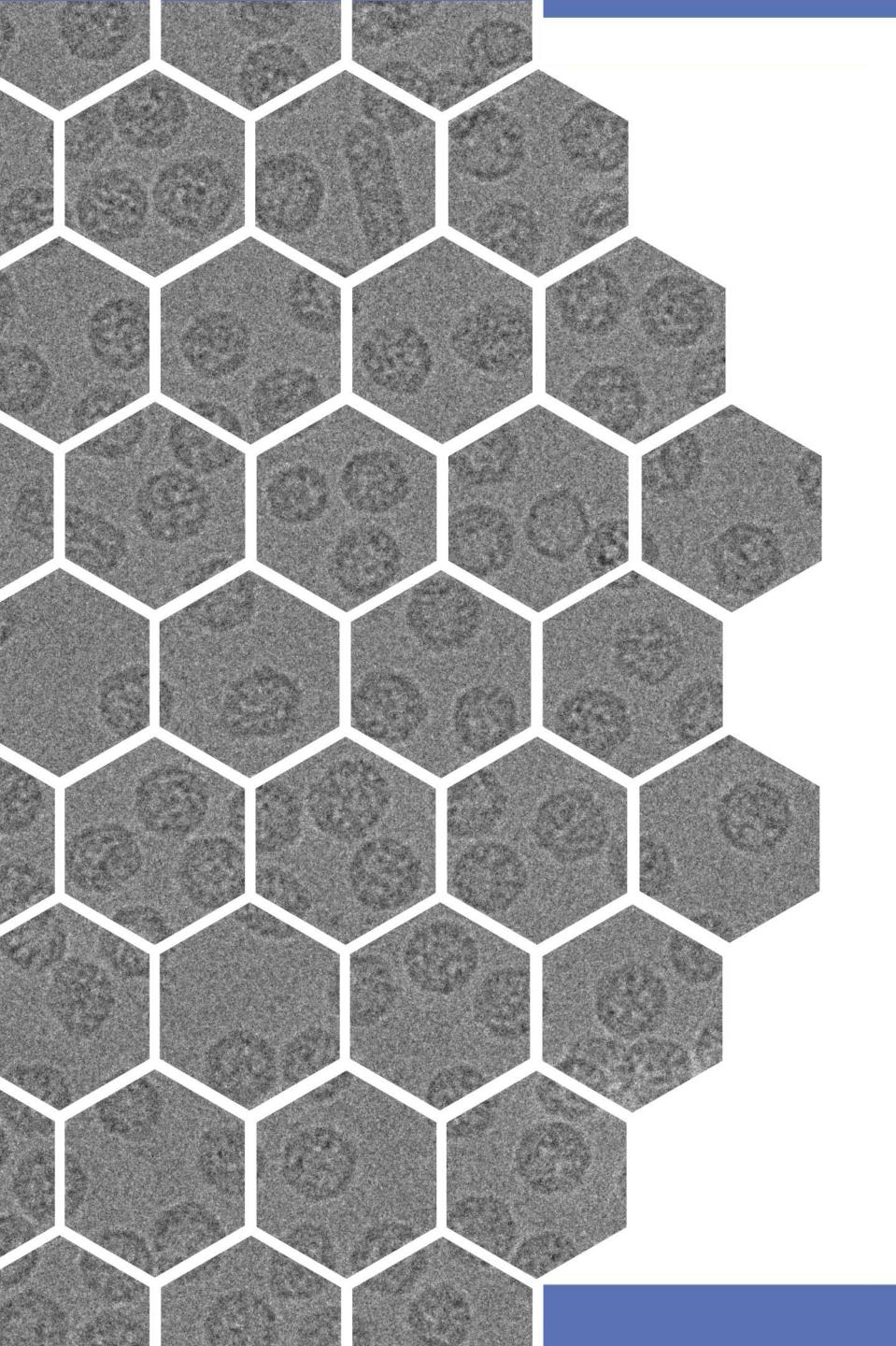
2024 Winter cryoEM course **Considerations for biological cryoEM**

January 29, 2024





Journal club and practical recap Considerations for biological cryoEM

- Overview
- Grids
- What happens to a sampleNewer methods



Course logistics: main topics

Section I: EM fundamentals Section 2 : EM crystallography Section 3 : Tomography Section 4 : SPA short course* March 11-15 Section 5 : Future perspectives





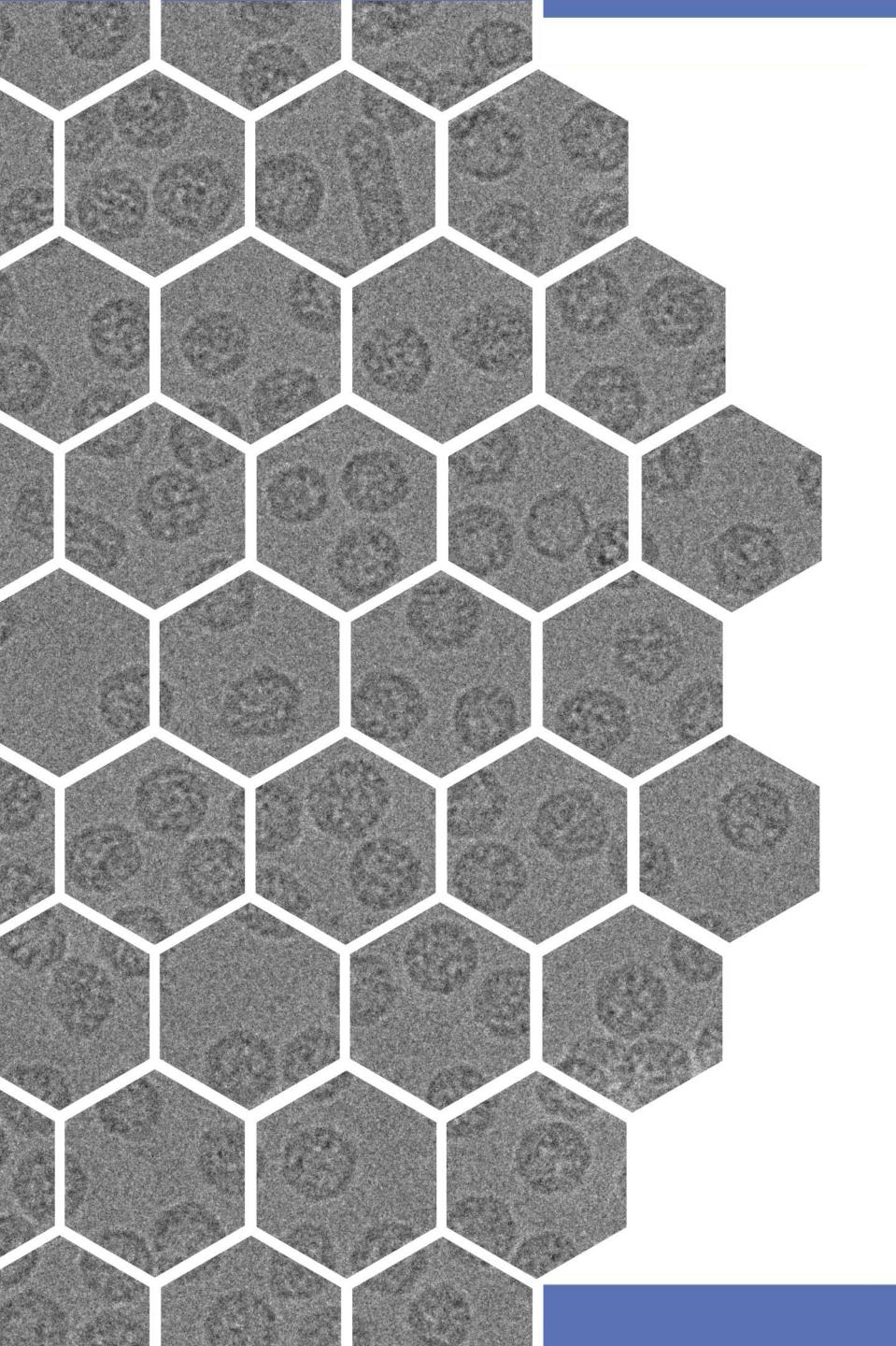
Course logistics: Wednesday practical

January 31, 2024

Sample preparation practical -standard vitrification -45 min first session -30 min second session





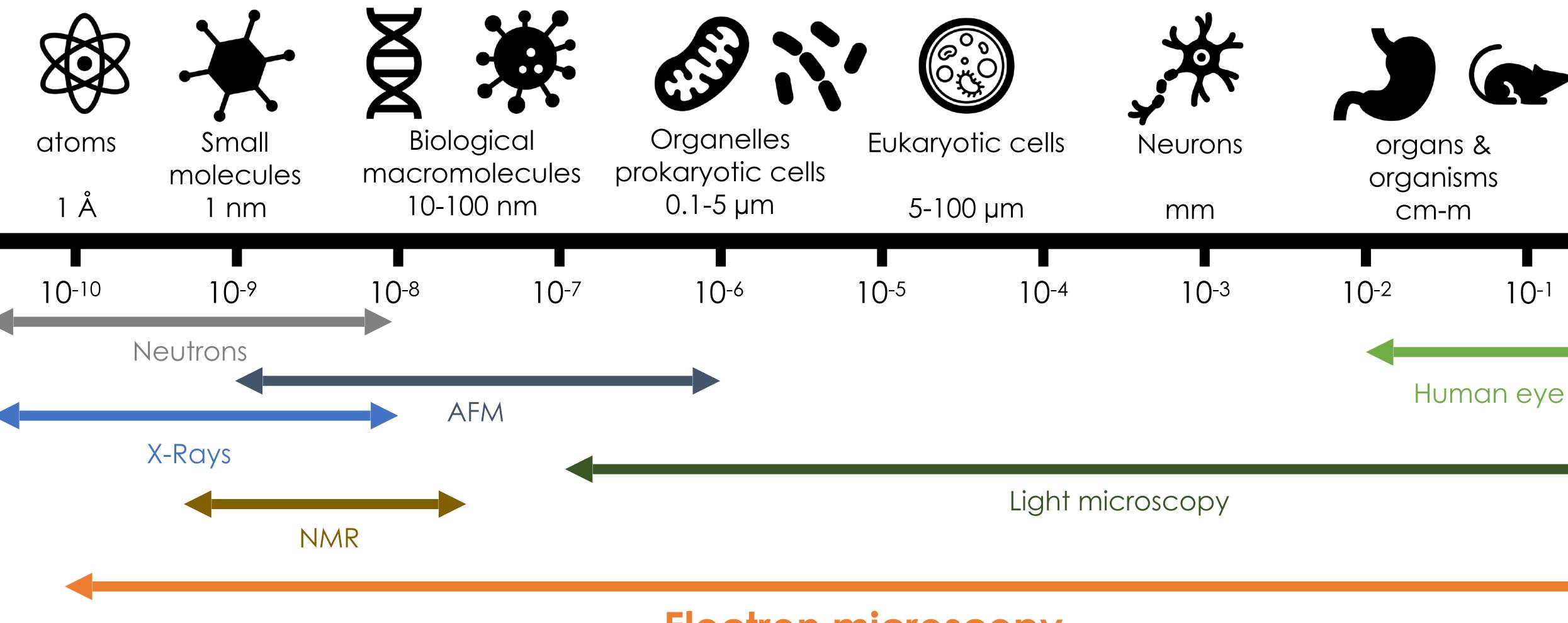


Journal club and practical recap Considerations for biological cryoEM

- Overview
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- What happens to a sampleNewer methods

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Scale of biology



doi: 10.2210/rcsb_pdb/goodsell-gallery-014



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Electron microscopy

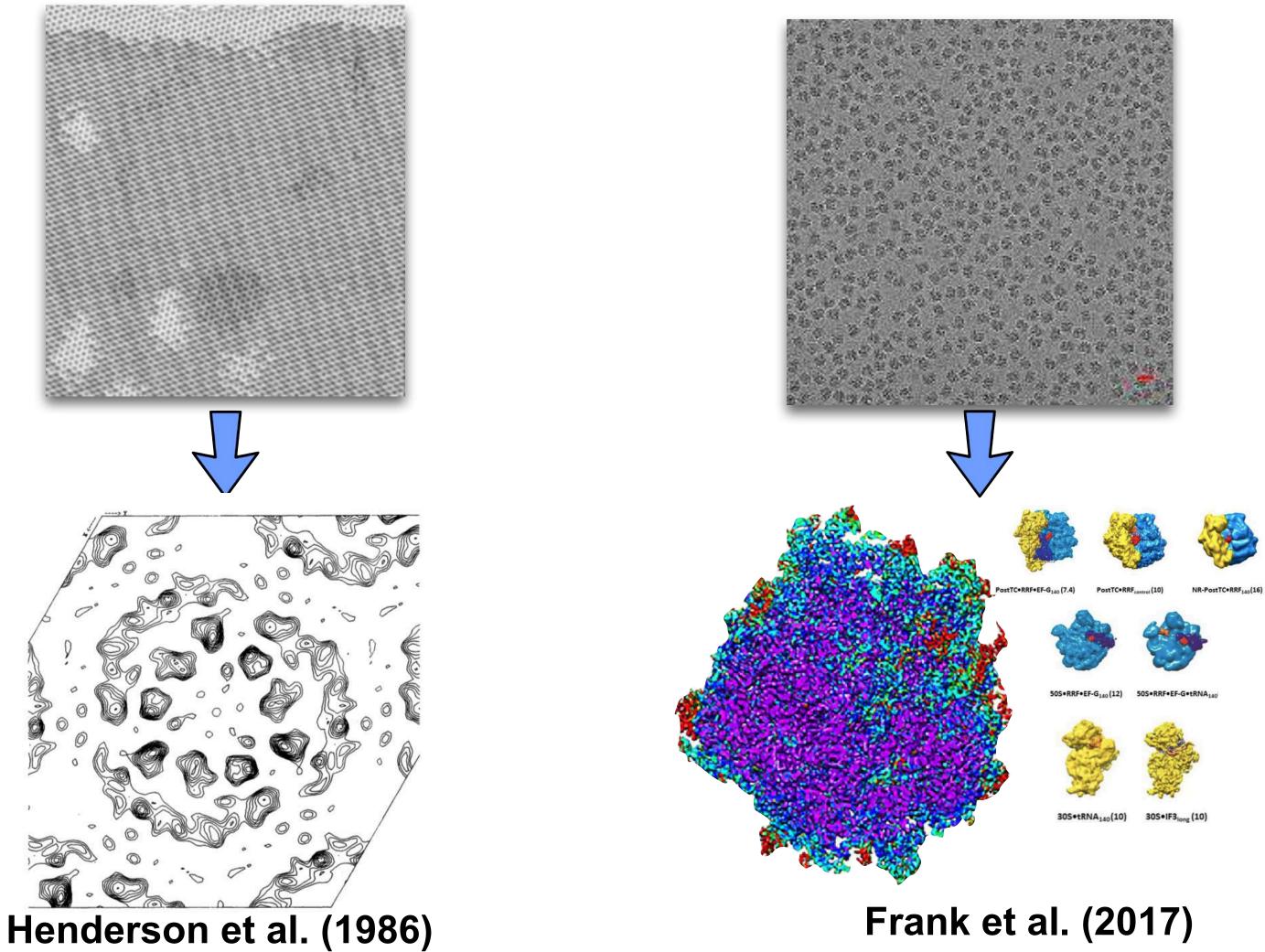






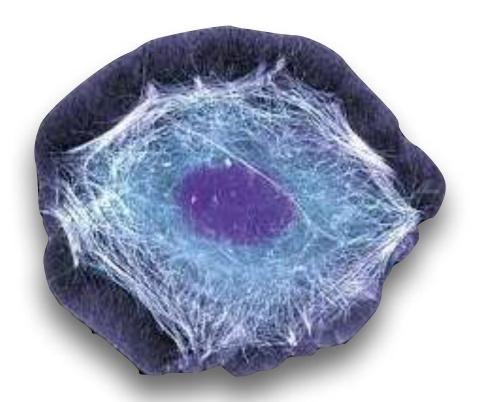
cryoEM: technology on the rise

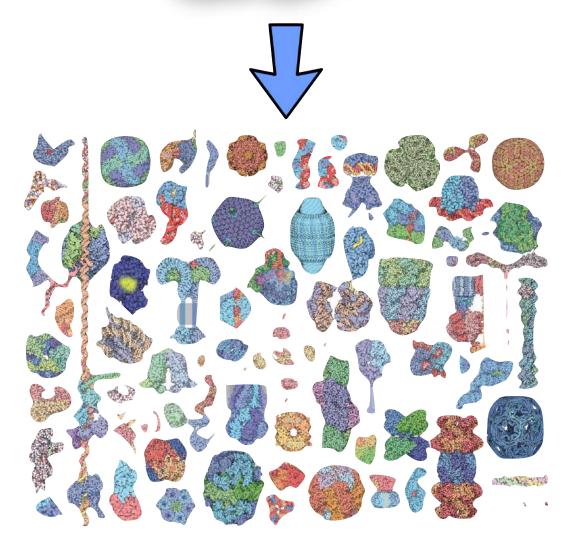
1986



2017

in progress





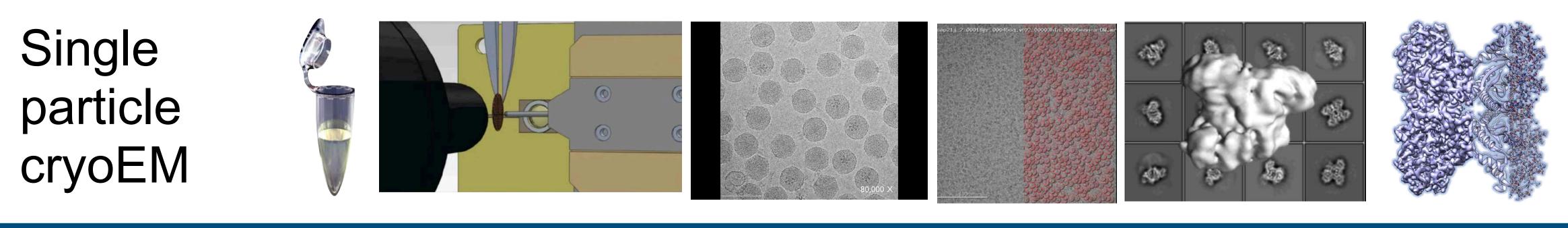
the next chapter

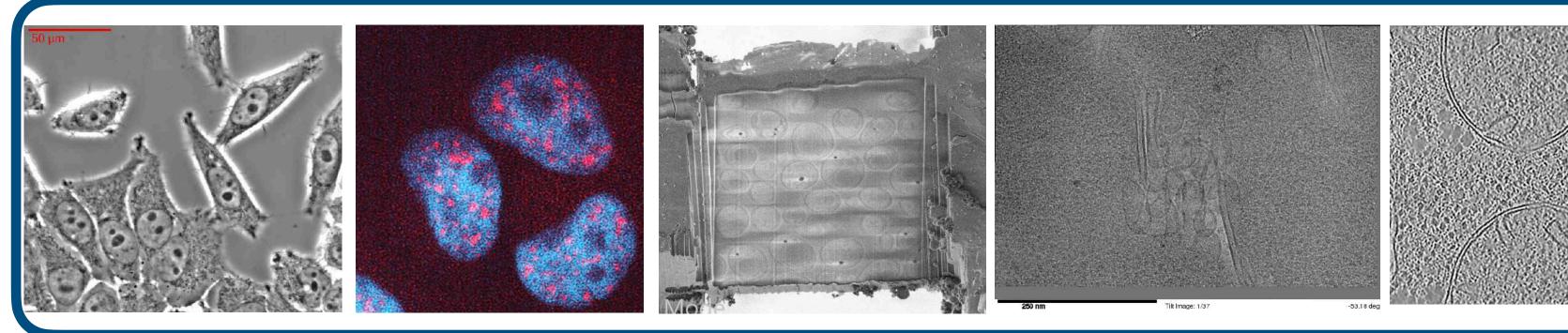
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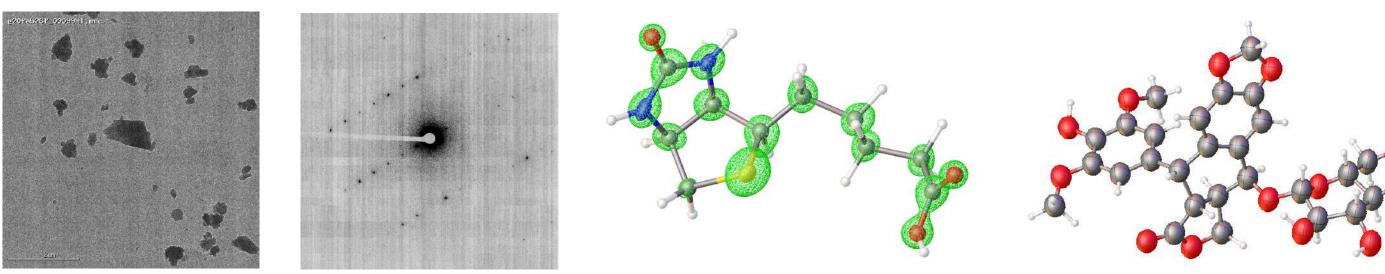
cryoEM: a technology on the rise





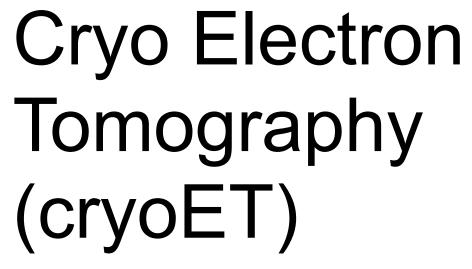
Micro crystal electron diffraction (microED)

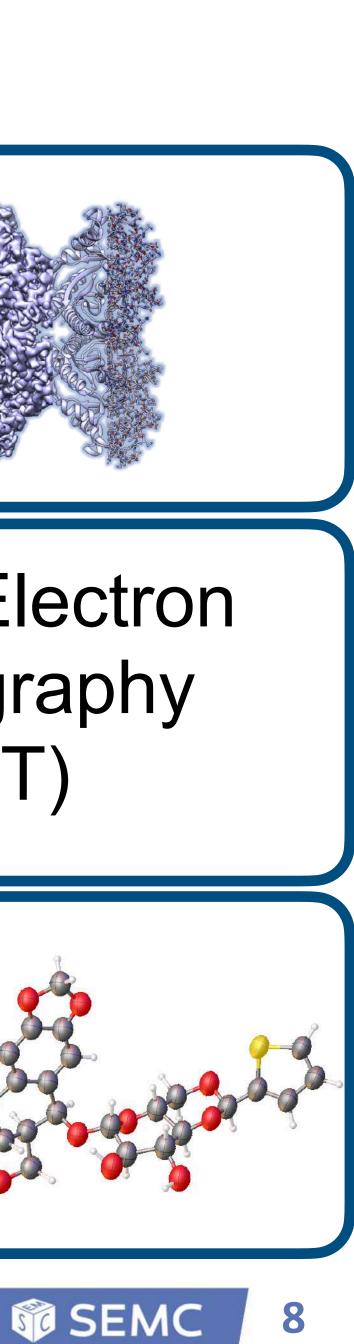




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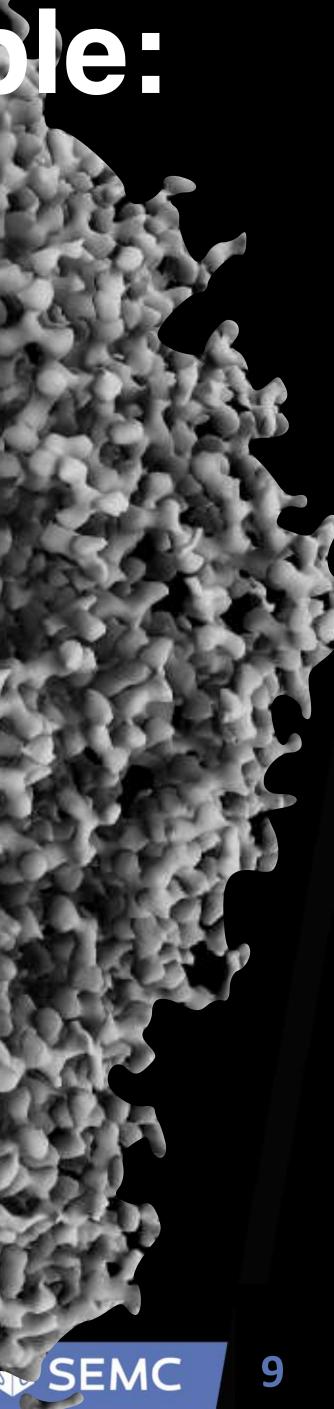
2 µm

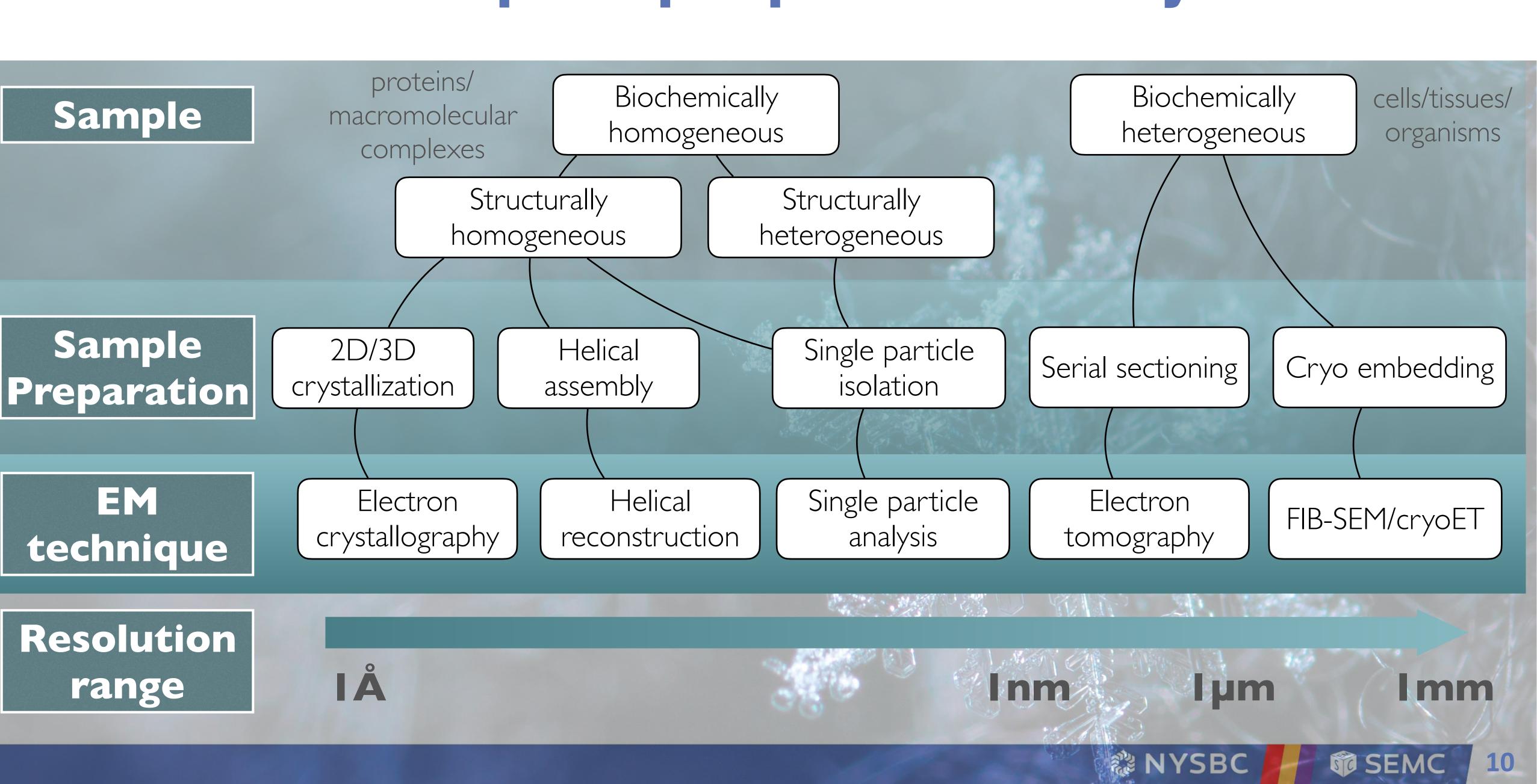


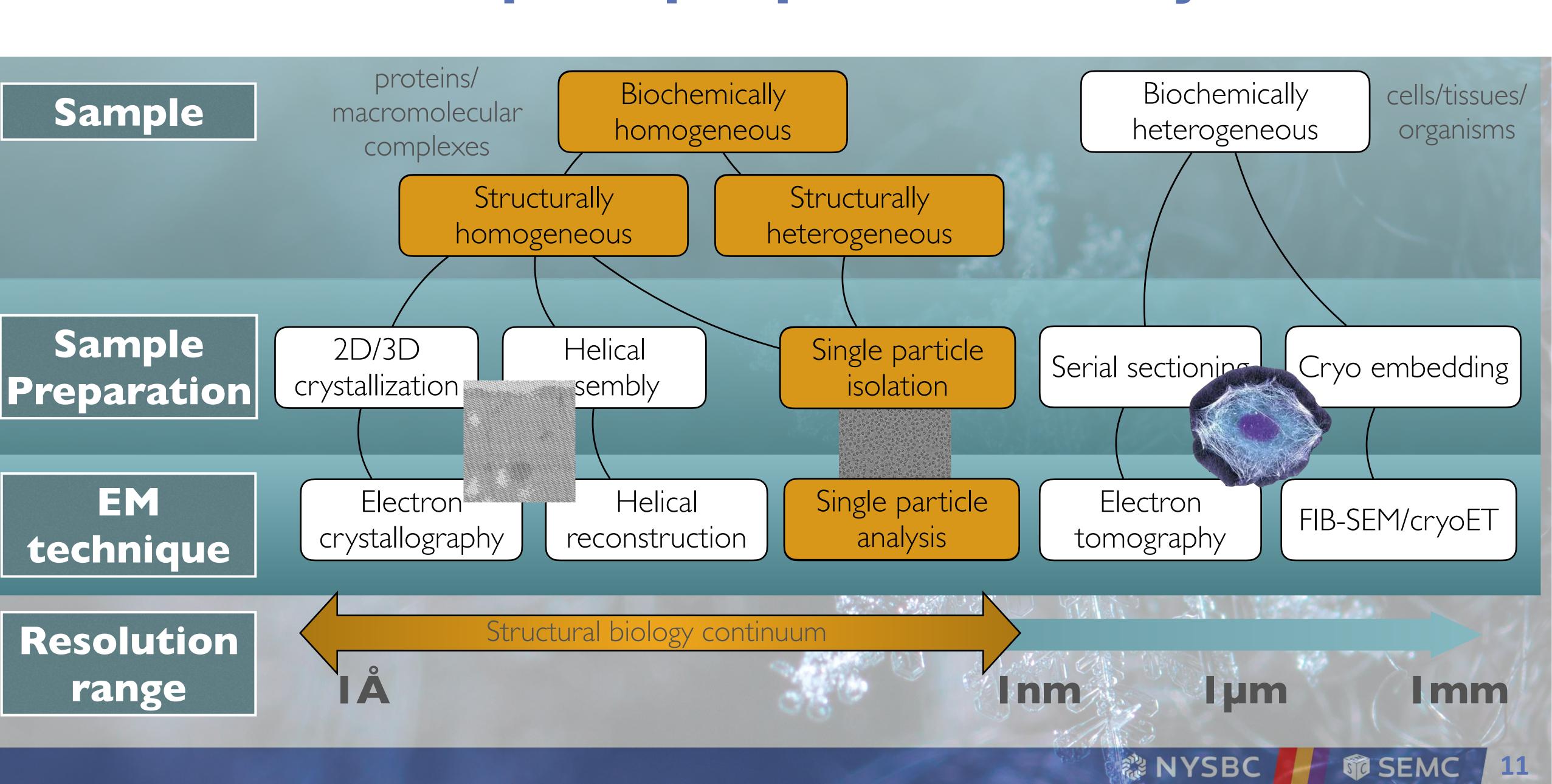


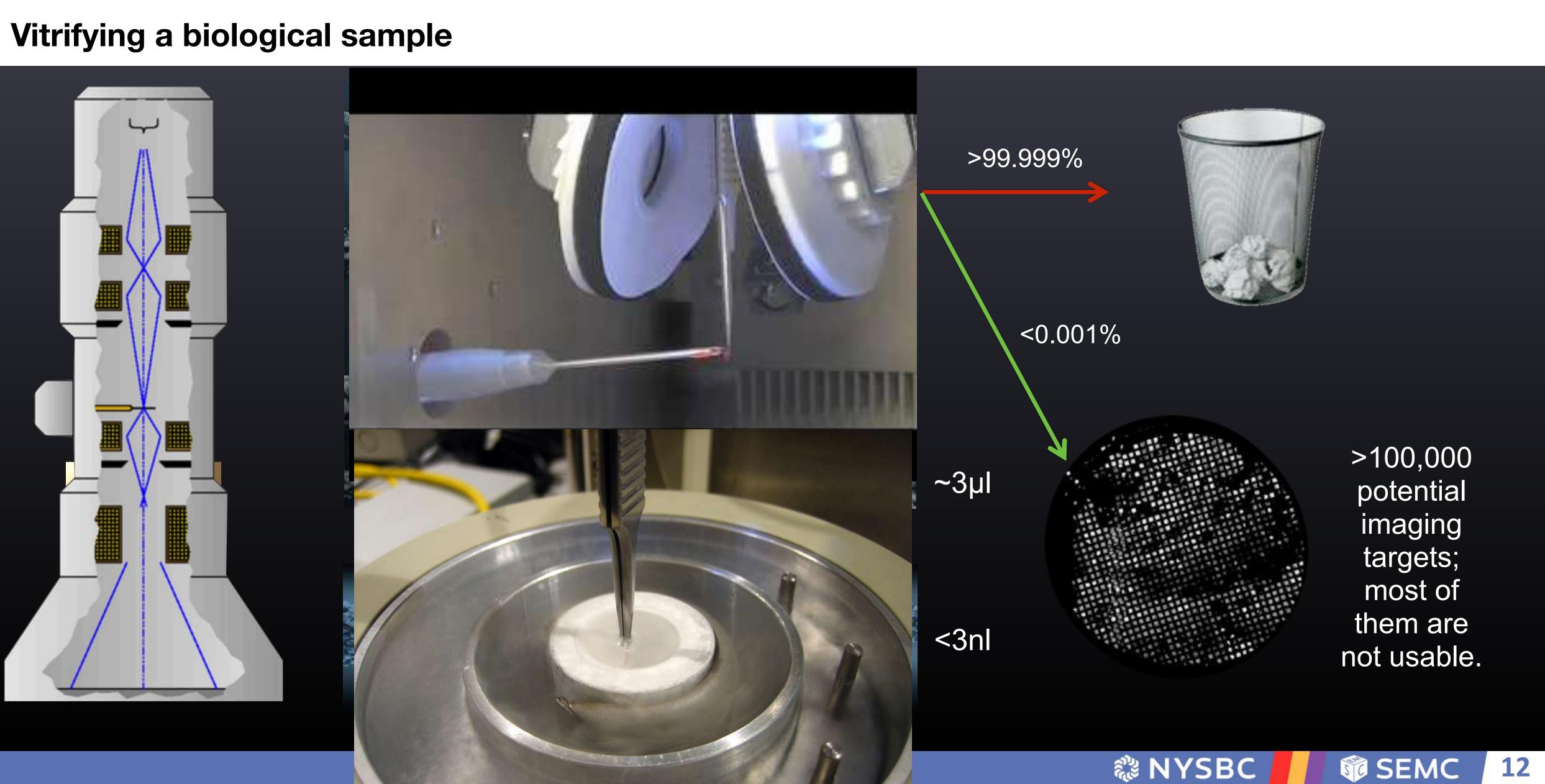
And true "atomic" resolution is possible:

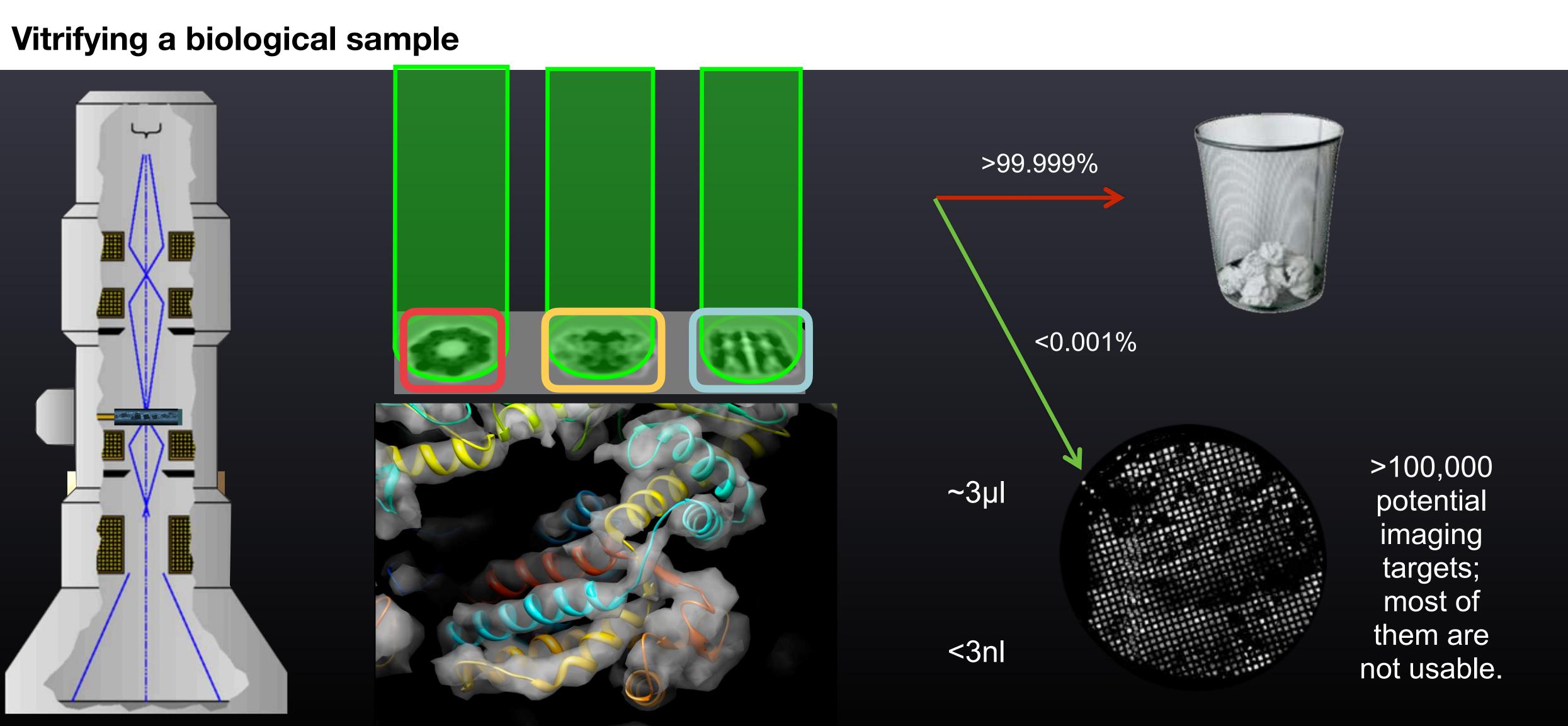
Nakane, et al. Single-particle cryo-EM at atomic resolution. Nature (2020).





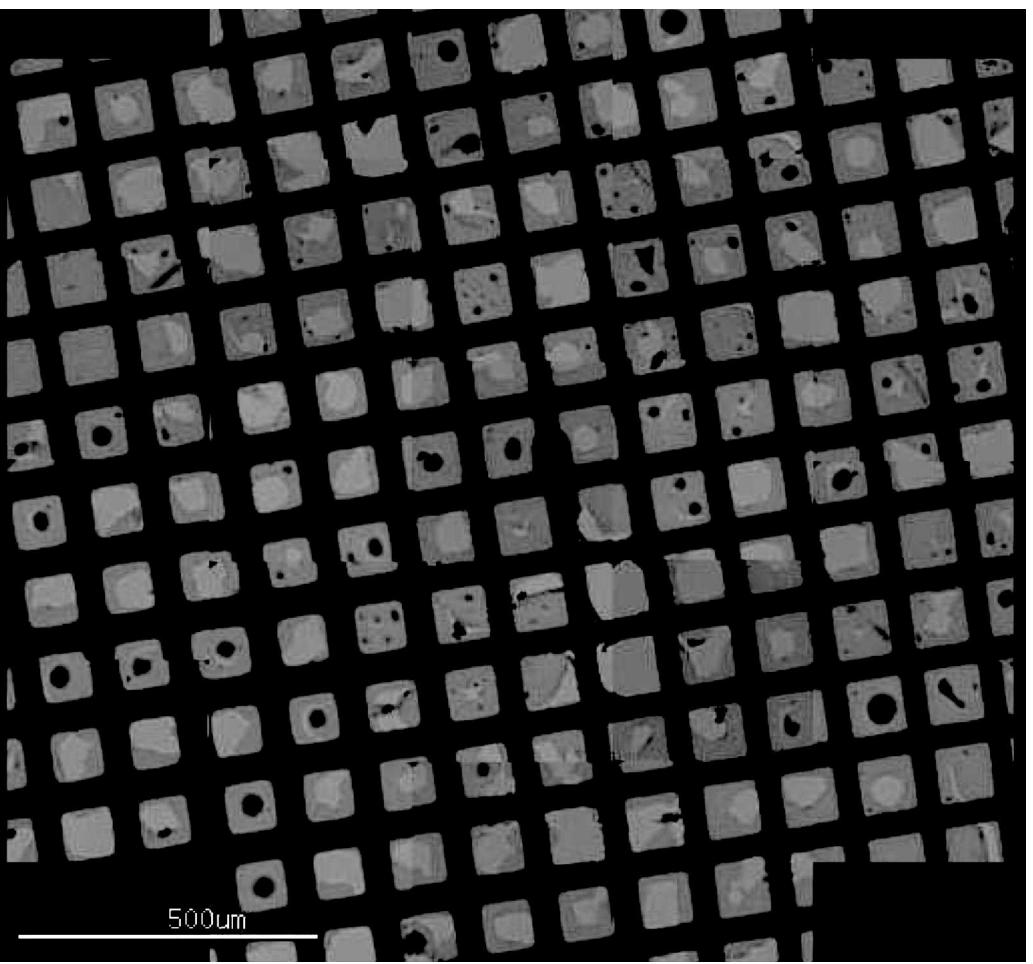




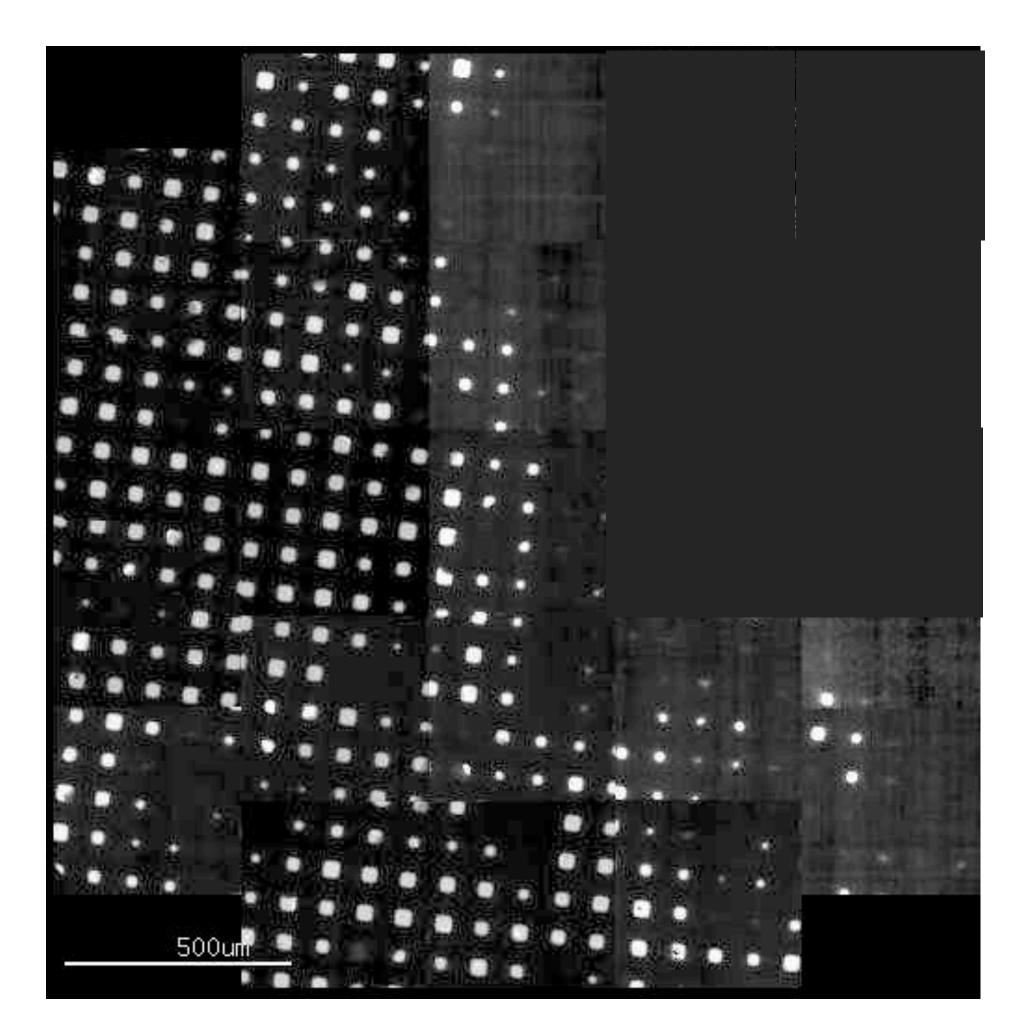


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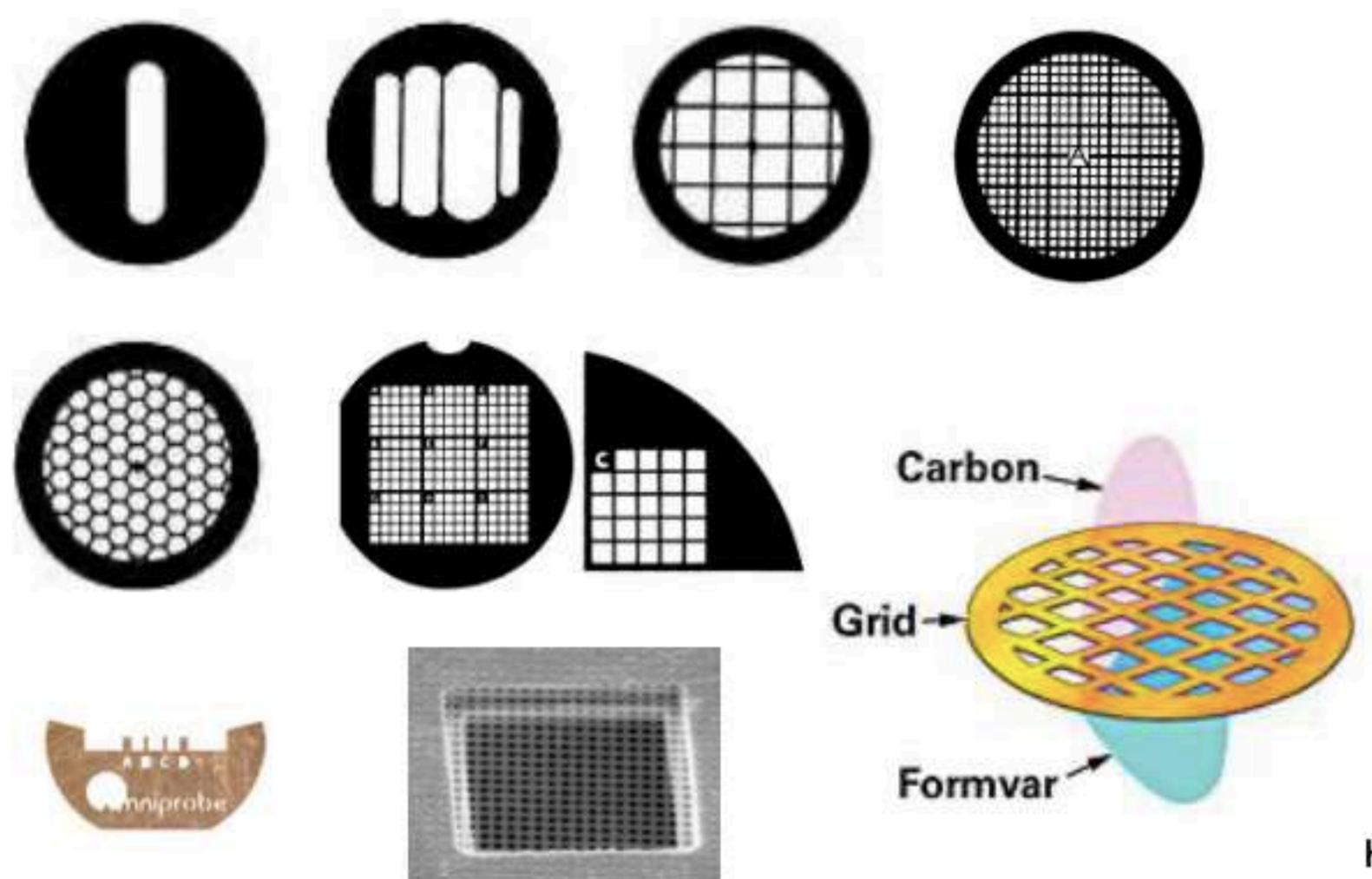




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Common Materials

Copper

Nickel

Gold

Aluminum

Molybdenum

Titanium

Stainless Steel

https://www.tedpella.com/grids_html/

🗱 NYSBC **© SEMC**









Rough grid parameters

Rim Width:		350-400µm.		
Thickness:		approximately 25µm thick.		
Diameter:		3.0 to 3.05mm		
Pitch:	ls 1"/mes	sh or 25.4mm/mesh		

Example 200 mesh pitch = $25.4/200 = 127\mu$ m

PELCO[®] Grid Size

Square Mesh	Pitch µm	Hole µm	Bar µm
50		508	
75		339	
100		254	
150		169	
200		127	
300		85	
400		64	
500		51	



% Trans-mission		
425	83	
284	55	
204	50	
125	44	
90	37	
54	31	
38	26	
28	23	





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Grid (Cu, Au, Mo, etc...) mesh

Foil (C, Au, etc...)

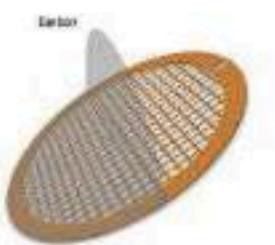
- Continuous
- lacy
- holey (hole size and spacing)







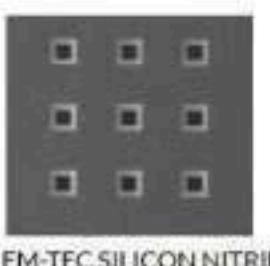




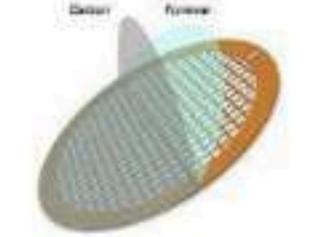
HOLEY CARBON SUPPORT FILMS



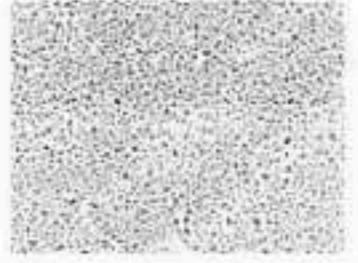
FORMVAR ONLY SUPPORT FILMS



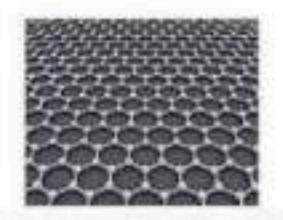
EM-TEC SILICON NITRIDE SUPPORT FILMS



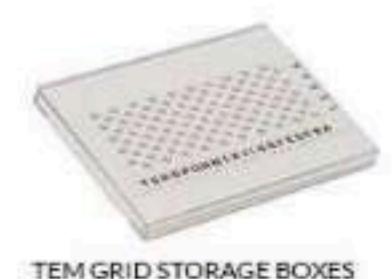
FORMVAR/CARBON SUPPORT FILMS



TEM CALIBRATION & TEST STANDARDS



EM-TEC GRAPHENE SUPPORT FILMS

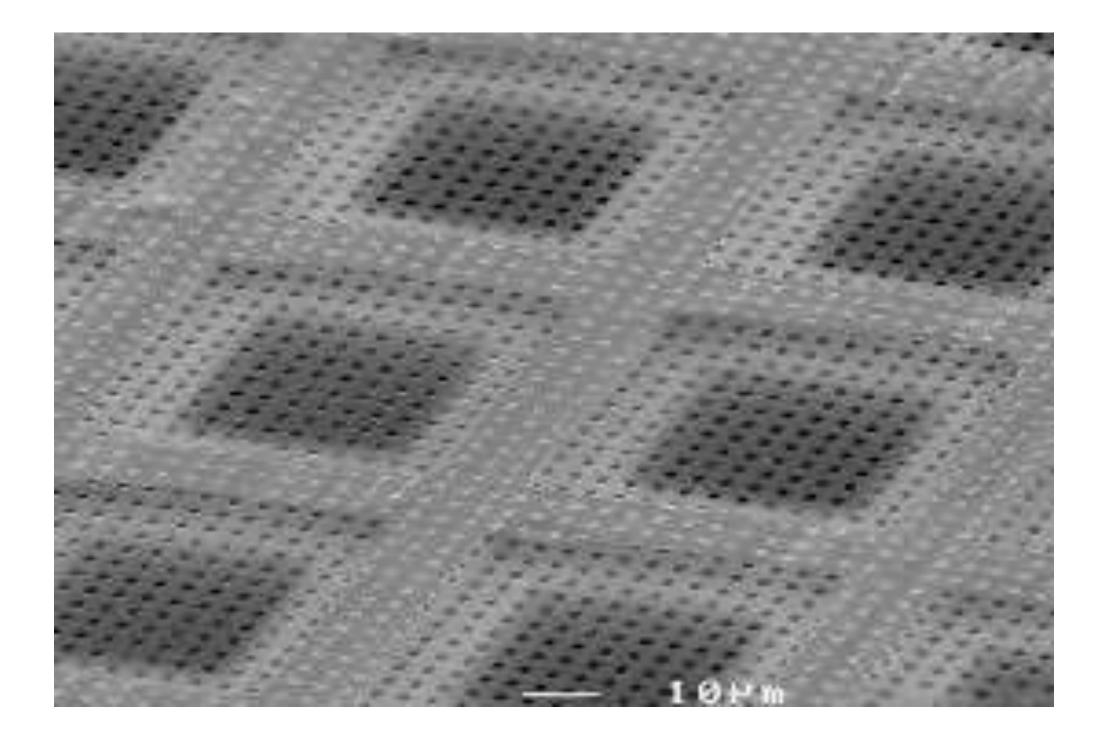


https://edgescientific.com/product-category/tem-supplies/tem-support-films/



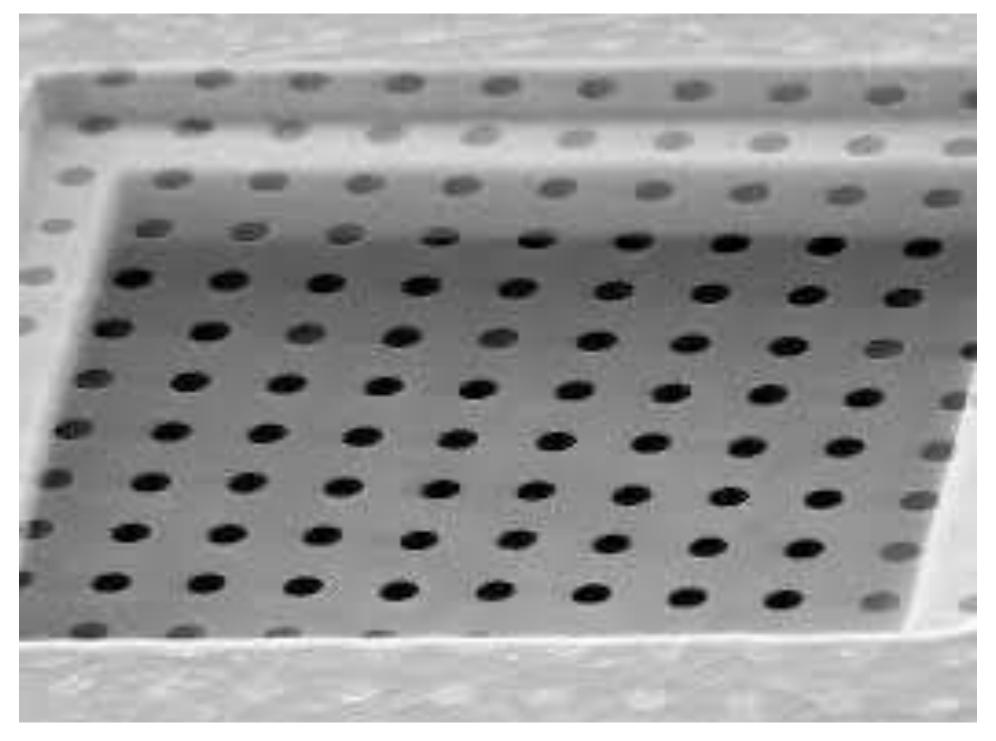


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• Protochips.com





Quantifoil.com



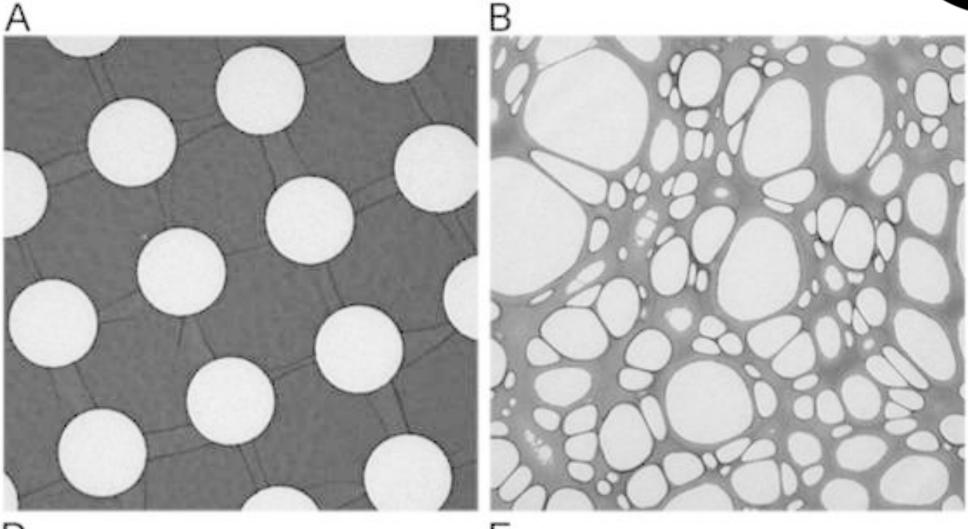


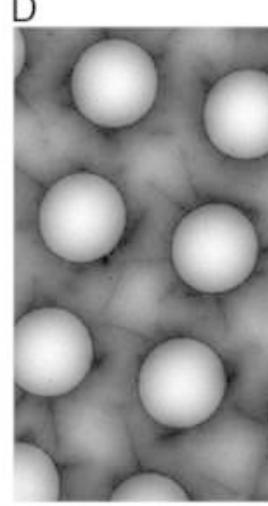




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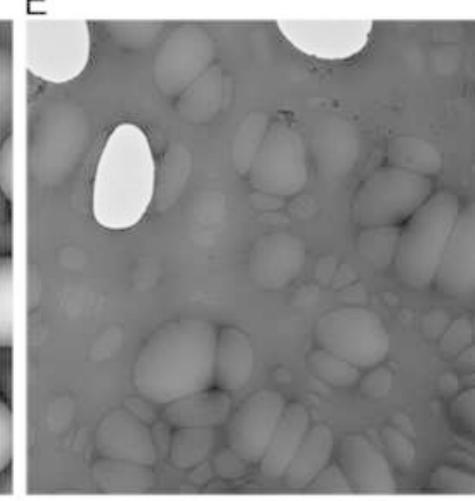
Cho, Hye-Jin & Hyun, Jae-Kyung & Kim, Jin-Gyu & Jeong, Hyeong & Park, Hyo & You, Dong-Ju & Jung, Hyun. (2013). Measurement of ice thickness on vitreous ice embedded cryo-EM grids: investigation of optimizing condition for visualizing macromolecules. Journal of Analytical Science and Technology. 4. 10.1186/2093-3371-4-7.

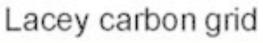


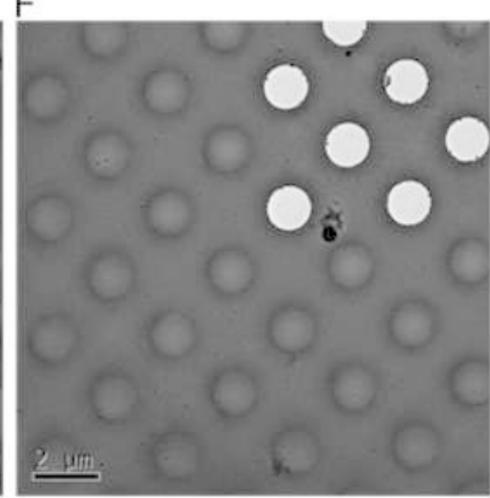


Quantifoil grid









C-flat grid

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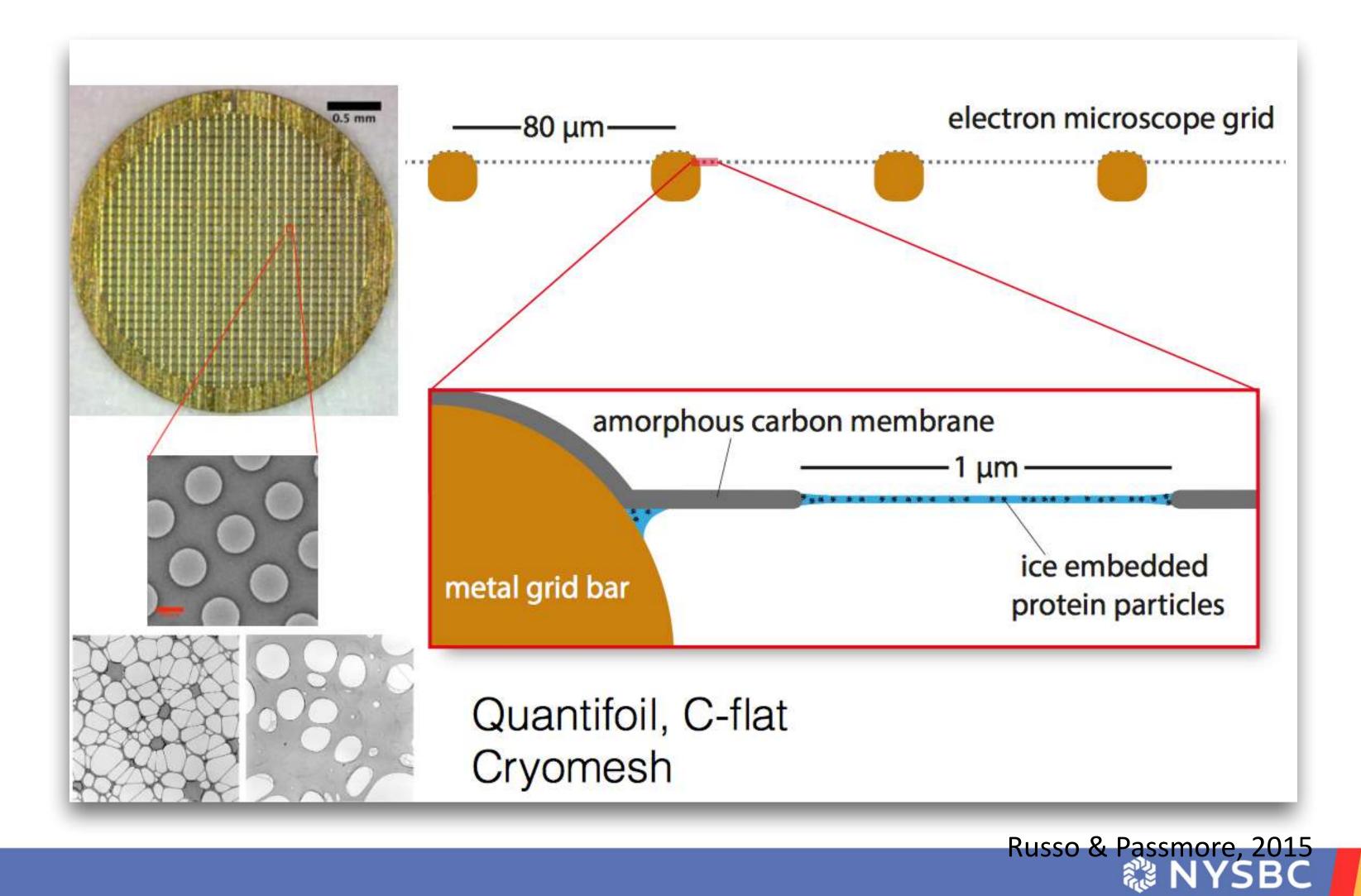








TERMINOLOGY





SEMC





TERMINOLOGY

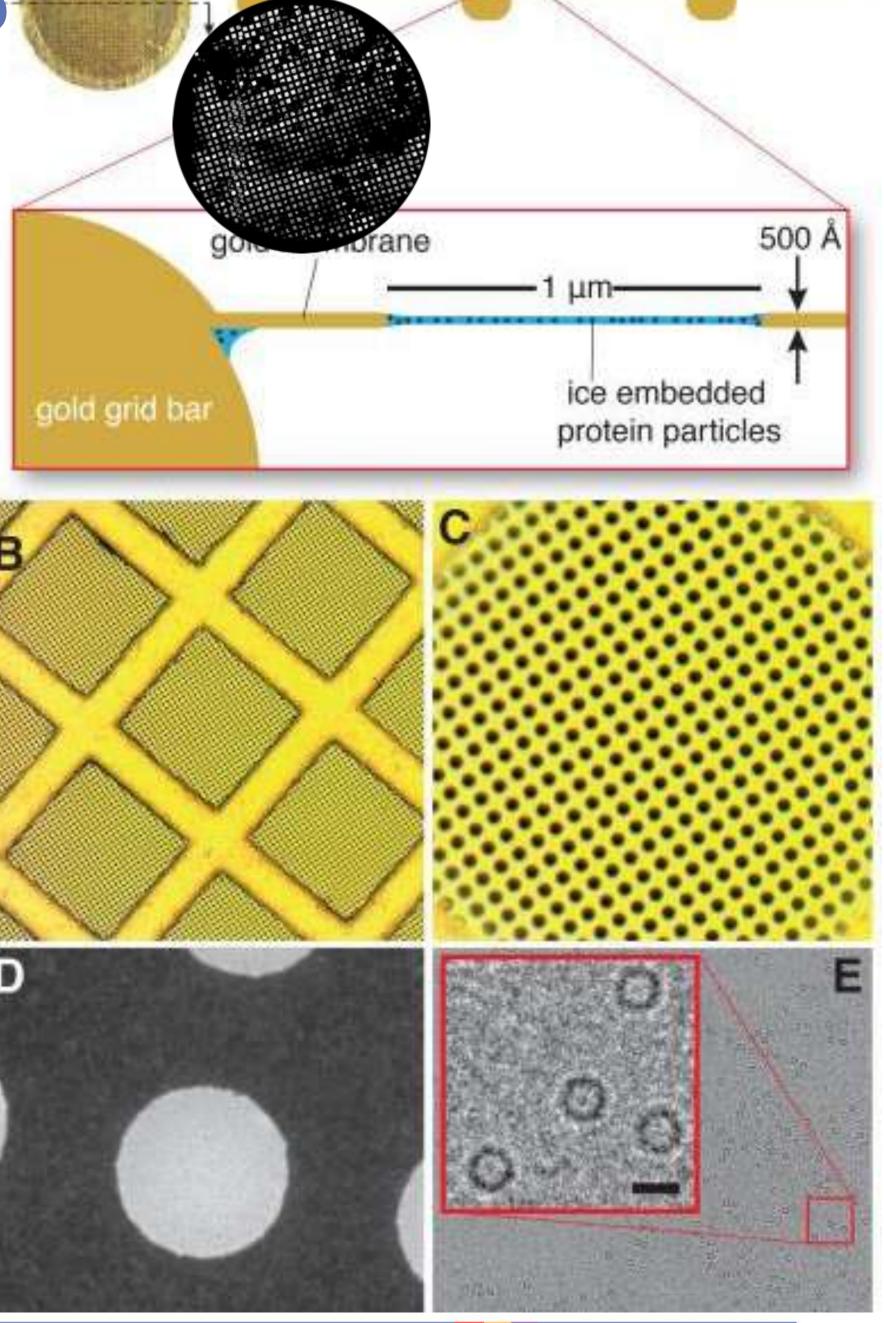
Holey gold foil on gold mesh grid

Advantages:

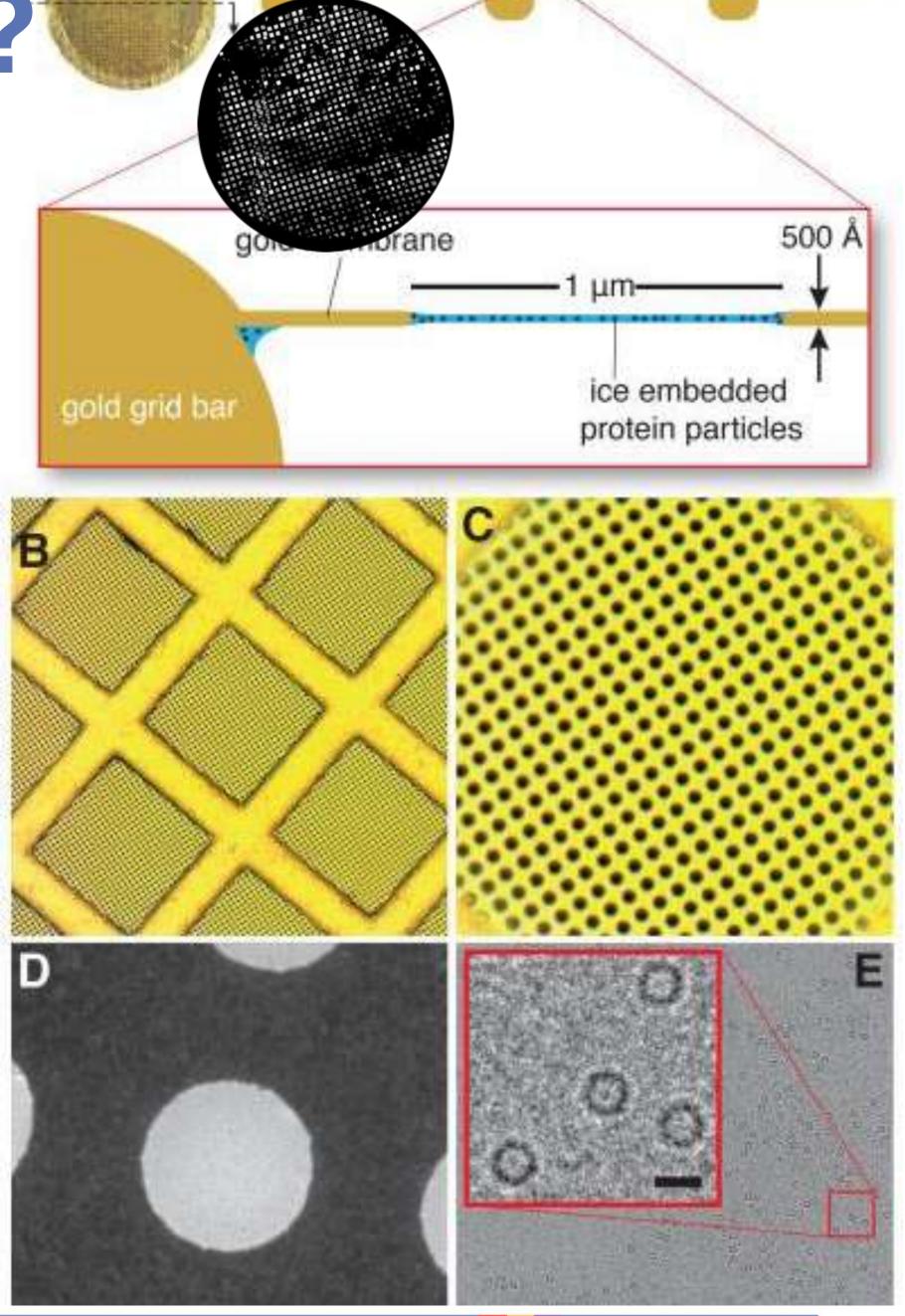
- Prevents differential thermal contraction when freezing
- Reduces beam-induced specimen movement
- Combined with direct detector technology allows for near atomic resolution

Disadvantages:

Difficult to find focus due to lack of amorphous substrate



Russo & Passmore, 2015



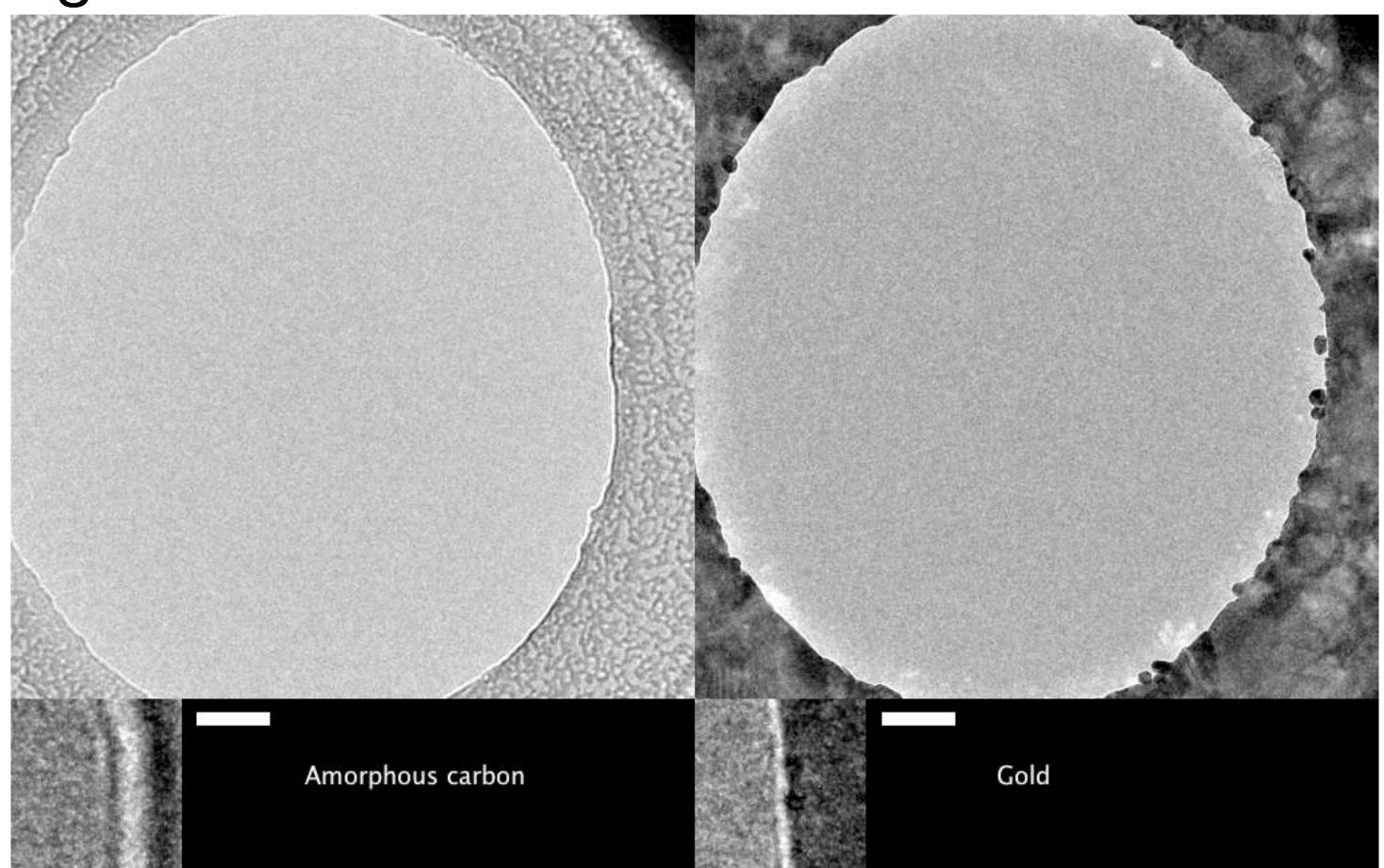
🗱 NYSBC







Gold grids





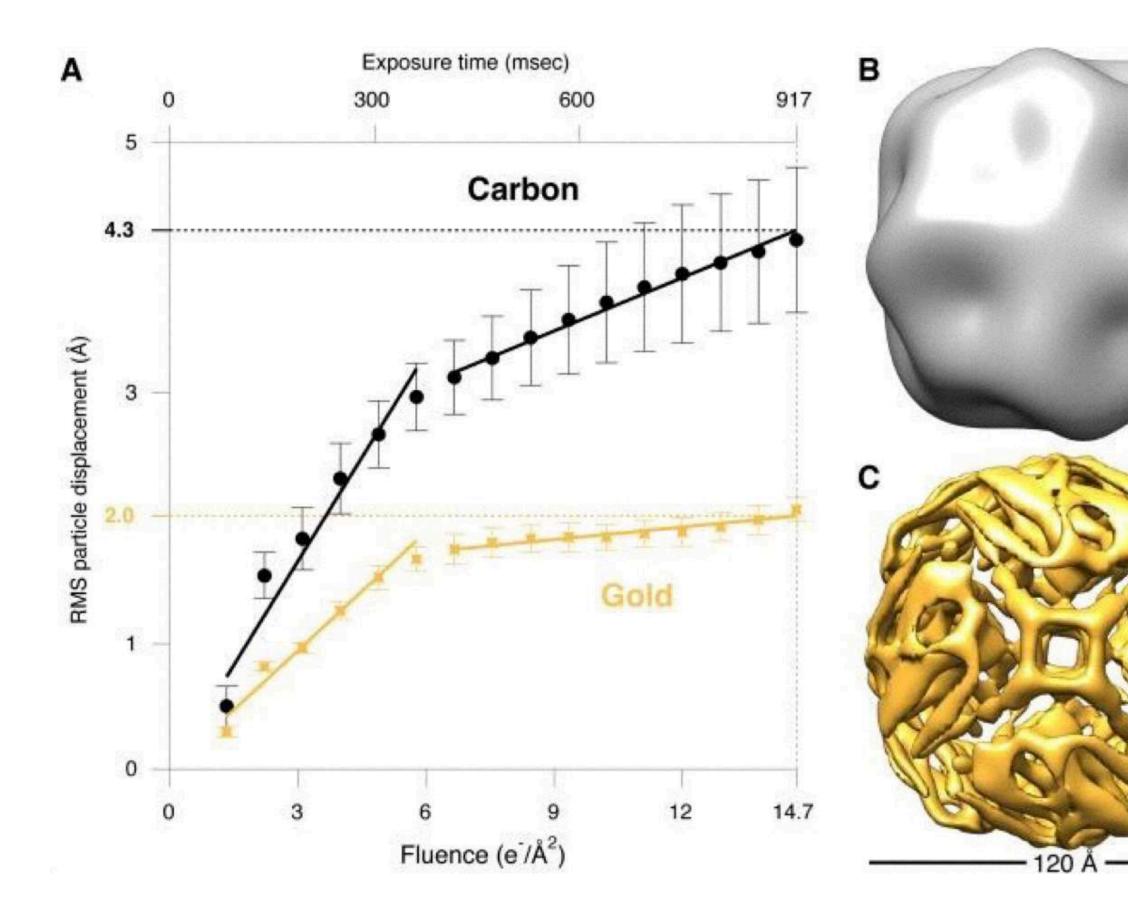
Russo & Passmore, 2015

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22



Gold grids





A. 80S ribosome movement during irradiation supported by amorphous carbon and gold using same imaging conditions.

Apoferritin density maps using same imaging conditions and identical processing for **B**. carbon and **C**. gold substrates. B. is at 25 Å and C. 8 Å resolution.

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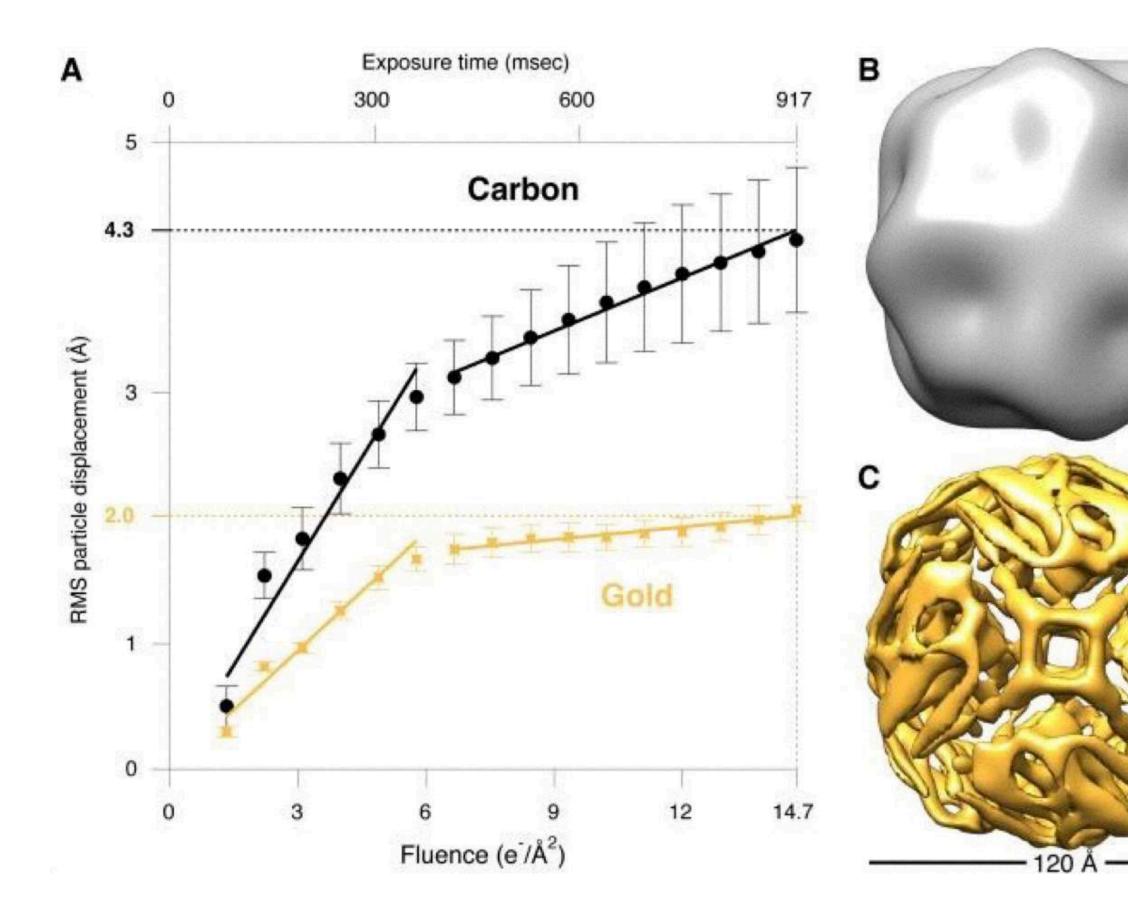
Russo & Passmore, 2015







Gold grids





A. 80S ribosome movement during irradiation supported by amorphous carbon and gold using same imaging conditions.

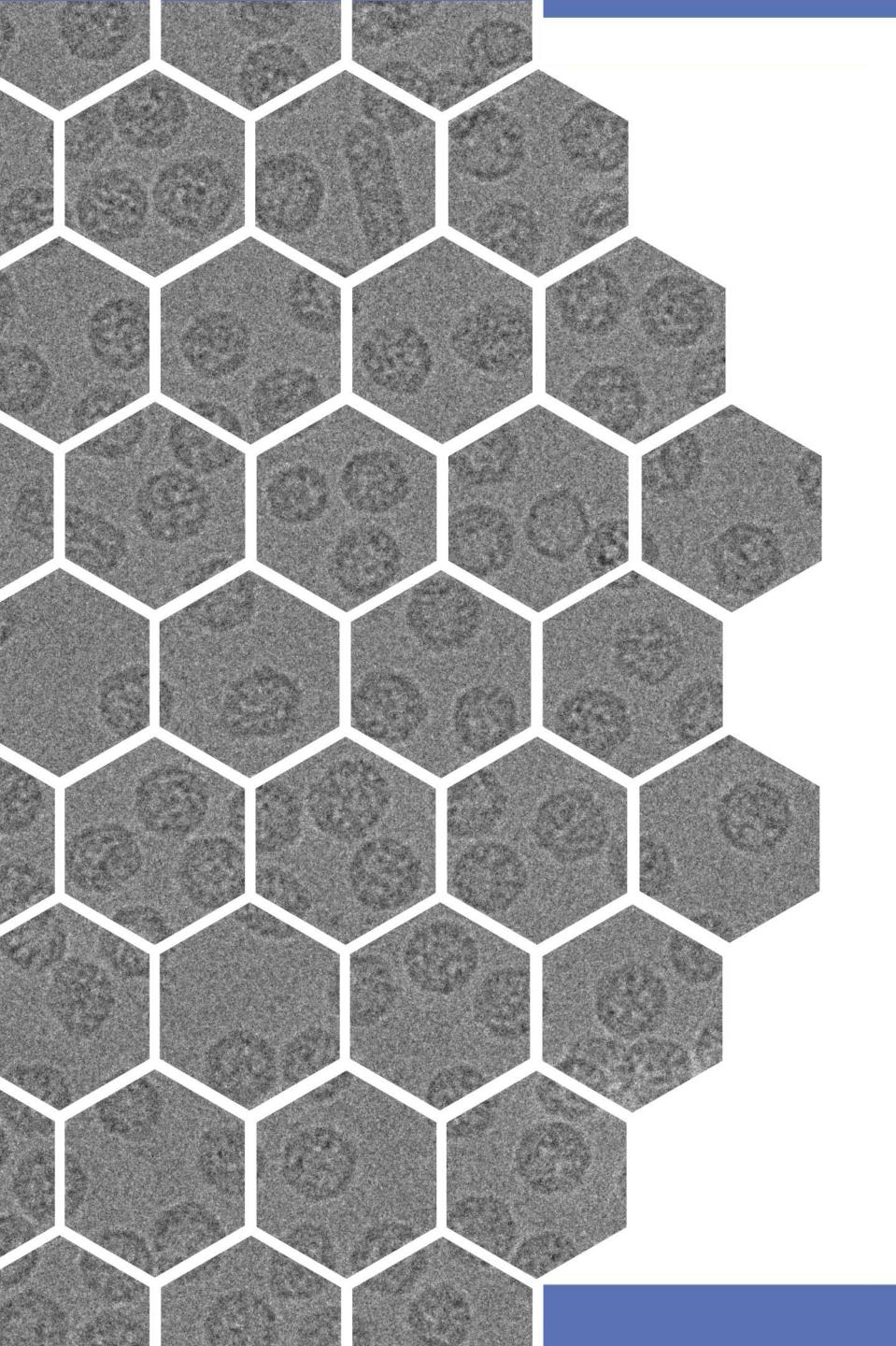
Apoferritin density maps using same imaging conditions and identical processing for **B**. carbon and **C**. gold substrates. B. is at 25 Å and C. 8 Å resolution.

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Russo & Passmore, 2015







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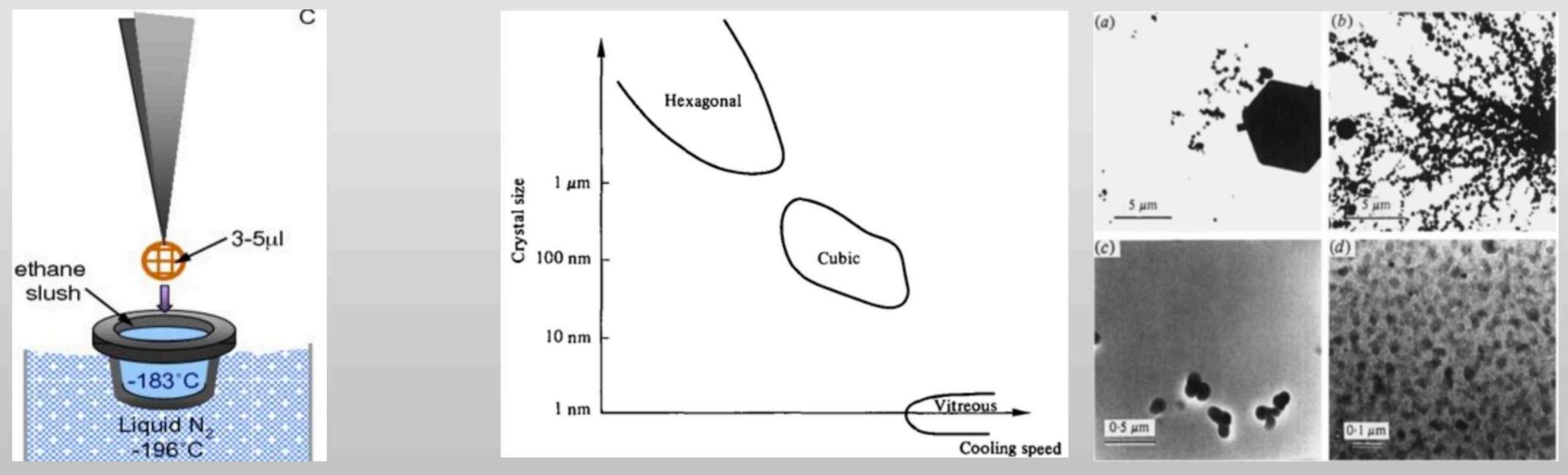






Vitrification process

- Liquid ethane is a suitable coolant.
- Liquid nitrogen boils on contact, which makes it a poor coolant for cryo-EM.
- Cooling speed faster than 10⁵-10⁶ 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

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Jacques Dubochet et al., 1988



How are samples prepared for cryoEM? Vitrification process



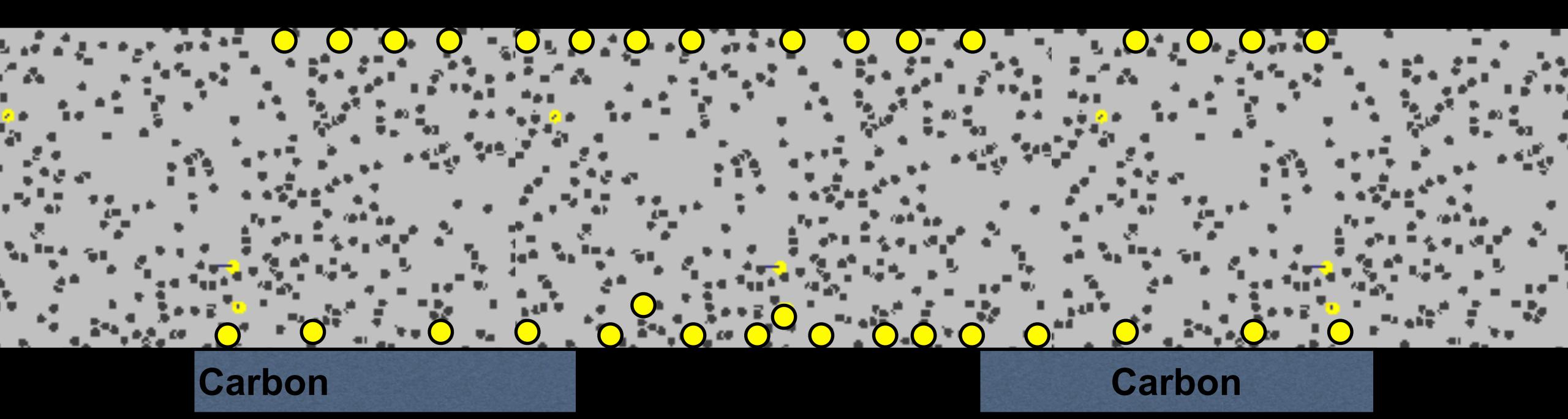
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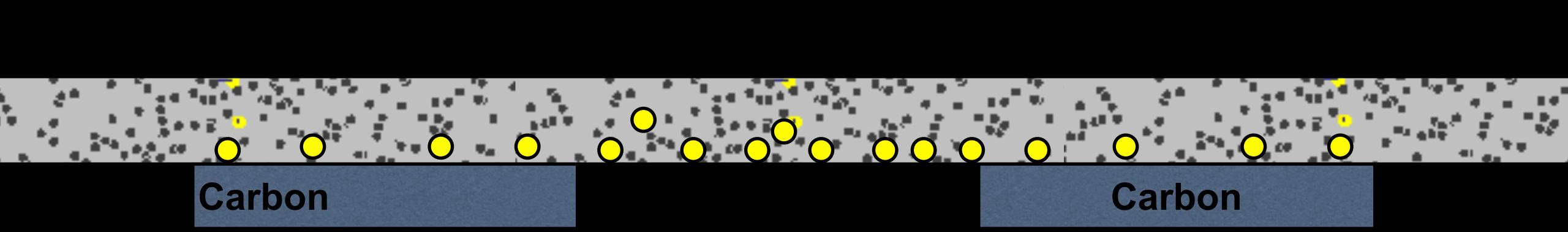


A hypothetical scenario during cryoEM grid preparation





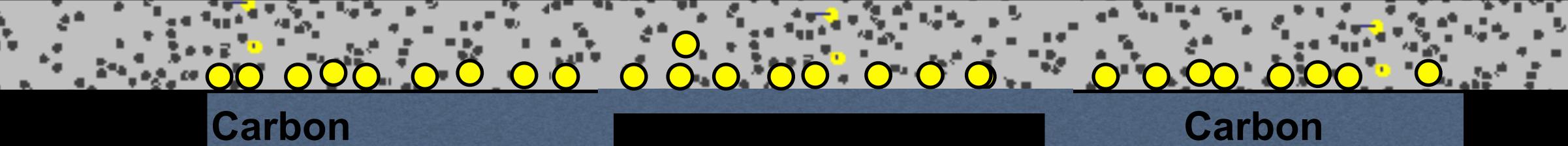
A hypothetical scenario during cryoEM grid preparation



http://weelookang.blogspot.com/2010/06/ejs-open-source-brownian-motion-gas.html

30

A hypothetical scenario during cryoEM grid preparation

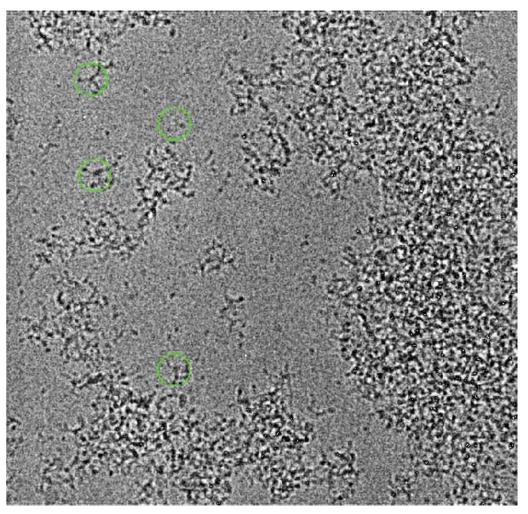


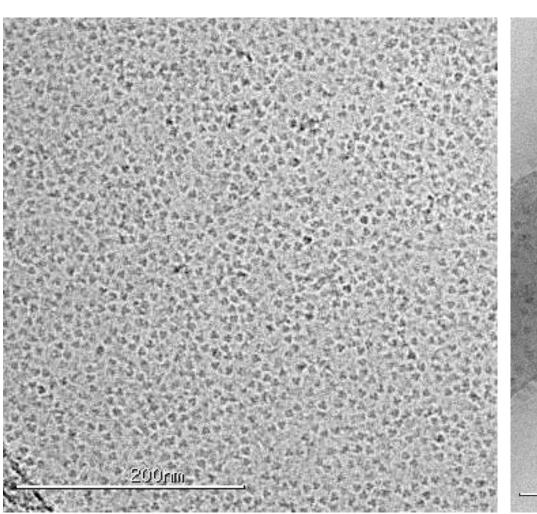
http://weelookang.blogspot.com/2010/06/ejs-open-source-brownian-motion-gas.html



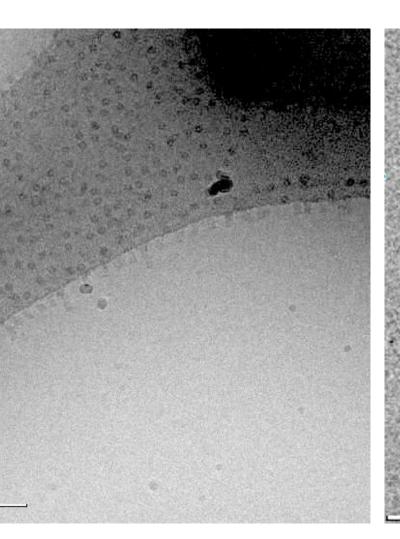


What issues arise?

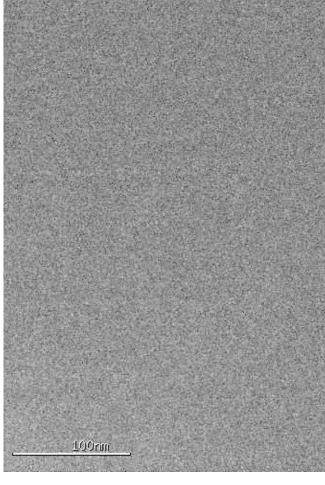




Aggregating in ice Preferred orientation



100mm

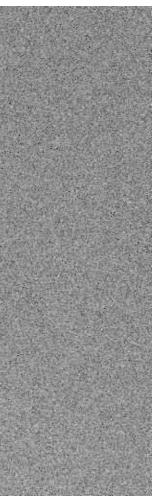


Particles not going into holes

200nm

Rejecting 90% of particles

Particles disappearing in ice

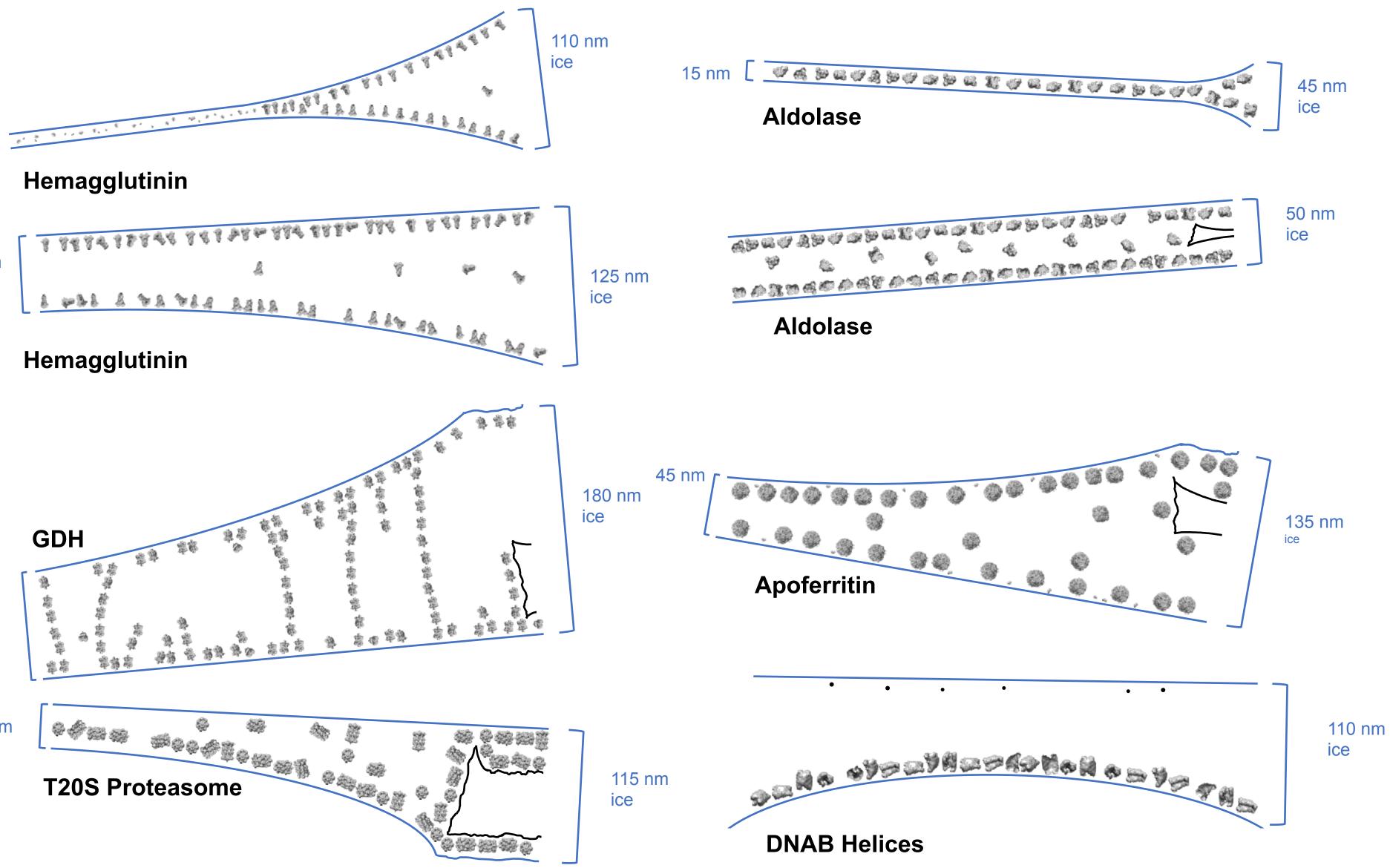


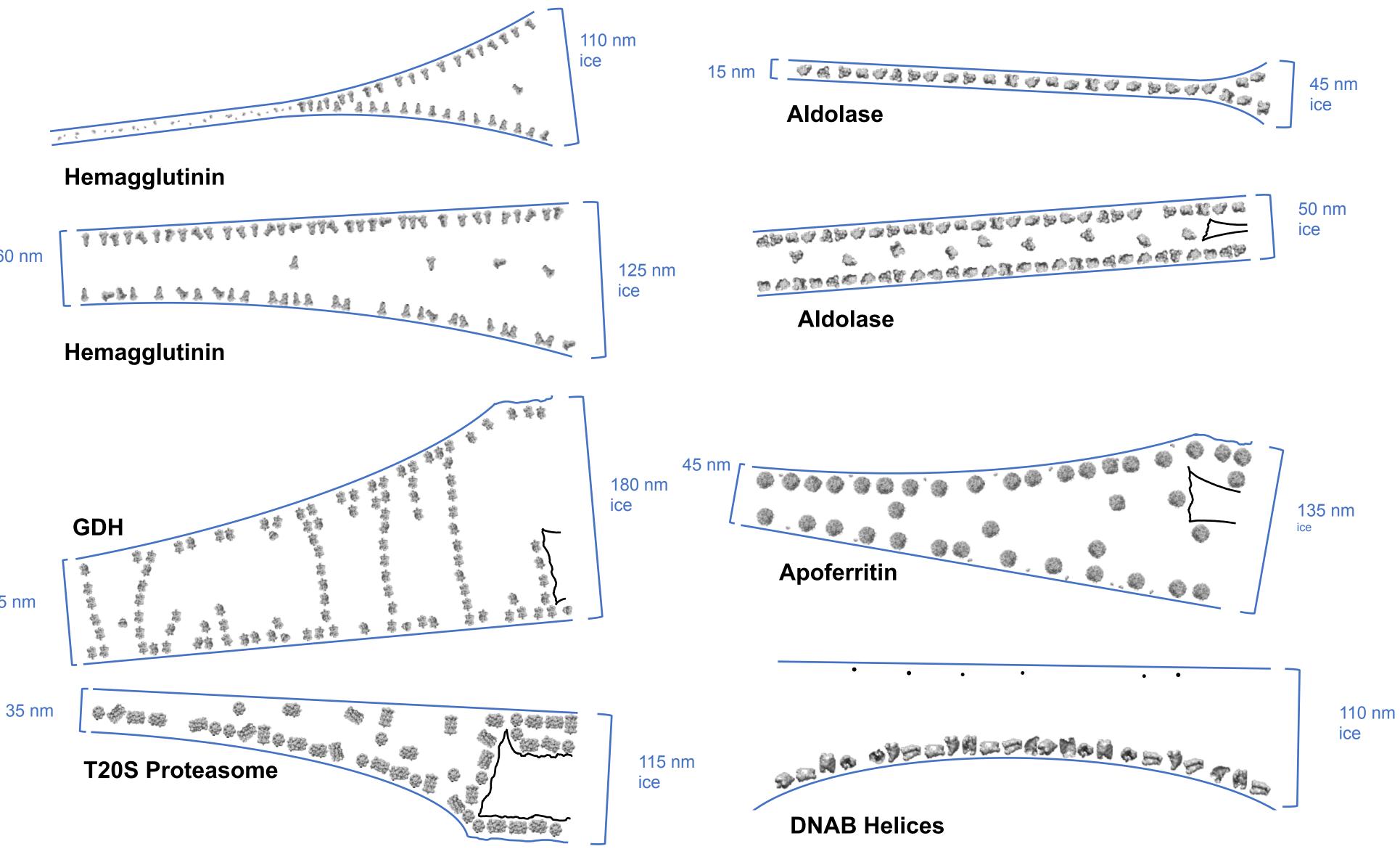






What issues arise?

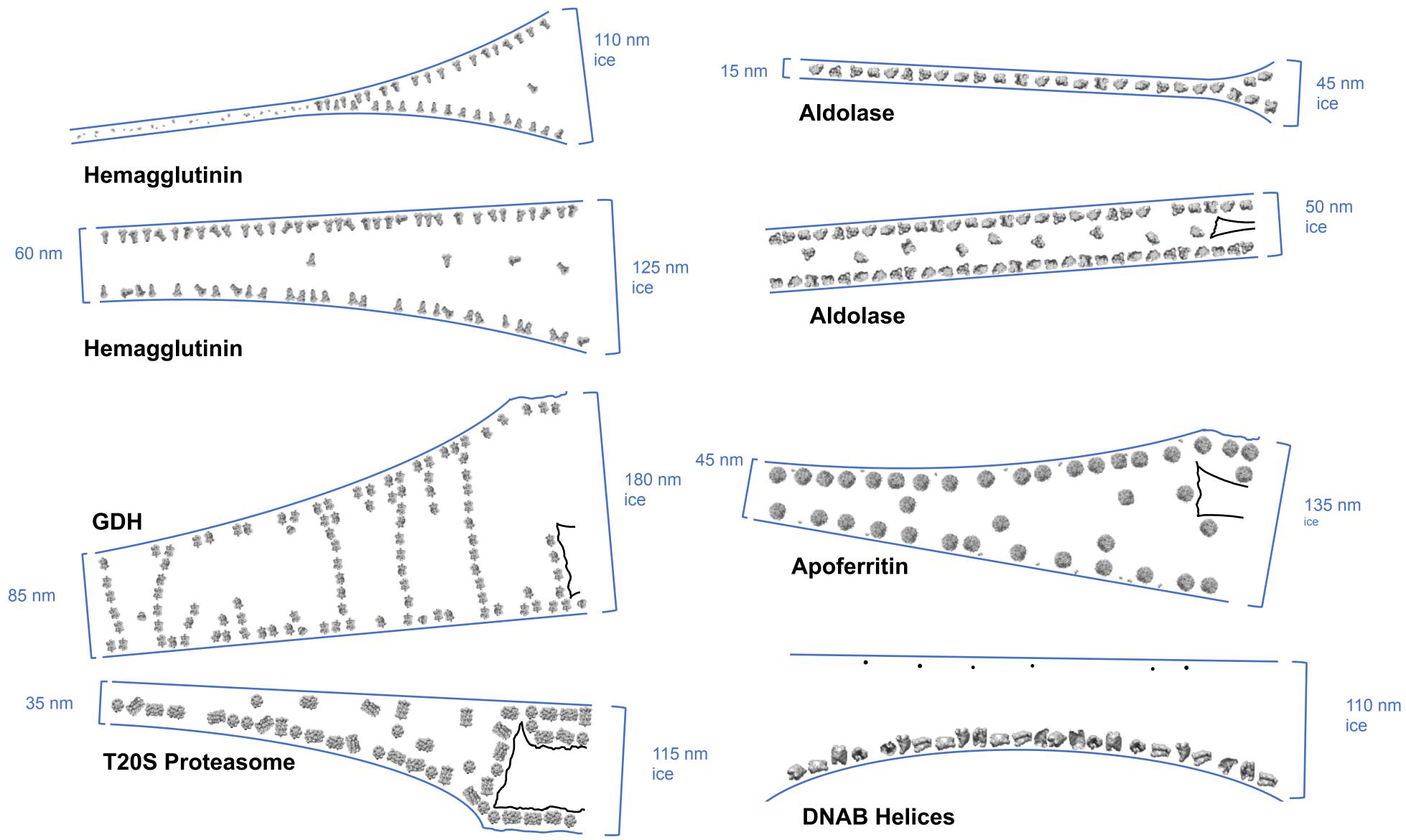




Noble AJ, et al. Routine single particle CryoEM sample and grid characterization by tomography. Elife. 2018;7.



Alex Noble



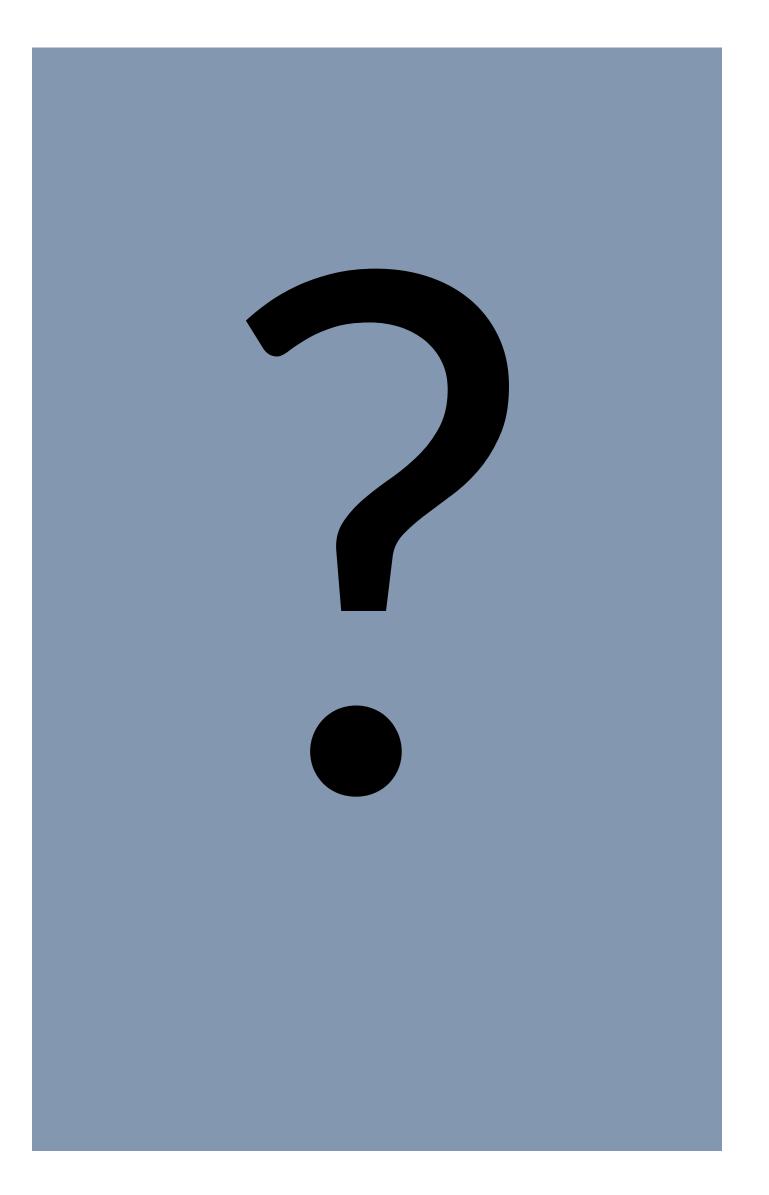
🗱 NYSBC 👘 SEMC







What issues arise?



Small protein

- VPP
- Thinner ice

Protein denaturation/Dissociation of protein complex

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- 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

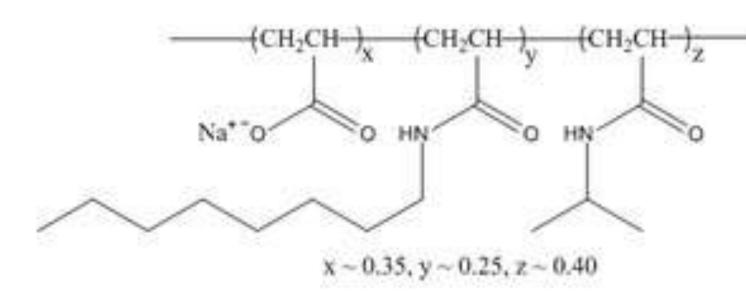
• Segmented analysis

Low concentration

- Multiple blots
- Affinity grids



Reagents for improving vitrification of Cryo-EM grids used in single particle analysis.



Molecular Formula: (C6.2H10.3O1.35N0.65Na0.35)35

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



35



Reagents for improving vitrification of Cryo-EM grids used in single particle analysis.

Surfactants and Cryoprotectants	Amount	Conc.	СМС	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-ß-D-Maltoside (DM)	100 µl	0.87% (w/v)	0.09% (w/v)	non-ionic detergent		
n-Decyl-α-D-Maltoside (DαM)	100 µl	0.46% (w/v)	0.08% (w/v)	non-ionic detergent		
n-Dodecyl-ß-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		
https://www.mitegen.com/product/cryo-em-vitrification-starter-kit/						

• [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 cryoelectron 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.









Reagents for improving vitrification of Cryo-EM grids used in single particle analysis.

PDB Release Date	PDB	Protein	Additive	
2020-01-08	6PWN	MscS mechanosensitive channel	0.01% f-OM	
2019-09-04	6KG7	Piezo2 mechanosensitive channel	0.65 mM f-FC8	
2019-08-28	6QTI	Nicotinamide nucleotide proton channel	0.05% CHAPS	
2019-08-07	6R7L	SecYEG translocon	0.2% f-OM	
2019-02-06	6E0H	TMEM16 scramblase	3 mM f-FC8	
2018-12-19	6N3Q	Sec protein-translocation channel complex	3 mM f-FC8	
2018-11-07	6H3I	Type 9 secretion system translocon	nslocon 1.5 mM f-FC8 or 0.7 mM f-ON	
2018-10-24	6DMR	TRPV5 ion channel	3 mM f-FC8	
2018-10-17	6D3R	CFTR	3 mM f-FC8	
2018-09-26	6HJR	Influenza Hemagglutinin	2% Octyl Glucoside	
2018-08-08	6FOO	Ryanodine receptor 1	0.2% f-OM	
2018-08-01	6CJQ	SthK CNG Potassium channel	3 mM f-FC8	
2018-05-23	5YX9	TRPC6 ion channel	0.5 mM f-OM	
2018-01-31	6C0V	P-Glycoprotein transporter ABCB1	3 mM f-FC8	
2017-12-27	6B5V	TRPV5 ion channel	3 mM f-FC8	
2017-12-13	6BPQ	TRPM8 channel	2% DMSO	

https://www.anatrace.com/Landing/2020/Mar20-Newsletter

Glaeser, RM, et al. (2017) Biophys Rep 3(1), 1-7.

Noble, AJ, et al. (2018) Nat Methods 15(10), 793-795.

Drulyte, I et al. (2018) Acta Crystallogr D Struct Biol 74(Pt 6), 560-571.

Chen, J, et al. (2019) J Struct Biol X Volume 1. DOI: 10.1016/ j.yjsbx.2019.100005

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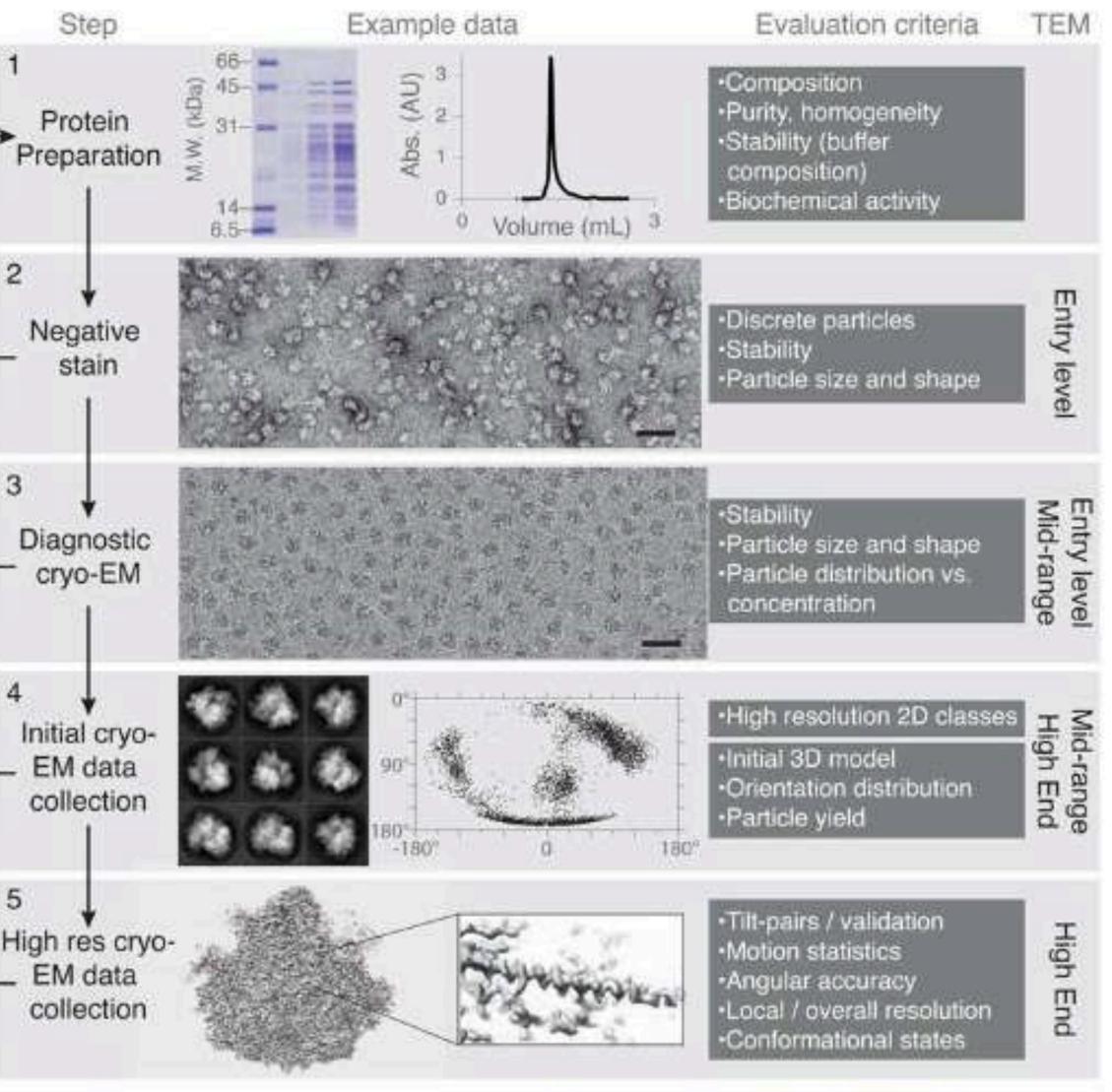


Preparing EM ready samples

THE OPTIMIZATION WORKFLOW

Structure determination by cryo-EM.

A systematic approach to 3D structure determination is shown. In the left column, the major steps are listed. Each step should be performed successively and only after one has been completed successfully should the scientist move onto the next step. In the second column, example data are shown for ribosomes (details in text). Scale bars on the micrographs are 500 Å. Each step should be evaluated with the criteria listed in the third column, returning to earlier steps for troubleshooting.



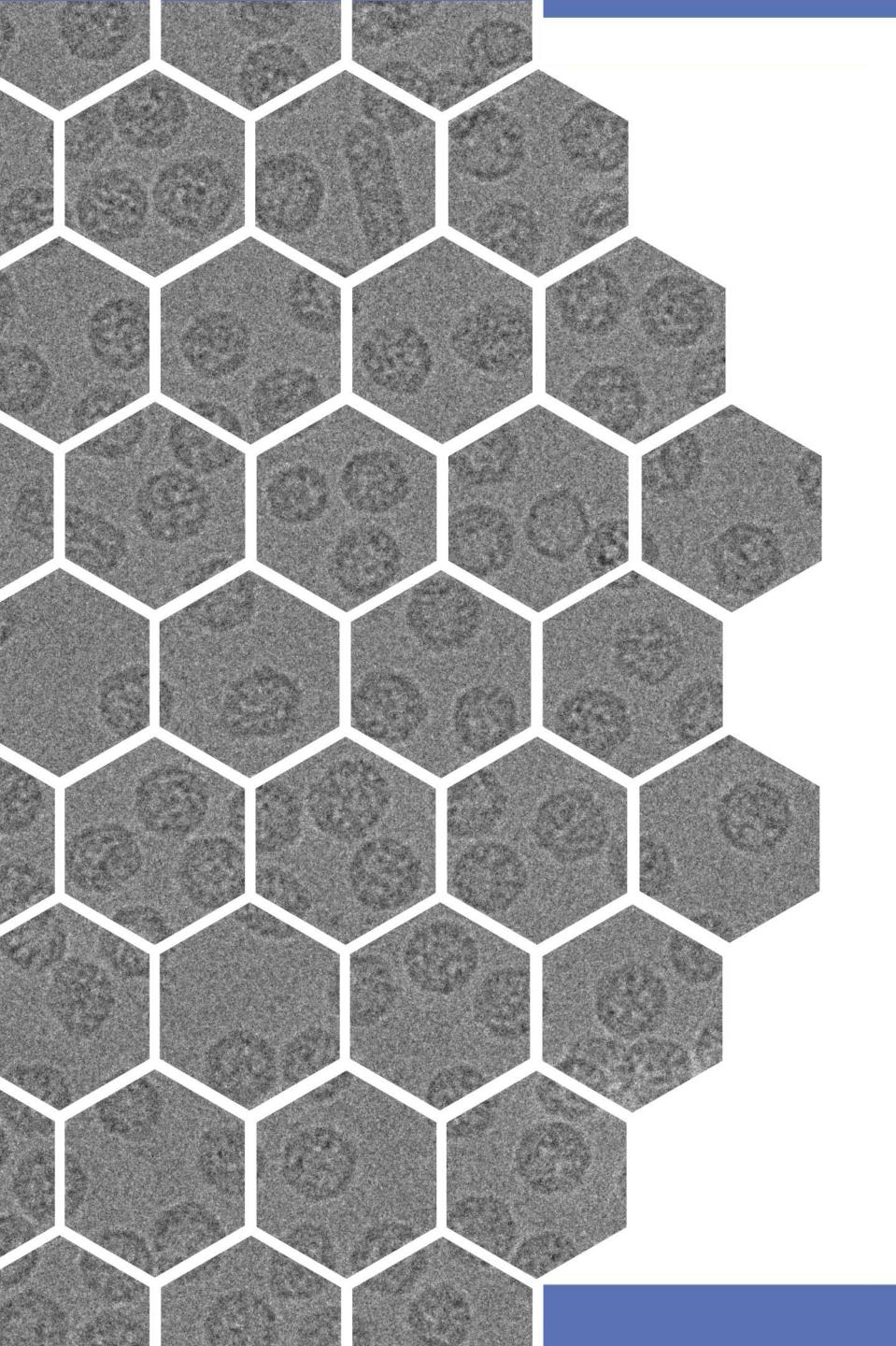
https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5140023/

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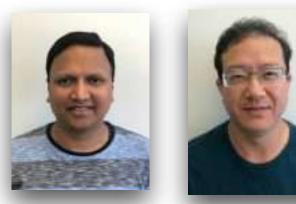






Improving Current CryoTEM Grid Preparation Methods

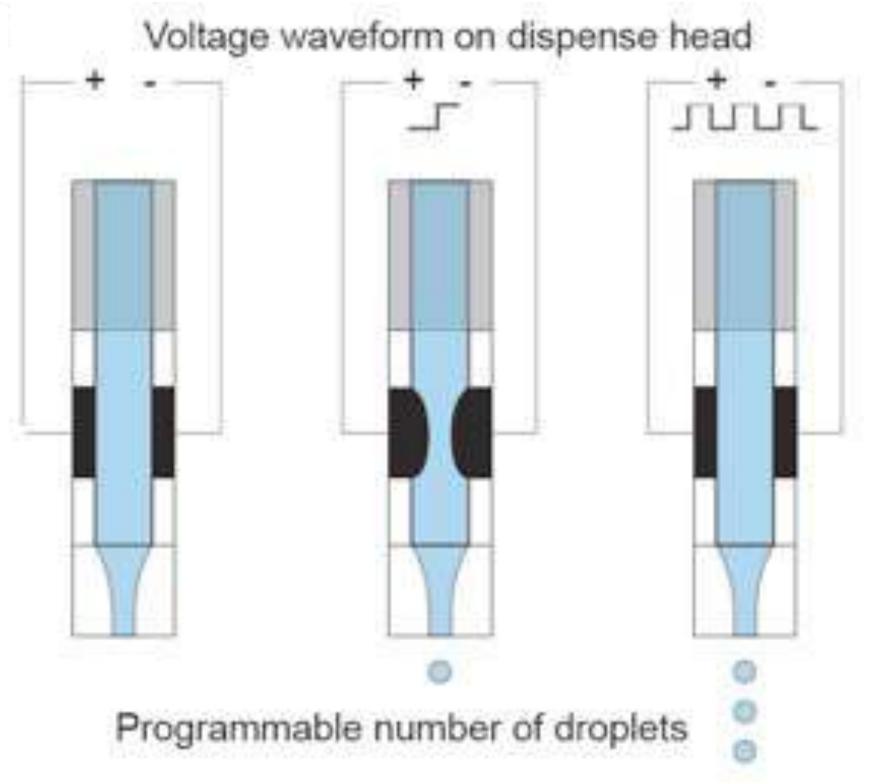
Dandey VP, Wei H, Zhang Z, Tan YZ, Acharya P, Eng ET, Rice WJ, Kahn PA, Potter CS, Carragher B. Spotiton: New features and applications. Journal of structural biology. 2018;202(2):161-9



Venkat Dandey

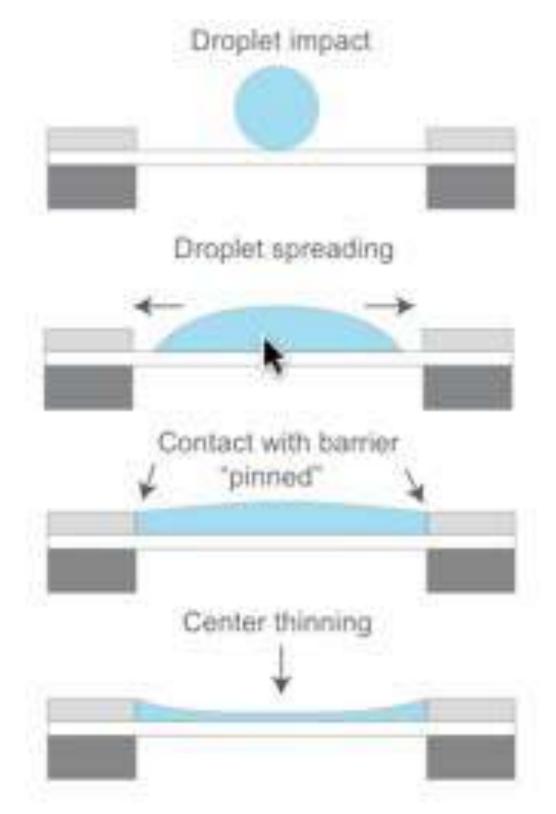
Hui Wei

Accurate pL dispensing





Thin films without blotting



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40

Improving Current CryoTEM Grid Preparation Methods

Dandey VP, Wei H, Zhang Z, Tan YZ, Acharya P, Eng ET, Rice WJ, Kahn PA, Potter CS, Carragher B. Spotiton: New features and applications. Journal of structural biology. 2018;202(2):161-9



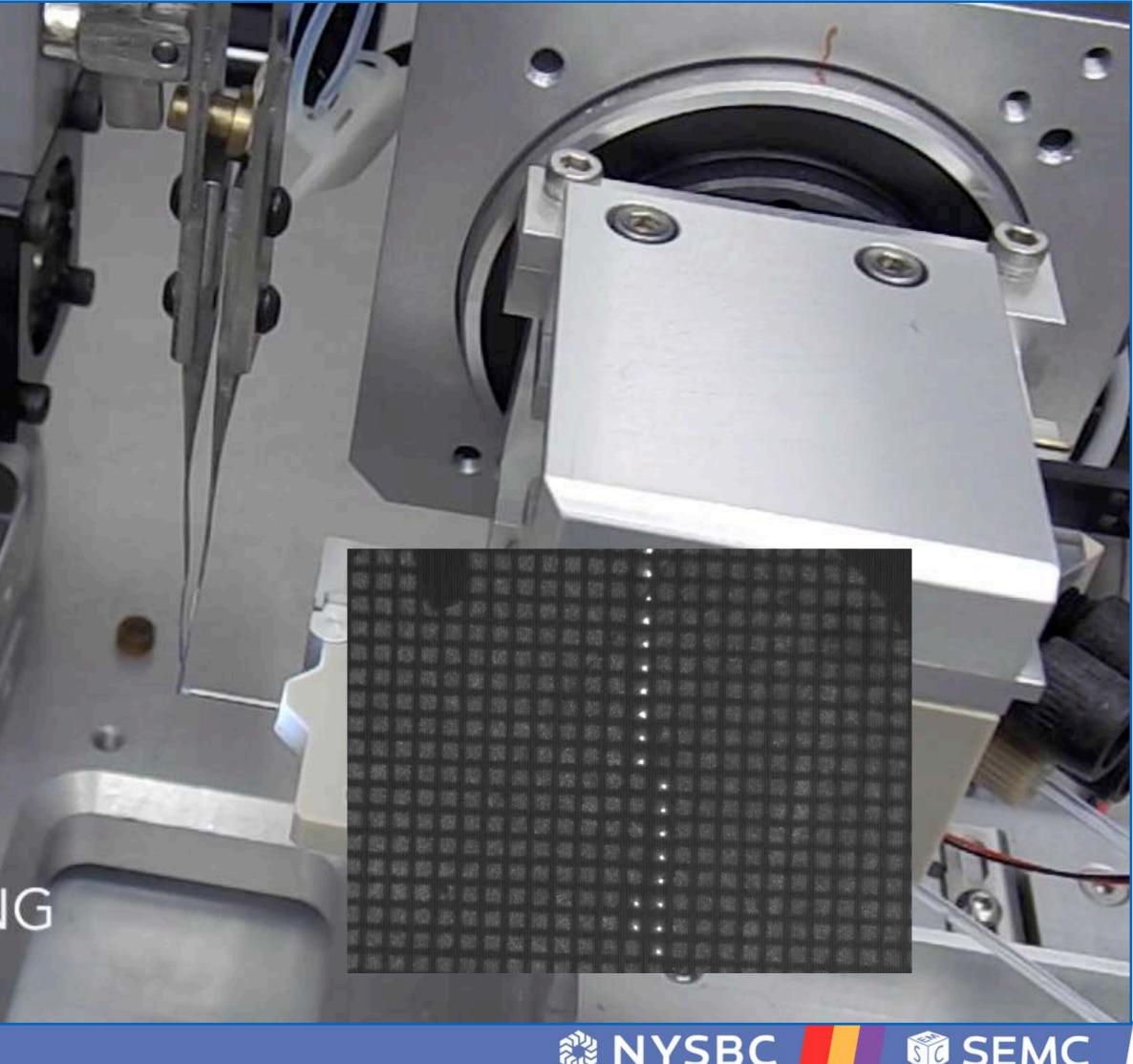


Venkat Dandey

Hui Wei







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Improving Current CryoTEM Grid Preparation Methods

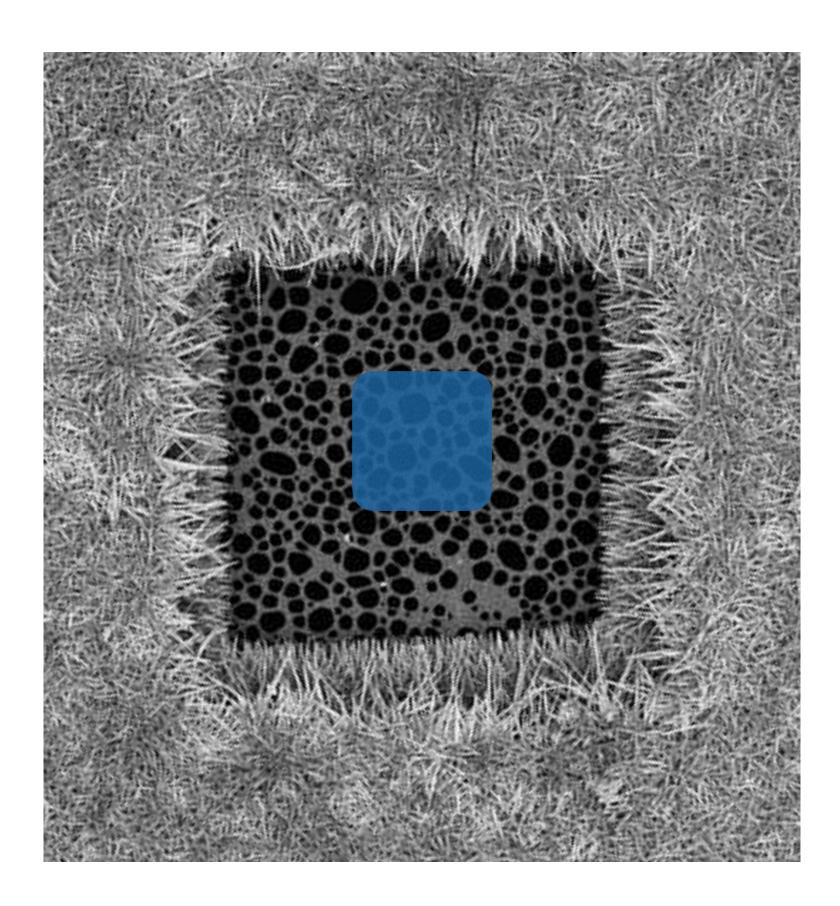
Wei H, Dandey VP, Zhang Z, Raczkowski A, Rice WJ, Carragher B, Potter CS. Optimizing "selfwicking" nanowire grids. J Struct Biol. 2018;202(2):170-4.



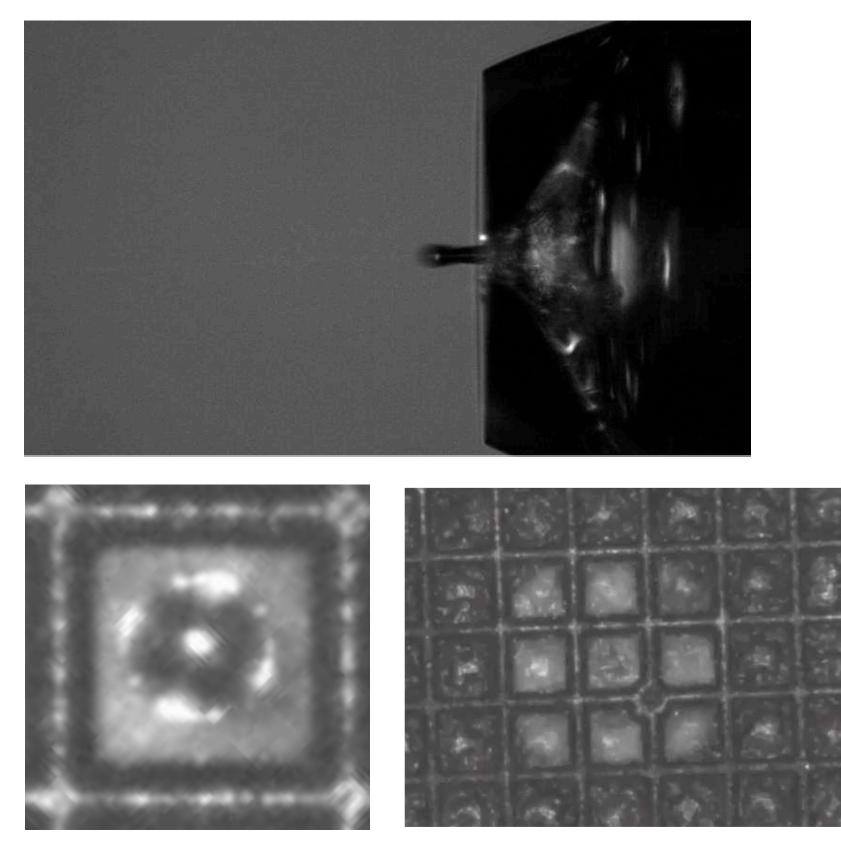


Venkat Dandey

Hui Wei







Single frame from loop

Video loop

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SEMC 42

Improving Current CryoTEM Grid Preparation Methods

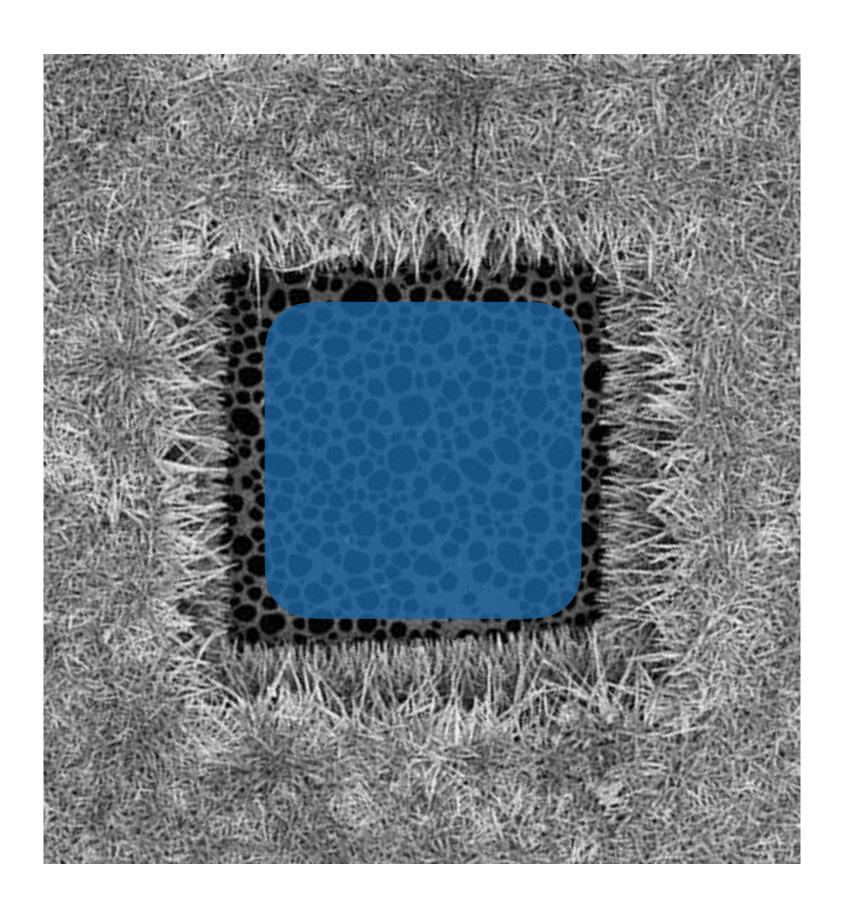
Wei H, Dandey VP, Zhang Z, Raczkowski A, Rice WJ, Carragher B, Potter CS. Optimizing "selfwicking" nanowire grids. J Struct Biol. 2018;202(2):170-4.



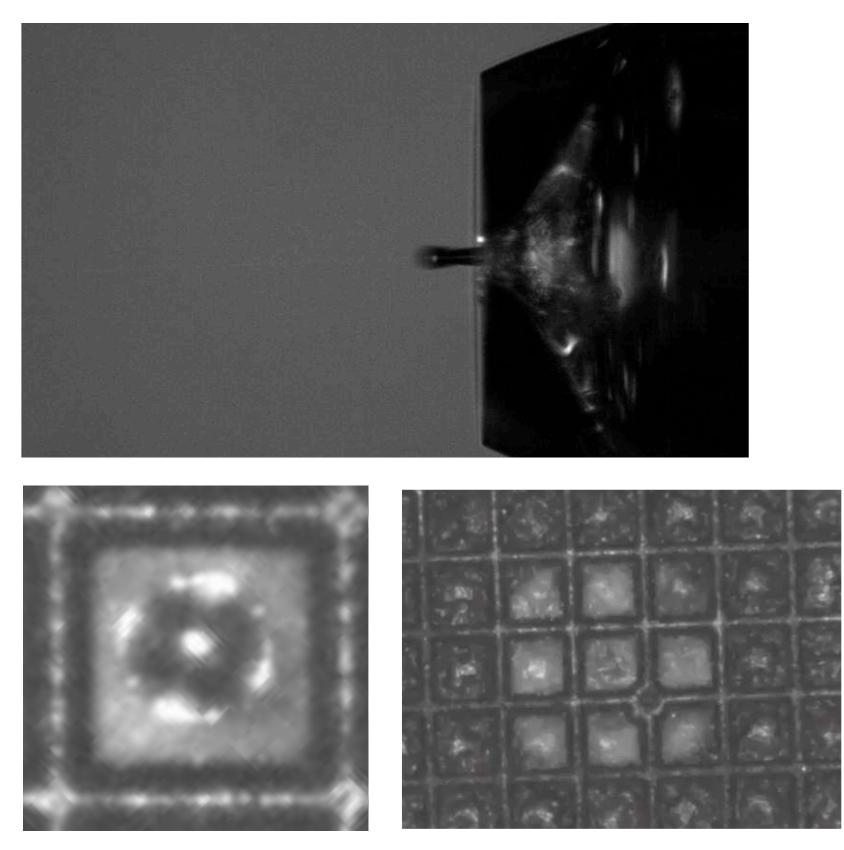


Venkat Dandey

Hui Wei







Single frame from loop

Video loop



43

Improving Current CryoTEM Grid Preparation Methods

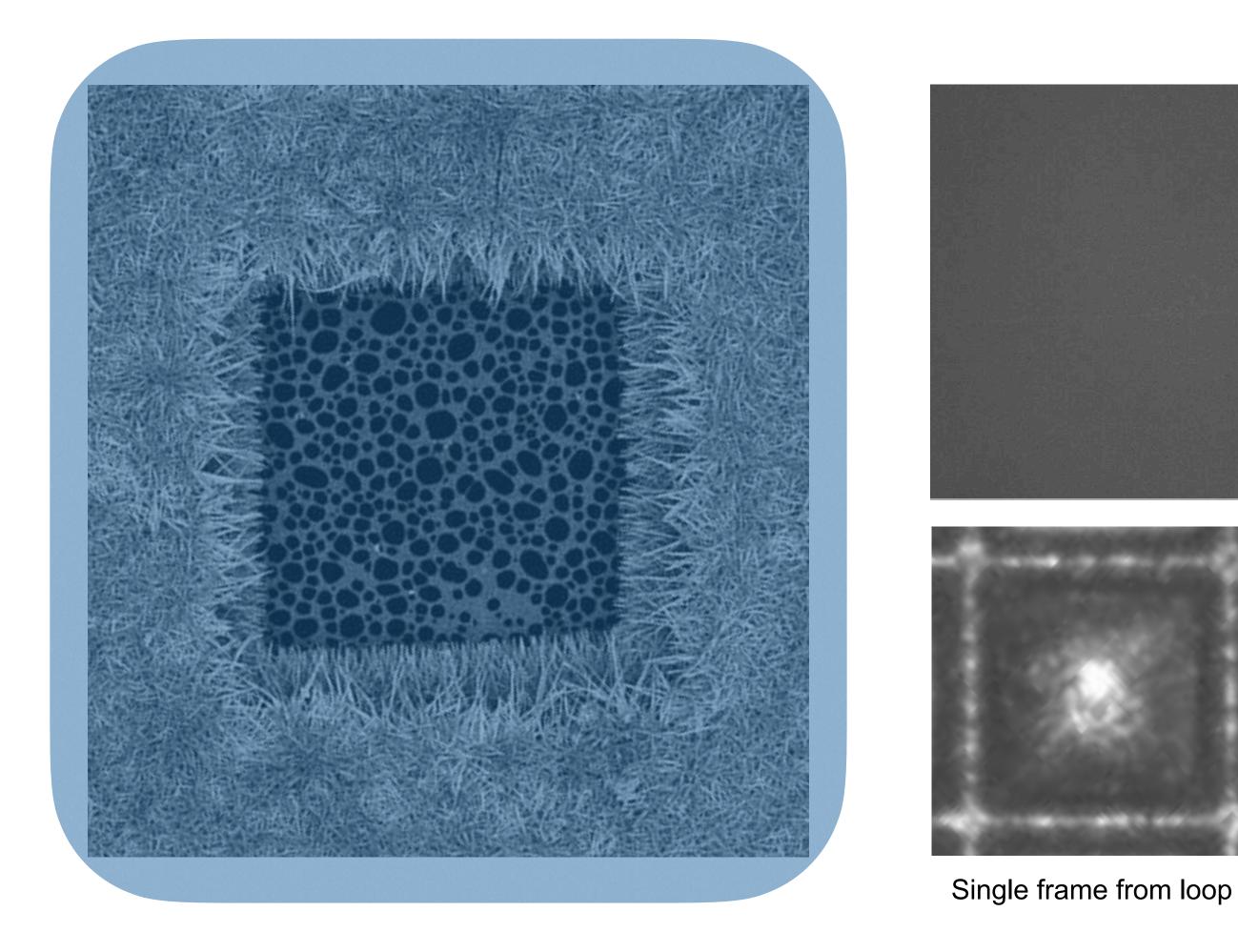
Wei H, Dandey VP, Zhang Z, Raczkowski A, Rice WJ, Carragher B, Potter CS. Optimizing "selfwicking" nanowire grids. J Struct Biol. 2018;202(2):170-4.



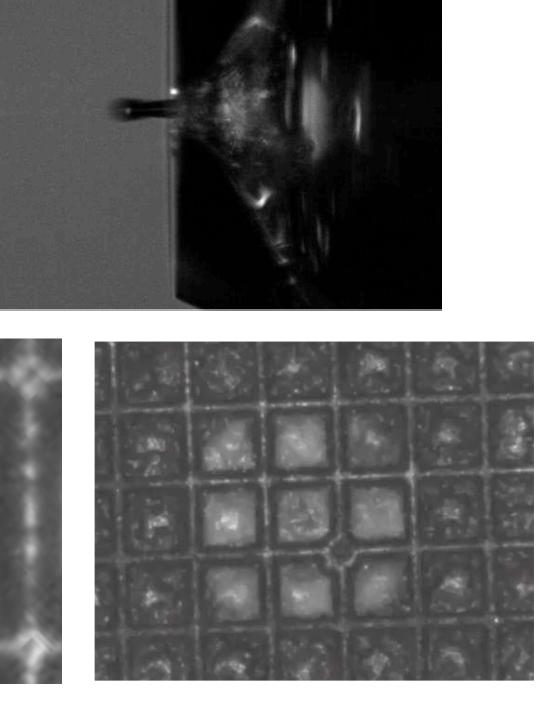


Venkat Dandey

Hui Wei







Video loop

NYSBC SEMC

44

Improving Current CryoTEM Grid Preparation Methods

Wei H, Dandey VP, Zhang Z, Raczkowski A, Rice WJ, Carragher B, Potter CS. Optimizing "selfwicking" nanowire grids. J Struct Biol. 2018;202(2):170-4.



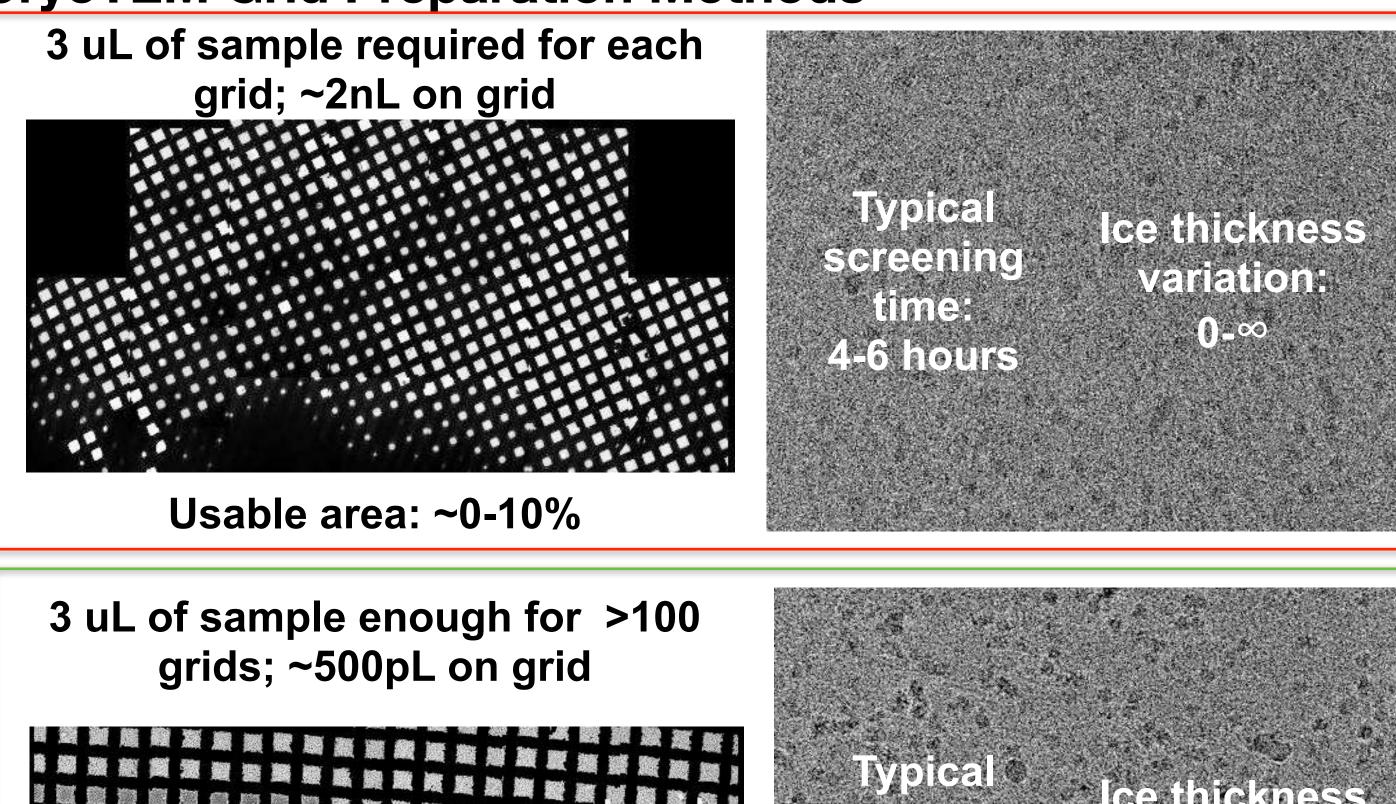


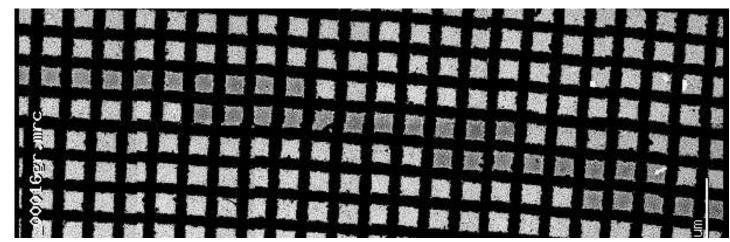
Venkat Dandey

Hui Wei

Vitrobot

Spotiton





Usable area: ~100%



screening time: 10 minutes

Ice thickness variation: $\sim 2X$



SEMC 45

What is chameleon?

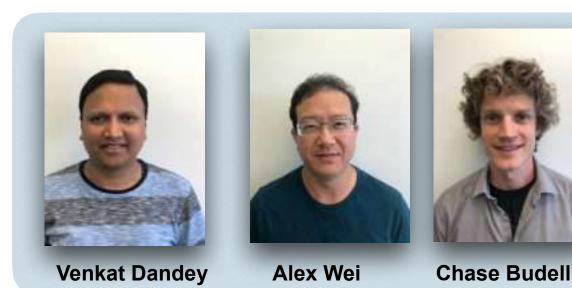
The Spotiton Project: Commercialization Chameleon: 2019 **Spotiton concept: 2011**







sptlabtech





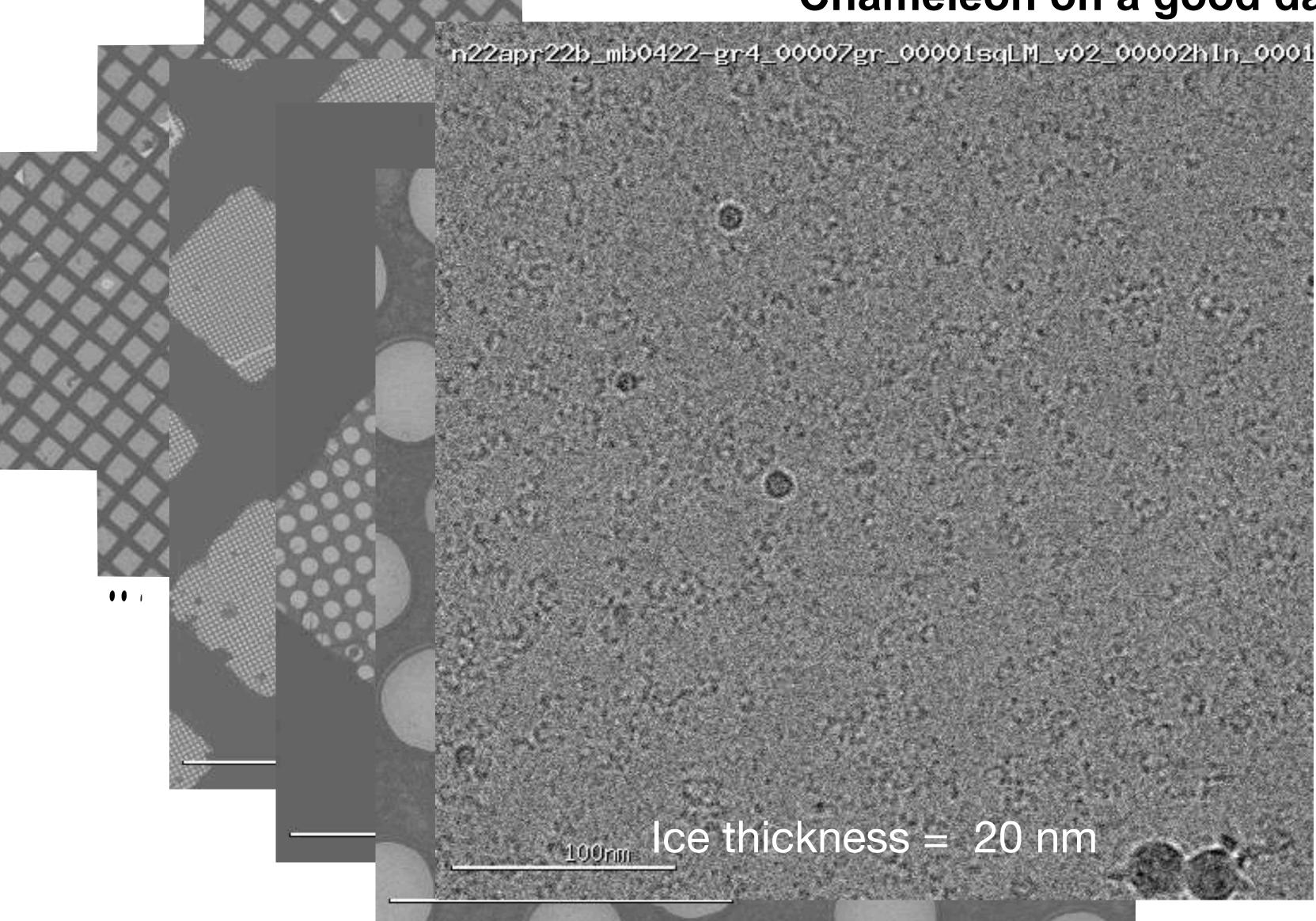
iii ttplabtech

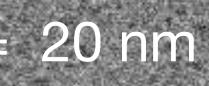






What is chameleon? Chameleon on a good day











Mahira Aragon

Alex Wei

Chase Budell



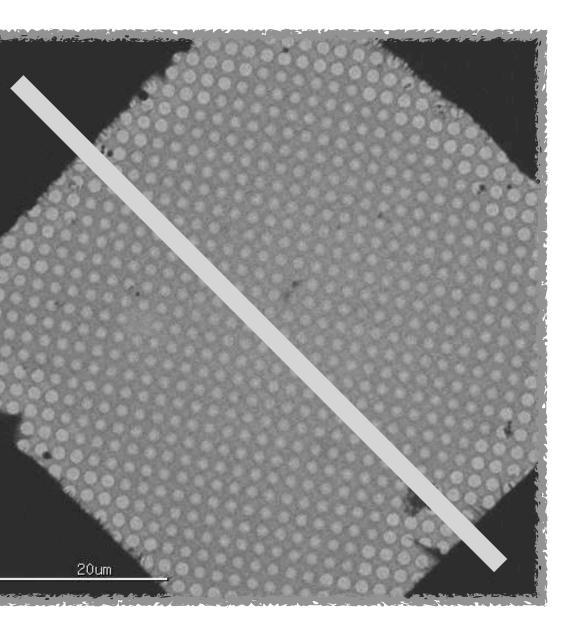


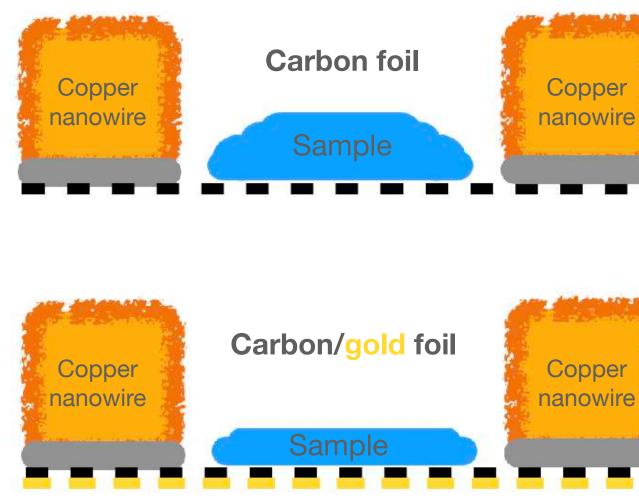


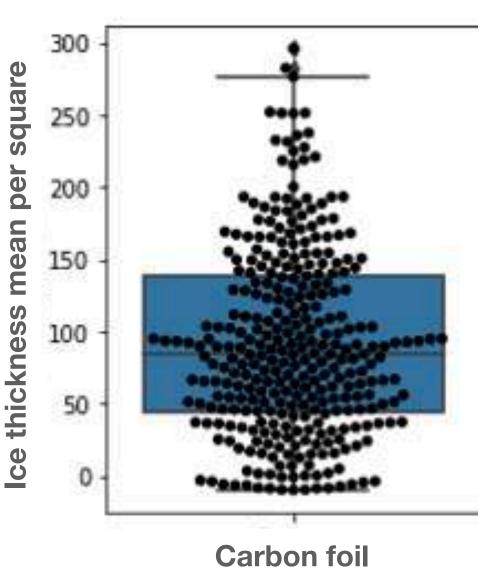


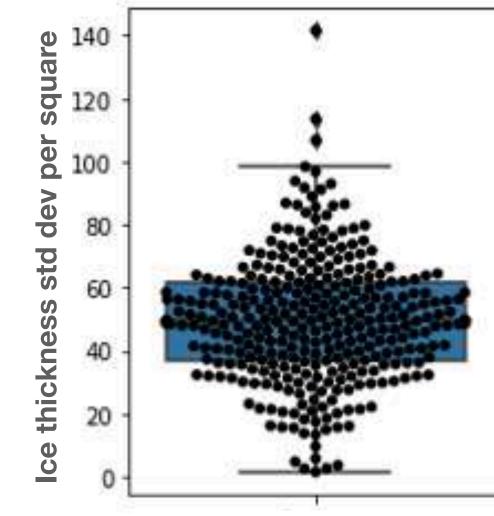


What is chameleon?



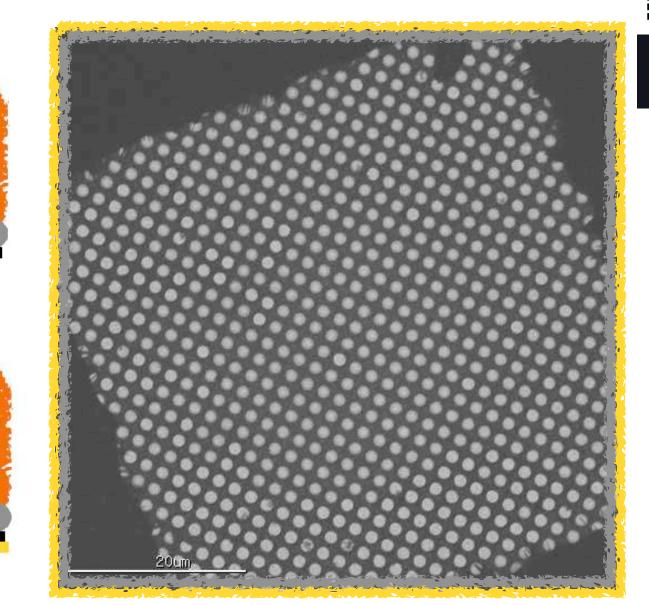




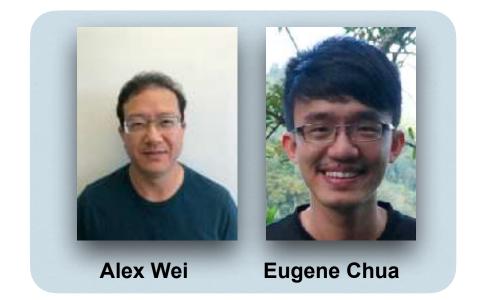


Carbon foil









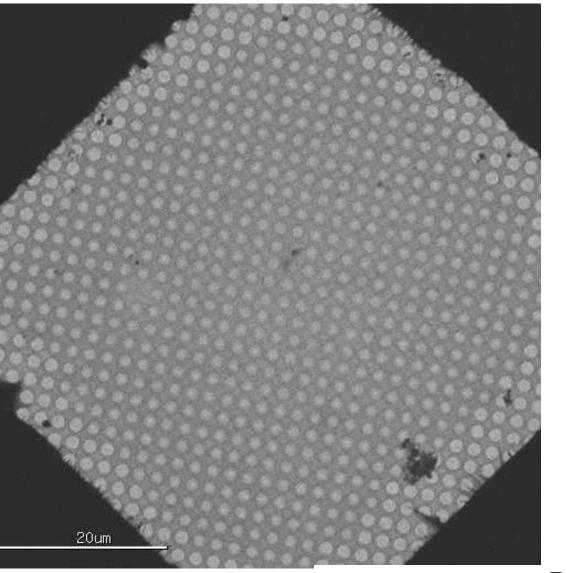
🗱 NYSBC

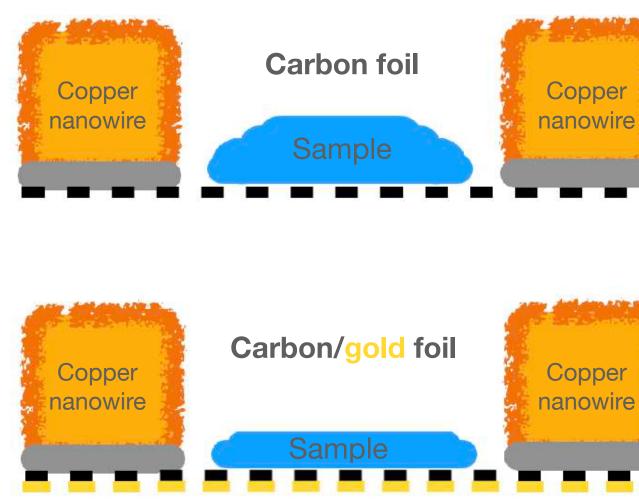






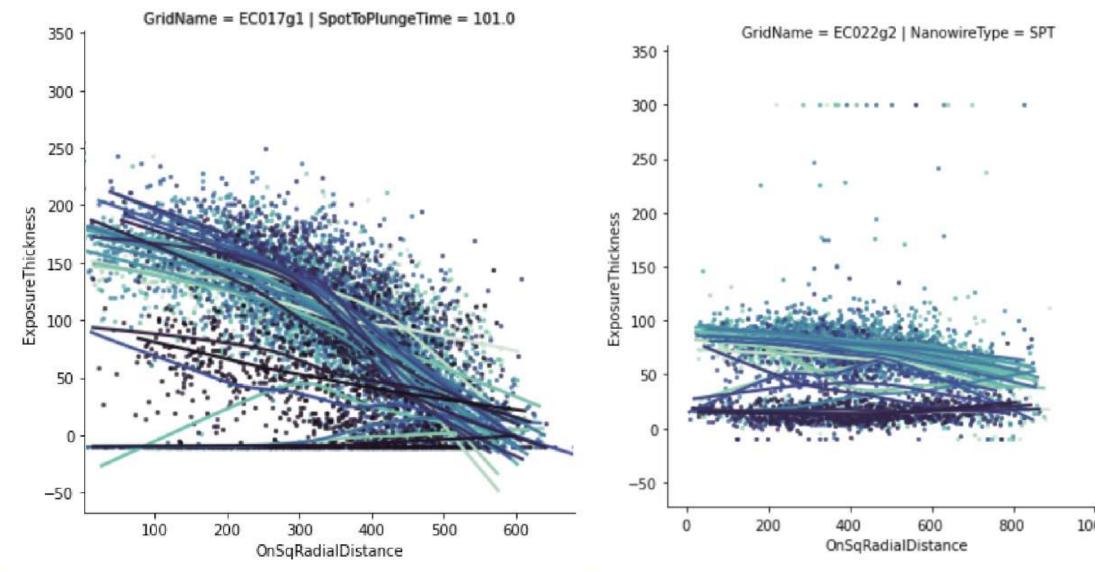
What is chameleon?



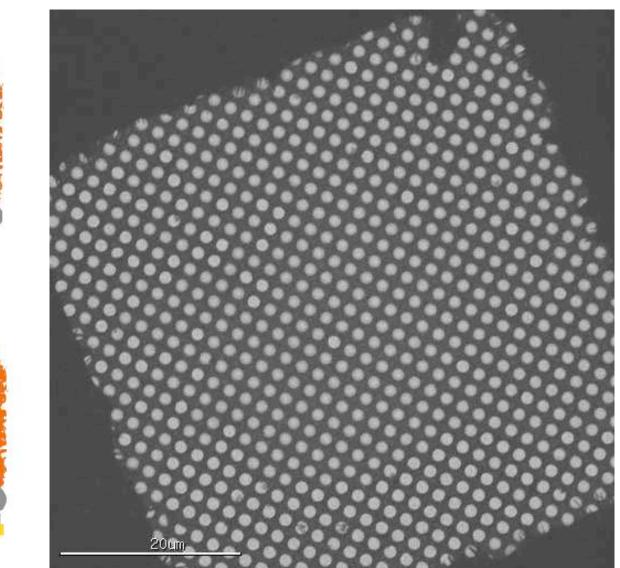


Batch 413







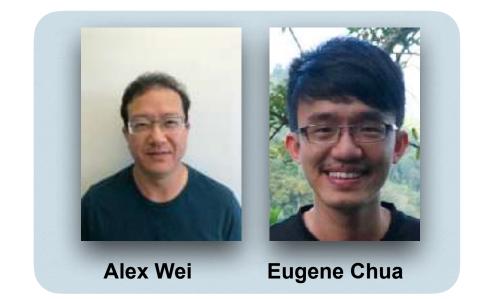




1000







🗞 NYSBC







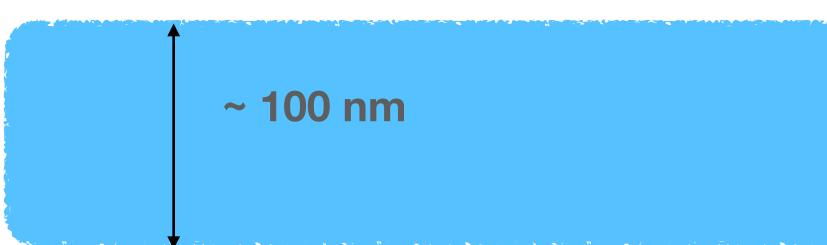


Grid geometry



Vitreous ice ideal -> typical thickness







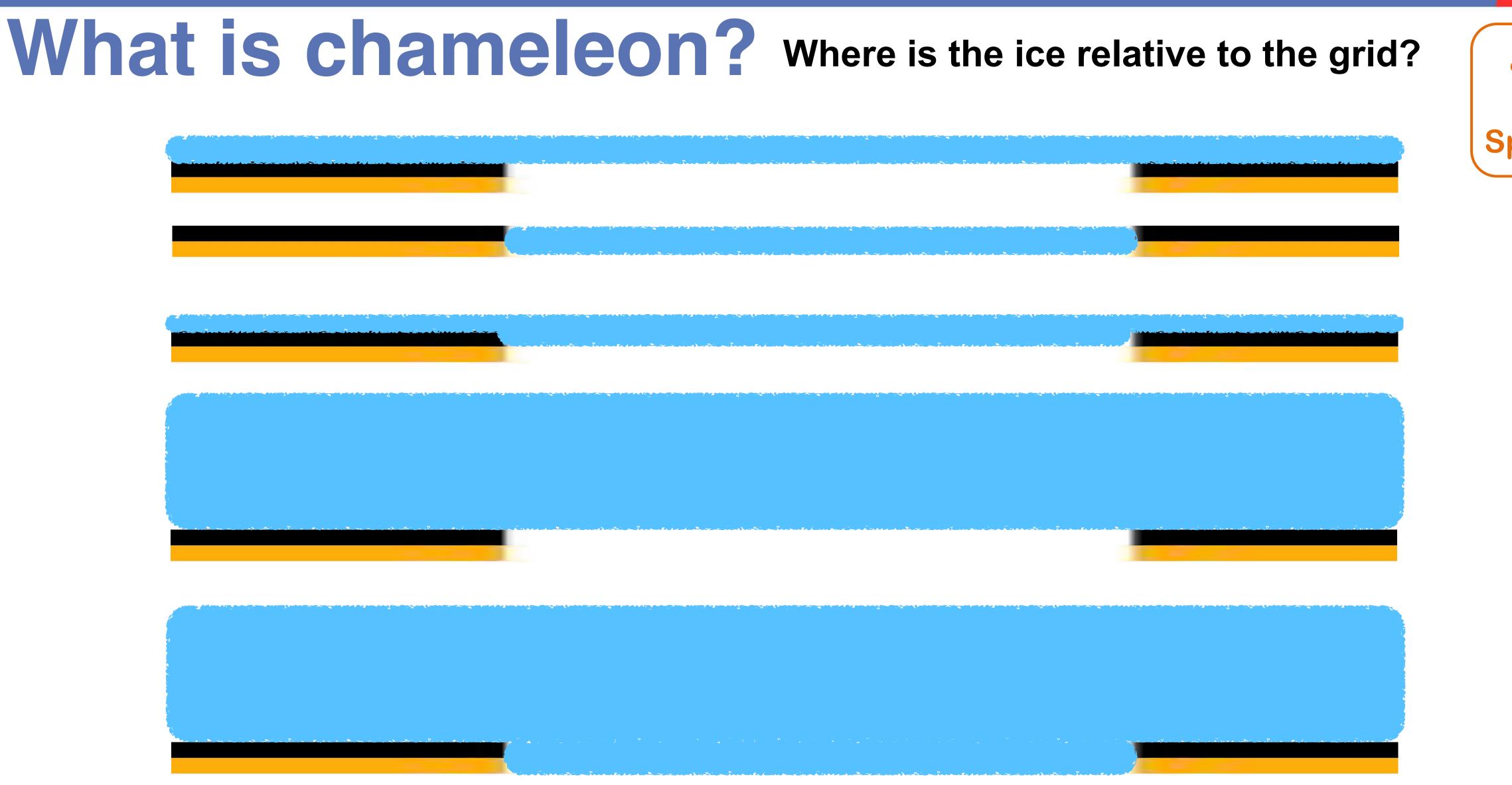
~1000 nm

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50



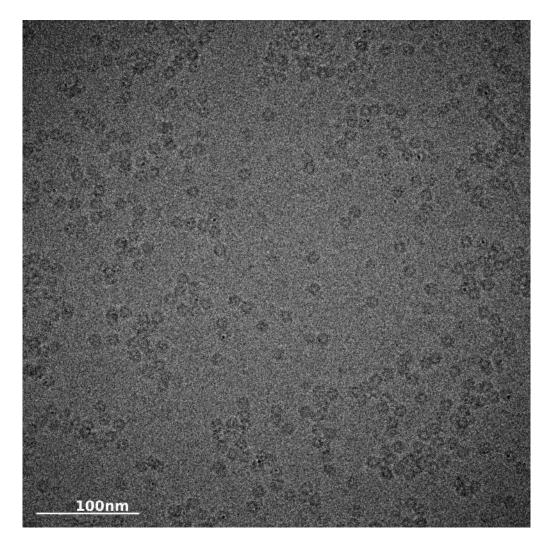


Does the substate determine this?

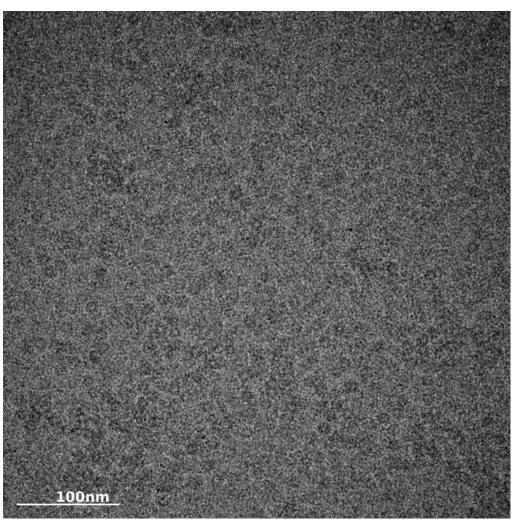




51



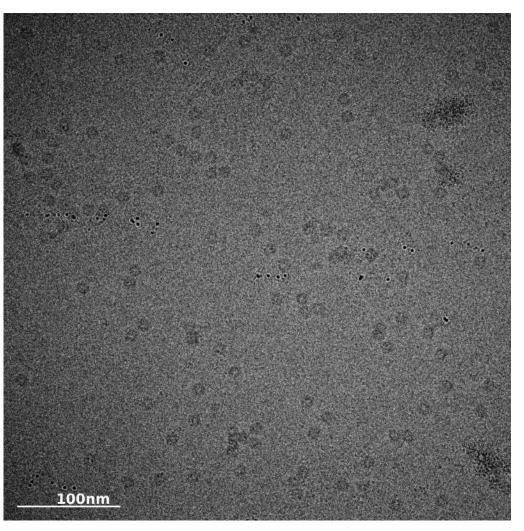
No graphene (8 mg/ml)



Graphene (8mg/ml)



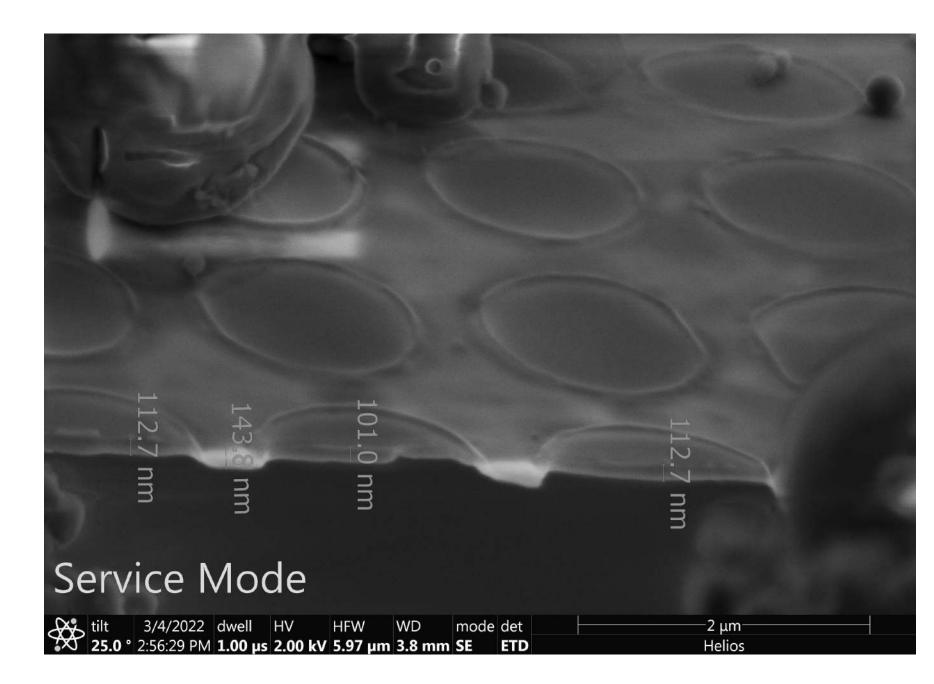
Add a graphene layer under the substrate



Graphene (0.8 mg/ml)

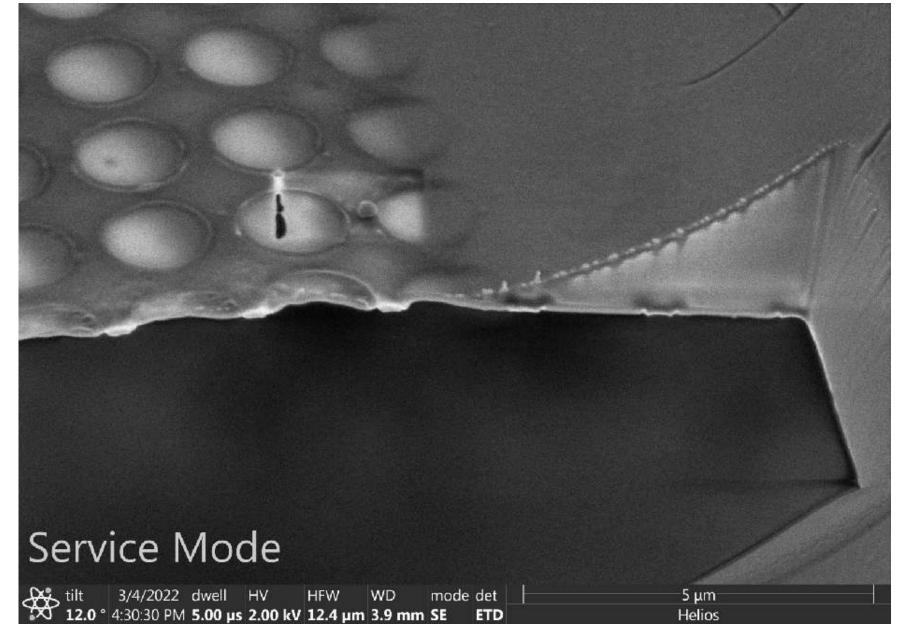






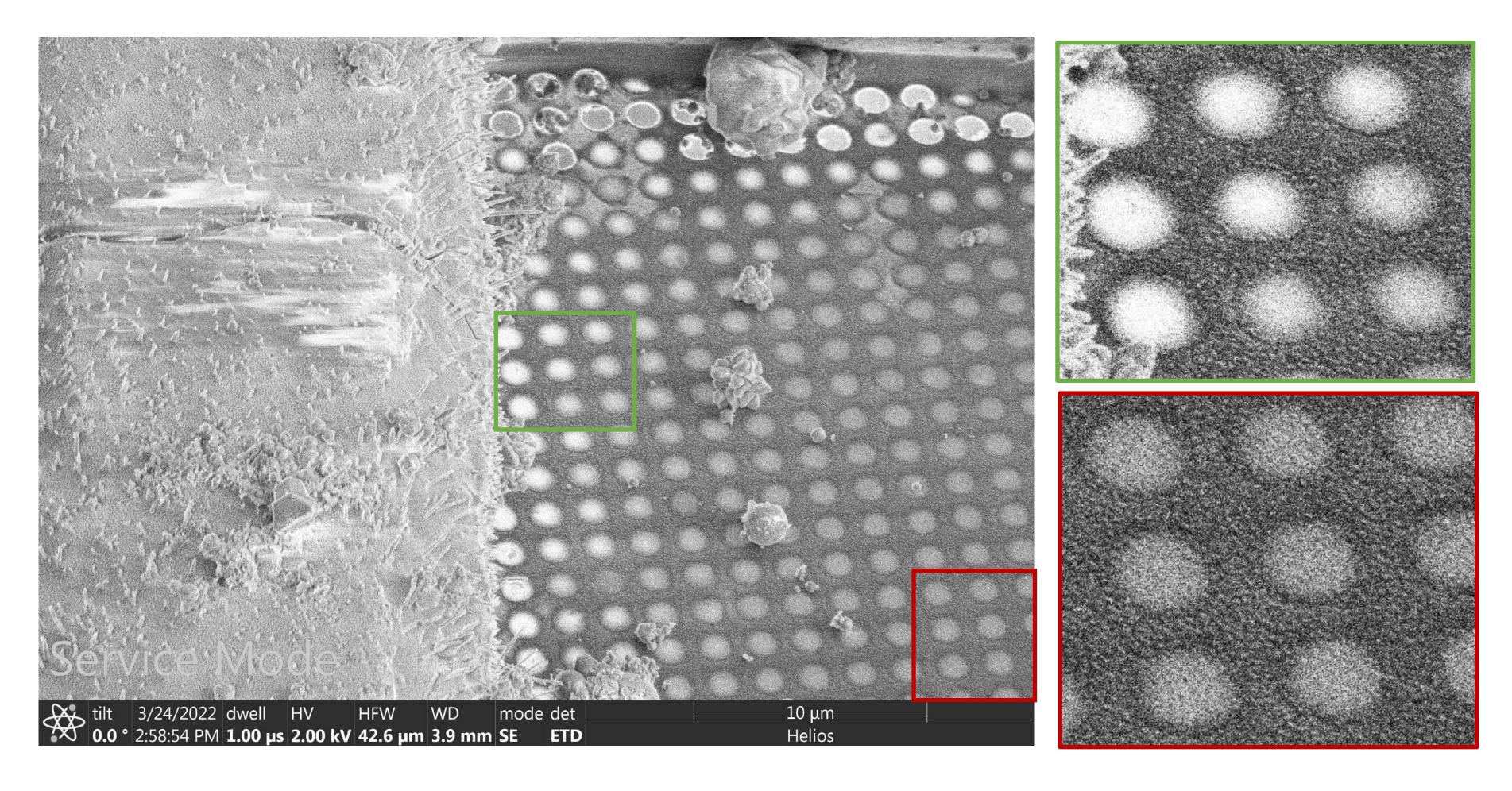


Cryo FIB-SEM





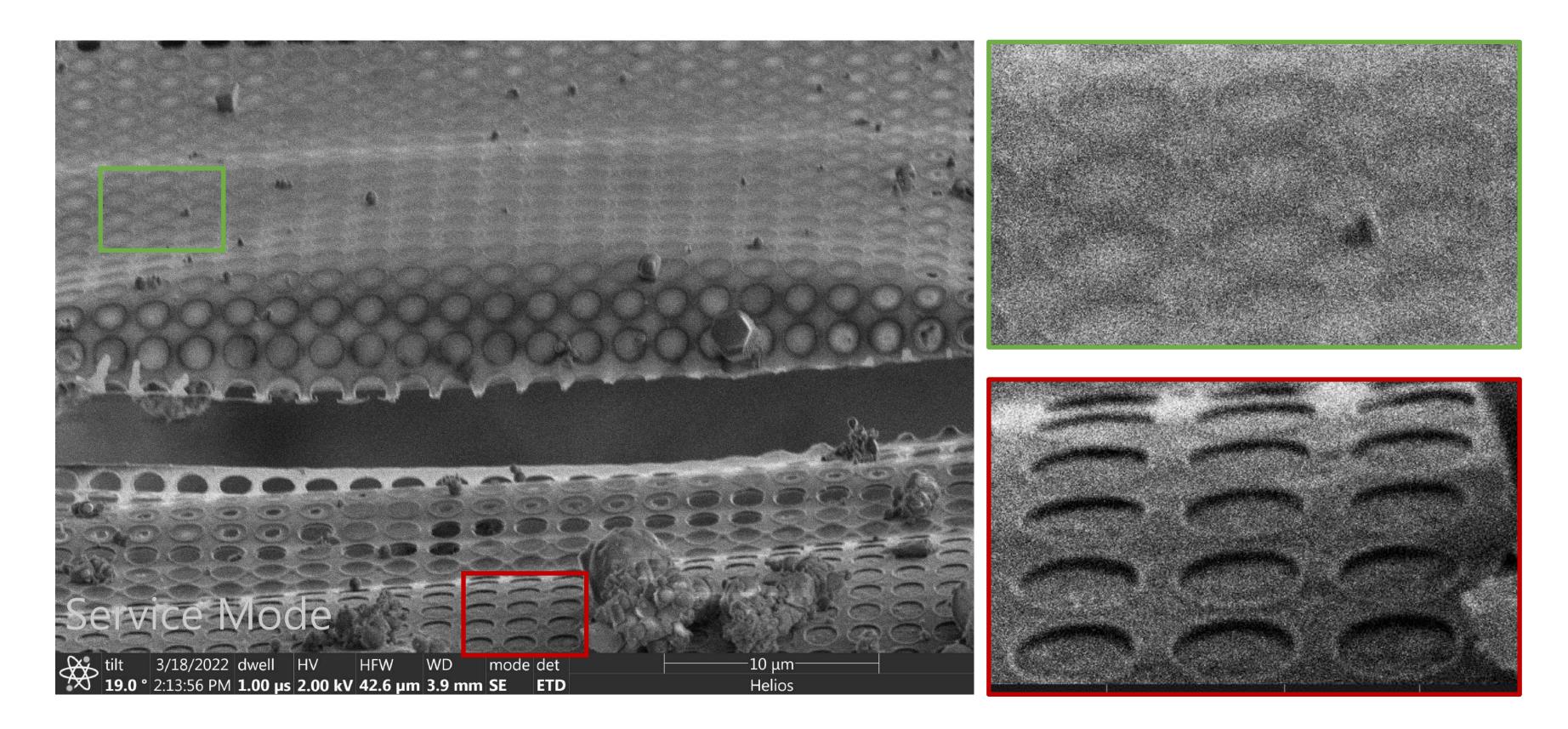
Sample application side of grid







Opposite side of sample application

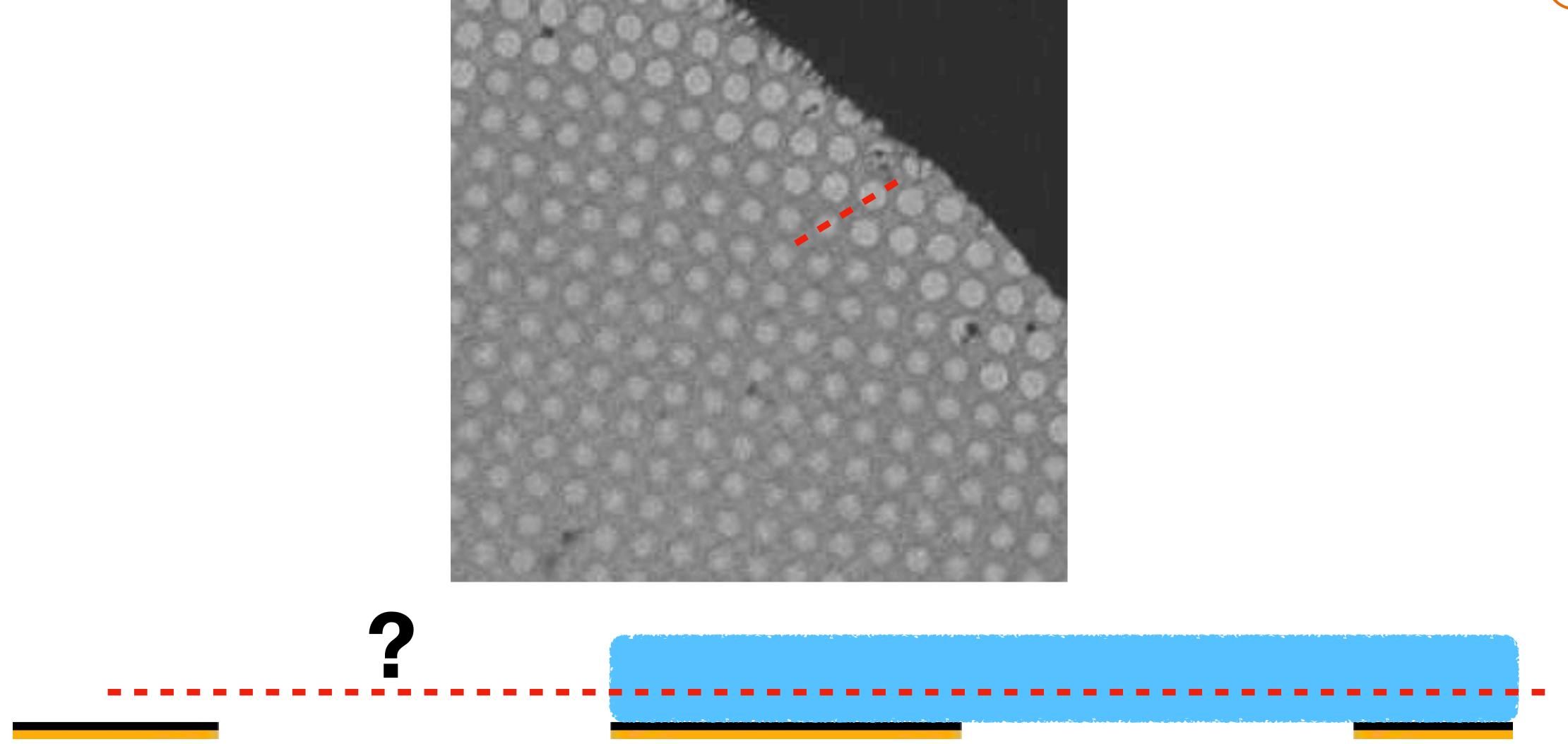








And why does this happen?

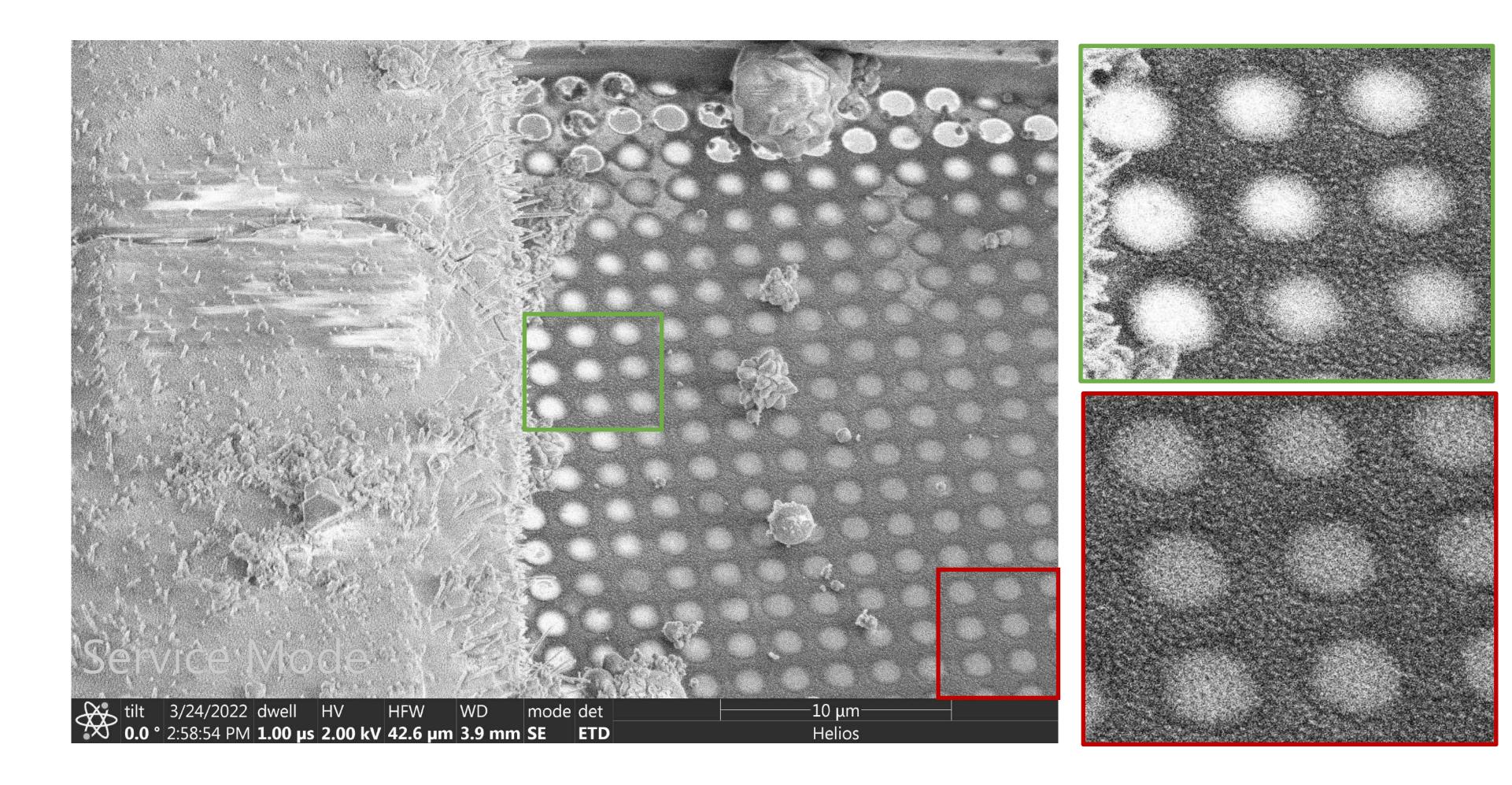


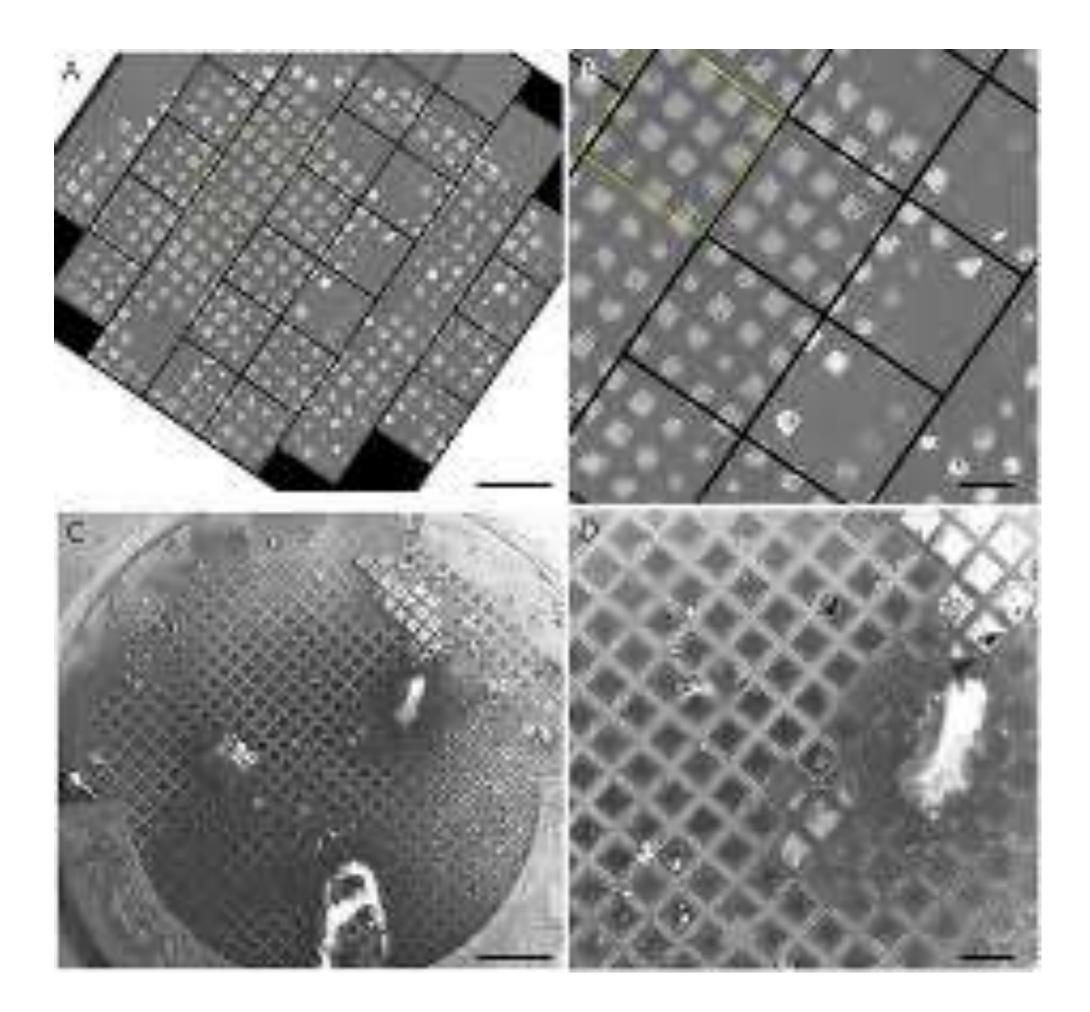




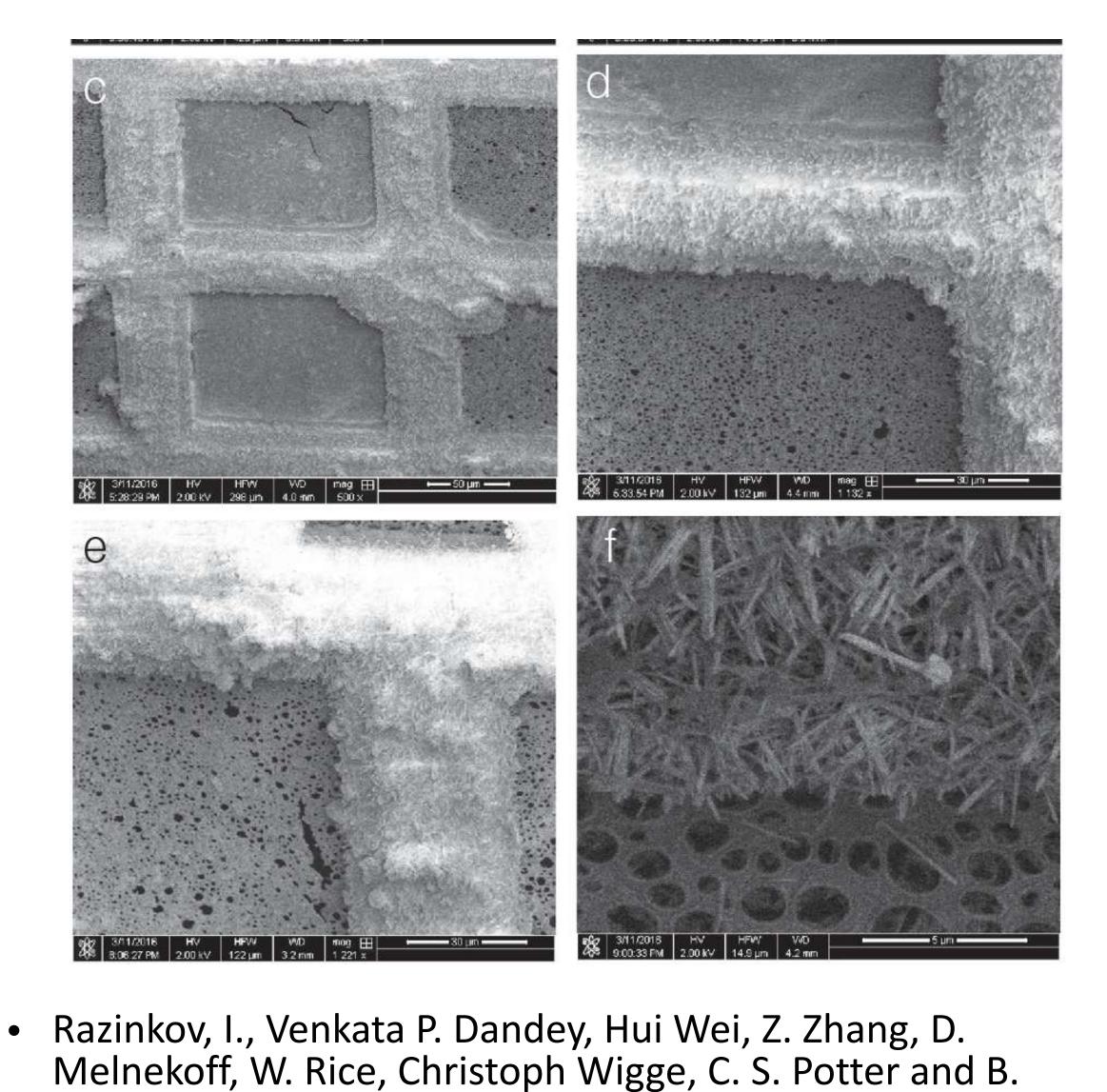


WHAT DOES A GRID LOOK LIKE?





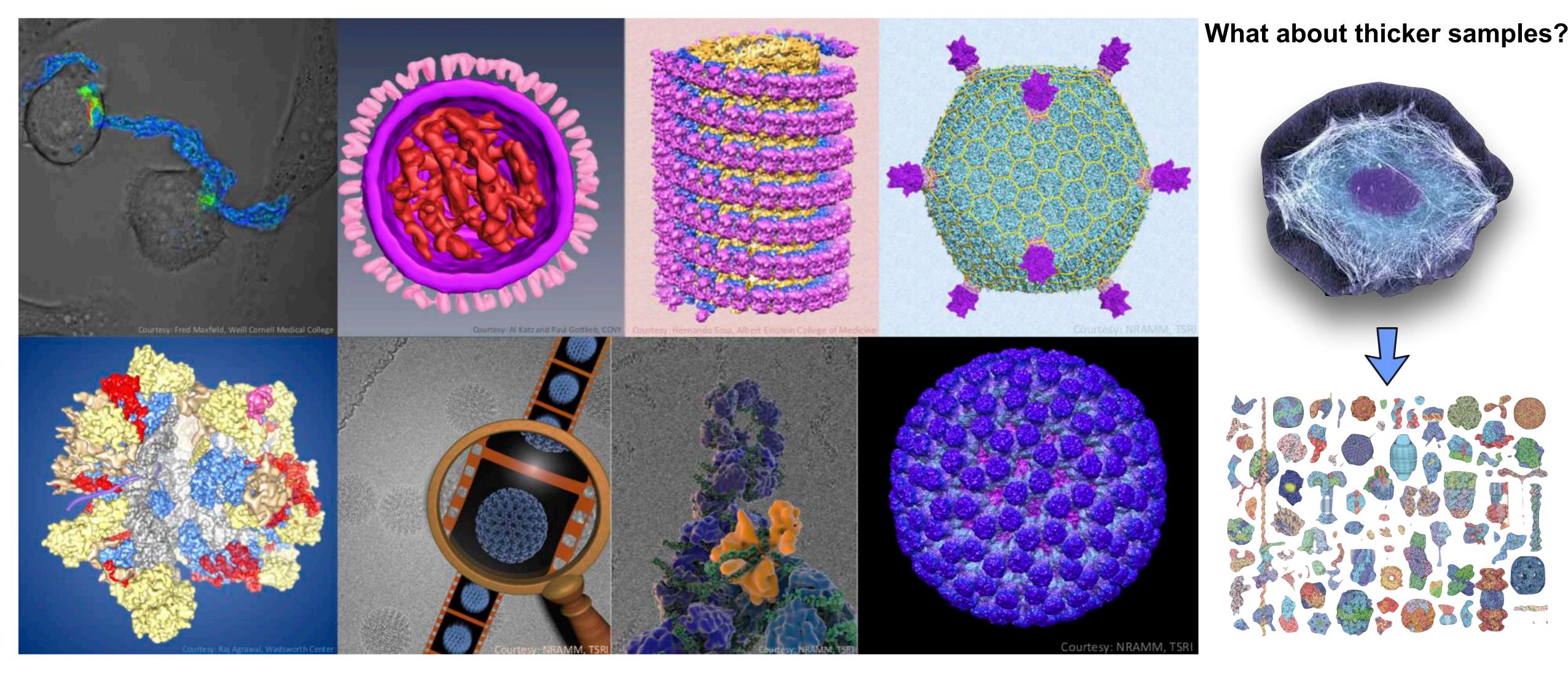
Schmidli, Claudio & Rima, Luca & Arnold, Stefan & Stohler, Thomas & • Syntychaki, Anastasia & Bieri, Andrej & Albiez, Stefan & Goldie, Kenneth & Chami, Mohamed & Stahlberg, Henning & Braun, Thomas. (2018). Miniaturized Sample Preparation for Transmission Electron Microscopy. Journal of Visualized Experiments. 2018. 10.3791/57310.



Carragher. "A new method for vitrifying samples for

cryoEM." Journal of structural biology 195 2 (2016): 190-198.

How are samples prepared for cryoEM?



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- - -







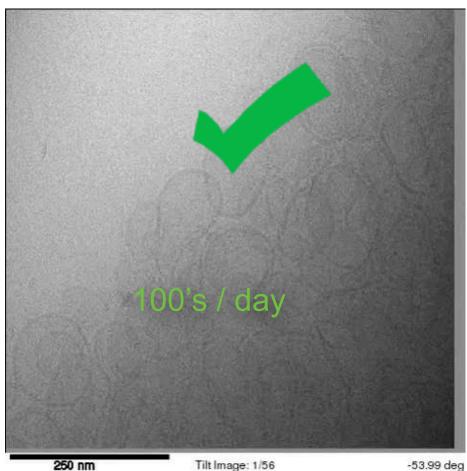
How are samples prepared for cryoEM?

Towards Automation for In Situ CryoEM

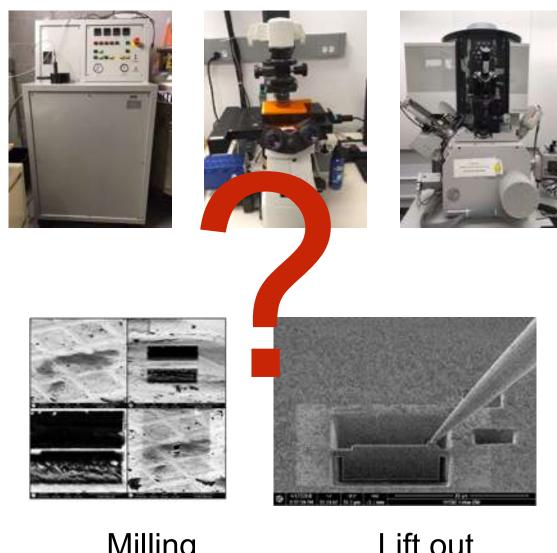


Sample





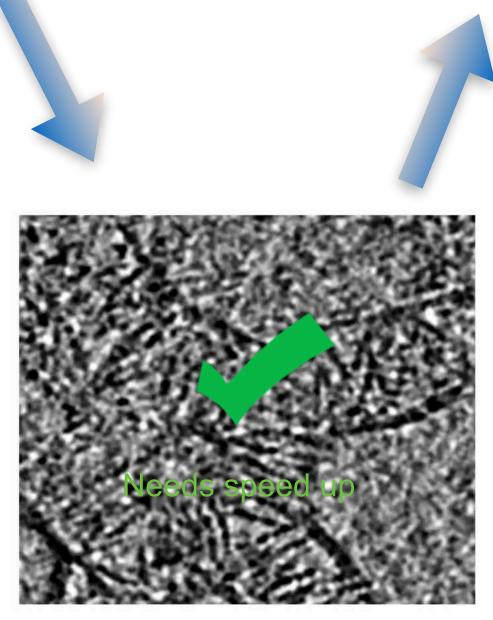
Automated Data Collection (Leginon, etc.)



Milling Grid preparation Lift out

Deep learning?

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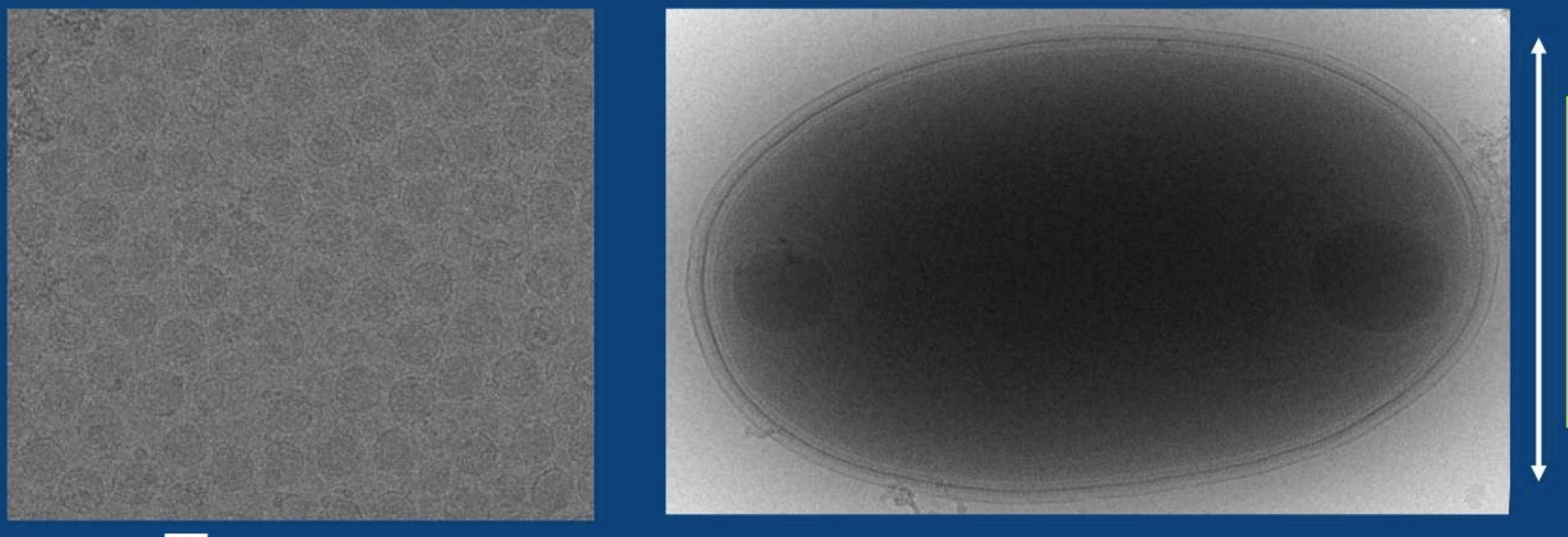


Streamlined Processing (Appion Protomo)





How are samples prepared for cryoEM? HOW THIN DOES THE SAMPLE NEED TO BE?



50 nm Bacteriophage (\$12)



E. coli, Salmonella, Cyanobacteria





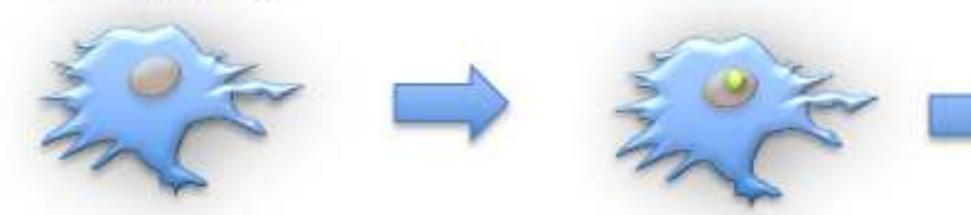


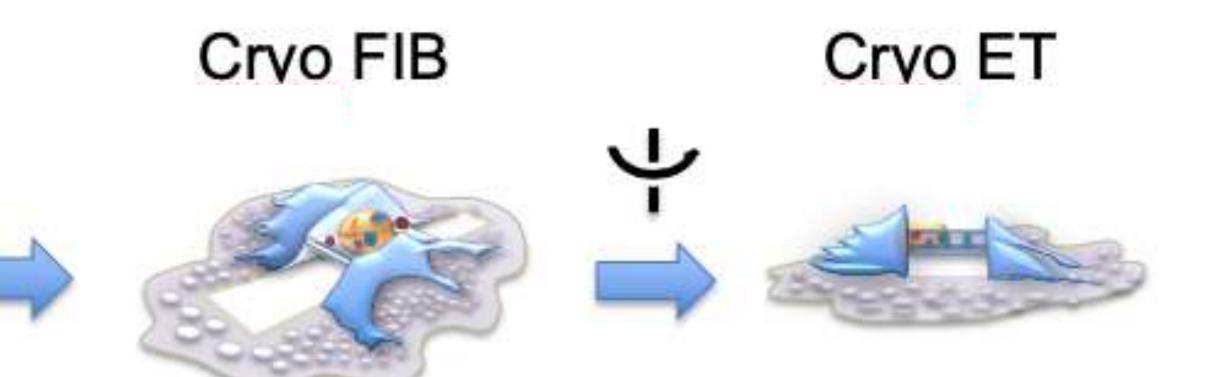
How are samples prepared for cryoEM?

CLEM workflow

Vitrified cell or tissue

Crvo LM











How are samples prepared for cryoEM? **STEP 1: Vitrify sample**

Plunge Freezing -Rapid freezing in liquid nitrogen (LN2)-cooled liquid ethane

High Pressure Freezing (HPF) -Rapid freezing at LN2 temps and high pressure



Leica Microsystems



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63



How are samples prepared for cryoEM? **STEP 1: Vitrify sample**

Sample: cells (adherent, suspension)

SEMC Equipment: Leica EM-GP, Gatan CP3, manual plunger

Bottlenecks: sample concentration, sample buffer, preferred orientation of sample on grid, plunge freezing parameters, grid mesh, support film, warming up, ice thickness, vitrification

Takeaways -Vitrify up to ~10 um of sample -One-sided blotting

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How are samples prepared for cryoEM? **STEP 1: Vitrify sample**

Sample: cells, tissue

SEMC Equipment: Wohlwend HPF Compact 01

Bottlenecks: sample amount, sample concentration, sample buffer, ice thickness, warming up, pressure, vitrification







How are samples prepared for cryoEM? **Takeaways STEP 2: Cryo-LM** -Do I have cells? Where? -Is my target fluorescing?

Sample: vitrified cells on bare or clipped grid

SEMC Equipment: Zeiss LSM 900 with Airyscan with Linkam Cryostage

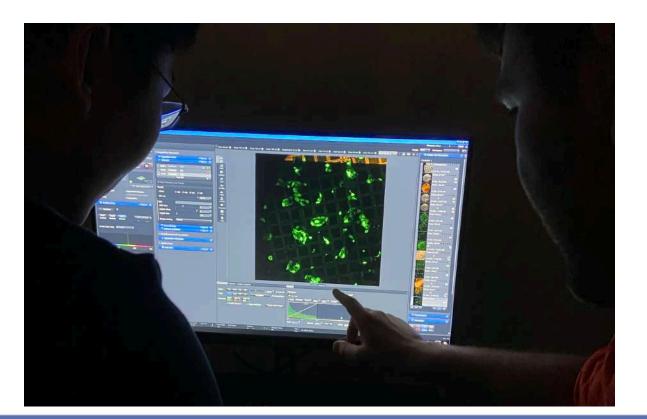
Bottlenecks: sample concentration, sample location, support film, laser damage, ice thickness, autofluorescence, warming up







Zeiss



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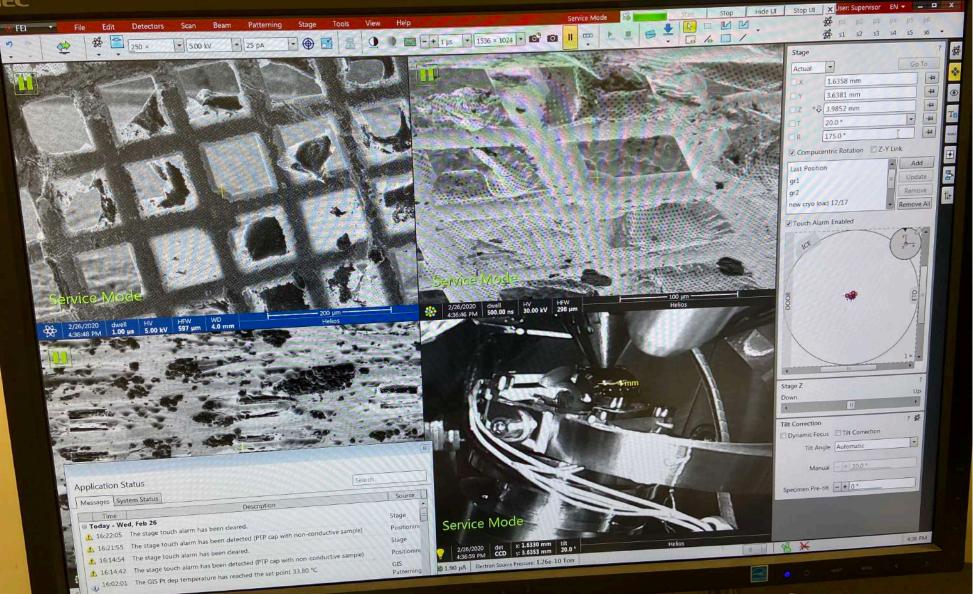




Sample: vitrified cells on a clipped grid

SEMC Equipment: Helios Nanolab G3 FIB-SEM, Leica VCT, VCM, & ACE

Bottlenecks: sample concentration, preferred orientation, imaging & milling parameters, grid mesh, support film, grid orientation, beam damage, charging, pole touches, dropping holder, image correlation, warming up, catching fire, ...

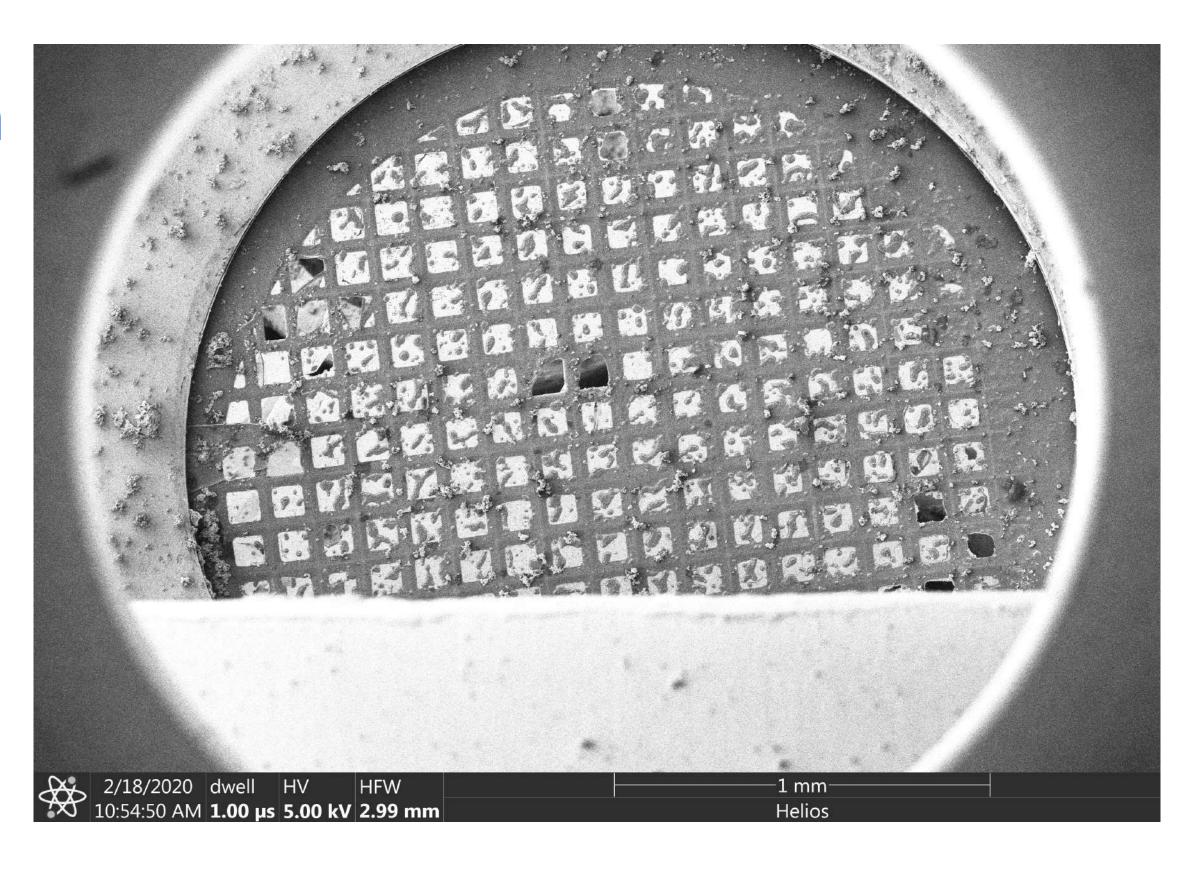


NYSBC





- 1) Grid overview/atlas images with electron and ion beam
- 2) GIS Platinum coat for 5 seconds
- 3) Mill large rectangular trenches on either side of each lamella site
- 4) Continue thinning lamella by milling on each side until <300 nm thickness

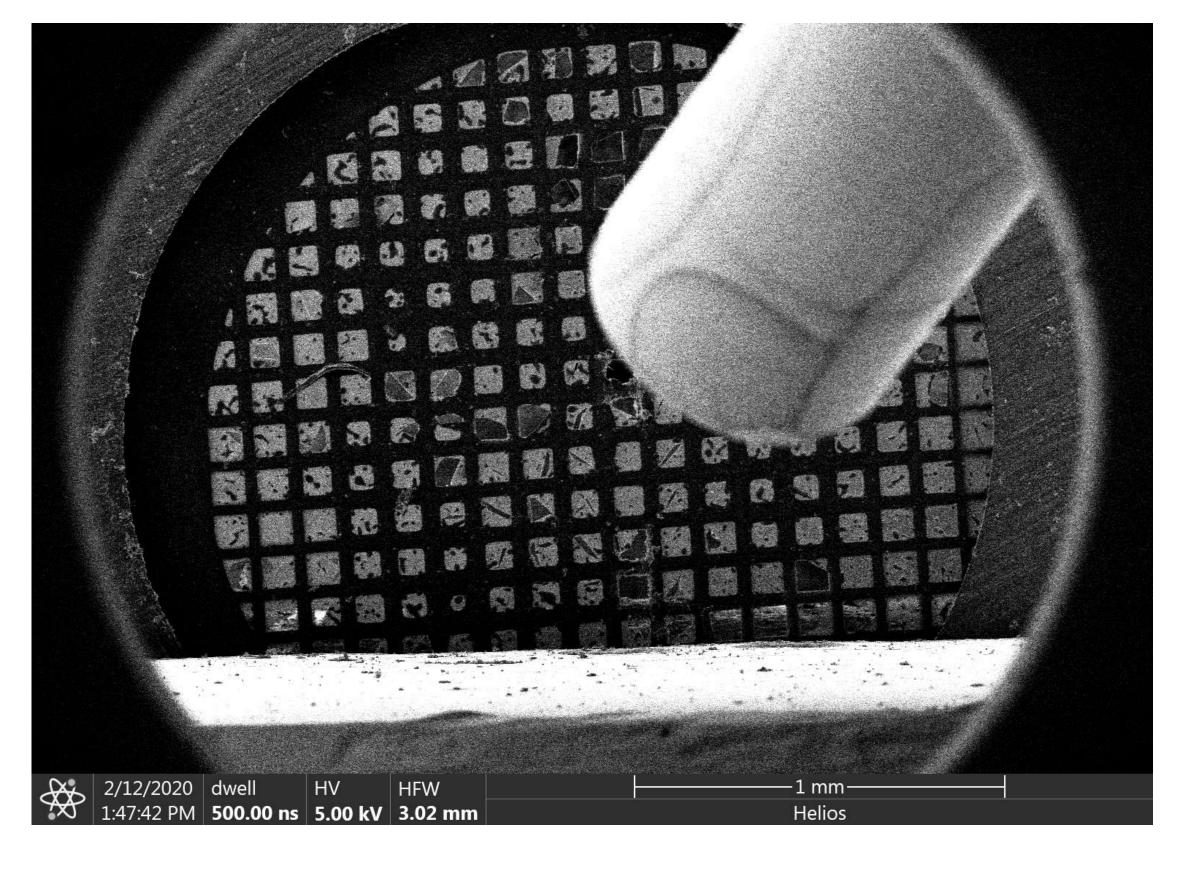


🗱 NYSBC **© SEMC**





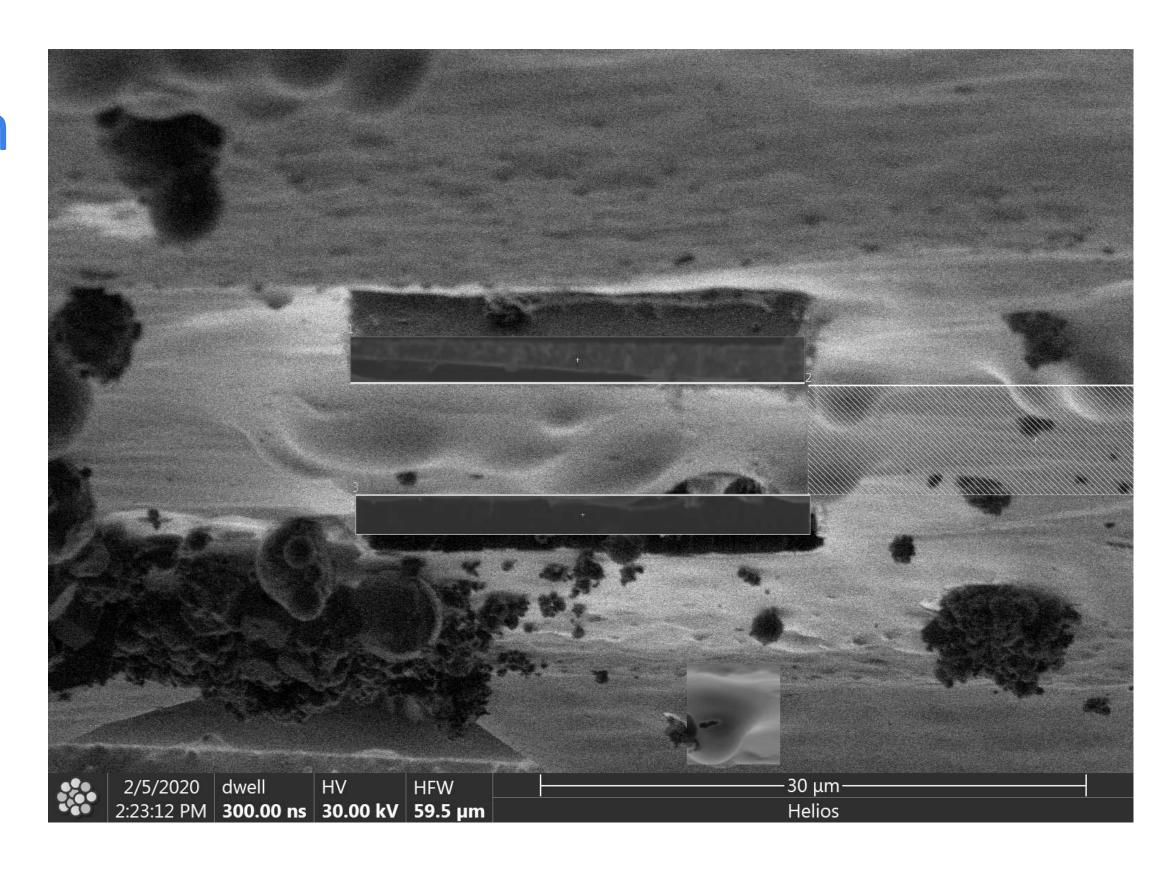
- 1) Grid overview/atlas images with electron and ion beam
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- 1) Grid overview/atlas images with electron and ion beam
- 2) GIS Platinum coat for 5 seconds
- 3) Mill large rectangular trenches on either side of each lamella site
- 4) Continue thinning lamella by milling on each side until <300 nm thickness

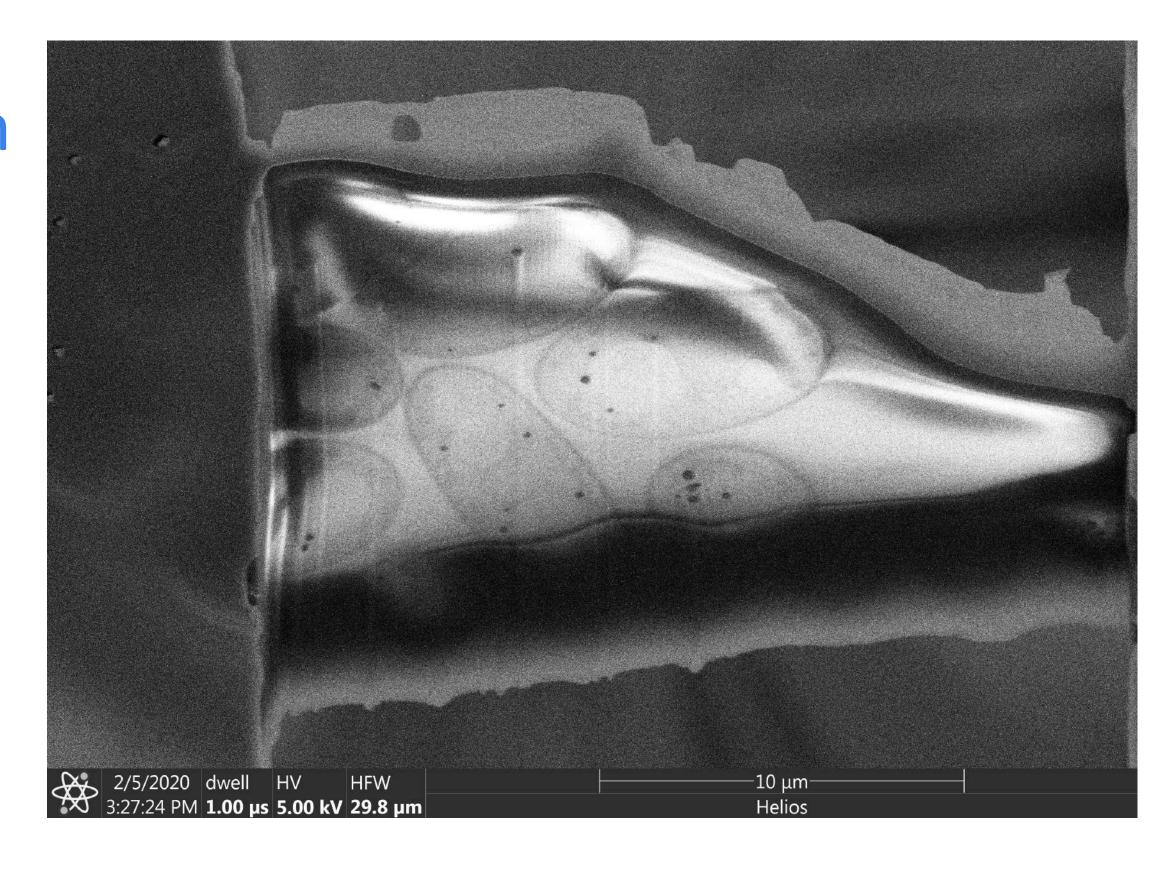


🗱 NYSBC





- 1) Grid overview/atlas images with electron and ion beam
- 2) GIS Platinum coat for 5 seconds
- Mill large rectangular trenches on either 3) side of each lamella site
- 4) Continue thinning lamella by milling on each side until <300 nm thickness

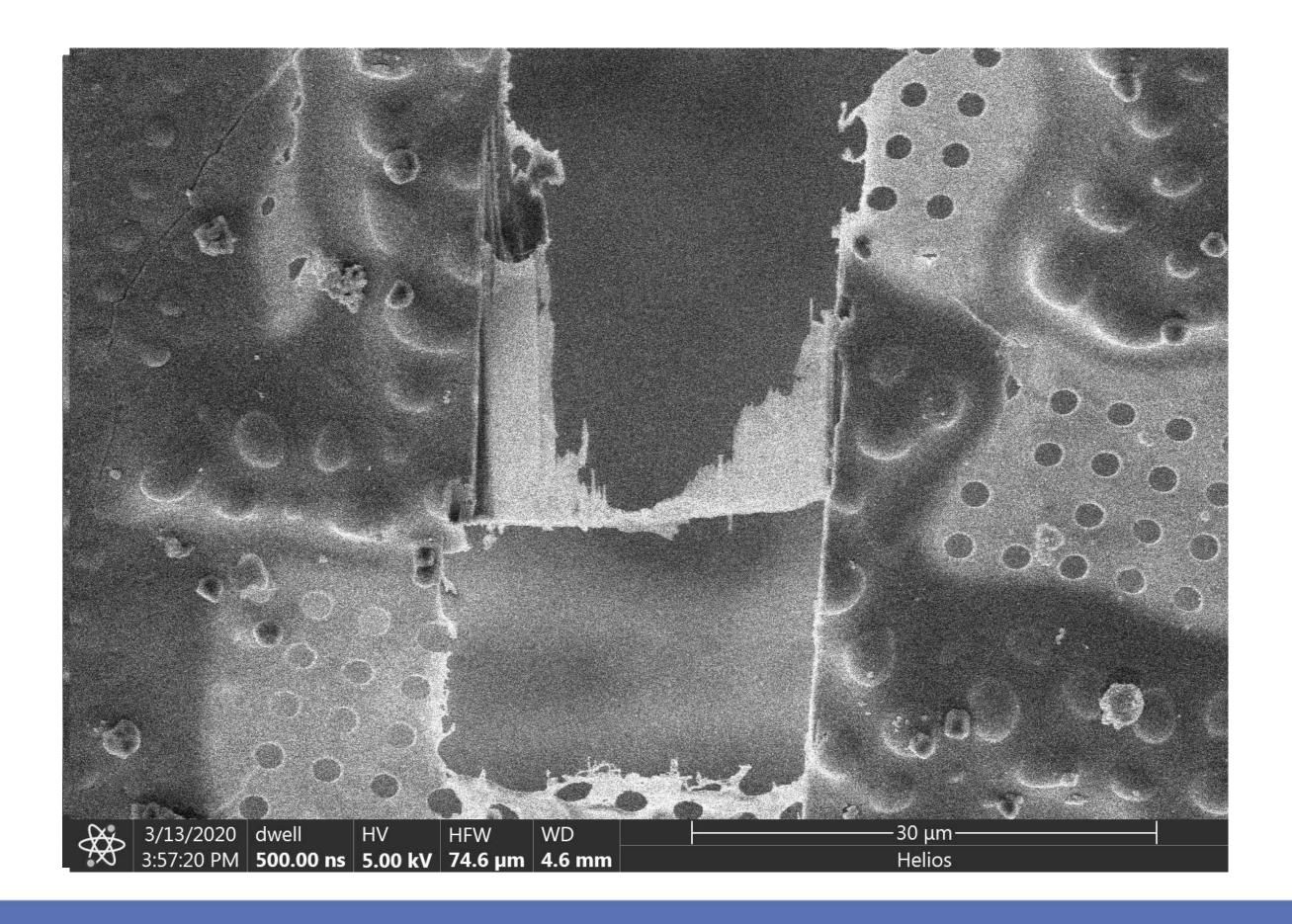


Regional NYSBC









Takeaways

-Large learning curve if done manually

-Many, many steps that all need to be completed successfully for experiment to work





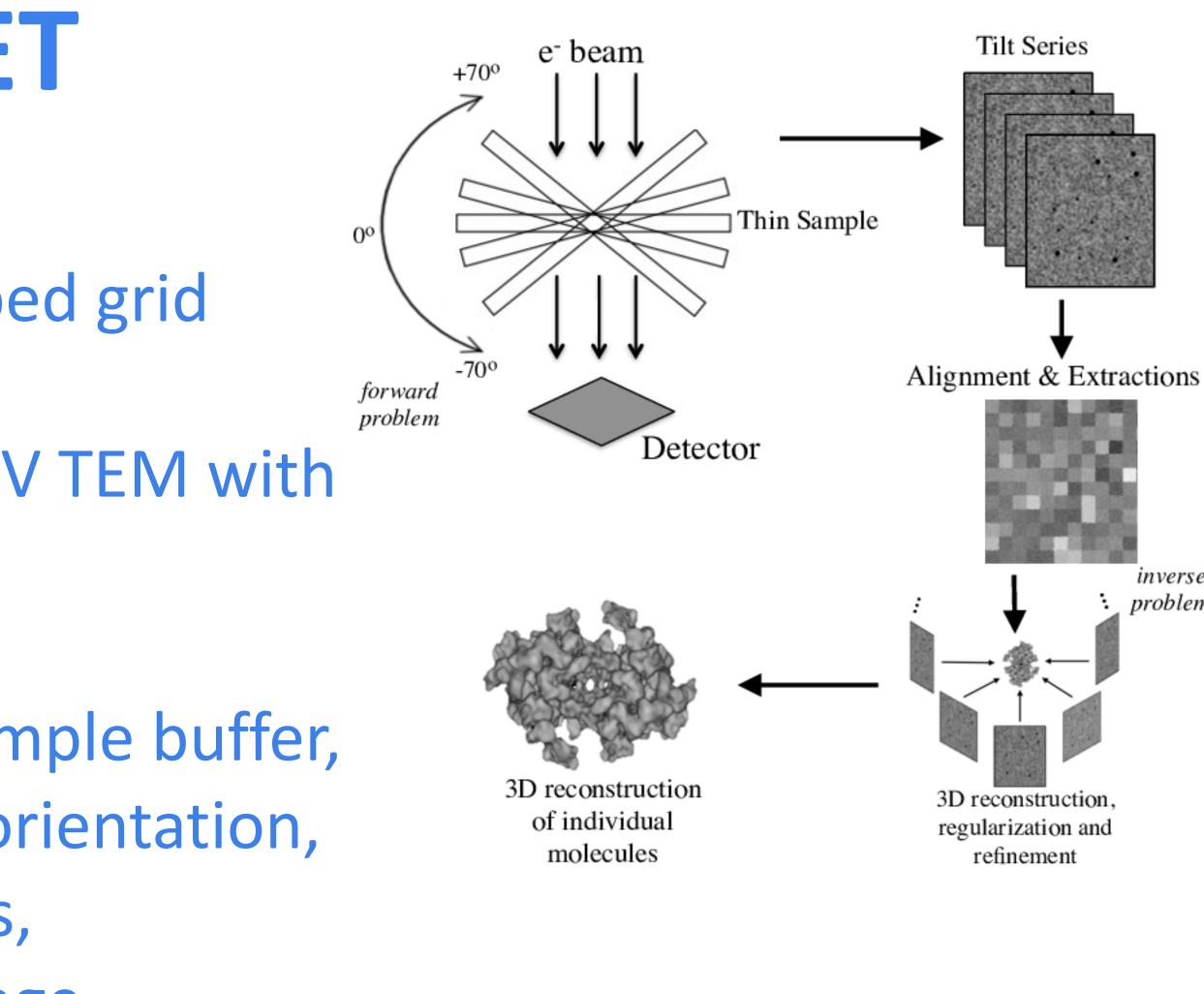


How are samples prepared for cryoEM? STEP 4: Cryo-ET e⁻ beam

Sample: whole cells or lamella on clipped grid

SEMC Equipment: TFS Titan Krios 300kV TEM with Energy filter + direct detector

Bottlenecks: sample concentration, sample buffer, preferred orientation, grid mesh, grid orientation, support film, warming up, ice thickness, vitrification, stage stability, beam damage, contamination, ...



NYSBC 🗱



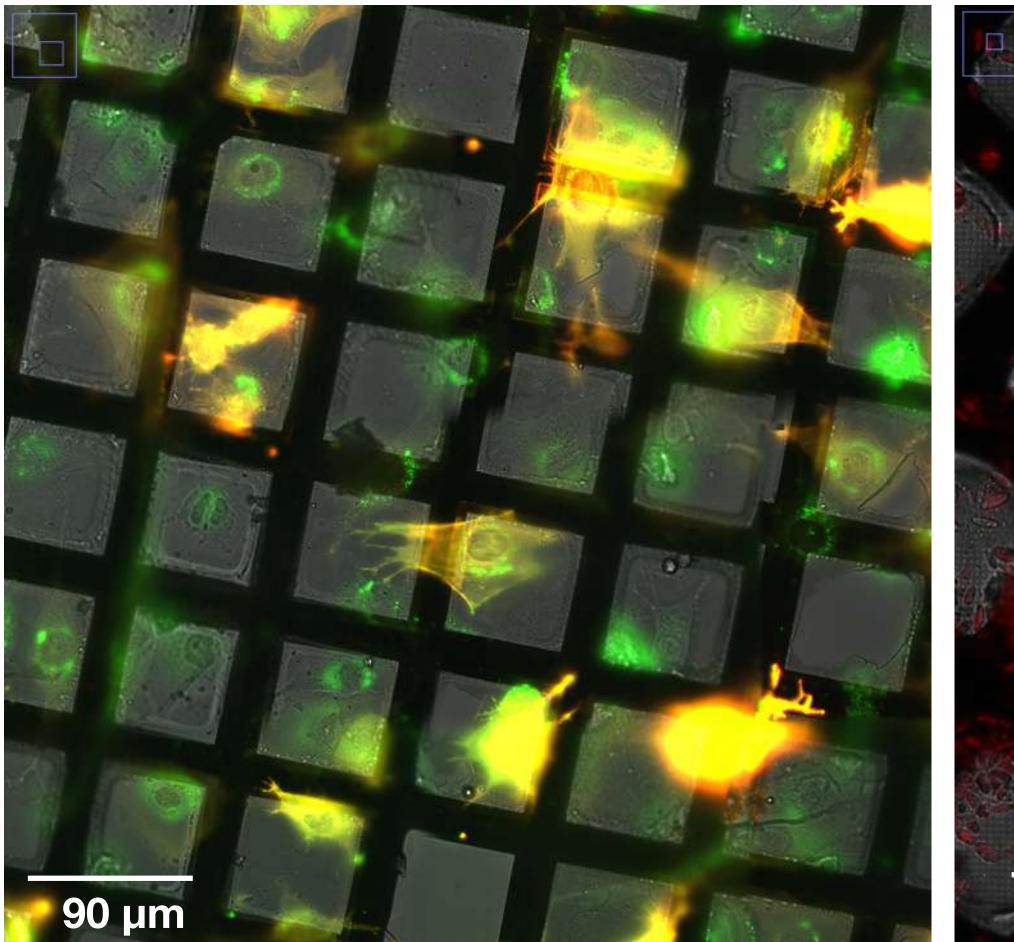




How are samples prepared for cryoEM?

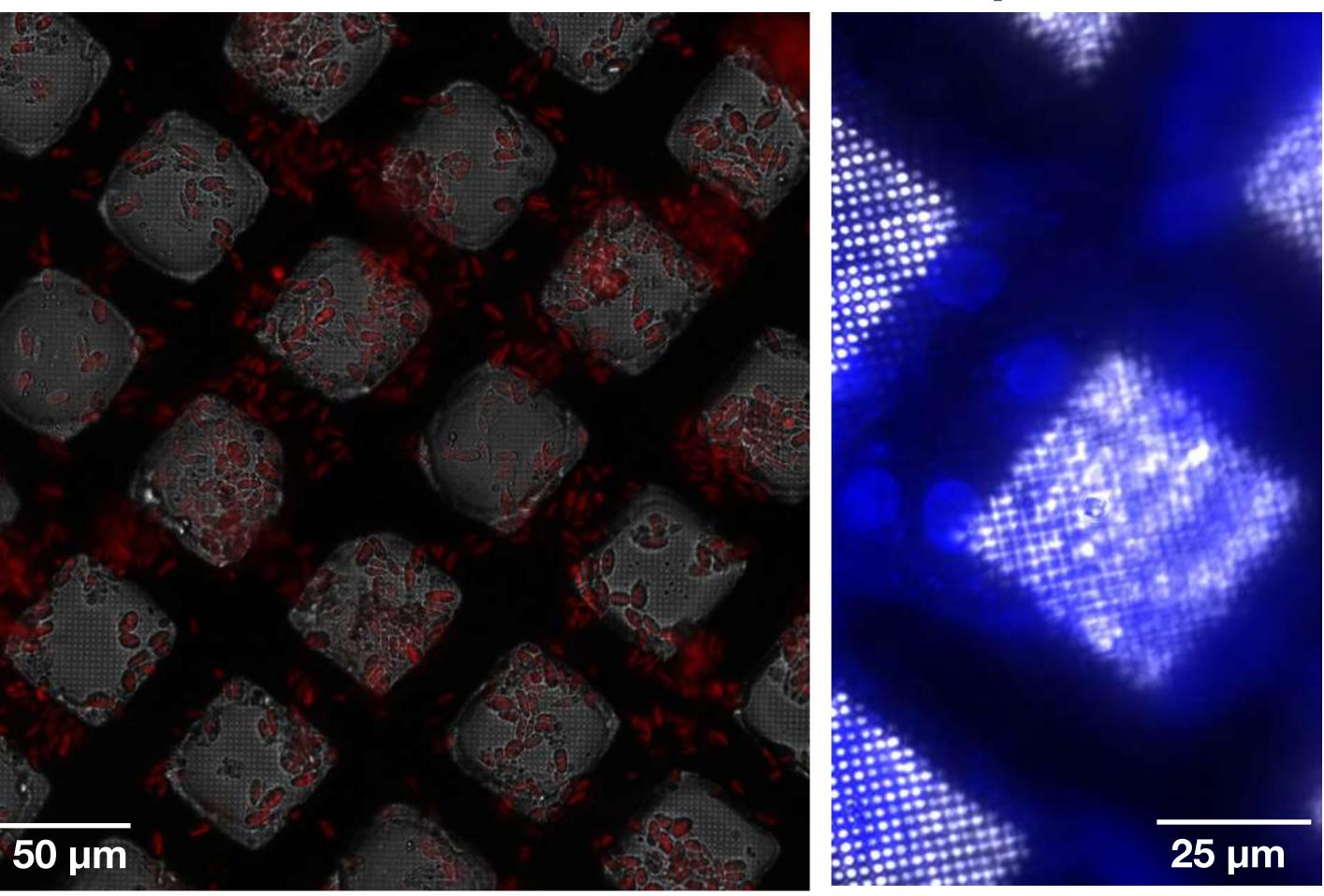
Mouse fibroblasts

Diatoms



Transfected adhesion signaling protein tagged with GFP (green) and F-tractin marker with mApple (red). Greg Alushin (RU)

Auto-fluorescence.



Microsporidia

Wei Dai (Rutgers) Gira Bhabha (NYU) **SEMC** 🗱 NYSBC

DAPI.



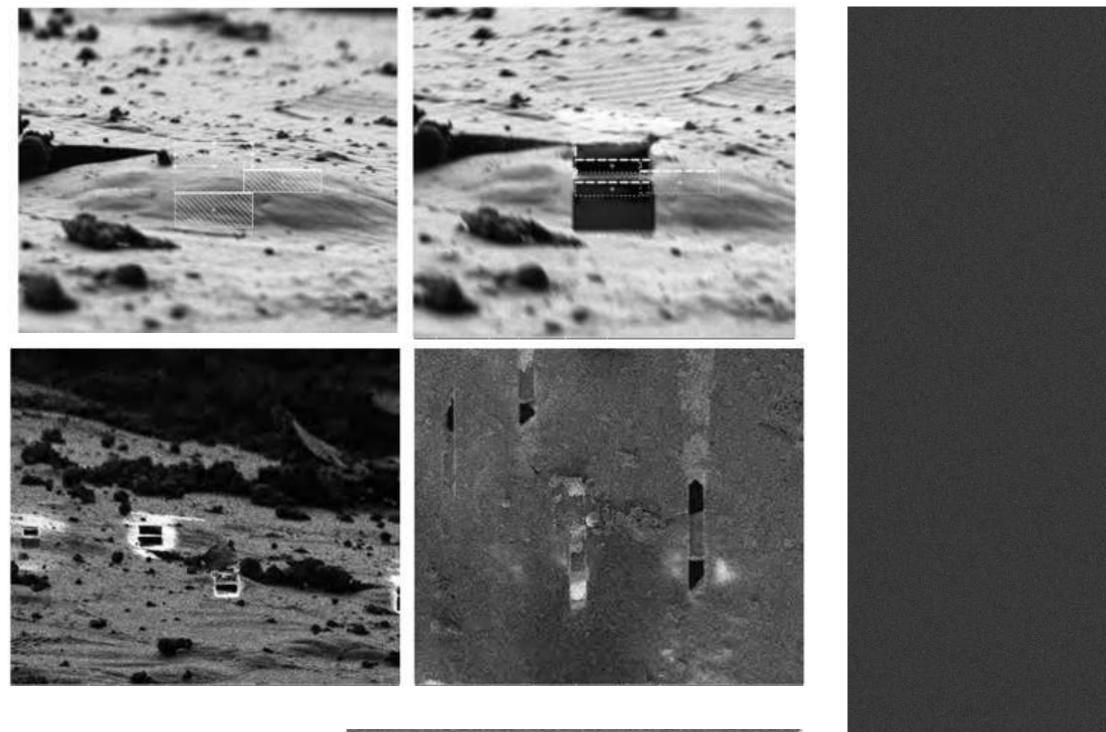
How are samples prepared for cryoEM?

Lamella

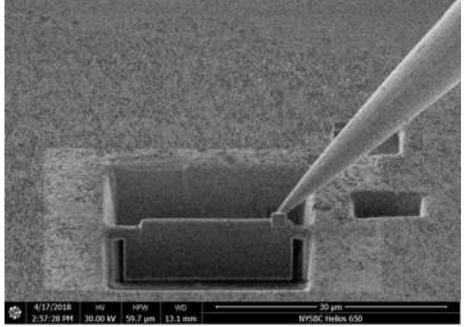
Rods

ΗV

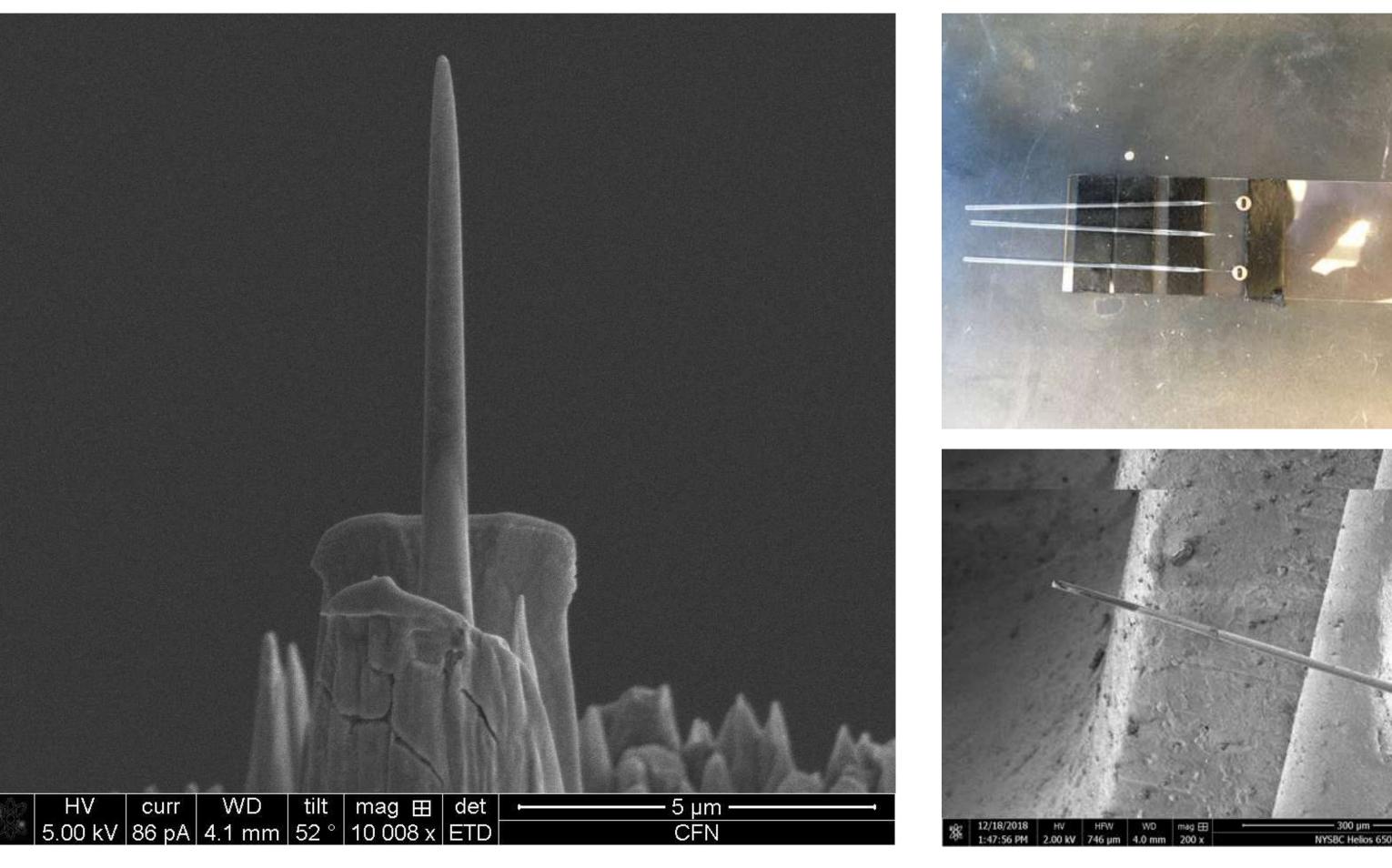
curr



with *lift-out*









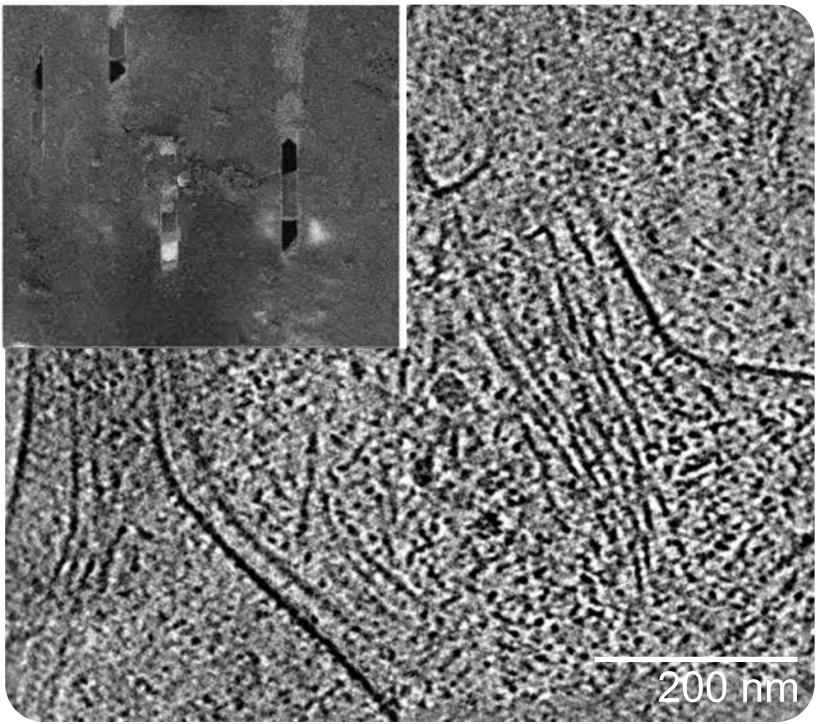


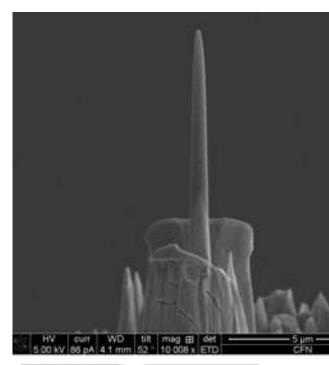






How are samples prepared for cryoEM? **Capillaries** Lamella Rods

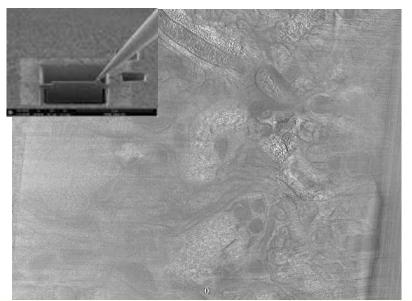




with *lift-out*

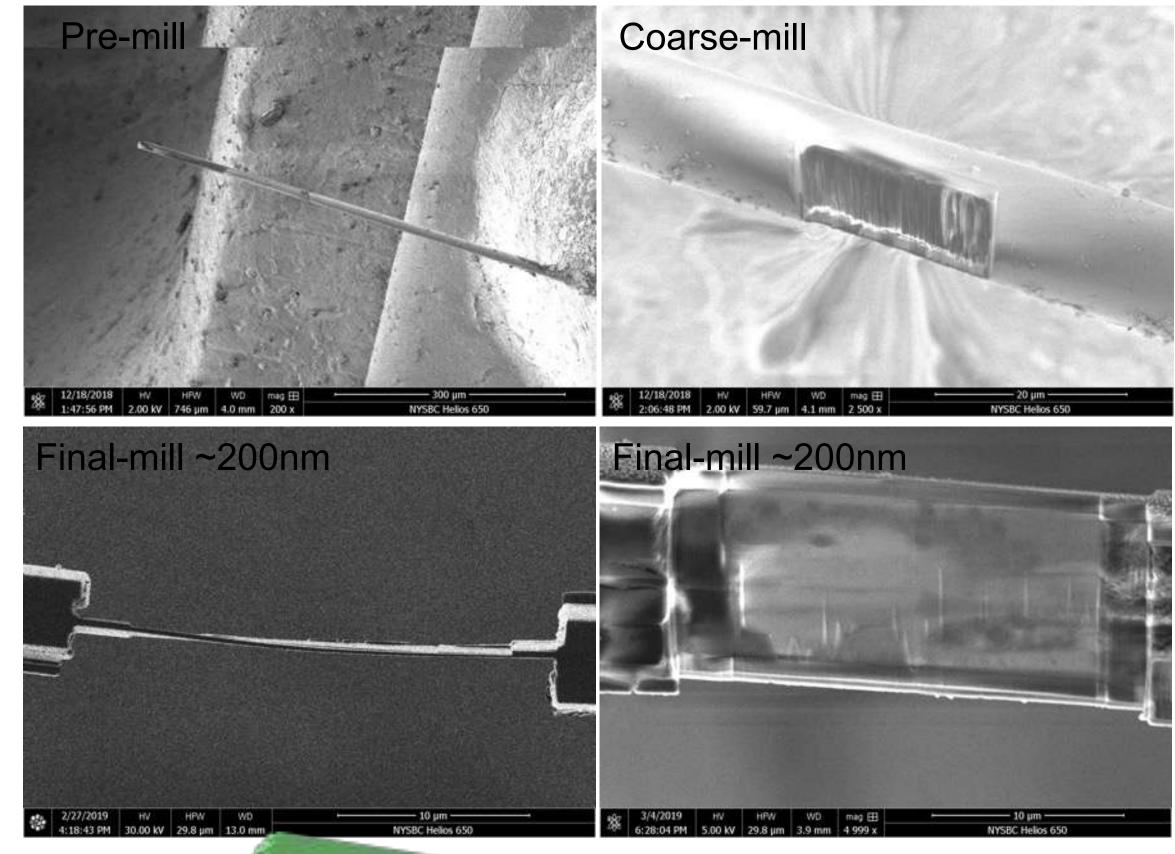
> Kotaro Kelley (NRAMM)

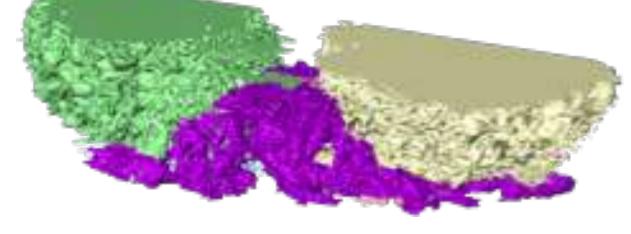
Zach Freyberg, (Univ. Pittsburg)



with Xin Group (BNL)

100 nm

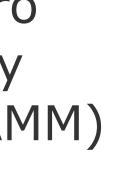




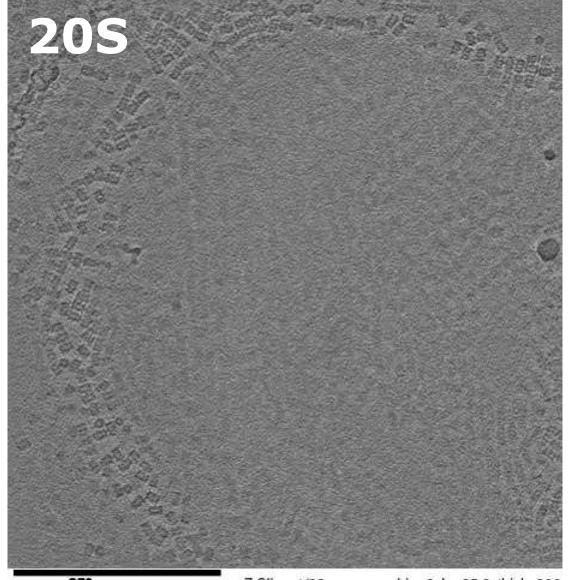
🗱 NYSBC

Kotaro Kelley (NRAMM)

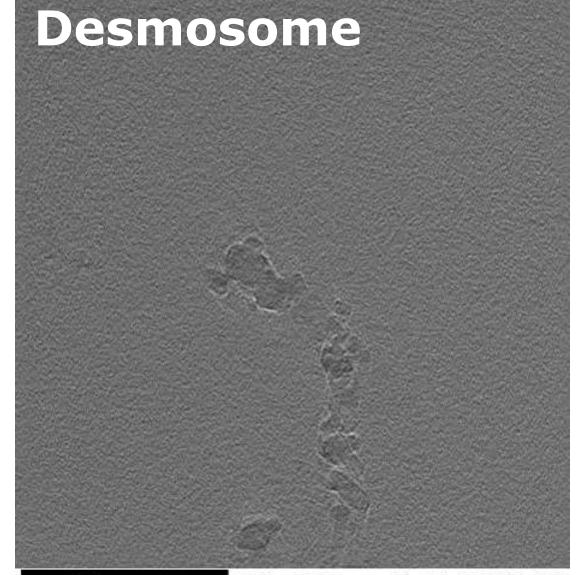




How are samples prepared for cryoEM?



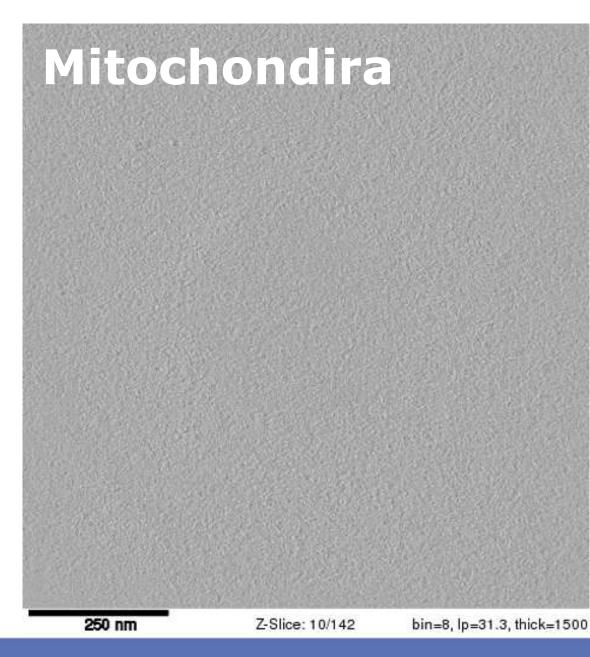
Alex Noble (NYSBC), Radostin Danev (MPI)



250 nm

Z-Slice: 1/28

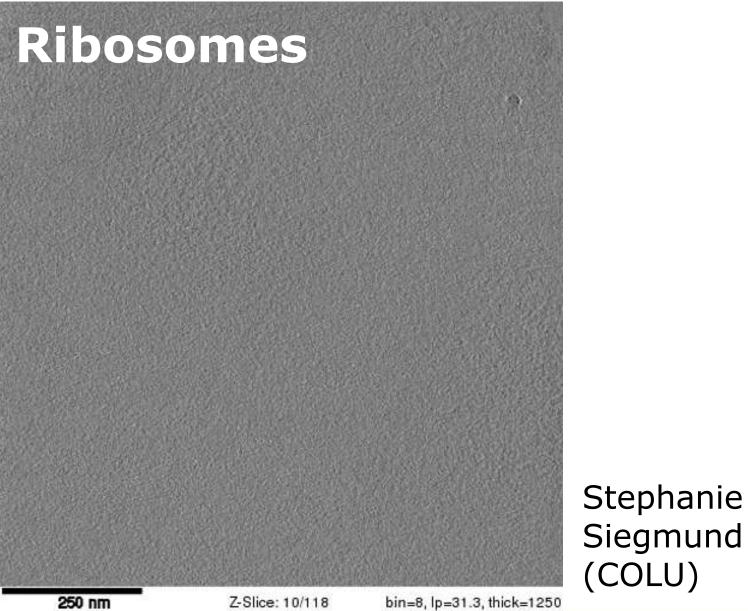
bin=8, lp=25.0, thick=200



Stephanie Siegmund (COLU)

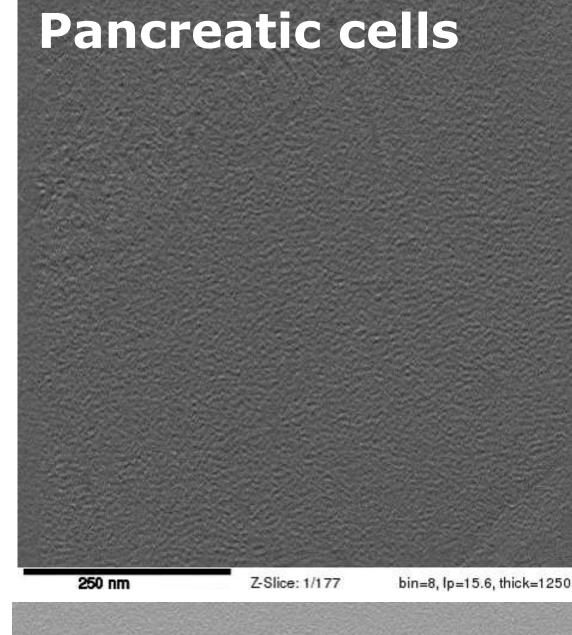
250 nm

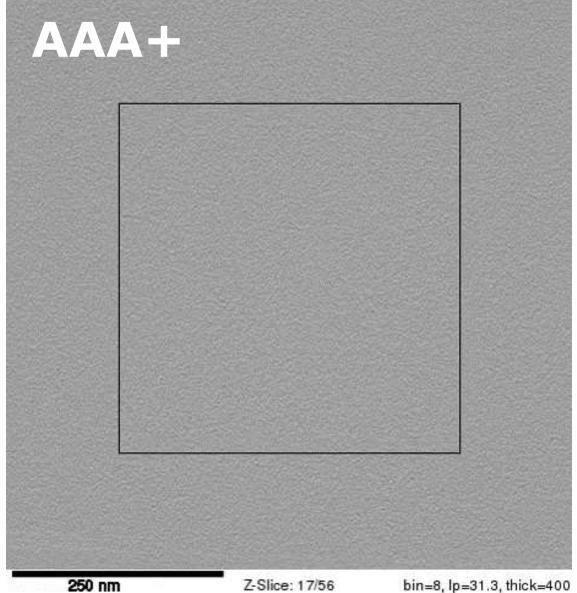
Z-Slice: 22/212



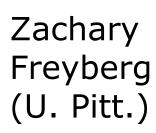
Julia Brasch (COLU)

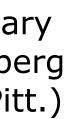
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NYSBC 🕄



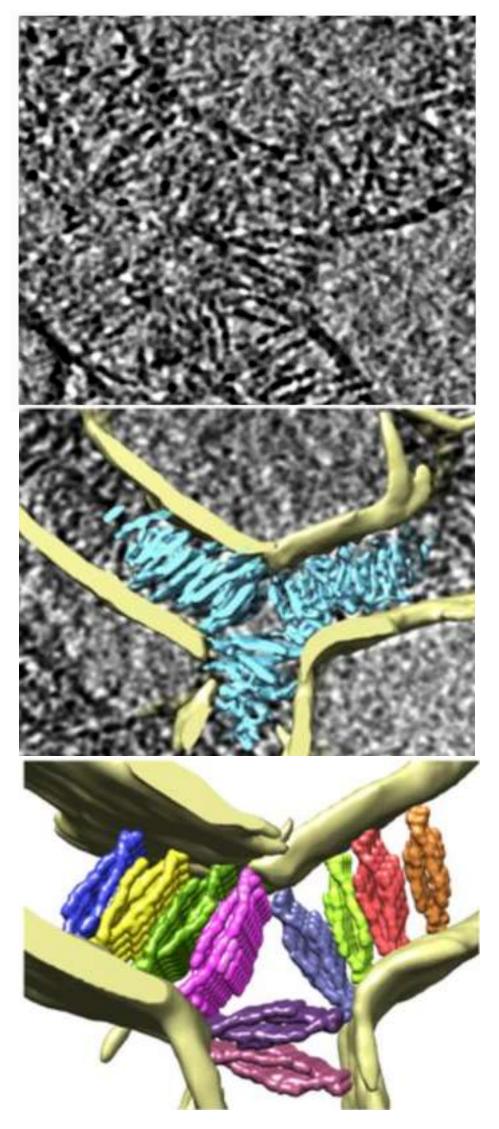


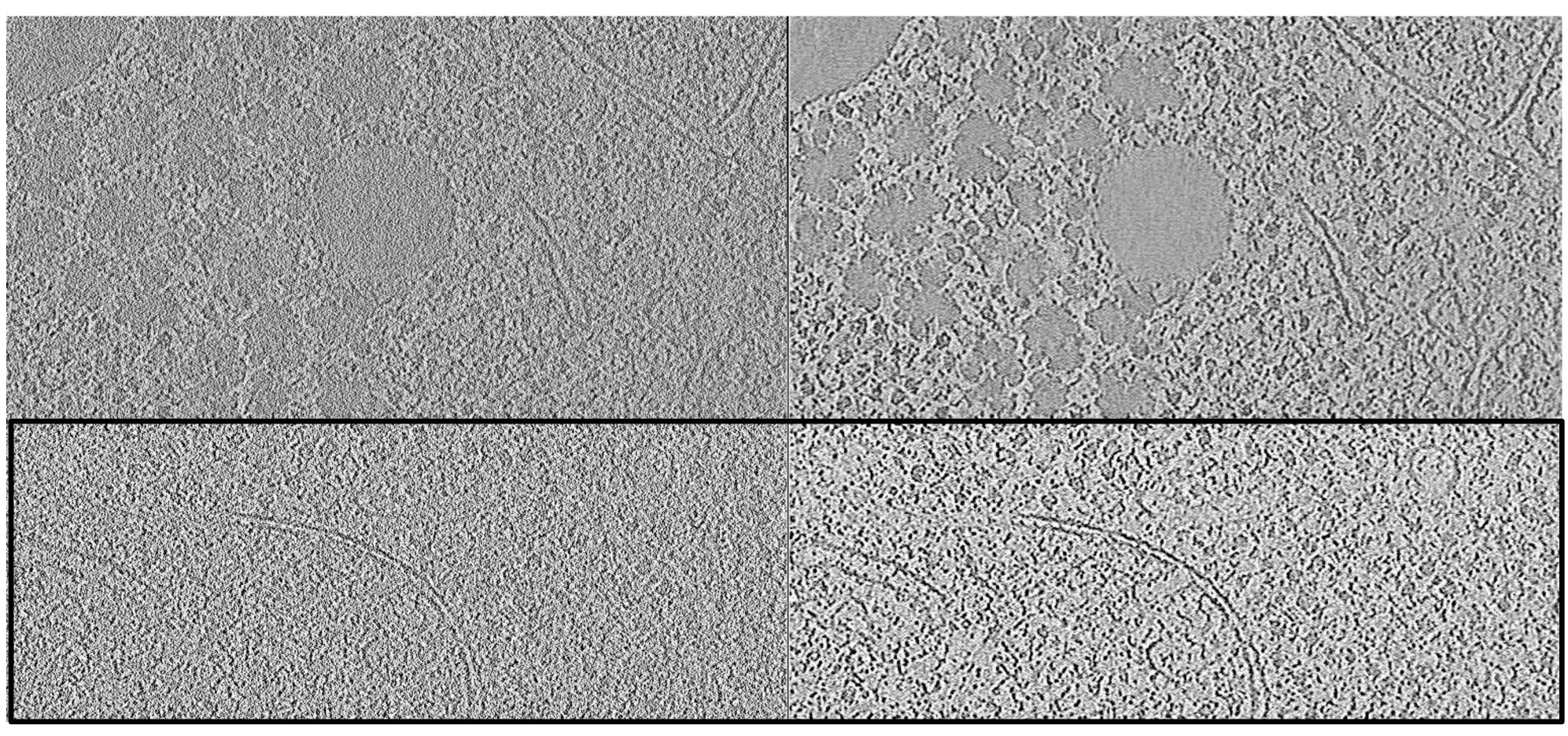


🗊 SEMC



How are samples prepared for cryoEM?





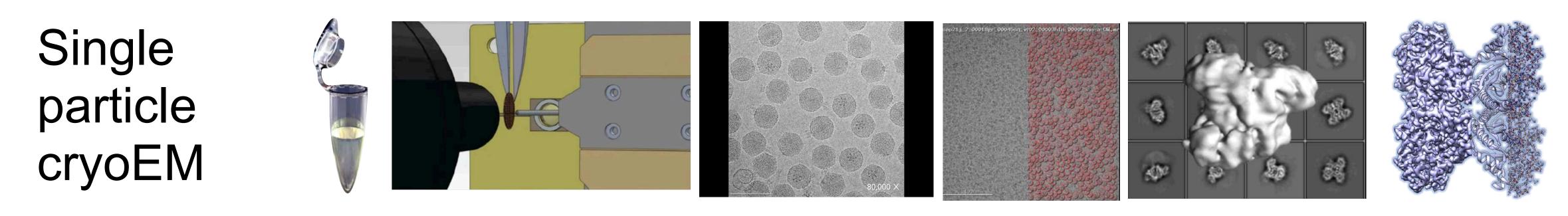
Micah Rapp (COLU)

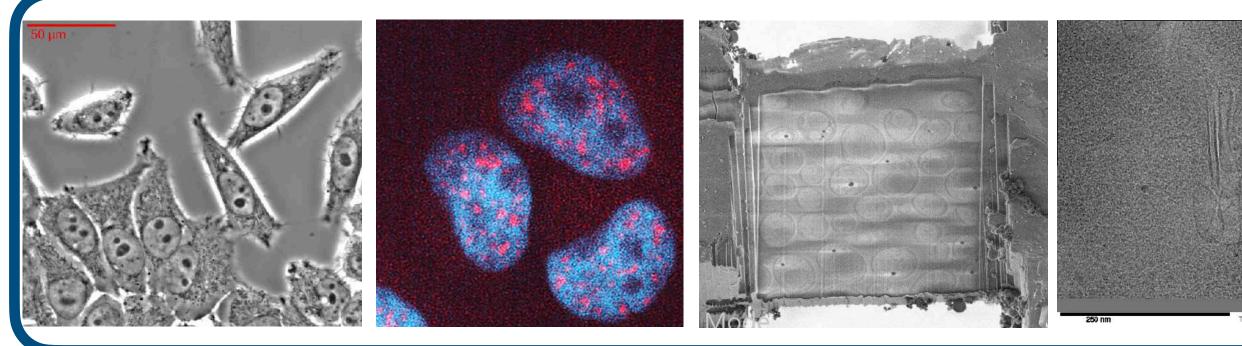
Kotaro Kelley (NRAMM)





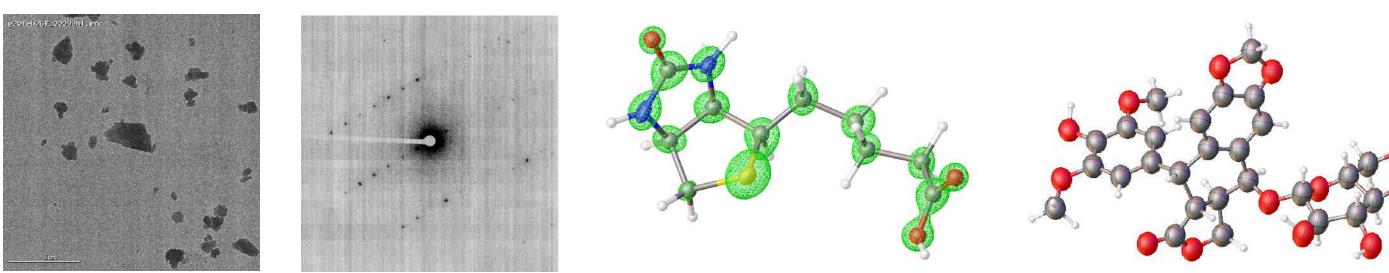
cryoEM: technology on the rise

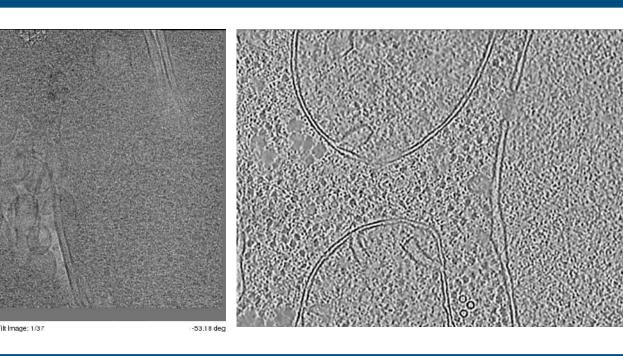




Micro crystal electron diffraction (microED)







Cryo Electron Tomography (cryoET)









TO BE CONTINUED

Questions?

