

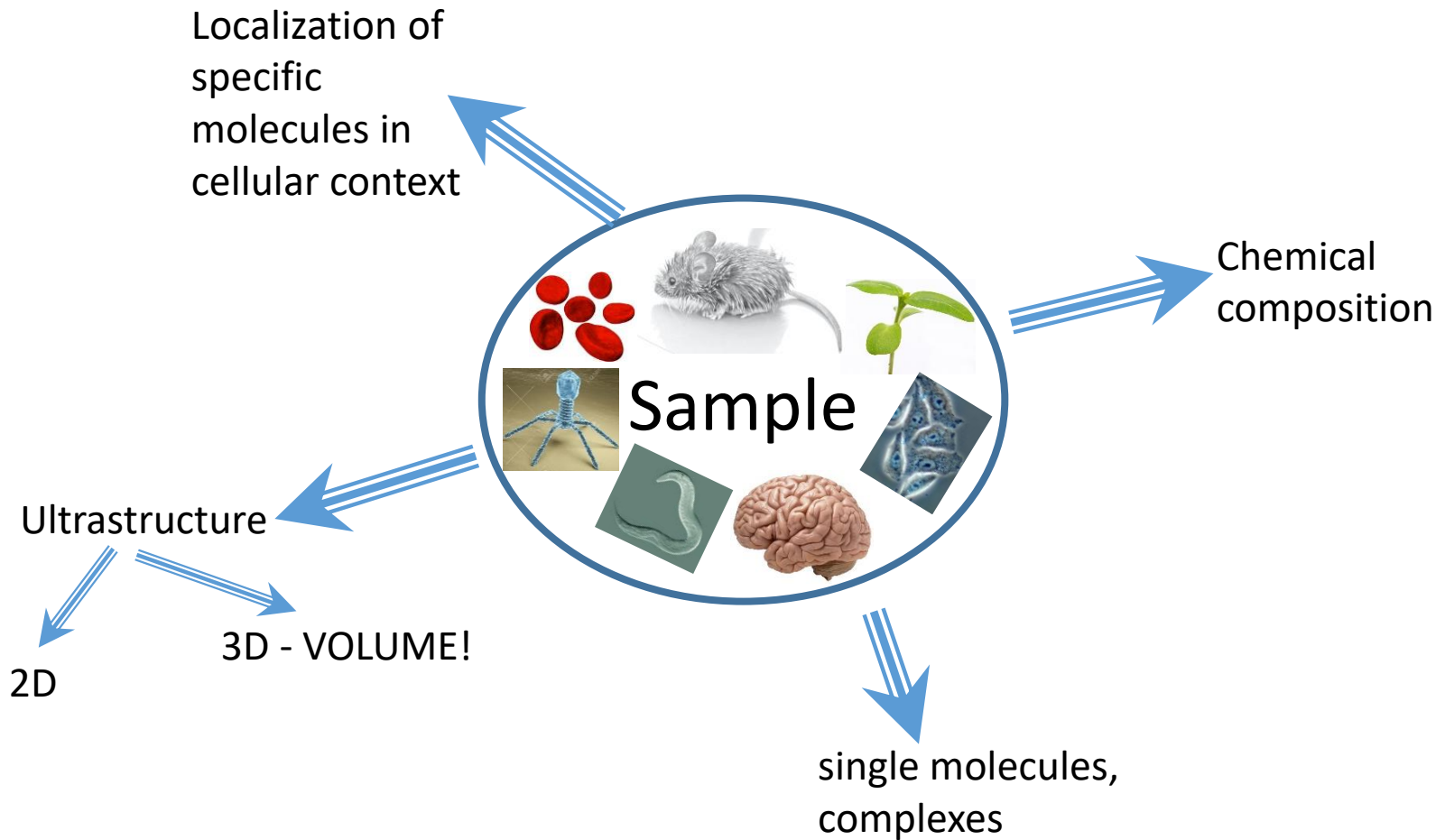
Advanced electron microscopy techniques

Vlada Filimonenko

Institute of Molecular Genetics

Electron Microscopy Core Facility



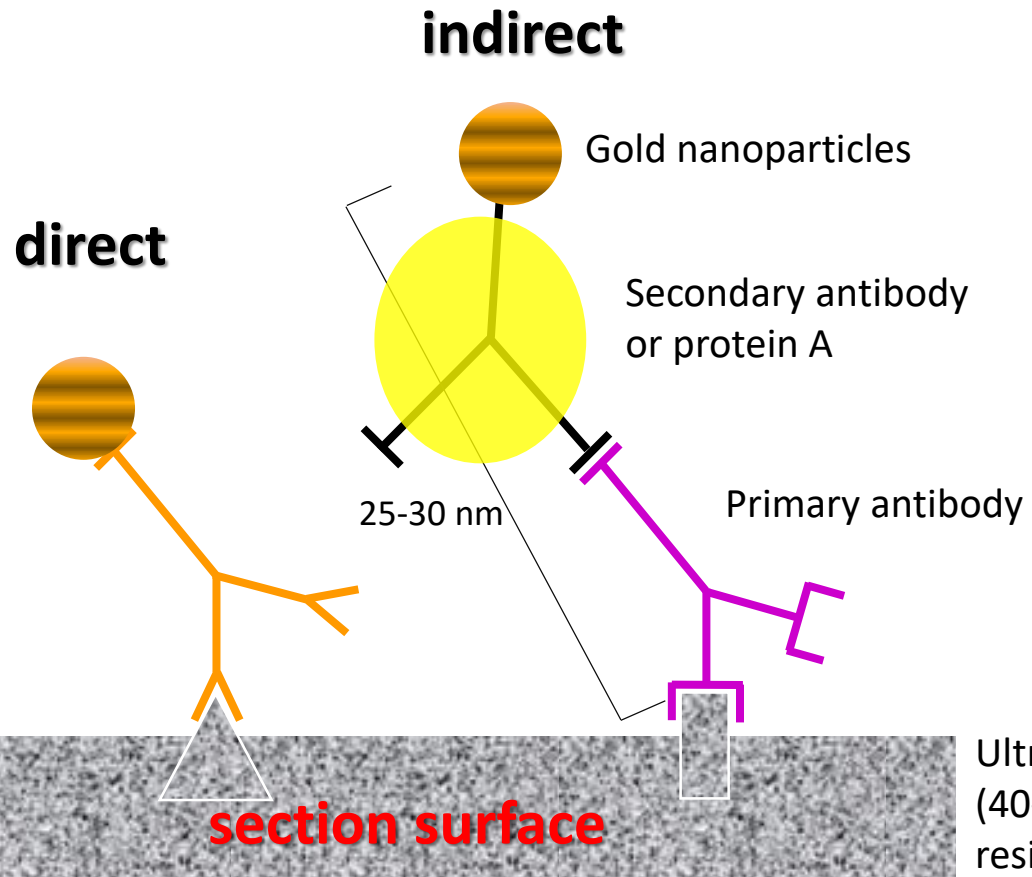
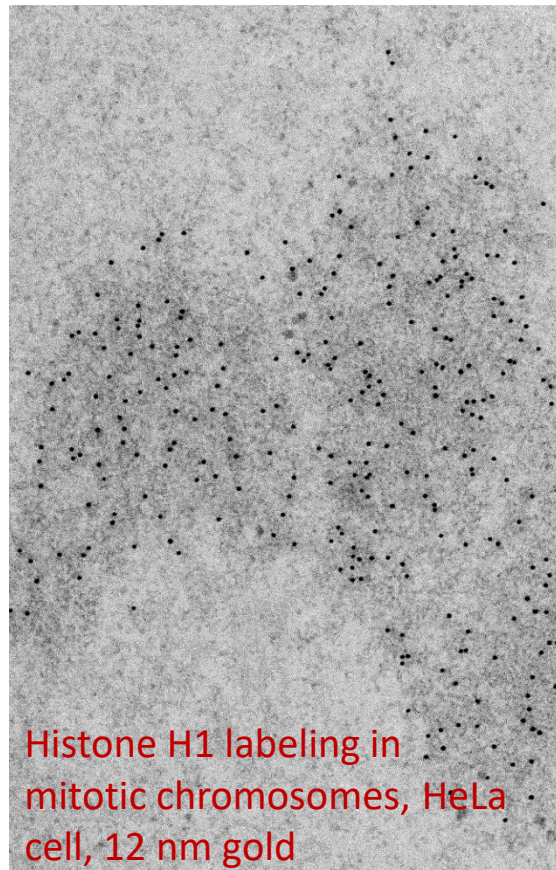


Correlative microscopy

CLEM, cryo-CLEM

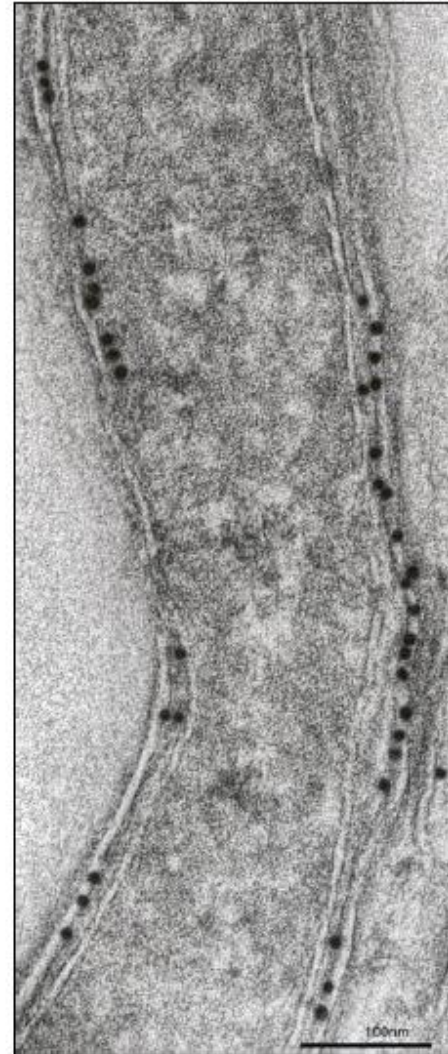
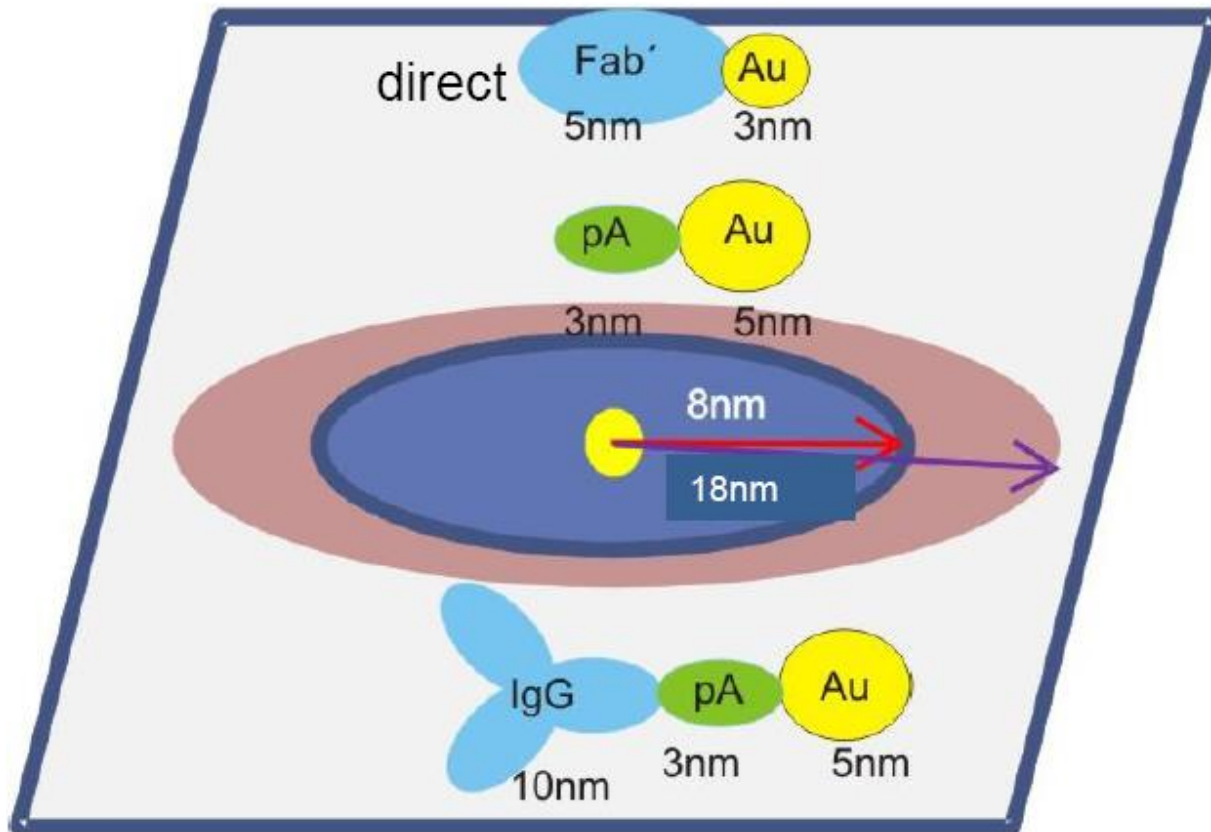
Labelling of specific molecules in cellular context

Immunogold labeling for electron microscopy



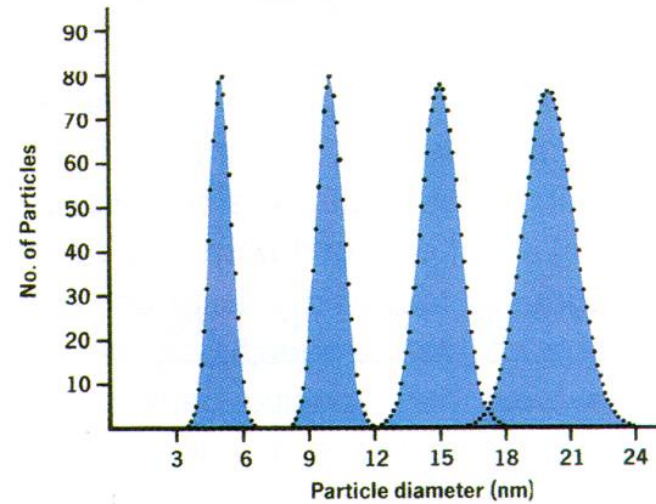
How can we improve the resolution?

antibody + gold

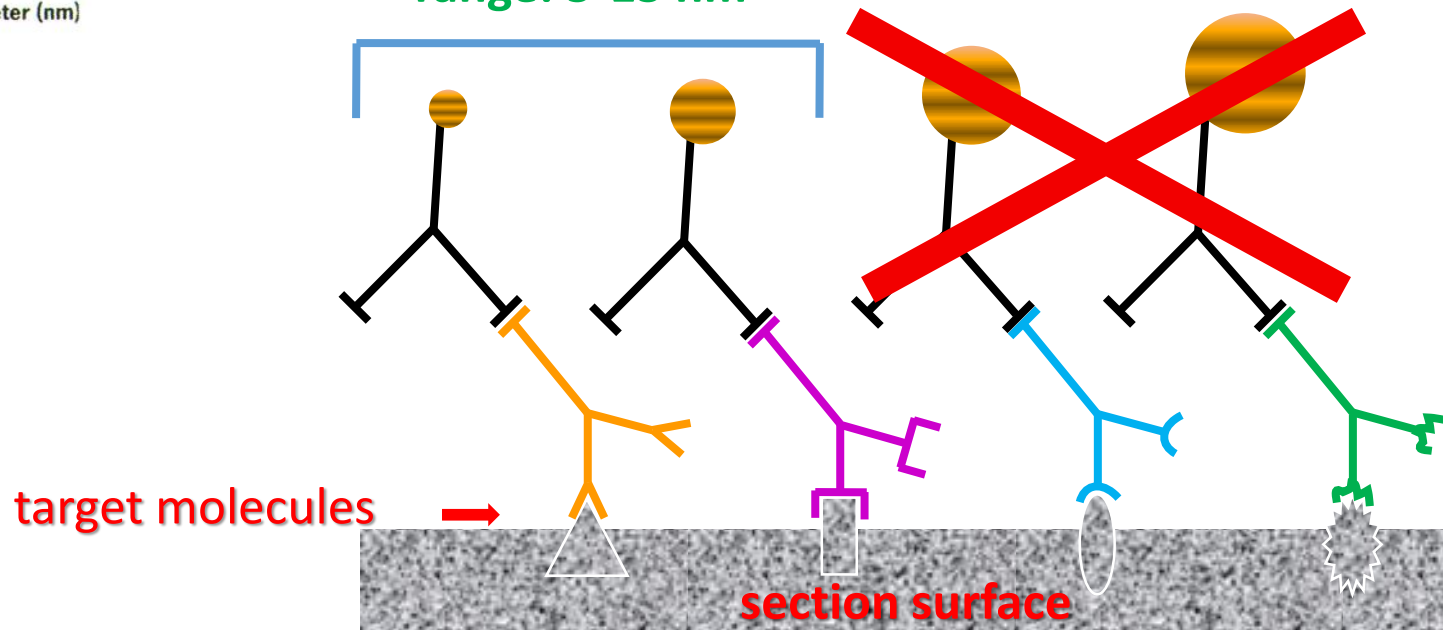


The best theoretical resolution: Fab fragments conjugated directly to very small Au particles giving „worstcase“- resolution of 8 nm radius
Specific IgG+ 5 nmAu/protein A = 18 nm

Gold particle size and multiple immunolabeling

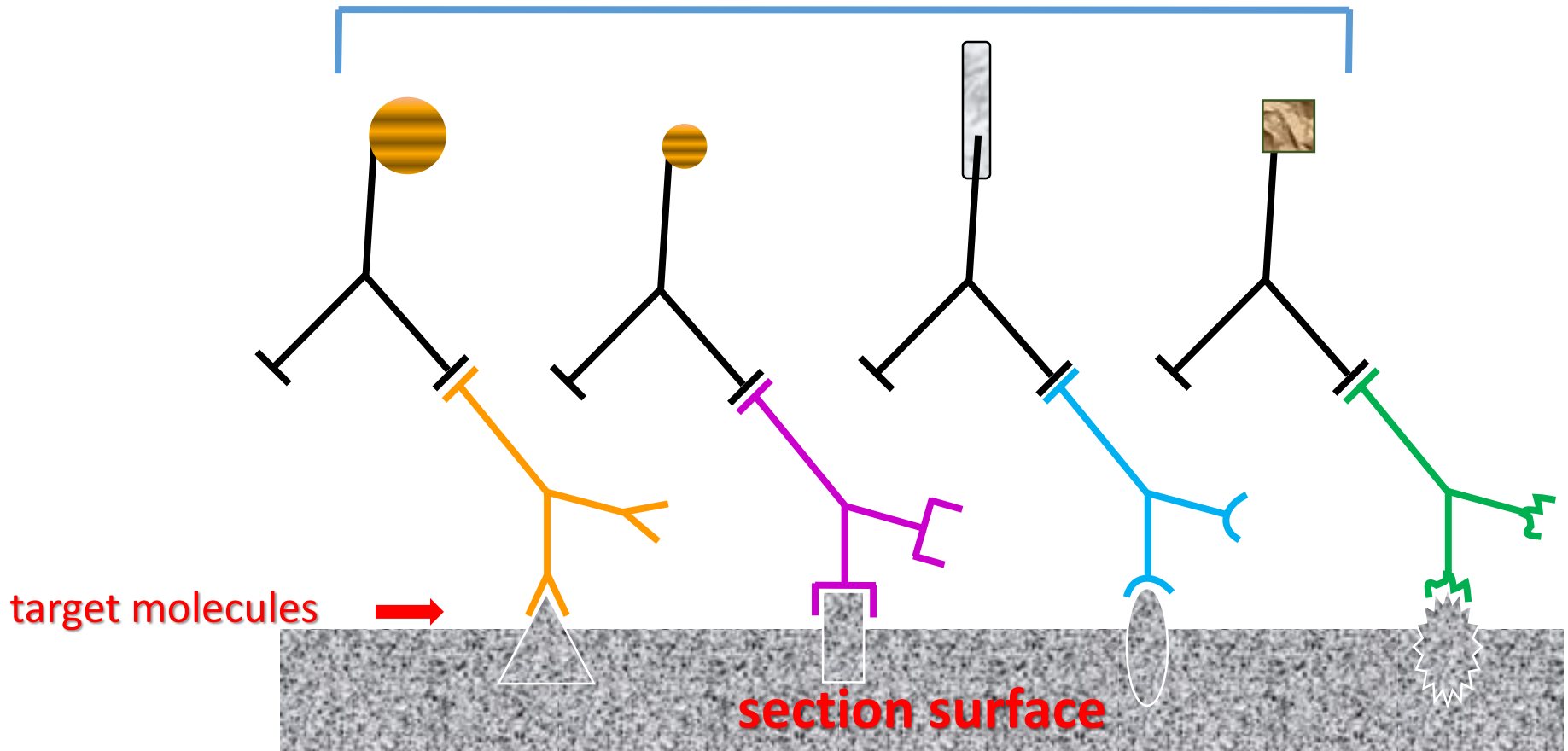


Optimal size
range: 5-15 nm

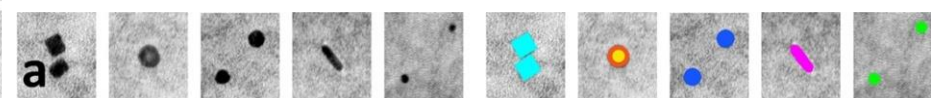
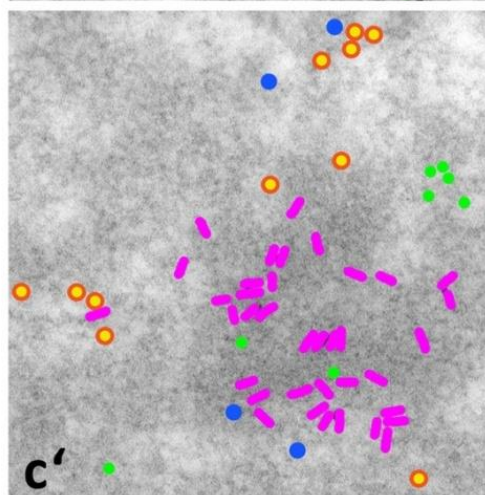
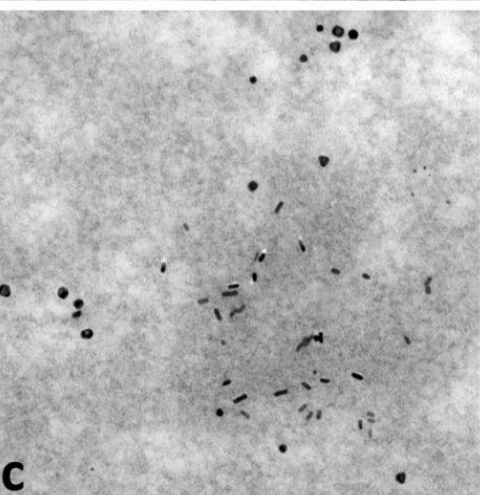
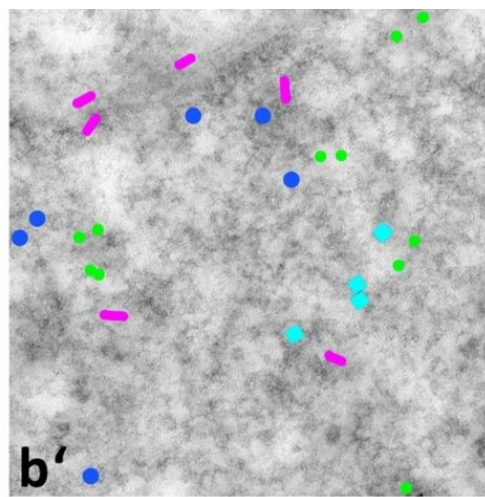
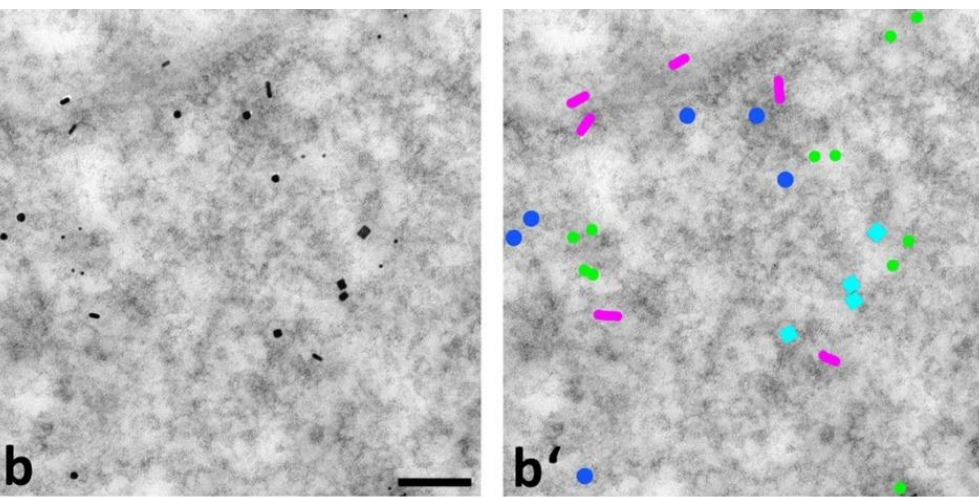


We need nanoparticles differing by other parameters

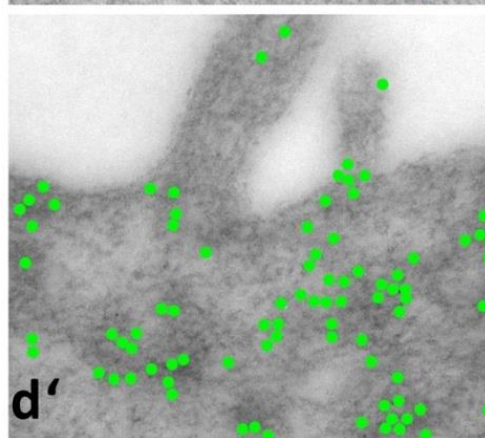
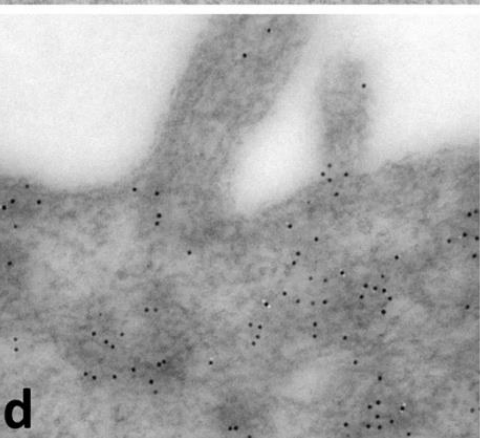
Optimal size range: 5-15 nm



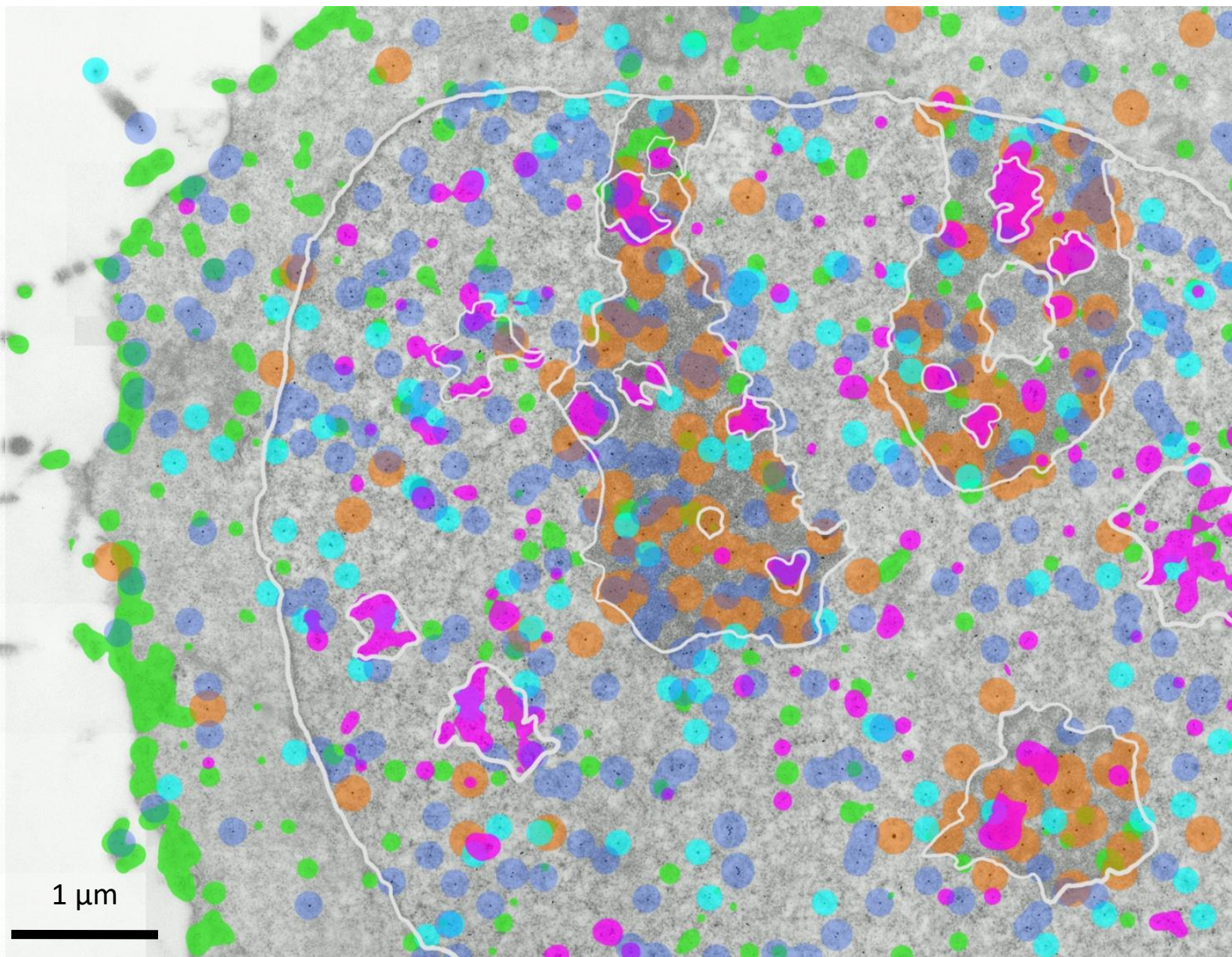
Immunolabeling of a biological sample



PIP2		AuNR
B23		AgAu
SMC2		Au12
Sm		PdC
actin		Au6



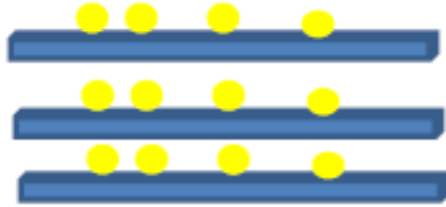
V. V. Philimonenko, A. A. Philimonenko, I. Šloufová, M. Hrubý, F. Novotný, Z. Halbhuber, M. Krivjanská, J. Nebesářová, M. Šlouf, and P. Hozák. Simultaneous detection of multiple targets for ultrastructural immunocytochemistry. *Histochem Cell Biol.* 2014; 141(3): 229–239



PIP2		AuNR
B23		AgAu
SMC2		Au12
Sm		PdC
actin		Au6

Methods of immunolabeling

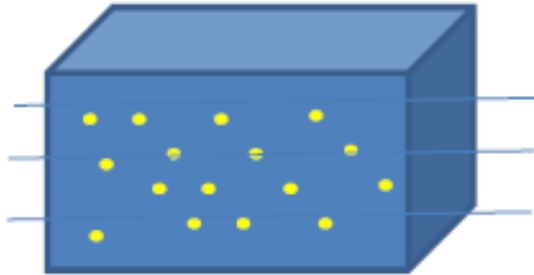
On-section



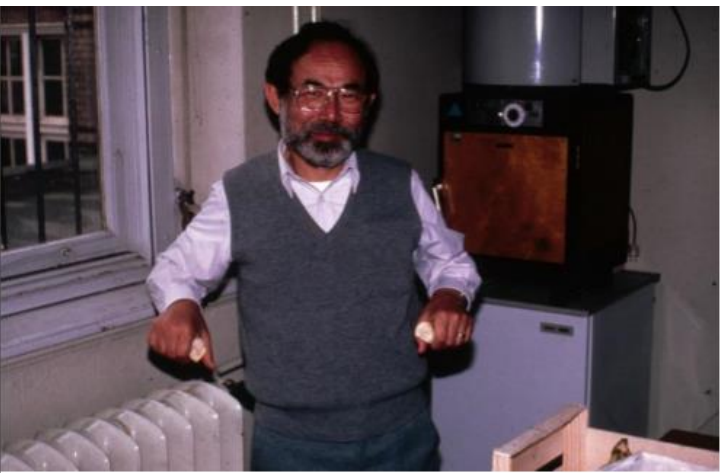
Post-embedding – on plastic section – lower efficiency

Tokuyasu method – thawed cryosections

In the volume

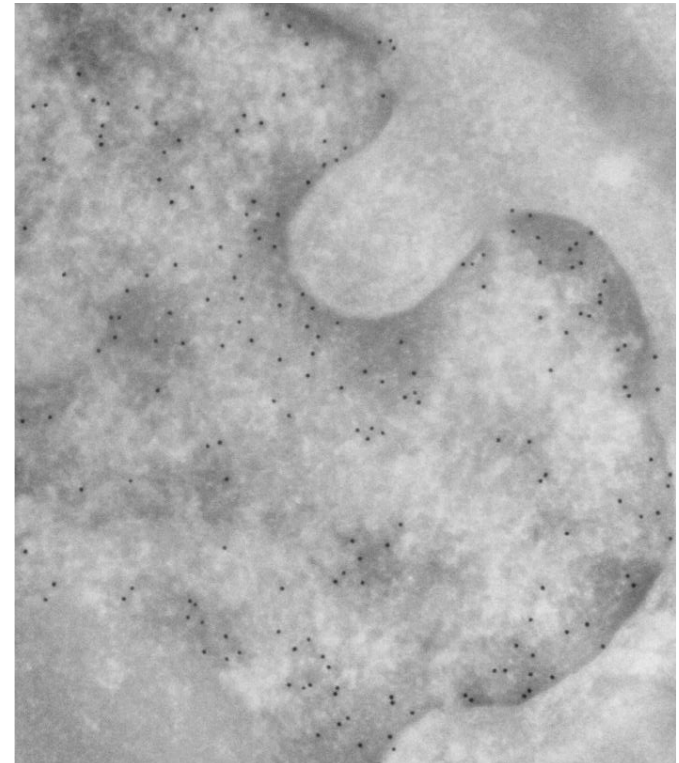
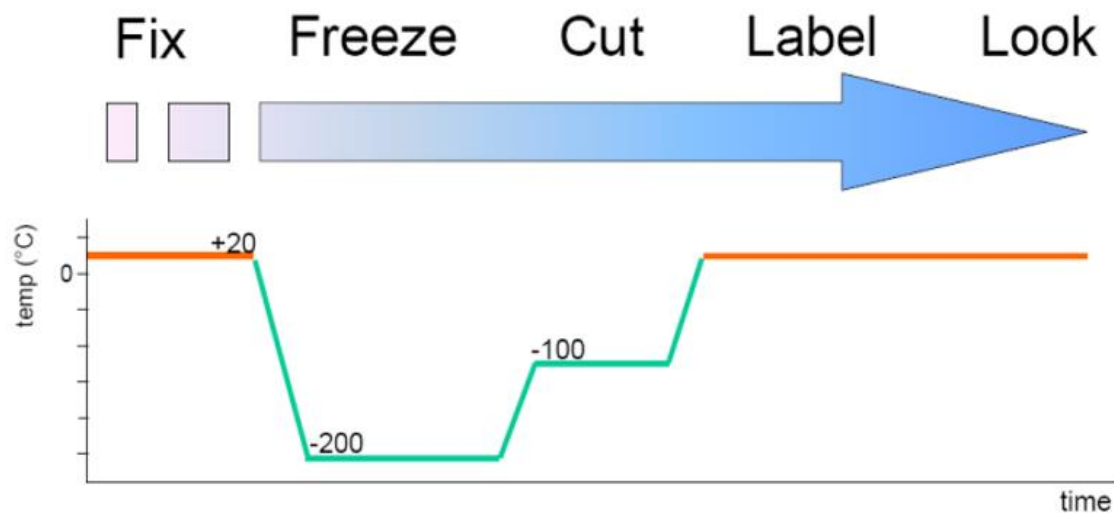


Pre-embedding – before embedding into the resin



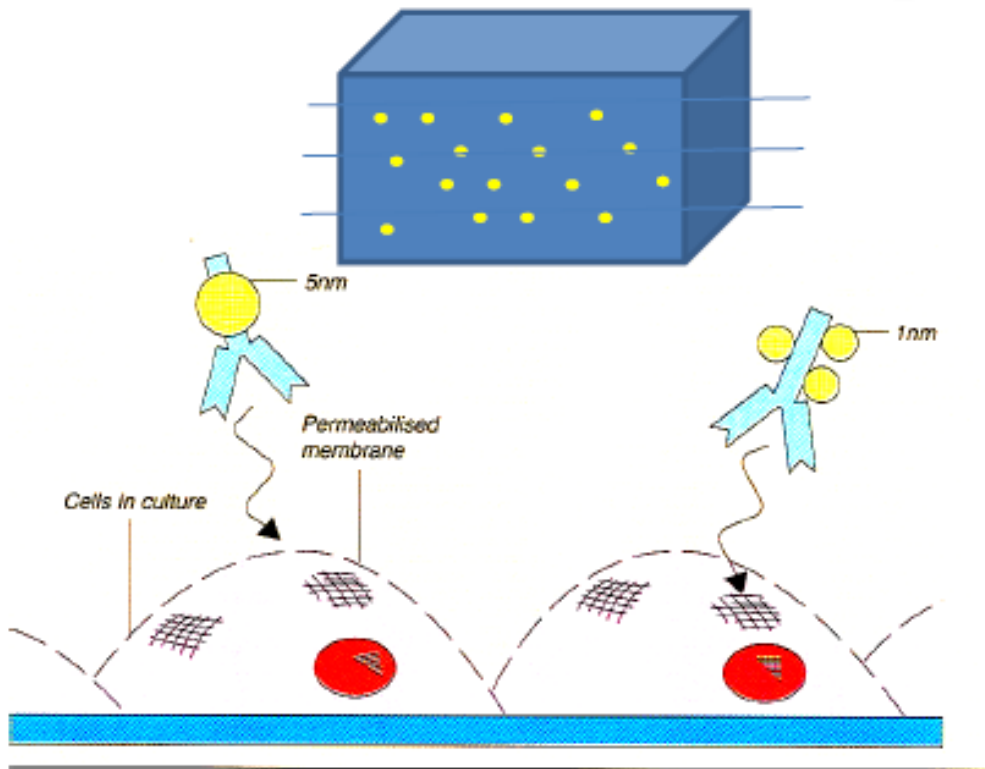
Tokuaysu technique

Tokuyasu, KT. 1973: A technique for ultracryotomy of cell suspensions and tissues. J Cell Biol. May;57(2):551-65,



Labelling of intracellular antigens

Pre-embedding



- + labelling throughout the whole section thickness
- + antigens are not denatured prior labelling

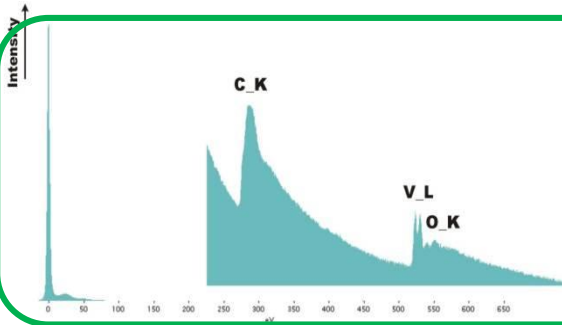
- antigen accessibility, permeabilization → cell ultrastructure, extraction of many components and risk of false negative results
- Larger gold conjugates ($>5\text{nm}$) diffuse unefficiently

HRP-coupled antibodies, GFP-anti GFP antibodies, fluorochrom/Ab, fluorescently conjugated Nanogold antibodies, ultrasmall gold conjugated / Ag enhancement

Analytical EM

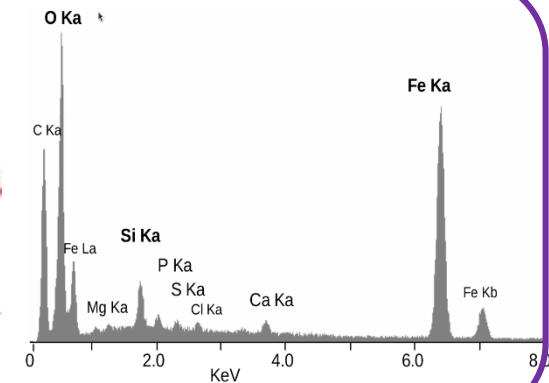
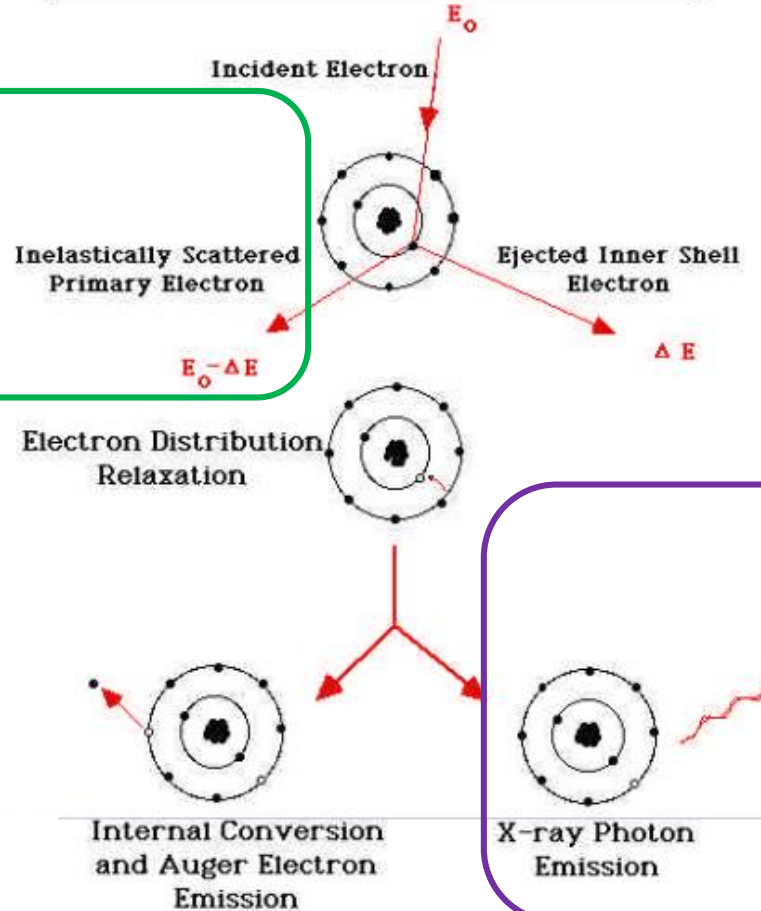
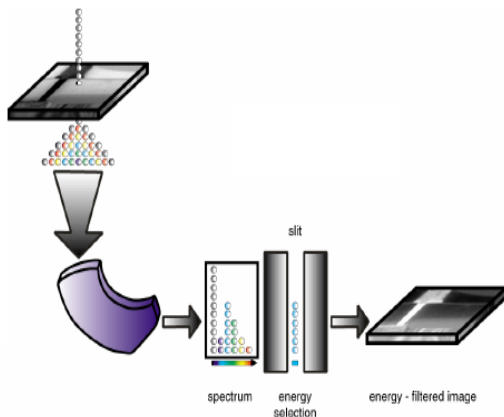
Electron Microscopy: element specific signals

Electron Excitation of Inner Shell Processes



EELS

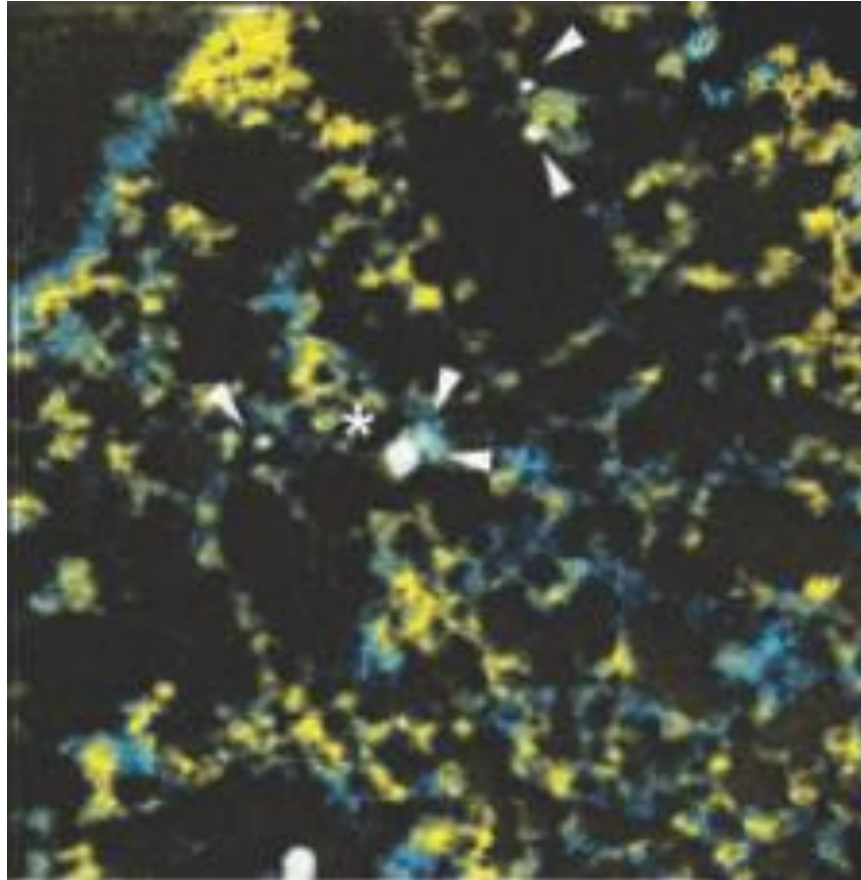
Electron-energy
loss spectroscopy



EDS

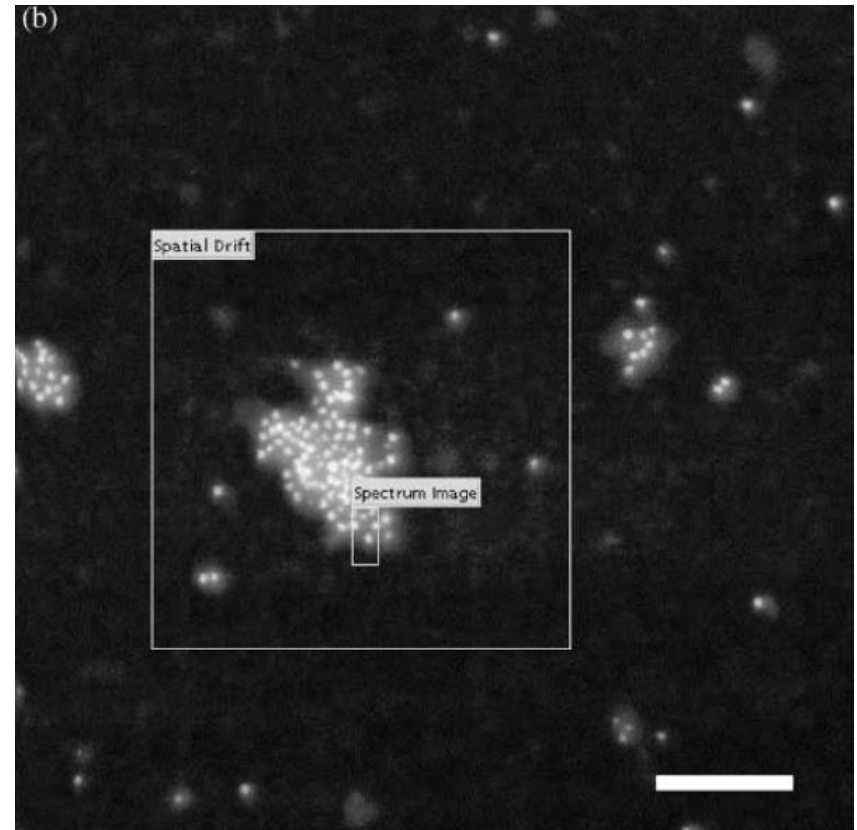
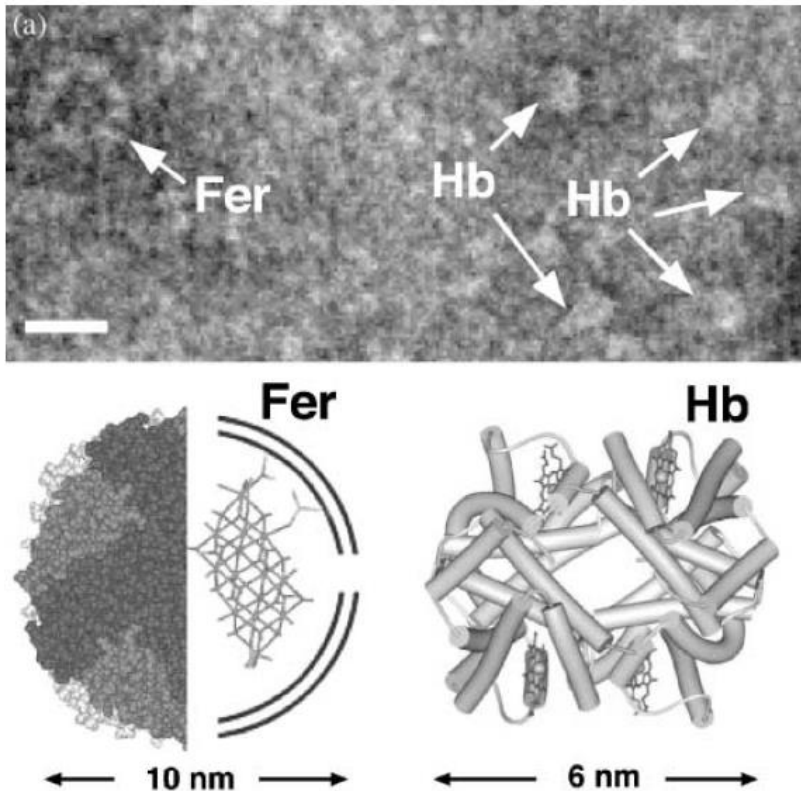
Energy-dispersive
X-ray spectroscopy

Mapping of protein-rich and nucleic-acid rich structures in the cell using EFTEM

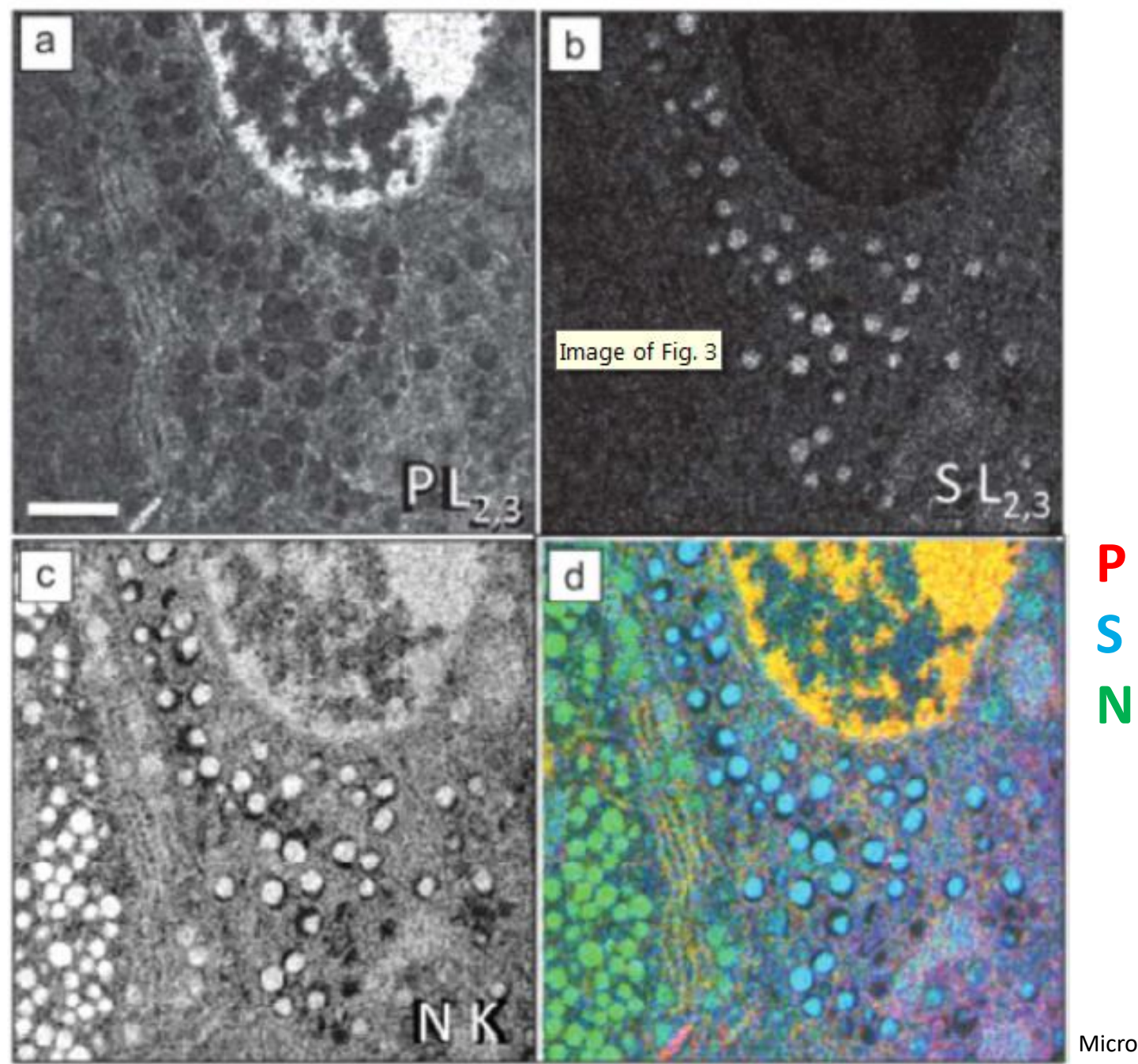


Ahmed K et al. PLoS One. 2010;5(5):e10531.

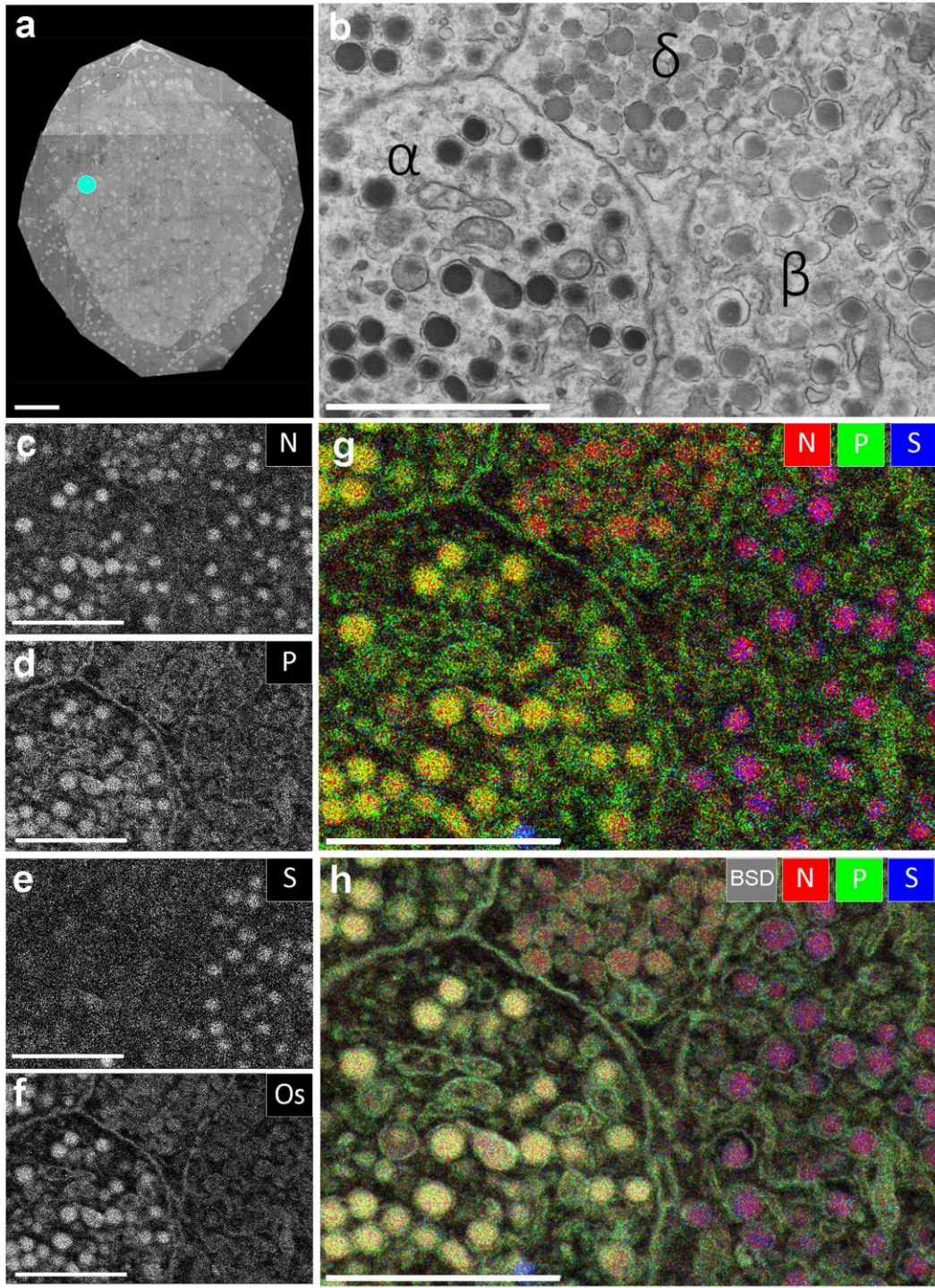
EELS mapping of single iron atoms in haemoglobin



EFTEM spectrum-imaging of unstained section of mouse pancreatic islet of Langerhans

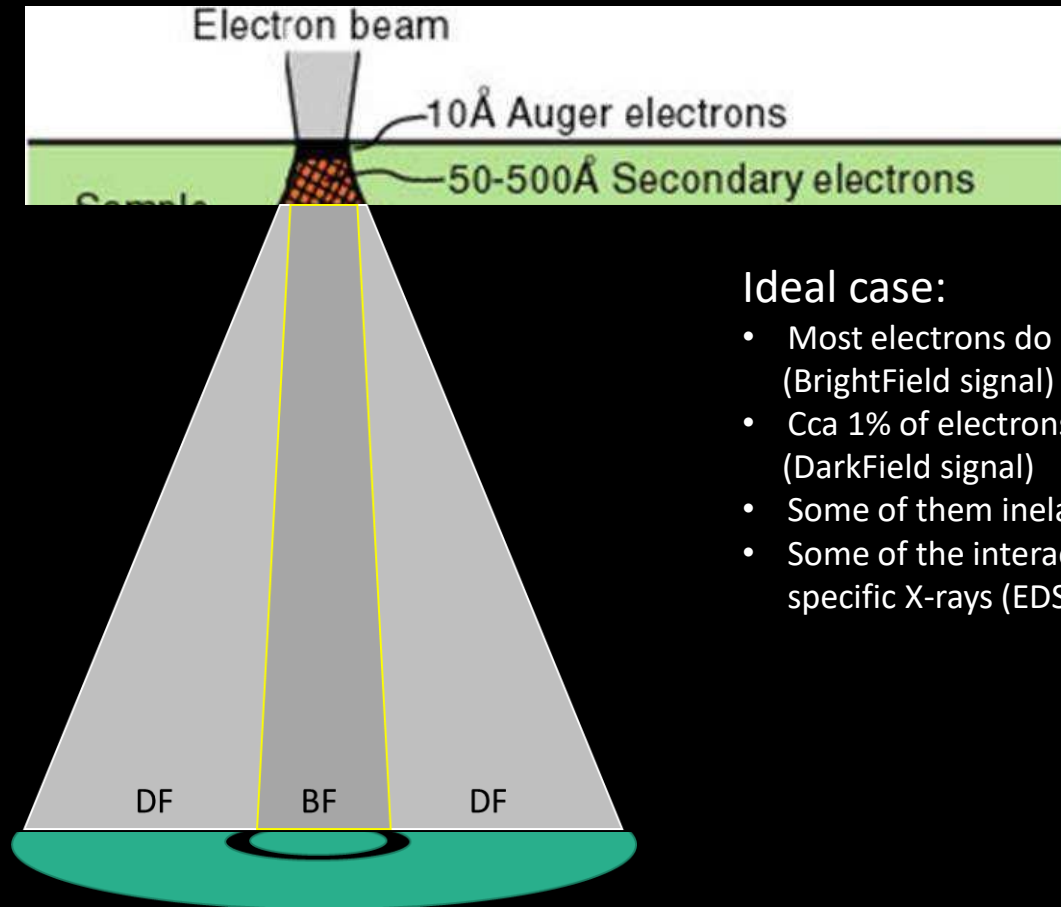


EDS mapping



mouse pancreatic
islet of Langerhans

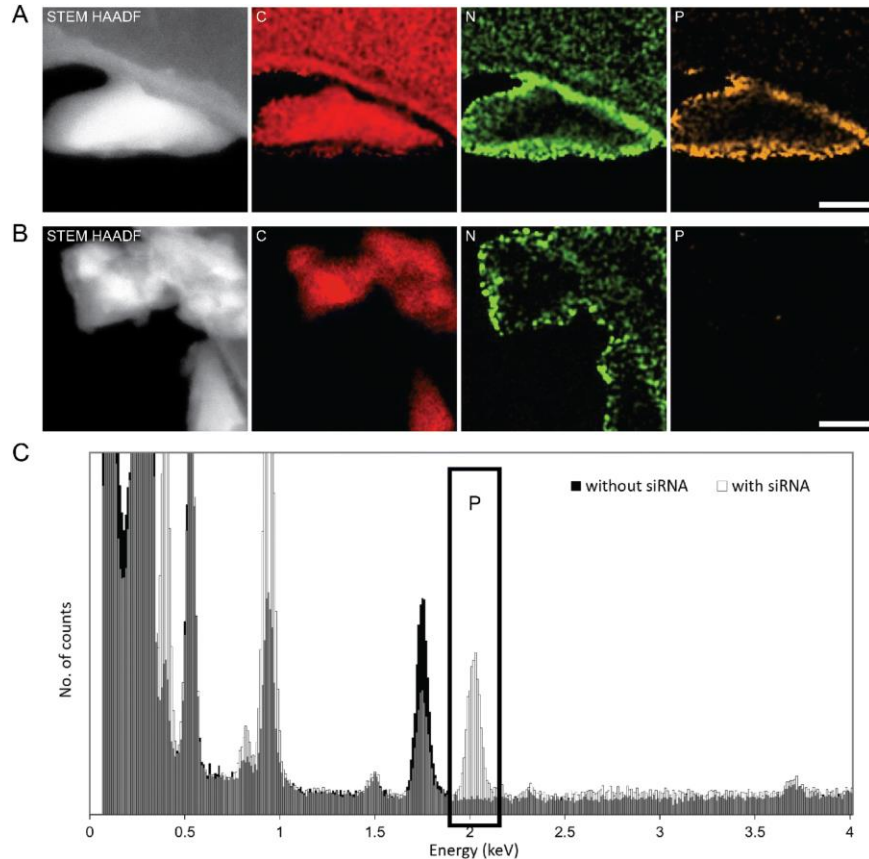
Spatial resolution of EDS



Ideal case:

- Most electrons do NOT interact at all (BrightField signal)
- Cca 1% of electrons interact once (DarkField signal)
- Some of them inelastically
- Some of the interactions produce specific X-rays (EDS signal)

Nanoparticle composition analysis by EDS



Nanodiamond particles with polymer coating to serve as siRNA delivery system

- Lacking direct proof for RNA binding to the polymer layer

EDS elemental mapping reveals:

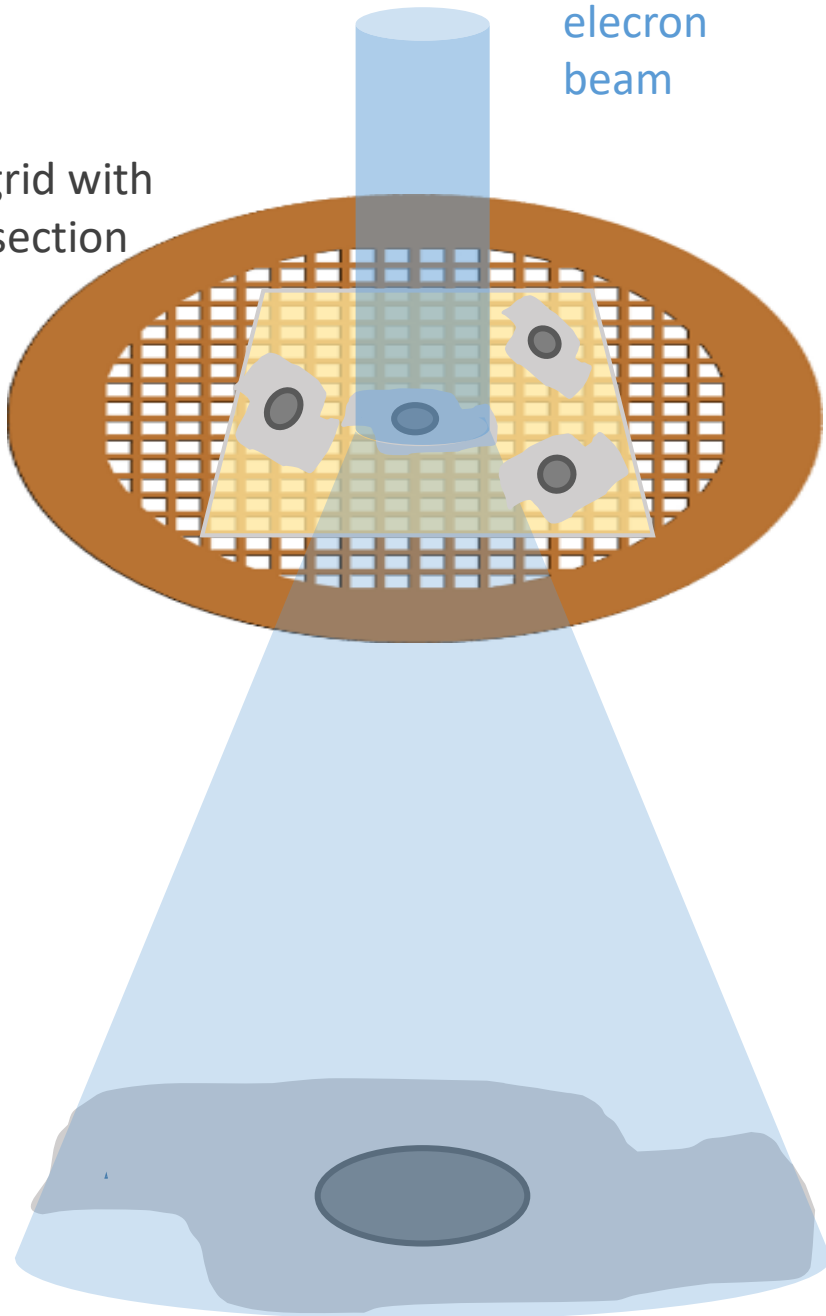
- Core-shell structure with N-containing polymer on surface
- Increase in N and presence of P after siRNA binding

Majer, J., M. Kindermann, D. Pinkas, D. Chvatil, P. Cigler and L. Libusova (2024). "**Cellular uptake and fate of cationic polymer-coated nanodiamonds delivering siRNA: a mechanistic study.**" *Nanoscale* **16**(5): 2490-2503.

Volume electron microscopy

electron
beam

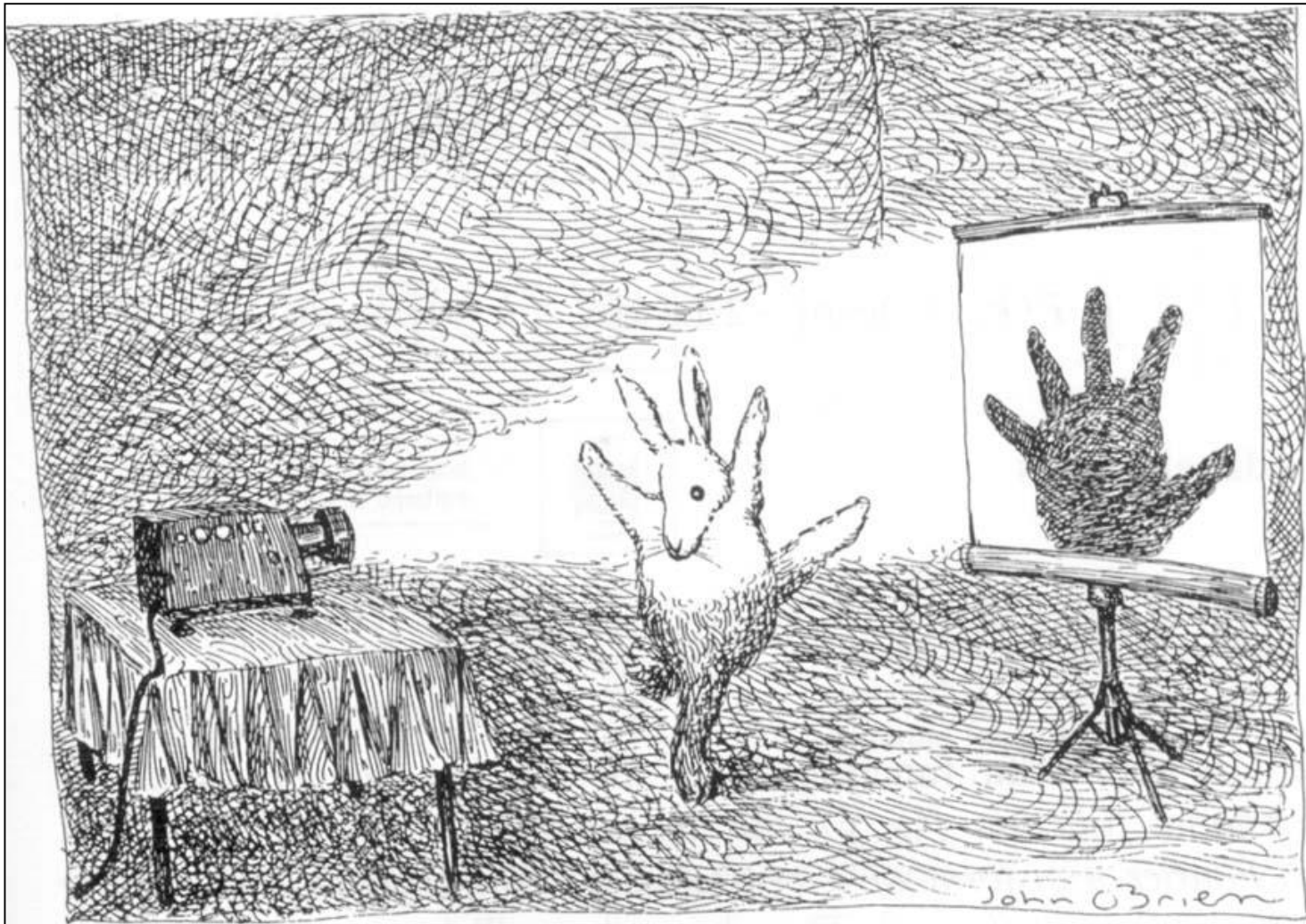
TEM grid with
resin section



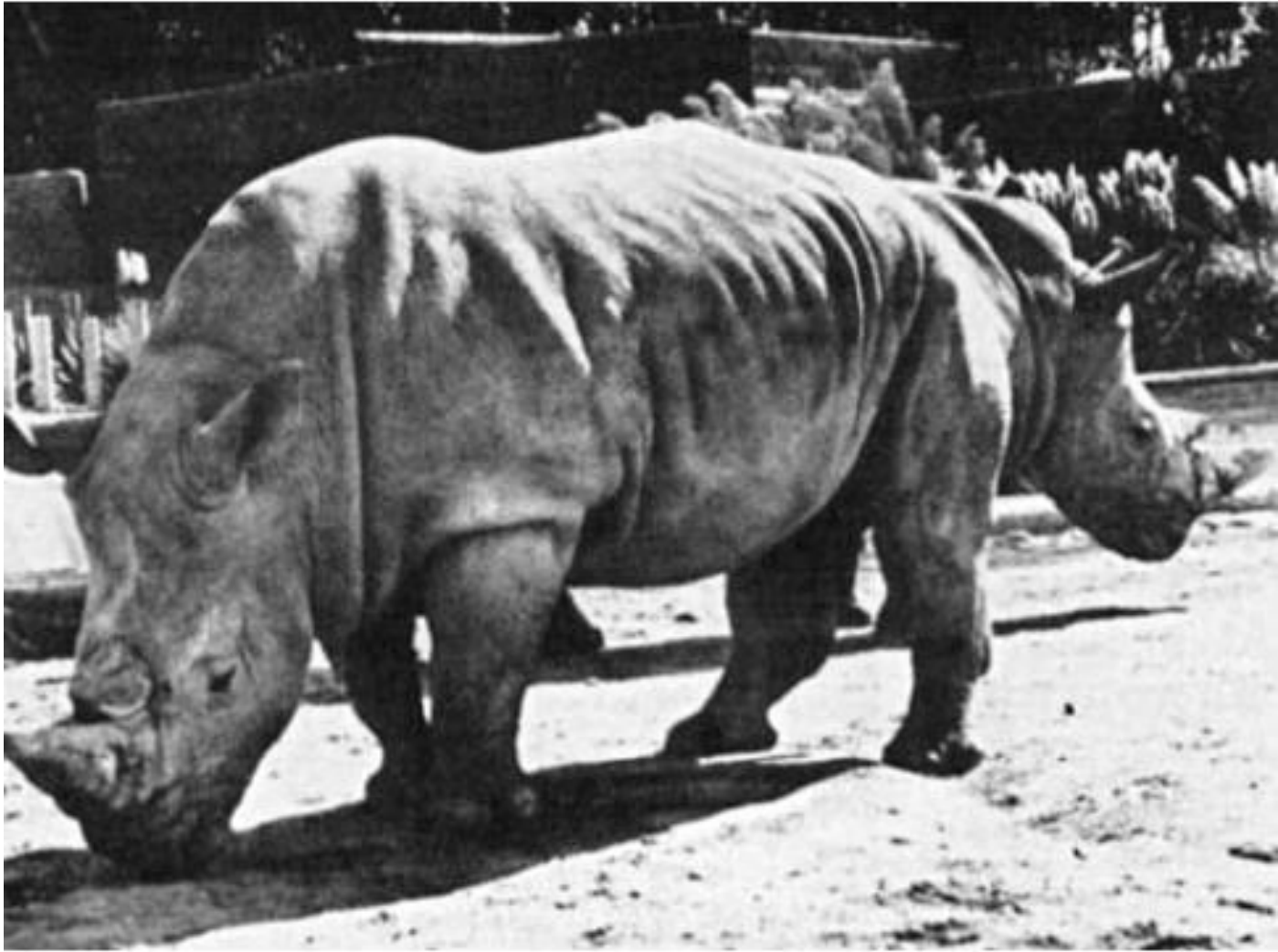
detector

Why do we need 3D information?

- Every TEM image is a 2D projection of a 3D/4D object

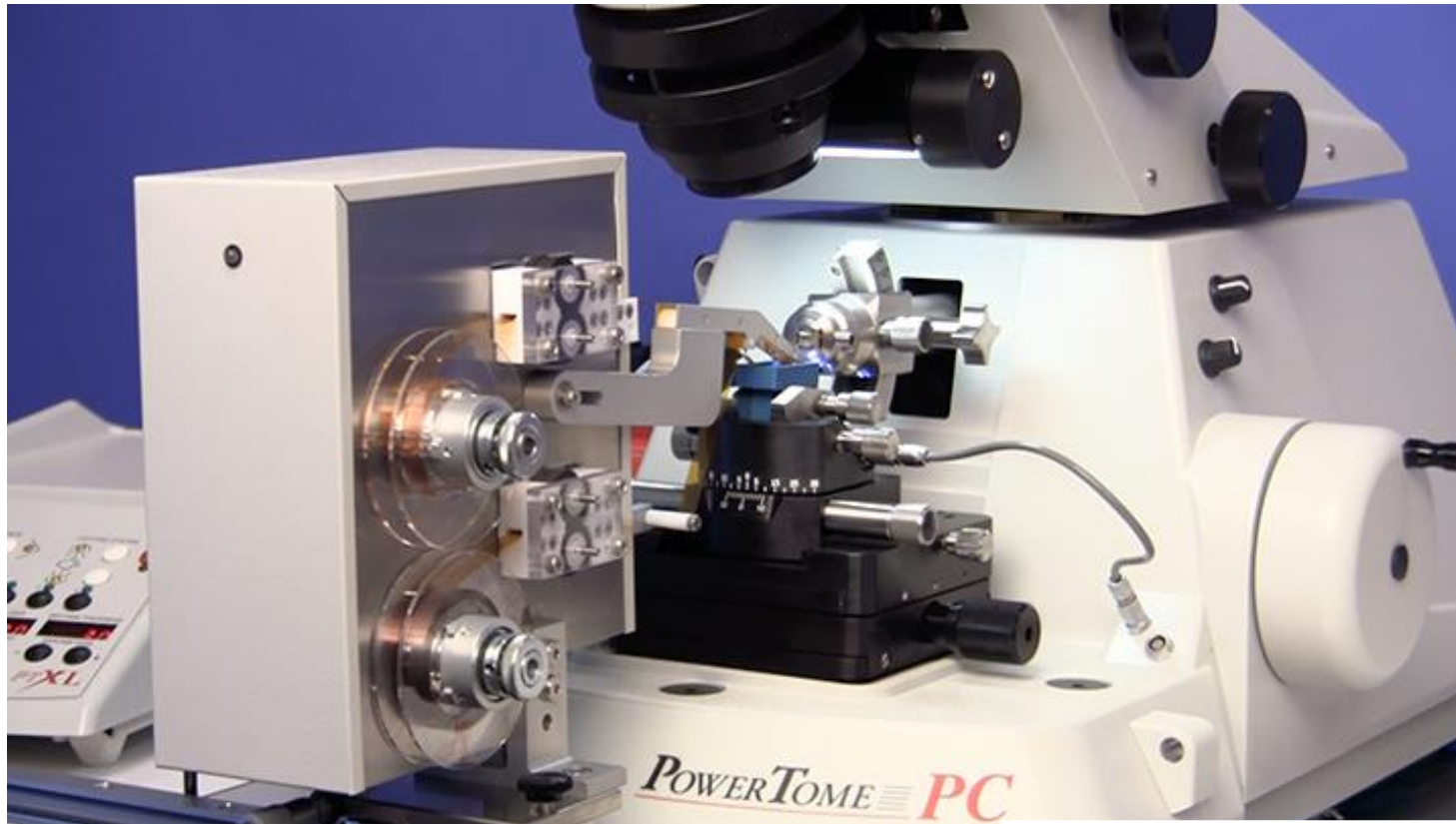


The TEM presents us with 2D images of 3D specimens, viewed in transmission. Our eyes and brain routinely understand reflected light images but are ill-equipped to interpret TEM images and so we must be cautious

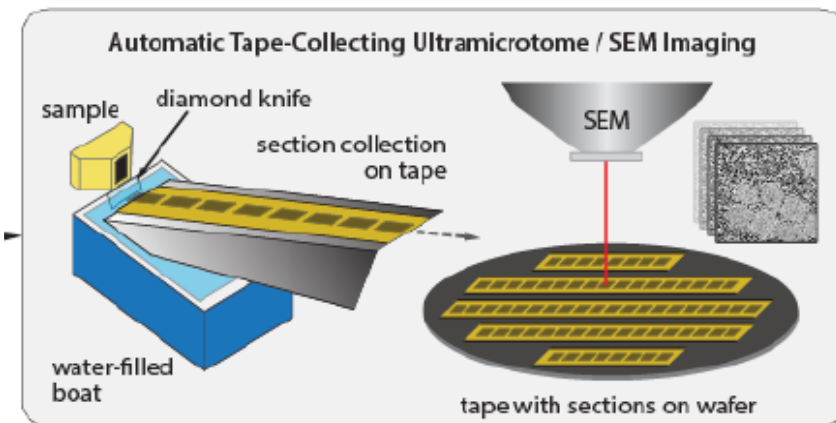


Photograph of two rhinos taken so that, in projection, they appear as one two-headed beast. Such projection artifacts in reflected-light images are easily discernible to the human eye but similar artifacts in TEM images are easily mistaken for 'real' features.

Array tomography



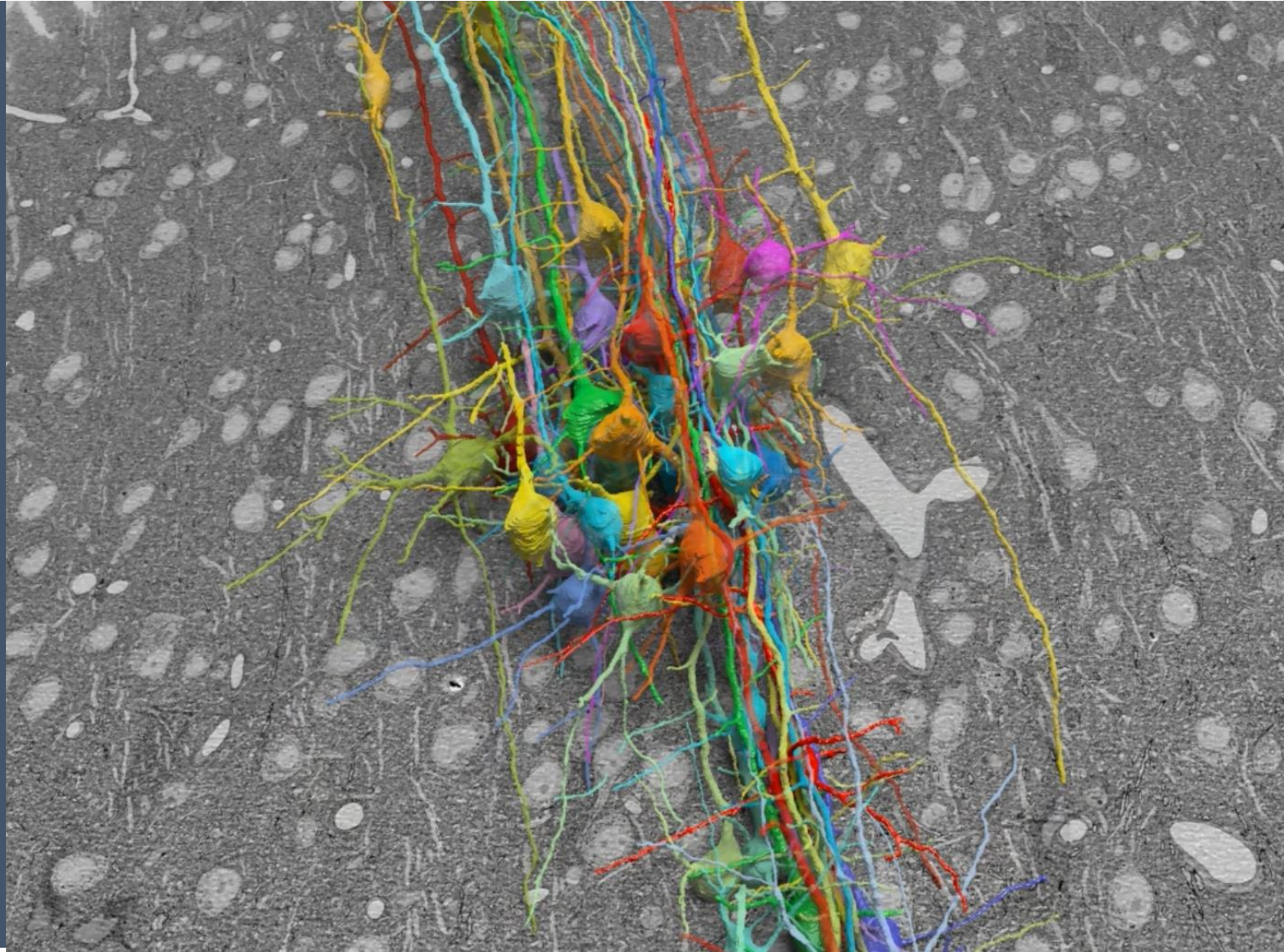
Thousands of ultra-thin sections with a thickness of 30nm can be automatically collected on 8mm wide Kapton tape for SEM imaging and subsequent 3-D reconstruction.

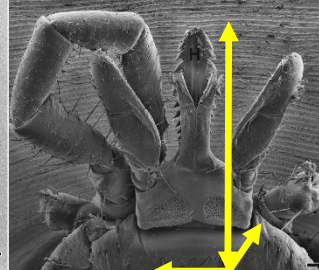
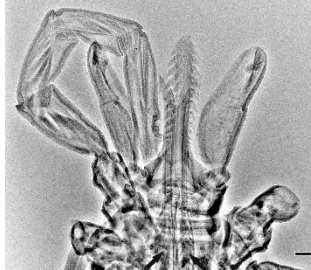
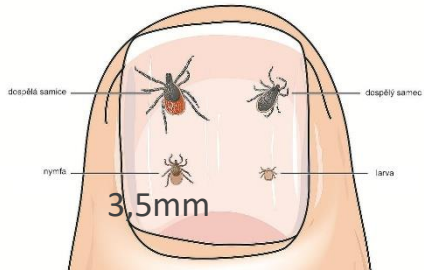


Large Volume High Resolution 3D SEM

ATUMtome und Atlas Array Tomography

the perfect team for large Volume SEM Imaging



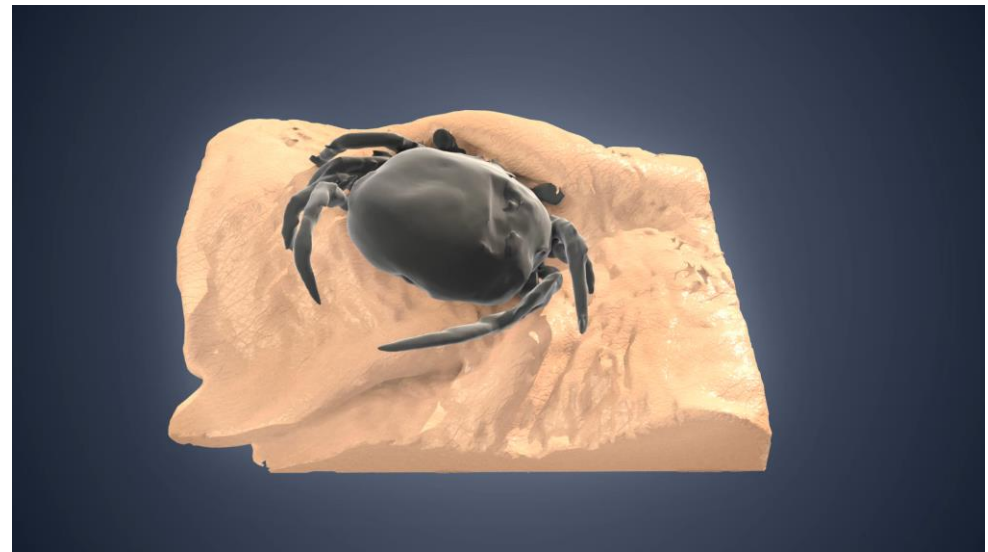
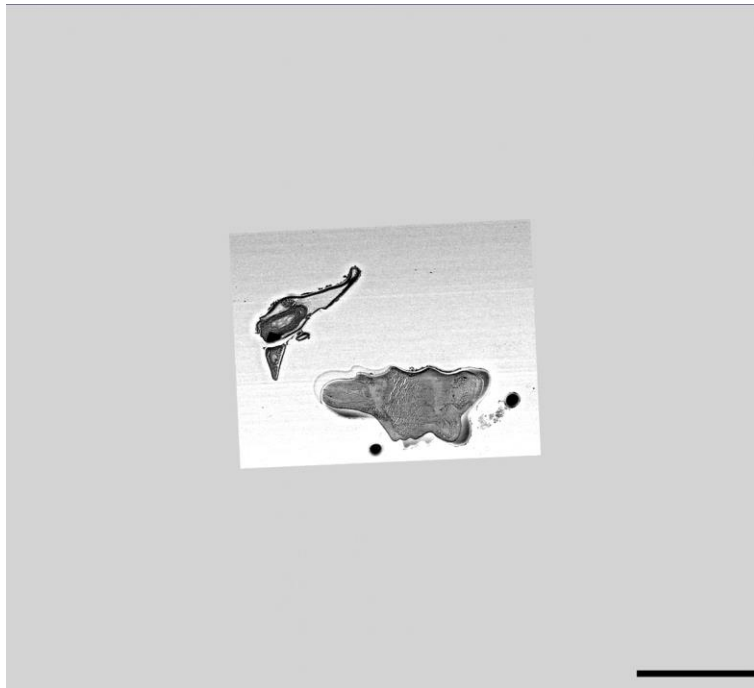


455 μm (5 μm / 500nm)

ARRAY TOMOGRAPHY

Reconstruction of the feeding apparatus, unfed nymph *Ixodes ricinus*

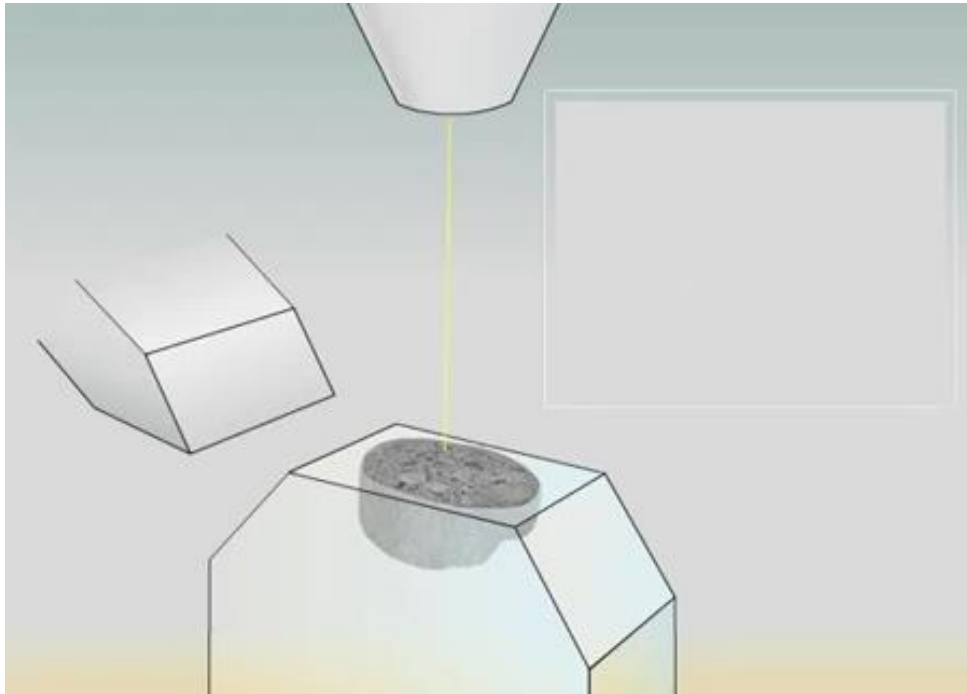
156x136 μm (39 nm)



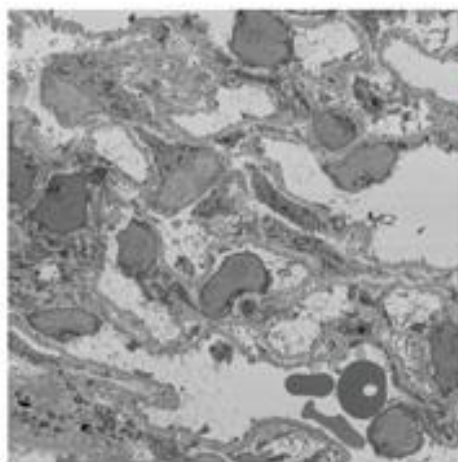
Vancova et al. 2020 SciRep

M. Vancova, LEM BC

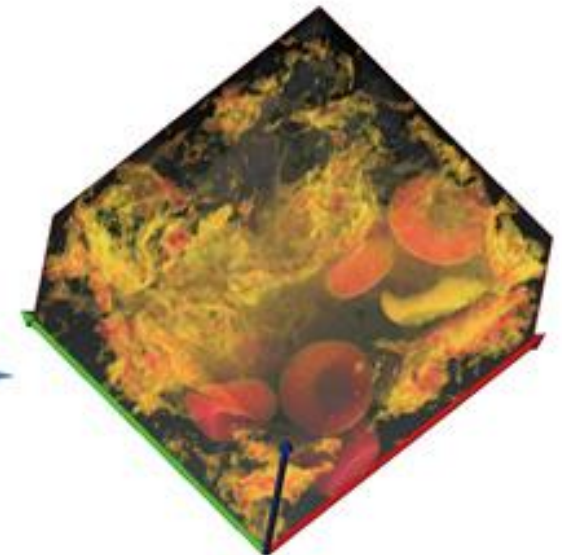
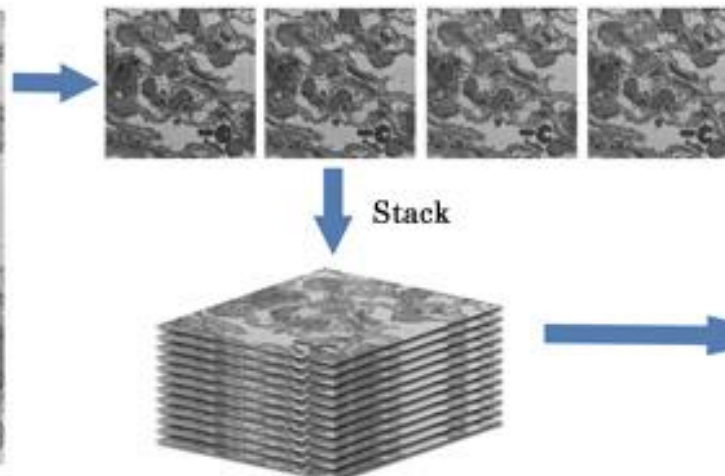
Serial block face SEM



GATAN website
3View System



Acquired image (backscattered electron image)
Specimen: Mouse kidney glomerular cells



Jeol company website; JSM-7200F/ 7800F

Large excretory organs of springtail – Minimal resin



SERIAL-BLOCK FACE SEM

SBEM strength:

- Larger ROI
- Tiles can be adjusted during the run
 - moved around
 - expanded
 - collapsed

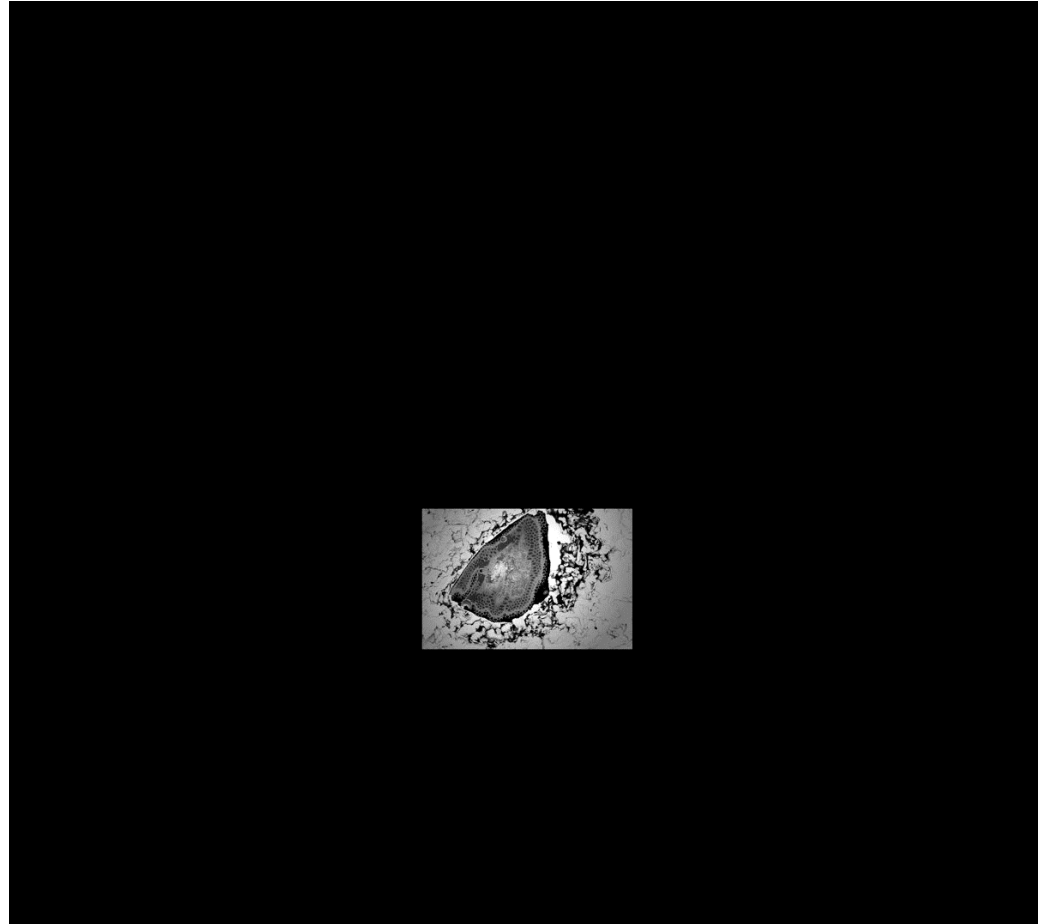
To note:

Acquisition time

Amount of data

Video shows approx every 5th slice (250 nm)

2 jumps to speed it up



Konopová, B., Týč, J. Minimal resin embedding of SBF-SEM samples reduces charging and facilitates finding a surface-linked region of interest. *Front Zool* **20**, 29 (2023).

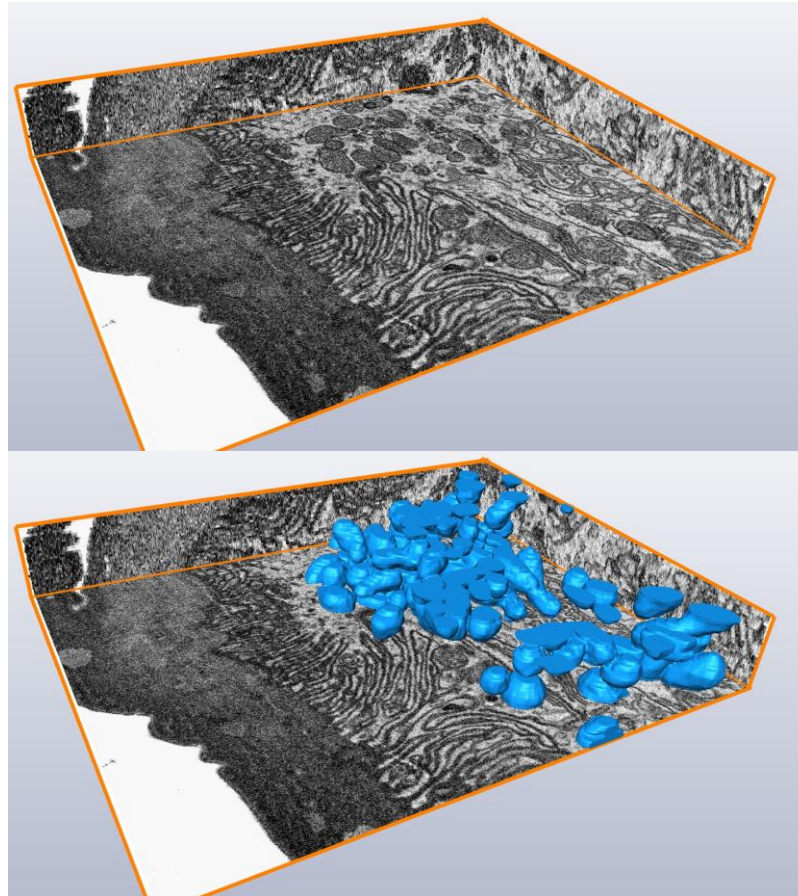
Large excretory organs of springtail

Run parametres	value
Pixel size	8 nm
Slice thickness	50 nm (part 25 nm)
Imaged area	170 x 170 μm
Total slices	2500
Total volume	0,002 mm ³
Total acquisition time	Approx. 1 month
Amount of Data acquired	8,2 Tb

Take home messages:

Large volumes are time and data storage demanding

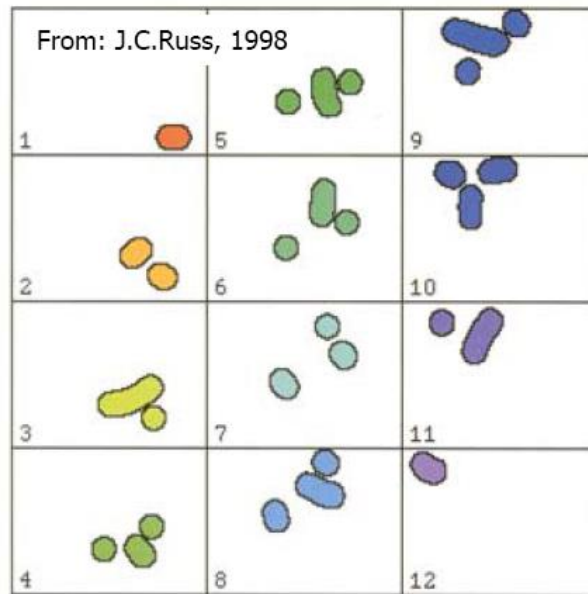
Resolution is enough to distinguish mitochondrial cristae and other fine membranous details



SERIAL-BLOCK FACE SEM

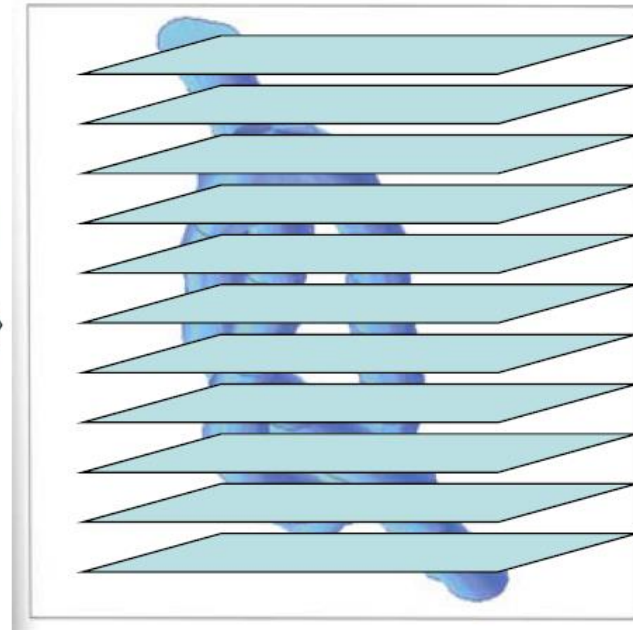


Problem of 3D serial sectioning: reconstruction of disordered microstructures



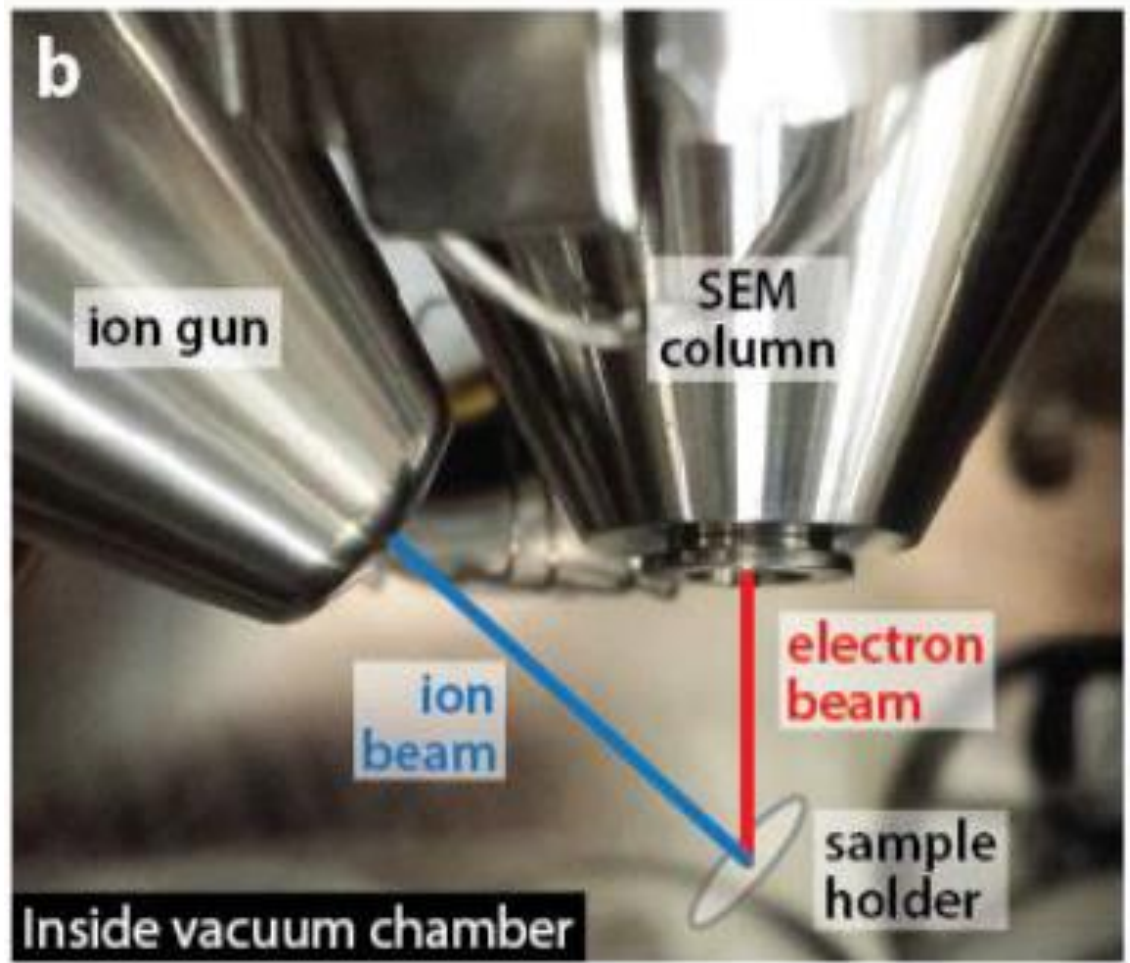
2D volume fraction

3D
→



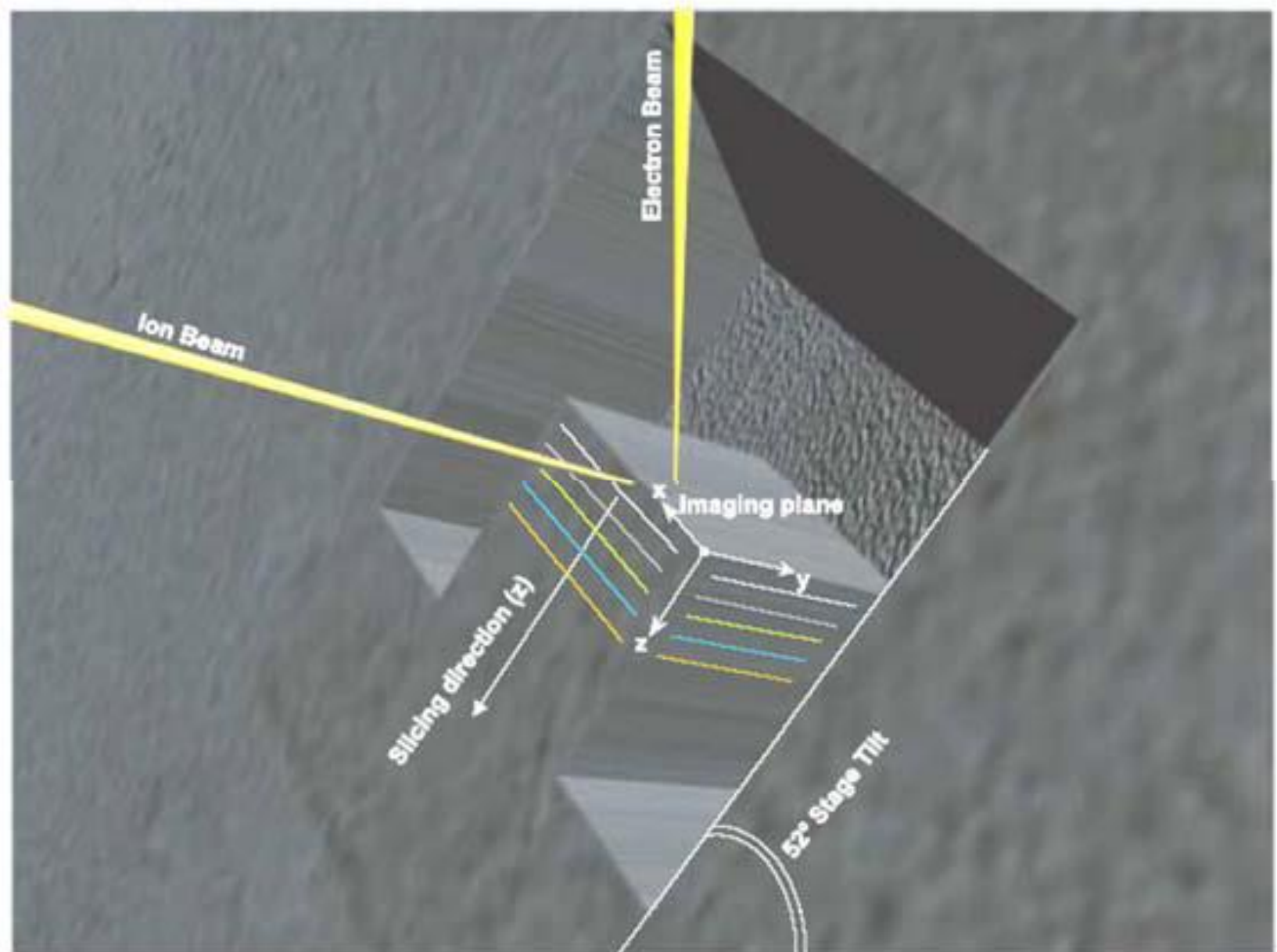
How many particles?
Shape?

FIB SEM = dual beam

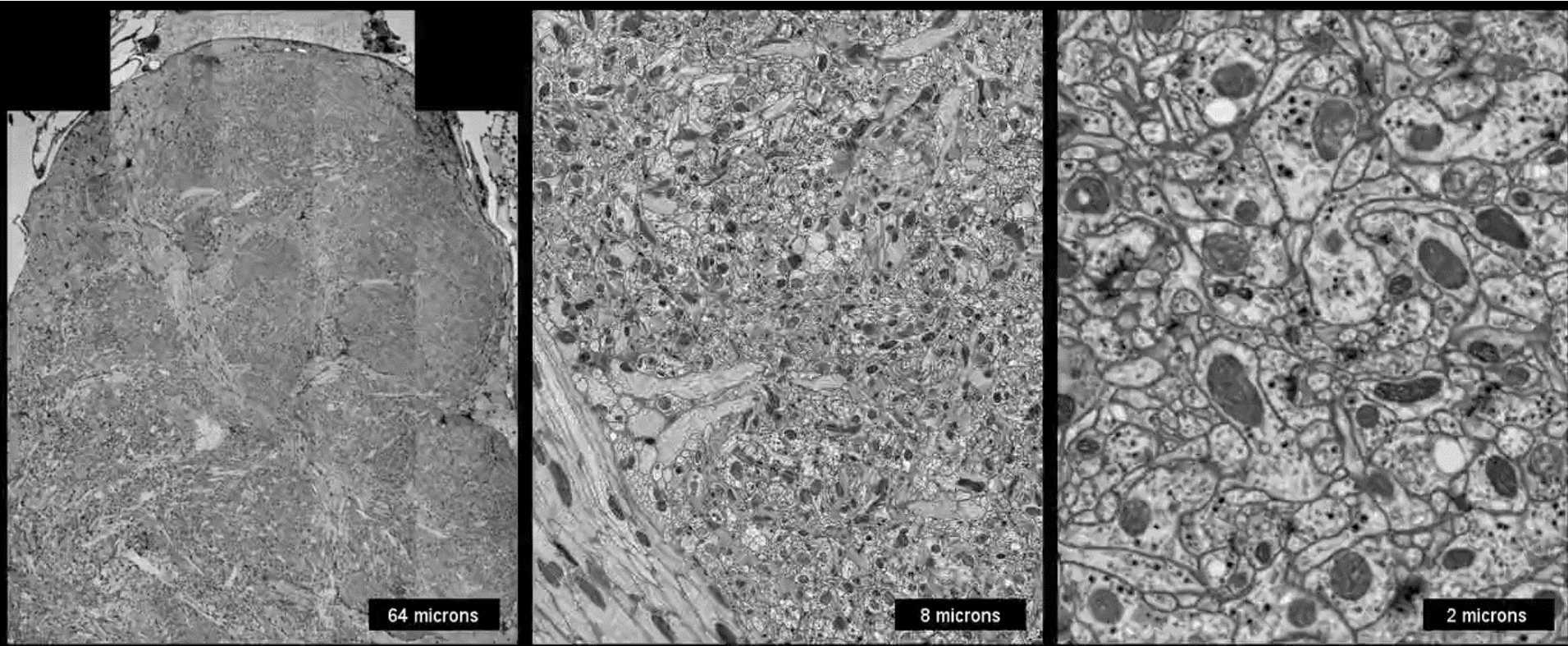


The sectioning is done by means of the focused ion beam working in the similar way as the scanning electron microscope.

FIB SEM = dual beam

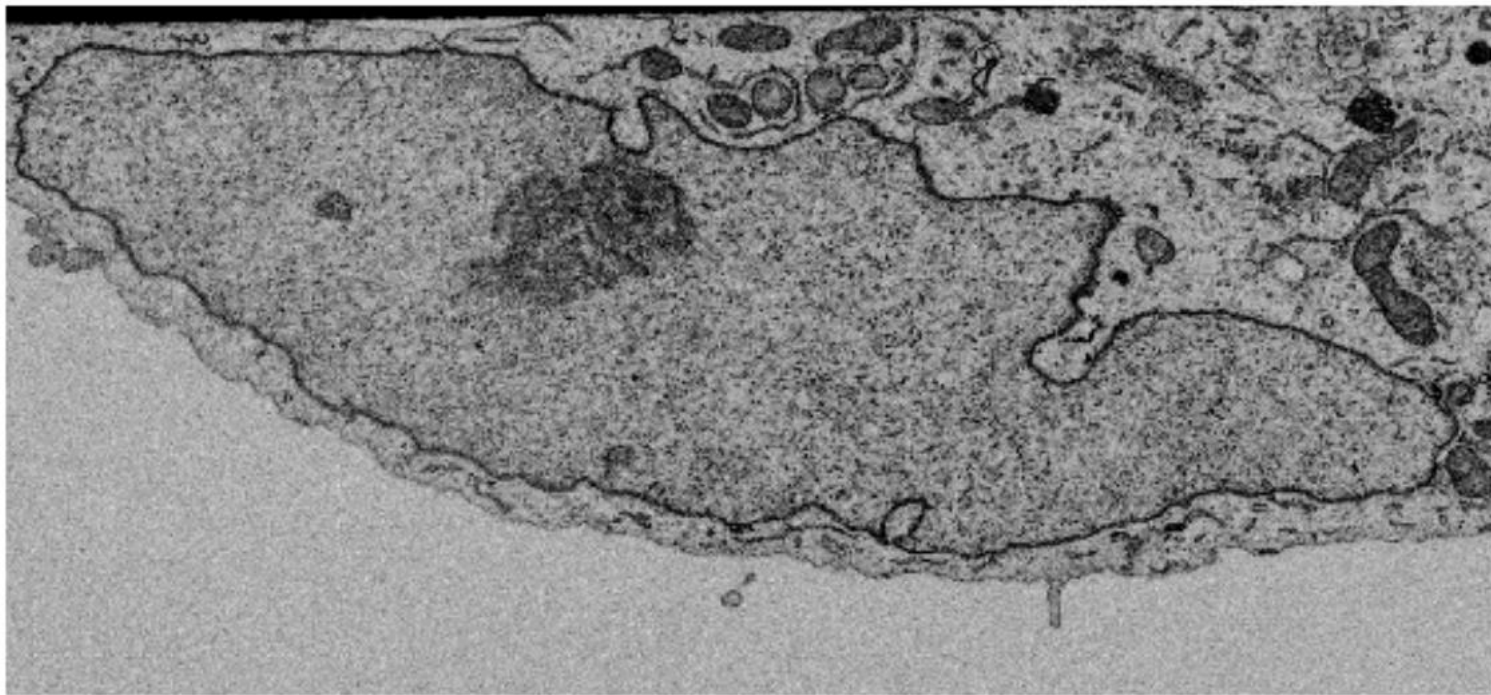


Drosophila brain imaged by FIB-SEM

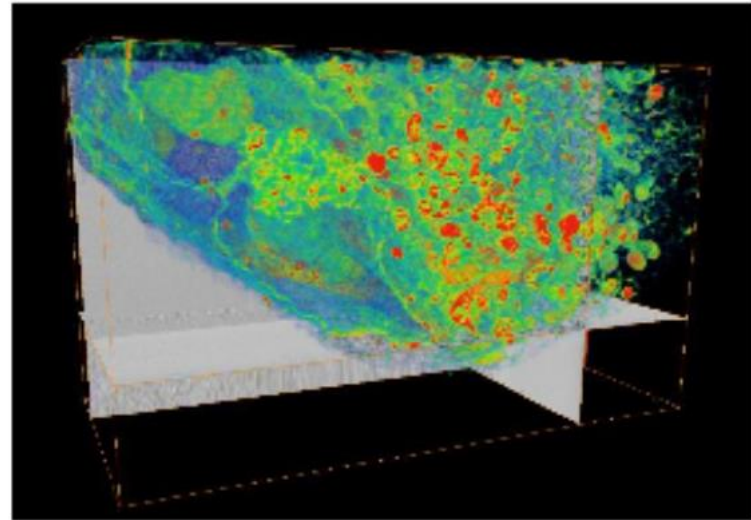


pixel sizes: 512nm, 64nm, and 16nm

[Hess Lab, Janelia Campus, HHMI](#)



A single slice image and a 3D visualization of the HT1080 cell. The data were acquired by **Dual-beam FEI Helios Nanolab 660 G3 UC** with in-lens backscatter electron detector (ICD) at 5 nm slice thickness and 3 nm pixel size. The dataset was post-processed with **Amira Software 6.2**. The sample was provided by **Dušan Cmarko (FFM CU)**.

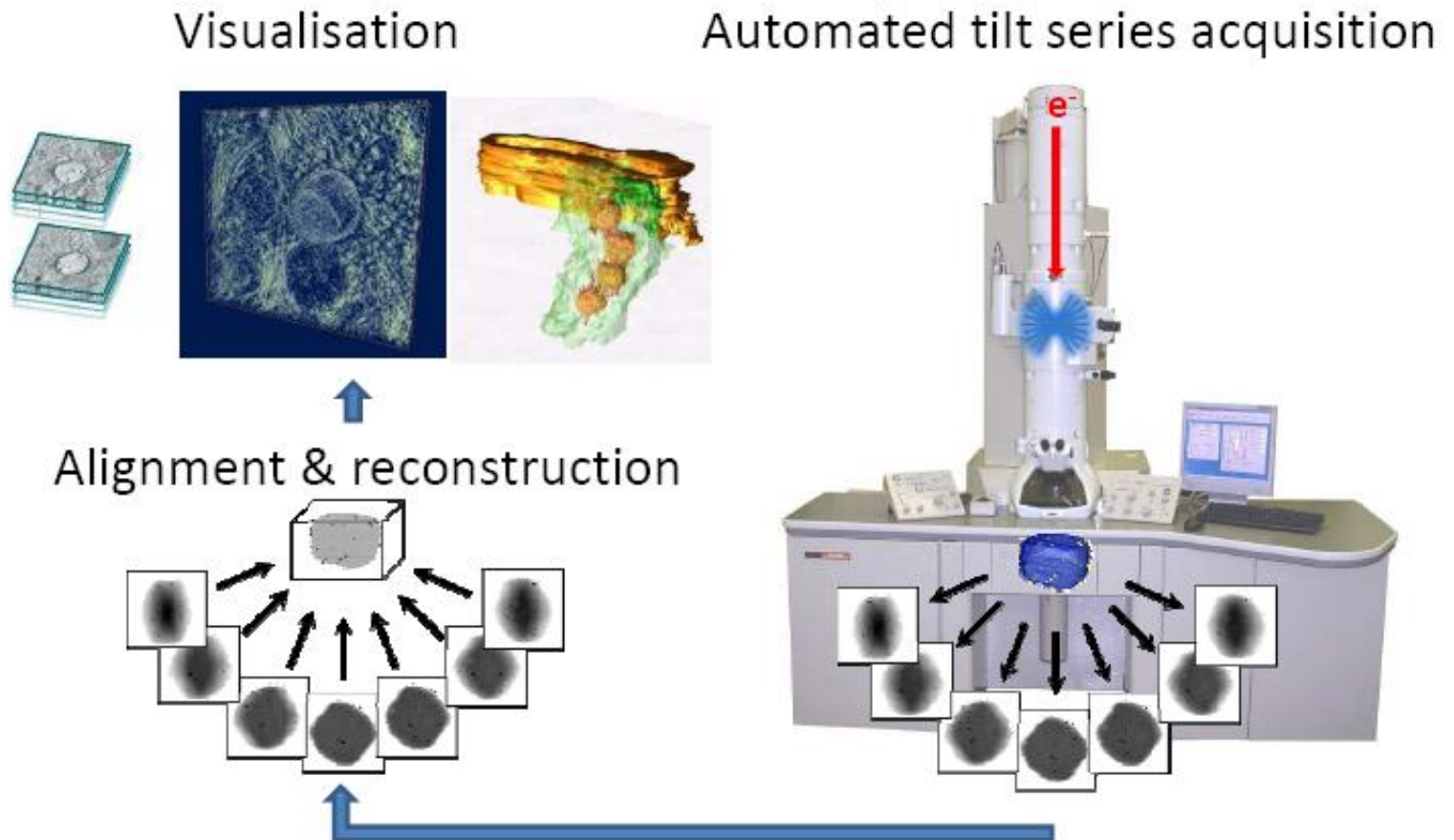


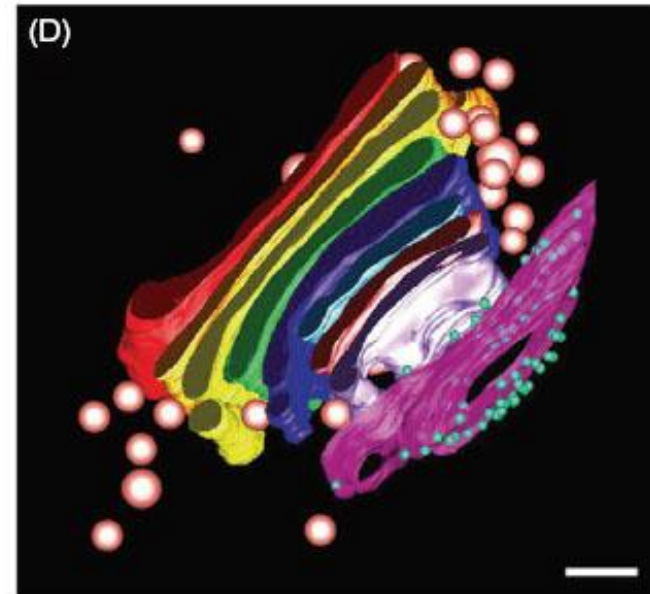
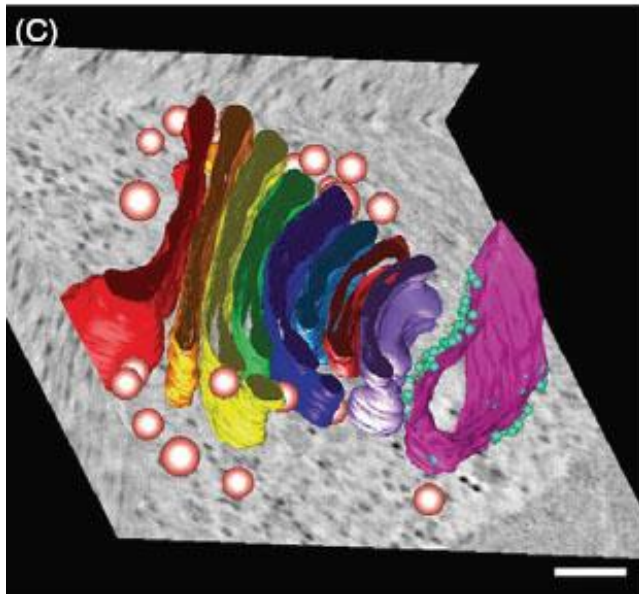
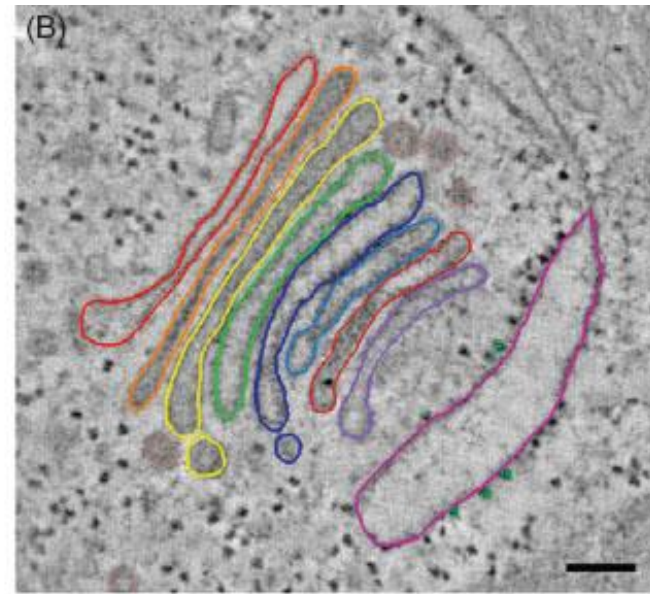
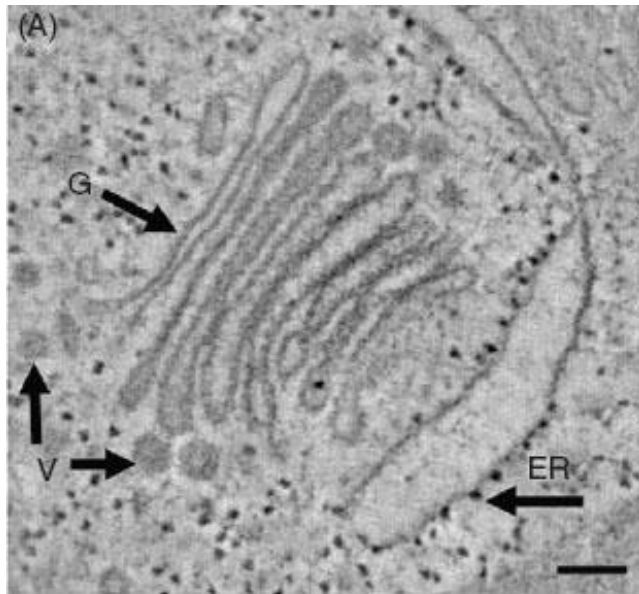
BIOCEV

M. Dalecká, currently IMG EM

Electron tomography

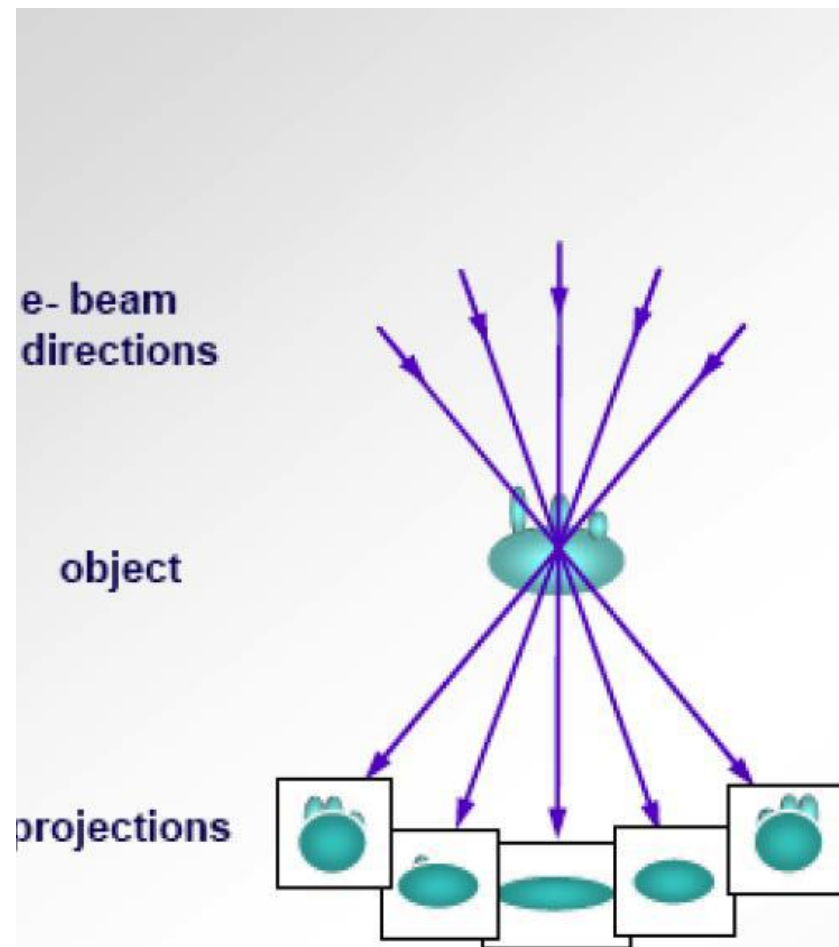
Is a method for volume reconstruction of an object from its projections





Cellular membranes in 3D
O'Toole 2010, Met. Cell Biol.

Electron tomography



IDEAL SITUATION:

As many images as possible from
ALL orientations (-90° to 90°)

RESOLUTION, d

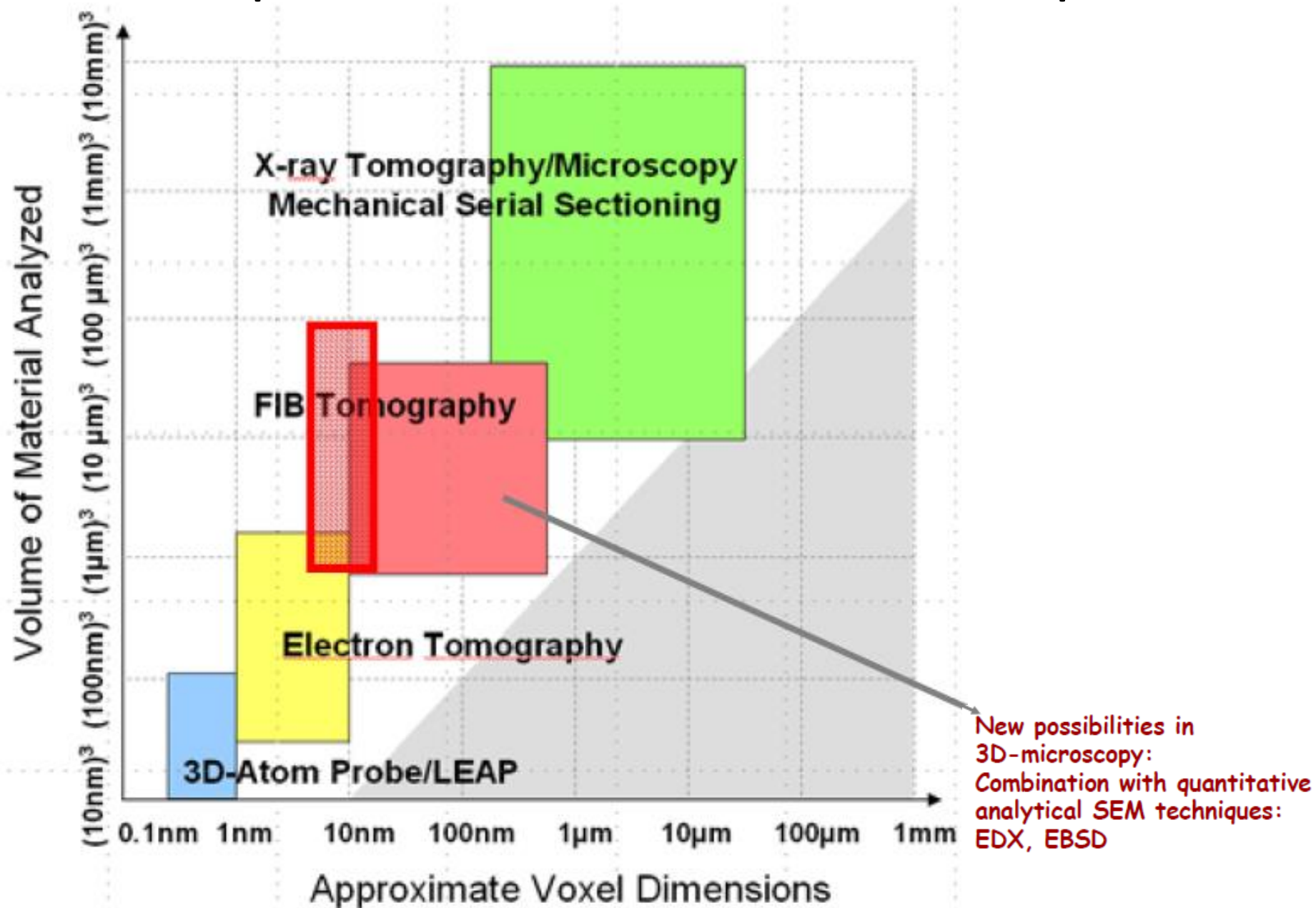
Crowther, DeRosier, and Klug, 1970

$$d = \pi \frac{D}{N}$$

D , diameter of the object
 N , number of projections

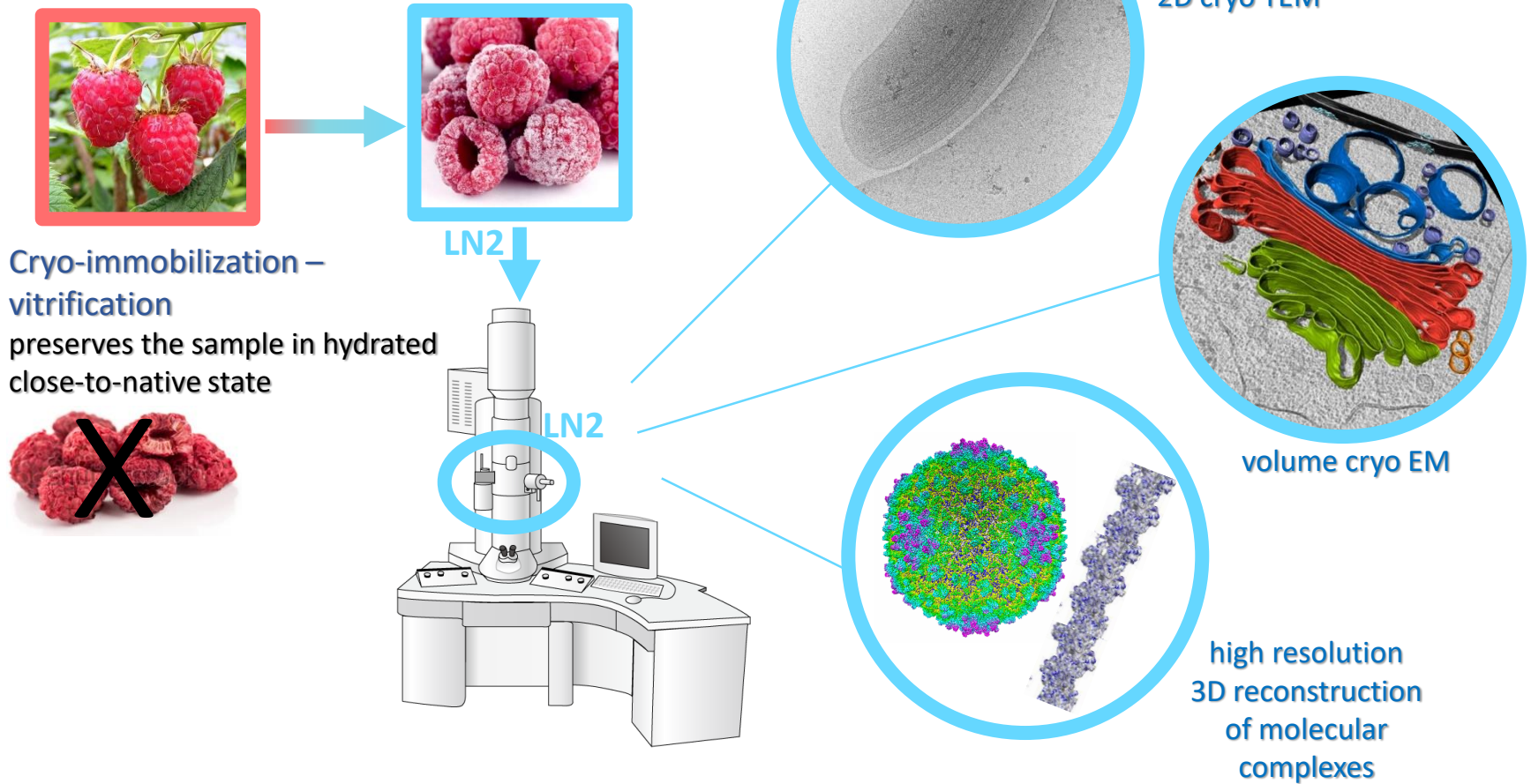
2 nm with 141 images
of a 100 nm object

Comparison of various 3D-techniques

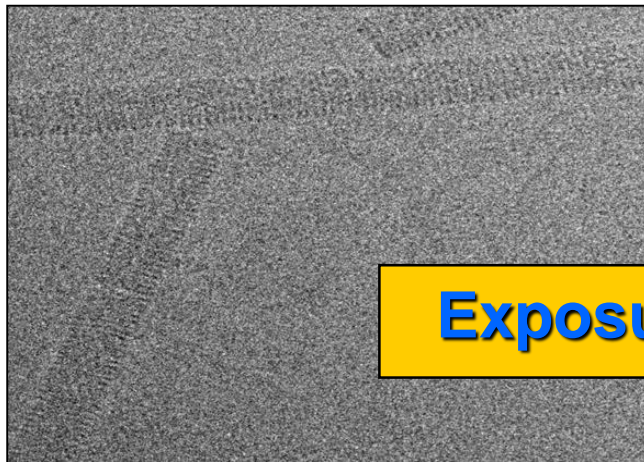
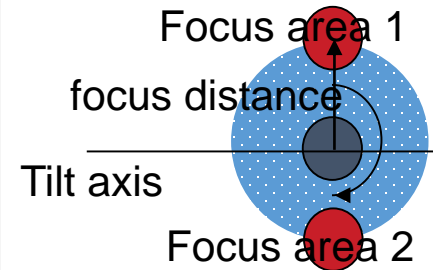
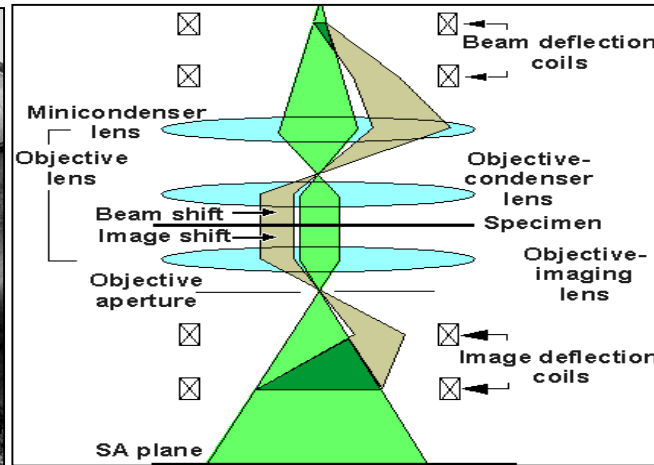
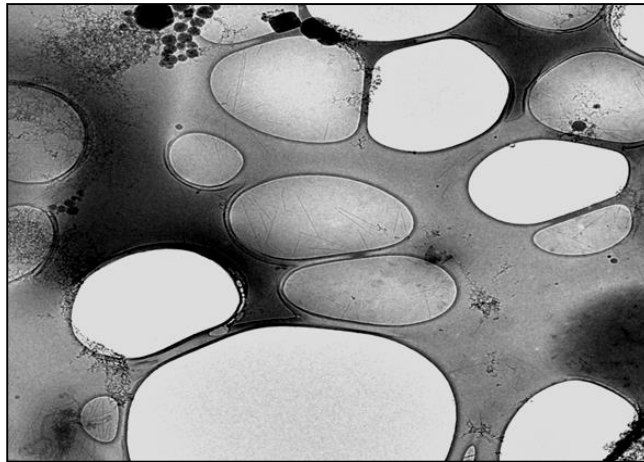


Cryo-TEM – low dose

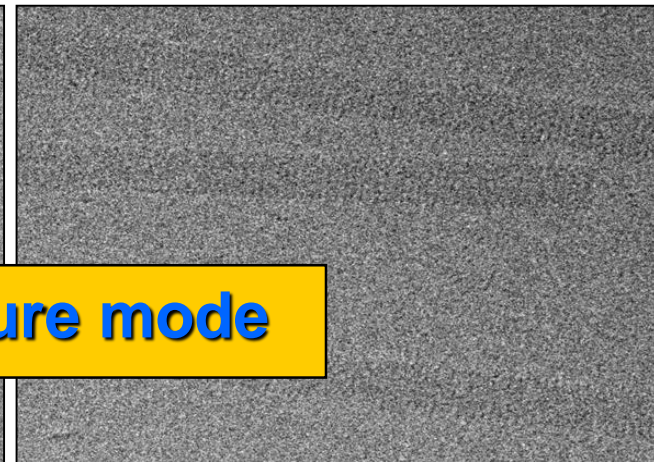
Cryo electron microscopy



Low-Dose / Cryo-TEM Principle



Exposure mode



Low dose

Low Dose Blank Peek

Status : LD on, Focus state 2

Search Focus Exposure

TEM Mi	1	2	TEM SA
3960x			43200x
Spot 5	Spot 5	Spot 2	
Int 62.26	Int 66.46	Int 66.46	
x 0.000 um	4.31 um		
y 0.000 um	156.8°		
		1.0 s	

Start Start

Expose Focus Series

Expose ☐ Use spotscan

☐ Dim Screen ☐ Series

Exposure time (sec) 1.0

Wait (sec) after plate in 0

☐ Pre-expose (sec) 0.0

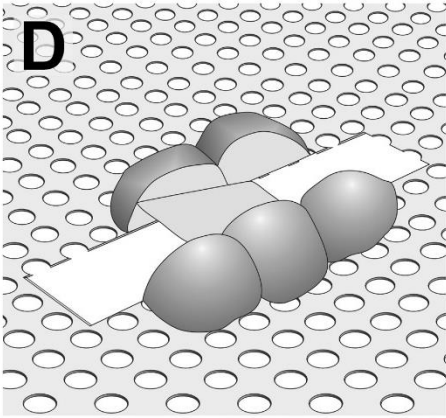
☐ Wait after pre-exposure 0.0

Tobacco Mosaic Virus (TMV)
Recorded on Slow-scan CCD using Low-Dose mode

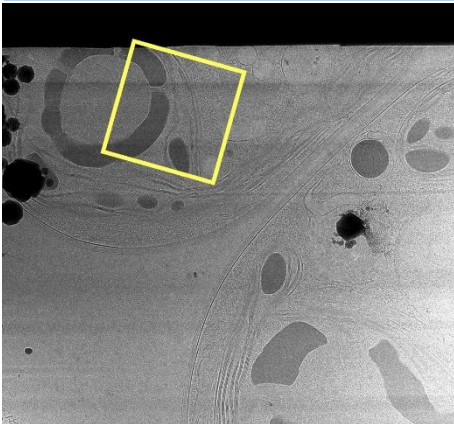
Cryo-ET of FIB-milled Lamella

lamella preparation
by FIB

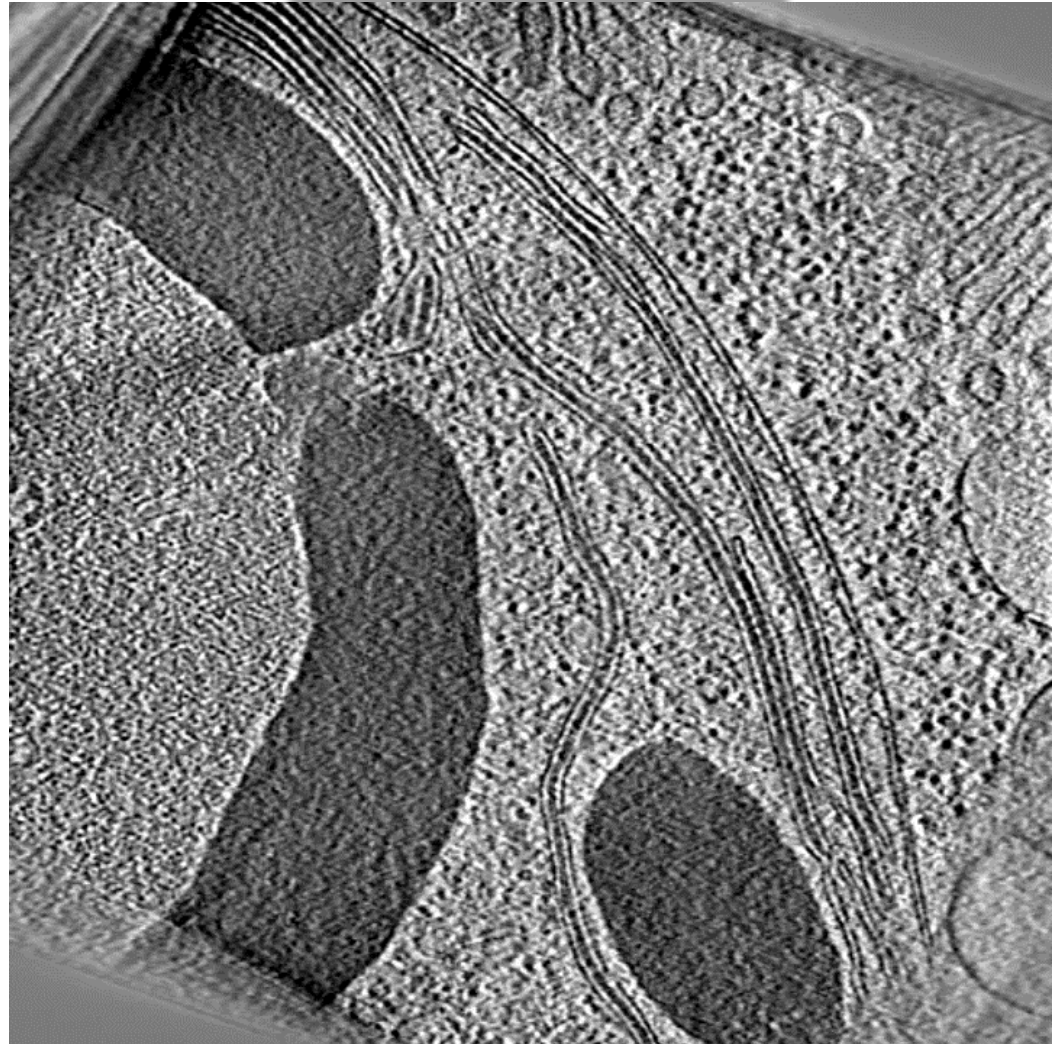
D



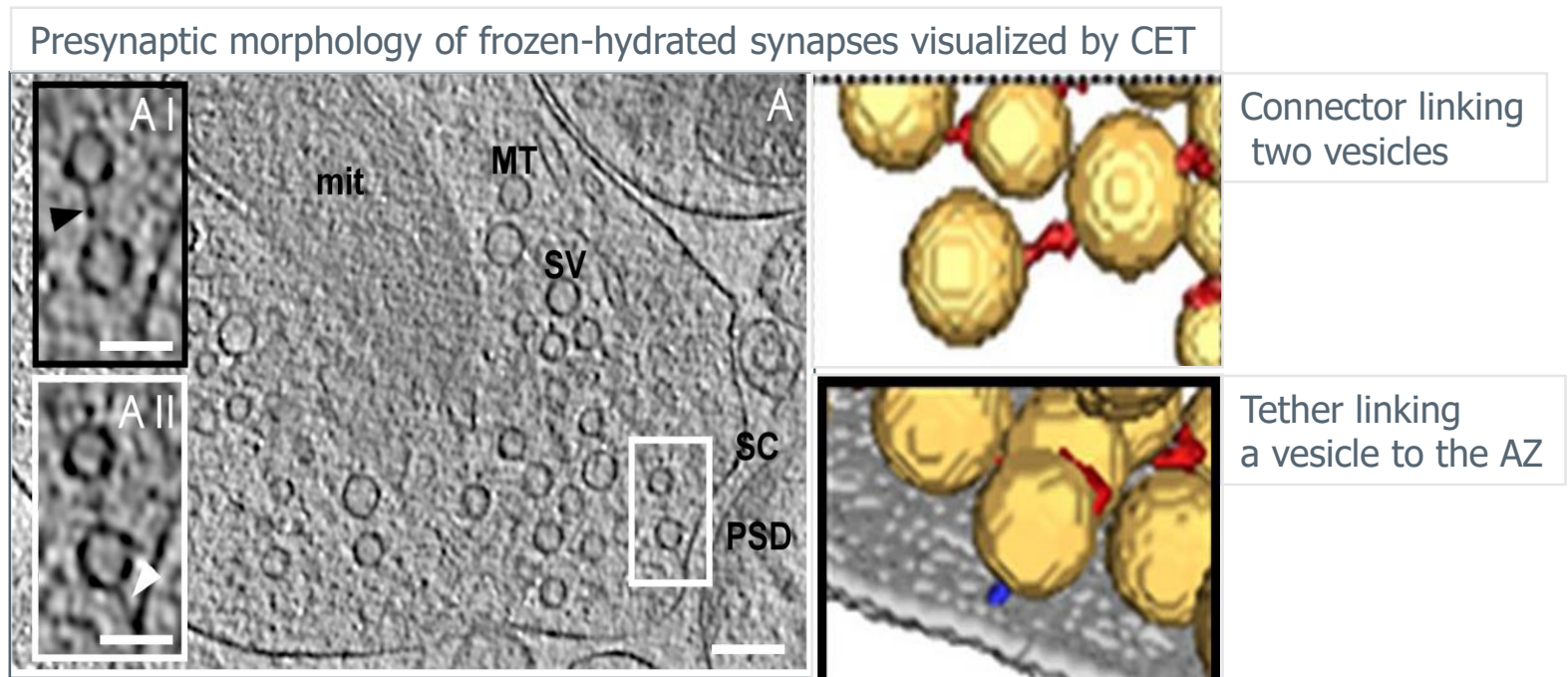
ROI selection in Cryo
TEM - Tomography



3D Reconstruction



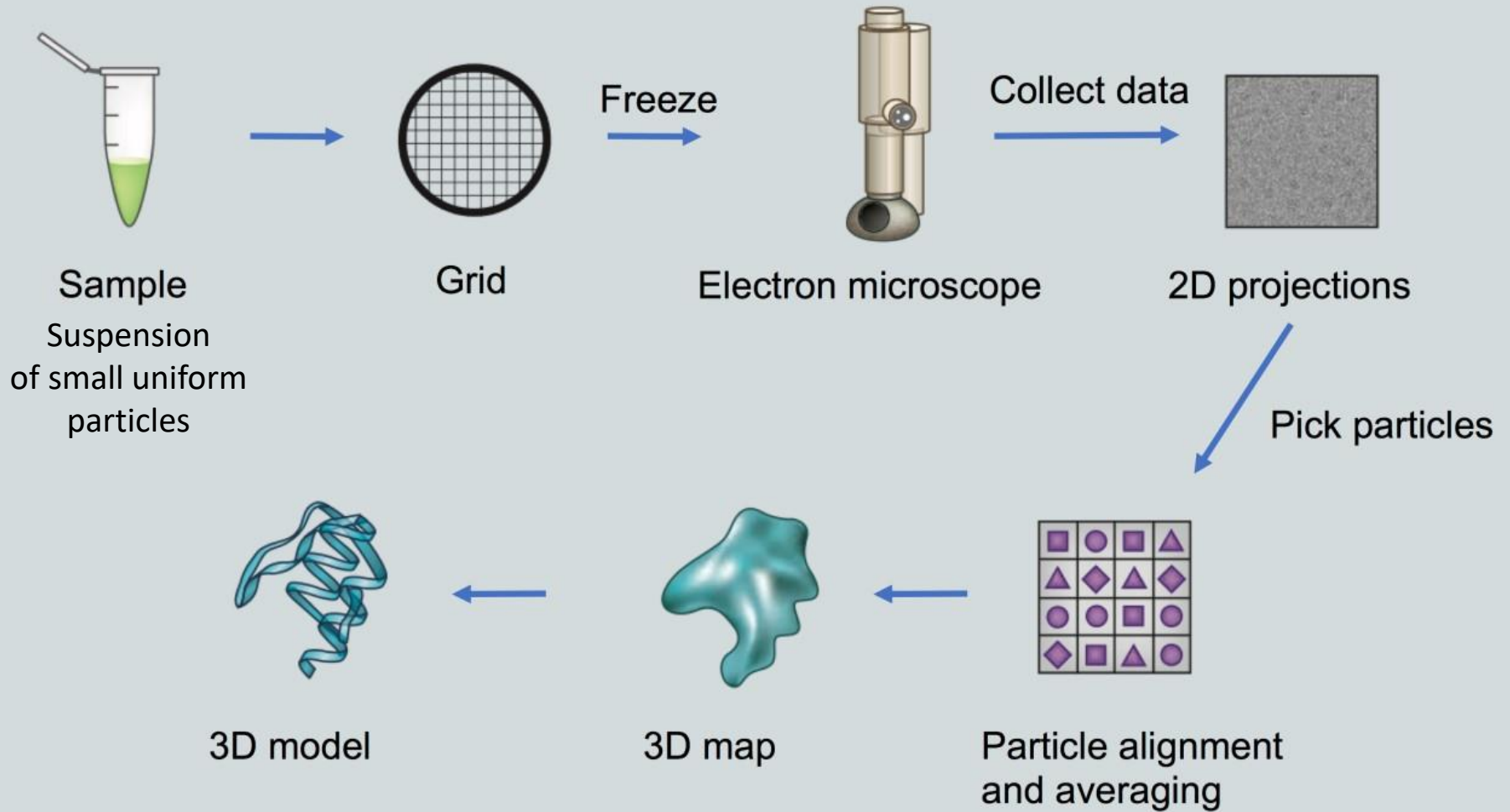
Structures visualized by Cryo-ET



Fernández-Busnadiego, R., Zuber B., Maurer, U.E., Cyrklaff, M., Baumeister, W., Lučić, V., 2010 *J. Cell Biol.* Vol. 188

Single particle analysis

SPA workflow



SPA workflow



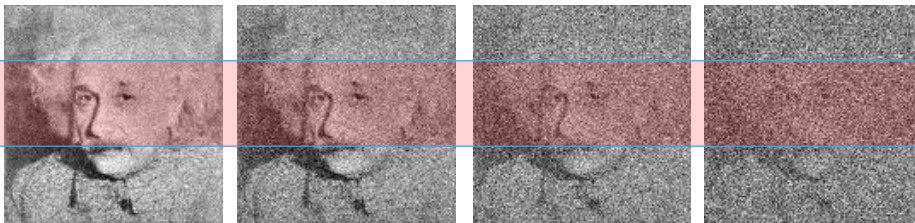
Protein purification



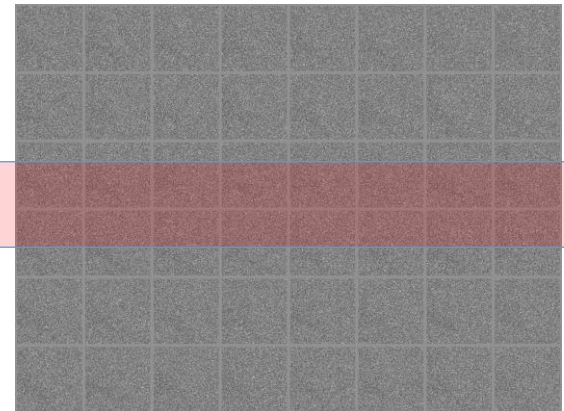
Protein vitrification



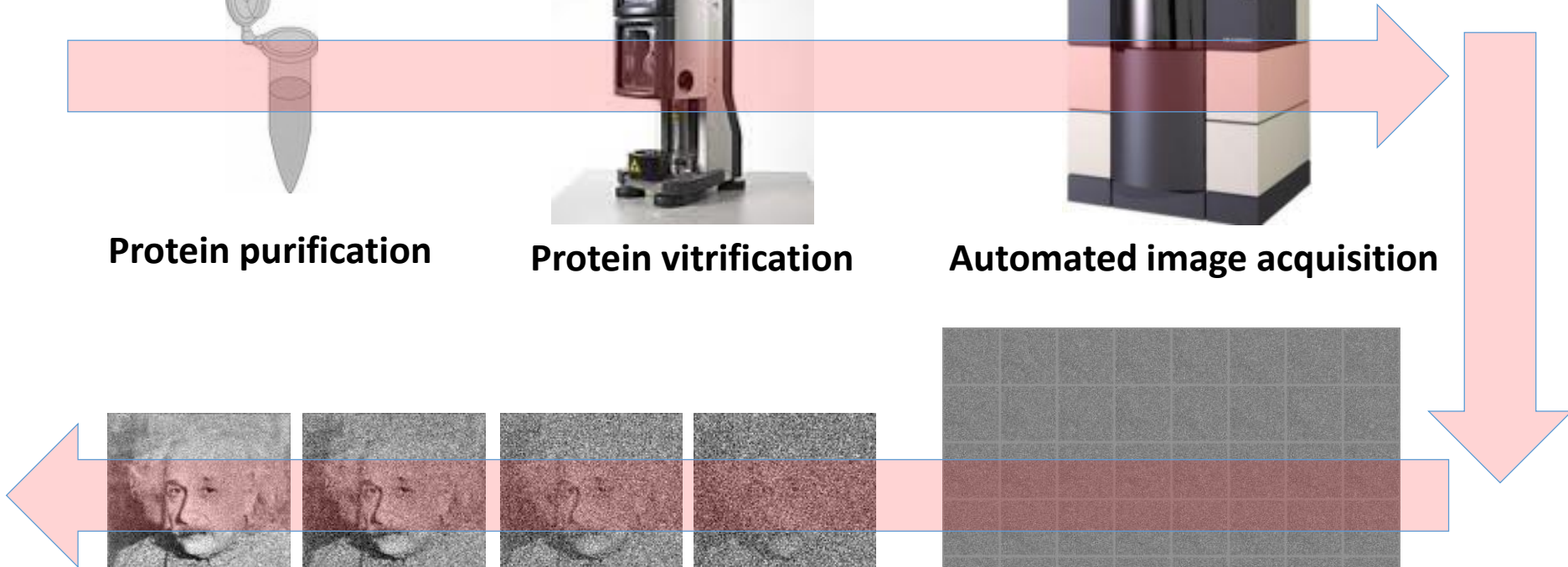
Automated image acquisition



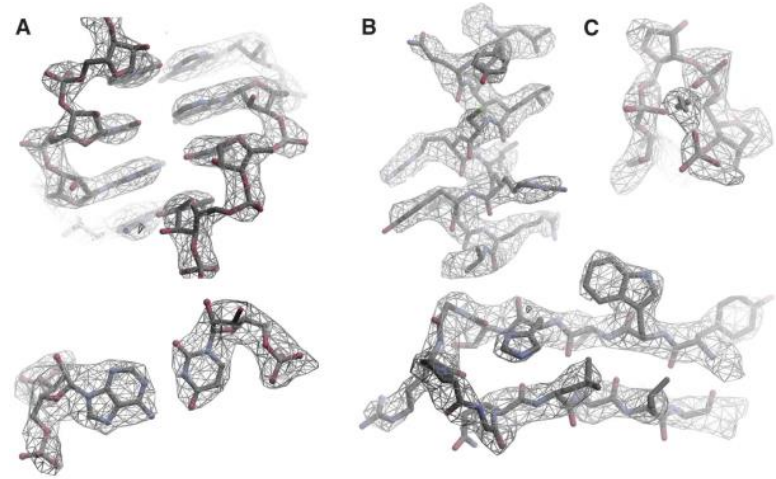
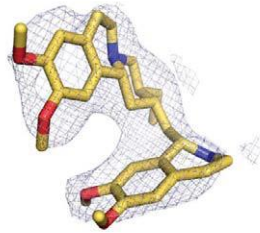
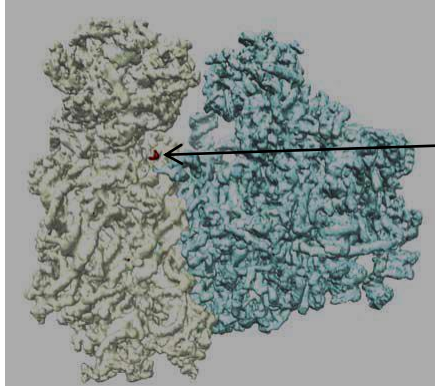
Iterative image processing



Particle classification

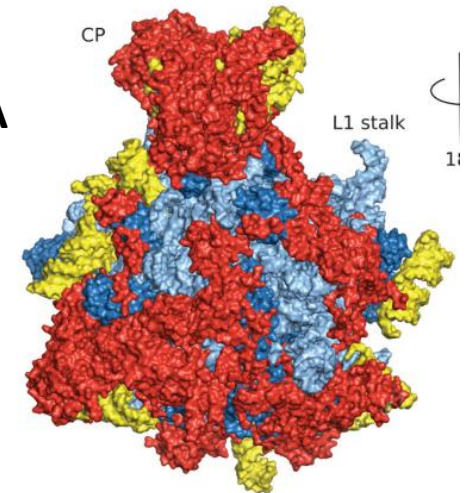


Ribosome (2014):

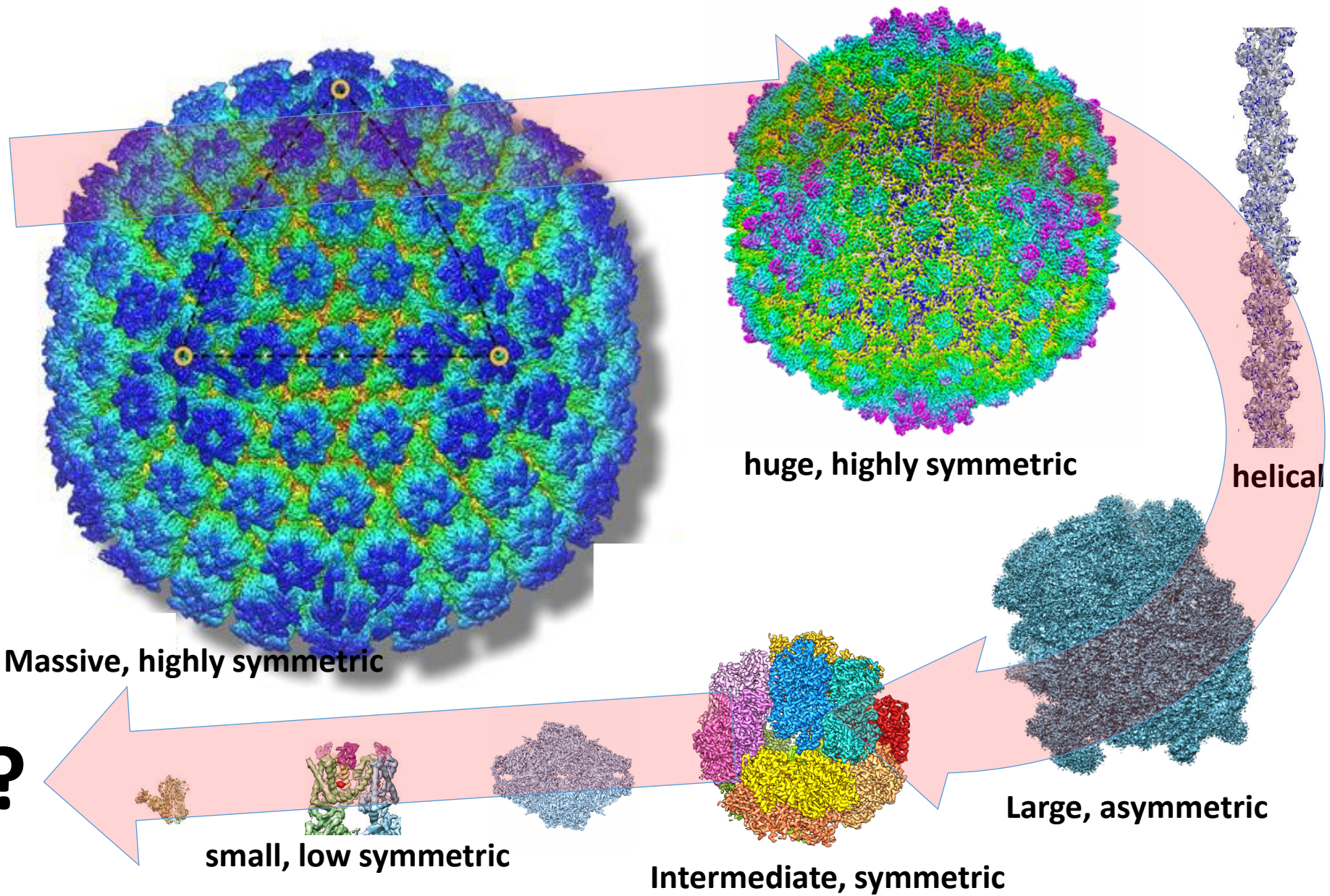


Full *de novo* model build

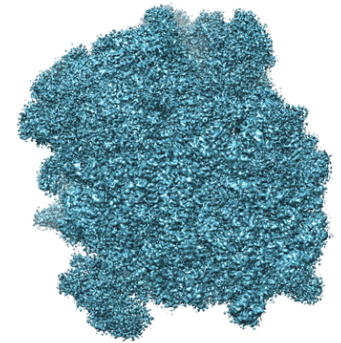
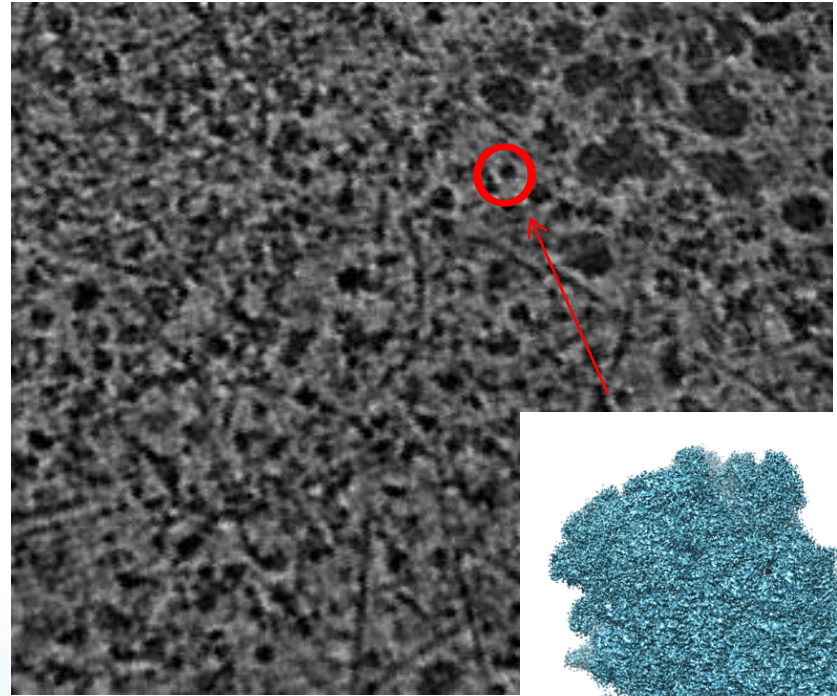
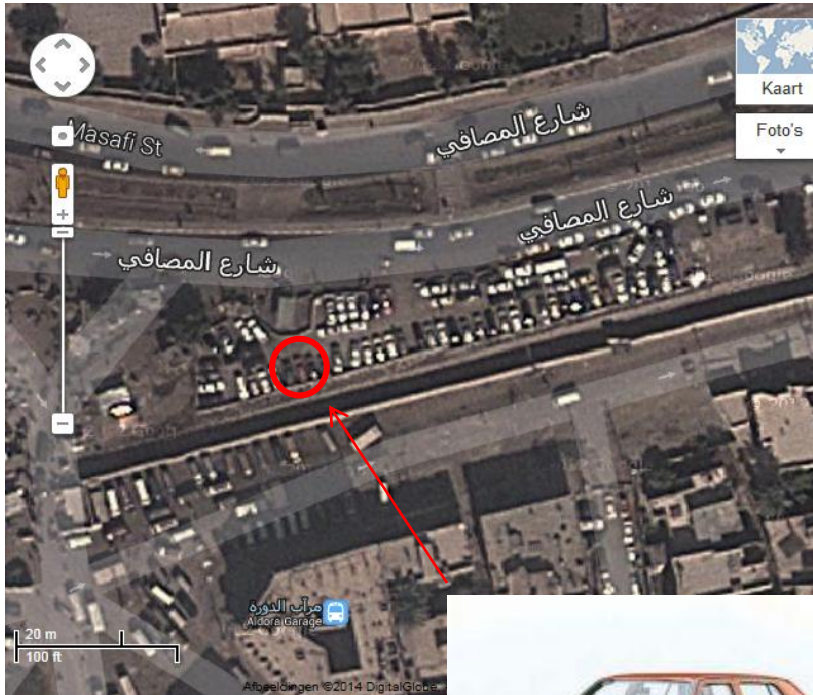
Routinely $\leq 4\text{\AA}$



- Hussain T, et al. **Cell** (2014) 159 pp. 597-607 (falcon)
Bischoff L, et al. **Cell Rep.** (2014) 9 pp. 469-475 (Falcon)
Arenz S, et al. **Molecular Cell** (2014) (falcon)
Brown A, et al. **Science** (2014) 346 pp. 718-722 (Falcon)
Greber BJ, et. Al. **Nature** (2014) (Falcon)
Shao S, et al. **Molecular Cell** (2014) 55 pp. 880-890 (Falcon)
Voorhees RM, et al. **Cell** (2014) 157 pp. 1632-1643 (Falcon)
Wong W, et al. **eLife** (2014) 3 (Falcon)
Fernandez IS, **Cell** (2014) 157 pp. 823-831 (Falcon)
Amunts A, **Science** (2014) 343 pp. 1485-1489 (Falcon)
Greber BJ, et al. **Nature** (2014) 505 pp. 515-519 (cover) (Falcon)

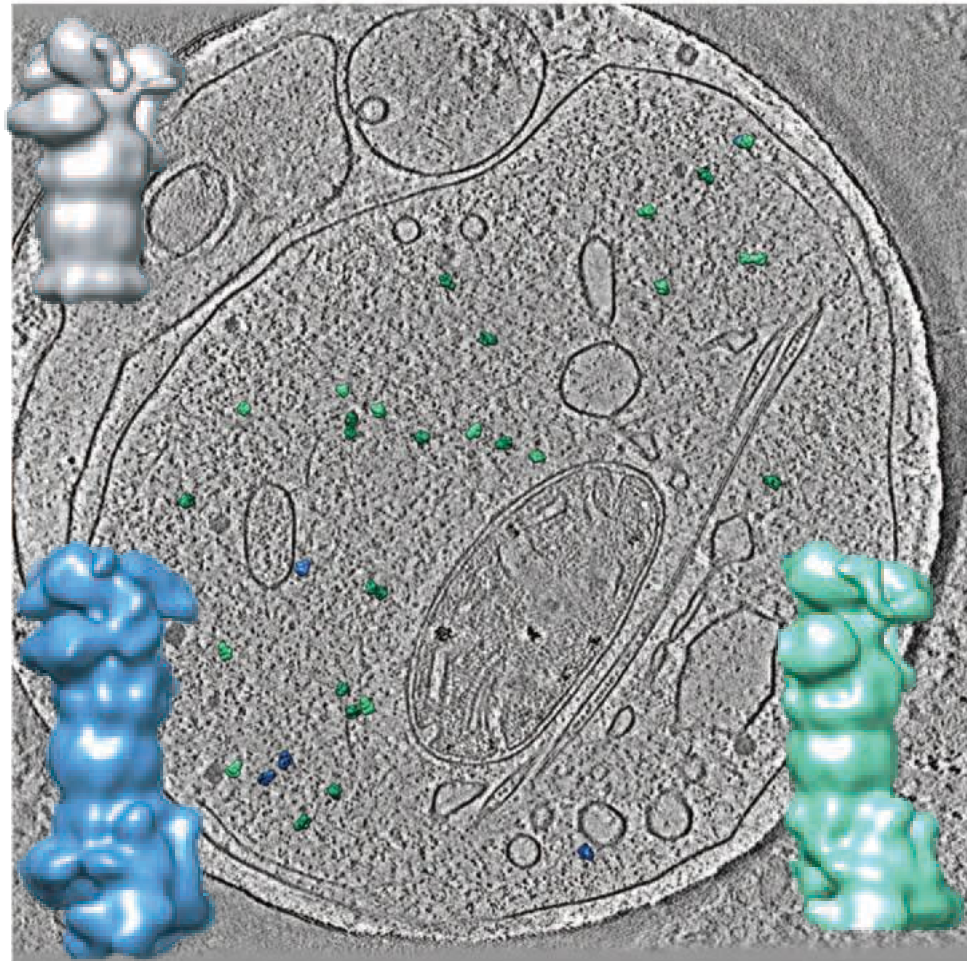


Protein in the context of the cell at intermediate resolution



Docking SPA protein into a tomogram

Cryo-ET of a neuron

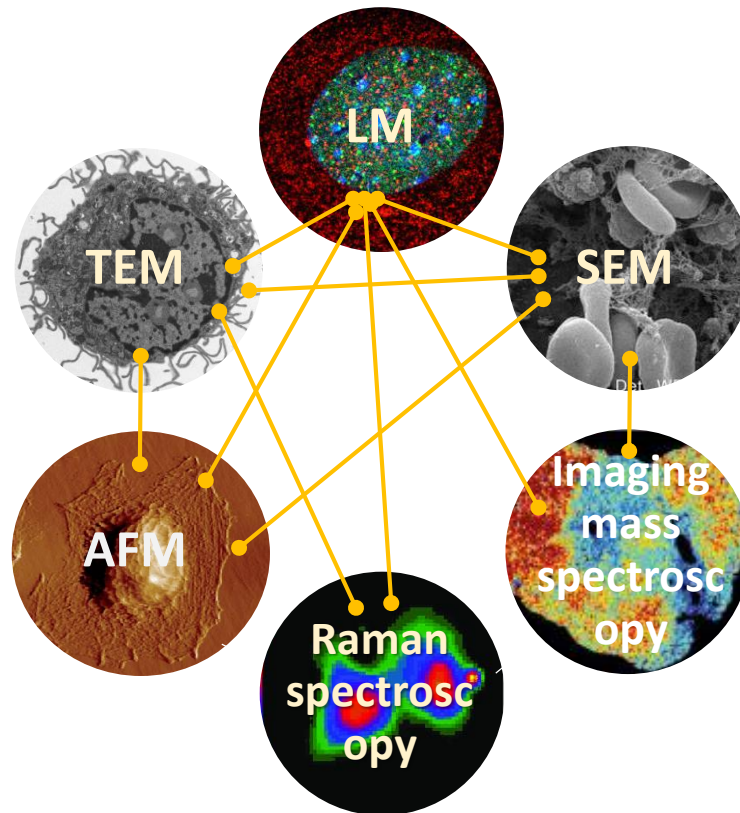


**Phase plate assisted sub-tomo averaging,
conformation classification and quantification
INSIDE the cell!**

Correlative microscopy

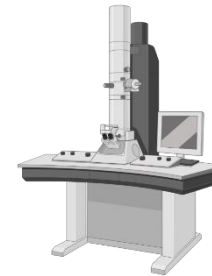
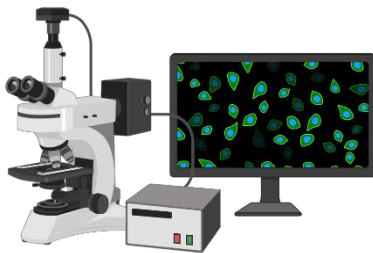
Correlative microscopy

combination of two or more **microscopic techniques** applied to **the same region** in a specimen



Examples:
LM + TEM
LM + SEM
AFM + TEM
LM + AFM
...

Correlative light and electron microscopy (CLEM)



Light Microscopy (widely used in biology)

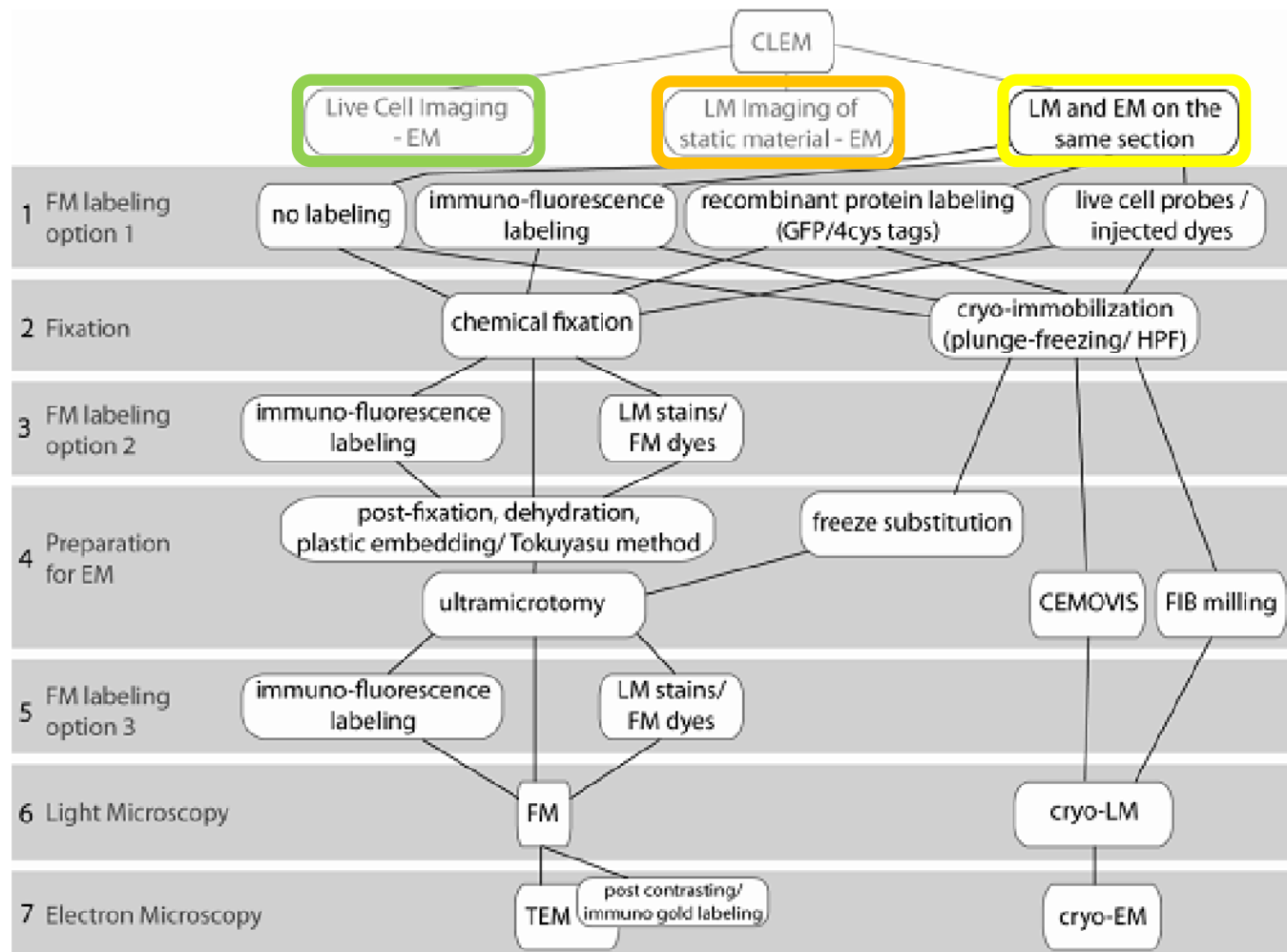
- Large field of view/overview
- Live cell imaging
- Short or no sample preparation
- Combination with fluorescent labels
- Resolution 200 – 20 nm
- Little cellular context

Electron Microscopy

- Excellent spatial resolution
- Provides the cellular context
- Extensive sample preparation

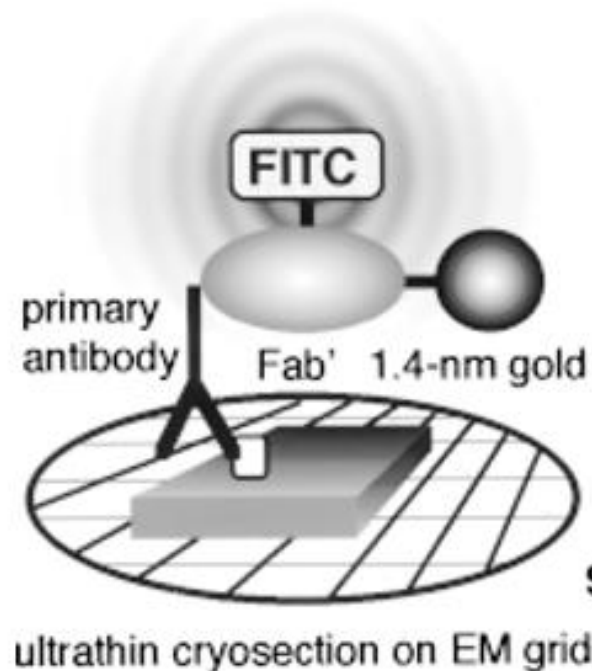
Challenges:

- Defining the precise position on the sample both in LM and EM
- Tailored sample preparation for the EM (preserve ultrastructure, fluorescence, antigenicity...)
- Fluorescent labels in LM to electron-dense labels in EM

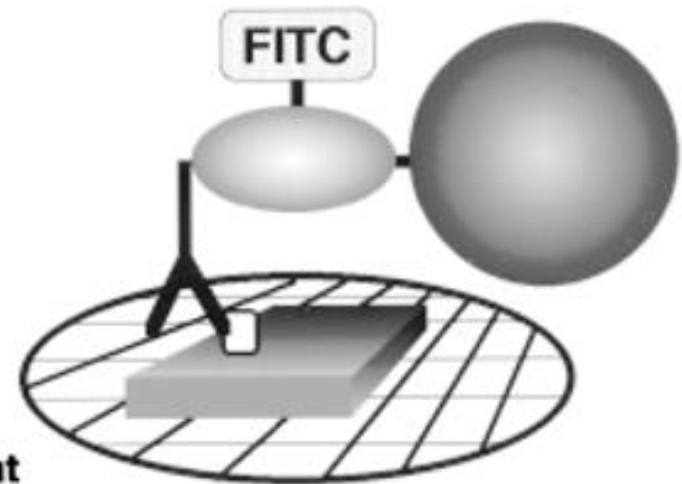


Immunolabeling using dual label

Fluorescence Microscopy



Electron Microscopy

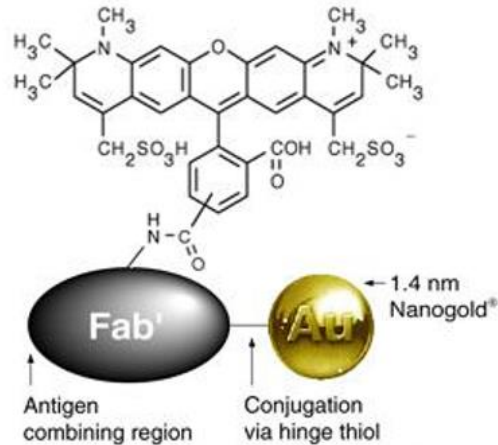


Silver Enhancement

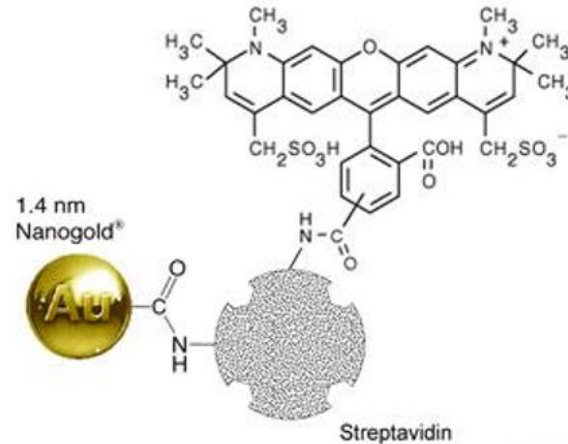
IMMUNOLABELING

Cav-1 localisation on the same ultrathin cryosection

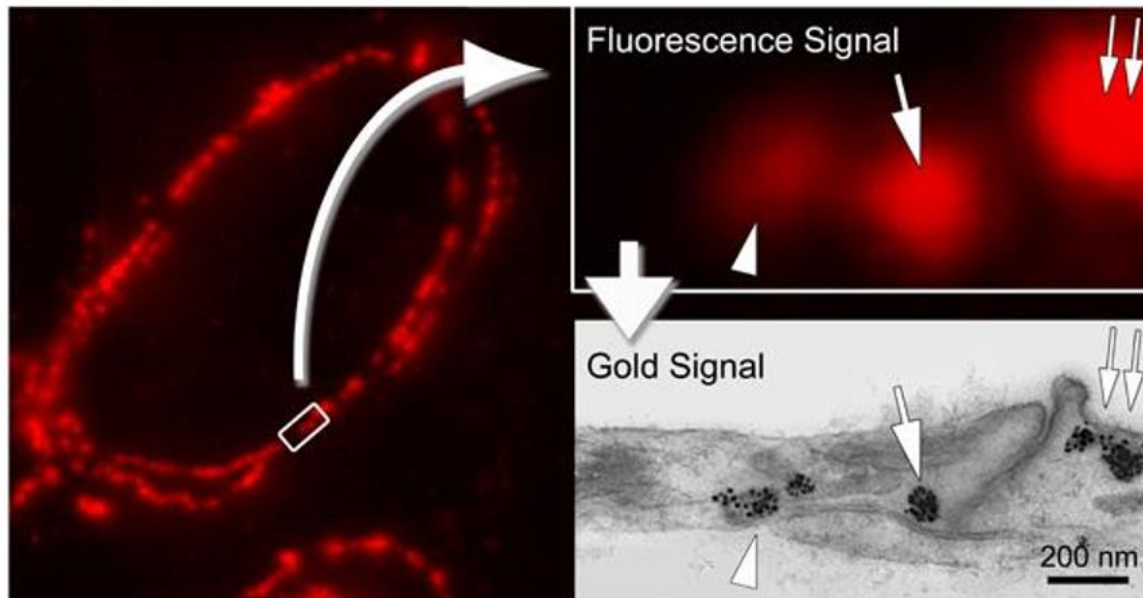
Alexa Fluor® 594 FluoroNanogold-Fab'



Alexa Fluor® 594 FluoroNanogold-Streptavidin



Correlative
fluorescence and
electron microscopic
immunolocalisation of
Caveolin 1 on the same
ultrathin cryosection.

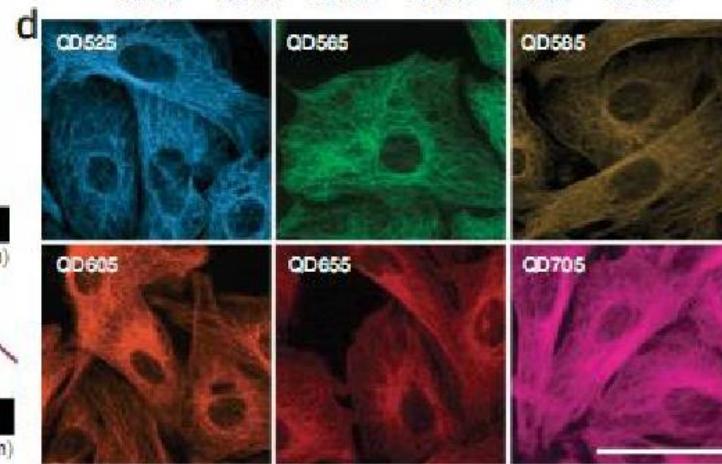
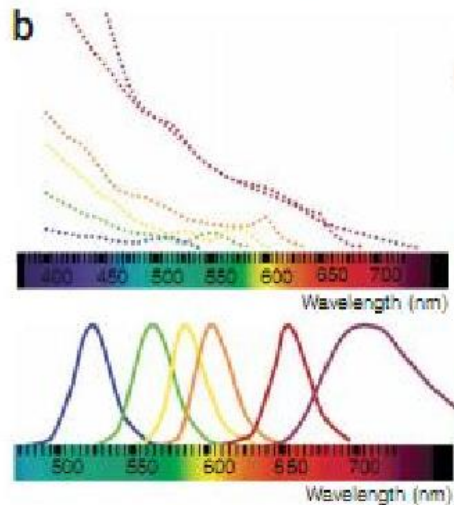
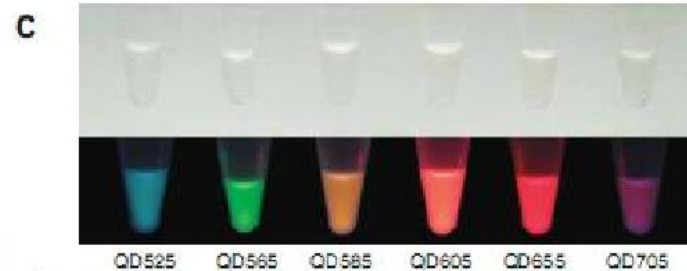
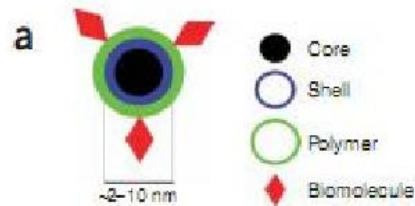
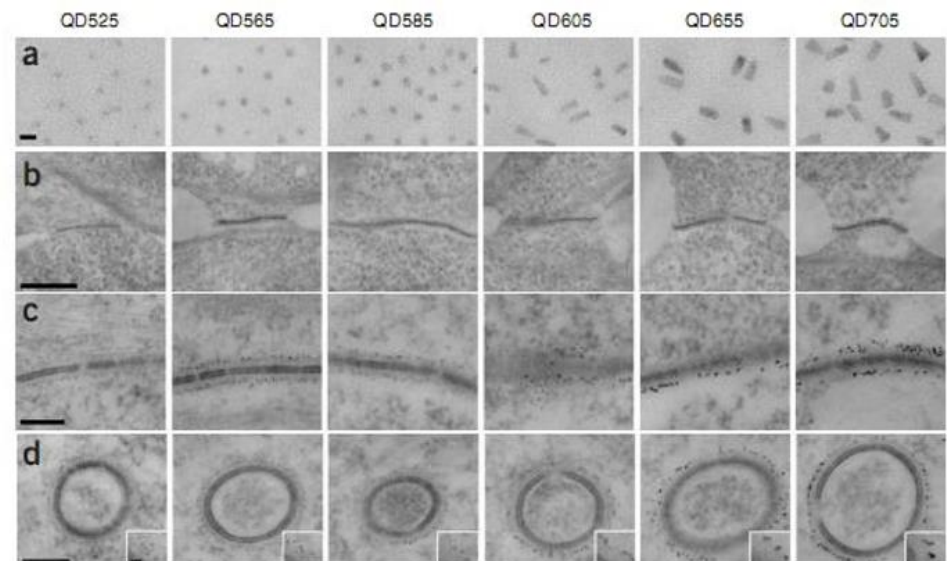


Labeled with
Nanoprobes Alex Fluro
594 FluroNanogold.

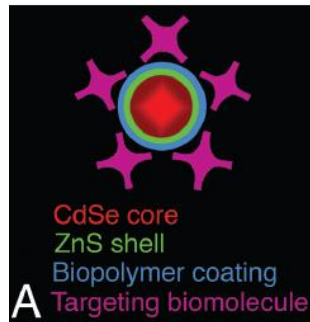
Silver enhancement
of immunogold is
REQUIRED

Quantum dots

- both fluorescent and electron dense
- different emission wavelength
- multilabeling analysis
- pre-embedding labeling (requires permeabilization)



Q-dots visualization by elemental mapping



Cd – M 4,5 403.7 eV N – K 401.6 eV

Se – L 2,3 1436 eV

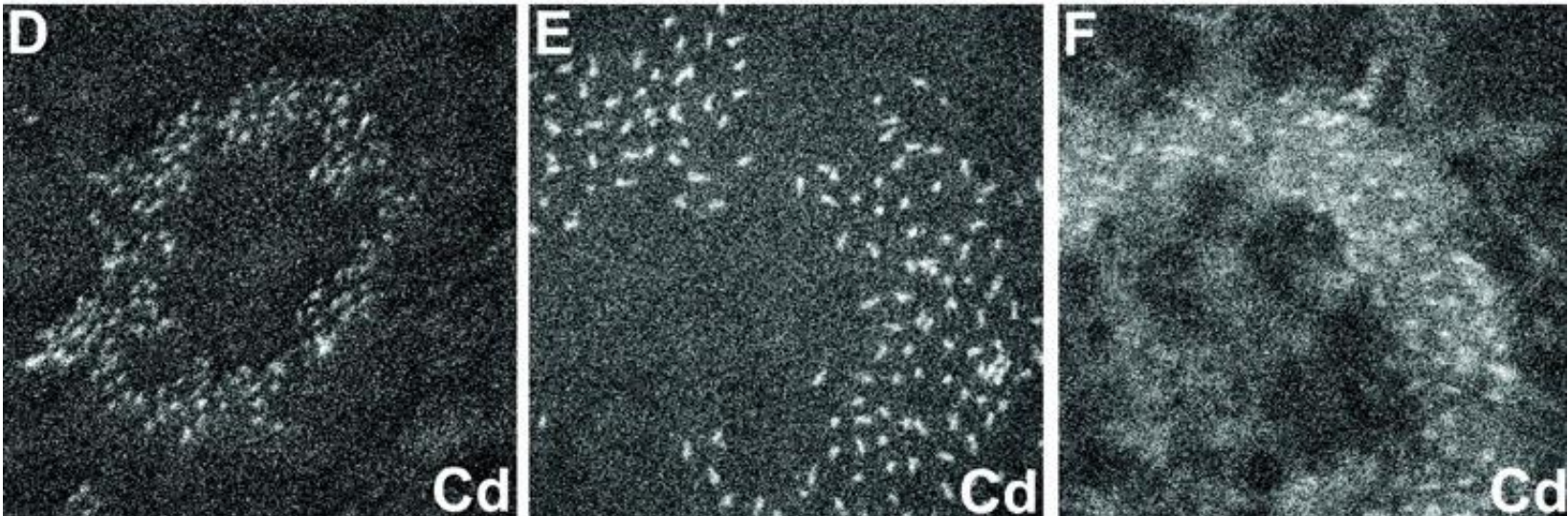
Te – M 4,5 577 eV

Zn – L 2,3 1020 eV

S – L 2,3 164.8 eV

N: 385 and 415 eV, Cd: 415 and 510 eV

EFTEM



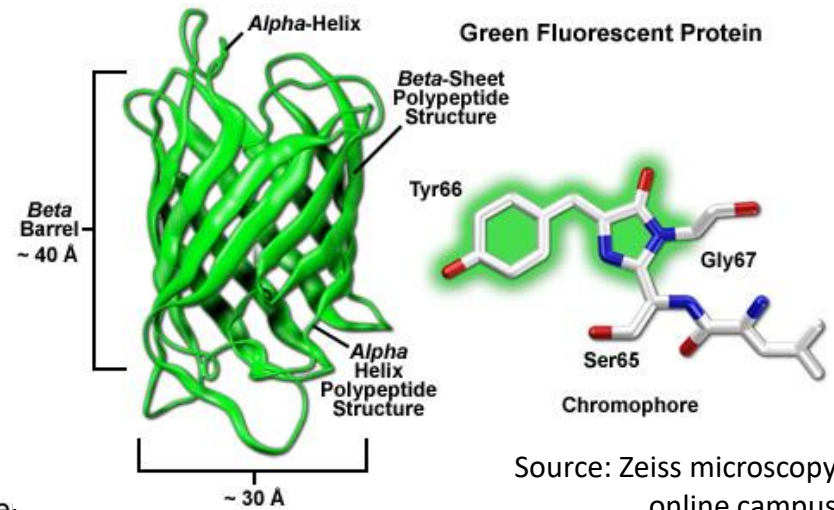
Genetically encoded fluorescent tags photooxidation

Green Fluorescent Protein (GFP)

Drawback: size (~30 kDa)

Bright fluorescence, well-established technology

Poor photo-oxidizer
Very low reactive oxygen yield
Size (~30 kDa)



Source: Zeiss microscopy
online campus

MiniSOG

size (~15 kDa)

Very good singlet oxygen
generator; Strong EM signal after
photo-oxidation.

Weak fluorescence
Bleaches rapidly

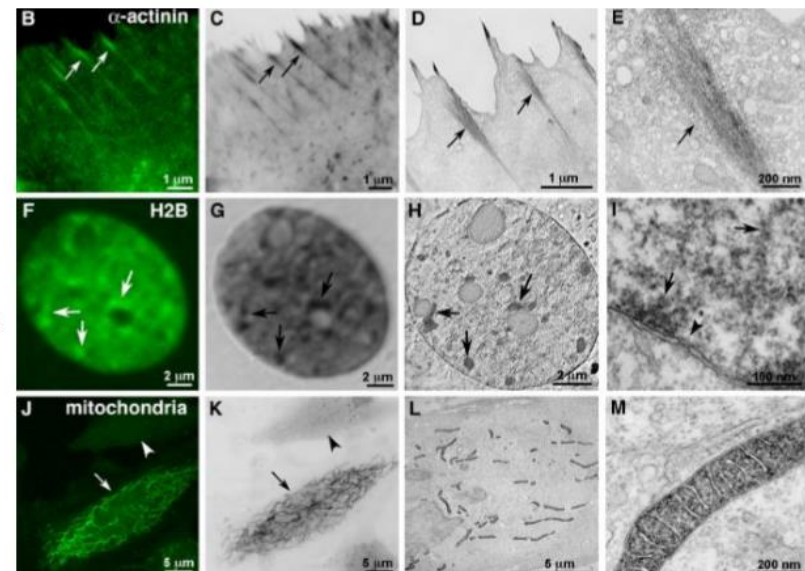
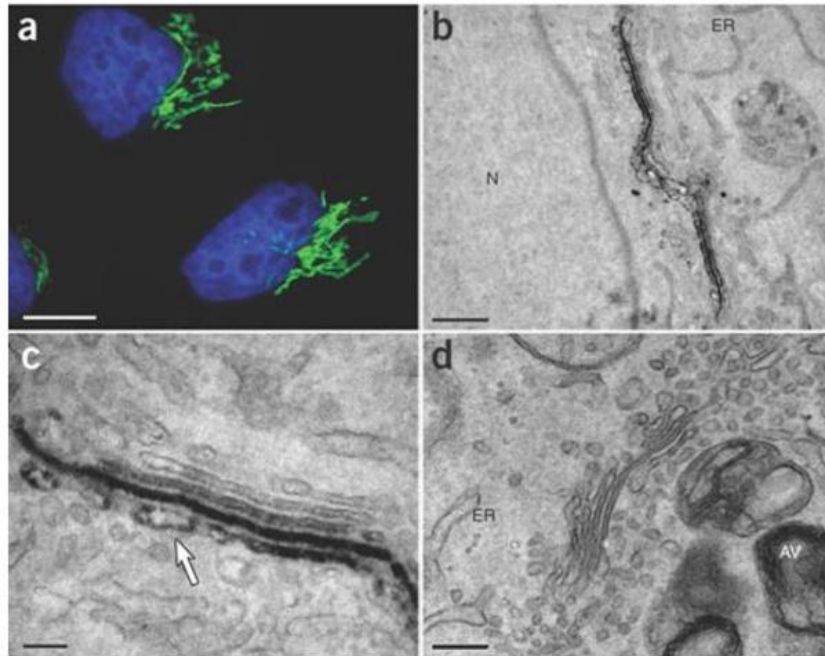
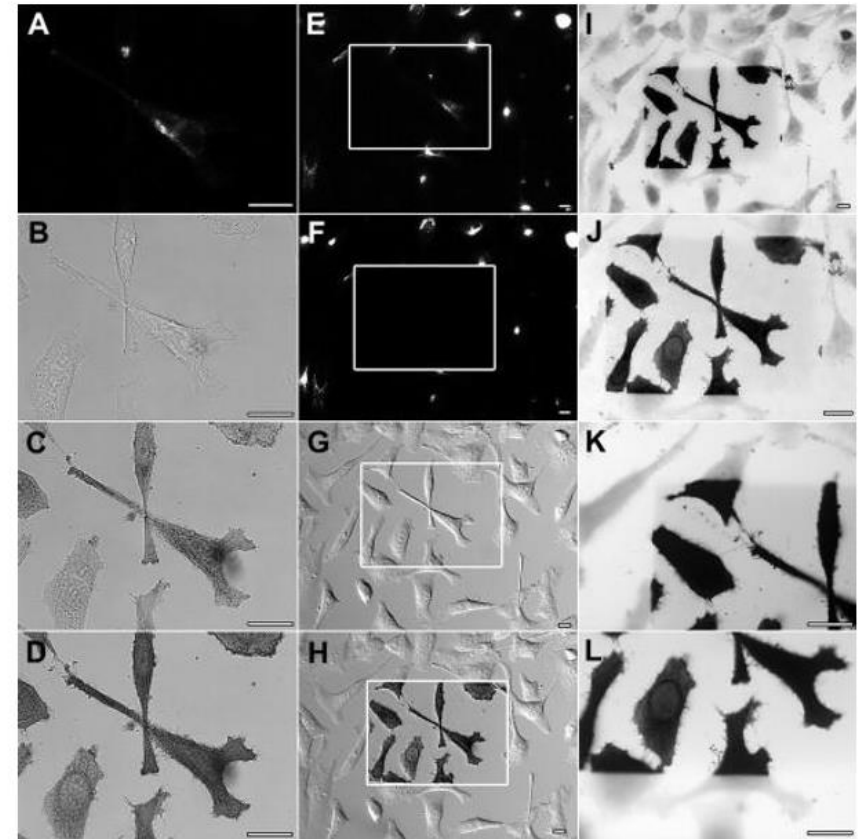


Photo-oxidation of GFP

+ DAB + light = precipitation



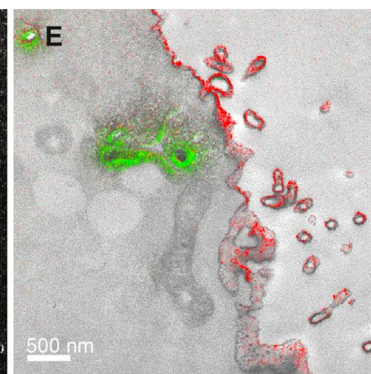
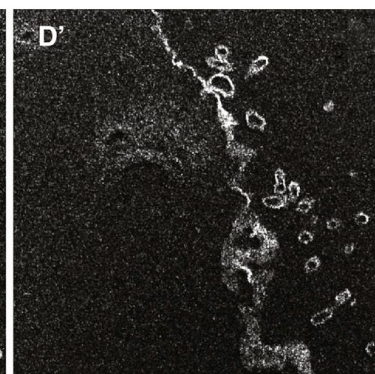
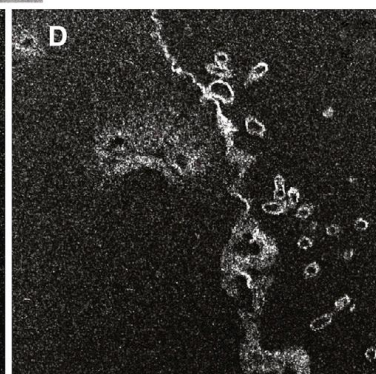
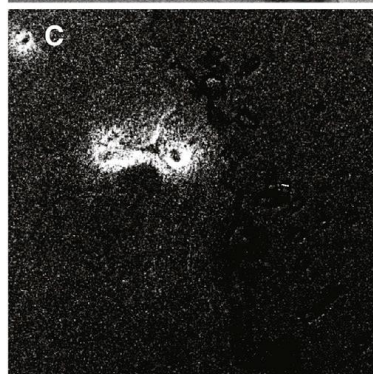
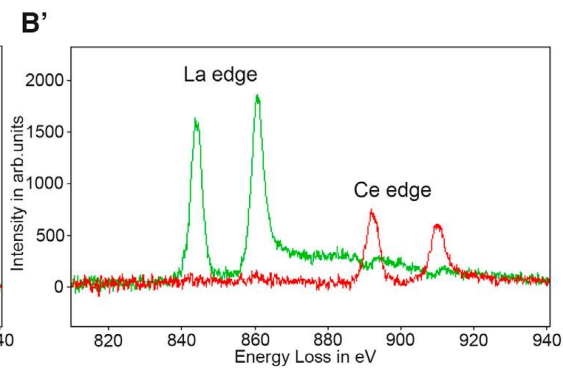
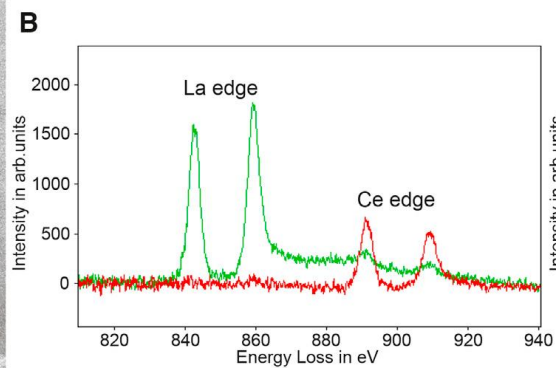
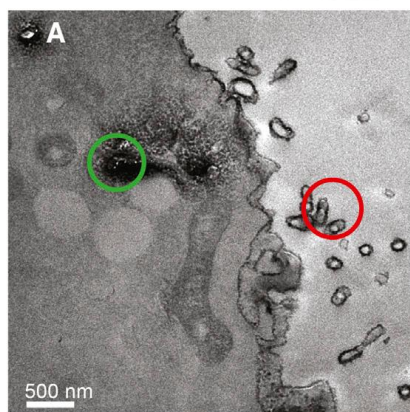
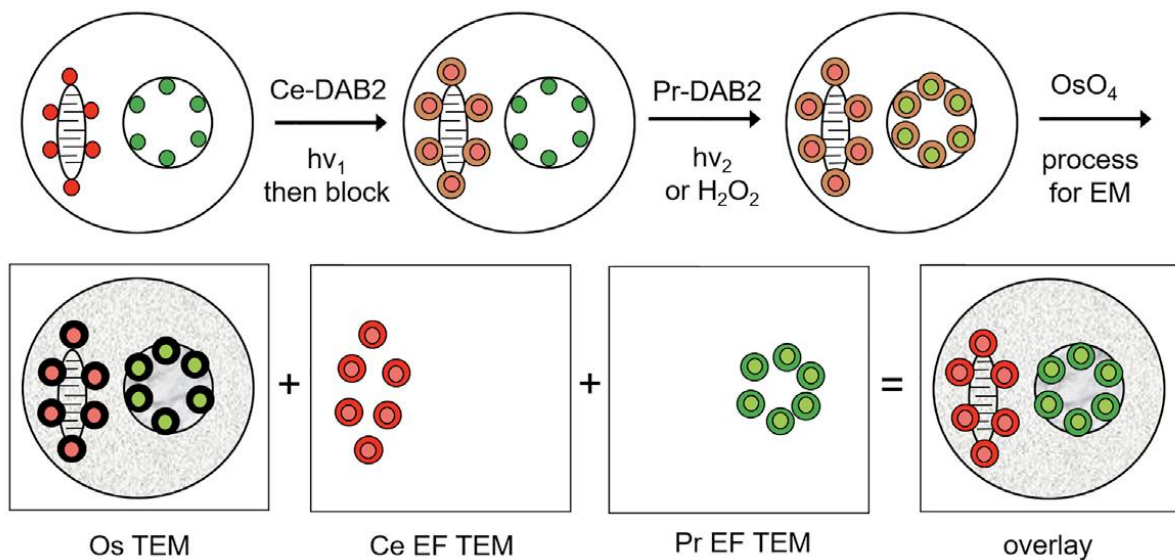
Correlative microscopy of GFP through photo-oxidation. HeLa cells expressing the Golgi-resident enzyme GalNAc-T2 fused to EGFP.



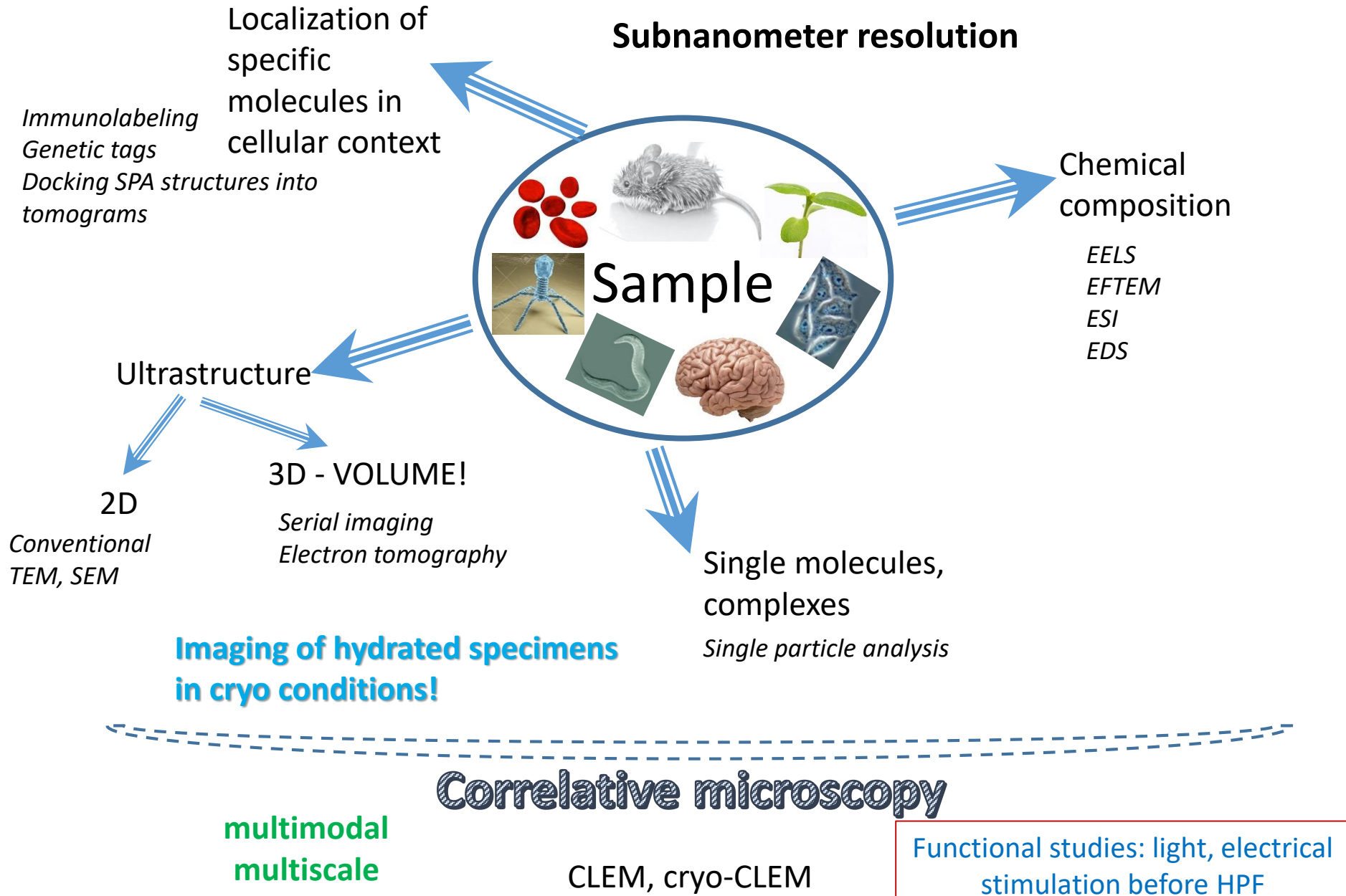
Bleaching process in higher (A–D) and lower magnification (E–H), and the same area after osmification and Epon embedding (I–L).

Fluorescent labels

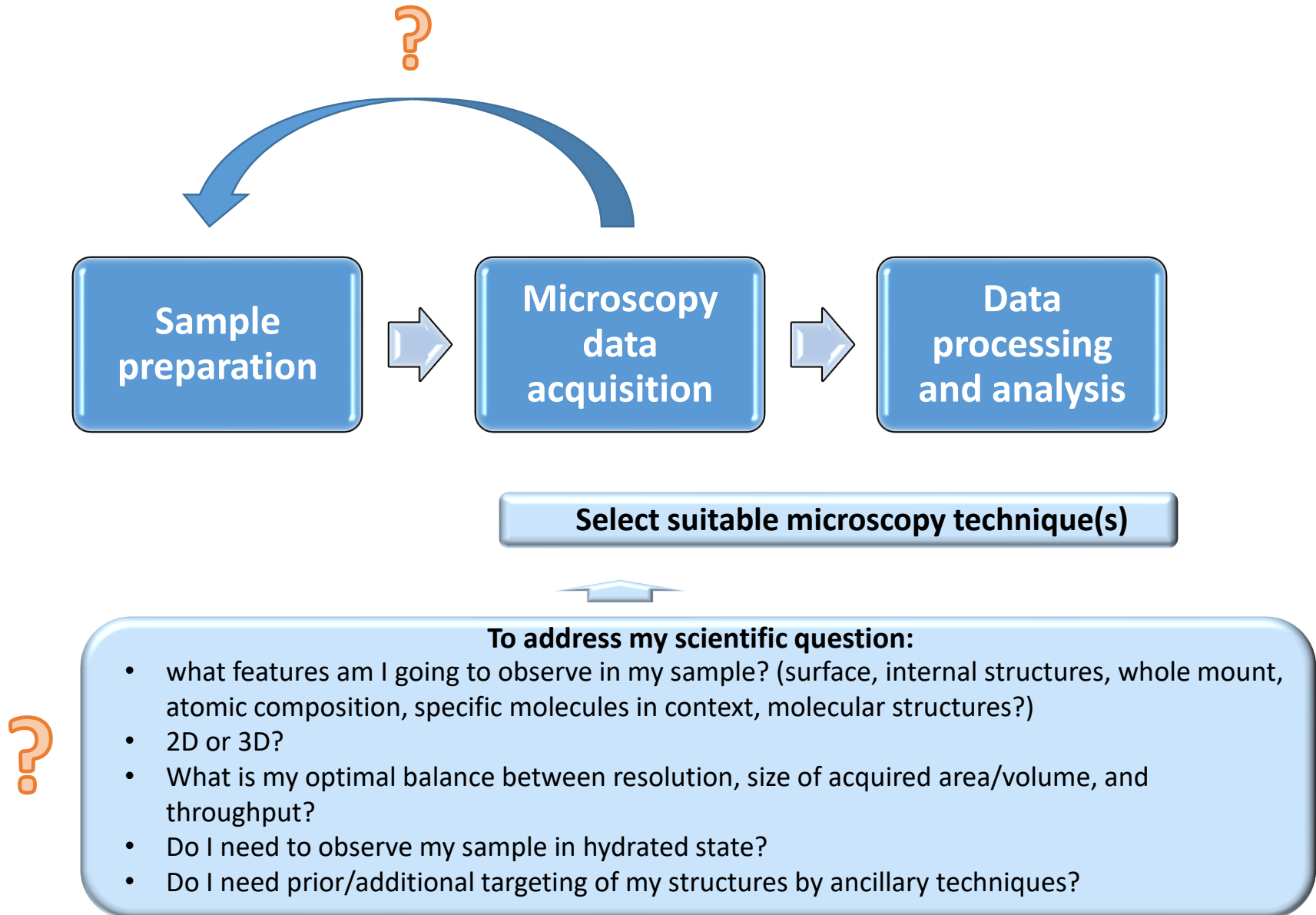
Electron-dense labels



Why electron microscopy?



- When you have several options, consider their advantages and disadvantages (quality of obtained information, cost, time and labor investments).
- Good to start with a more simple method for first tests, screenings



The Electron Microscopy Core Facility

Contact: Vlada Filimonenko
vlada@img.cas.cz



CzBI Call for Financial Support

- Call opens: **12 September 2024**
 - Application deadline: **15 October 2024**
 - Result announcement: **by 30 December 2024**
 - Project execution period: **1 January 2025 - 31 December 2025**
- <https://www.czech-bioimaging.cz/czbi-user-support>

