicina, U-Chile |



## Advanced Course

# Procesamiento de Imágenes y Bioseñales I & II

### Primavera, 2012

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**OVERVIEW**

**Part I**

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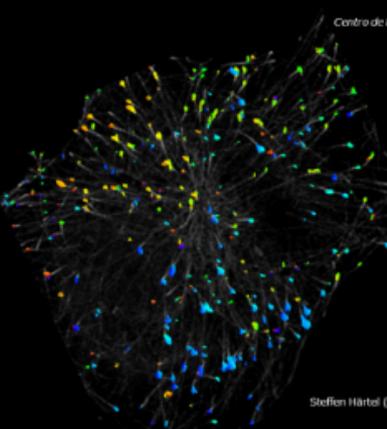
**PROFESSORS**

Universidad de Chile  
FCFM  
Centro de Modelamiento Matemático

Takeshi Asahi

Facultad de Medicina  
SCIAN-Lab / CCDAI

Dr. Steffen Härtel  
Dr. Enzo Brunetti  
Dr. Omar Ramirez  
Dr. Victor Castañeda  
Dr. Mauricio Cerda  
Dr. Jairo Ralli  
Jorge Jara  
Luis Brenner  
Felipe Santibañez



**ACTIVITIES**

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- Bibliographic Seminars
- Hands-on Microscopy
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Steffen Härtel (shartel@med.uchile.cl)  
Sandra de la Fuente  
Pamela Weber

Register @  
Escuela de Postgrado  
Facultad de Medicina  
Universidad de Chile

More information @  
[www.scian.cl/mim](http://www.scian.cl/mim)

Informaciones en:

-> [www.scian.cl](http://www.scian.cl)

-> [www.scian.cl/mim](http://www.scian.cl/mim)

Notas:

Prácticos (25%)

Seminarios (25%)

Examen Final (50%)

5 grupos a 3:

- Seminarios Prácticos

20 min presentación y pyd

- Seminarios Bibliográficos

20 min presentación +

20 min preguntas y discusión

### PROFESORES PARTICIPANTES (INDICAR UNIDADES ACADÉMICAS)

#### **ICBM | Facultad de Medicina, U-Chile y BNI**

Dr. Steffen Härtel, Director SCIAN-Lab, Programa de Anatomía y Biología del Desarrollo (PABD), Instituto de Neurociencia Biomédica (BNI), Instituto de Ciencias Biomédicas (ICBM), [shartel@med.uchile.cl](mailto:shartel@med.uchile.cl)

Dr. Enzo Brunetti, Laboratorio Neuro-sistemas, [enzo@neuro.med.uchile.cl](mailto:enzo@neuro.med.uchile.cl)

Dr. Omar Ramírez, SCIAN-Lab, PABD, [ojanor@gmail.com](mailto:ojanor@gmail.com)

Dr. Víctor Castañeda, SCIAN-Lab, PABD, [vcastane@gmail.com](mailto:vcastane@gmail.com)

Dr. Mauricio Cerda, SCIAN-Lab, PABD, [mcerda@med.uchile.cl](mailto:mcerda@med.uchile.cl)

Dr. Jarno Ralli, SCIAN-Lab, PABD, [jarno@ralli.fi](mailto:jarno@ralli.fi)

Dr.(c) Jorge Jara, SCIAN-Lab, PABD, [jjaraw@gmail.com](mailto:jjaraw@gmail.com)

Ing. Luis Briones, SCIAN-Lab, PABD, [lbriones@med.uchile.cl](mailto:lbriones@med.uchile.cl)

Ing. Felipe Santibáñez, SCIAN-Lab, PABD, [fsantibanez@med.uchile.cl](mailto:fsantibanez@med.uchile.cl)

#### **Centro de Modelamiento Matemático | Facultad de Ciencias Físicas y Matemáticas, U-Chile**

Dr. Takeshi Asahi, Laboratorio de Modelamiento en Imágenes Científicas y Visualización (MOTIV) y Centro de Modelamiento Matemático (CMM), [tasahi@dim.uchile.cl](mailto:tasahi@dim.uchile.cl)



Basic Science

R&D

Human Capital Formation  
Medical Informatics

Basic Science

FONDECYT  
CONICYT

FONDEF  
CONICYT

DAAD / DFG

ICM

50 M\$Ch  
100 kUS\$ / year

110 M\$Ch  
220 kUS\$ / year

60 M\$Ch  
120 kUS\$ / year

60 M\$Ch  
120 kUS\$ / year

PI  
PostDocs  
PhD - students  
Master - students  
Undergraduate  
Research – Assistants  
Technicians

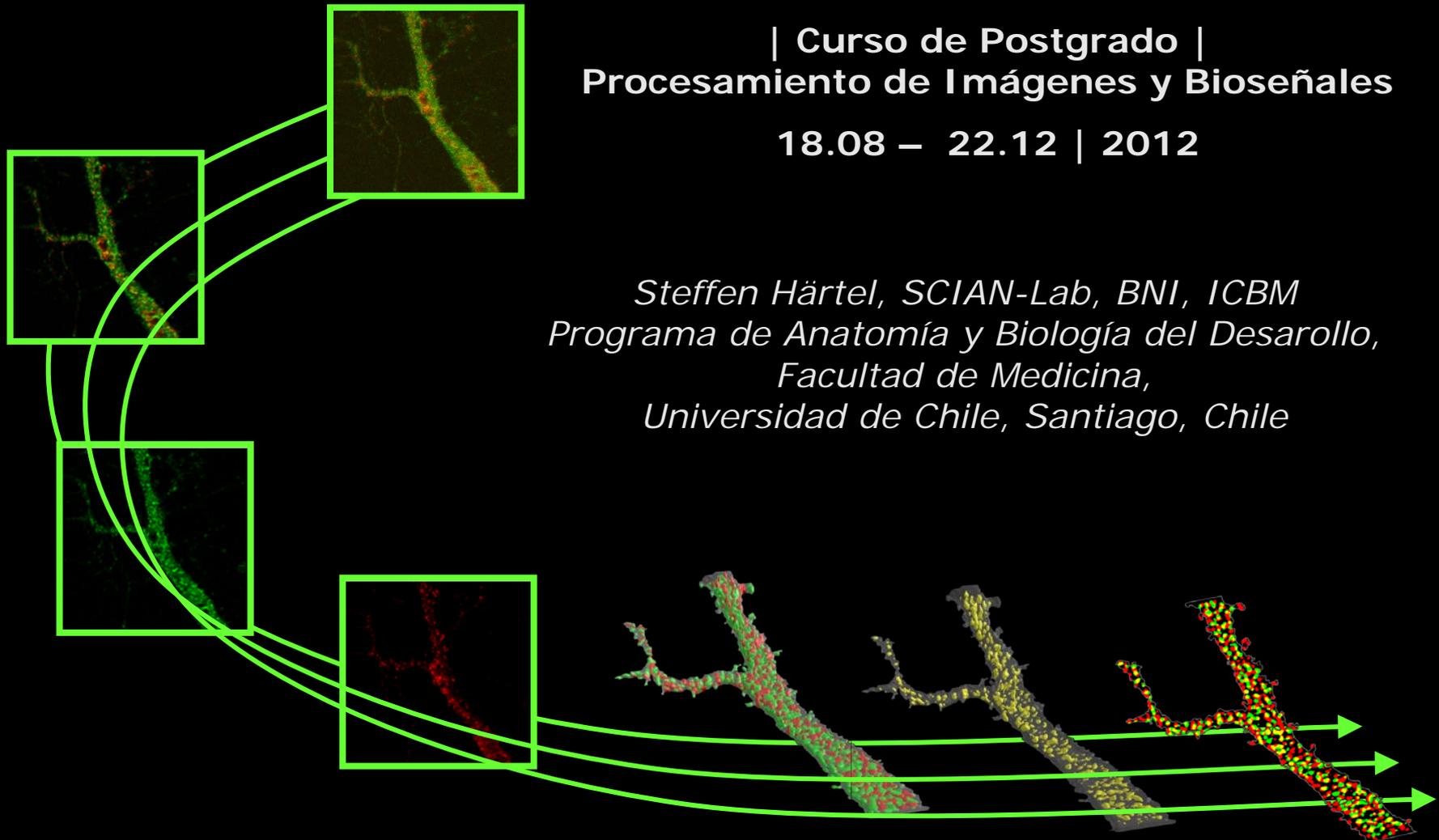
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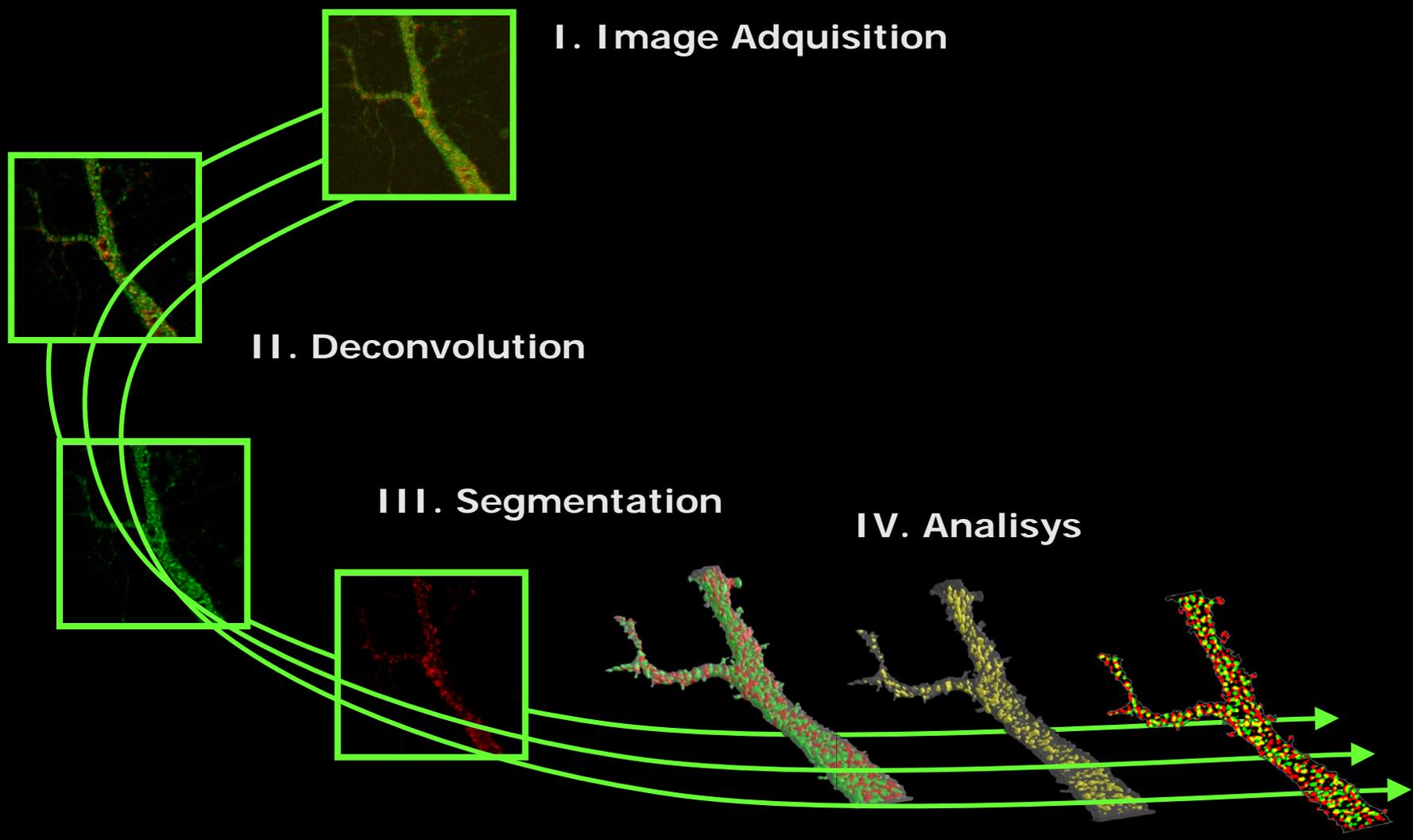
Biophysics  
Biology / Computer Sc / Electrical Engineer  
Computer Sc / Electrical Engineer  
Medical Technology / Electrical Engineer  
Computer Sc  
Medicine / Computer Sc / Electrical Engineer / Biology  
Biotechnology / Labtechnician / Administration

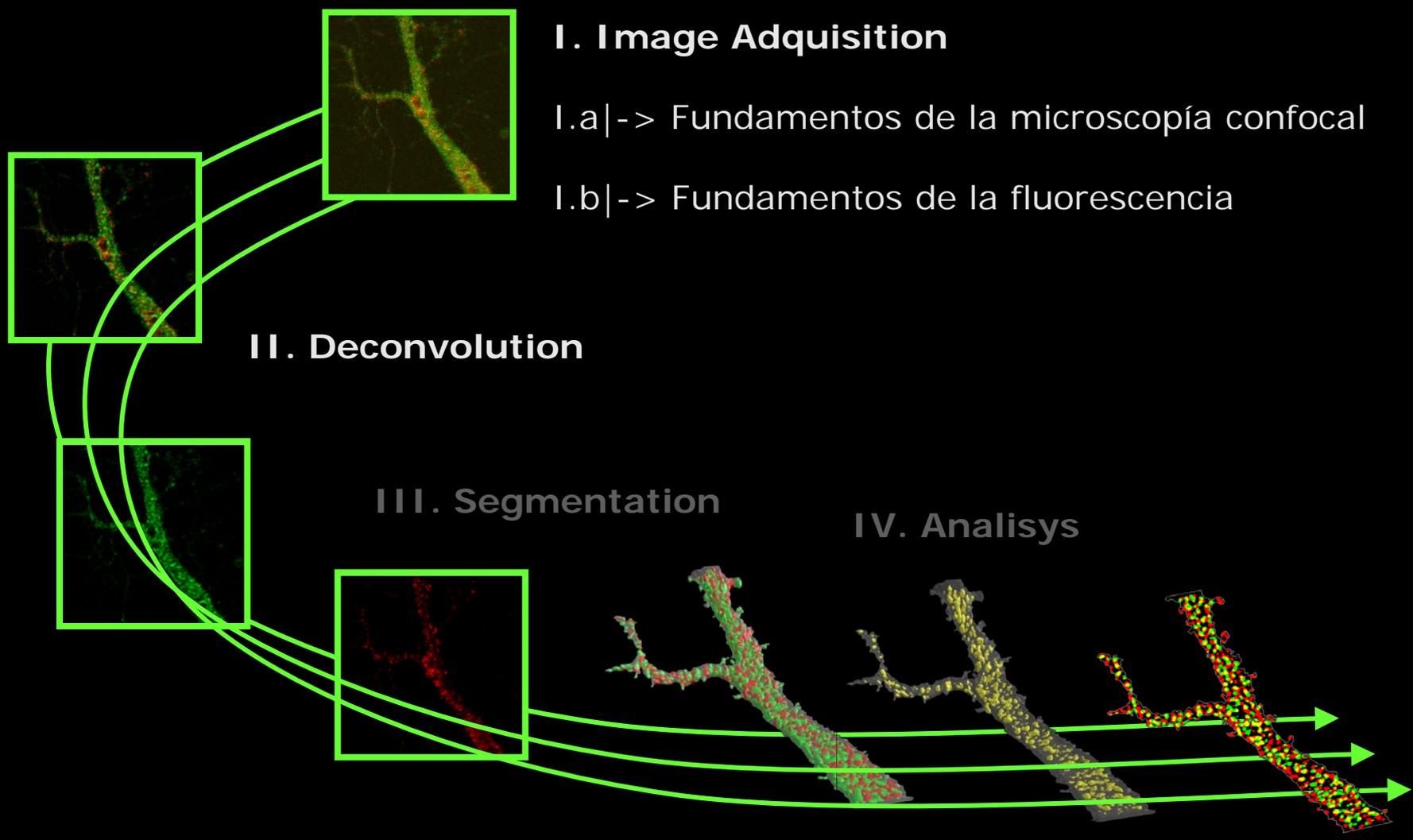


| **Curso de Postgrado** |  
**Procesamiento de Imágenes y Bioseñales**  
**18.08 – 22.12 | 2012**

*Steffen Härtel, SCIAN-Lab, BNI, ICBM  
Programa de Anatomía y Biología del Desarrollo,  
Facultad de Medicina,  
Universidad de Chile, Santiago, Chile*









**“It is very easy to answer many of these fundamental biological questions. You just look at the thing !**

**Make microscopes a hundred times more powerful and many problems of biology would be made very much easier.”**

**Richard Feynman (1918-1988)**

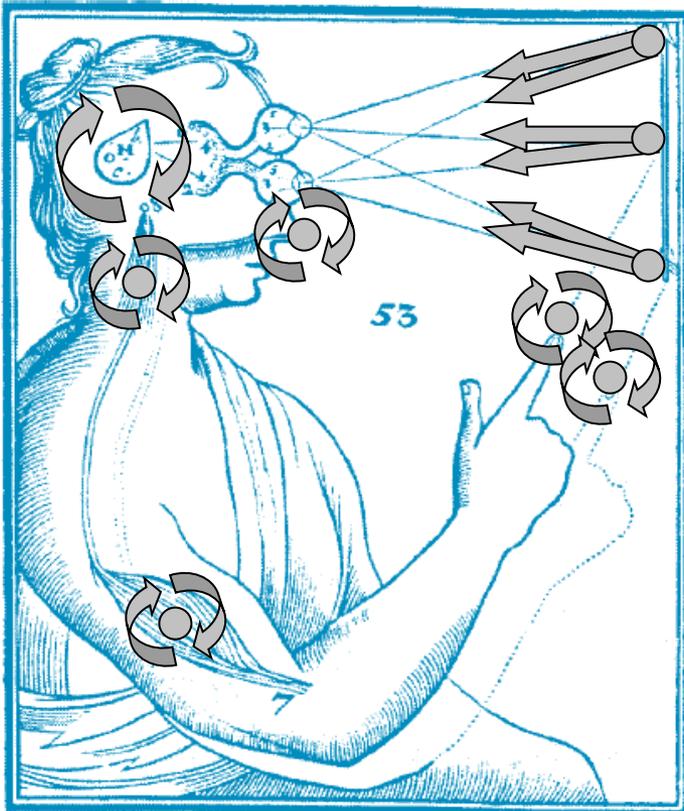


**René Descartes (1596-1650)**

**... just look at the thing ...  
¿ Human visual perception ?**

***Treatise of man (~ 1637)***

***Passions of the soul (~ 1649)***



*glandula pinealis* / pineal organ

A combination of ...

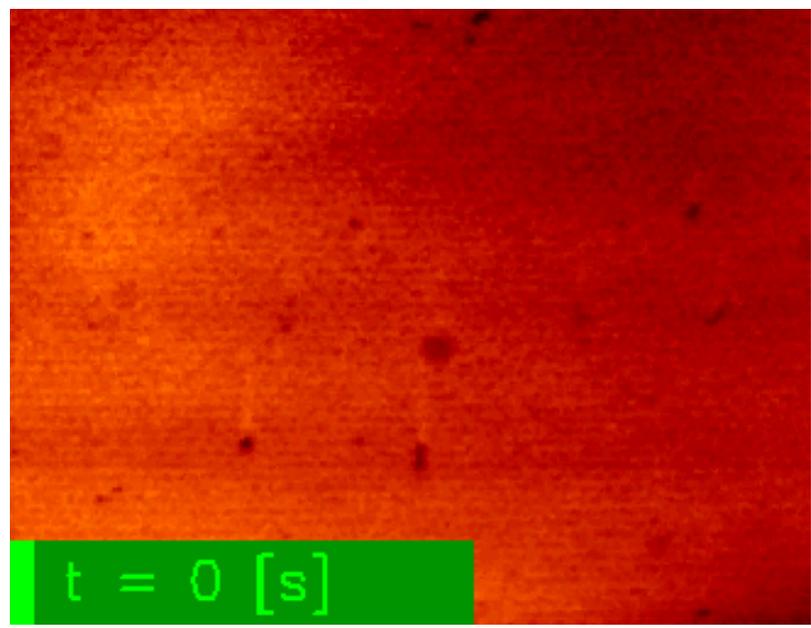
1 | direct signals ...

2 | signals from other senses ...

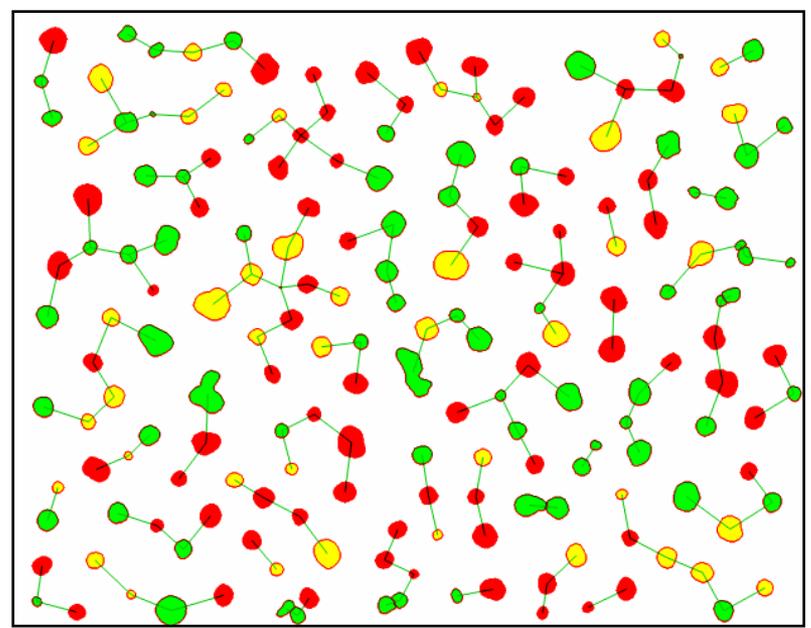
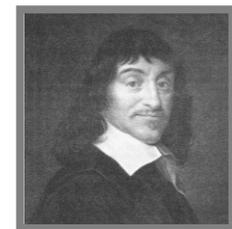
3 | feedback loops ...

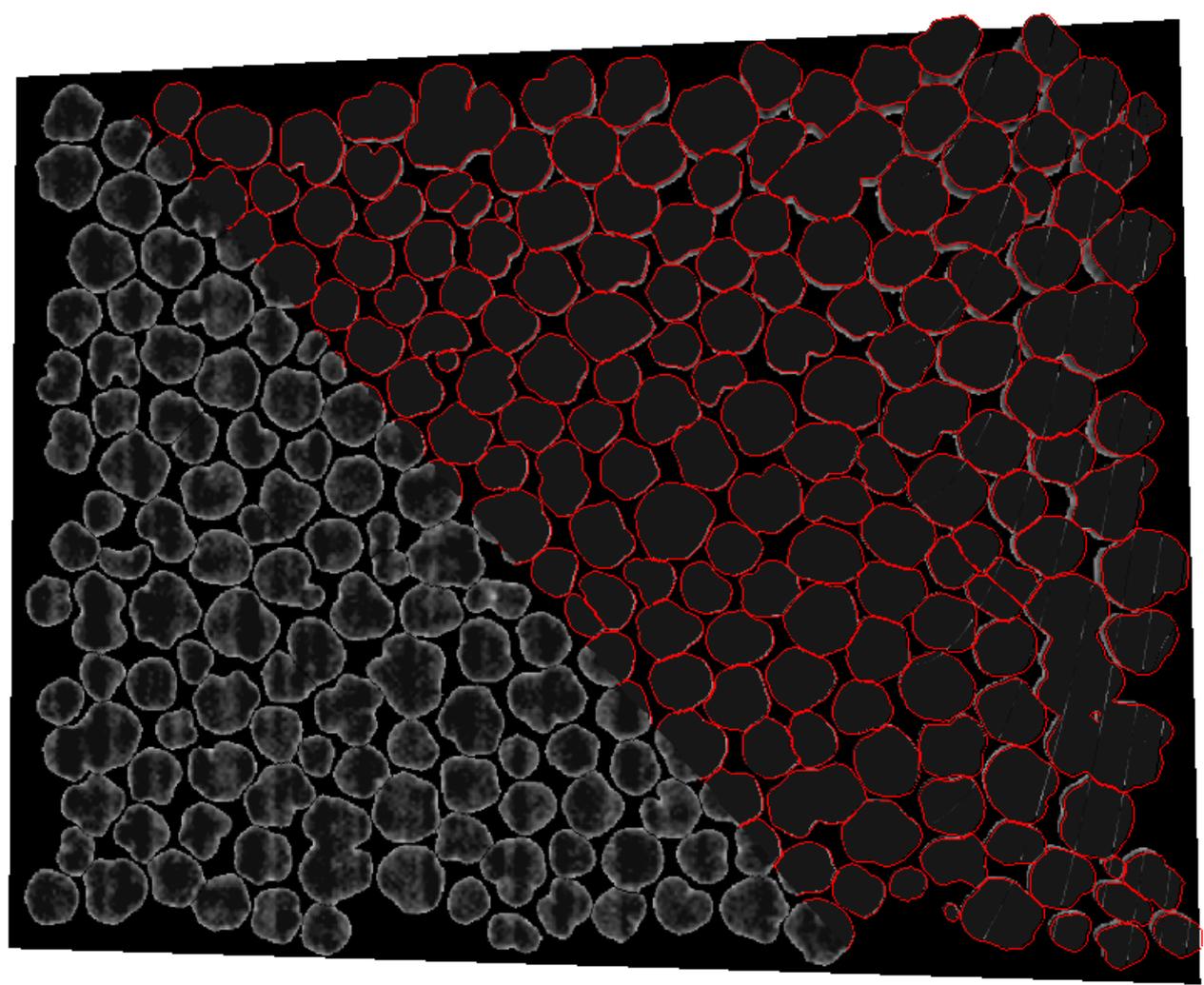
... produce a symbolic  
representation of an object.

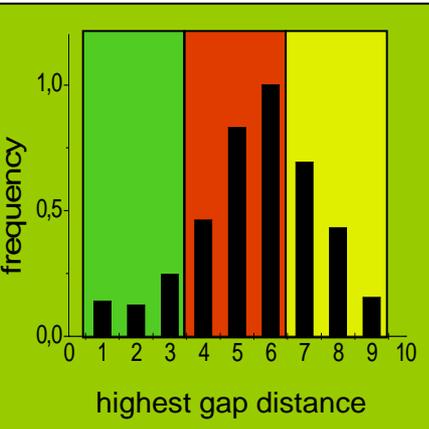
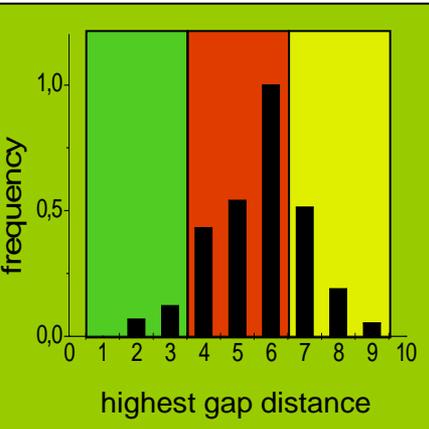
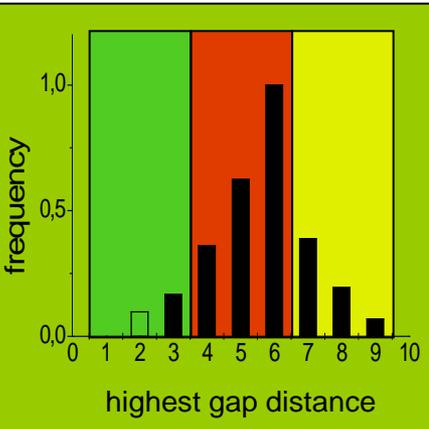
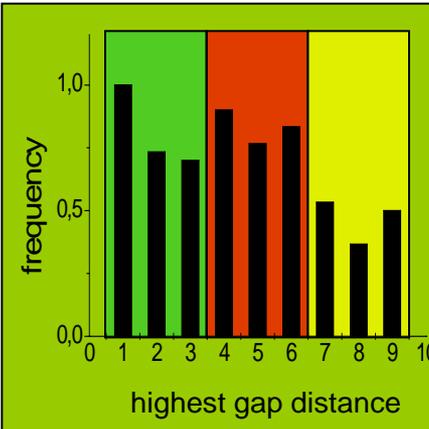
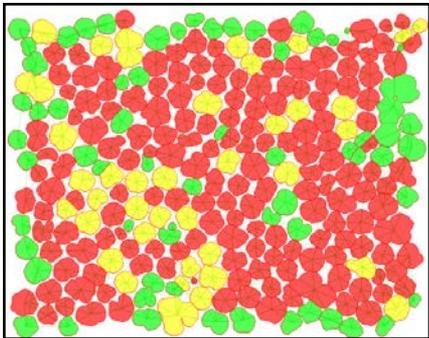
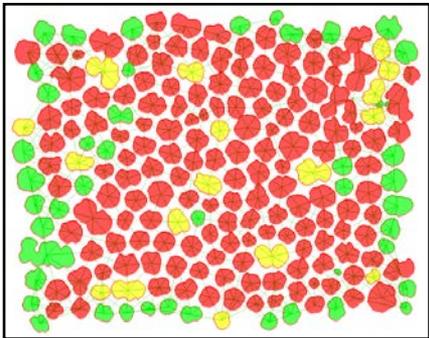
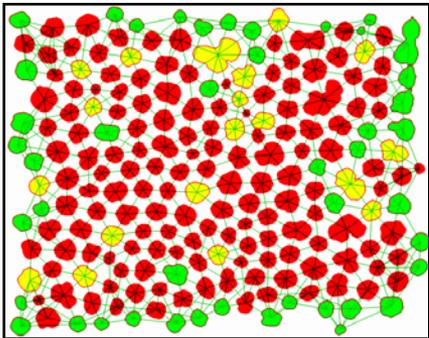
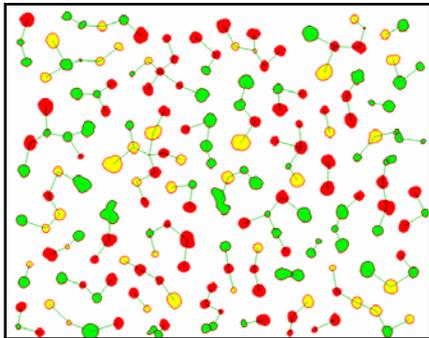
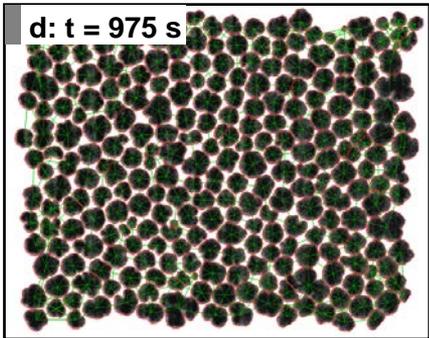
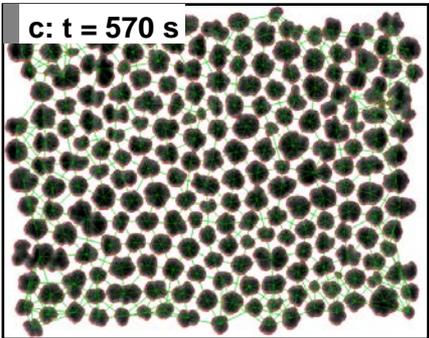
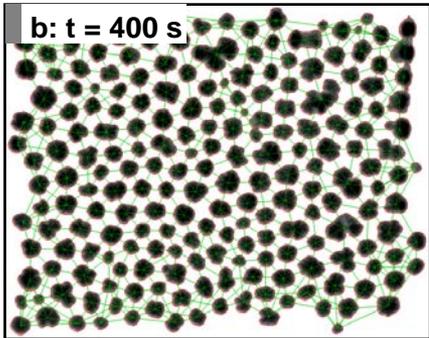
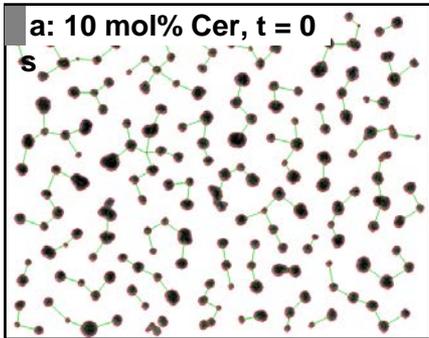
| Best resolution in  $t$  &  $x$  ...



|| Humas vs machine vision

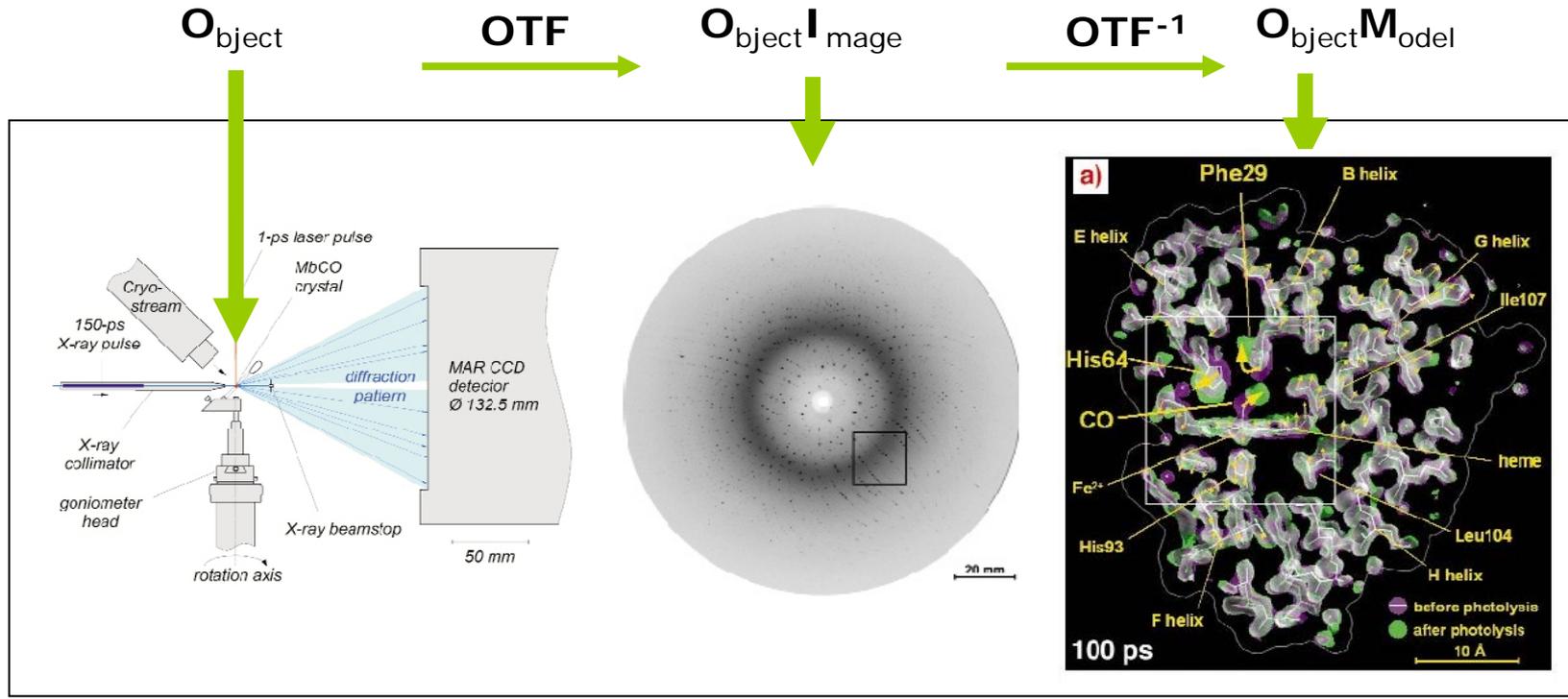






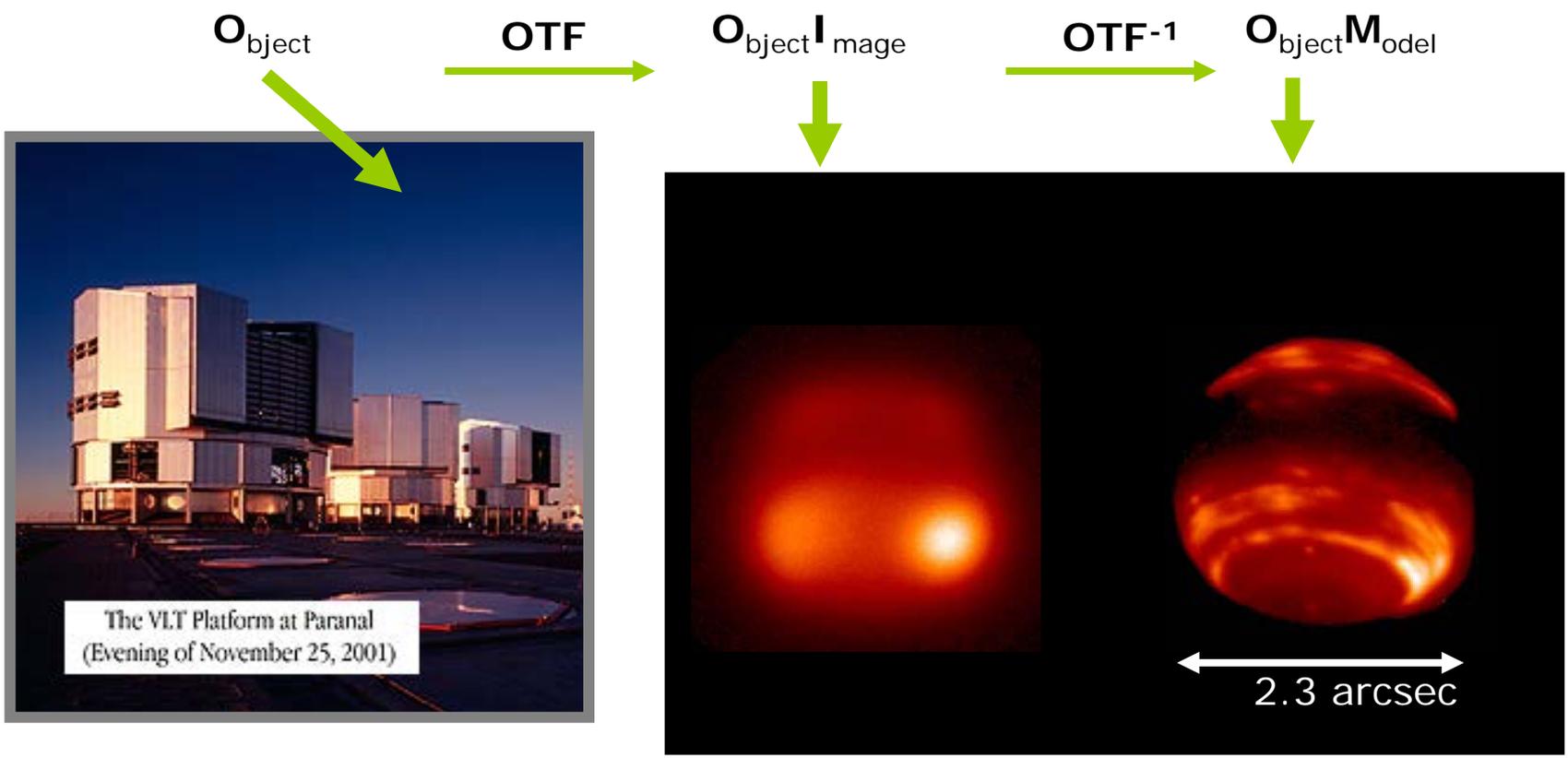


OTF: Object/Optical Transfer Function  
 Myoglobin in Action | Picosecond Laue Crystallography Diffraction Data  
 Schotte et al (2003) Science



<http://www.youtube.com/watch?v=InKIBZYarzM>

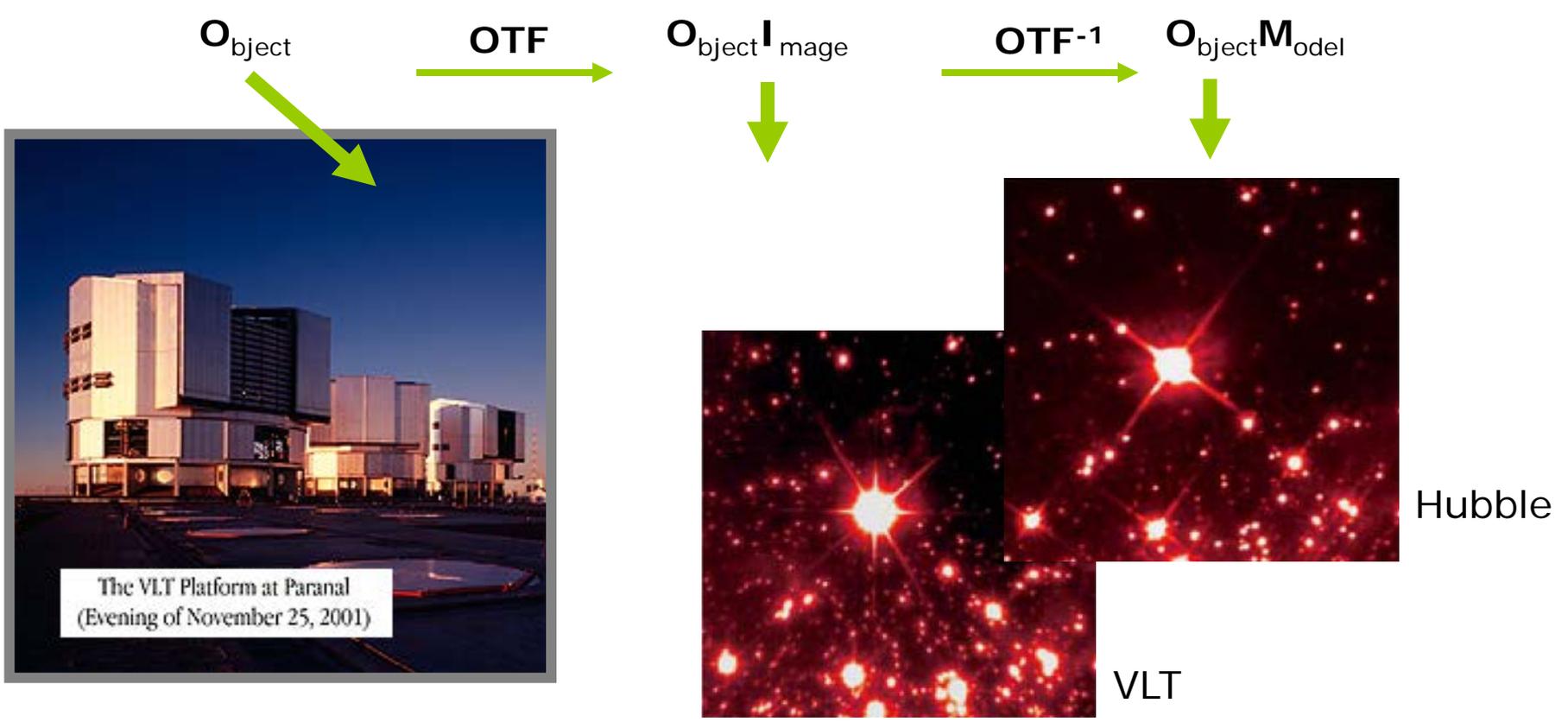
Diffraction Limited Resolution for a 10m telescope  $\sim \lambda/D \sim 0.01$  arcsec is limited to  $\sim 0.5$  arcsec by the turbulent atmosphere.



The VLT Platform at Paranal (Evening of November 25, 2001)

2.3 arcsec

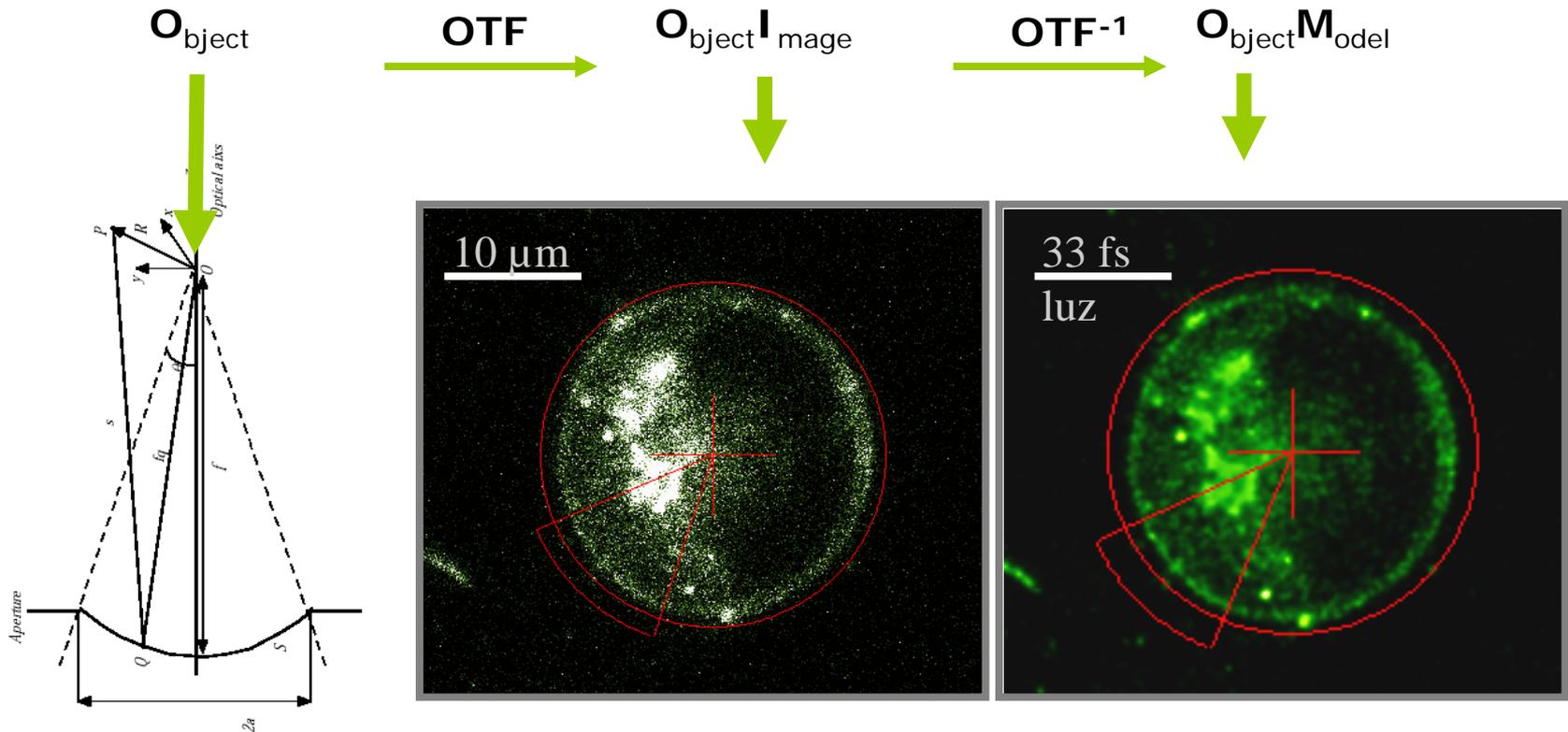
Diffraction Limited Resolution for a 10m telescope  $\sim \lambda/D \sim 0.01$  arcsec is limited to  $\sim 0.5$  arcsec by the turbulent atmosphere.



The VLT Platform at Paranal (Evening of November 25, 2001)

# Confocal Microscopy | From Geometric Optics to Diffraction Theory

Diffraction: The deviation of an electromagnetic wavefront from the path predicted by geometric optics when the wavefront interacts with a physical object such as an opening or an edge.





Seeing is believing? Alison J. North, *The Journal of Cell Biology*, Vol. 172, No. 1, January 2, 2006 9–18

The objective lens is the most critical component of a microscope and yet few researchers grasp the differences between specific objective classes.

Keep the acquisition settings constant between specimens to be compared quantitatively and particularly between sample and control.

“When you employ the microscope, shake off all prejudice, nor harbour any favorite opinions; for, if you do, ‘tis not unlikely fancy will betray you into error, and make you see what you wish to see.” Henry Baker, chapter 15, “Cautions in viewing objects” of *The Microscope Made Easy, 1742*.

“Remember that truth alone is the matter that you are in search after; and if you have been mistaken, let not vanity seduce you to persist in your mistake.” Henry Baker, *The Microscope Made Easy, 1742*.

## Seeing is believing? A beginners' guide to practical pitfalls in image acquisition

Imaging can be thought of as the most fun part of experiments. You see something; you report what you see. If only things were truly this simple. Modern imaging technology has brought about a revolution in the kinds of questions we can approach, but this comes at the price of increasingly complex equipment. Moreover, in an attempt to market competing systems, the microscopes have often been inappropriately described as easy to use and suitable for newcomers. Inadequate understanding of the experimental manipulations and equipment setup leads to the introduction of errors during image acquisition. In this feature, I review some of the most common practical pitfalls faced by researchers during image acquisition, and how they can affect the interpretation of the experimental data.

This article is targeted neither to the microscopy guru who pushes forward the frontiers of imaging technology nor to my imaging specialist colleagues who may wince at the overly simplistic comments and lack of detail. Instead, this is for beginners who gulp with alarm when they hear the

word “confocal pinhole” or sigh as they watch their cells fade and die in front of their very eyes time and time again at the microscope. Take heart, beginners, if microscopes were actually so simple then many people (including myself) would suddenly be out of a job!

All data are subject to interpretation. Deliberate scientific fraud exists, but in modern microscopy a far greater number of errors are introduced in complete innocence. As an example of a common problem, take colocalization. Upstairs in the lab, a researcher collects a predominantly yellow merged image on a basic microscope, naturally interpreted as colocalization of green and red signals. But on the confocal microscope, there is no yellow in the merged image.

**“When you employ the microscope, shake off all prejudice, nor harbour any favorite opinions; for, if you do, ‘tis not unlikely fancy will betray you into error, and make you see what you wish to see.” Henry Baker, chapter 15, “Cautions in viewing objects” of *The Microscope Made Easy, 1742*.**

How can this be? Many factors contribute. Here, I take the reader through the imaging process, from sample preparation to selection of the imaging and image-processing methods. Throughout, we will be on the lookout for problems that can produce misleading results, using colocalization as the most common example. Because one

short article cannot be an exhaustive “how to” guide, I have also assembled a bibliography of a few highly recommended textbooks and microscopy web sites, which readers should consult for more extensive treatments of the critical issues introduced here.

### Sample preparation

“Garbage in = garbage out” is the universal motto of all microscopists. A worrying tendency today is to assume that deconvolution software or confocal microscopes can somehow override the structural damage or suboptimal immunolabeling induced by poor sample preparation. The importance of appropriate fixation, permeabilization, and labeling methods for preserving cellular morphology or protein localization is well known to electron microscopists (Hayat, 2000), but often underestimated in optical microscopy (Fig. 1).

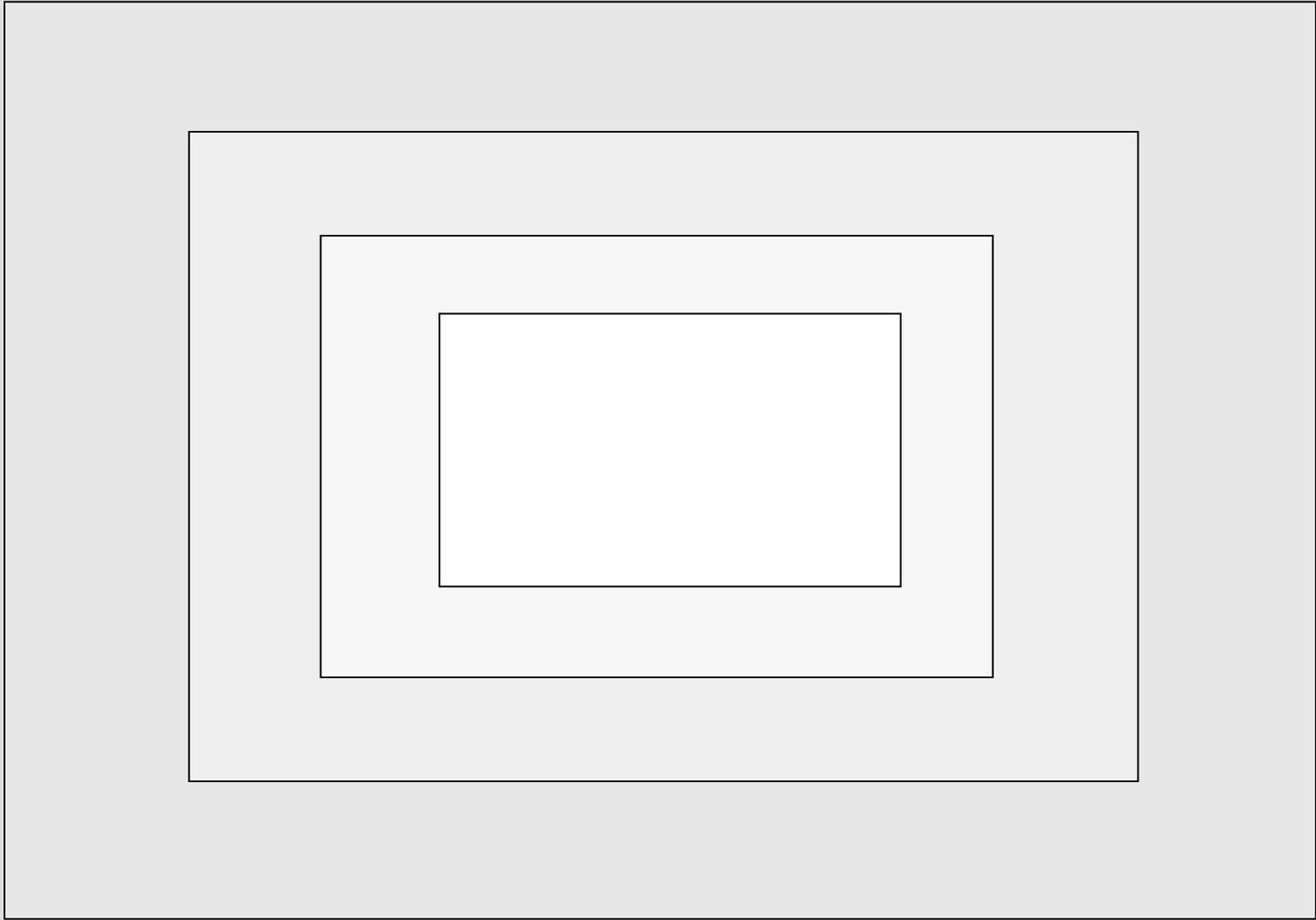
Many labs use one standardized protocol for labeling with all antibodies, irrespective of whether the targets are membrane- or cytoskeleton-associated, nuclear or cytosolic. However, inappropriate fixation can cause antigen redistribution and/or a reduction in antigenicity. It is therefore important to test each antibody on samples fixed in a variety of ways, ranging from solvents such as methanol to chemical cross-linking agents such as paraformaldehyde and glutaraldehyde (for protocols see Bacallao et al., 1995; Allan, 1999), although glutaraldehyde fixation often reduces antigenicity and increases background and autofluorescence. Consult textbooks for notorious pitfalls: phalloidin labeling is incompatible with methanol fixation, while microtubules are inadequately fixed by formaldehyde. Moreover, certain cell types, such as yeast cells, require specialized fixation protocols (Hagan and Ayscough, 1999).

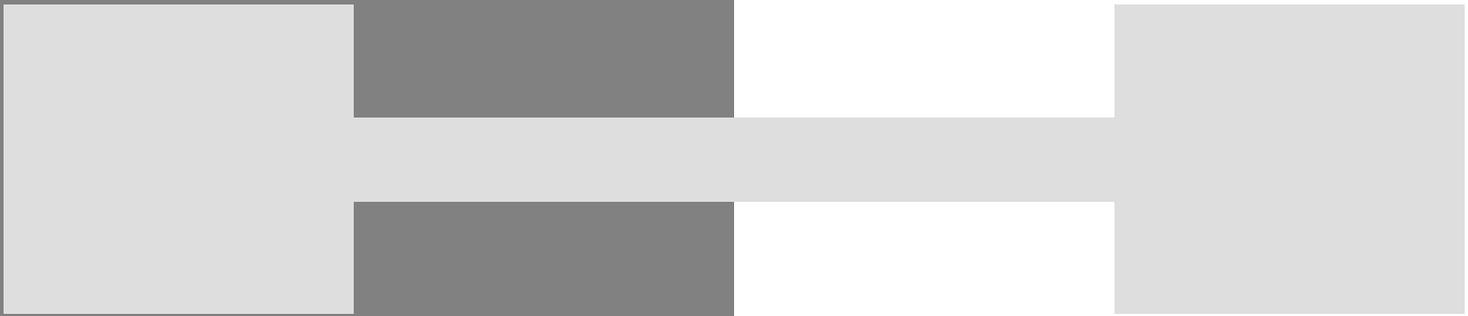
Permeabilization is also critical in achieving a good compromise between antigen accessibility and ultrastructural integrity. Specific detergents will produce different effects (for example, Triton treatment produces smaller holes in

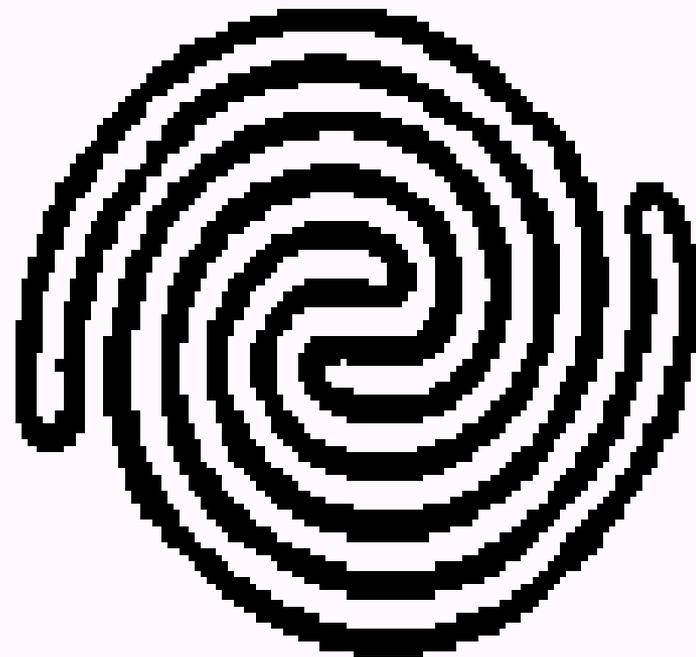
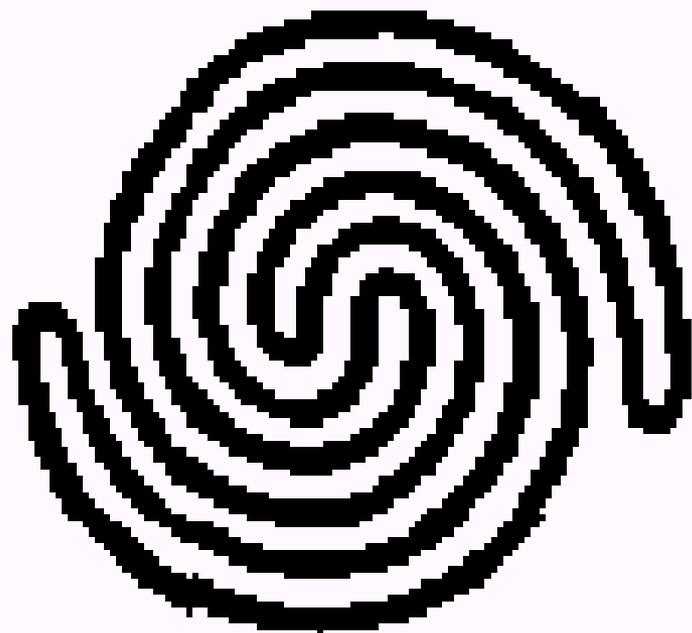
JCB | FEATU

THE JOURNAL OF CELL BIOLOGY

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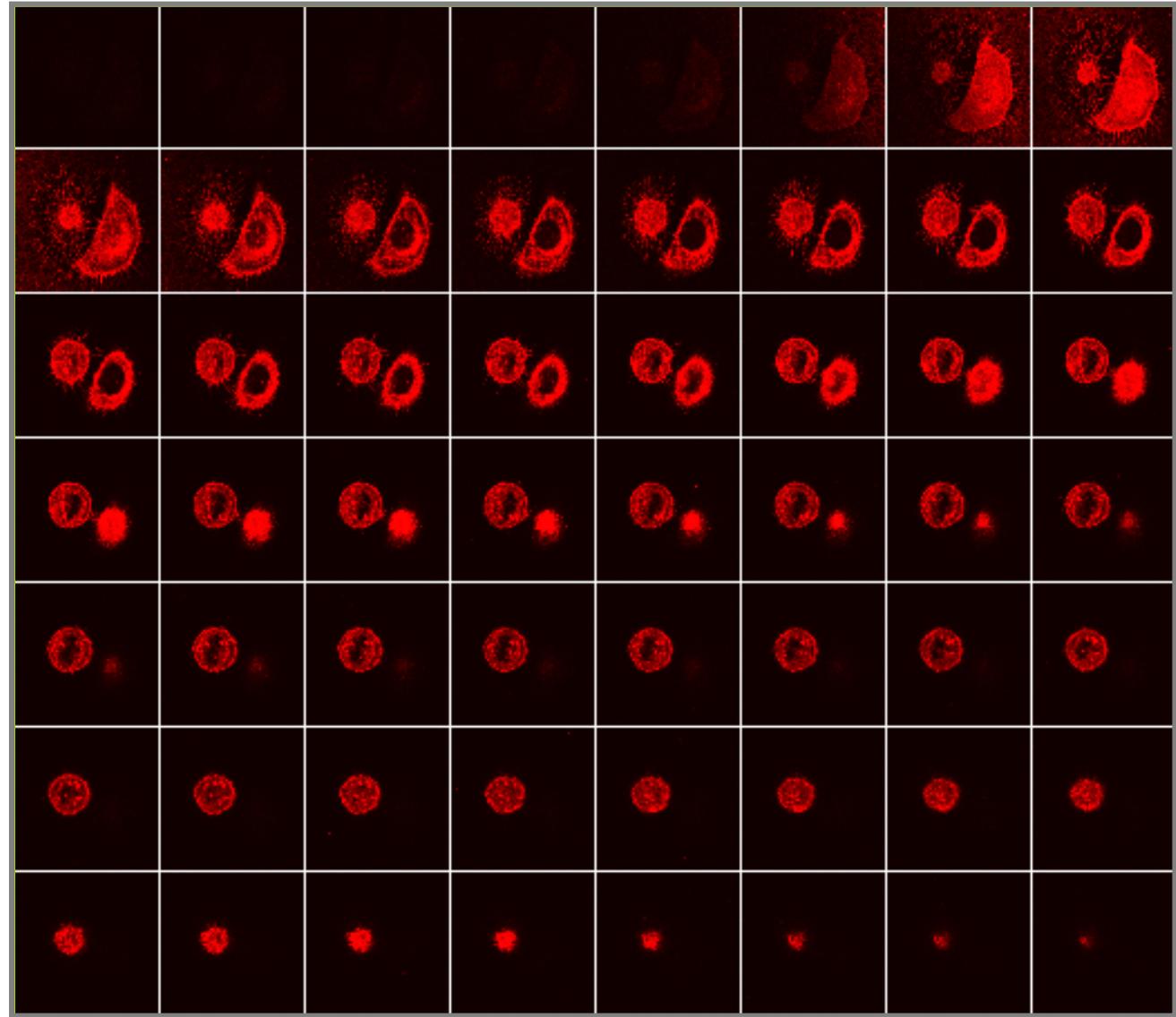




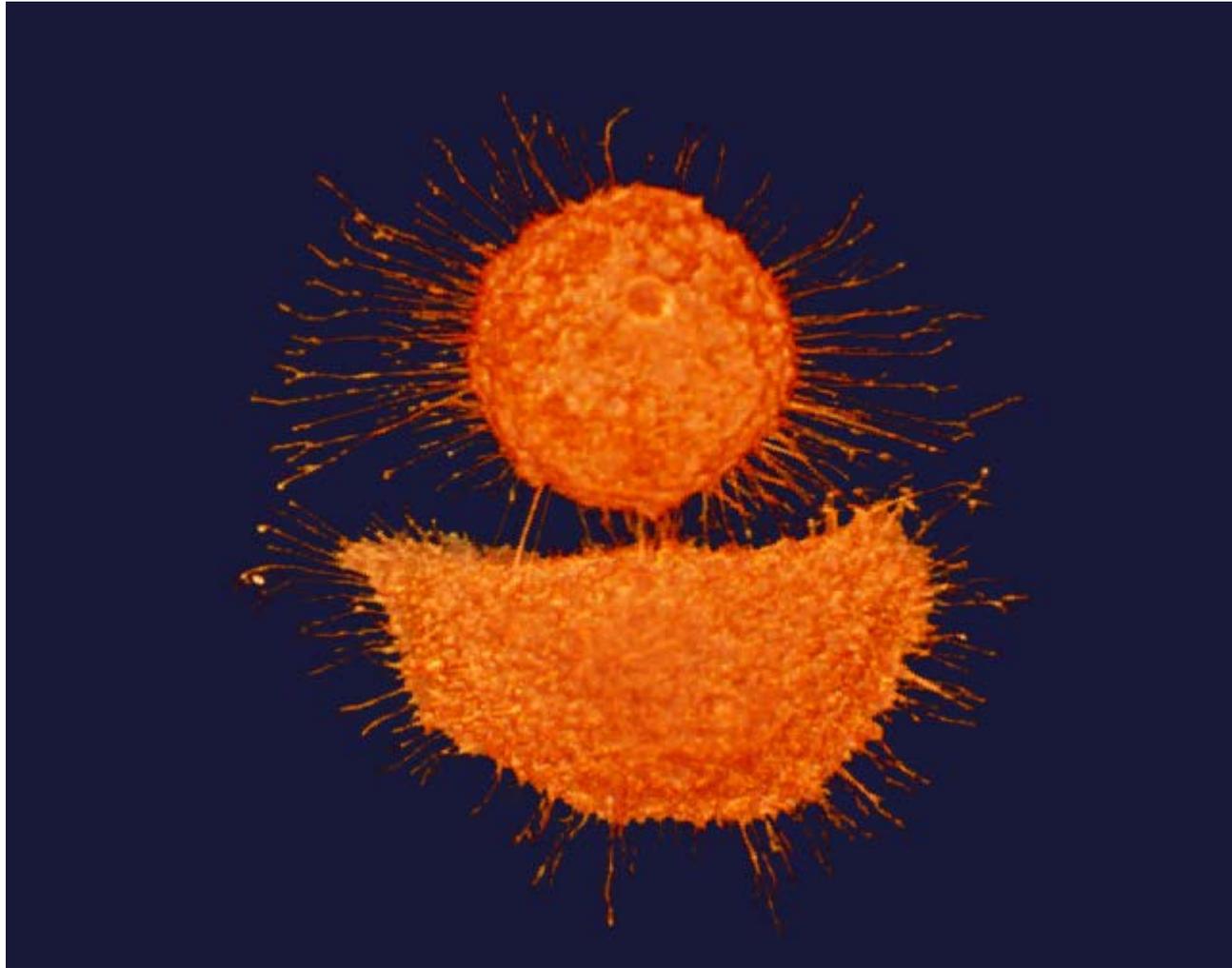
**Hans Janssen (1595), ....  
Galileo Galilei (1610)**



# | - > Microscopy



# | - > Microscopy



Luminescencia:



- *Fluorescencia*  $\Delta t \sim 10^{-8}s$
- *Fosforescencia*  $\Delta t \sim 10^{-3}-10^0s$

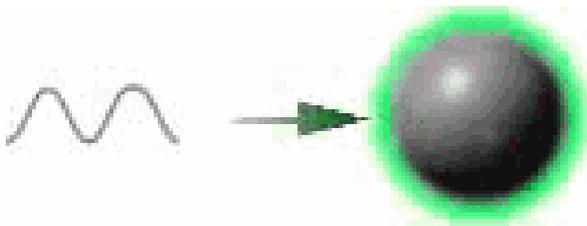
Interacciones ...

- intra- e inter moleculares ...

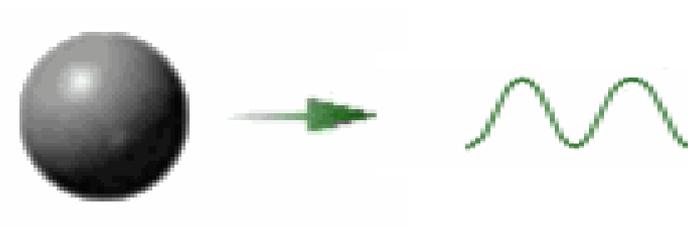
producen cambios ...

- espectrales
- tiempos de vida
- polarización
- intensidad ...

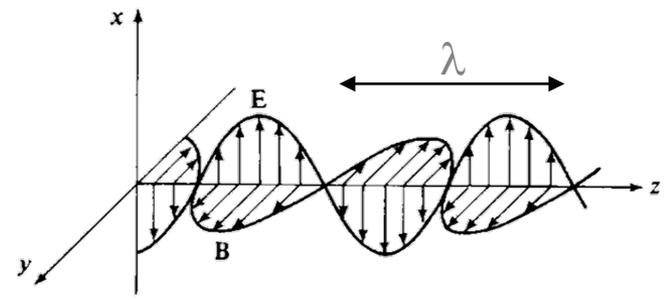
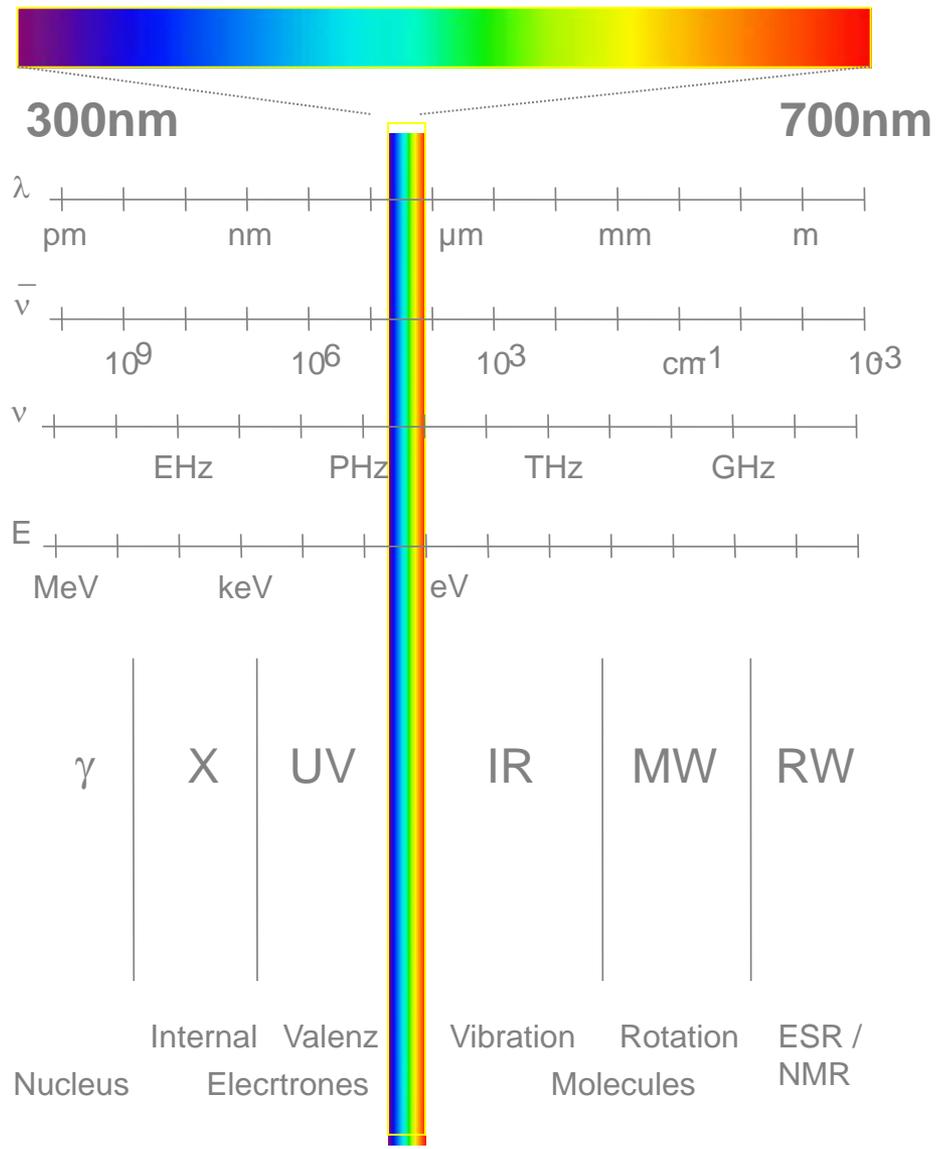
←  $\Delta t$  →



*Absorción / Excitación*



*Emisión*



## Energía de un fotón: (~1-5eV)

$$E = h \nu = hc\lambda^{-1} \quad | \quad c = \nu \lambda$$

$\nu$ , frecuencia [ $\text{s}^{-1}$ ]  
 $h$ , constante de Planck [ $6.626 \cdot 10^{-34} \text{ Js}^{-1}$ ]  
 $\lambda$ , longitud de onda [m]  
 $c$ , velocidad de luz [ $\sim 3 \cdot 10^8 \text{ ms}^{-1}$ ]

## Energía molecular:

$$E = E_{\text{rot}} + E_{\text{vib}} + E_{\text{el}}$$

1 :  $10^3$  :  $50 \cdot 10^3$

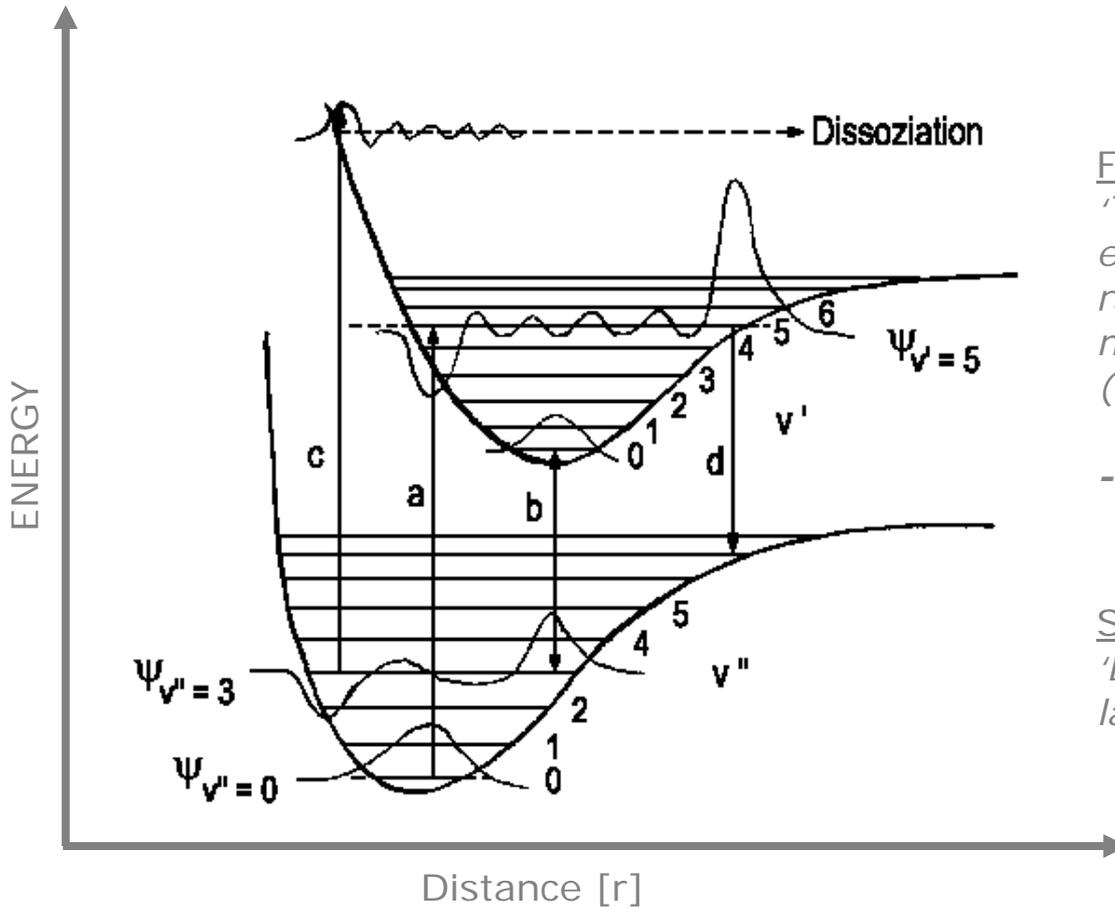
## Energía térmica:

$$E = k T \quad (\sim 2.5 \cdot 10^{-2} \text{ eV}, T \sim 20^\circ\text{C})$$

$k$  = Constante de Boltzmann ( $0.86 \cdot 10^{-4} \text{ eV/K}$ )



# | -> Franck Condon



Frank Condon :

*'Transiciones entre niveles electrónicos ocurren mucho más rápido que movimientos de núcleos moleculares.'*

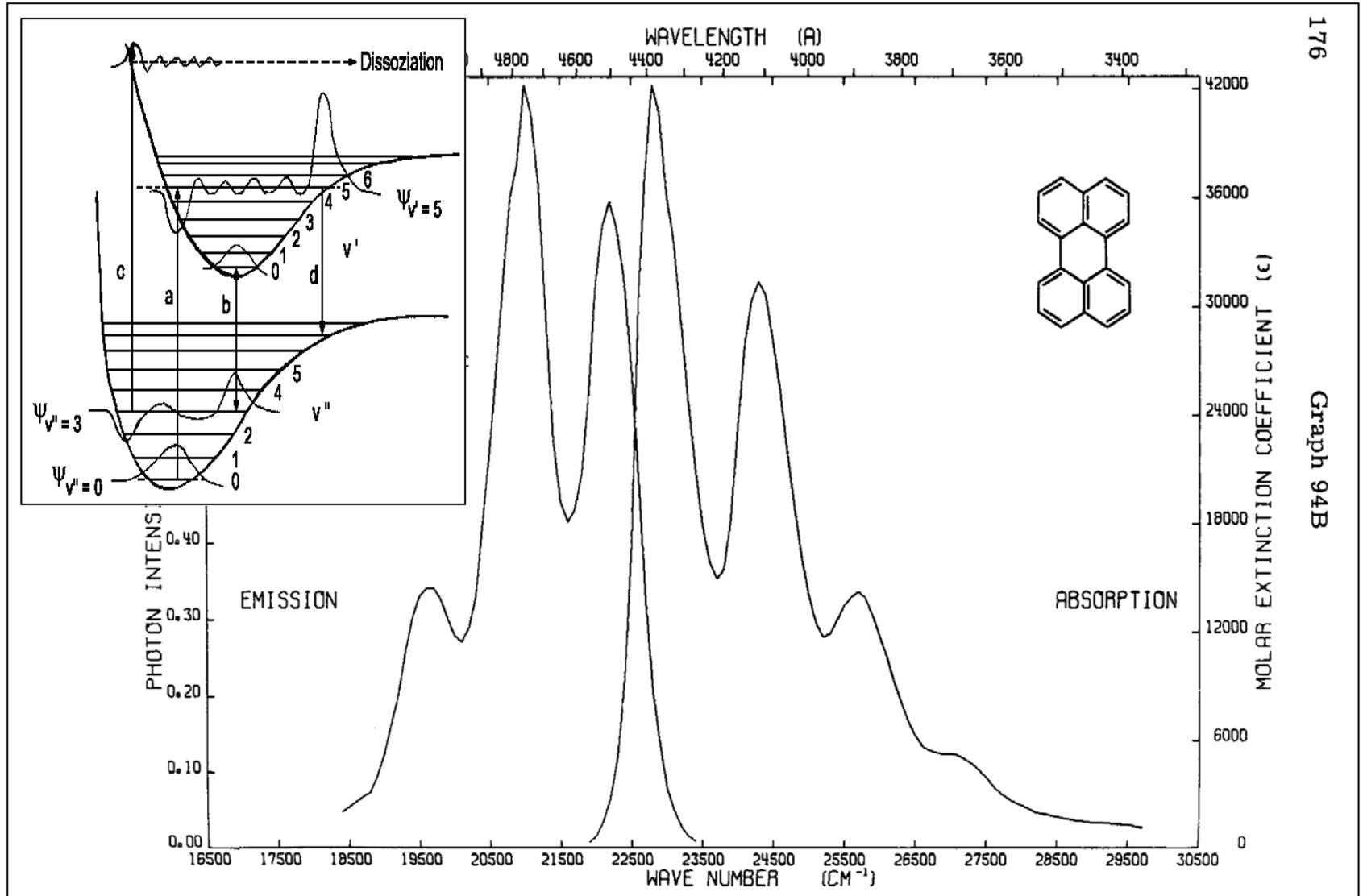
*(masa<sub>electron</sub> / masa<sub>atom</sub>: 1 : 2000)*

**-> Mirror Image Rule**

Stokes (Shift) :

*'La energía de emisión es menor a la energía de excitación.'*

# | - > Mirror Image Rule



## “Principles of Fluorescence Spectroscopy” Chapter 1

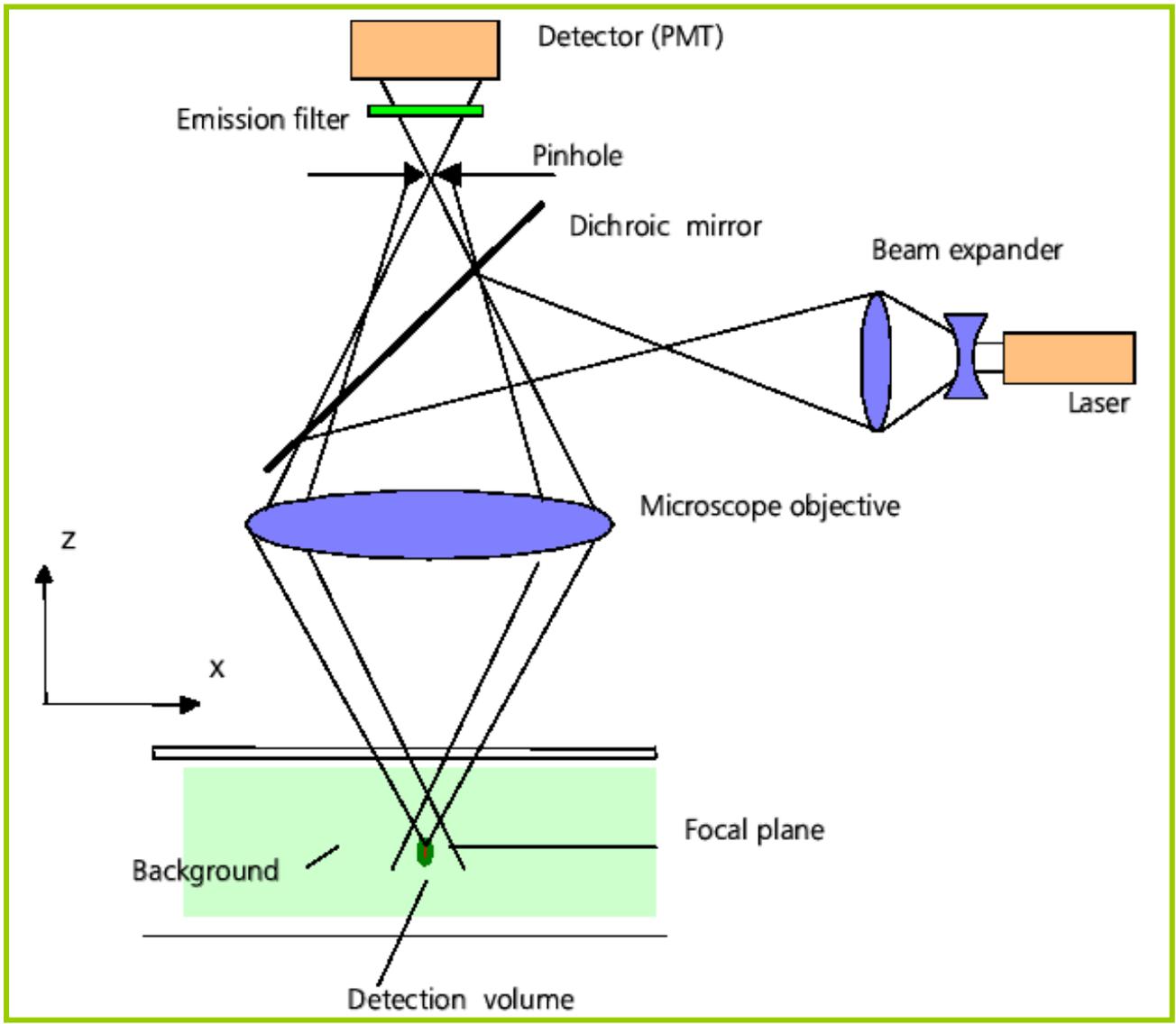
| - > Polarisation ...

| - > Steady-State and Time-Resolved Fluorescence...

| - > Resonance Energy Transfer ...

| - > Quenching, Bleaching ...

# | -> Diffraction limited Microscopy





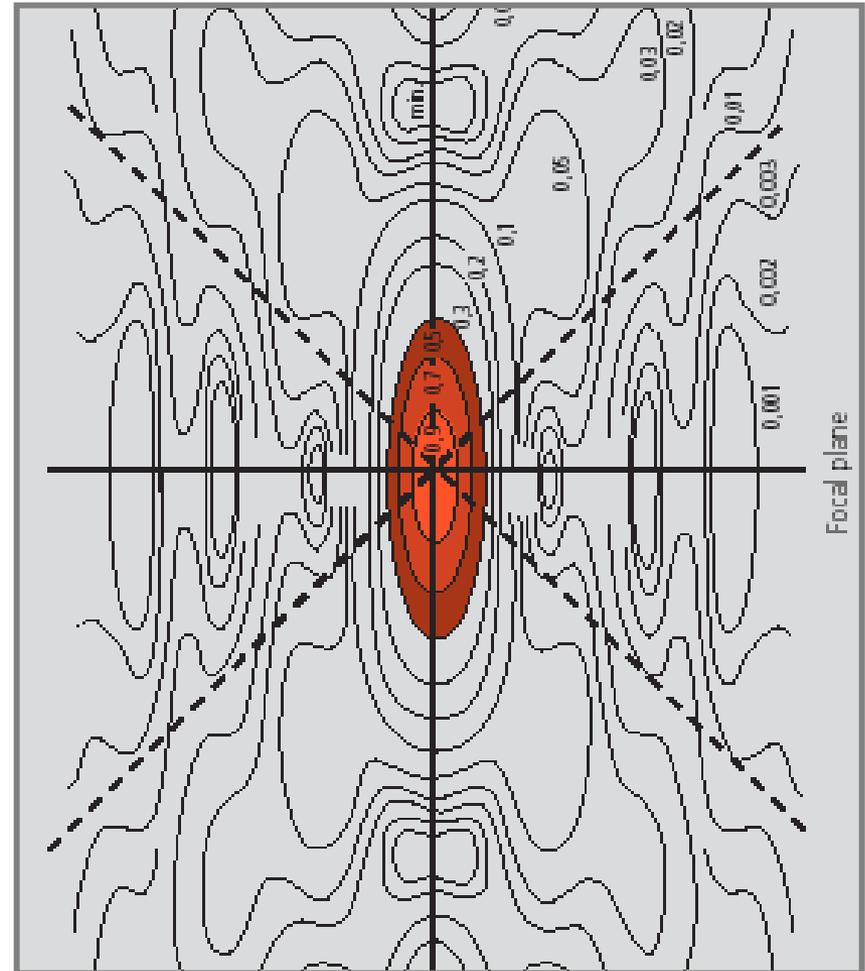
Óptica no-geométrica /  
Teoría de difracción

$$\text{PSF} = |U|^2 = f(J_0)$$

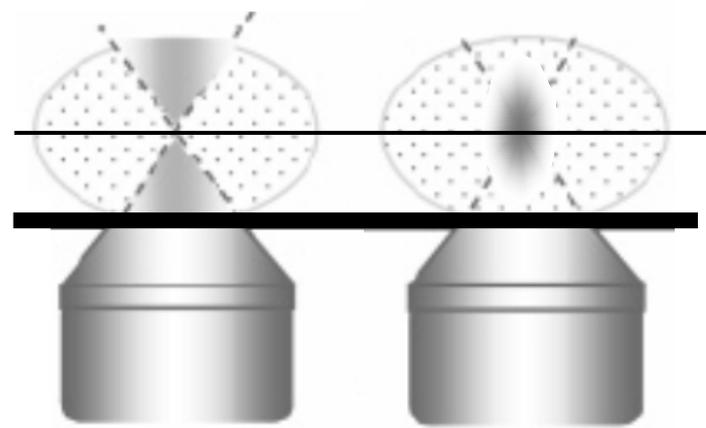
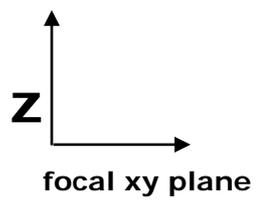
$U$ , Integral de Difracción de Kirhoff

$J_0$ , Serie de funciones de Bessel

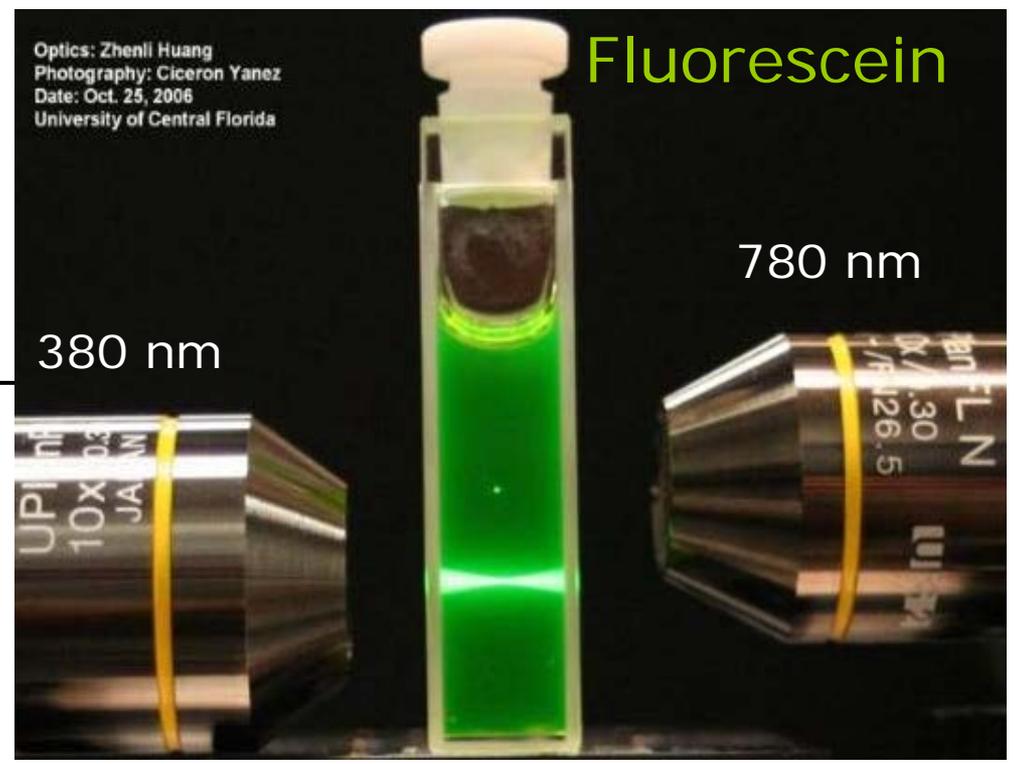
*(Born & Wolf, Principles of Optics, 6th edition 1988,  
Pergamon Press)*



| Best localization: confocal microscopy

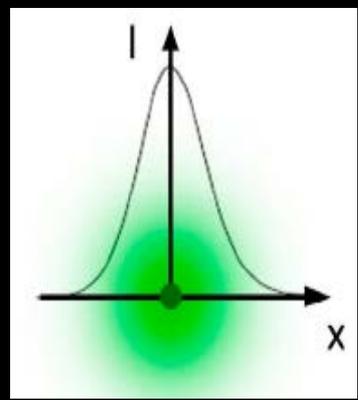
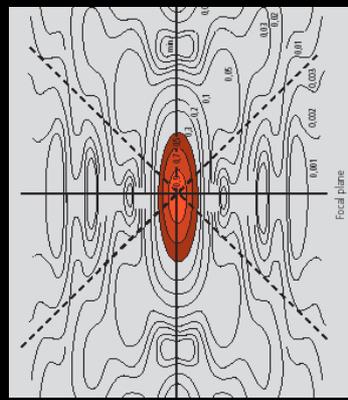
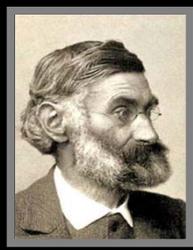


convencional / confocal

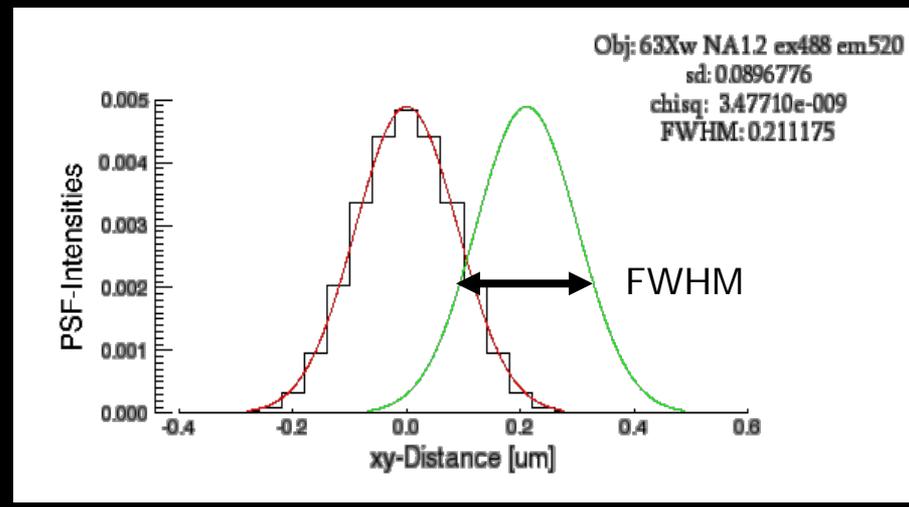


# | - > Diffraction limited Microscopy

Ernst Abbe  
1840 - 1905



Resolution  $\sim \lambda / 2 \sim$  Full Width at Half Maximum (FWHM)



# | -> Beyond diffraction

M Goepfert-Mayer  
1906-1972

M Gustafson  
1960-2011

S Hell  
MPI Göttingen  
BIOQUANT Hdg

E Betzig  
Janelia Farm



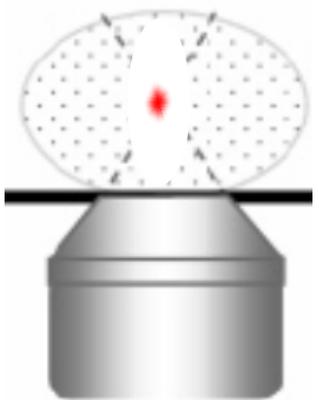
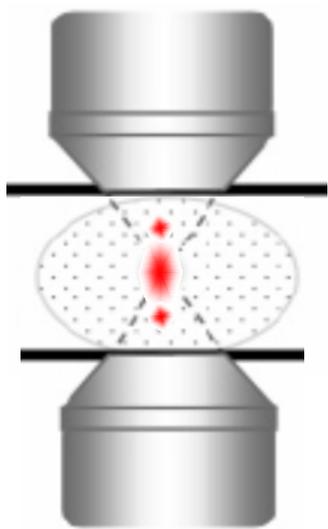
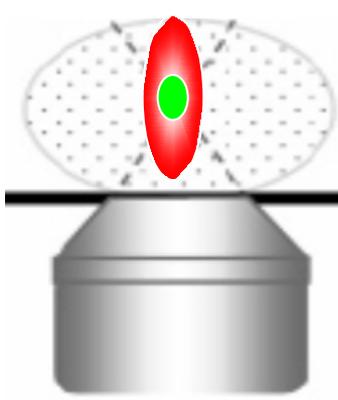
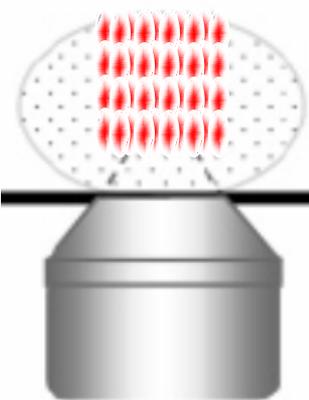
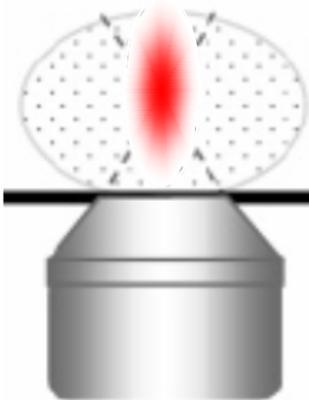
FWHM(xy) ~  $\lambda/2$

~  $\lambda/4$

~  $\lambda/\infty$

~  $\lambda/4$

~  $\lambda/100$



2-photon

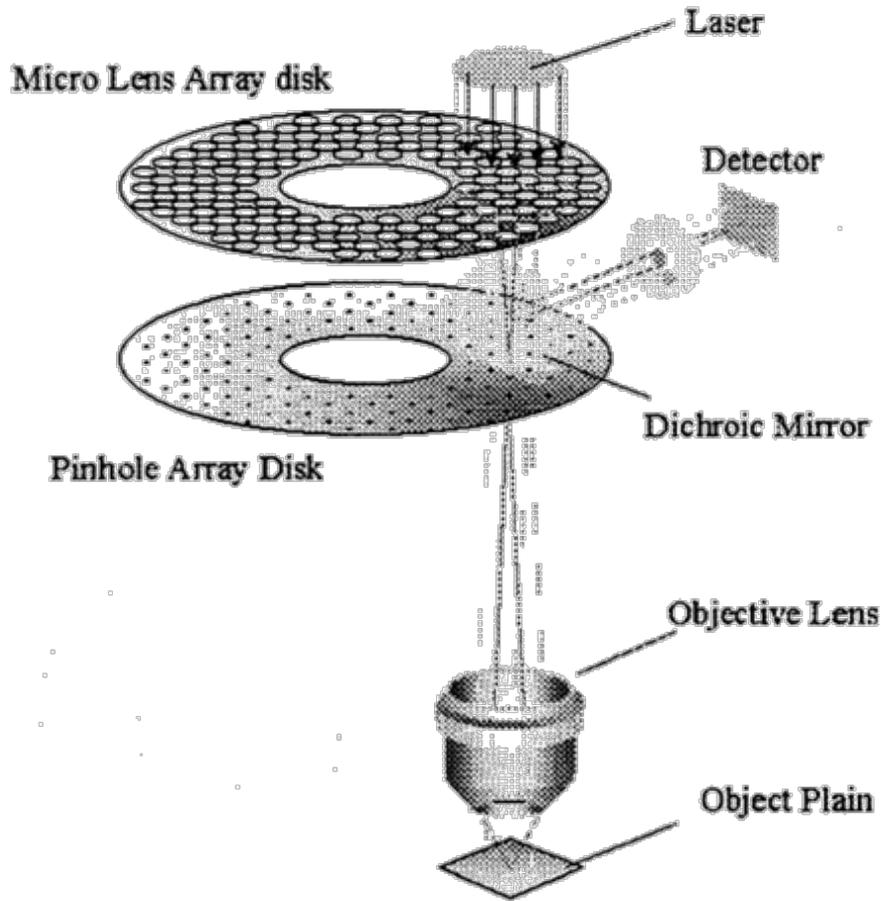
SIM

STED

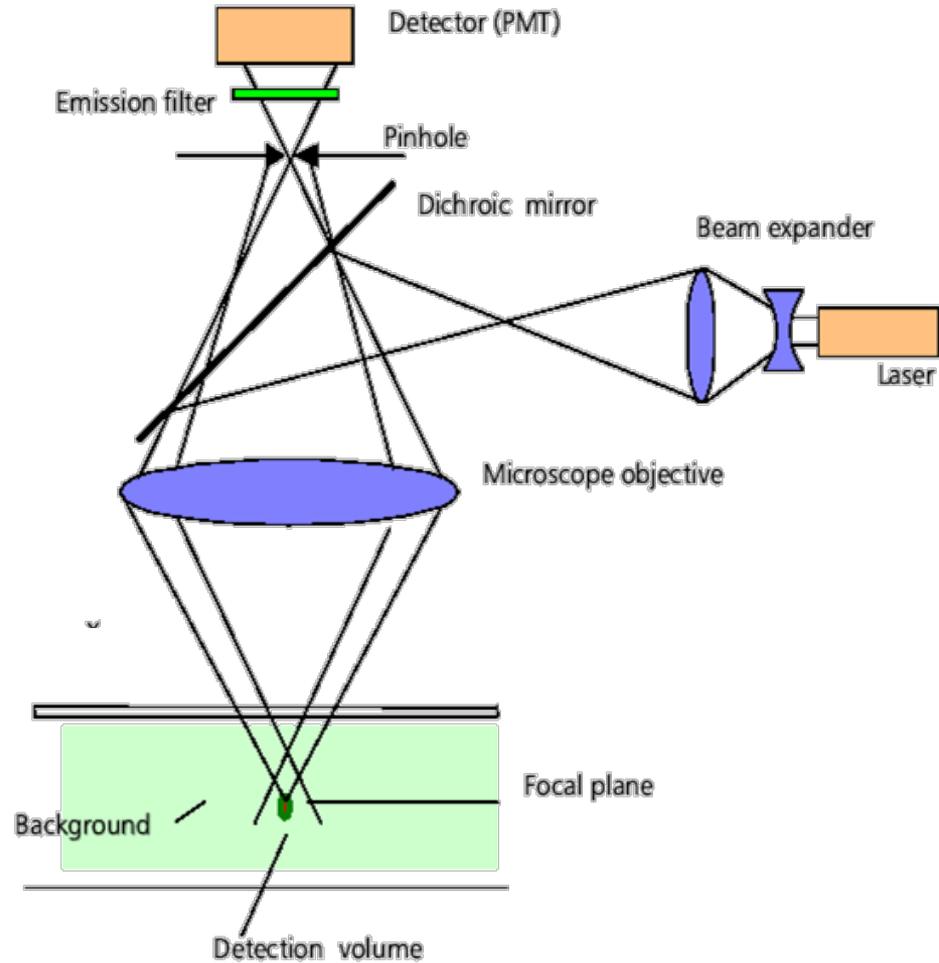
4- $\pi$

PALM

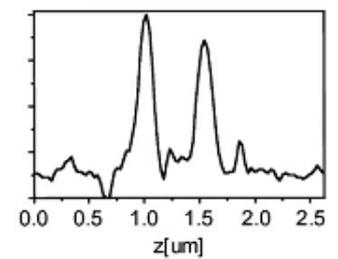
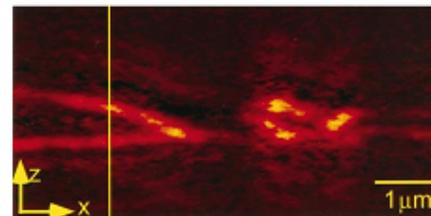
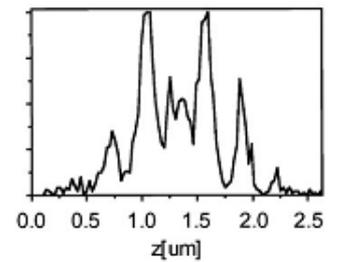
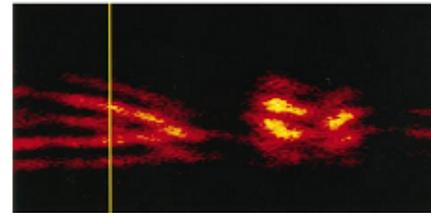
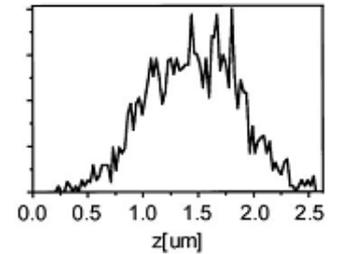
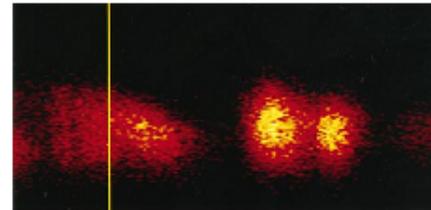
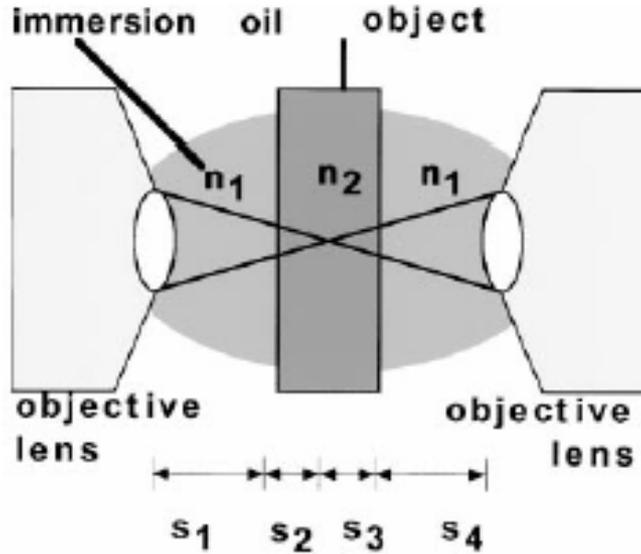
# | - > Spinning Disk



**spinning disk**



**confocal**



## 4Pi-Confocal Imaging in Fixed Biological Specimens

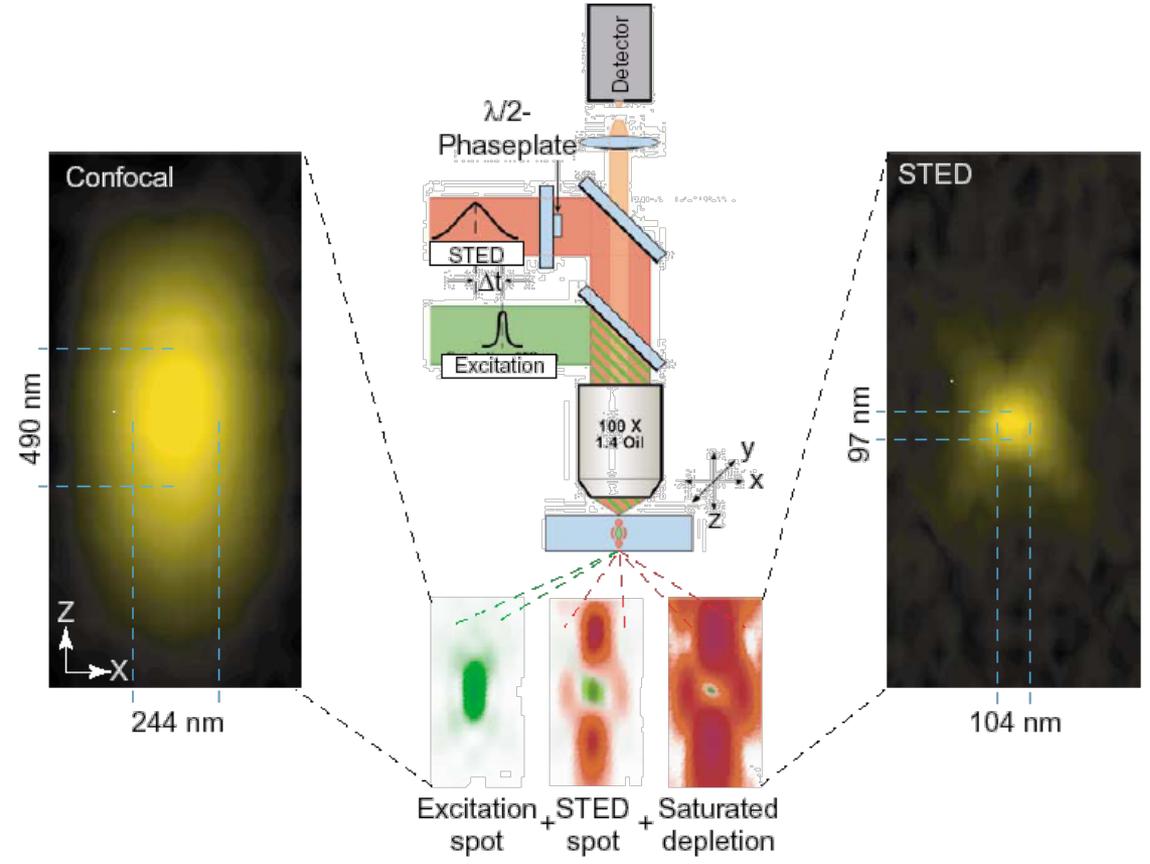
Martin Schrader,\* Karsten Bahlmann,\* Günter Giese,# and Stefan W. Hell\*  
 Biophysical Journal Volume 75 October 1998 1659-1668

# STED Microscopy

Stimulated

Emission

Depletion



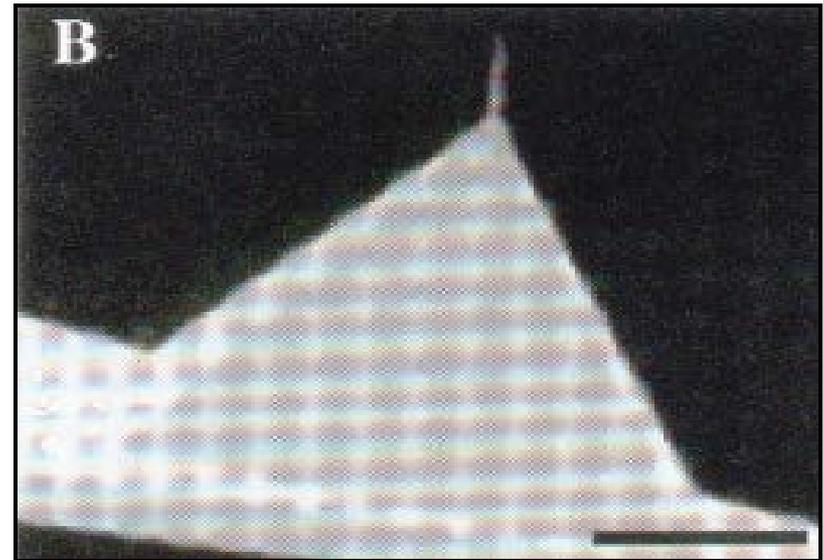
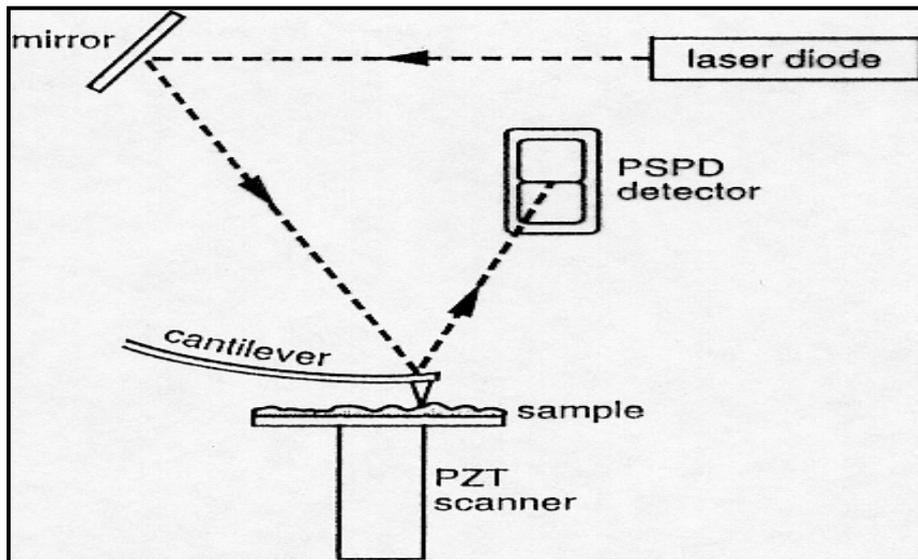
## Concepts for nanoscale resolution in fluorescence microscopy

Stefan W Hell\*, Marcus Dyba<sup>1</sup> and Stefan Jakobs<sup>2</sup>

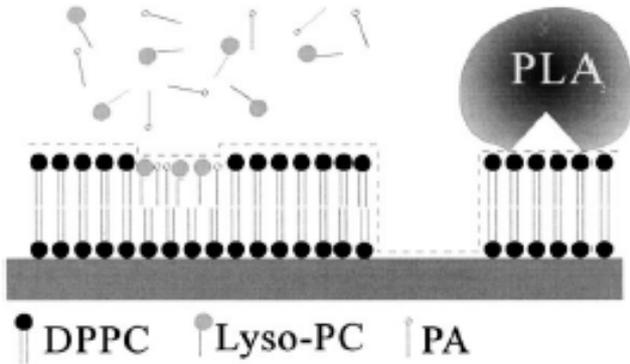
Current Opinion in Neurobiology 2004, 14:599-609

AFM allows the investigation of structural and functional properties of biomolecules in liquid environments, by a unique combination of :

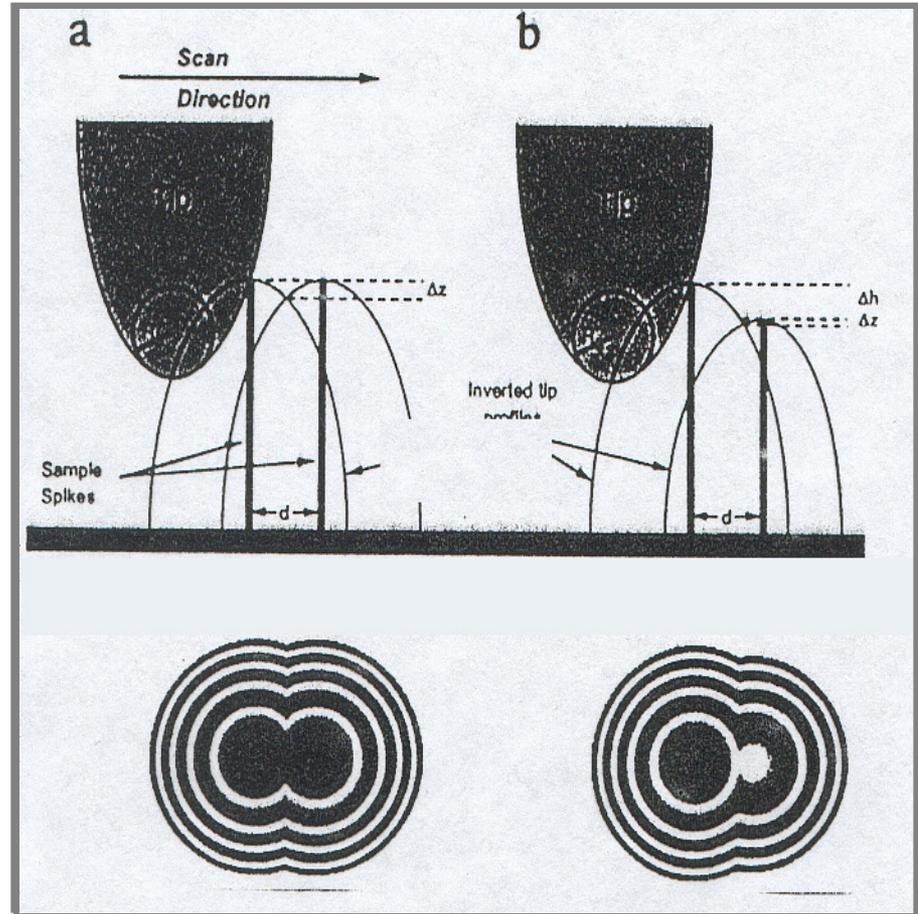
- *subnanometer* spatial resolution
- *millisecond* temporal resolution
- *piconewton* force sensitivity



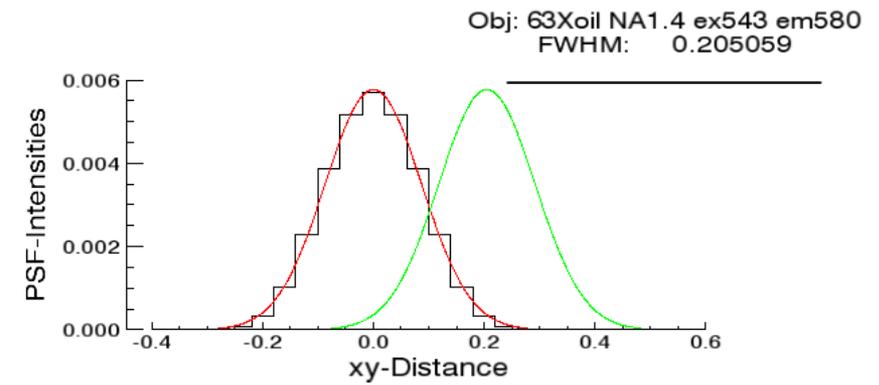
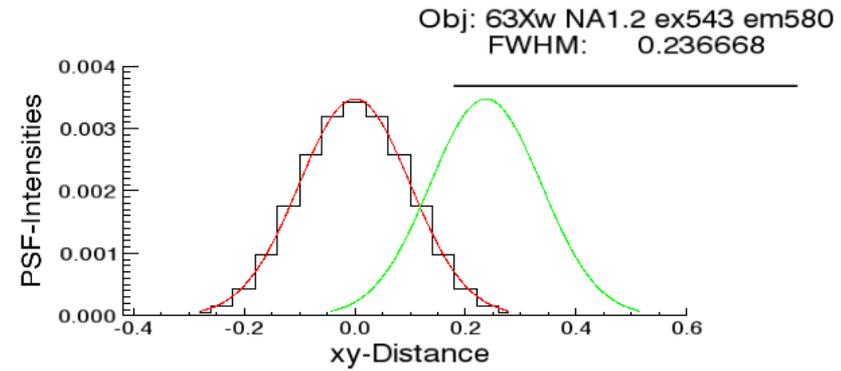
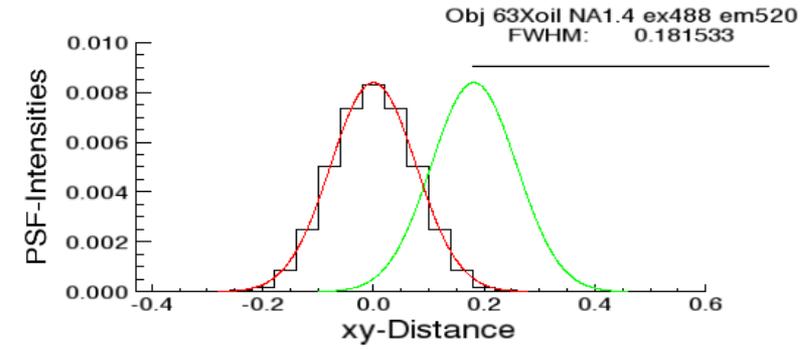
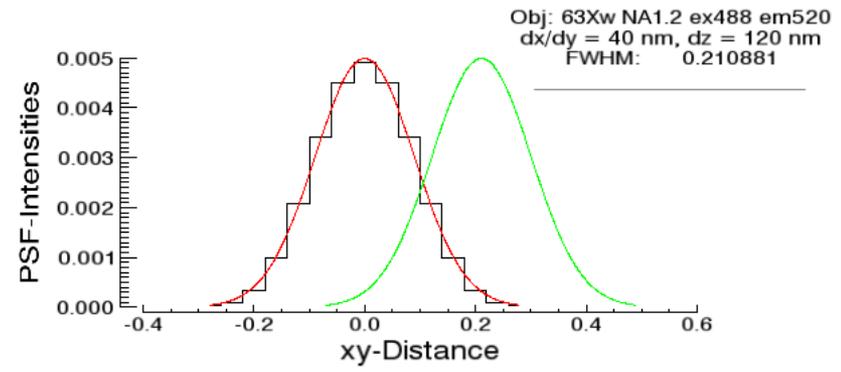
# | -> Atomic Force Microscopy



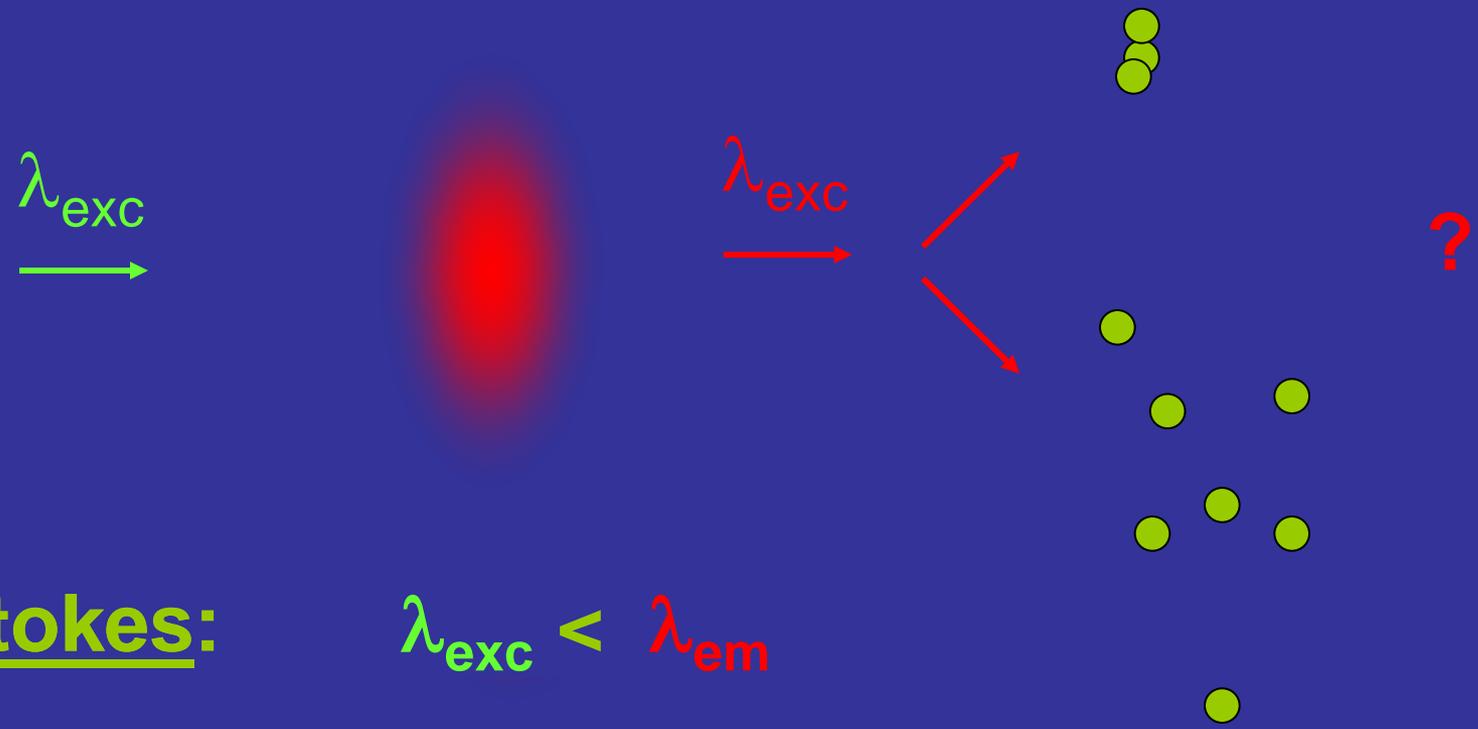
*M Grandbois et al. (1998) Biophys J.*



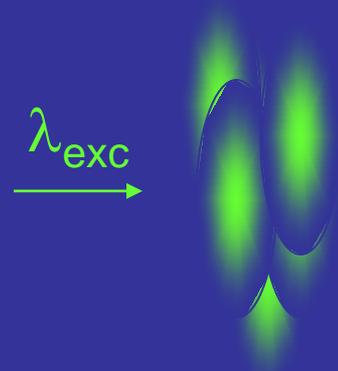




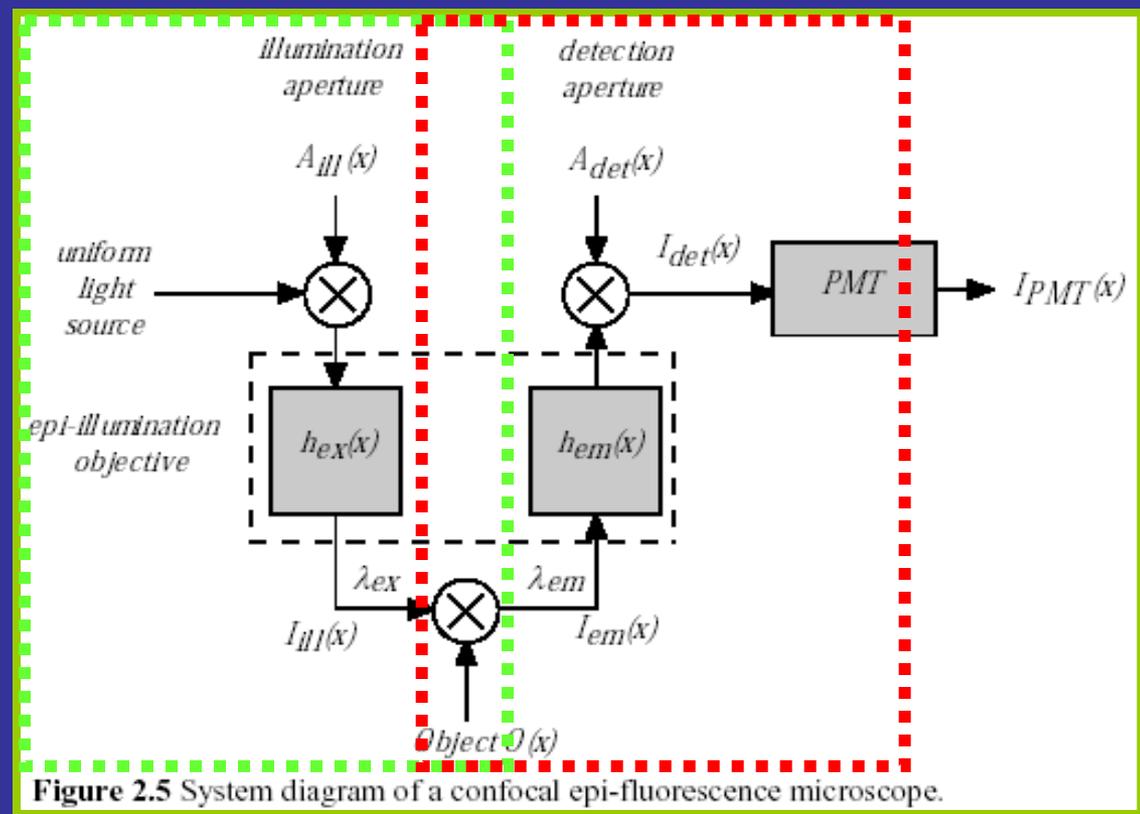
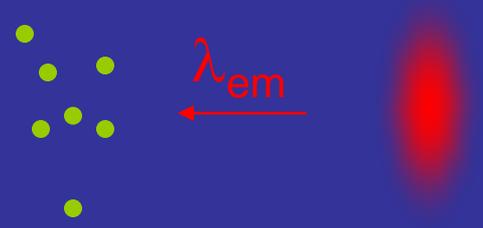
# | -> Convolution



# | - > Convolution

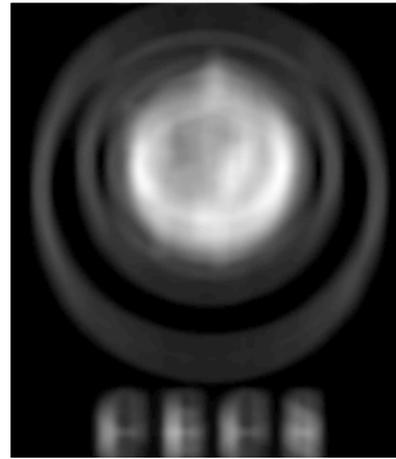
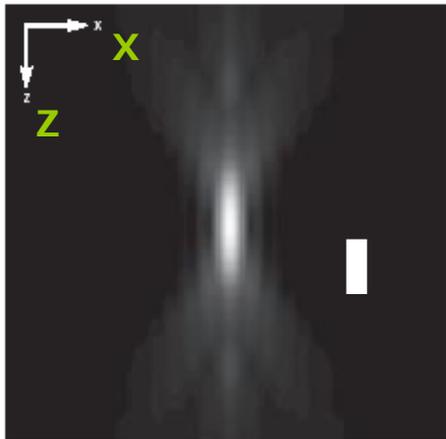


Stokes:  $\lambda_{exc} < \lambda_{em}$   
 $n(\lambda_{exc}) > n(\lambda_{em})$



**Figure 2.5** System diagram of a confocal epi-fluorescence microscope.

# | -> Convolution

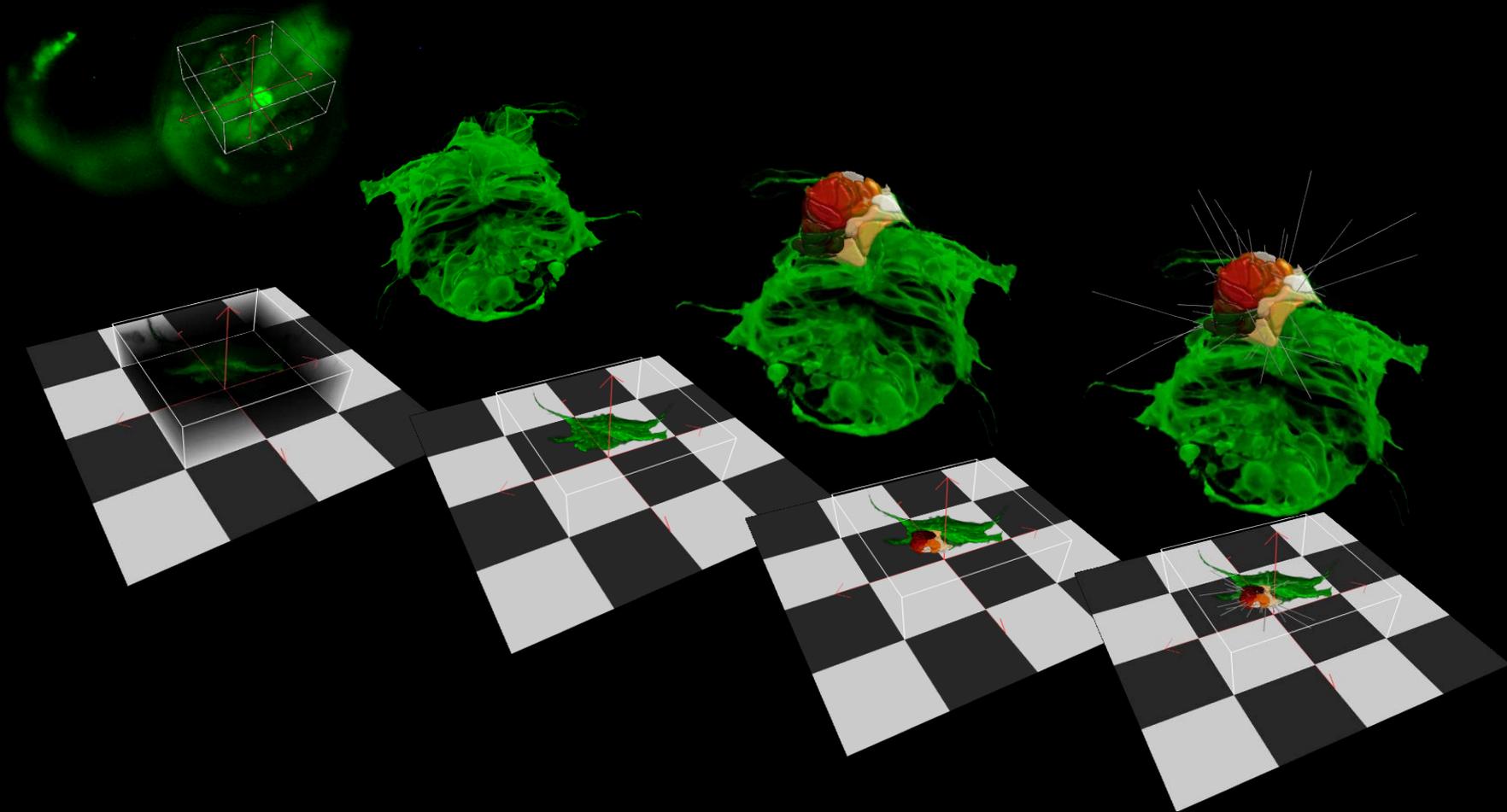


*PSF:*

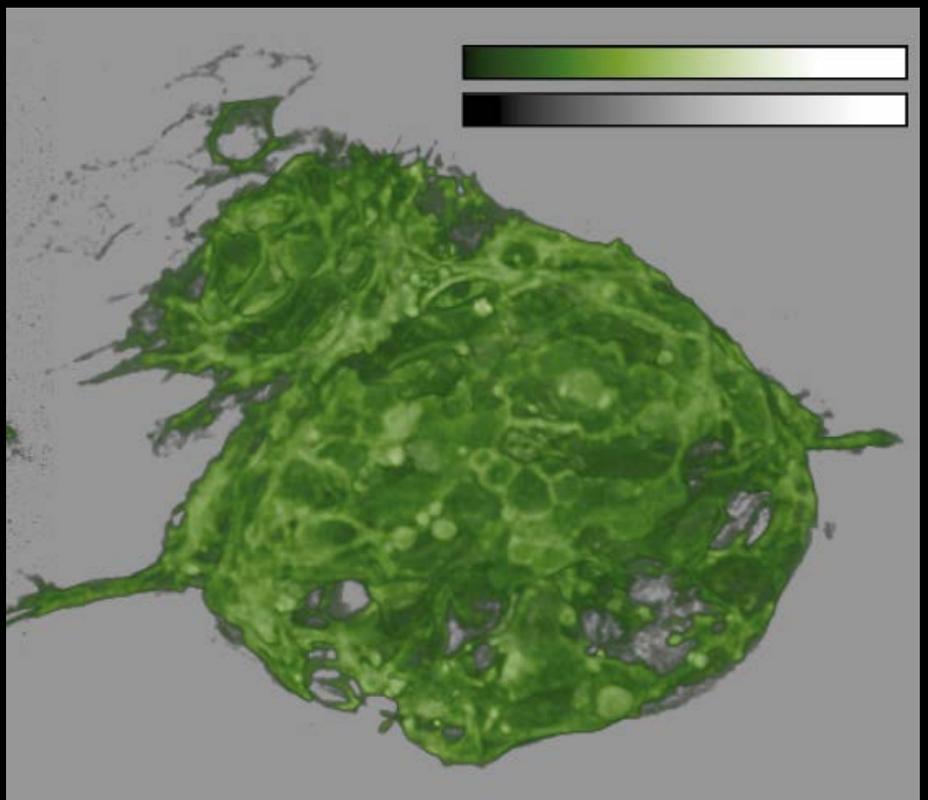
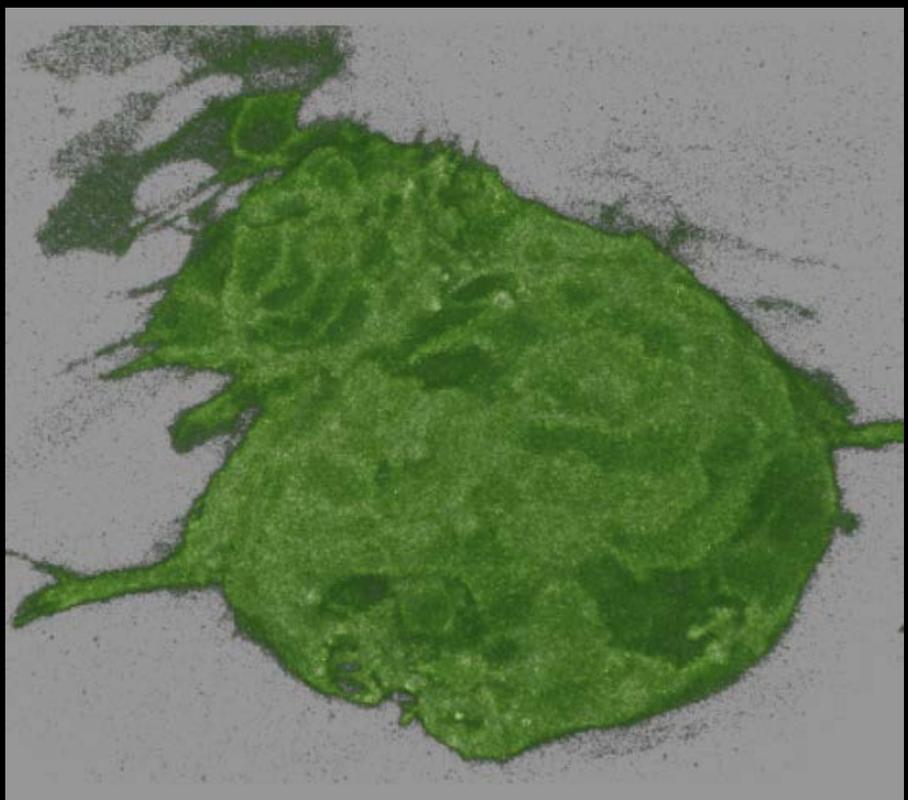
$\Delta xy \sim 500 \text{ nm}$  |  $\Delta z \sim 1500 \text{ nm}$



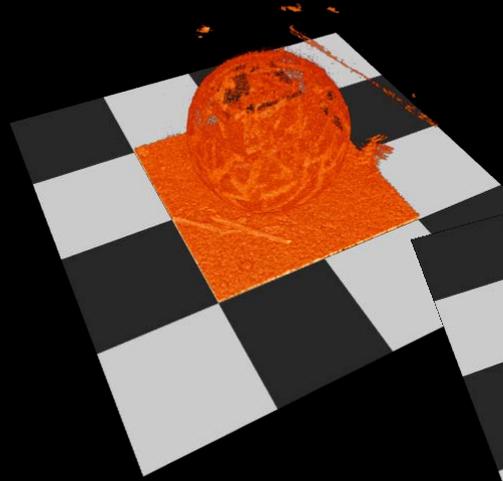
## Transgenic *flh::GFP*



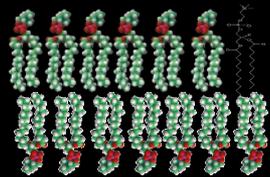
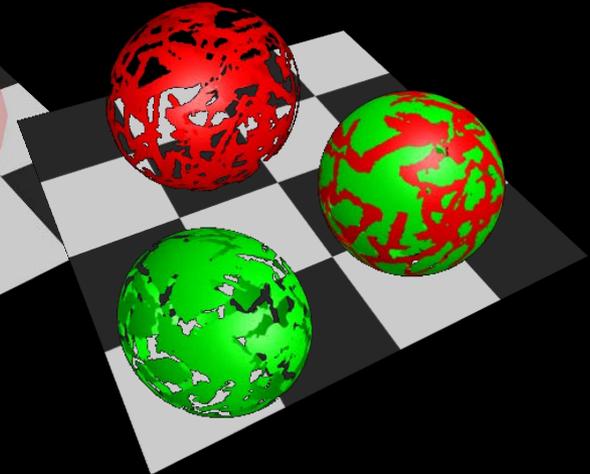
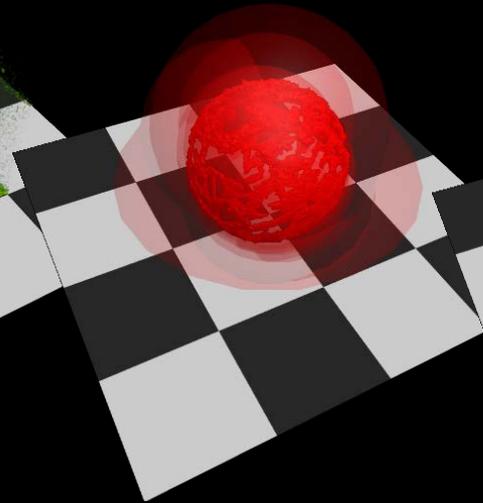
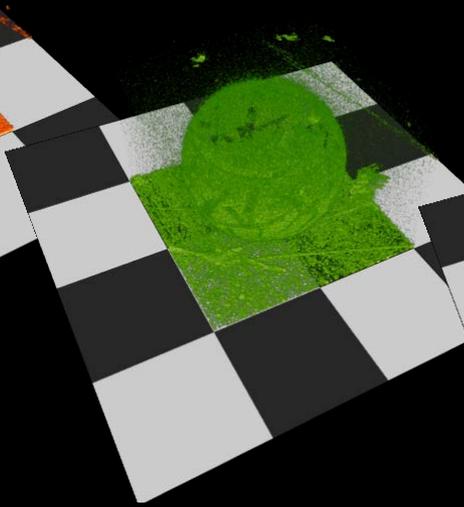
# | - > Deconvolution



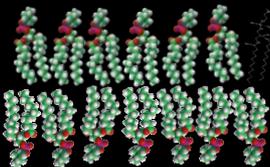
**DPPC**



**DLPC**

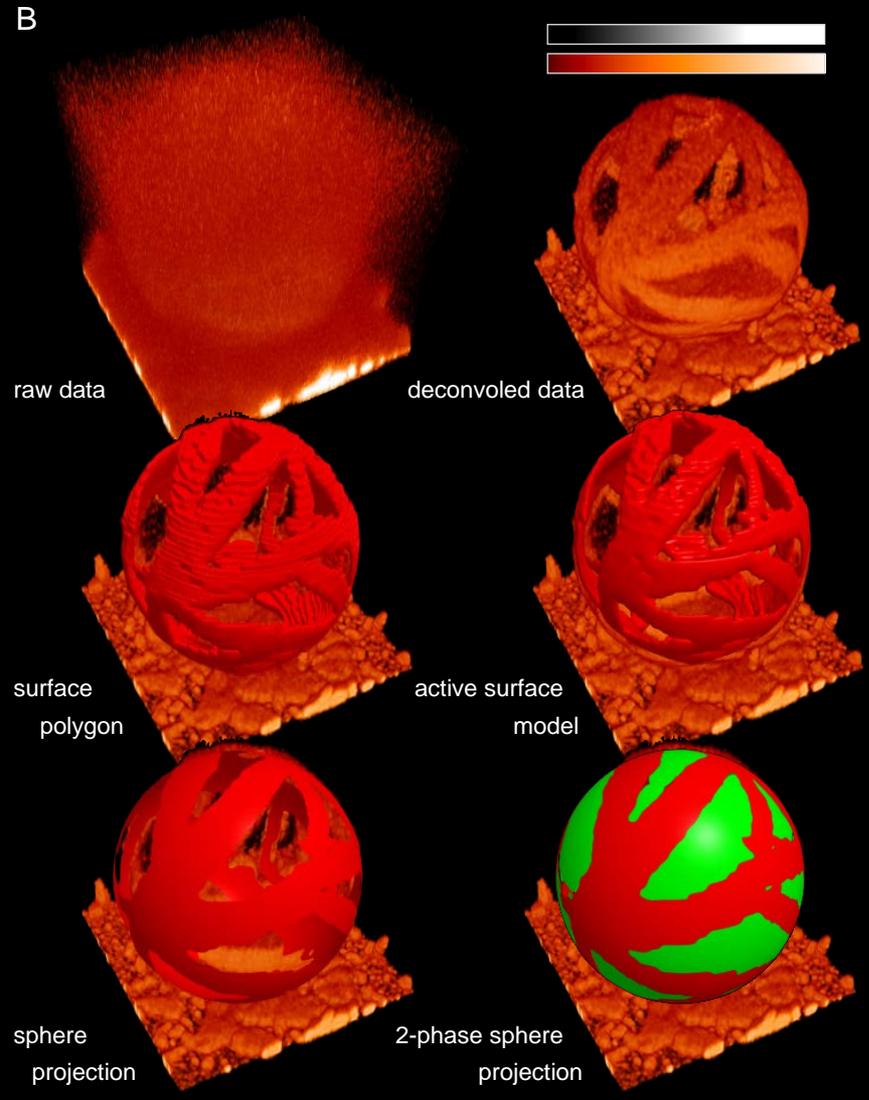
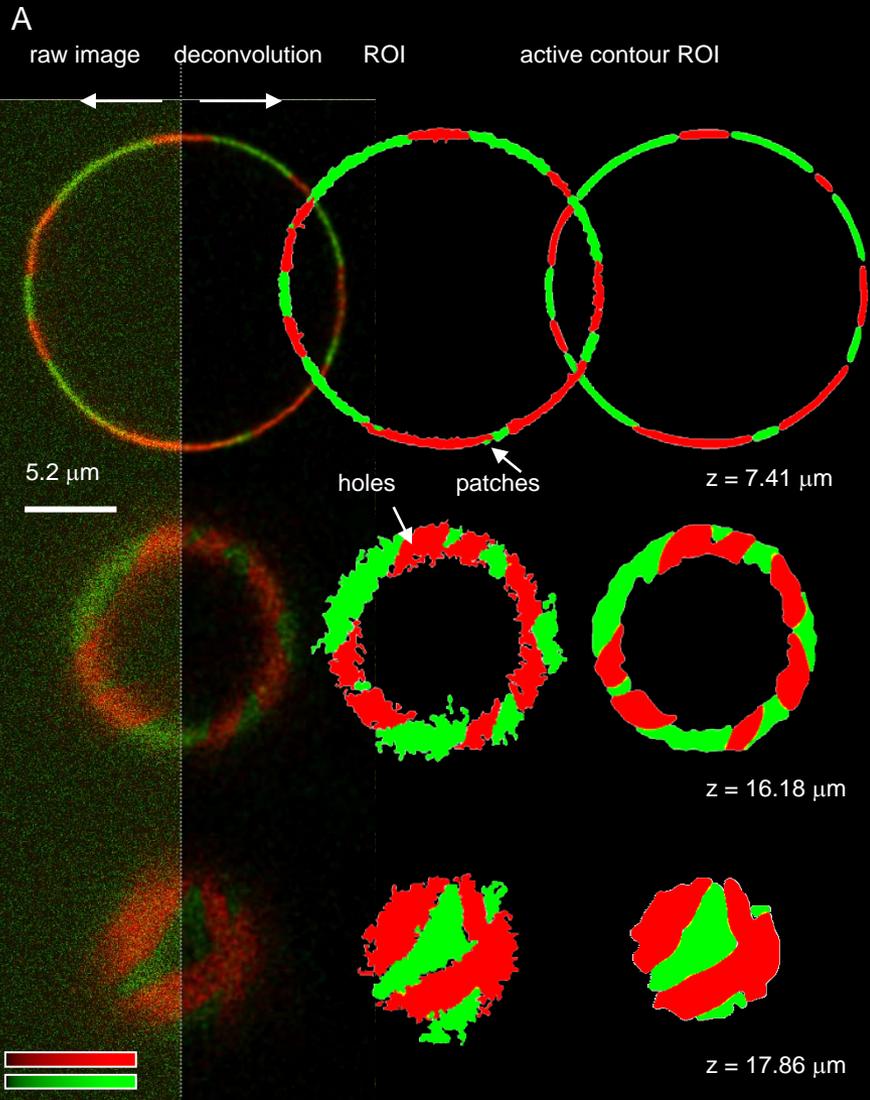


At  $T = 20\text{ }^{\circ}\text{C}$ : **Solid Gel or  $S_0$  phase is formed by DPPC 16:0 (1,2-Dipalmitoyl-*sn*-Glycero-3-Phosphocholine) + 0.5 mol% DiI $_{18}$ .**

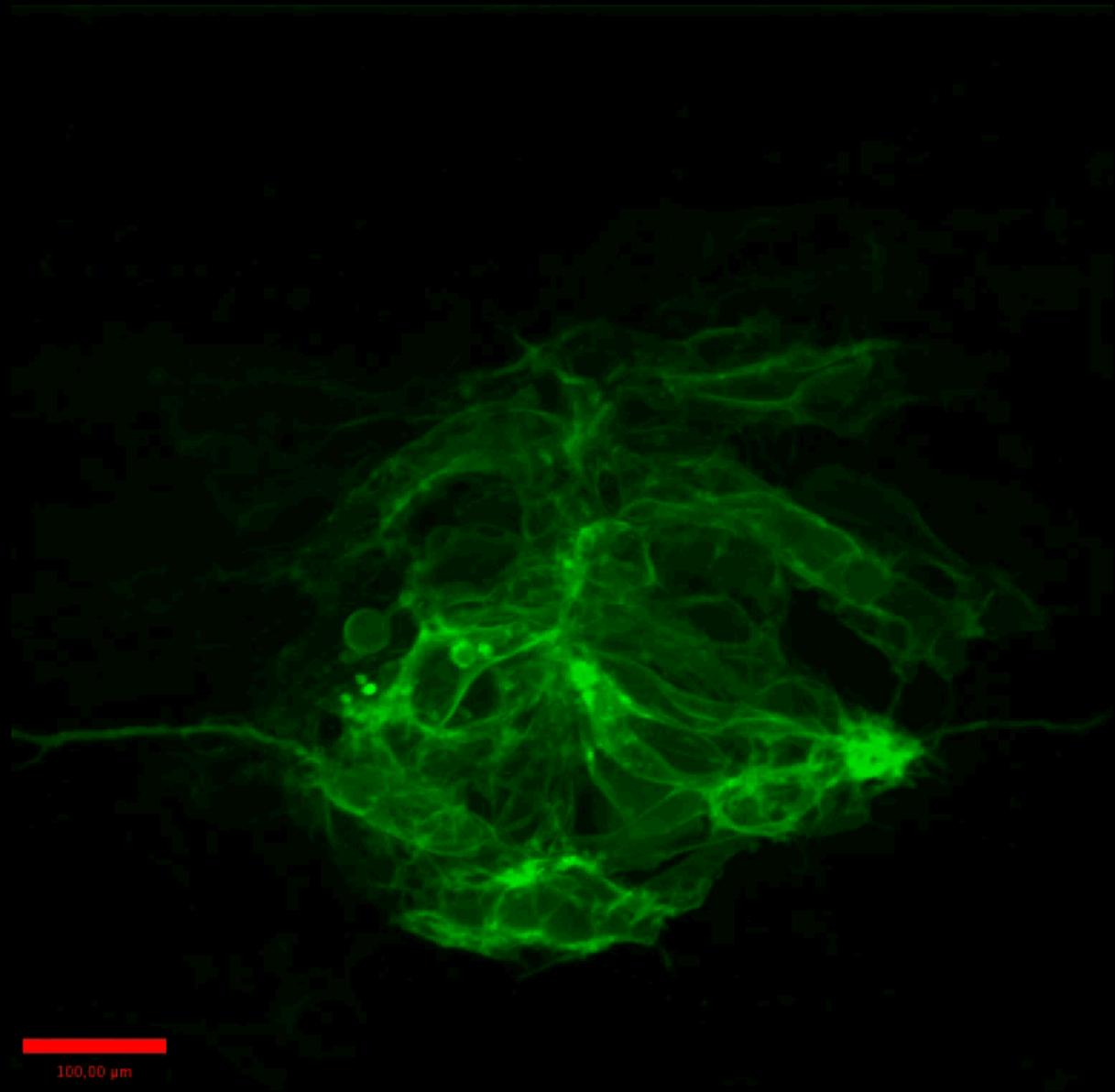


At  $T = 20\text{ }^{\circ}\text{C}$ : **Fluid phase is formed by DLPC 12:0 PC (1,2-Dilauroyl-*sn*-Glycero-3-Phosphocholine) + 0.5 mol% BODIPY-PC.**

# | - > Deconvolution



# | - > Deconvolution



*PSF: Point Spread Function*

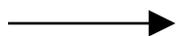
$$N(\mathbf{PSF}(x, y, z) \otimes \mathbf{f}(x, y, z) + \mathbf{b}(x, y, z)) = \mathbf{I}(x, y, z)$$

*f: Object Function*

*b: Offset Function*

*I: Image Matrix*

*N: Noise Function*



## Calculator

[Numerical aperture](#)

[Excitation wavelength](#)

 (nm)

[Emission wavelength](#)

 (nm)

[Number of excitation photons](#)

[Backprojected pinhole radius](#)

 (nm)

[B.P. distance between pinholes](#)

 Only for Nipkow disks (μm)

[Lens medium refractive index](#)

[Specimen medium refractive index](#)

[Acquisition depth](#)

 (μm)

Calculate also PSF

- confocal
- widefield
- nipkow
- 4Pi

Select one

*PSF: Point Spread Function*

*f: Object Function*

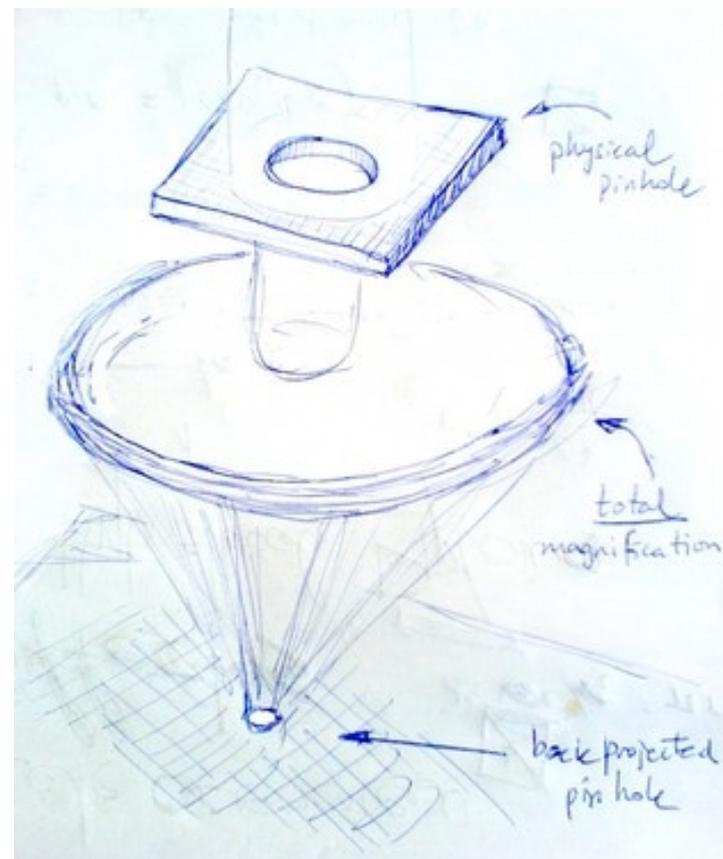
*b: Offset Function*

*I: Image Matrix*

*N: Noise Function*

$$N(\mathbf{PSF}(x, y, z) \otimes \mathbf{f}(x, y, z) + \mathbf{b}(x, y, z)) = \mathbf{I}(x, y, z)$$

**Backprojected  
confocal pinhole**



<http://support.svi.nl/wiki/NyquistCalculator>

*PSF: Point Spread Function*

*f: Object Function*

*b: Offset Function*

*I: Image Matrix*

*N: Noise Function*

$$N(\mathbf{PSF}(x, y, z) \otimes \mathbf{f}(x, y, z) + \mathbf{b}(x, y, z)) = \mathbf{I}(x, y, z)$$

### Biorad

- [Biorad MRC 500, 600 and 1024](#)
- [Biorad Radiance](#)

### Leica

- [Leica confocals TCS 4d, SP1 and NT](#)
- [Leica confocal SP2](#)
- [Leica confocal SP5](#)

### Nikon

- [TE2000-E with the C1 scanning head](#)

### Olympus

- [Olympus FV300](#)
- [Olympus FV500](#)
- [Olympus FV1000](#)

### Zeiss

- [Zeiss LSM410 inverted](#)
- [Zeiss LSM510](#)



Literature: eg. Noise Theory and Application to Physics: Philippe Réfrégier, Springer

**PSF:** Point Spread Function

**f:** Object Function

**b:** Offset Function

**I:** Image Matrix

**N:** Noise Function

- Black Body Irradiation  
(Poisson)

- Detector Noise  
(Gauss)

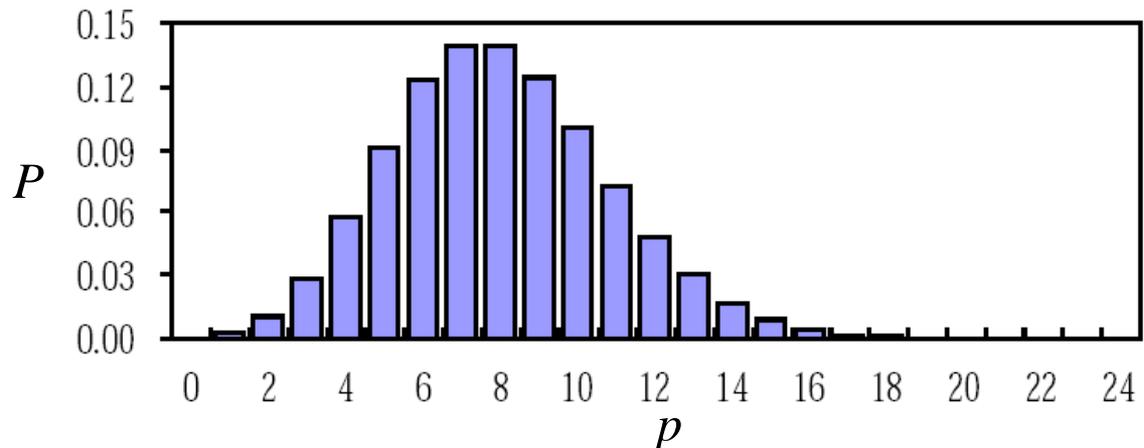
$$\mathbf{N}(\mathbf{PSF}(x, y, z) \otimes \mathbf{f}(x, y, z) + \mathbf{b}(x, y, z)) = \mathbf{I}(x, y, z)$$

$$P(p, \mu) = \frac{\mu^p}{p!} \cdot e^{-\mu}$$

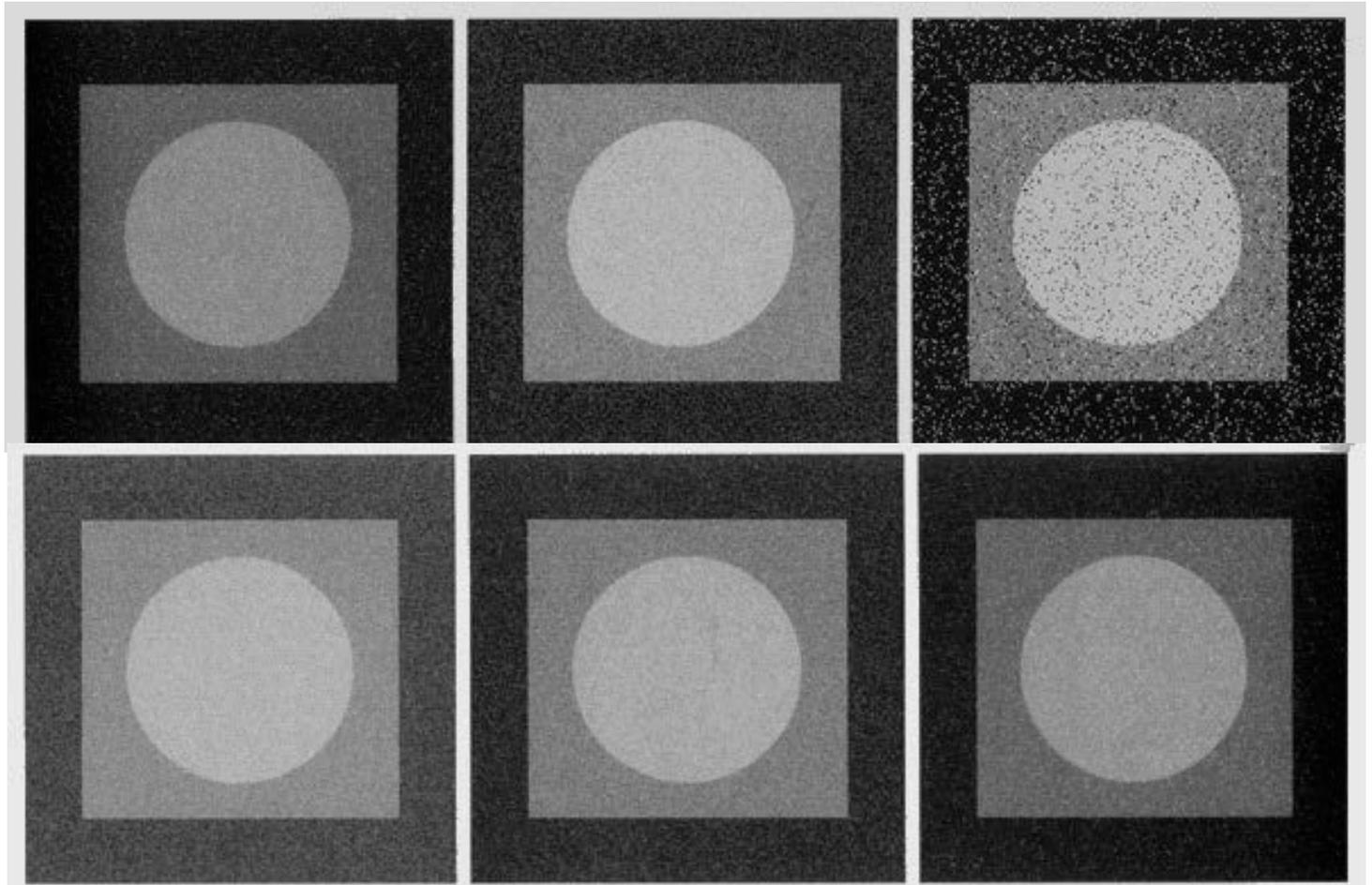
$$1. \bar{p} = \mu = \sigma^2, sd = \sigma = \sqrt{\bar{p}} = \sqrt{\mu}$$

$$2. \text{counting} : \bar{p} \pm \sqrt{\bar{p}}$$

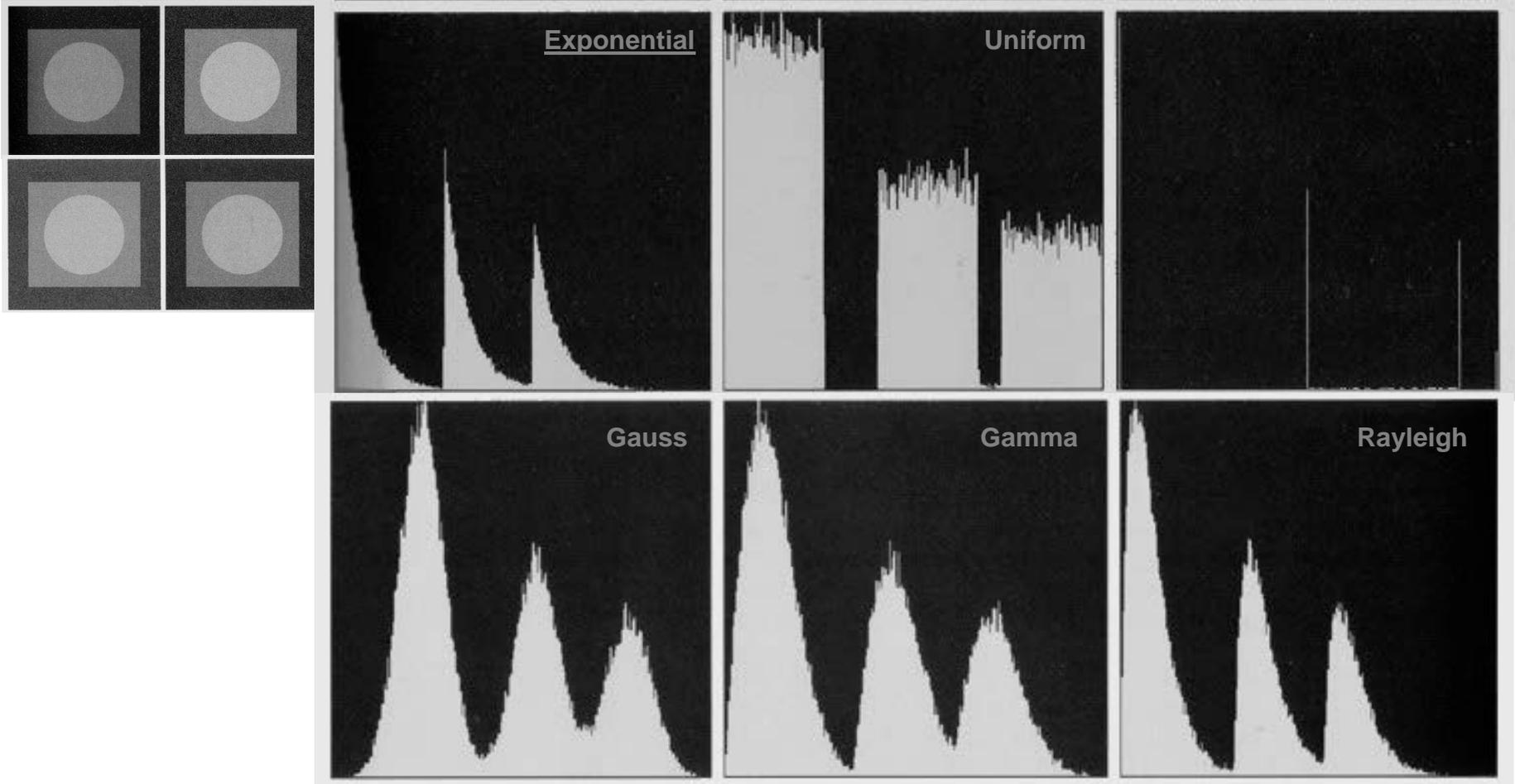
$$3. \text{Poisson}(\text{discrete}) \rightarrow \text{Gauss}(\text{continuous}) : \mu \rightarrow \infty$$



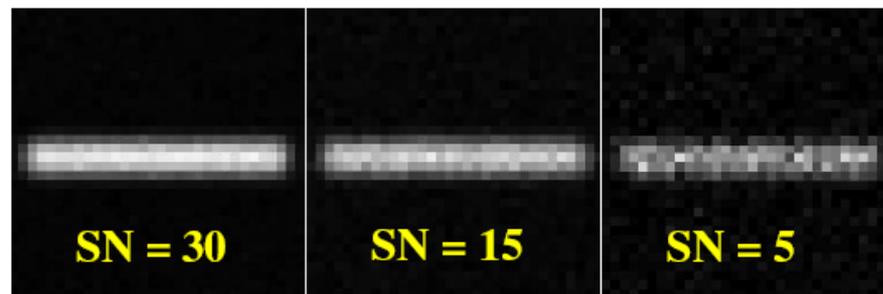
# | -> Noise



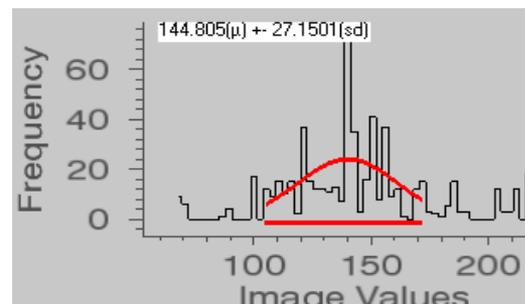
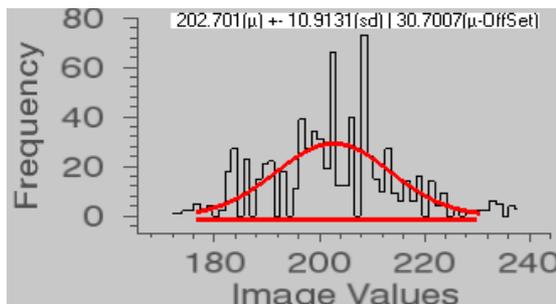
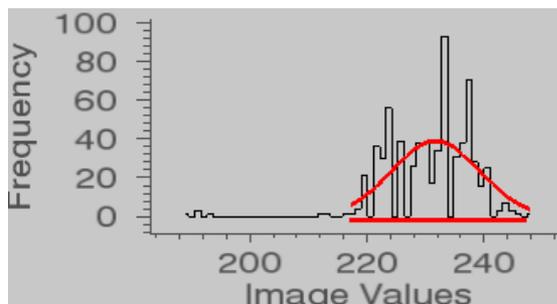
# | -> Noise



The Signal to Noise ratio (SN) is a number not always easy to estimate. The easiest way to obtain some figures is to look at the textures of bright areas in your object image. In the figure at left you see examples of such textures obtained from originally the same object image to which various levels of poisson noise were added.



**Figure 9. Images with different generated noise levels**

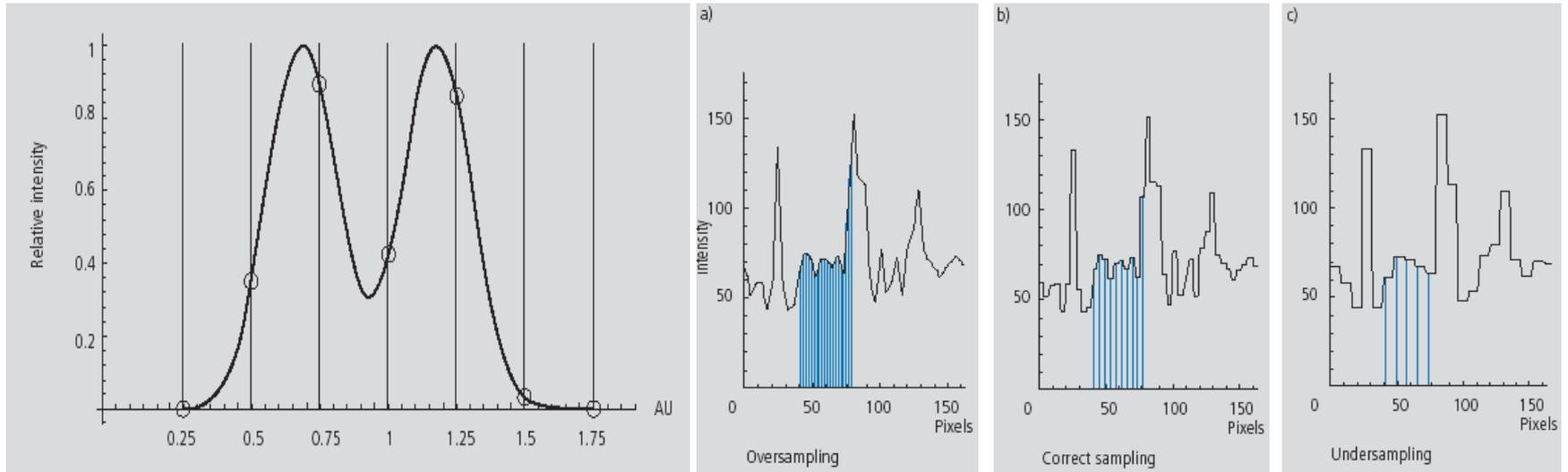


$$SNR = \frac{\bar{I}}{\sigma} = \frac{\bar{I}}{\sqrt{\sigma^2}} = \frac{229}{7.5}$$

$$SNR = \frac{\bar{I}}{\sigma} = \frac{\bar{I}}{\sqrt{\sigma^2}} = \frac{200}{10}$$

$$SNR = \frac{\bar{I}}{\sigma} = \frac{\bar{I}}{\sqrt{\sigma^2}} = \frac{139}{27}$$

# | -> Nyquist /Shannon Theorem



- Undersampling loses structures.
- Oversampling waists memory/computation time.

The 'Nyquist /Shannon Theorem' or 'Sampling Theorem' for the digital sampling of analogue signals suggests a Nyquist rate  $NR \geq 2v$  ?

**! Diffraction theory calculates lateral NR ~ 20 pixel/ $\mu\text{m}$ (~50 nm/pixel) !**  
**... axial NR ~ (~150 nm/pixel)**

*PSF: Point Spread Function*

*f: Object Function*

*b: Offset Function*

*I: Image Matrix*

*N: Noise Function*

$$N(\text{PSF}(x, y, z) \otimes f(x, y, z) + b(x, y, z)) = I(x, y, z)$$



## Calculator

[Numerical aperture](#)

[Excitation wavelength](#)

 (nm)

[Emission wavelength](#)

 (nm)

[Number of excitation photons](#)

[Backprojected pinhole radius](#)

 (nm)

[B.P. distance between pinholes](#)

 Only for Nipkow disks (μm)

[Lens medium refractive index](#)

[Specimen medium refractive index](#)

[Acquisition depth](#)

 (μm)

Calculate also PSF

- confocal
- widefield
- nipkow
- 4Pi

Select one

**PSF:** Point Spread Function

**f:** Object Function

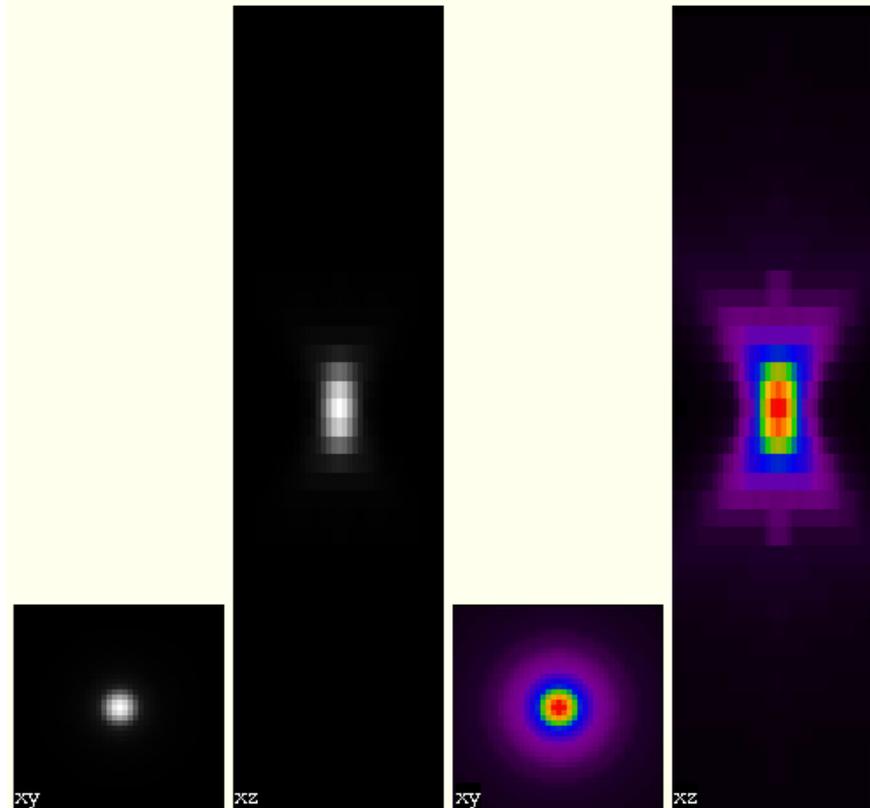
**b:** Offset Function

**I:** Image Matrix

**N:** Noise Function

$$N(\mathbf{PSF}(x, y, z) \otimes \mathbf{f}(x, y, z) + \mathbf{b}(x, y, z)) = \mathbf{I}(x, y, z)$$

[Nyquist sampling](#) (x,y,z in nm): 46, 46, 165



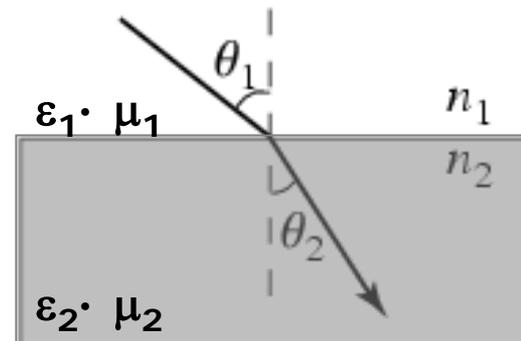
**Index of refraction:**  $n = (\epsilon \cdot \mu)^{1/2} = c/v,$

$\epsilon$  electric permittivity and  $\mu$  magnetic permeability.

## Snell's Law:

$$\sin \theta_1 n_1 = \sin \theta_2 n_2$$

- 1.518 [Zeiss Oil]
- 1.33 [Water]
- 1.0008 [Air]



Refractive Index:

$$RI = n_1/n_2 = v_2/v_1$$

Snell's Law:

$$\sin\theta_1 n_1 = \sin\theta_2 n_2$$

$$n = n(\lambda) !$$

- 1.518 [Zeiss]
- 1.33 [Water]
- 1.0008 [Air]

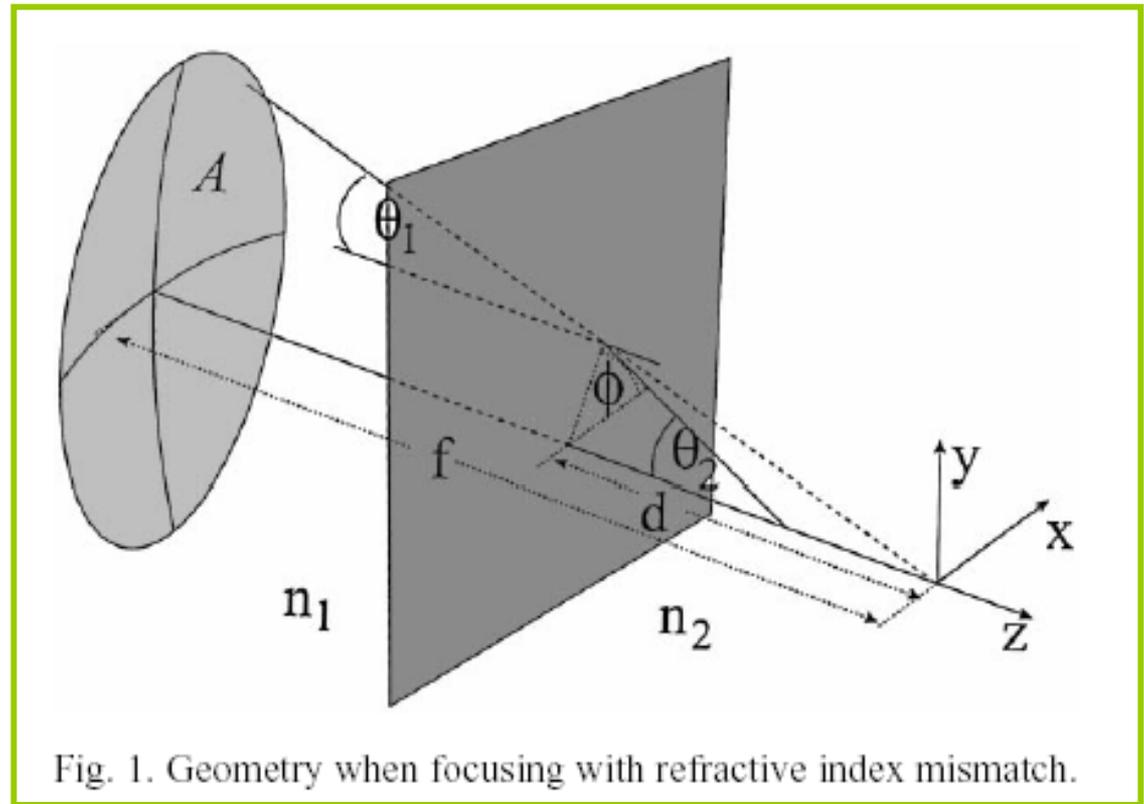
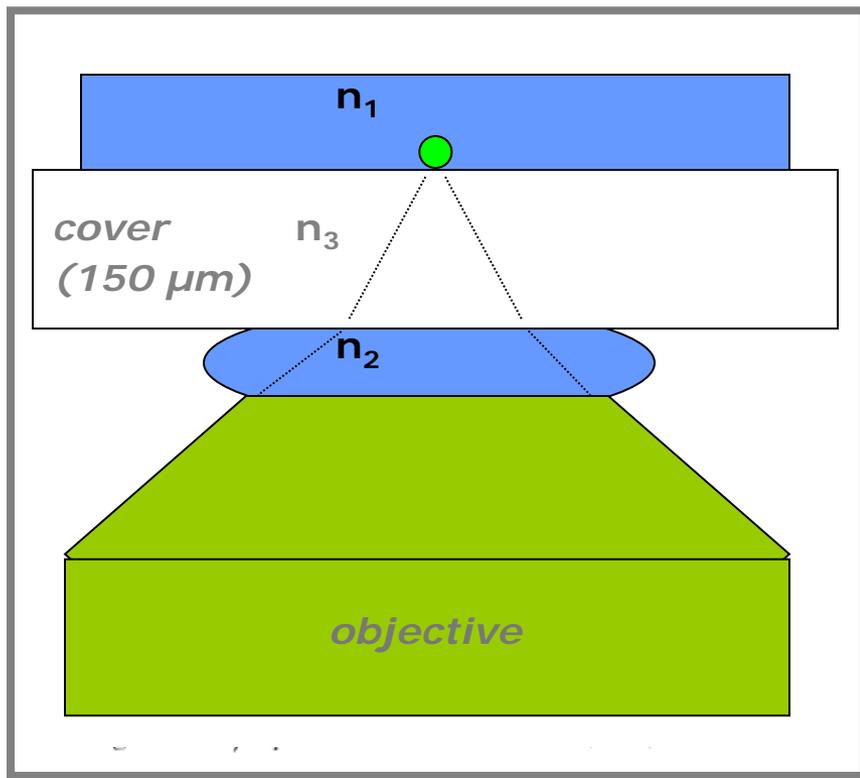


Fig. 1. Geometry when focusing with refractive index mismatch.

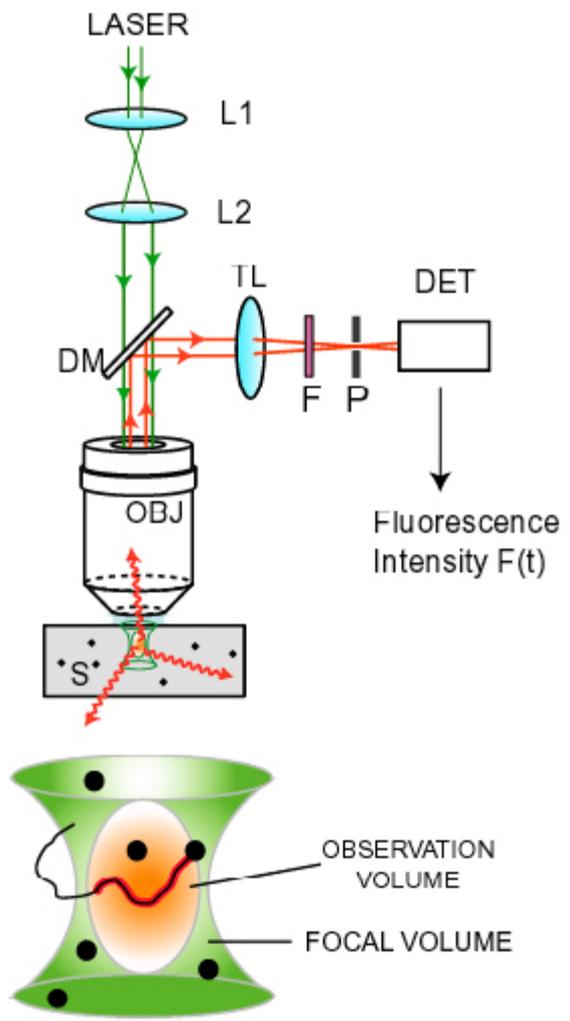
(Egner et al 1998)

● Micro-esfera:  $\varnothing = 6 \mu\text{m}$



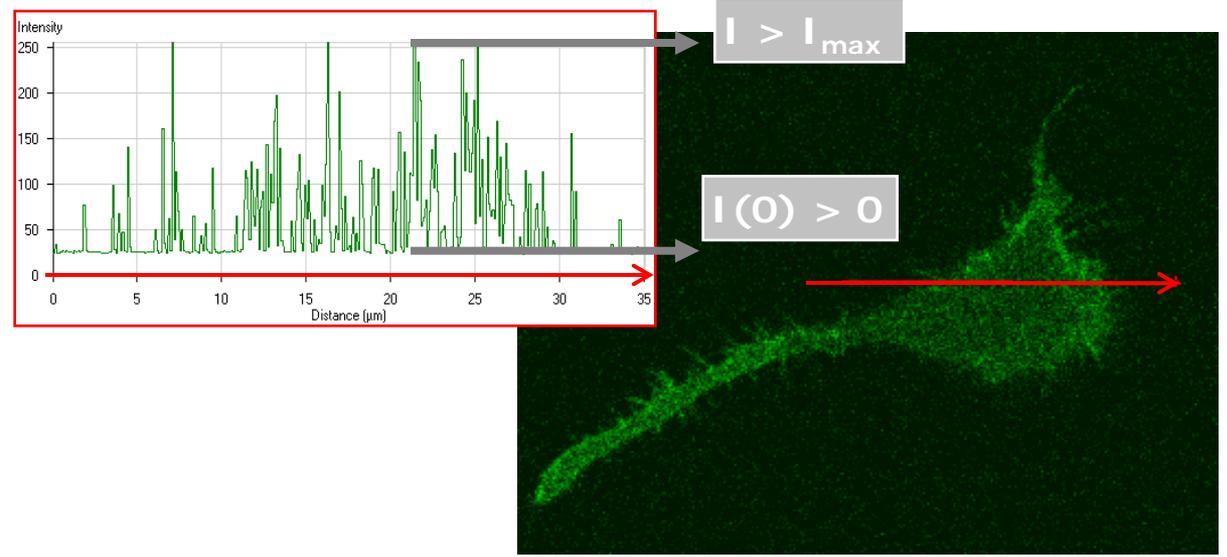
*agua/aceite* -- *aceite/aceite*  
 $n_1 \neq n_2$        $n_1 = n_2$

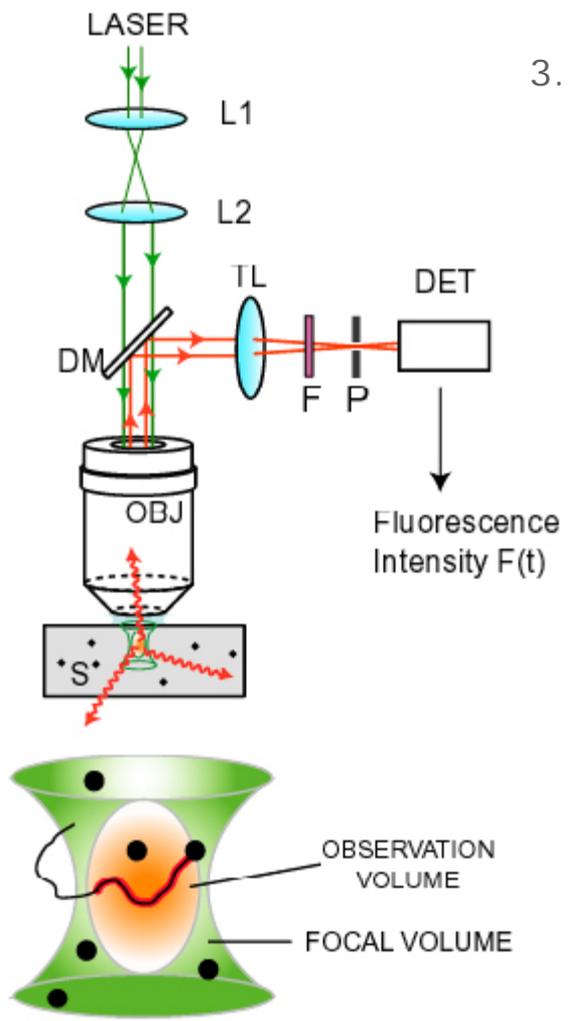
Ley de Snell:  $n_i \cdot \sin\theta_i = n_k \cdot \sin\theta_k$   
 $n = n(\lambda)$  !



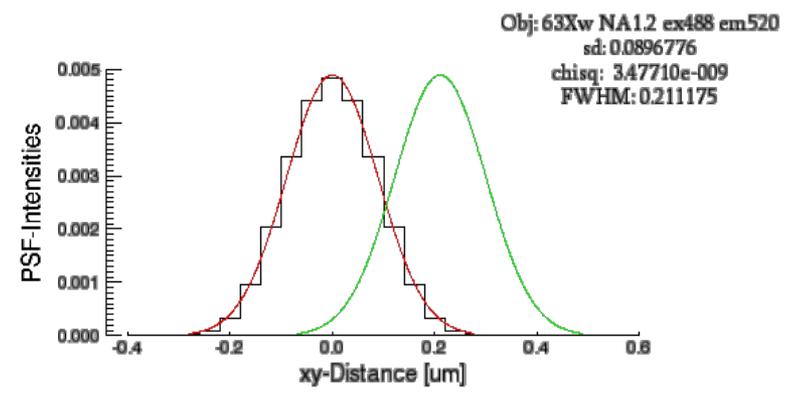
The observation volume (femtoliter) defined by the Point Spread Function must be considered as a mini-spectrofluorimeter.

1. You need to consider the Offset  $I(0)$  in order to calibrate your signal  $I(0) \geq 0$  !
2. Never saturate the signal:  $I \leq I_{max}$  (255 for 8 bit) !

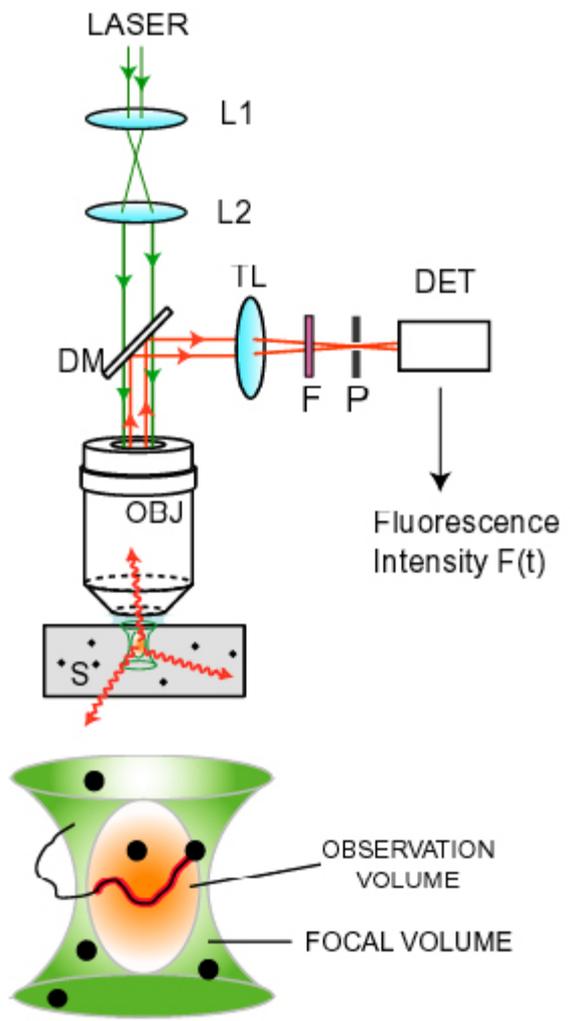




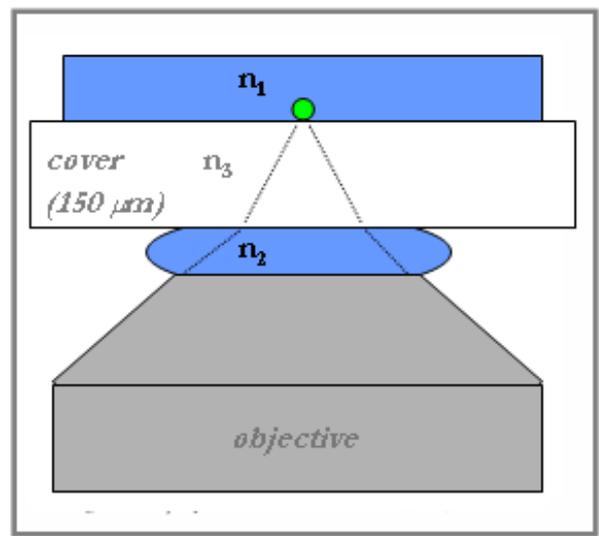
- You need to consider sampling distances in  $\Delta x$  and  $\Delta y \approx 50$  nm and  $\Delta z \approx 150-300$  nm for later deconvolution, or calculate the explicit sample distances @ <http://support.svi.nl/wiki/NyquistCalculator>



- Use the right immersion setup !  
 $n_1 = n_2$  !  
 Keep refractive index / index of refraction constant !



● Micro-esfera:  $\varnothing = 6 \mu\text{m}$



*agua/aceite* -- *aceite/aceite*  
 $n_1 \neq n_2$        $n_1 = n_2$   
 Ley de Snell:  $n_1 \cdot \sin\theta_1 = n_k \cdot \sin\theta_k$   
 $n = n(\lambda)$  !

