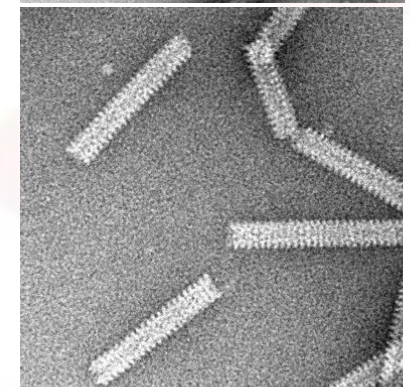
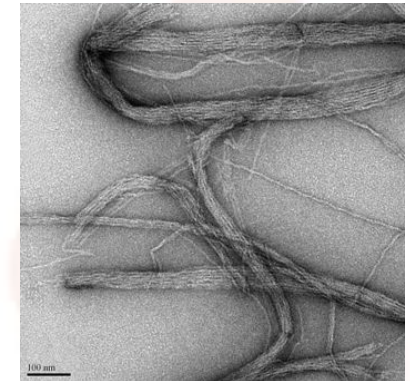


Sample preparation for Structural studies by EM



Réseau National de Formation en
Biologie Structurale Intégrative

Guy Schoehn IBS/UVHCI/ISBG
Guy.schoehn@ibs.fr



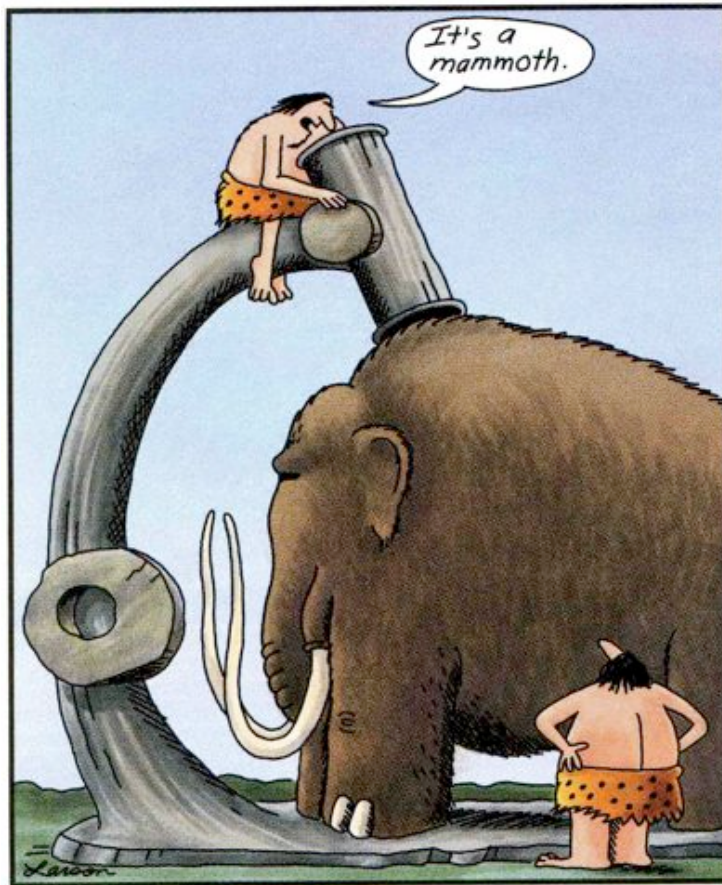
Preparation of biological samples for electron microscopy

- They have to:
- resist to the vacuum and to the electrons
 - stay in their native form



Preparation of biological samples for electron microscopy

They have to: - being thin enough



Early microscope



- Negative staining
- Cryo-electron microscopy
- Metal shadowing

The two mains techniques

Negative staining

Advantages :

Fast

Small amount of protein
(concentration 0.01-0.1mg/ml,
few μ l)

Small proteins are visible

High contrast

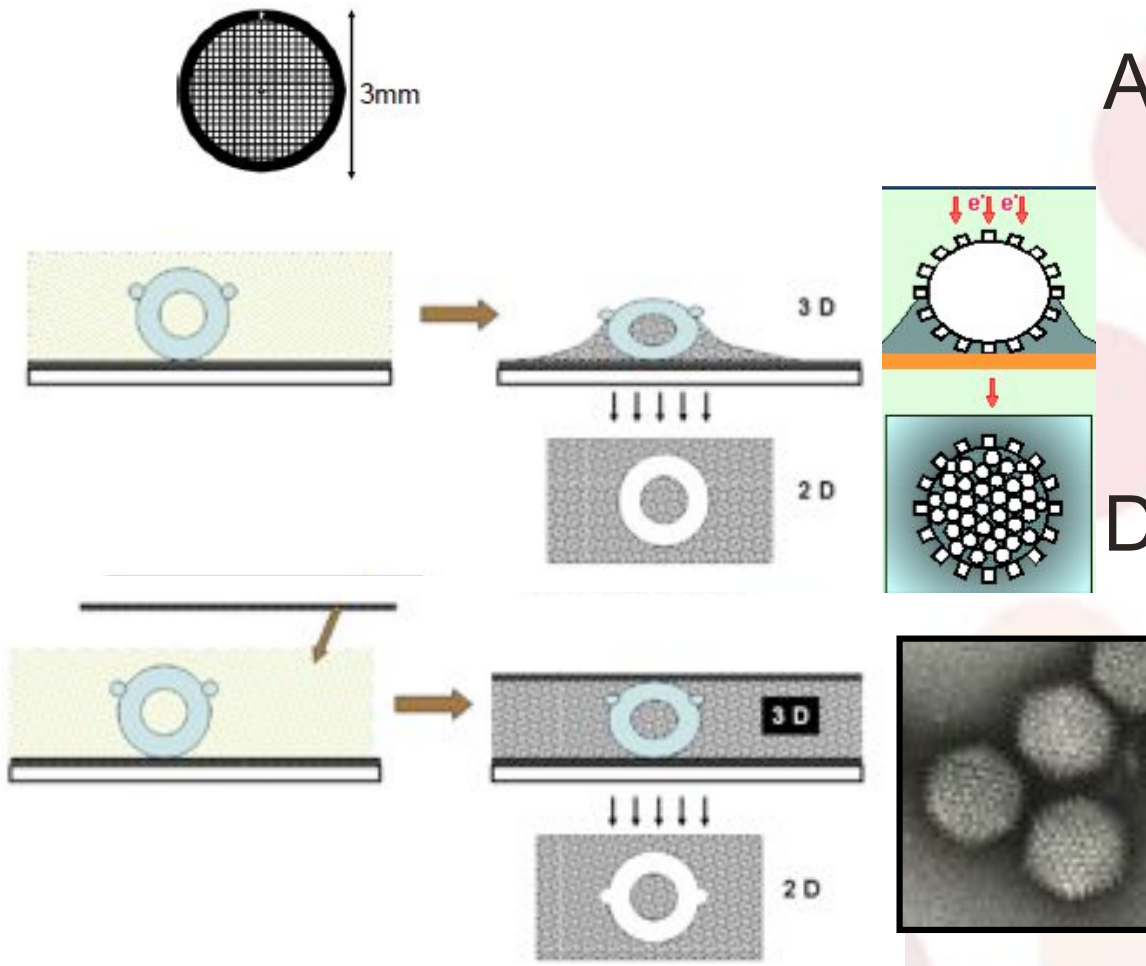
Disadvantages :

Artefacts due to the stain

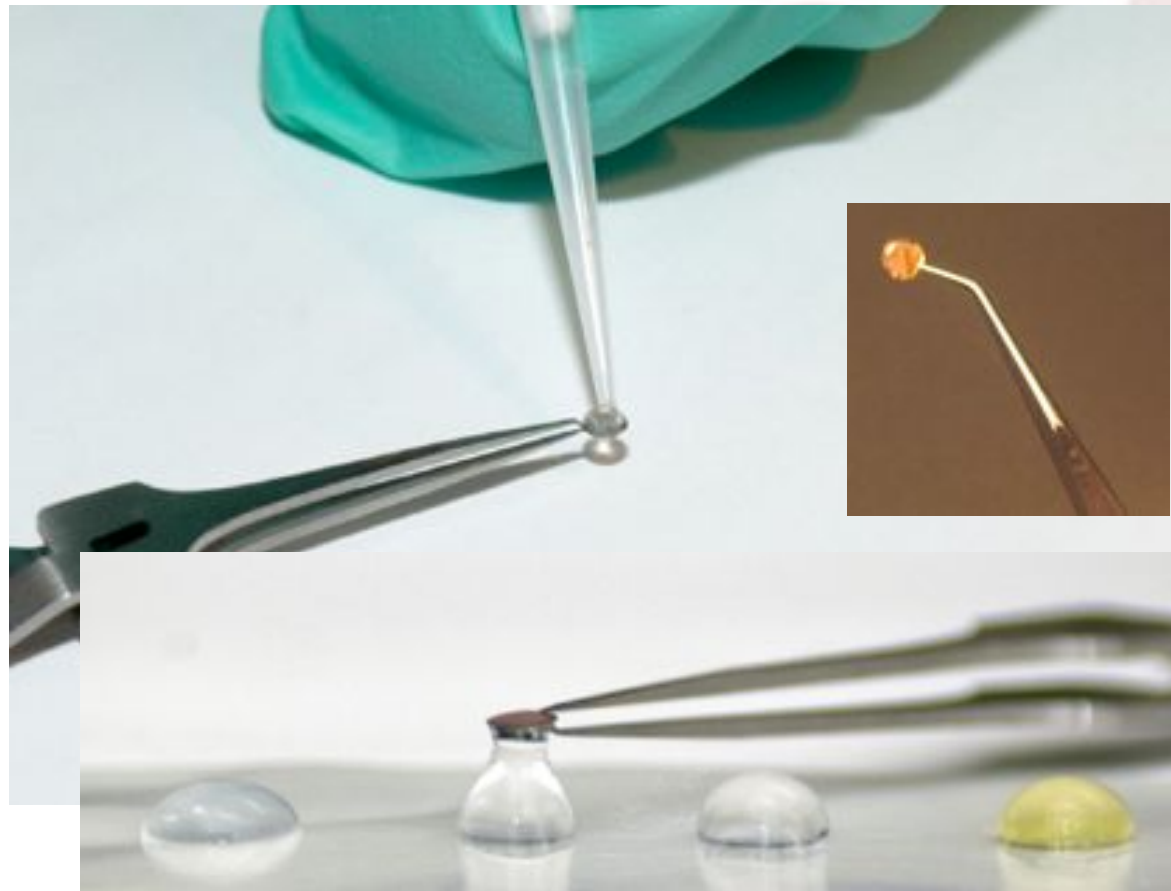
Flattening and drying

Footprint of the sample in the
heavy atom

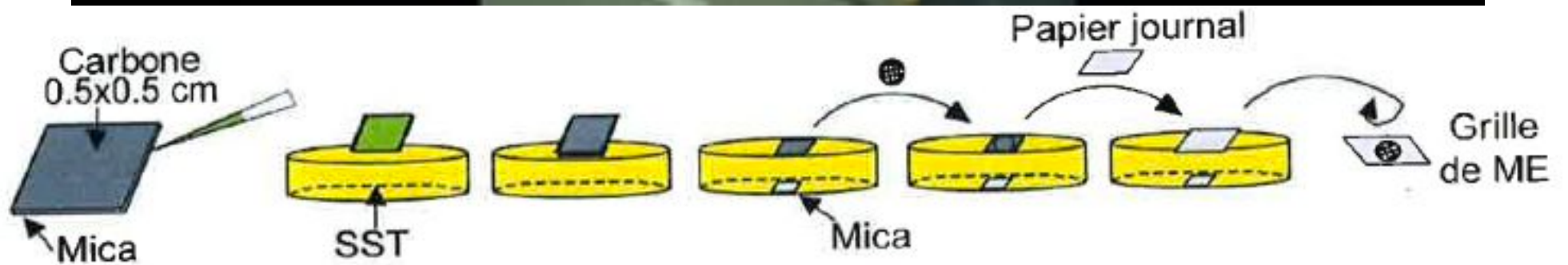
Low resolution (15 Å)



Preparation method



Preparation method



Stains:

Uranyl formate (finer grain, but unstable)

Uranyl acetate (coarser grain but stable)

- Higher contrast and radiation resistant
- but acidic (pH 4)
- It also function as fixative !

Sodium silicotungstate

Sodium phosphotungstate (thin layer of stain)

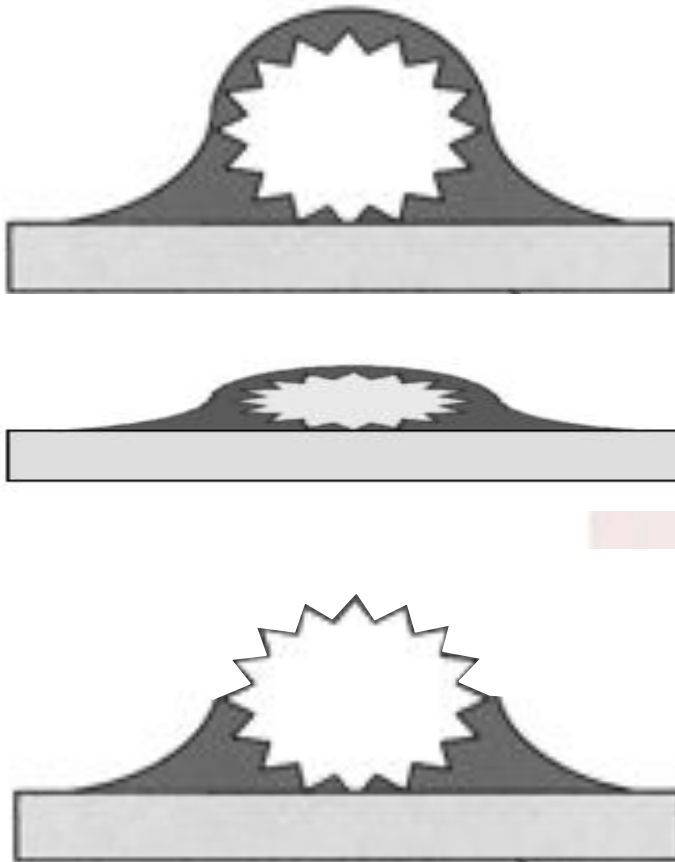
Ammonium molybdate (thick layer of stain)

Aurothioglucose

- - neutral pH
- - but lower contrast and less radiation-resistant

What we obtain in NS

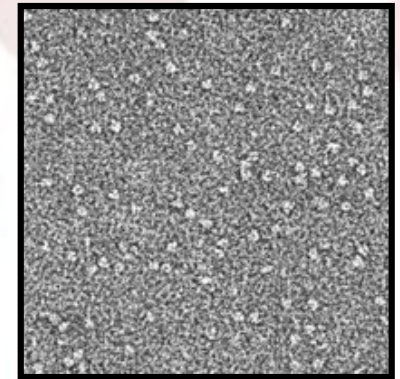
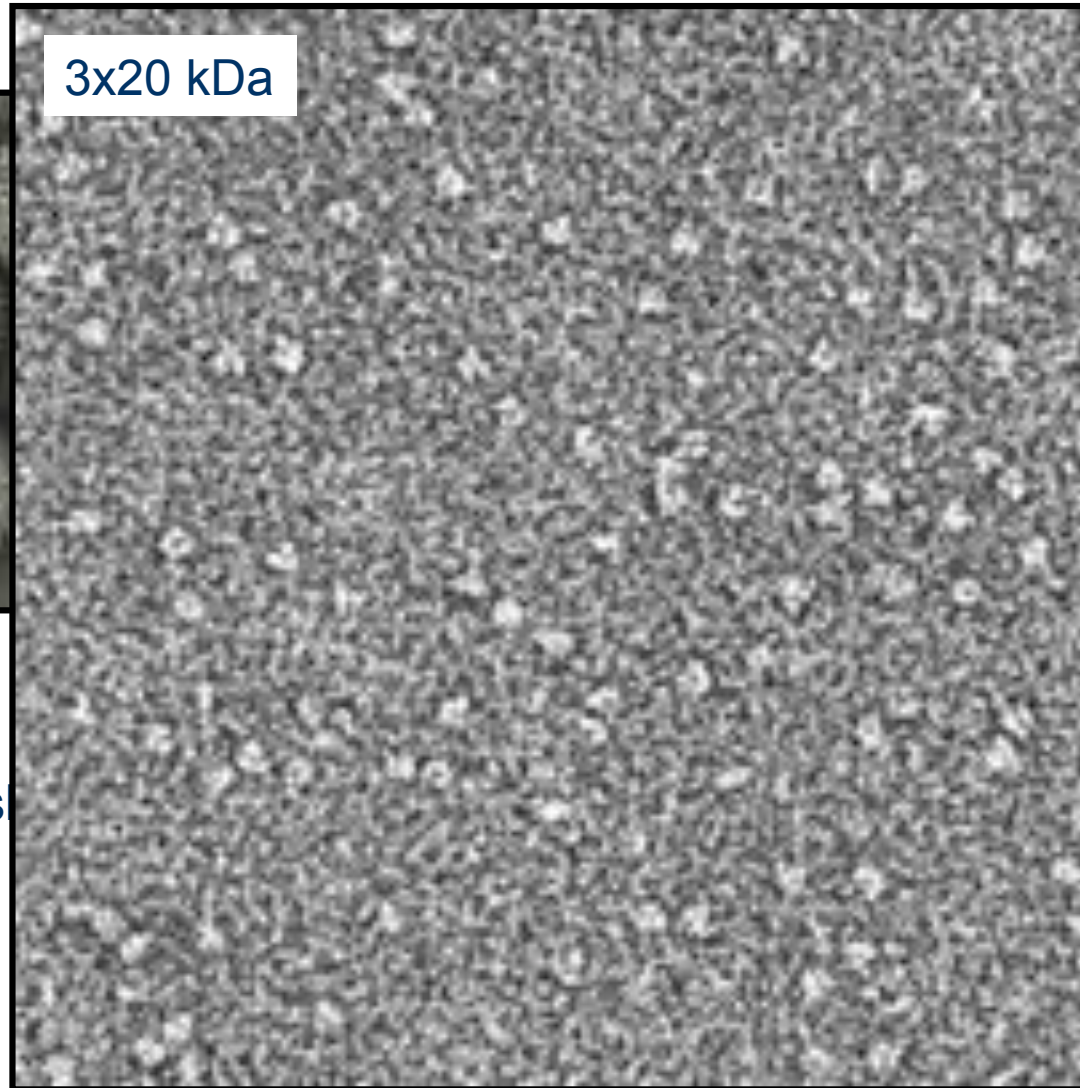
Negative stain



Examples of negatively stained samples



Meas
Pitch



Tetrahedral
aminopeptidase
480 kDa

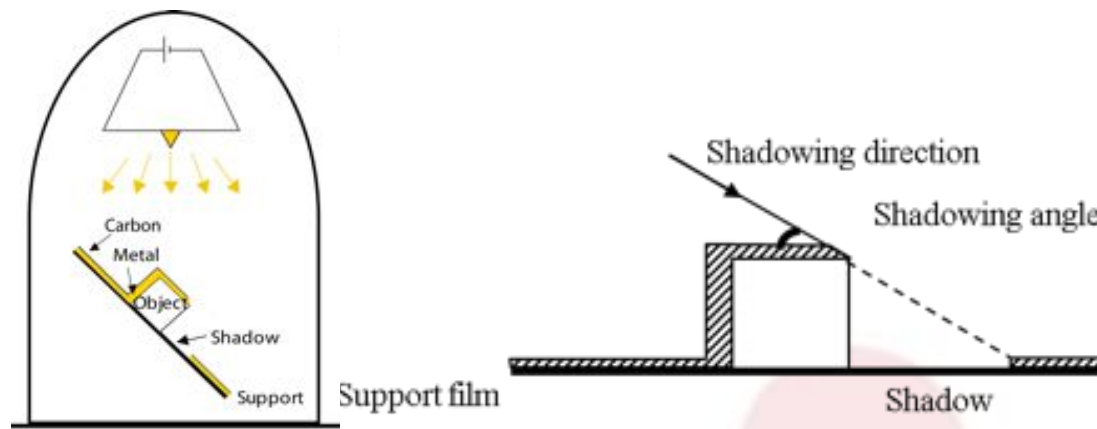


Negative staining

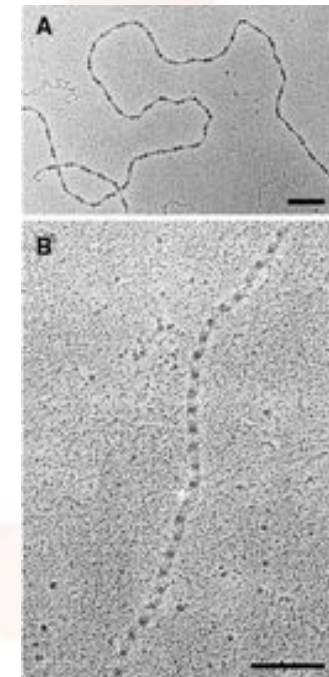
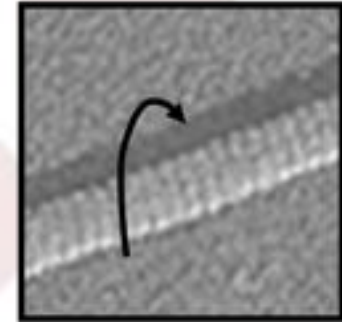


Still very usefull prior to cryo EM in most of the cases

Shadowing



- Aim :
 - Visualize the surface of the particles
 - Handiness determination
- Technique :
 - Heavy atom evaporation
 - The sample is tilted in the evaporator
 - The tilted sample can also be rotated



Negative staining and shadowing



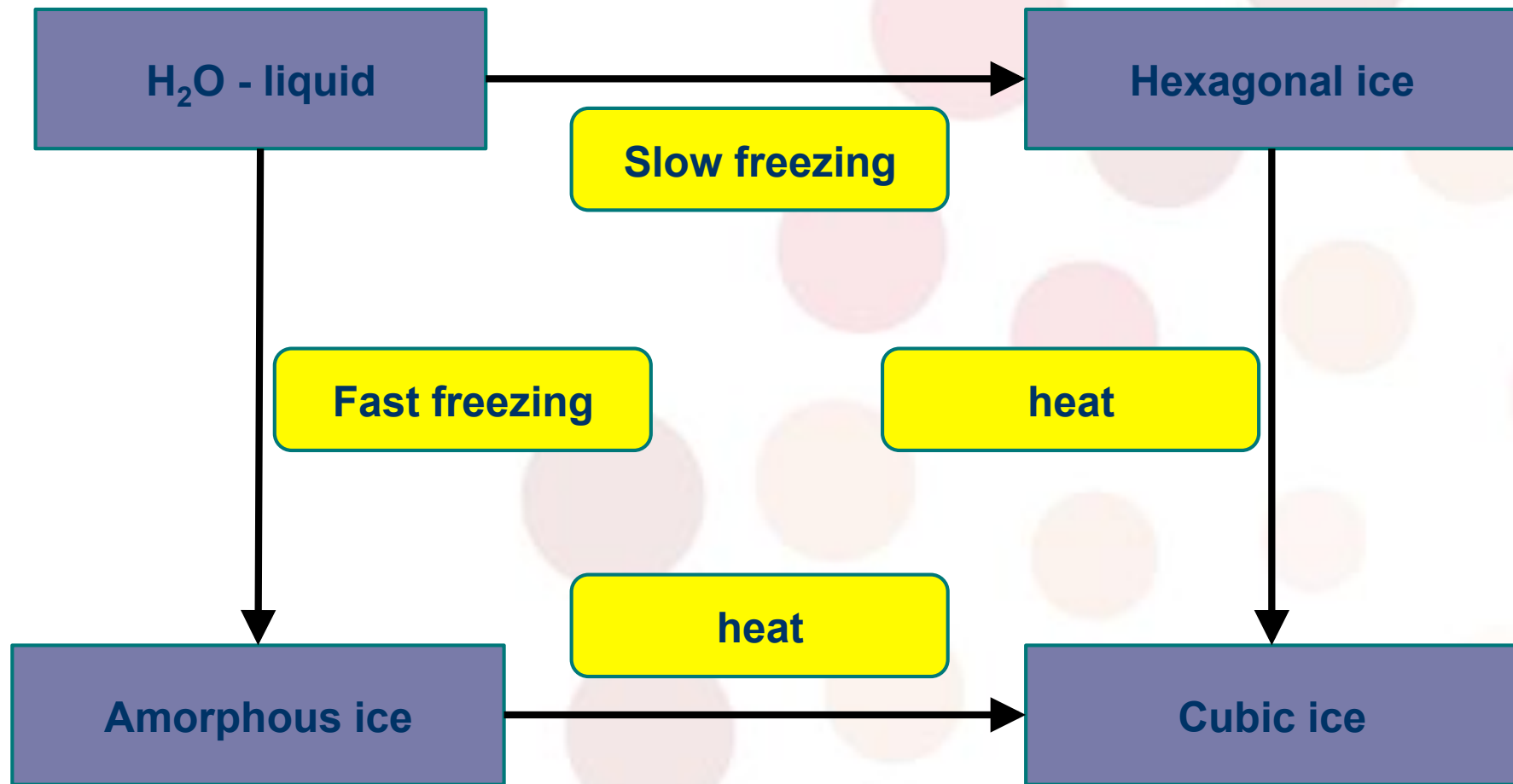
What we obtain

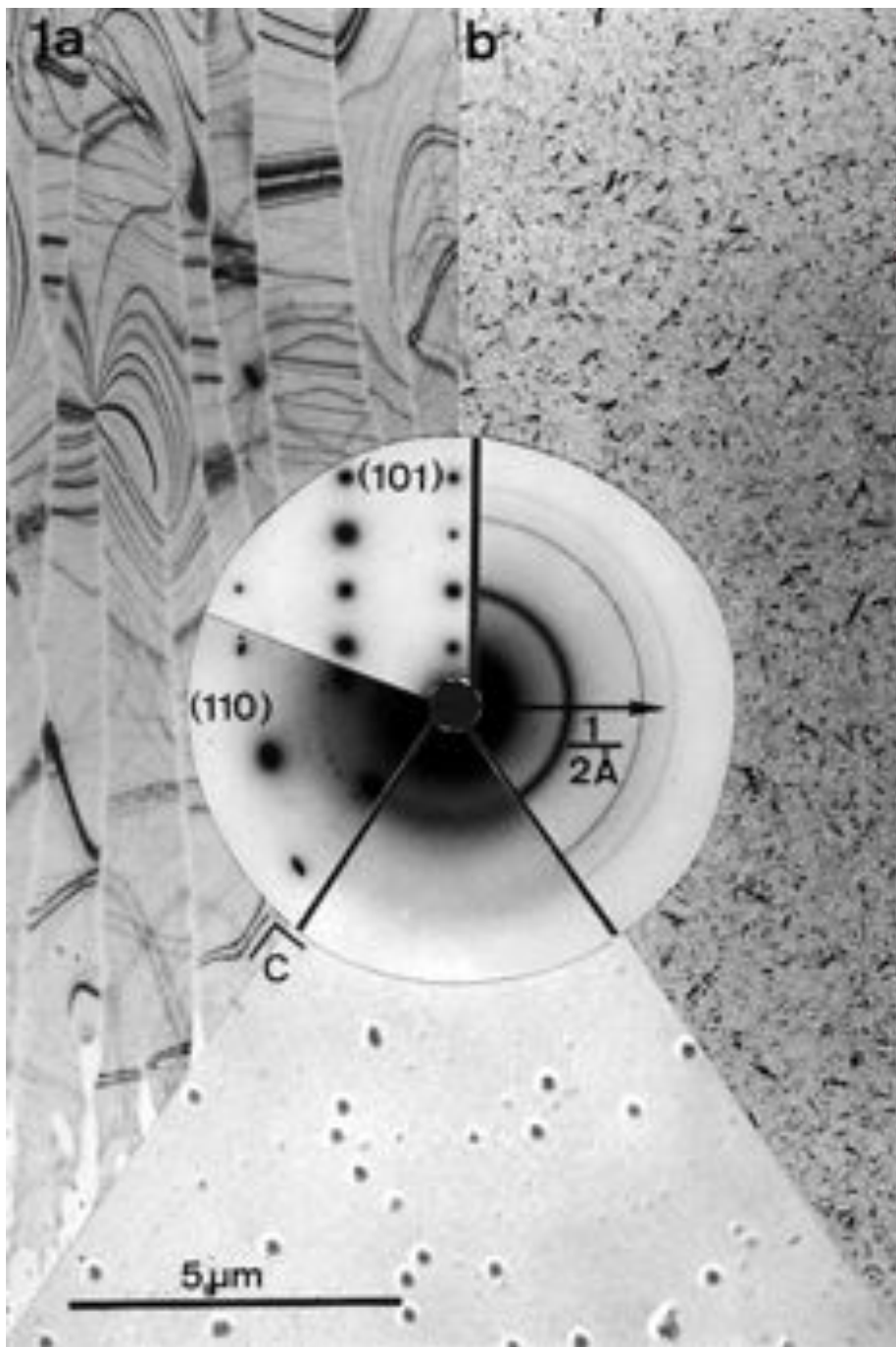


What we would like to have

For real structural work one need something else

Vitrification

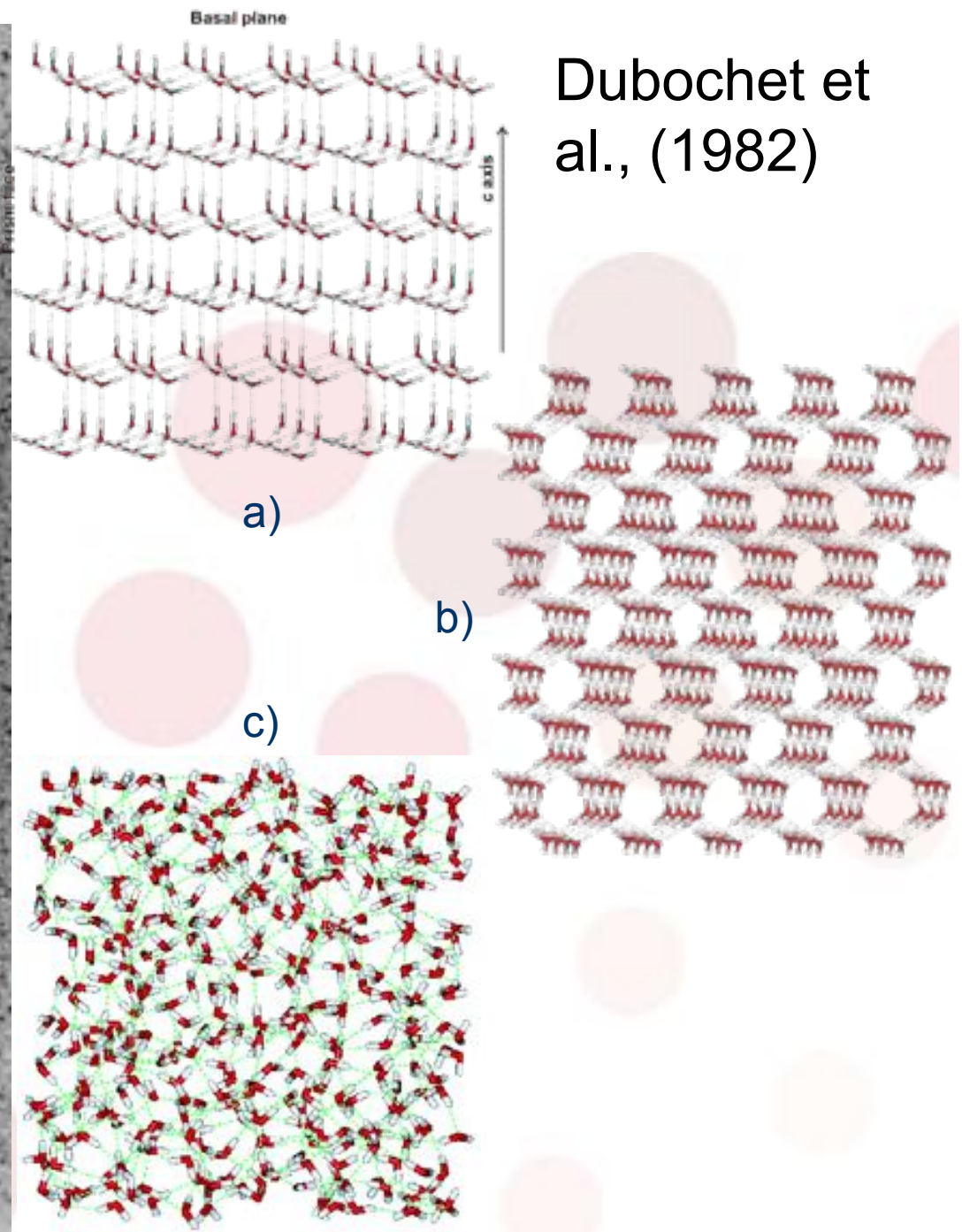




a) hexagonal

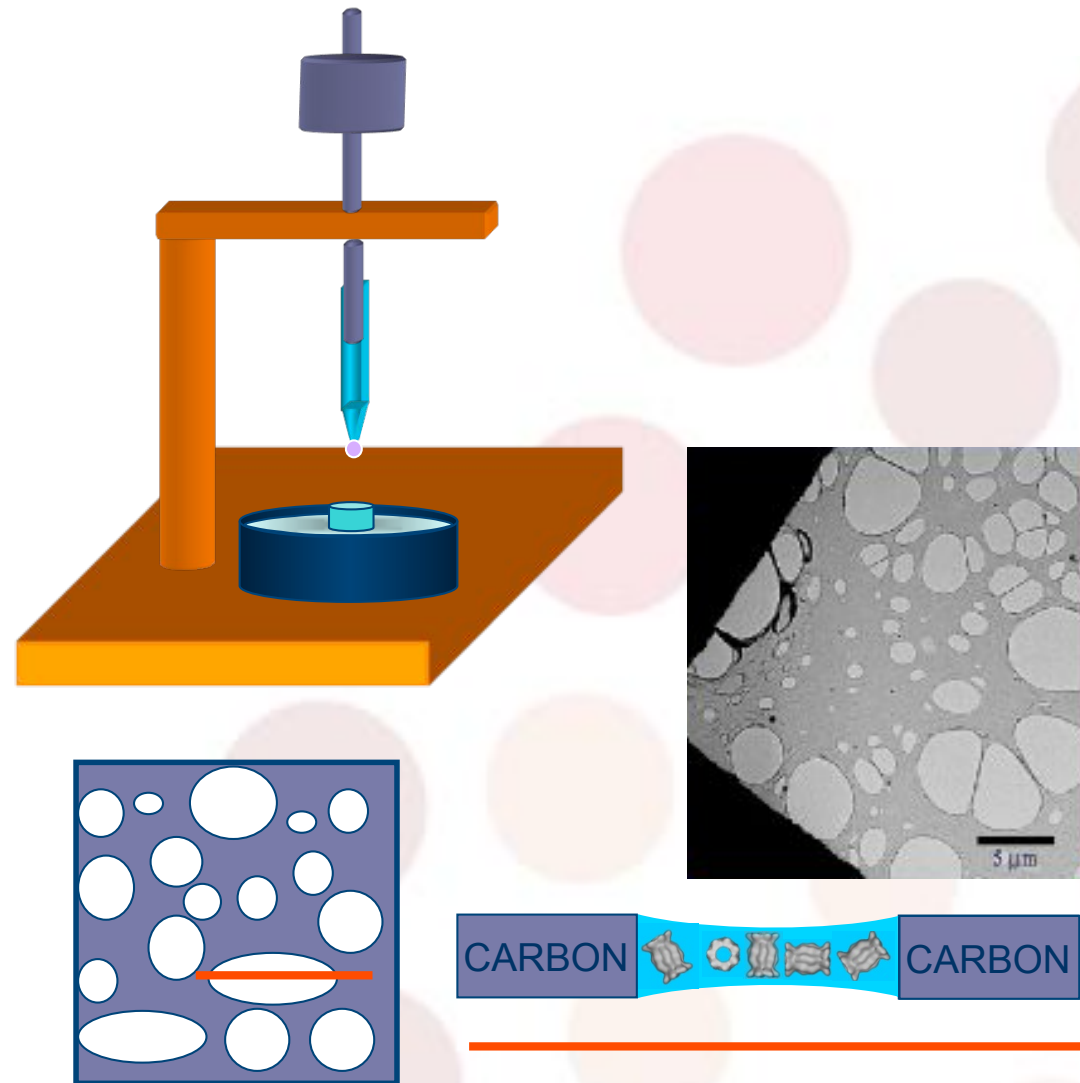
b) cubic

c) amorphous



Dubochet et al., (1982)

Cryo Electron Microscopy



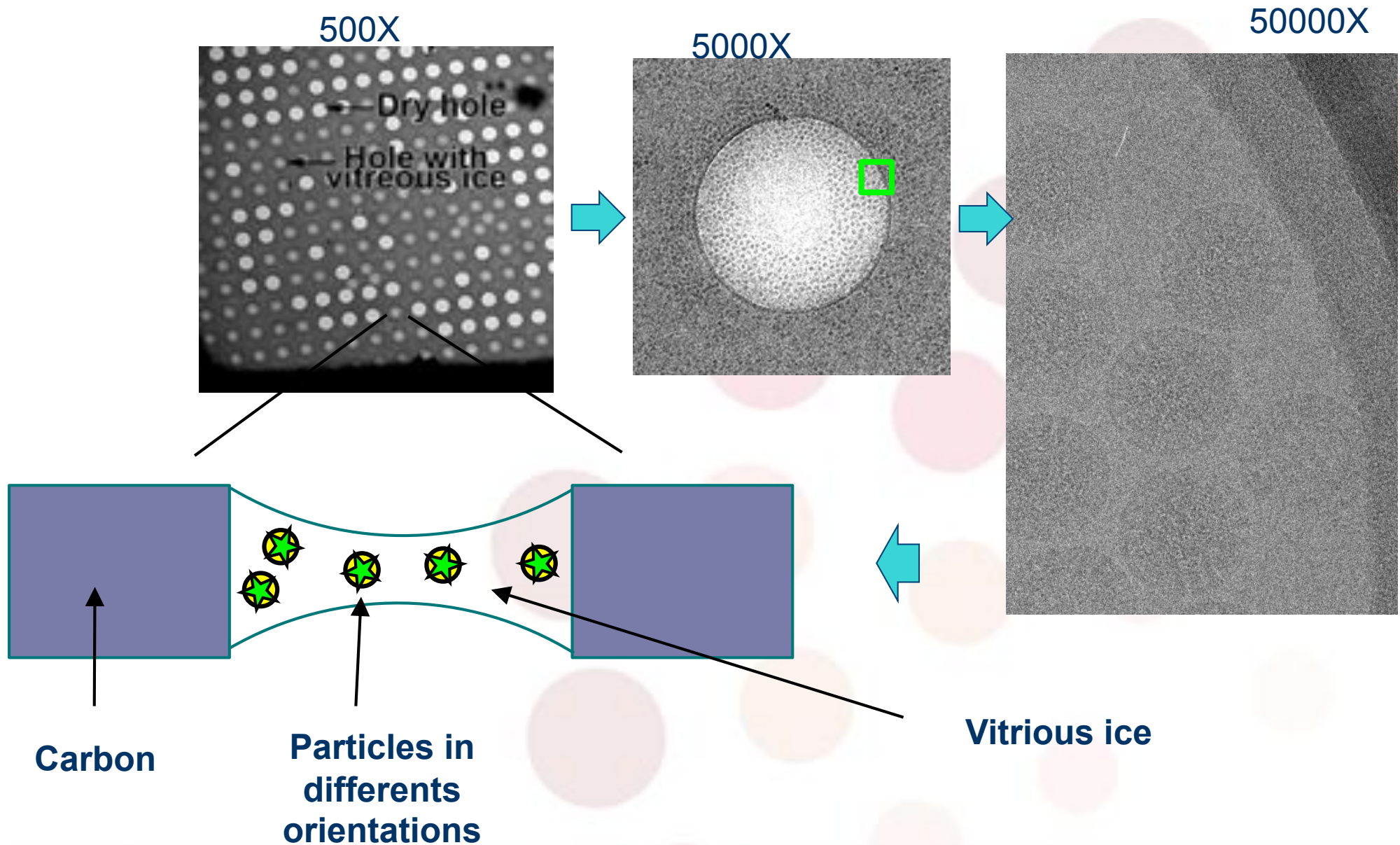


The freezing instrument





How does a frozen grid look like?



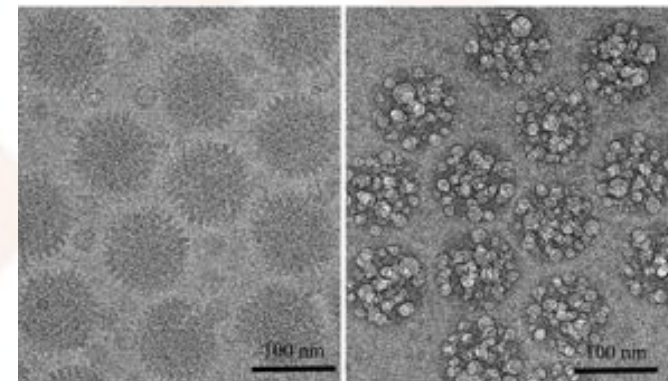
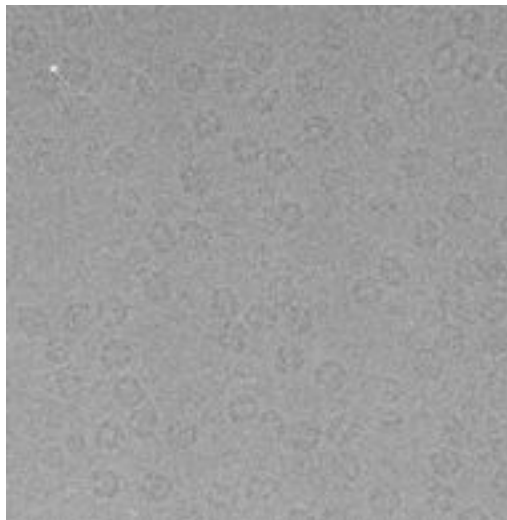
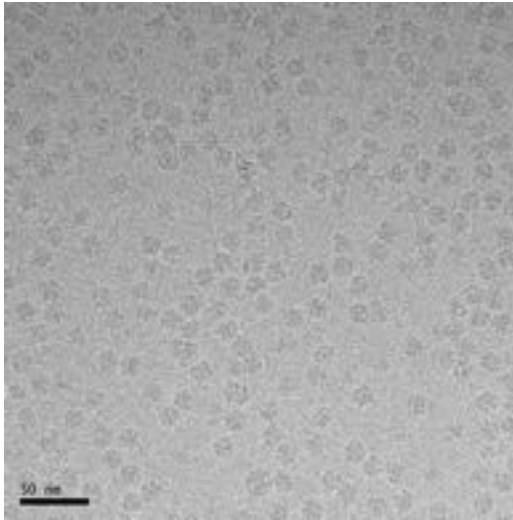
Cryo: advantages / disadvantages

Advantages :

- Native state of the sample
- High resolution
- Small amount of sample

Disadvantages :

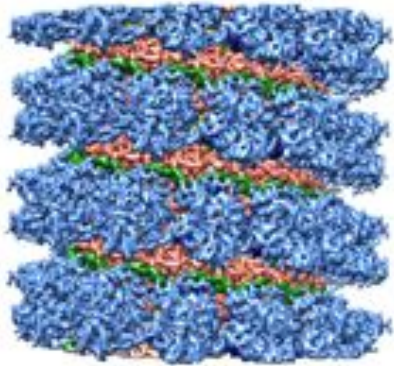
- Low contrast
- High sensitivity against electrons
- Size of the sample
- More expensive



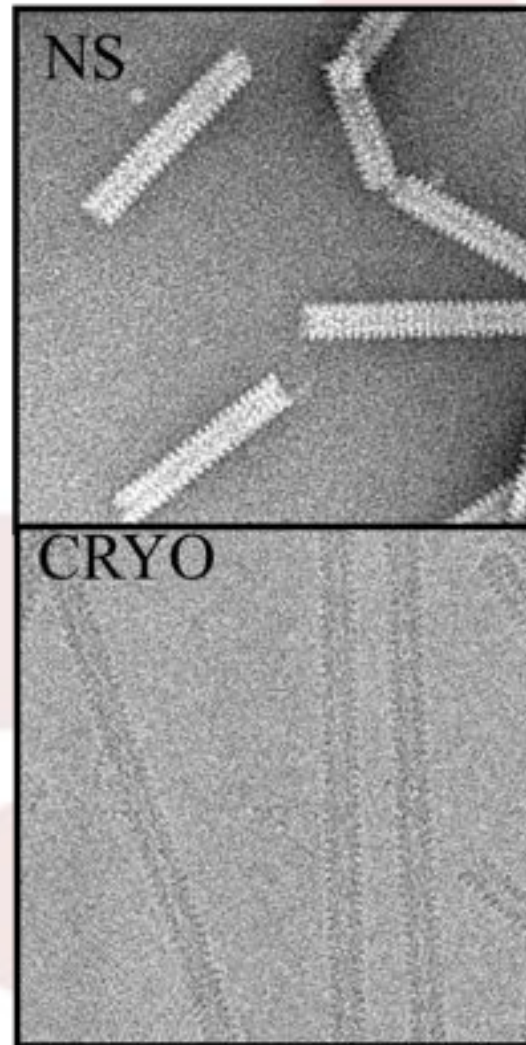
Negative staining vs. cryo



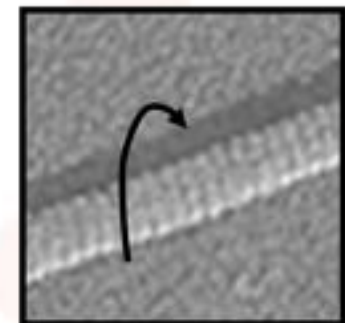
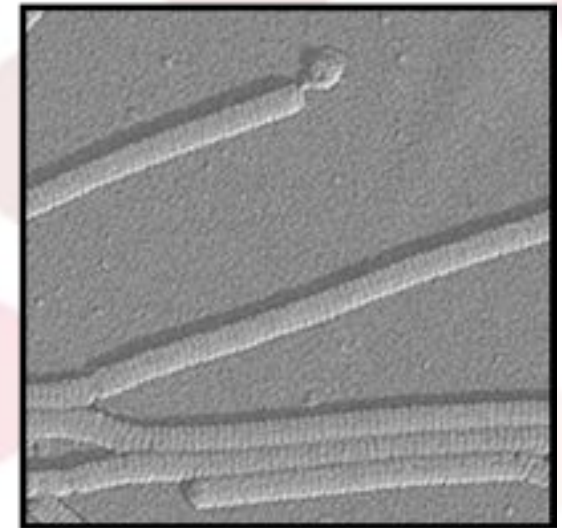
Comparison between negative staining, cryo and shadowing



4.3 Å, Gutsche et al., Science 2015



Shadowing



What is a good cryo grid?

Good amorphous ice

- not crystalline ice
- no « leopard skin » pattern
- no contamination

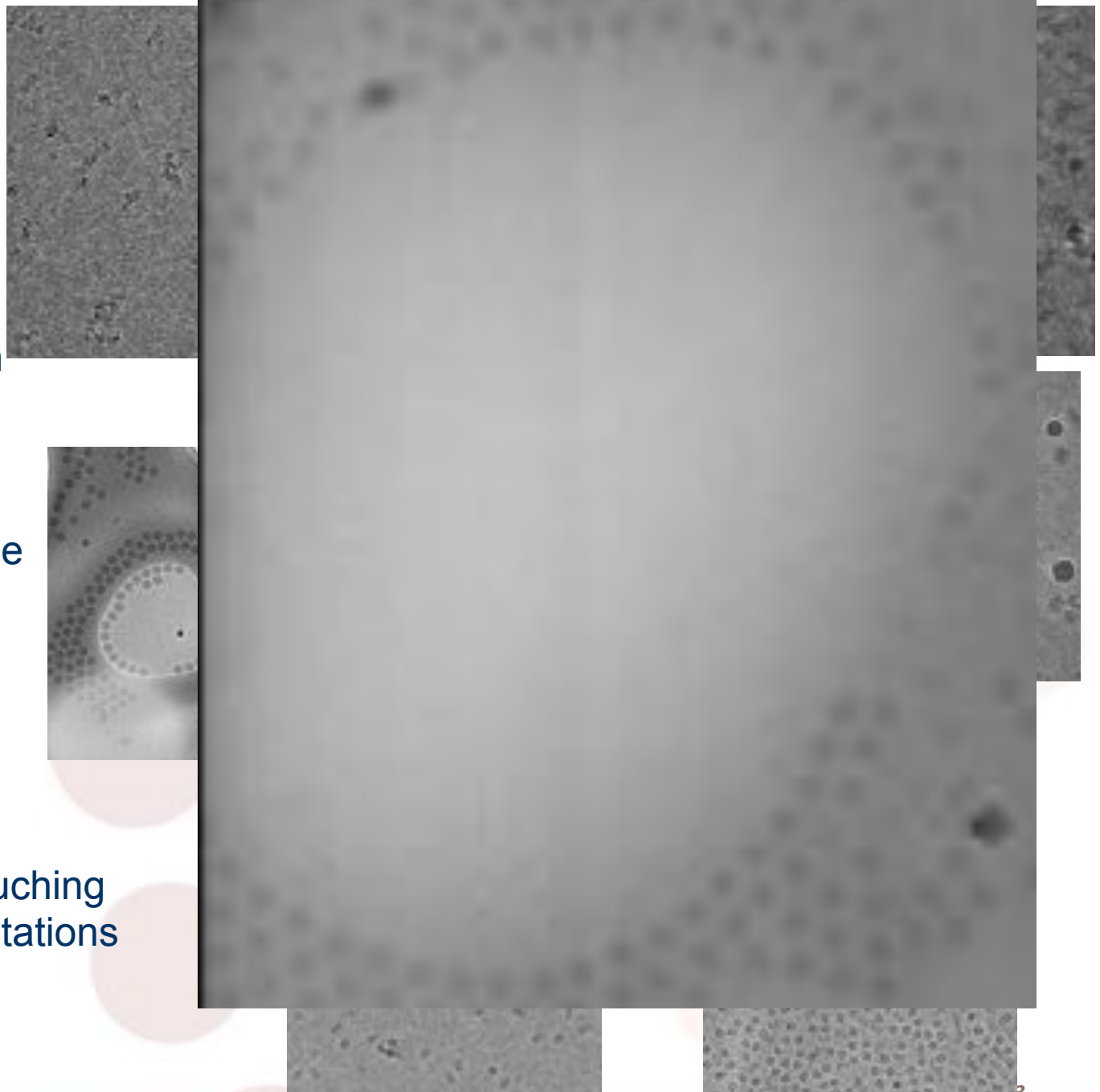
Appropriate ice thickness

- typically as thin as possible

Clearly visible particles

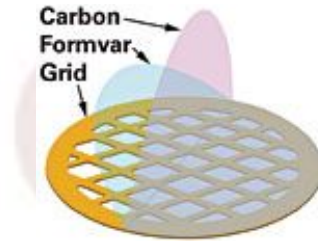
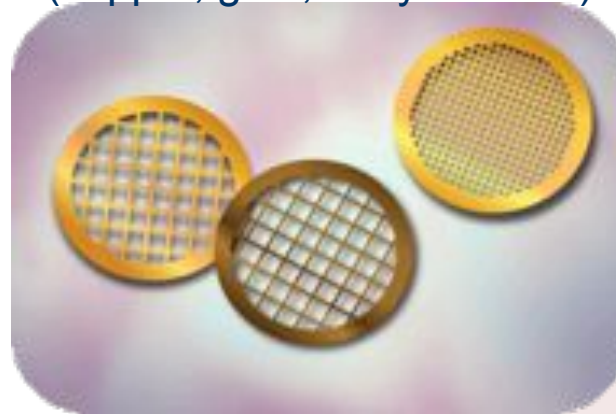
Good particle distribution

- in holes
- dense but particles not touching
- randomly distributed orientations



Variables in grid preparation

EM grid
(copper, gold, molybdenum)



Grid

- type of grid and substrat
- batch and age of grid
- glow discharging

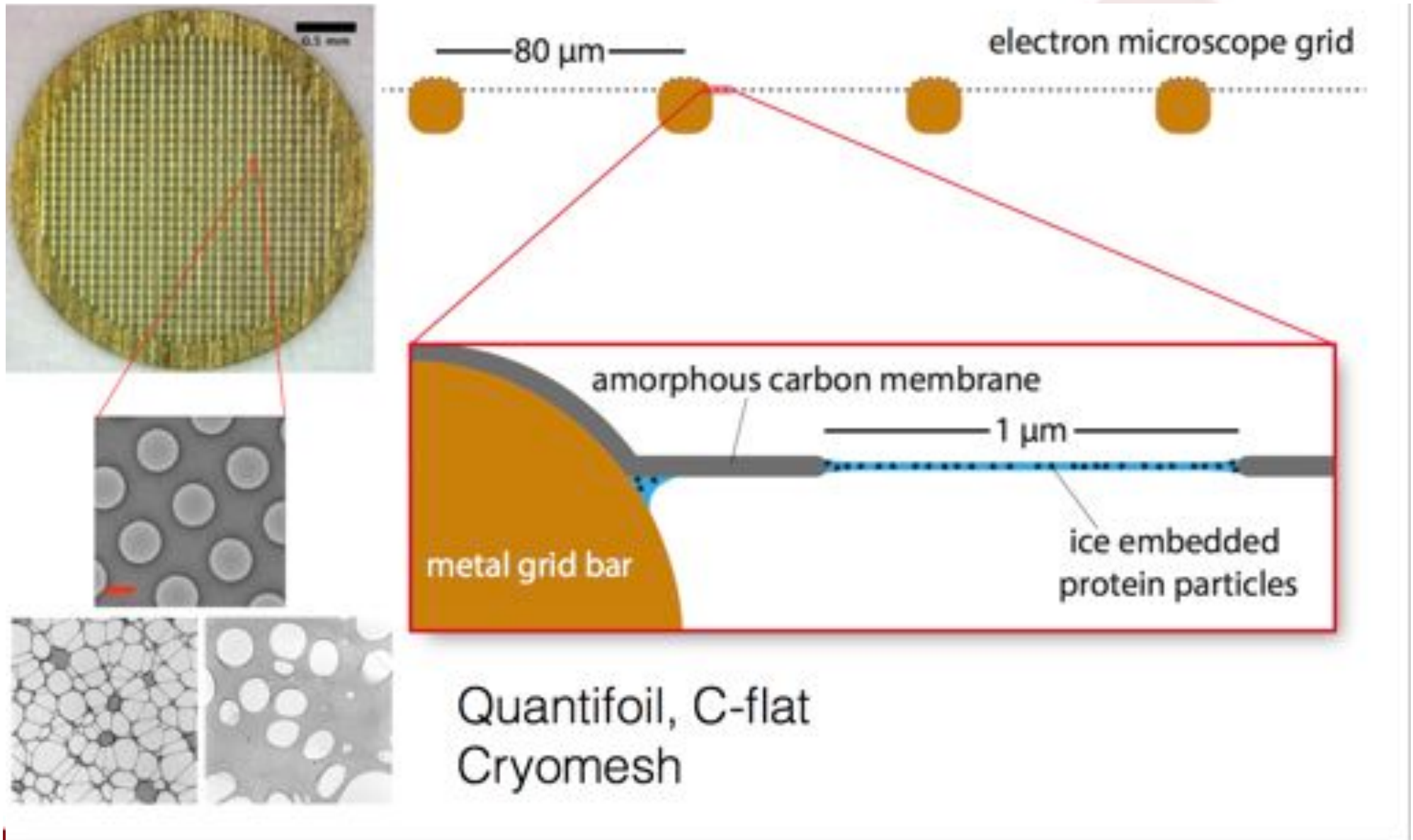
Sample

- concentration
- buffer composition
- detergent

Freezing

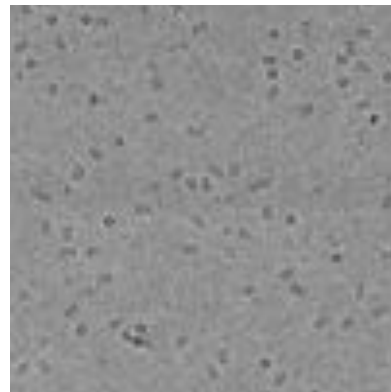
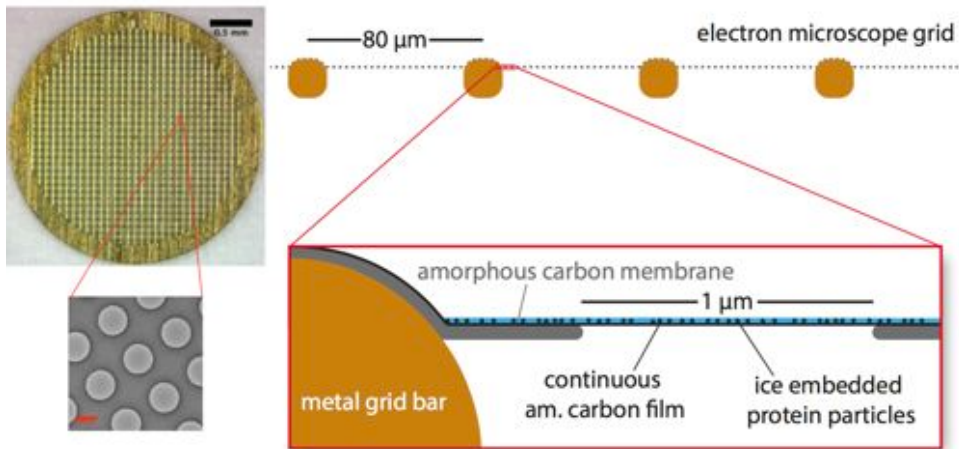
- blotting time
- single or double side blotting
- waiting time
- multiple sample application

The different substrats on grid

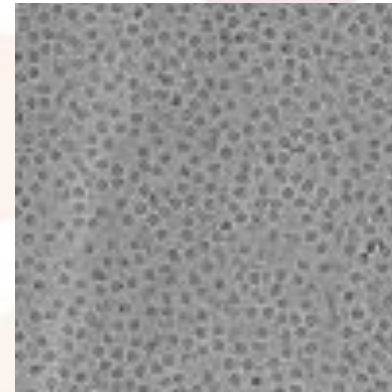
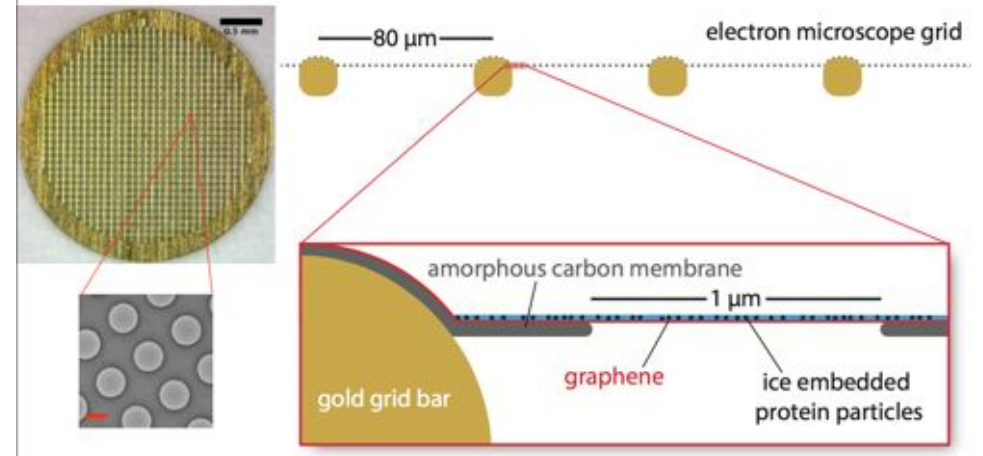


The different substrats on grid

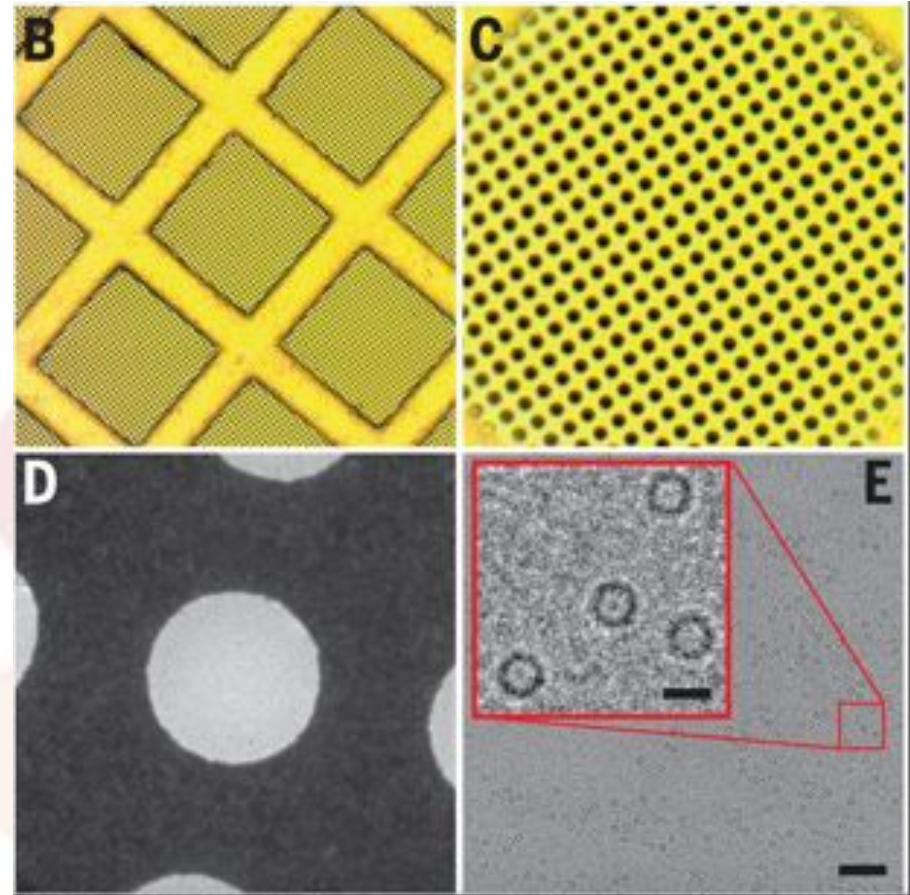
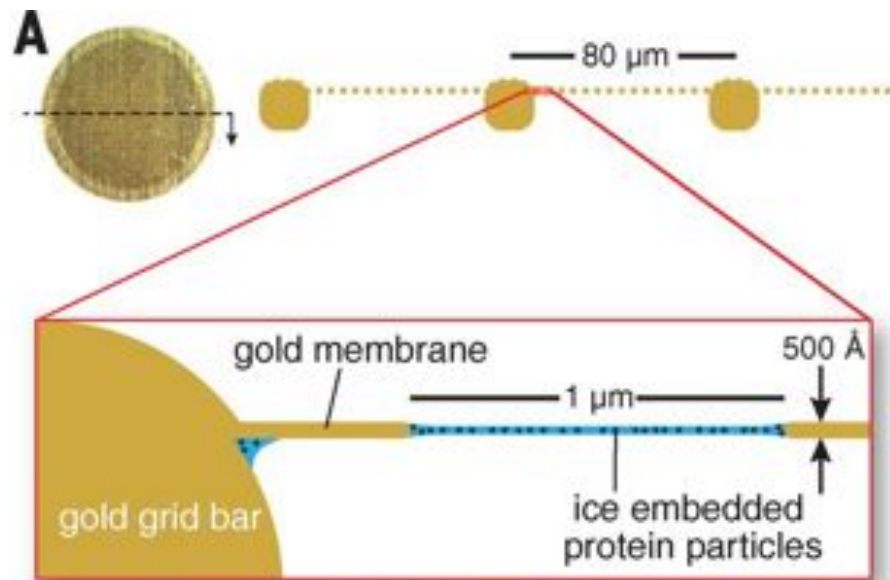
Traditional substrates for cryo-EM



Graphene substrates for cryo-EM



The different substrats on grid



Grid

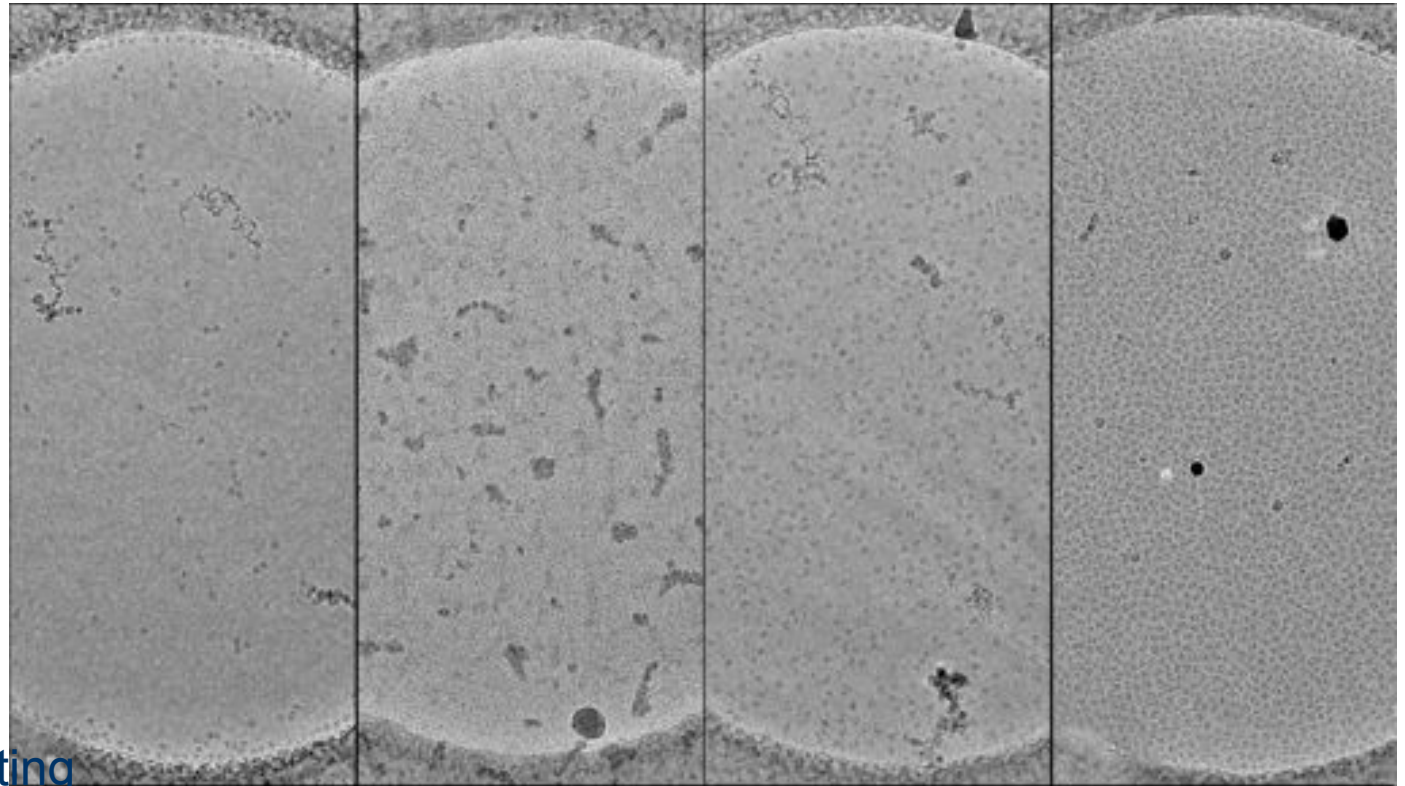
- type of grid and substrat
- batch and age of grid
- glow discharging

Sample

- concentration
- buffer composition
- detergent

Freezing

- blotting time
- single or double side blotting
- waiting time
- multiple sample application



Buffer composition

Buffer composition

Density of protein : 1.36 g/cm³

Density of water : 1 g /cm³

Density of glycerol : 1.26 g/cm³

Beware of high concentrations of

-glycerol

-sugars

-salts

-detergents

Grid

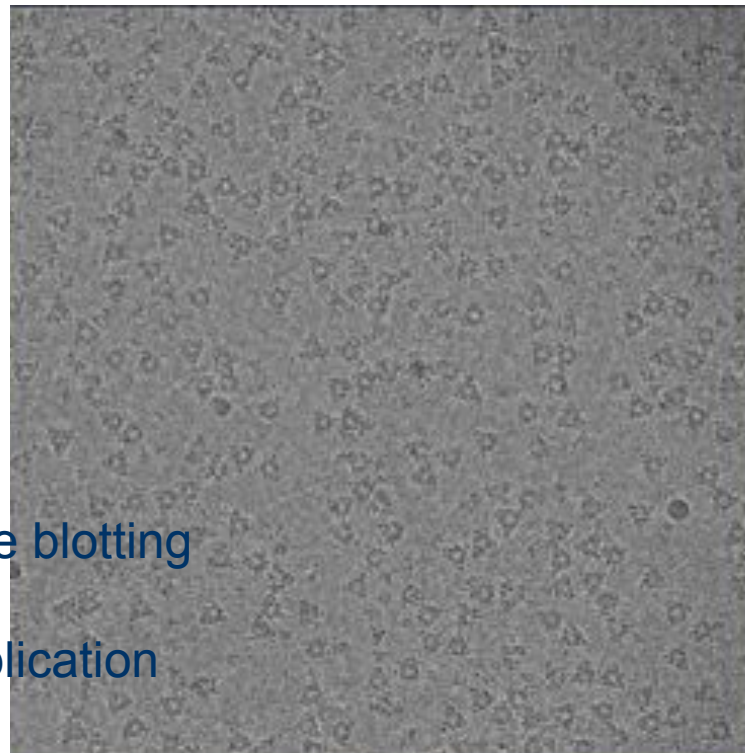
- type of grid and substrat
- batch and age of grid
- glow discharging

Sample

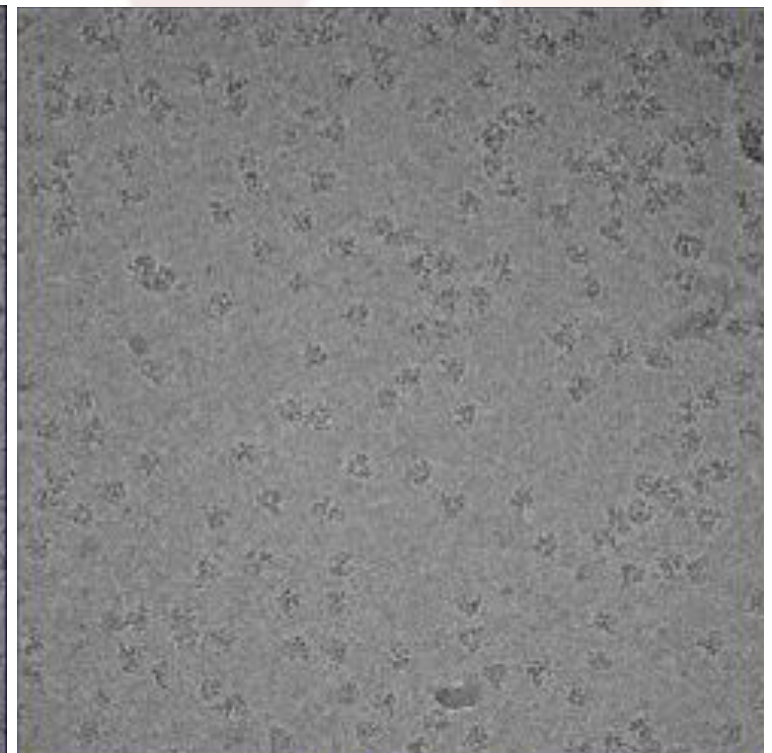
- concentration
- buffer composition
- detergent

Freezing

- blotting time
- single or double side blotting
- waiting time
- multiple sample application



No detergent



Detergent

Freezing device

Home made



€

Gatan



€€

FEI



€€€

Grid

- type of grid and substrate
- batch and age of grid
- glow discharging

Sample

- concentration
- buffer composition
- detergent

Freezing

- blotting time
- single or double side blotting
- waiting time
- multiple sample application

What is good for cryo EM?

Size

The bigger the better

200 kDa currently for near-atomic resolution

Symmetry

The higher the better

Pseudo symmetry can be a problem

Shape

Globular better than extended

« extra features » highly beneficial

Homogeneity

The more homogenous the better but heterogeneity now more manageable due to

- Better image quality
- New software development

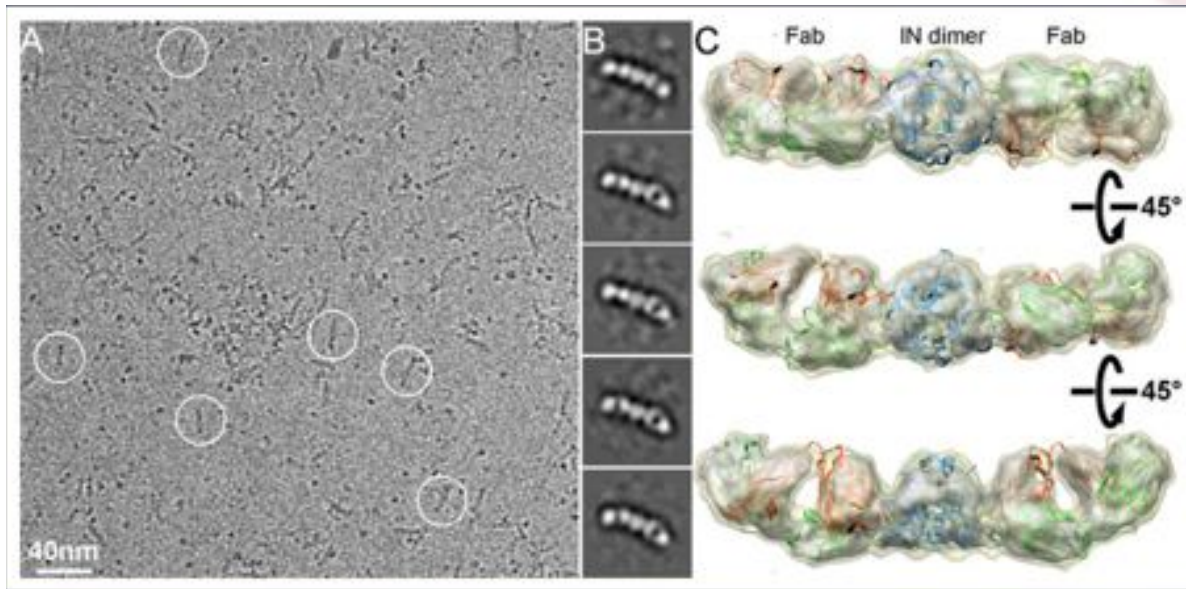
Size

The bigger the better

200 kDa currently for near-atomic resolution

Symmetry

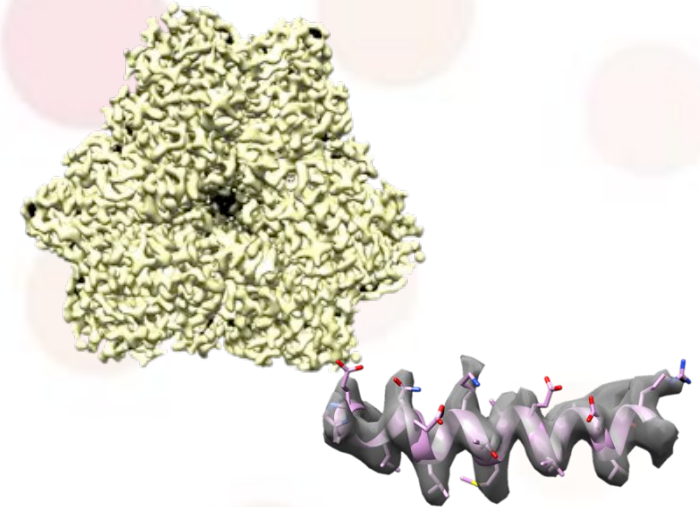
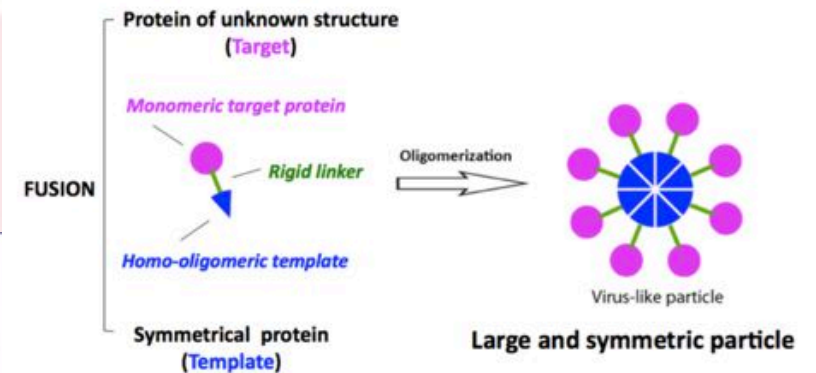
The higher the better



HIV-1 integrase dimer (65 kDa) in complex with two Fabs (total of 165 kDa).

10 Å resolution. Wu et al., 2012 Structure 20: 582-592.

What is good for cryo EM?



Tetrahedral
Amminopeptidase
480 kDa

What is good for cryo EM?

Shape

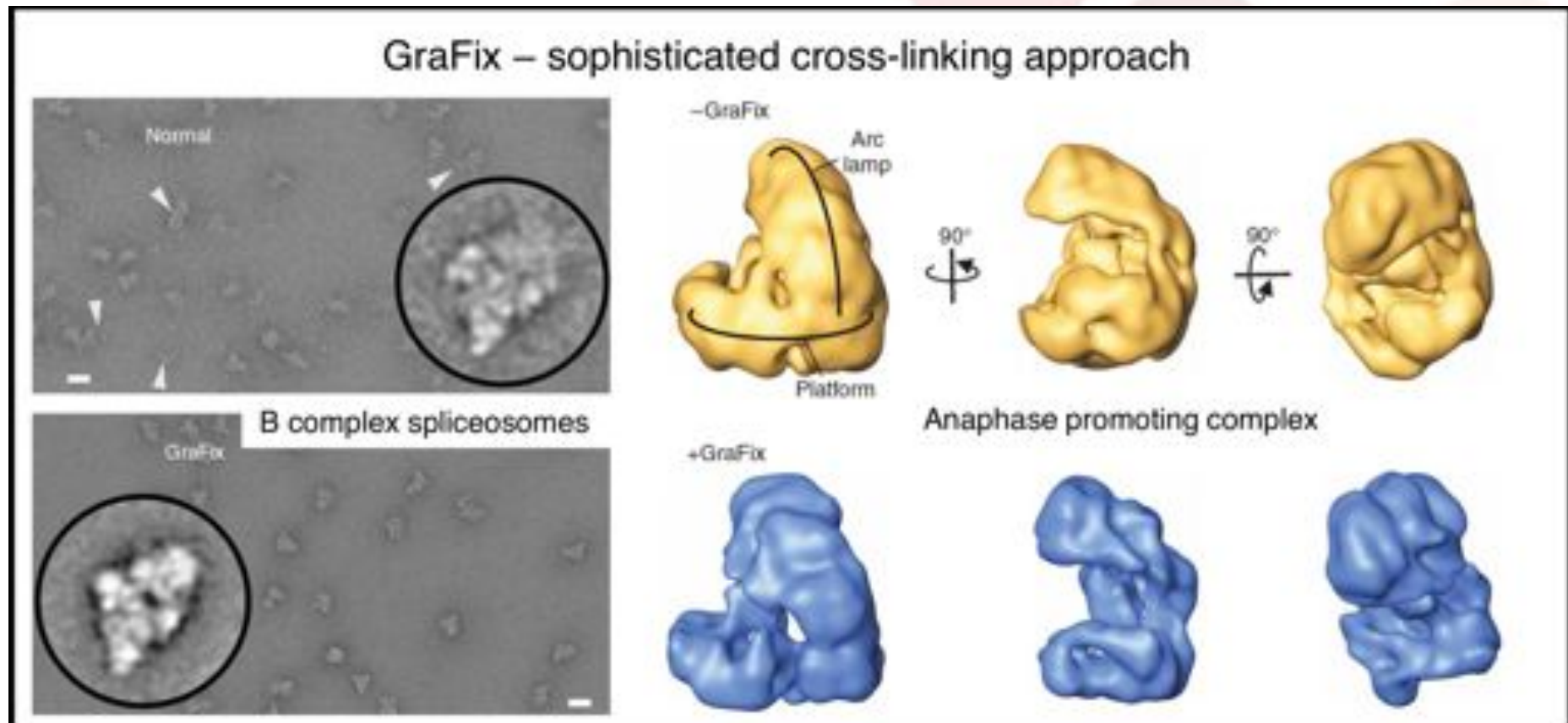
Globular better than extended

« extra features » highly beneficial

Homogeneity

The more homogenous the better but heterogeneity now more manageable due to

- Better image quality
- New software development



Kastner et al., (2008) Nat. Methods 5:53-55.

A lot of parameter can be changed

But start with the easier setting and if it doesn't work then try the more sophisticated one

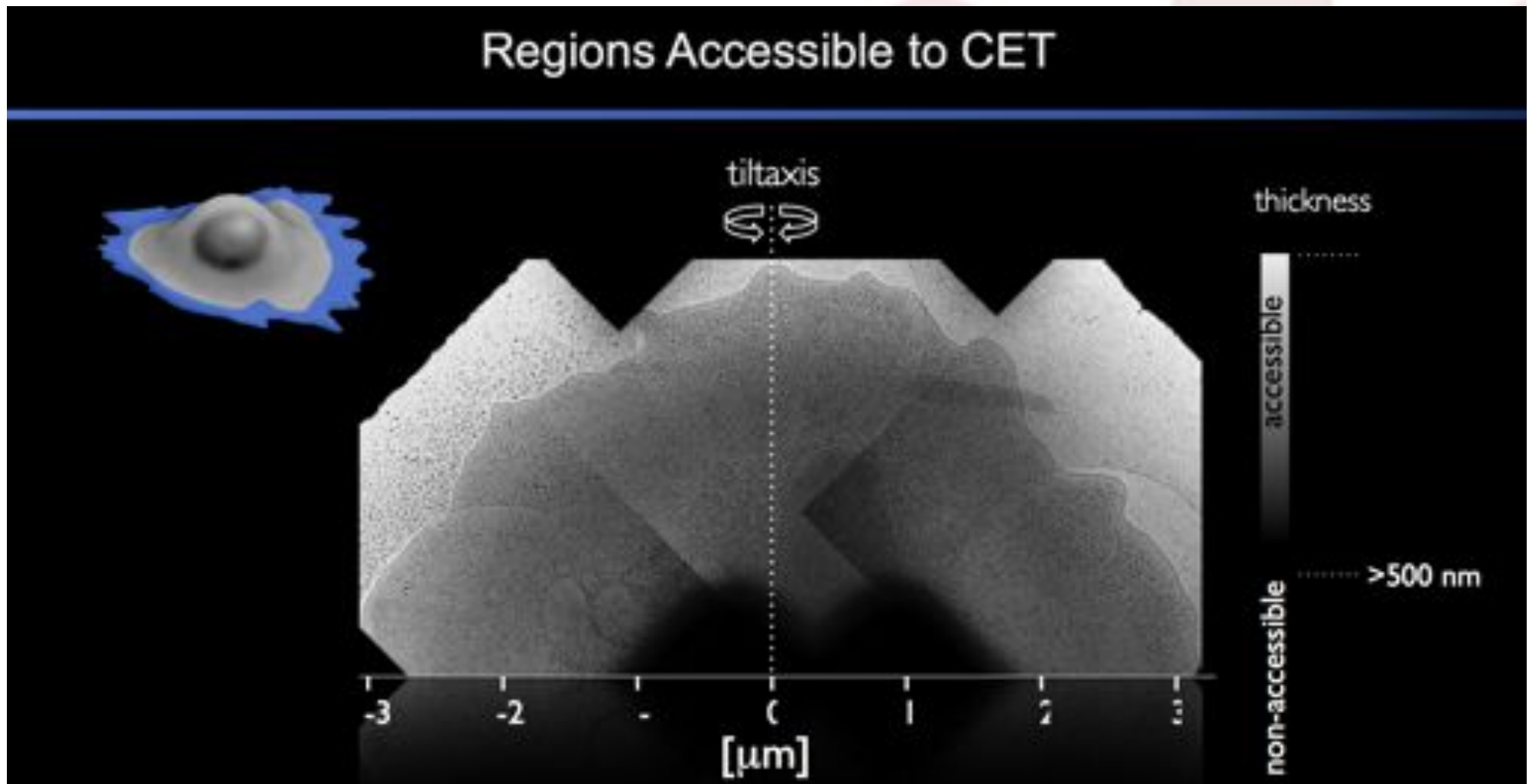
What can we freeze on the grid?



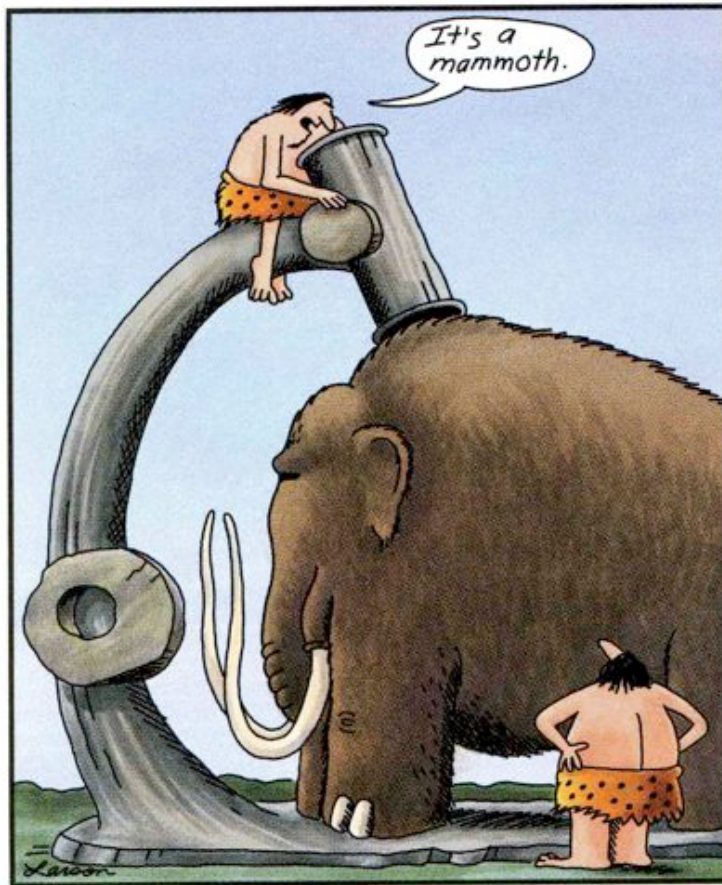
Bacterial Cells
~0.5 μm

Small Cells
~2 μm

Mammalian Cells
~5-10 μm



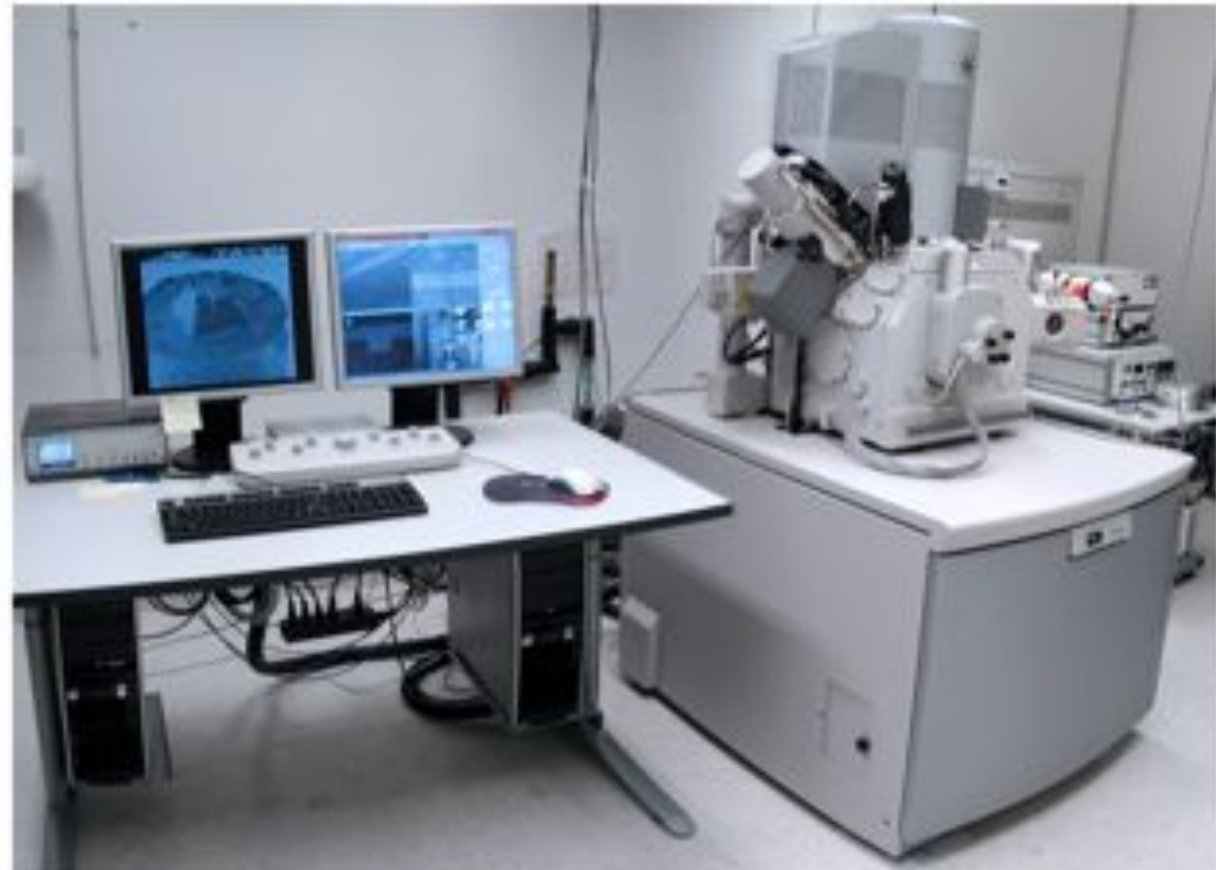
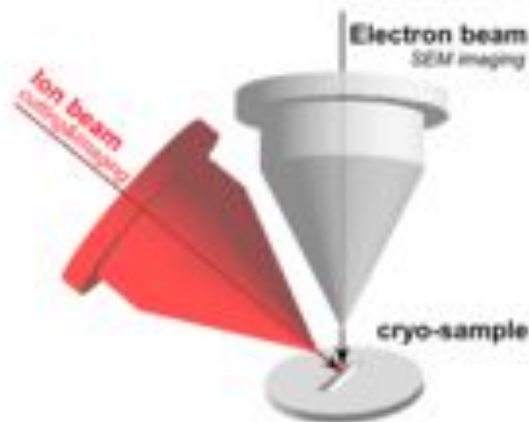
« Thick samples »



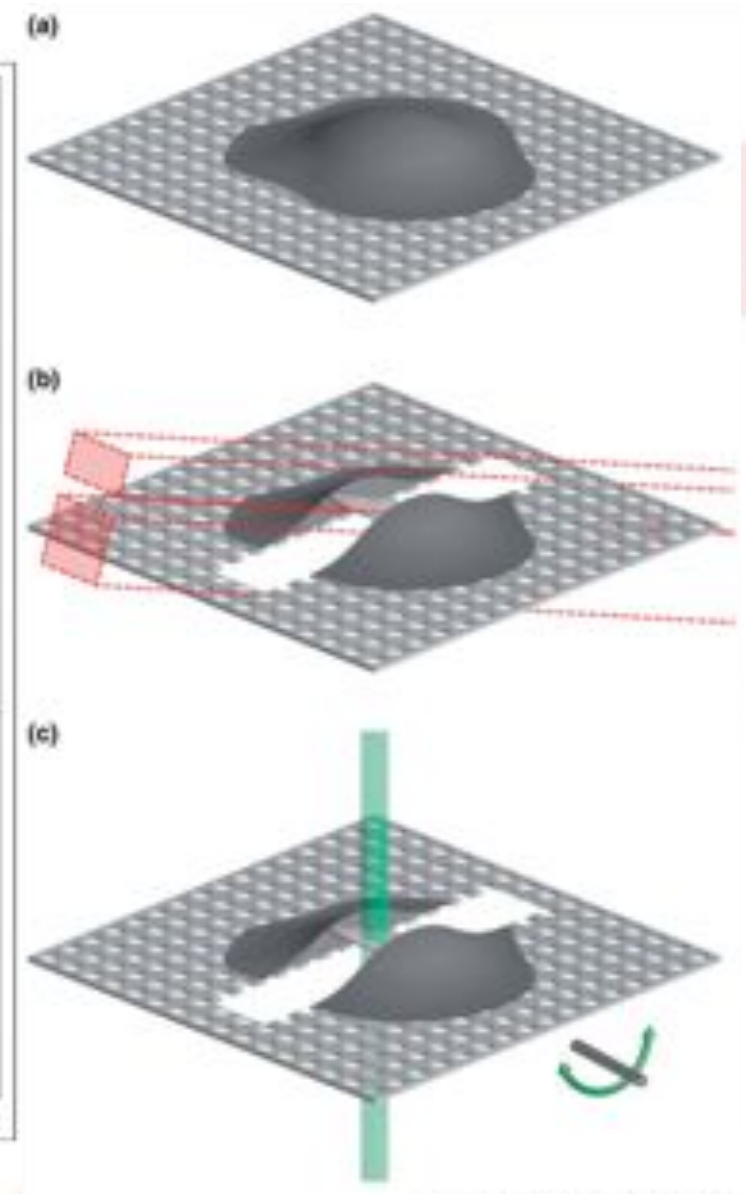
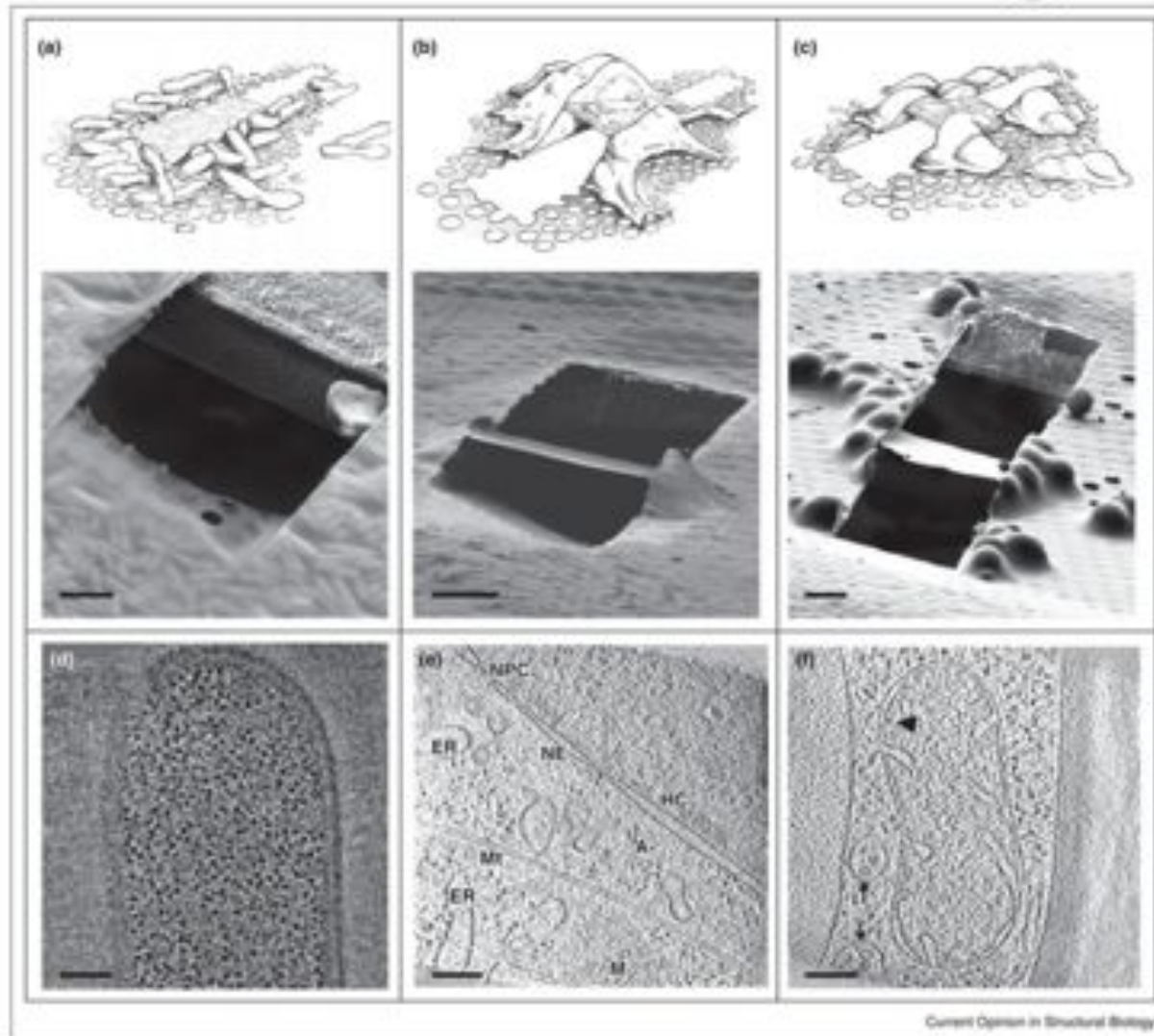
Early microscope

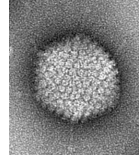
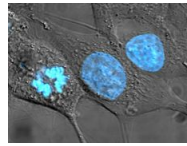


Focused Ion Beam – FIB

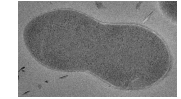


Focused ion beam milling





EM on Cellular sample



Cryo fixation

HPM100



Chemical fixation



- Fixation
(to stop all
cellular process)

- « Rigidification »

CEMOVIS



-FC7

Freeze> Tokuyasu
substitution



RT gelatine
embedding

RT resin
embedding

PLT



RT
sectionning



Cryosections
Immunogold
UC7-FC7



RT
sectionning



RT
sectionning



- Sectionning

-Visualization

Cryo-TEM

TEM

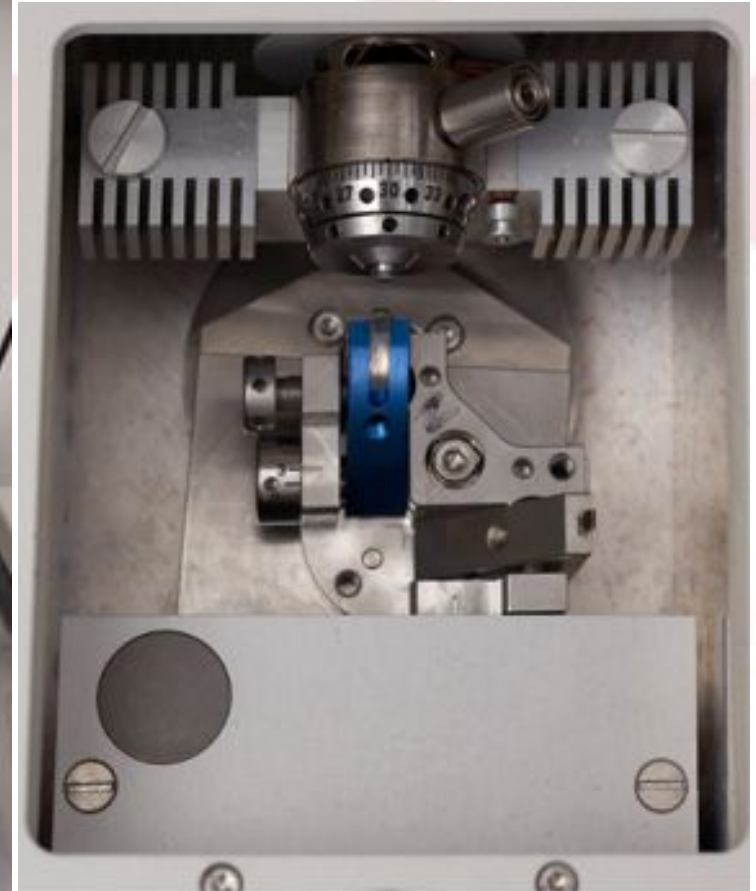
TEM

TEM

TEM

| | | | | | |
|-----------------|-------|-----|--------|-----|----|
| Ultrastructure | +++++ | +++ | + / ++ | ++ | ++ |
| Immunolabelling | - | ++ | +++ | +/- | ++ |
| Difficulty | +++ | ++ | ++(+) | + | + |

The ultra microtome



The high pressure freezing device



Cellular sample

Chemical fixation

RT resin embedding

- Dehydration
 - Impregnation
 - Polymerization
- 4 days

RT sectionning UC7

- Sectionning
- 1 day

TEM

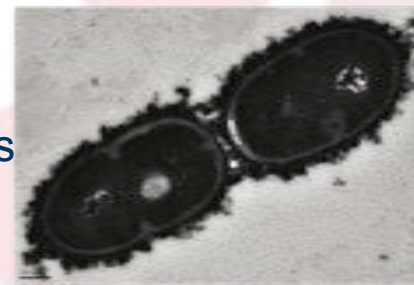
- Imaging
- 1(2) day(s)

| | |
|-----------------|-----|
| Ultrastructure | ++ |
| Immunolabelling | +/- |
| Difficulty | + |

Classical embedding

- No specific instrument needed
- Easy
- Good contrast

• D39_{WT}



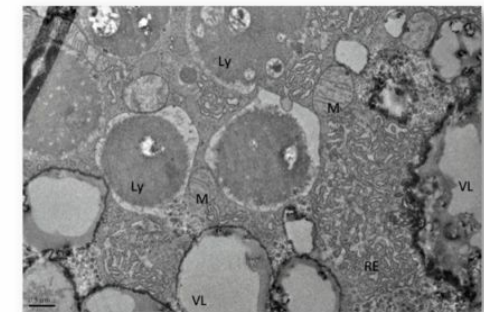
• D39_{ΔAil/ΔPhdD}



S. pneumoniae, RR + EPON, T



Mitochondria in
HeLa cell,



M : Mitochondrie / RE : Reticulum Endoplasmique / VL : Vesicule Lipidique
Ly : Lysosomes / AP : Autophagosomes

Drosophila larvae, fat body

Cellular sample

Chemical fixation

Tokuyasu
RT gelatine
embedding

Gelatine
Cryo-protection
Freezing

1 day

Cryosections
Immunogold
UC7-FC7 IGL

Sectionning

1 day

TEM

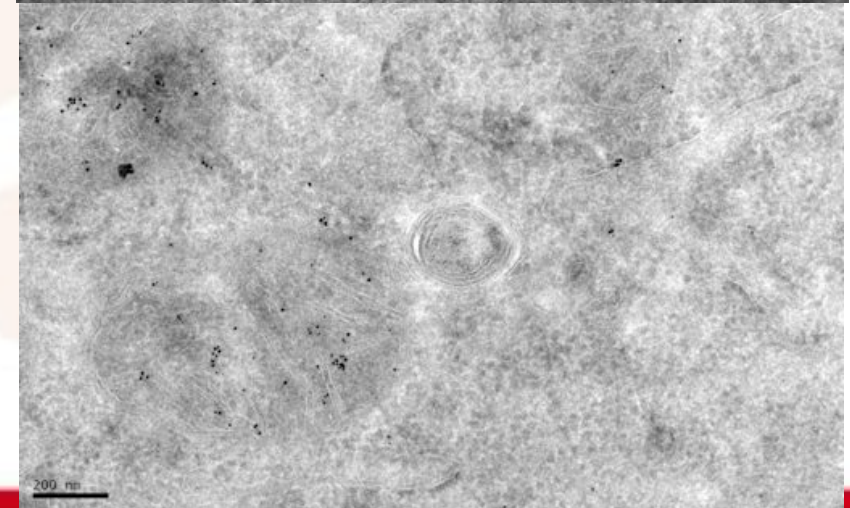
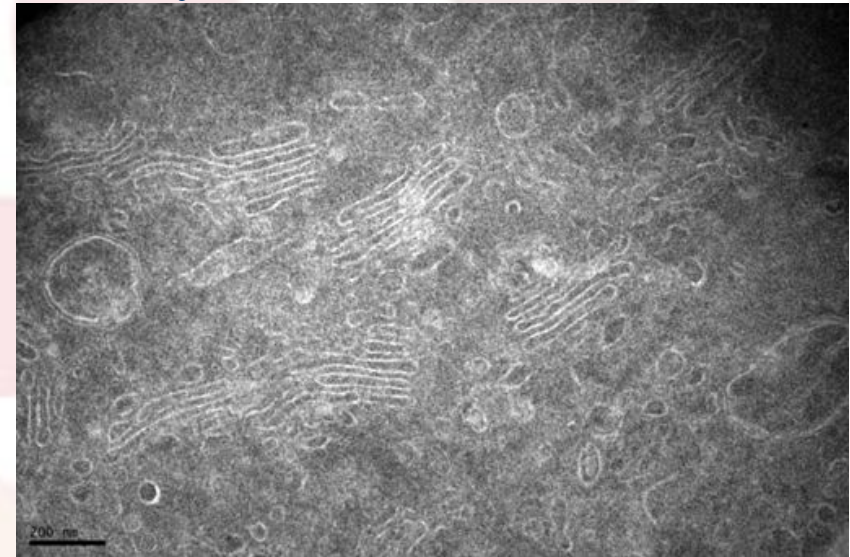
Imaging

1 (2) days

| | |
|-----------------|--------|
| Ultrastructure | + / ++ |
| Immunolabelling | +++ |
| Difficulty | ++(+) |

Tokuyasu for immunogold

- Fast
- Good structure preservation
- Best for immunogold
- Cryosections



Cellular sample

Cryo fixation



Freeze substitution

- Substitution
 - Impregnation
 - Polymerization
- 5-10 days

RT sectionning

- Sectionning
- 1 day

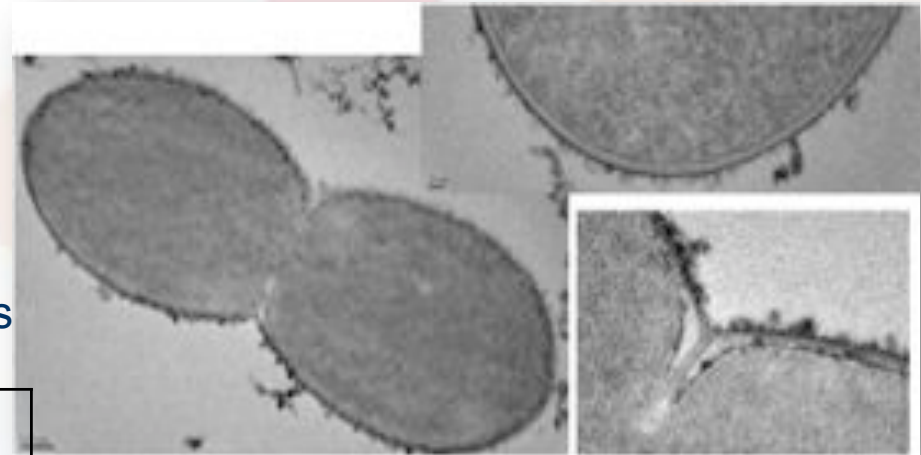
TEM

- Imaging
- 1(2) days

| | |
|-----------------|-----|
| Ultrastructure | +++ |
| Immunolabelling | ++ |
| Difficulty | ++ |

HPF & freeze substitution

- HPF (HPM100) & FS (AFS2)
- Automated process
- RT sections
- Immuno-compatible (HM20)
- Good structure preservation



S. pneumoniae, R6

Cellular sample

Cryo fixation

HPM100

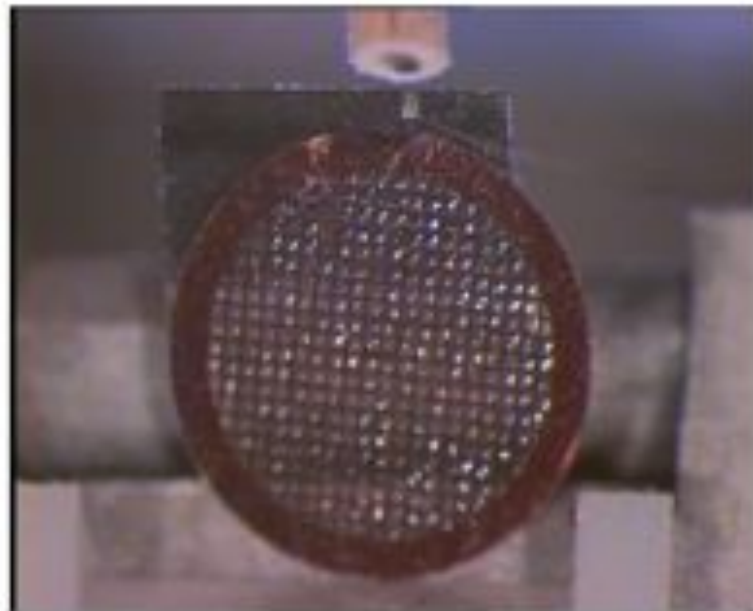


HPF device

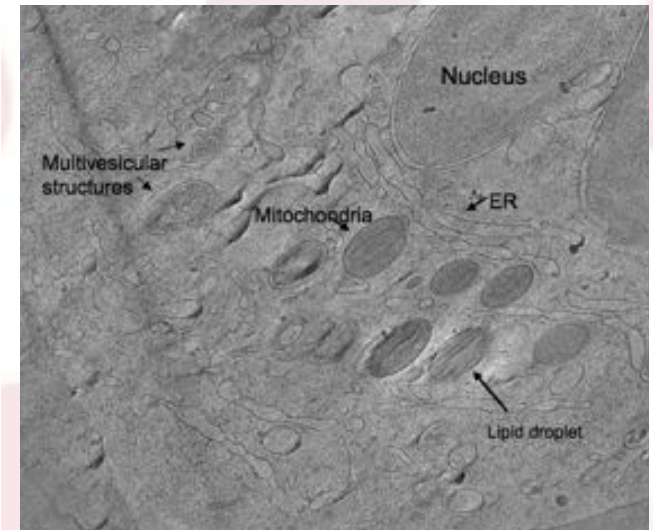
CEMOVIS

UC7-FC7

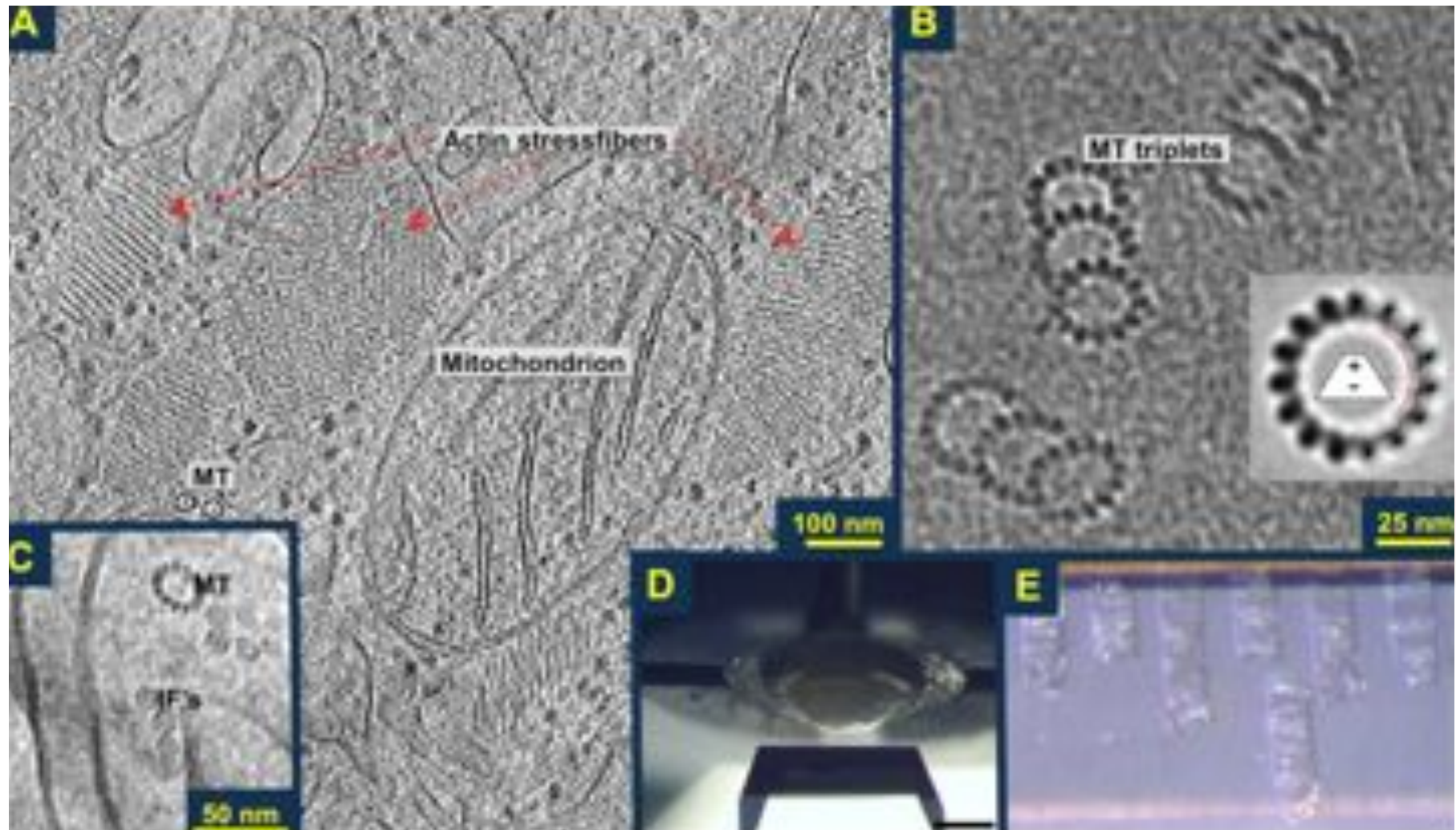
Cryo-TEM



picture by Peter Peters



Example of Cemovis

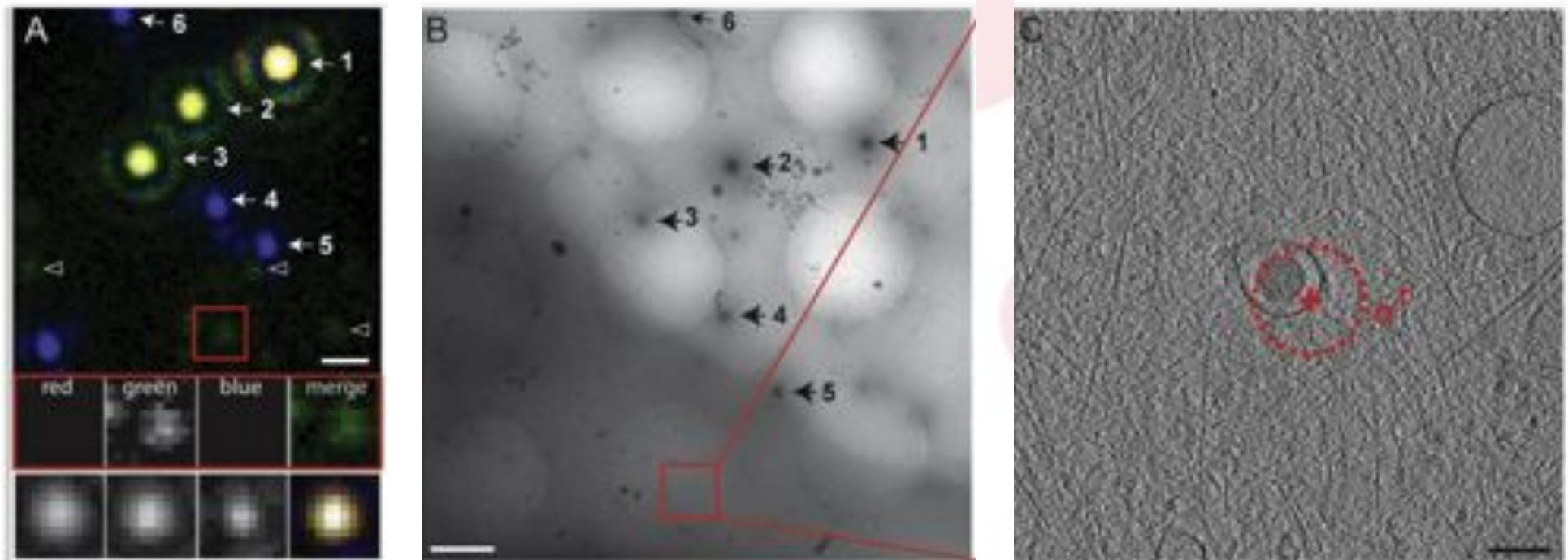


Very difficult

Correlative light electron microscopy

Locate an event by light microscopy and then find the same event in the EM to image it at higher mag

Use of probe that are electron dense and fluorescent



The sample preparation step is crucial (as well as the purification and data collection steps)

Some other techniques exist for sample preparation:

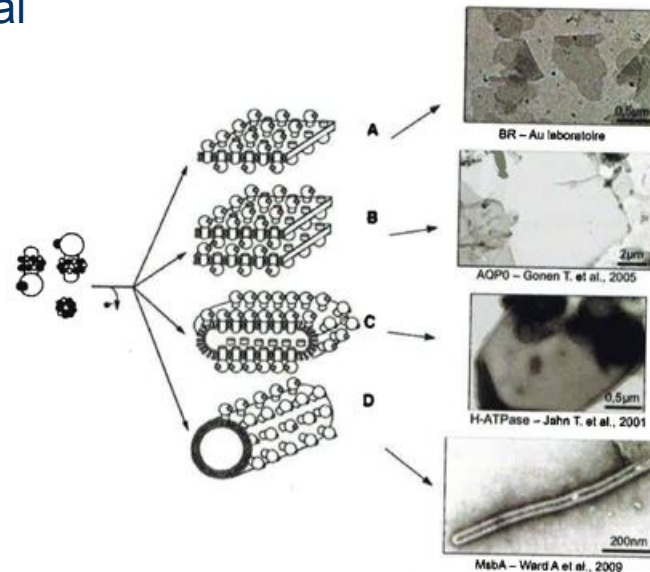
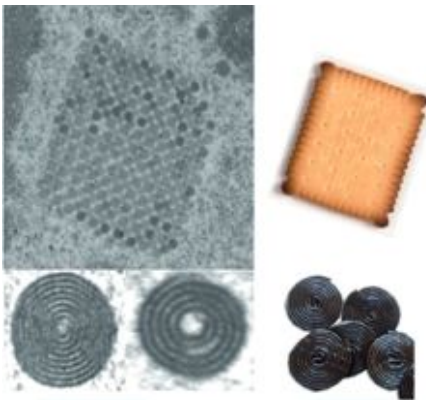
Time resolved EM

Cryo-negative staining

Specificity for tomography

For 2D crystal

Etc...



Questions????

