

# CRYOEM 001 : EM SAMPLE PREPARATION

NCCAT Embedded Training — Master Class series

September 30, 2020

NATIONAL CENTER FOR  
CRYOEM ACCESS & TRAINING



New York Structural  
Biology Center

SIMONS ELECTRON  
MICROSCOPY CENTER





# CRYOEM 001 : SINGLE PARTICLE MASTERCLASS

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Introduction to cryoEM: SPA

Building a cryoEM toolkit

EM compatible samples

EM support films and grids

Sample preparation

Tools of the trade:

microscopes and detectors

Microscope operations

Data collection strategies

Data assessment & QC

Data processing:

cryoEM IT infrastructure

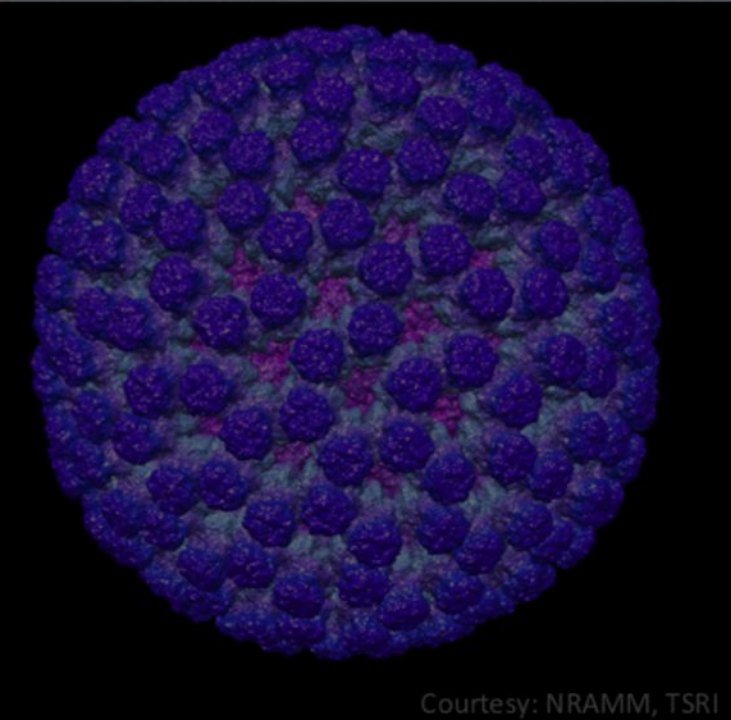
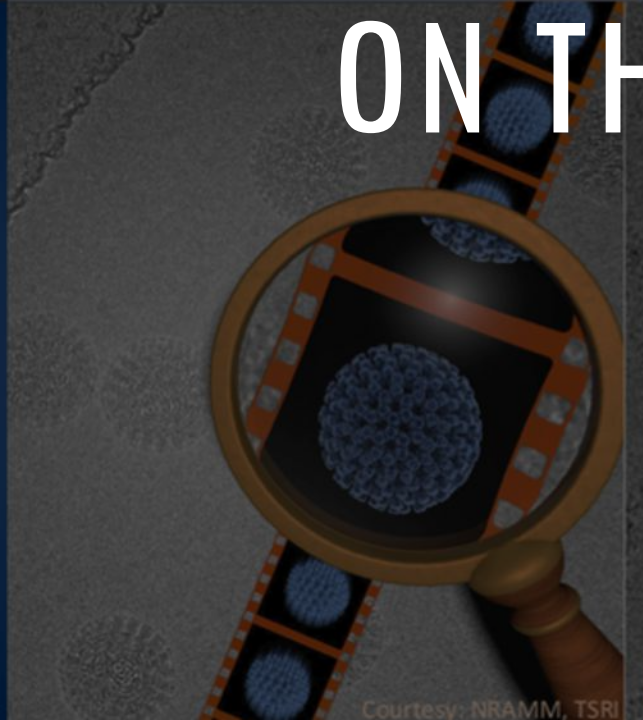
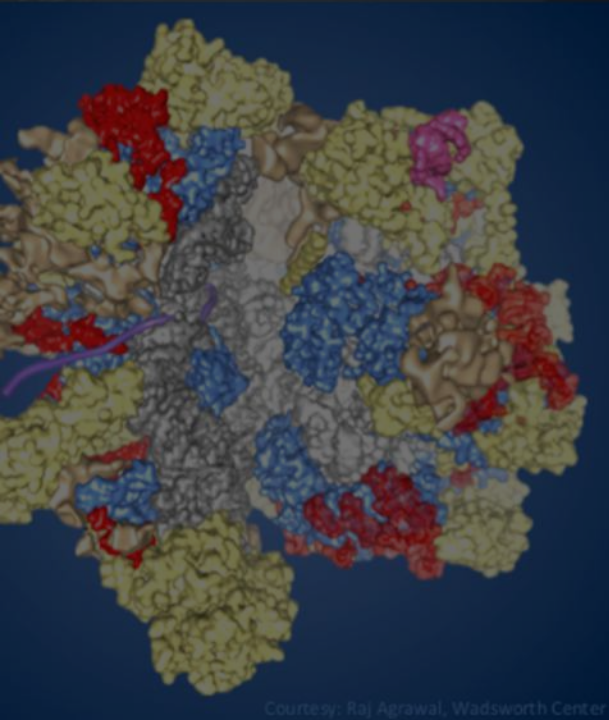
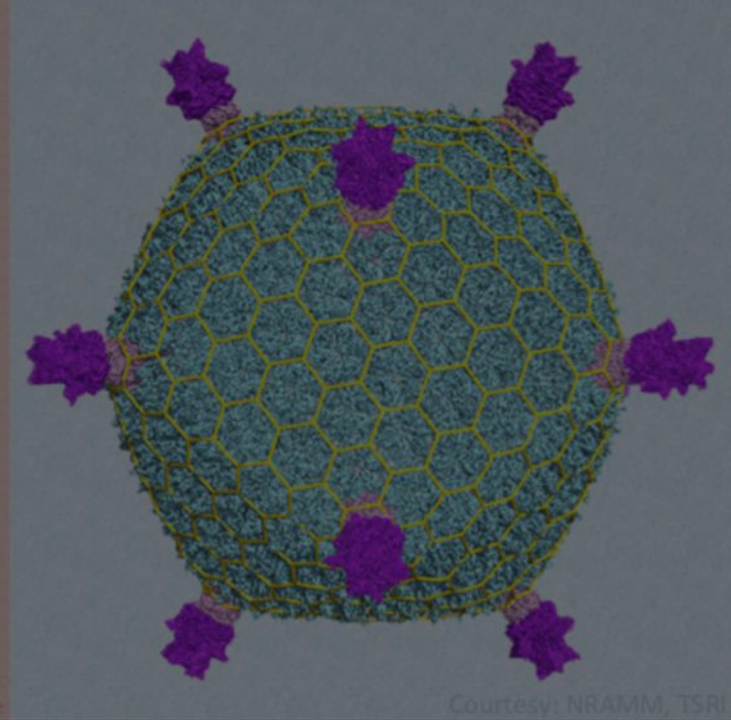
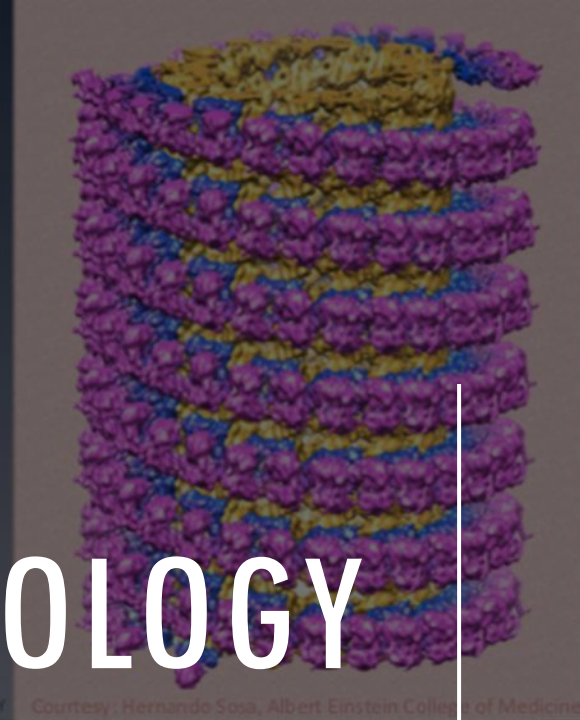
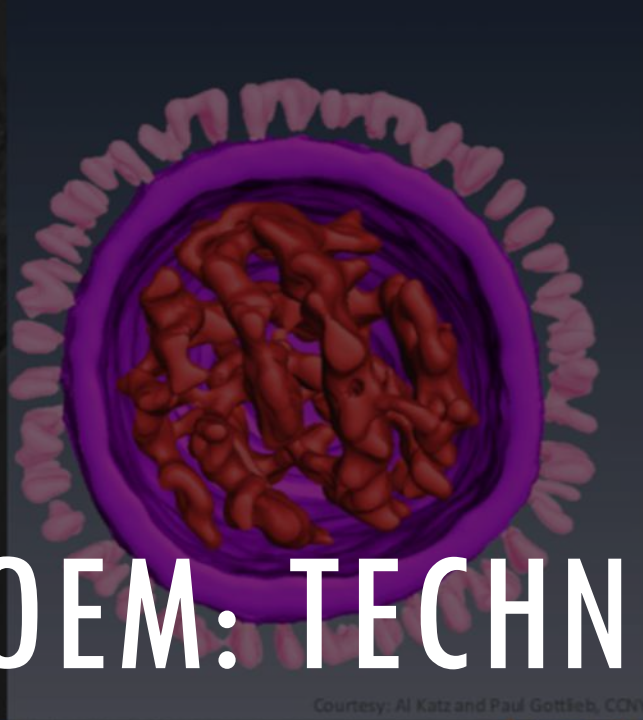
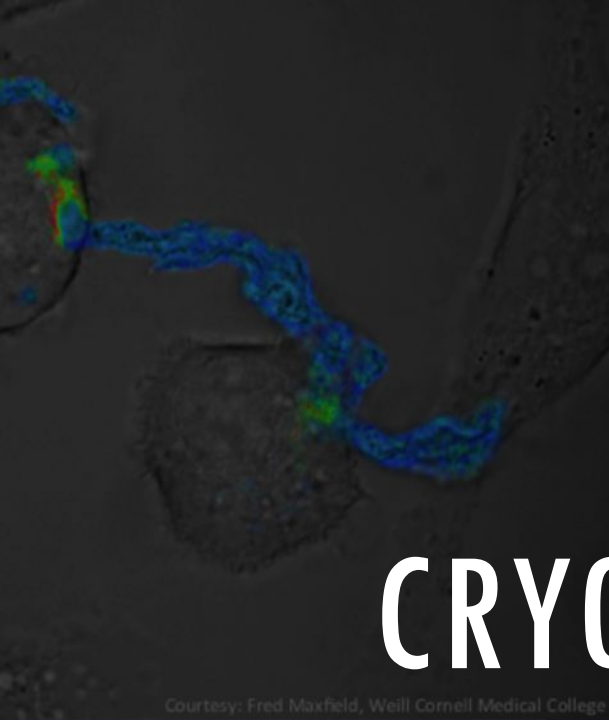
On-the-fly feedback

3D Reconstruction

Visualization and validation



# CRYOEM: TECHNOLOGY ON THE RISE





# HOW ARE SAMPLES PREPARED?

## Sample

proteins/  
macromolecular  
complexes

Biochemically  
homogeneous

Biochemically  
heterogeneous

cells/tissues/  
organisms

Structurally  
homogeneous

Structurally  
heterogeneous

## Sample preparation

2D/3D  
crystallization

Helical  
assembly

Single particle  
isolation

Serial sectioning

Cryo embedding

## EM technique

Electron  
crystallography

Helical  
reconstruction

Single particle  
analysis

Electron  
tomography

FIB-SEM/cryoET

## Resolution range

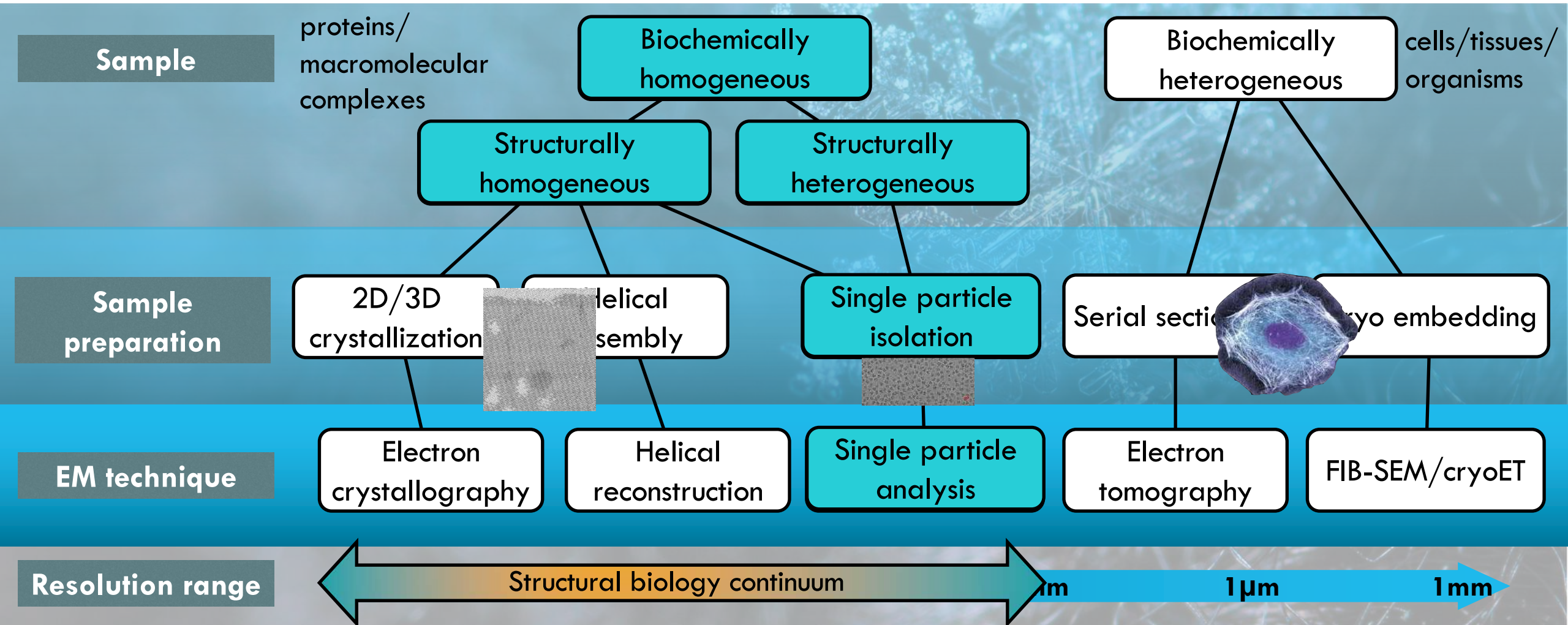
1Å

1nm

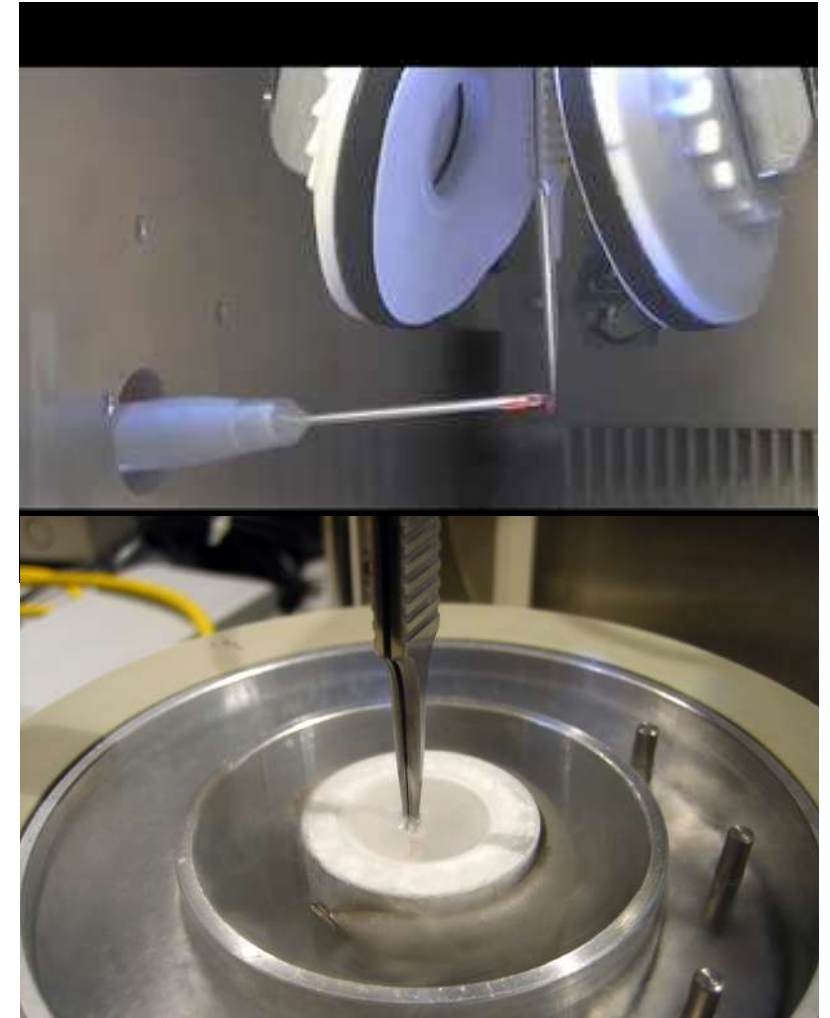
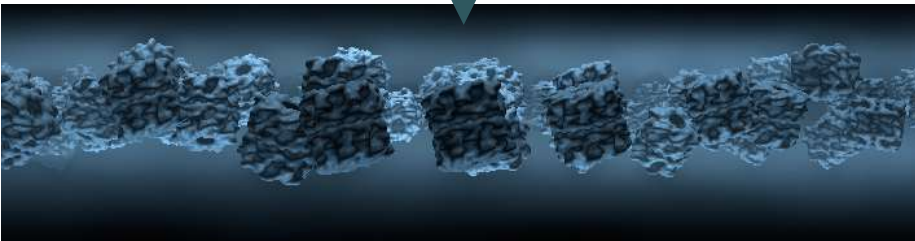
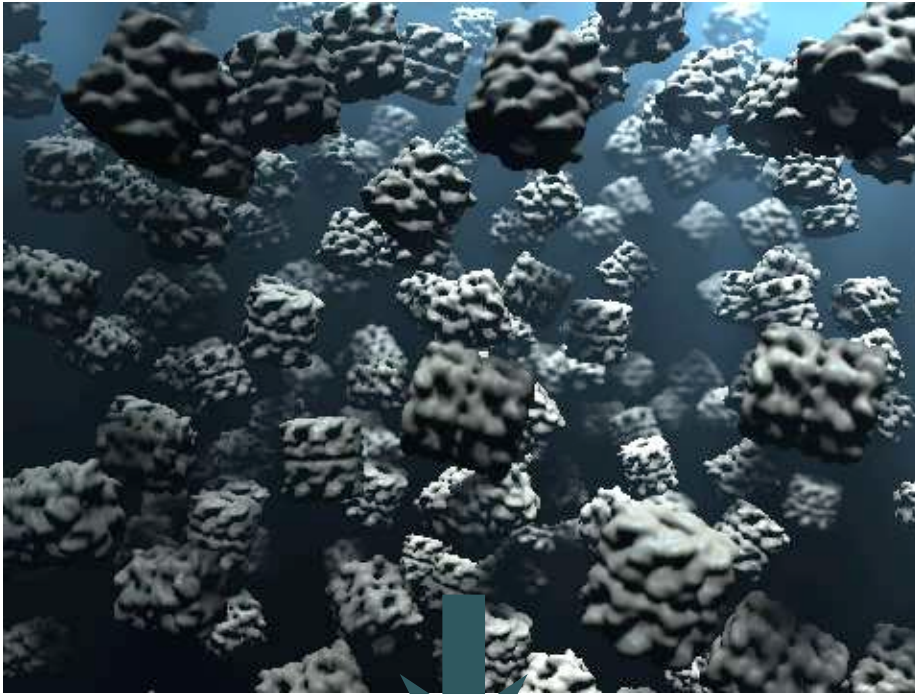
1µm

1mm

# HOW ARE SAMPLES PREPARED?



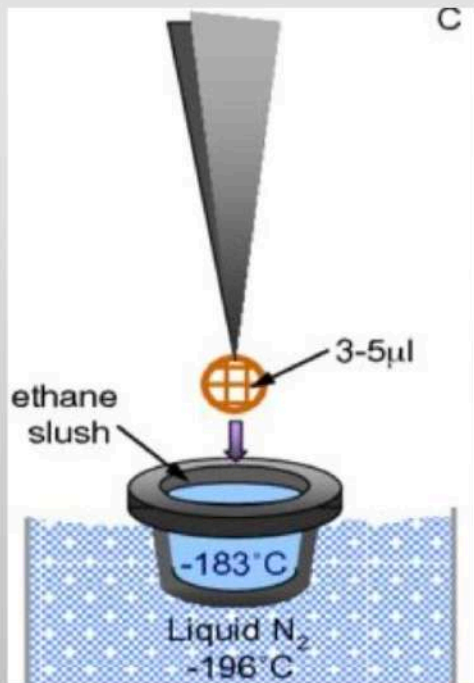
# PLUNGE FREEZING



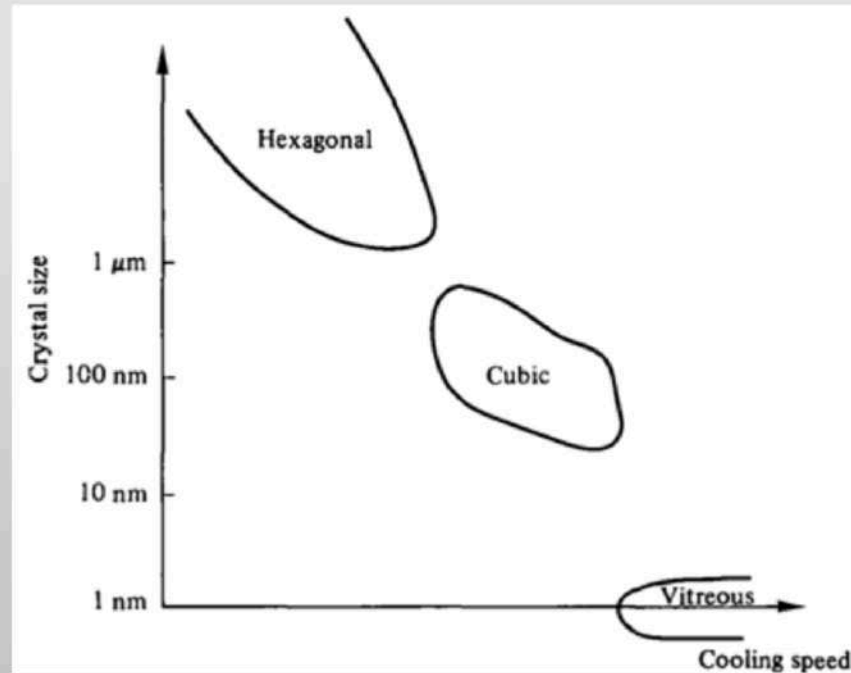


# PLUNGE FREEZING

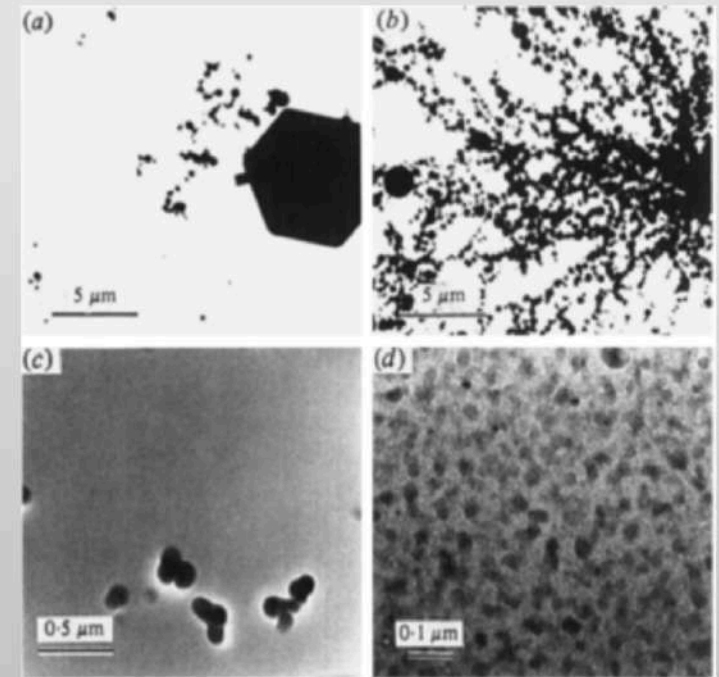
- Liquid ethane is a suitable coolant.
- Liquid nitrogen boils on contact, which makes it a poor coolant for cryo-EM.
- Cooling speed faster than  $10^5$ - $10^6$  K/s ensure the formation of vitrified ice.



Setup of liquid ethane  
(Image from Wen Jiang)



Cooling speed &  
forms of ice



Different forms of ice contamination

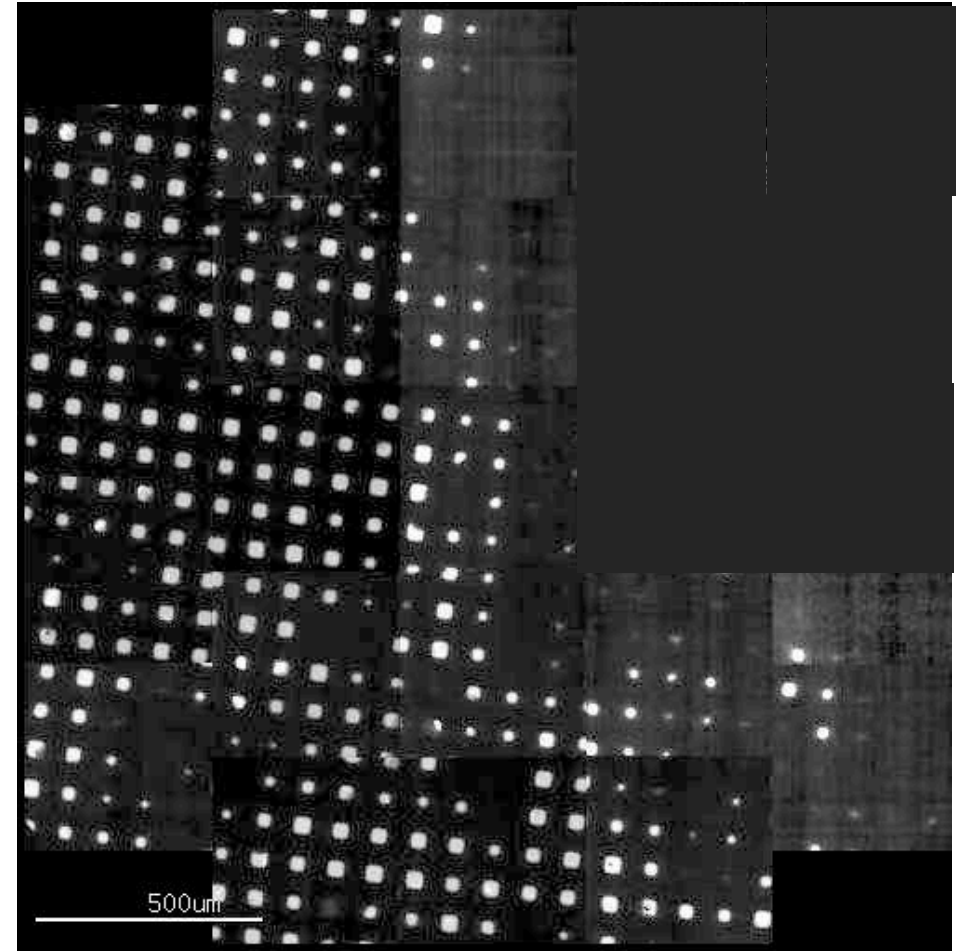
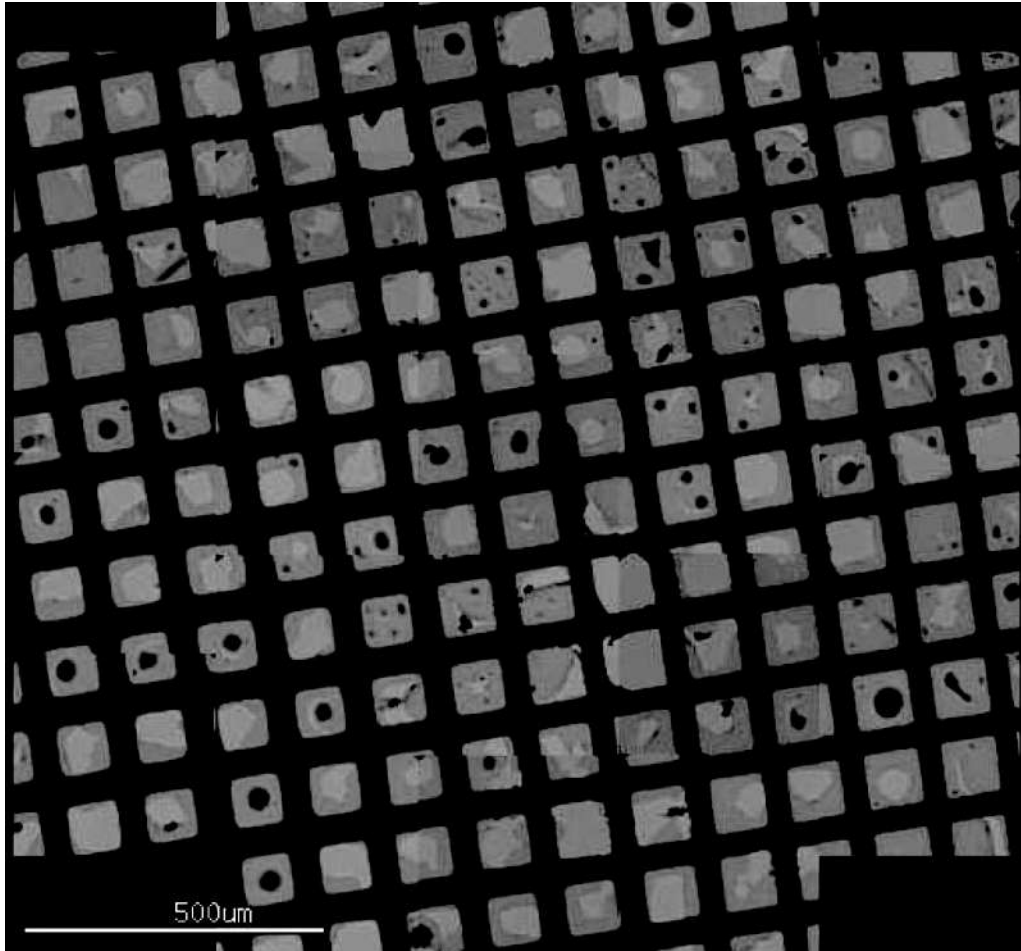


# PLUNGE FREEZING



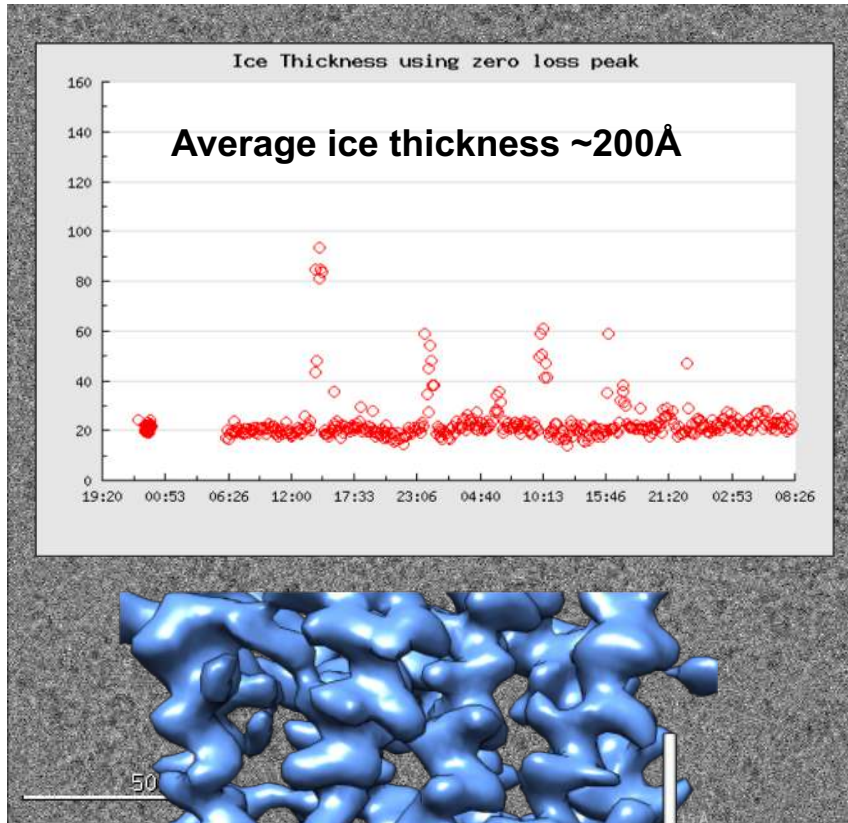


# WHAT DO GRIDS LOOK LIKE?

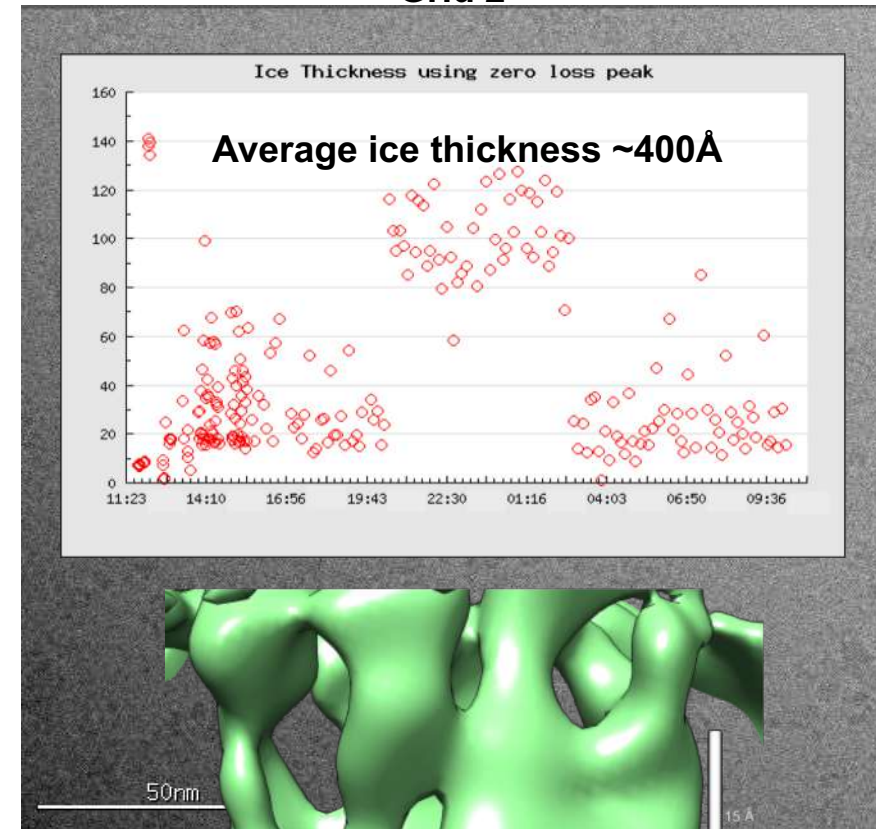


# GRID PREPARATION IS A CHALLENGE

Grid 1



Grid 2

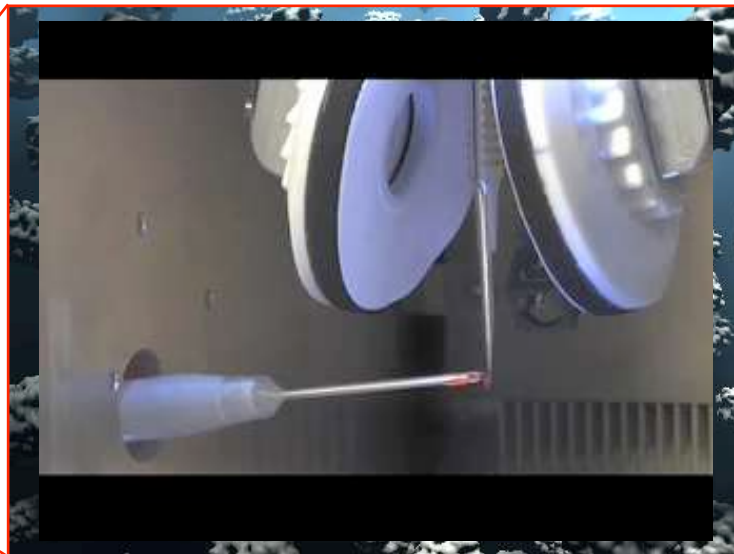


Yong Zi Tan

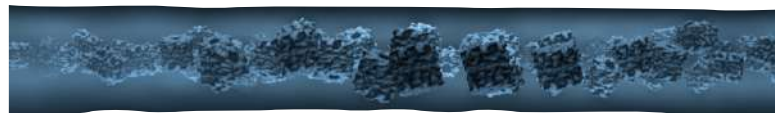
45 kDa integral membrane  
protein + Fab



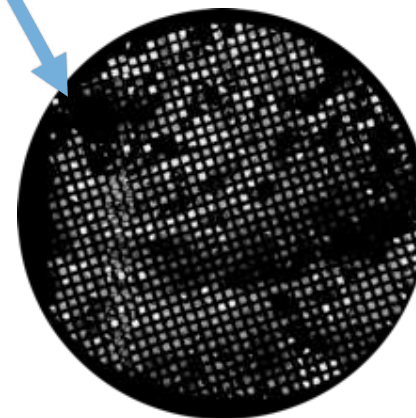
# GRID PREPARATION IS A CHALLENGE



~3 $\mu$ l



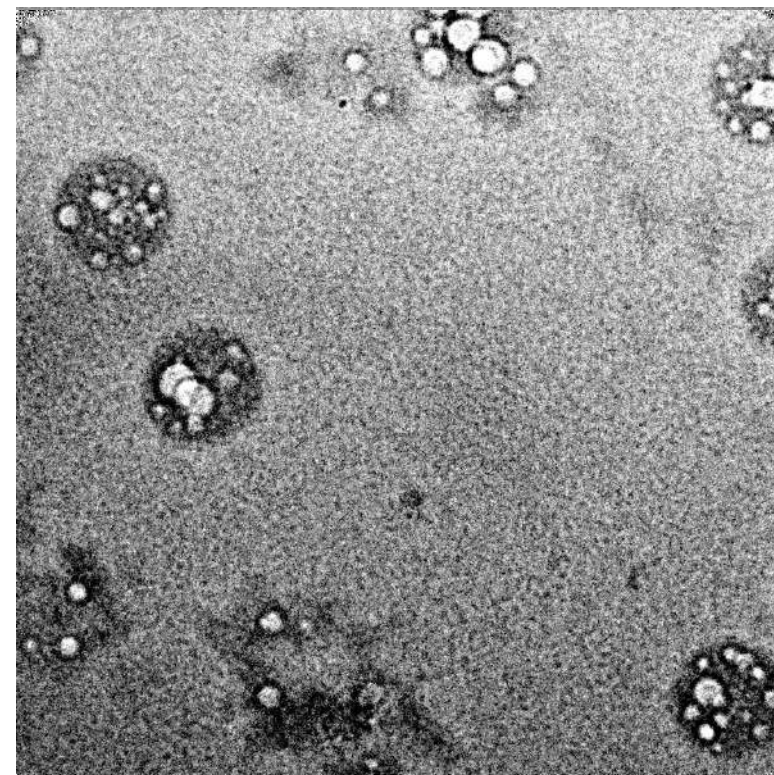
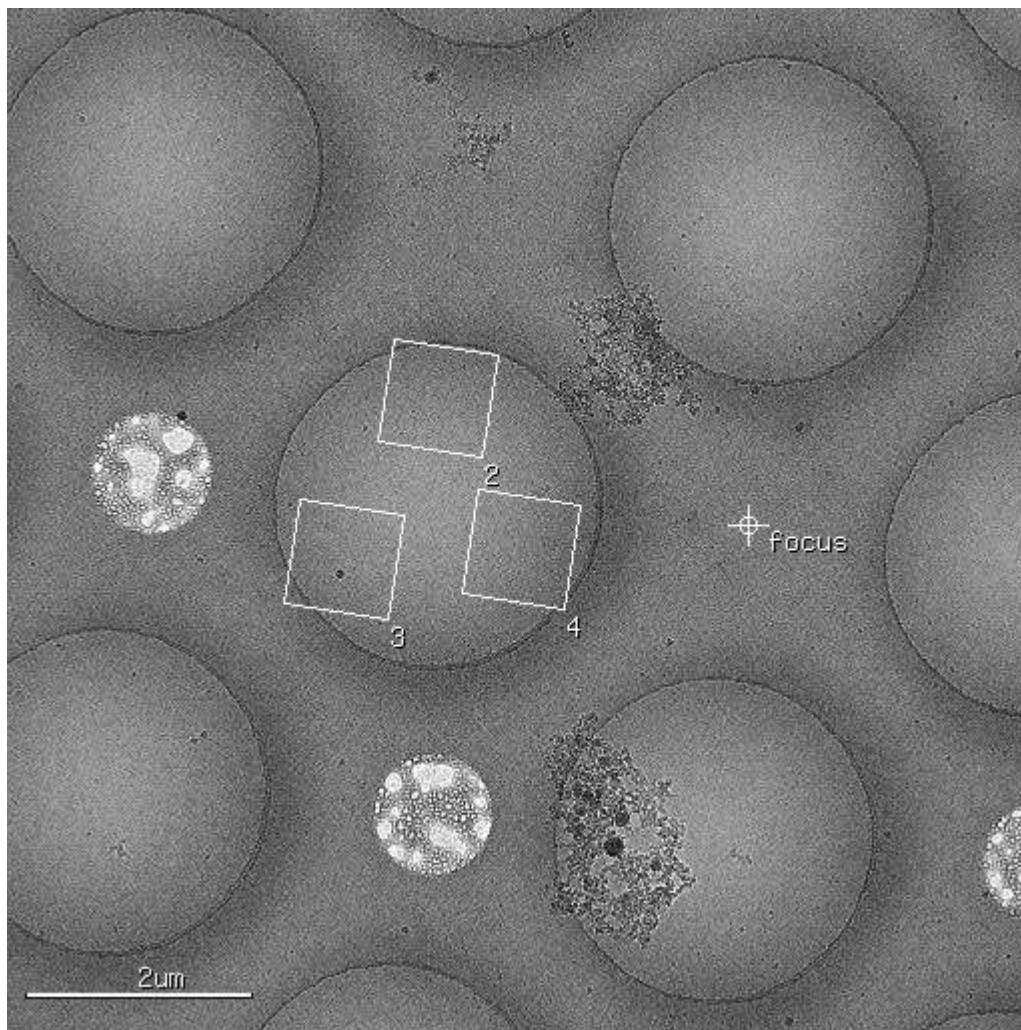
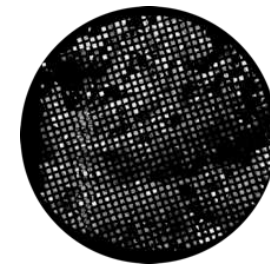
<3nl



>100,000 potential imaging targets; most of them are not usable.

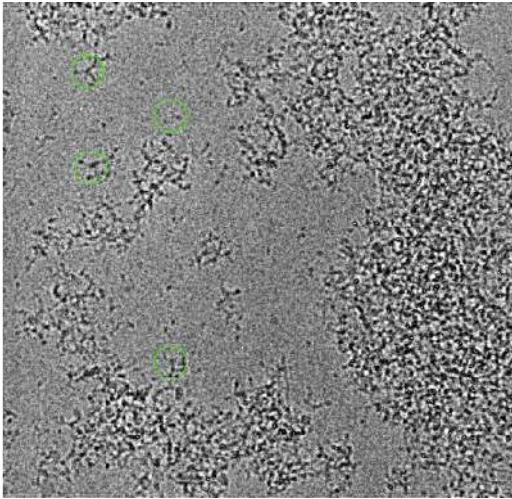
Graphics courtesy Gabe Lander

# LOW DOSE IMAGING

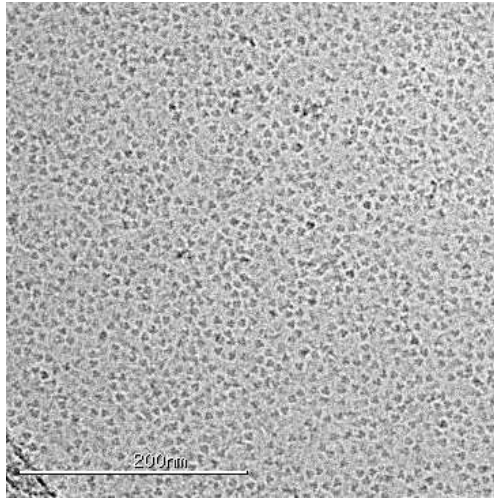




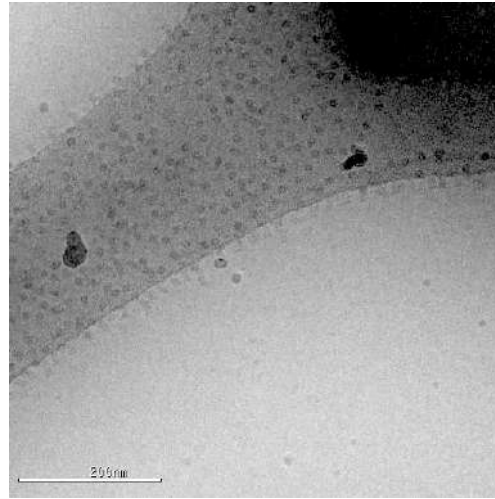
# WHAT ISSUES ARISE?



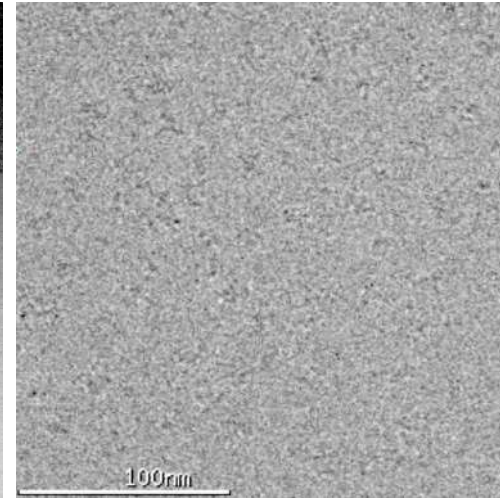
Aggregating in ice



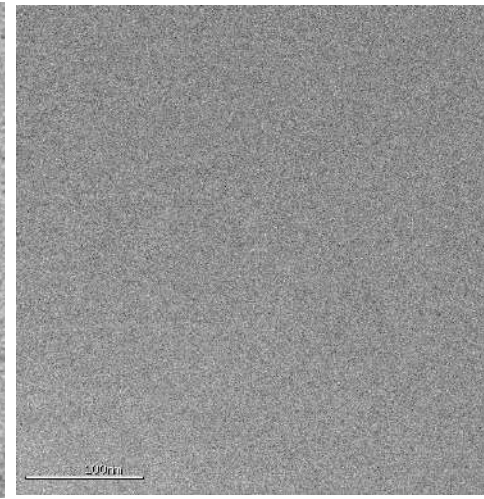
Preferred orientation



Particles not going into holes



Rejecting 90% of particles



Particles disappearing in ice

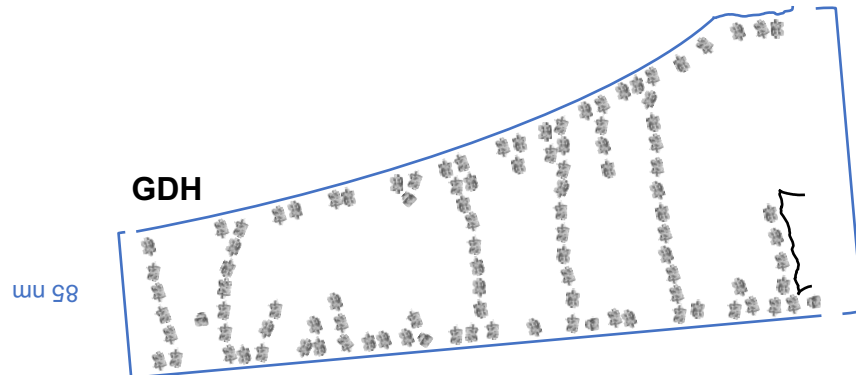
# WHAT ISSUES ARISE?



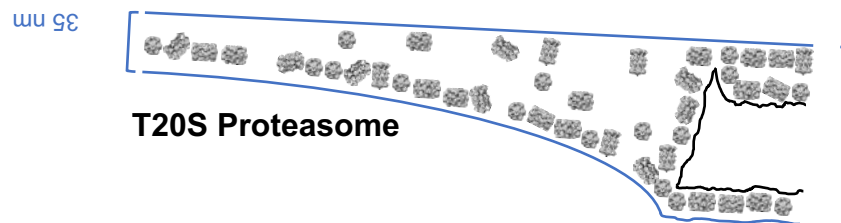
Hemagglutinin



Hemagglutinin



GDH



T20S Proteasome

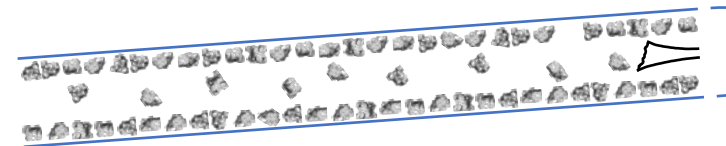
110 nm  
ice



Aldolase

45 nm  
ice

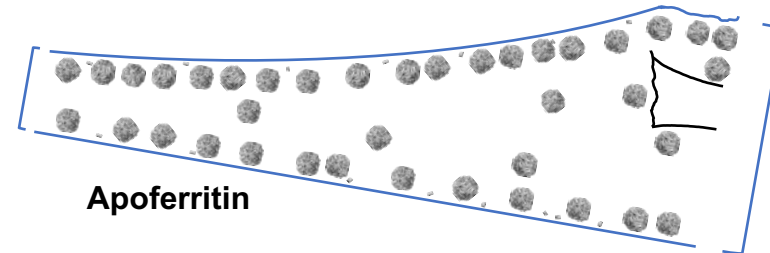
125 nm  
ice



Aldolase

50 nm  
ice

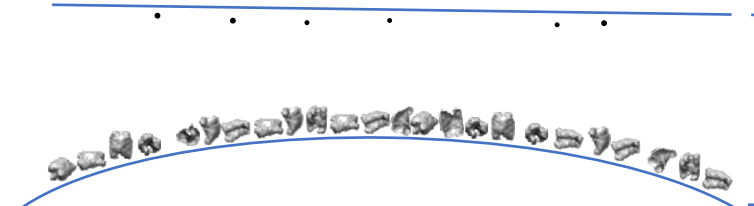
180 nm  
ice



Apoferritin

135 nm  
ice

115 nm  
ice



DNAB Helices

110 nm  
ice

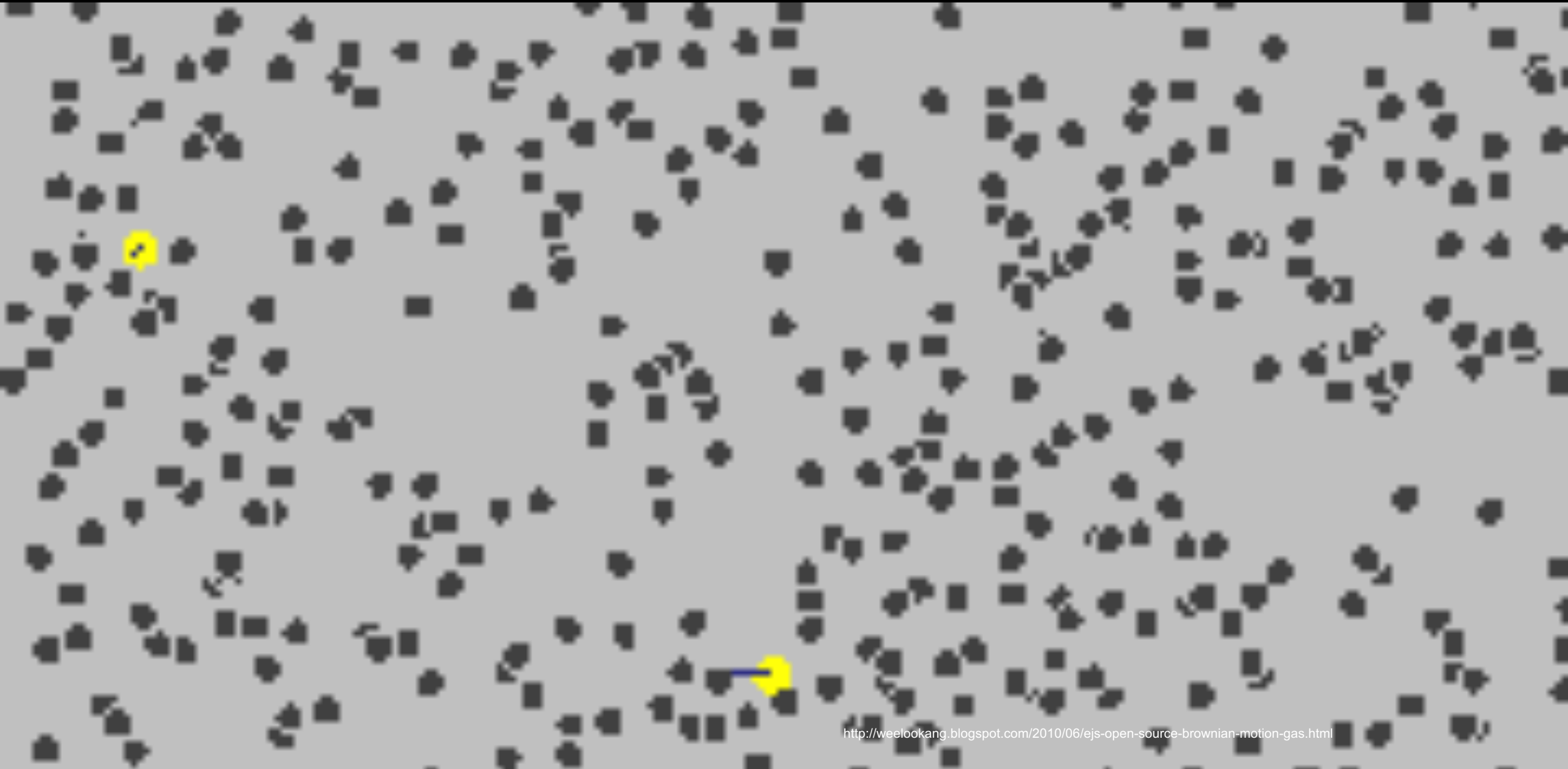
Noble AJ, et al.  
Routine single  
particle CryoEM  
sample and grid  
characterization  
by tomography.  
Elife. 2018;7.



Alex Noble

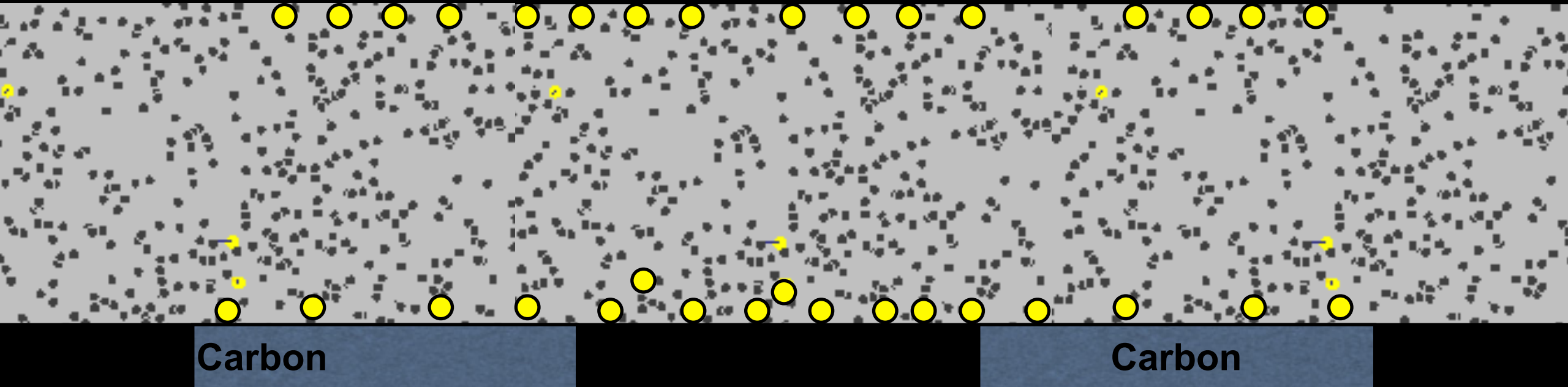


# What happens to samples during vitrification?



# What happens to samples during vitrification?

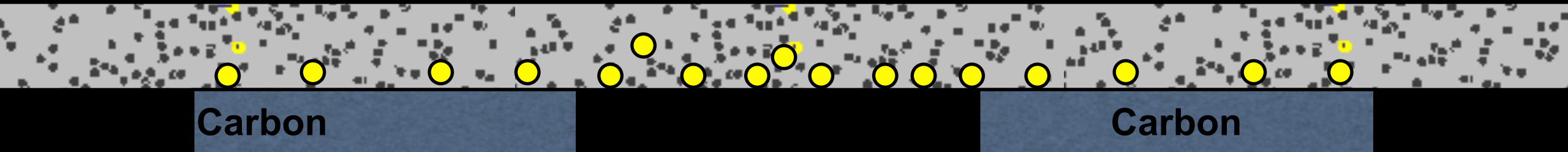
A hypothetical scenario during cryoEM grid preparation

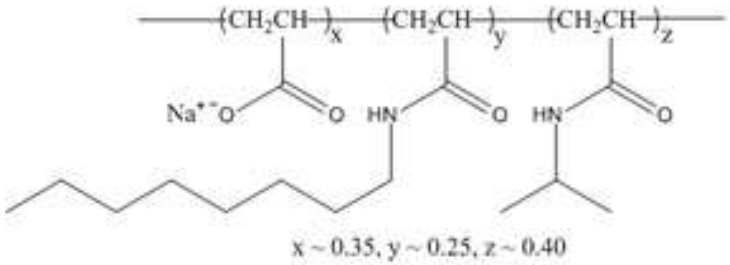




# What happens to samples during vitrification?

A hypothetical scenario during cryoEM grid preparation





Molecular Formula:  
(C<sub>6.2</sub>H<sub>10.3</sub>O<sub>1.35</sub>N<sub>0.65</sub>Na<sub>0.35</sub>)<sub>35</sub>

Molecular Weight:  
approx. 8 kDa

CAS#: 1423685-21-5

# Amphipol A8-35

A short amphipathic polymer that is specifically designed for membrane protein stabilization. The surfactant possesses a very high affinity for the transmembrane surfaces and allows to solubilize membrane proteins in a detergent-free aqueous solution



# REAGENTS FOR IMPROVING VITRIFICATION OF CRYO-EM GRIDS USED IN SINGLE PARTICLE ANALYSIS.

Surfactants and Cryoprotectants	Amount	Conc.	CMC	Class
Fluorinated Octyl Maltoside (FOM)	100 µl	0.41% (w/v)	0.07% (w/v)	non-ionic detergent
Hexadecyl-trimethyl-ammonium Bromide (CTAB)	100 µl	0.34% (w/v)	0.03% (w/v)	cationic detergent
n-Decyl-β-D-Maltoside (DM)	100 µl	0.87% (w/v)	0.09% (w/v)	non-ionic detergent
n-Decyl-α-D-Maltoside (DaM)	100 µl	0.46% (w/v)	0.08% (w/v)	non-ionic detergent
n-Dodecyl-β-D-Maltoside (DDM)	100 µl	0.09% (w/v)	0.01% (w/v)	non-ionic detergent
Sodium Deoxycholate	100 µl	1.66% (w/v)	0.17% (w/v)	anionic detergent
Triton X-100	100 µl	0.15% (w/v)	0.01% (w/v)	non-ionic detergent
Tween 20	100 µl	1% (w/v)	0.01% (w/v)	non-ionic detergent
CHAPSO	100 µl	2.5% (w/v)	0.5% (w/v)	zwitterionic detergent
Amphipol A8-35	100 µl	5% (w/v)		anionic surfactant
Glycerol	1 ml	30% (w/v)		cryoprotectant

- [1] Noble *et al.* (2018) Routine Single Particle CryoEM Sample and Grid Characterization by Tomography. DOI: 10.7554/eLife.34257.
- [2] Thonghin *et al.* (2018) Cryo-electron microscopy of membrane proteins. *Methods* **147**:176.
- [3] Drulyte *et al.* (2018) Approaches to altering particle distributions in cryo-electron microscopy sample preparation. *Acta Cryst. D* **74**:560.
- [4] Glaeser *et al.* (2017) Opinion: hazards faced by macromolecules when confined to thin aqueous films. *Biophys Rep* **3**:1.
- [5] Gatsogiannis *et al.* (2016). Membrane insertion of a Tc toxin in near-atomic detail. *Nat. Struct. Mol. Biol.* **23**:884.
- [6] Efremov *et al.* (2015) Architecture and conformational switch mechanism of the ryanodine receptor. *Nature* **517**:39.

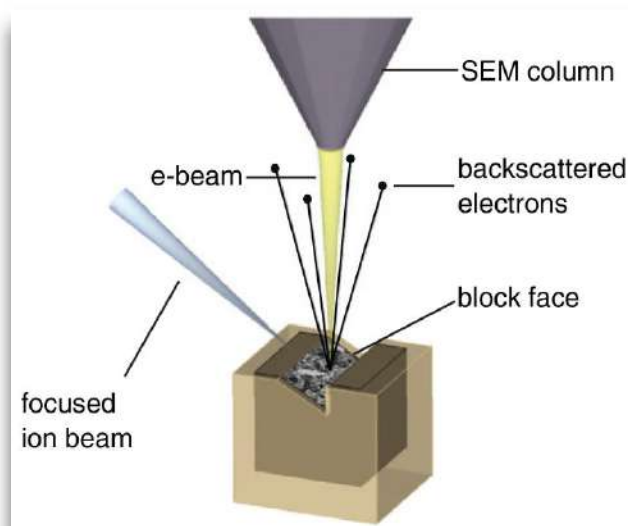
<https://www.mitegen.com/product/cryo-em-vitrification-starter-kit/>

# FIB/SEM VS THIN SECTION SAMPLE PREP

- Chemical fixation
- Staining
  - En bloc, enhanced contrast and electrical conductivity
- Dehydration
- Embedding
- Au/Pd coat
  - Conductivity

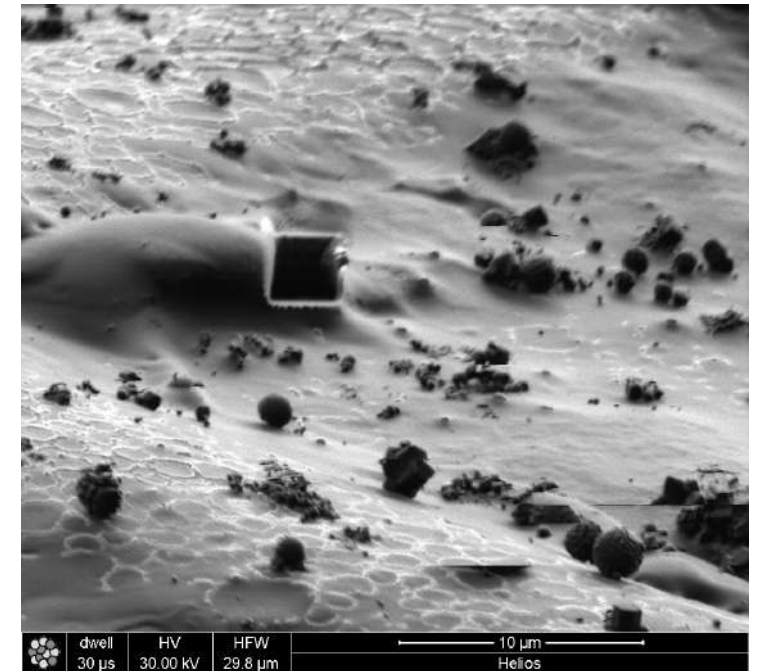
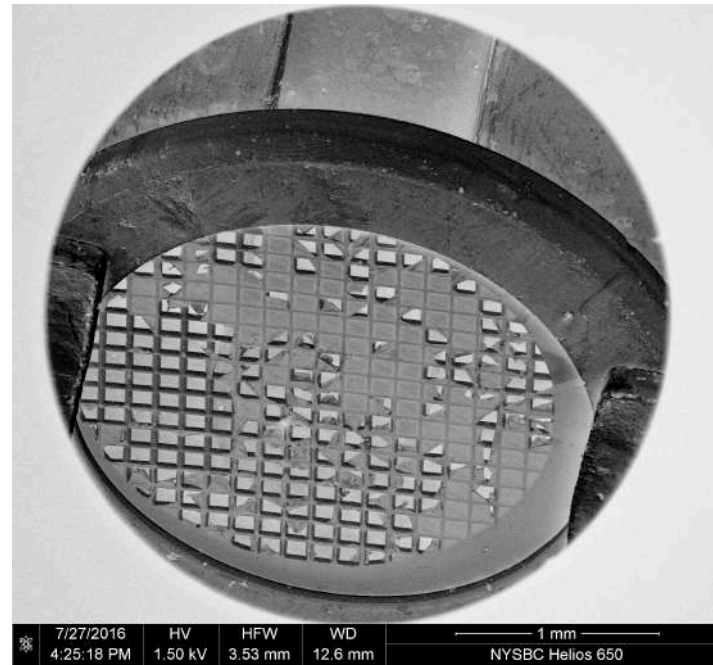
Cryofixation: High pressure freezing  
Dehydration: Freeze substitution

- Chemical fixation
- Dehydration
- Embedding
- Sectioning
- Staining

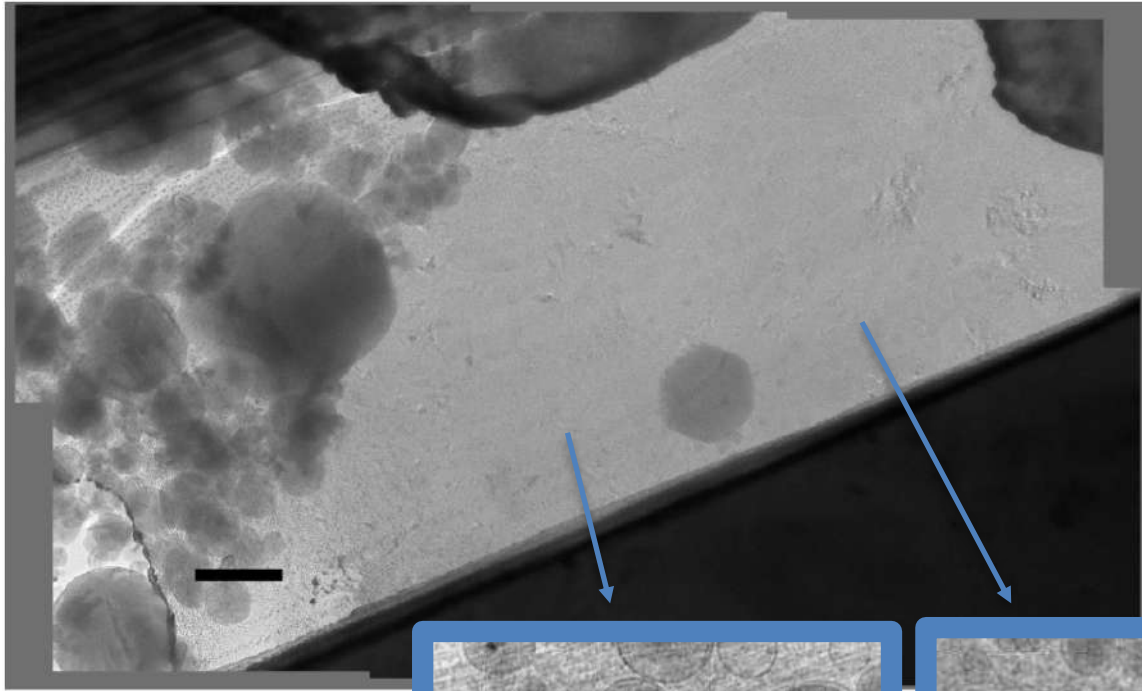




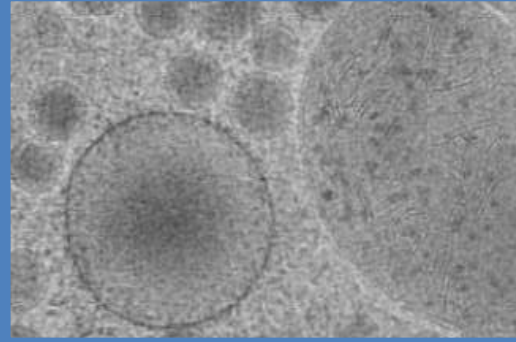
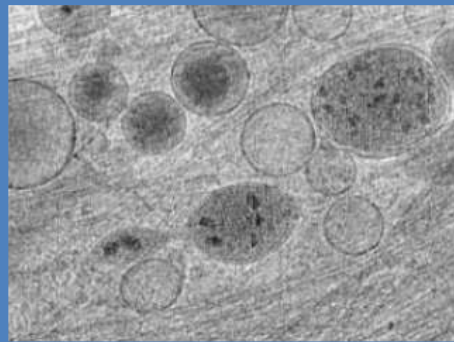
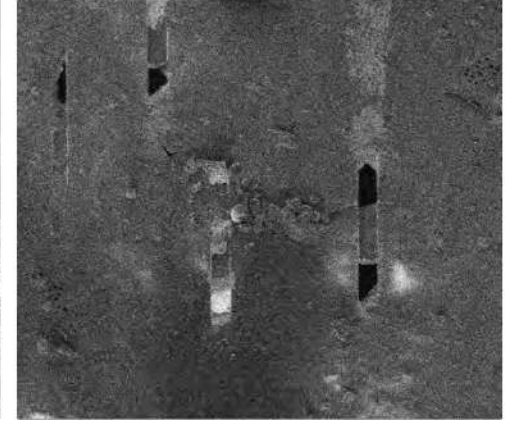
# CRYO FIB MILLING



# CRYO FIB MILLING

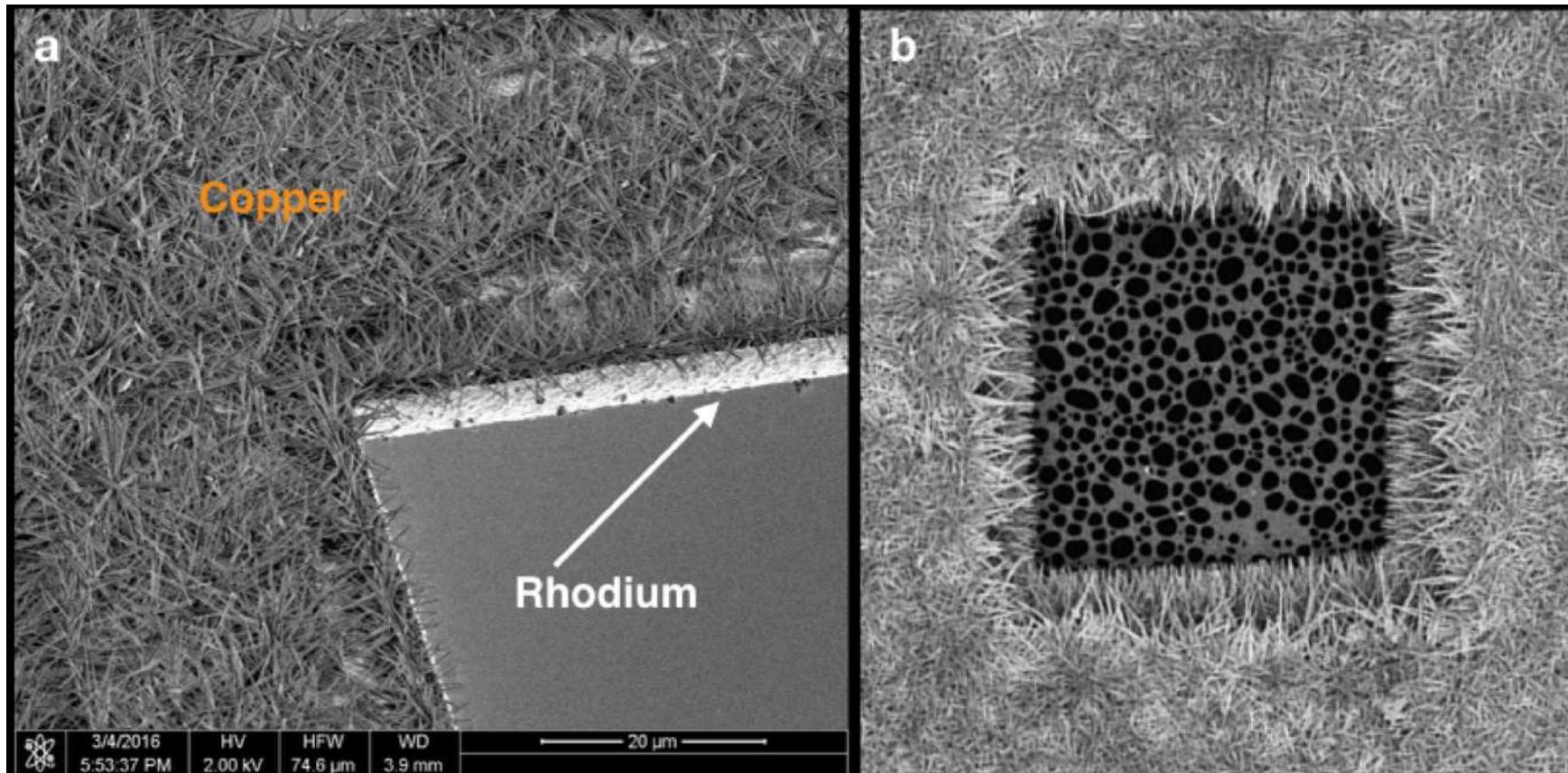


Scale bar: 1  $\mu\text{m}$

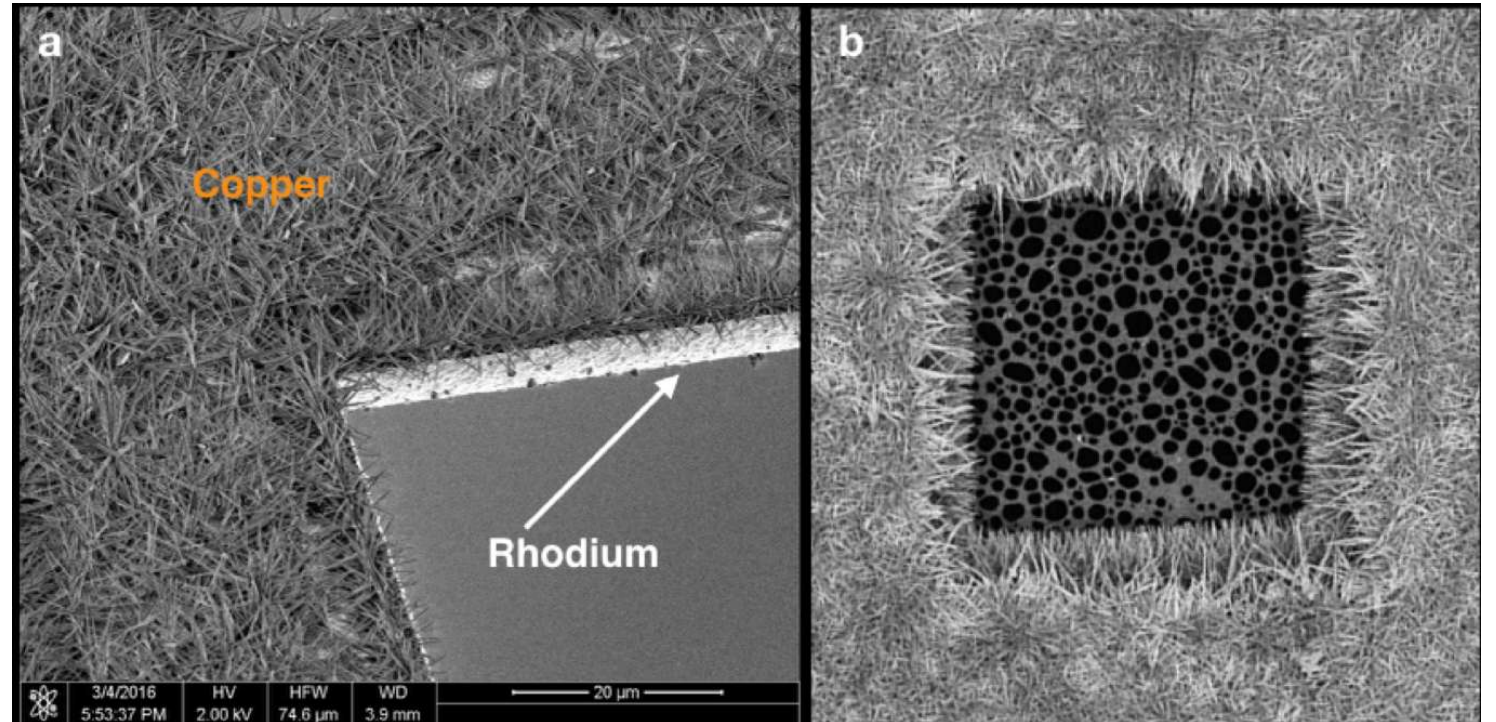




# BLOT FREE VITRIFICATION

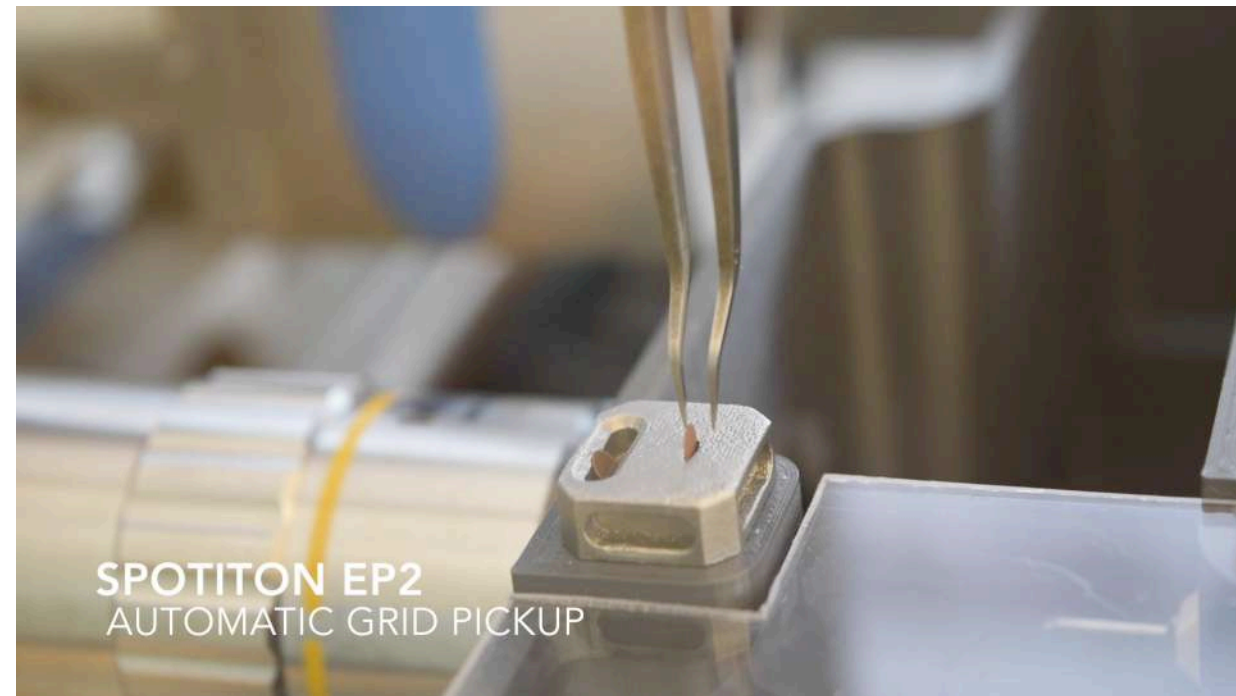
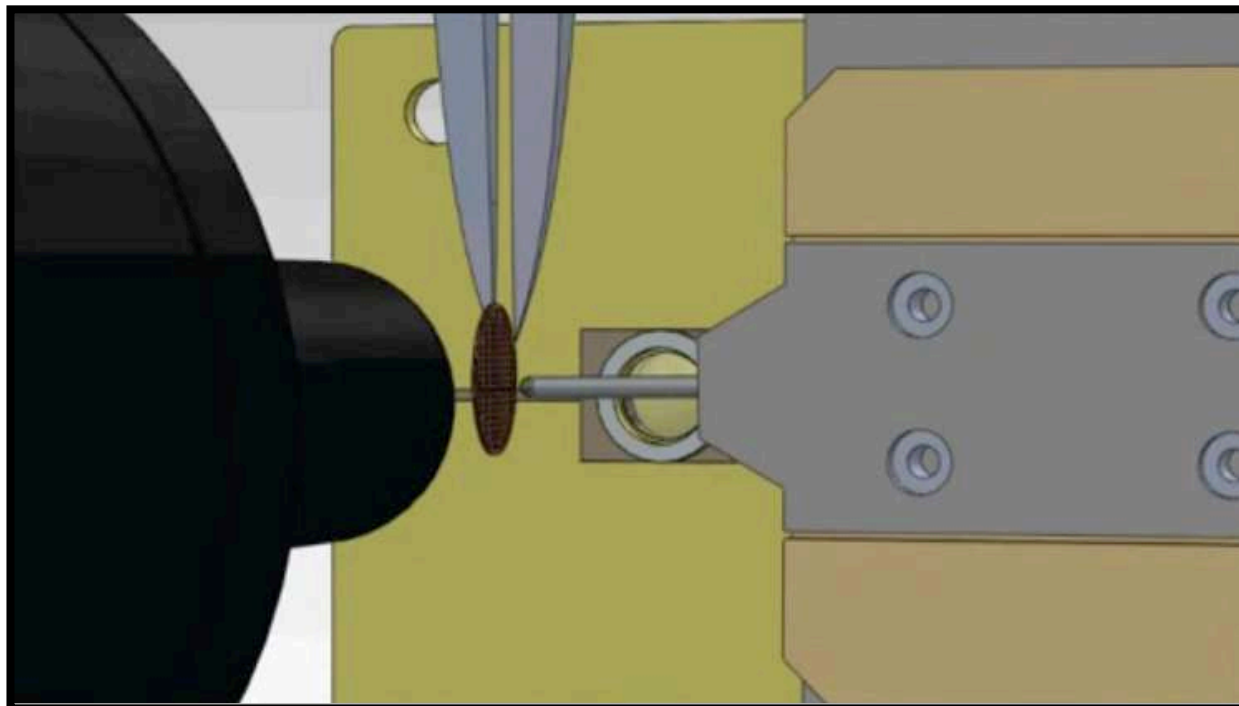


# SPOTITON | CHAMELEON





# SPOTITON | CHAMELEON



# WHAT NEXT?

## cryoEM 001 : Single Particle Masterclass

1. Building a cryoEM toolkit
2. EM compatible samples
3. EM support films and grids
4. Sample preparation
5. Tools of the trade:  
microscopes and detectors
6. Microscope operations
7. Data collection strategies
8. Data assessment & QC
9. Data processing:
  - cryoEM IT infrastructure
  - On-the-fly feedback
  - 3D Reconstruction
10. Visualization and validation



# DIFFICULT SPECIMENS

## Small protein

- VPP
- Thinner ice

## Protein denaturation/Dissociation of protein complex

Continuous carbon film

Graphene oxide

Cross-linking (GraFix)

## Preferred orientation

Tilt stage

Cross-linking

Detergent

Glow-discharging conditions

Support film (Graphene oxide)

Image analysis (3D classification)

## Flexibility

Focused classification (subtraction)

Multibody refinement

## Filamentous protein

- Segmented analysis

## VI. Low concentration

Multiple blots

Affinity grids