# Cryo sample preparation for cryoET investigations

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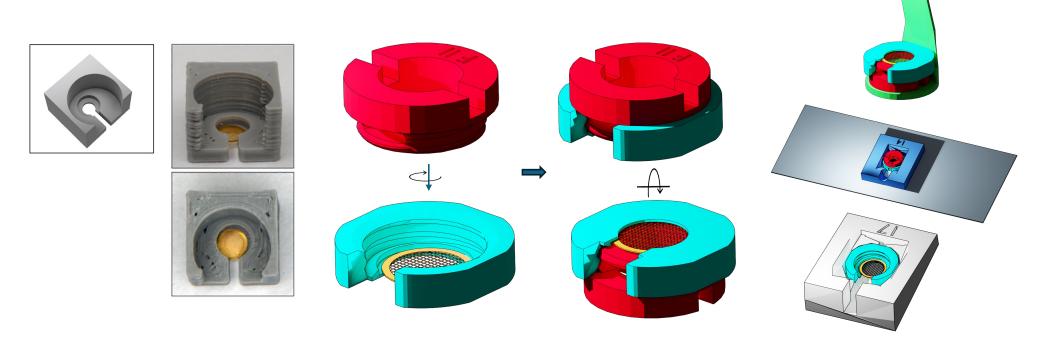
2025 Tomo Shortcourse - NYSBC

#### Goals

- 1. Get the cells on grids
  - Do the cells need to be milled (thin enough already)?
  - Do the cells need to be positioned on the grid (between bars)?
  - Do the cells need to be shaped (patterns of adhesive molecules)?
- 2. Freeze the samples
  - Do the samples have the right geometry to plunge-freeze?
    - · Thickness and total mass
  - Do the samples need cryoprotectants?
- 3. Locate the targets for milling/CryoET
  - Labeling prior to freezing if needed

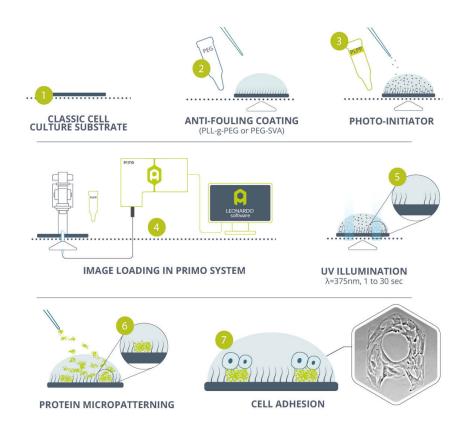
# Preparing Grids for cell growth

- If cells are to be grown directly on the grid, the normal carbon support might not be sufficiently stiff or resilient
  - Evaporate ~20-25 nm carbon on top, and baked overnight at 60° C
- Handling the grids prior to and during culturing can lead to breaking and bending
  - 3-D printed cases let you handle without directly touching the grids until freezing

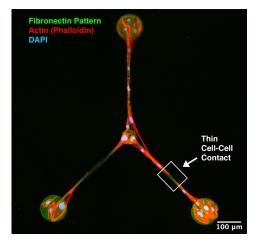


Fäßler, F., Zens, B., Hauschild, R., & Schur, F. K. M. (2020). 3D printed cell culture grid holders for improved cellular specimen preparation in cryo-electron microscopy. Journal of Structural Biology, 212(3), 107633.

### Positioning cells for milling: PRIMO



- PRIMO system uses a DMD (digital micromirror device) to illuminate an arbitrary pattern with UV light using the microscope's objective lens
- 1. Coat the grid in a non-cell adhesive layer (e.g. PLL-PEG)
- 2. Add the UV-sensitive photo-activator (PLPP)
- 3. Illuminate the grid using the user-supplied pattern
- 4. Add cell-adhesive molecule (fibronectin, PLL, etc)
- 5. Seed the cells
  - May need to play with the concentration

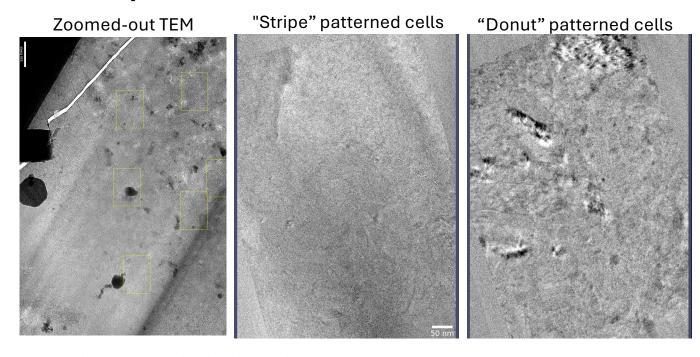


Source: https://www.alveolelab.com/technology/micropatterning-surface-functionalization/

#### Vitrification of Cells

- Need to cool down the entire sample fast enough to prevent crystalline ice formation
  - 290 K to 90 K: ~20 μs
  - ~10<sup>6</sup> degrees/second
- Heat needs to transfer from the sample to the cooling medium, and away then from the sample to allow for more heat to move from more interior in the sample.
  - 1. Transfer within sample to interface
  - 2. within cooling medium away from sample
- Limits the thickness of plunge-frozen samples to less than 1 µm
  - Total mass on grid also affects freezing
- Once ice starts, excess heat from heat of formation also needs to be removed!
- Below ~ -40° C, homogenous ice formation begins
  - · Doesn't need nucleation centers, spontaneously forms
  - The small ice crystals form don't affect freeze-substitution, but cryo-ET is harmed

#### Incomplete vitrification of cells



- "Burning" pattern indicative of incomplete vitrification can be seen zoomed-out TEM
- Appearance changes with tilt angle due to crystal planes of the crystalline water
- Lowers the S/N of the back-projected tomogram
  - · Distorts the structure of embedded proteins?
- Counter with
  - 1. Cryoprotectants
  - 2. High-pressure freezing

### Cryoprotectant agents

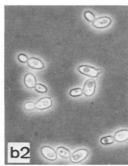
#### 1. Cell permeable

 Most commonly used is glycerol – can affect cellular structure

#### 2. Cell impermeable

- Only affects freezing in media surrounding cells
- Heat of ice formation doesn't need to be removed





0% Glycerol

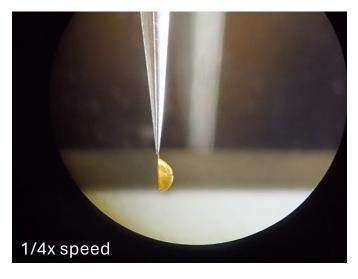
9% Glycerol

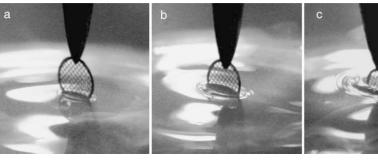
Cryoprotectant	Degassed	Vitrification	Background
Native	-	Incomplete	None
15% PVP in PBS	-	Incomplete	Acceptable
15% PVP in 0.1 M PB	-	Incomplete	Acceptable
15% BSA in 0.1M PB	-	Incomplete	High
20% Dextran/5% Sucrose in PBS	-	Complete	High
20% Dextran/5% Sucrose in PBS	+	Complete	High
20% Dextran in PBS	+	Incomplete	Acceptable
10% Dextran in PBS	+	Incomplete	Acceptable
10% Dextran in PB	+	Complete	Acceptable
5% Dextran in PB	+	Incomplete	Acceptable
Native	+	Incomplete	None
10% BSA in medium	+	Incomplete	Acceptable
20% BSA in medium	+	Complete	High

Niedermeyer, W., Parish, G. R., & Moor, H. (1977). Reactions of yeast cells to glycerol treatment alterations to membrane structure and glycerol uptake. *Protoplasma*, 92(3–4), 177–193. Zens, B., Fäßler, F., Hansen, J. M., ...., & Schur, F. K. M. (2024). Lift-out cryo-FIBSEM and cryo-ET reveal the ultrastructural landscape of extracellular matrix. *Journal of Cell Biology*, 223(6).

### Liquid-Ethane Plunge-freezing







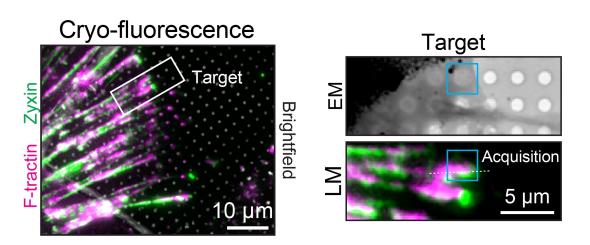
0 μs 500 μs 1000 μs

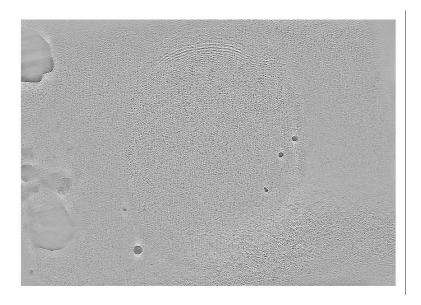
- One sided blotting does not disturb the sample on the grid (Only method on Leica GP)
- Blotting from one side means the liquid has to move through the layer of carbon using capillary action
- The entire sample needs to cool within ~10-20 ms to ensure that ice is vitreous (not crystalline)
- Highly dynamic cooling
  - Liquid ethane touches bottom of grid before top
  - Movement of liquid ethane?
  - Kinetics of heat transfer within cellular sample?
  - Kinetics of transport of heat within liquid ethane?
  - Grids bars as radiators or just extra mass?

Kasas, S., Dumas, G., Dietler, G., Catsicas, S., & Adrian, M. (2003). Vitrification of cryoelectron microscopy specimens revealed by high-speed photographic imaging. Journal of Microscopy, 211(1),

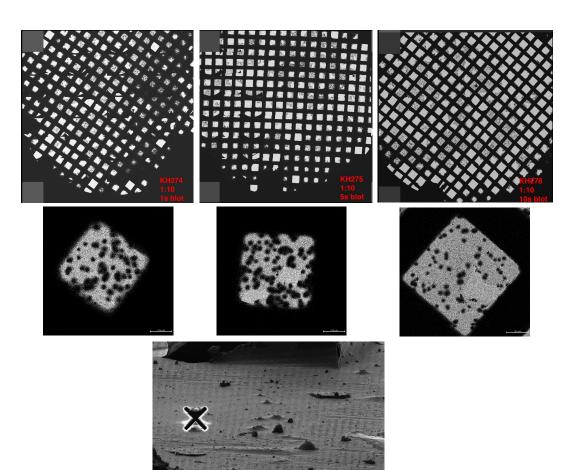
# Test Case: Plunge-Freezing with No Milling

- If area of interest is thin enough to be transparent without milling
  - Will ice will be vitreous after plunge freezing?
  - Positioned in center of grid square?
  - Blot time?
- Blotted 7s and was able to acquire good tomograms in edge of the cell

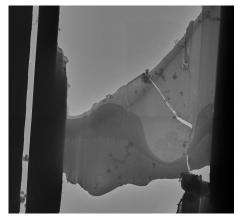




# Test Case: Unpatterned Bacteria for Plunge-Freezing



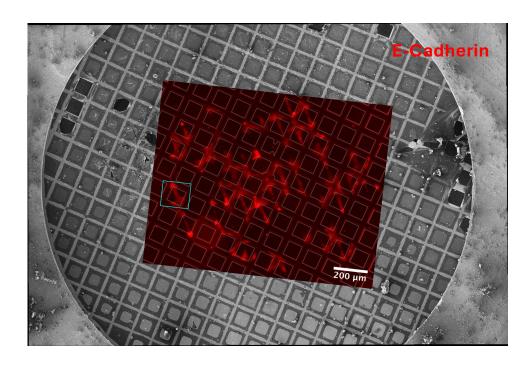
- Bacterial samples were added and different blotting times tested
  - No patterning, just plasma cleaning and direct addition
  - No cryoprotectant
  - 10s blotting removed almost all the surrounding buffer
- Cells visible in the SEM as mounds
- All the cells were vitreous
- Where buffer remained, some non-vitreous ice

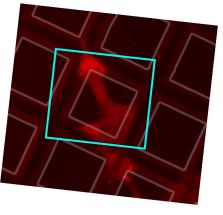




Test Case: Patterned Stripes of Cells for Plunge-

Freezing

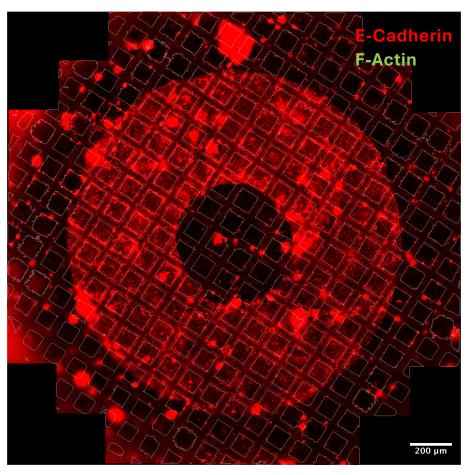


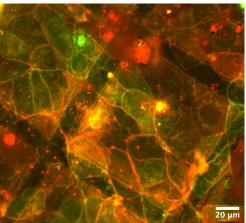


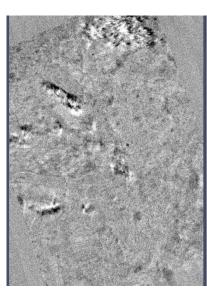


- "Stripes" of fibronectin were patterned with PRIMO, and EPH4 (mouse mammary epithelial) cells were seeded and allowed to grow 24h
- Plunge-frozen on Leica GP
  - 15s blot time
- Cells had non-vitreous ice, although tomogram reconstructions were usable
- Not physiologically relevant, cells didn't form mature cell:cell junctions

# Test Case: Patterned "Donuts" of Cells for Plunge-Freezing





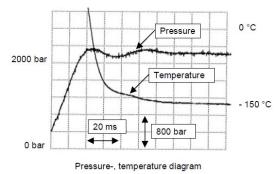


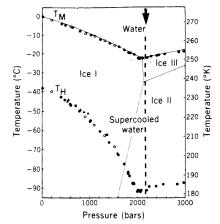
- Patterned a "donut" to let the EPH4 cells grow in a sheet, while still allowing the water the drain
  - Added 0.05% detergent (FOM) prior to plunge freezing on GP
  - Blotted 15s
  - $\sim$ 0.5  $\mu$ m water left on top of the cells
- Inside of cells was not vitreous!

### High Pressure Freezing



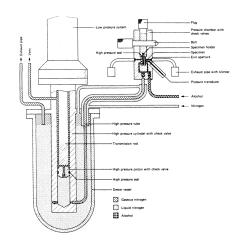
- Increase the pressure to ~ 2000 Mbar prior to freezing
- This slows the formation of crystalline ice
  - · Increases viscosity of water
  - Prevents the expansion needed to create crystalline ice
- Put the sample between two brass "planchettes"
- Planchettes are squeezed by high-pressure LN<sub>2</sub>, cushioned by a layer of alcohol
- After pressurizing (~15 ms), the alcohol is pushed out and the LN2 directly contacts the planchettes
- Cools at  $\sim 10^3$ - $10^4$  degree/s





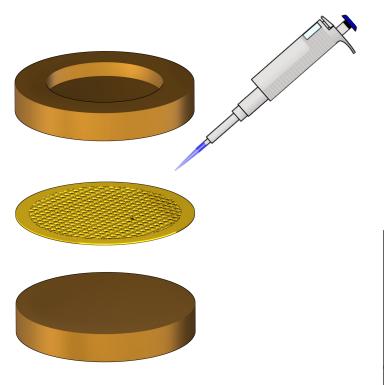




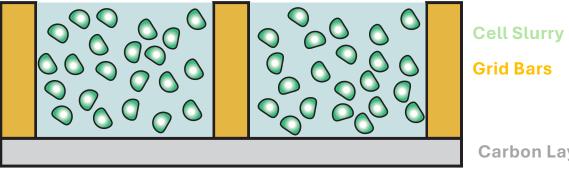


Dahl, R., & Staehelin, L. A. (1989). High-pressure freezing for the preservation of biological structure: Theory and practice. Journal of Electron Microscopy Technique, 13(3), 165–174

### High-pressure freezing – Suspended Sample



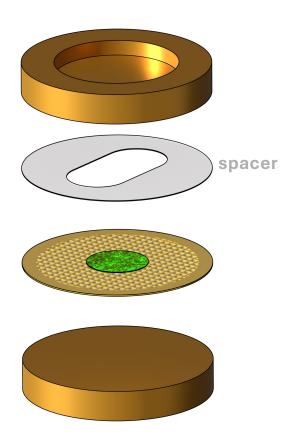
- Concentrate sample with cryoprotectant into a "slurry"
- Deposit sample onto the upside-down grid
  - Space between bars act as small wells
  - Gold grids ~ 25 µm thick
- Sandwich between two brass planchettes
  - Polished and 1-hexadecene coated to facilitate removal
- High-pressure Freeze
- Sample should be everywhere, can mill top part with "Waffle Method"



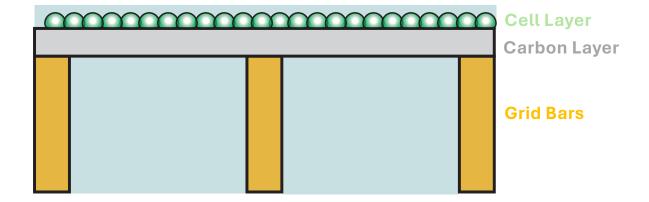
**Carbon Layer** 

Kelley, K., Raczkowski, A. M., ..., G., Potter, C. S., Carragher, B., & Noble, A. J. (2022). Waffle Method: A general and flexible approach for improving throughput in FIB-milling. Nature Communications, 13(1), 1857

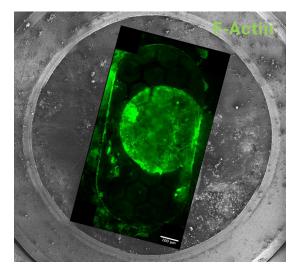
### High-pressure freezing – Adhered Sample

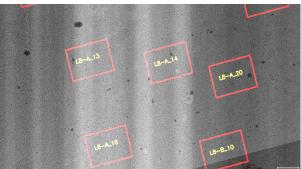


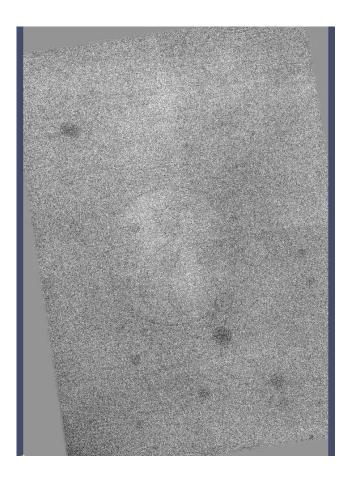
- · Grow cells directly on the grid
- Add an aperture grid as a spacer to keep the cells from being squeezed
  - Copper spacer: ~ 35 μm
  - gold spacer: ~ 25 μm
  - Steel spacer ~ 10 µm
- Sandwich between two brass planchettes
  - Polished and 1-hexadecene coated to facilitate removal
- High-pressure Freeze
- Thickness of the media layer is dependent on the spacer type



# Test Case: Patterned Sheets of Cells for High-Pressure Freezing



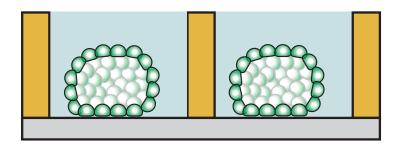


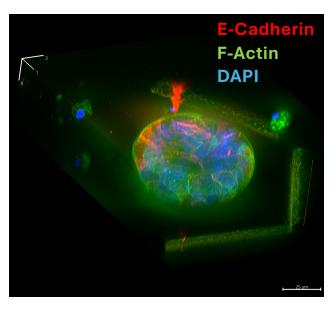


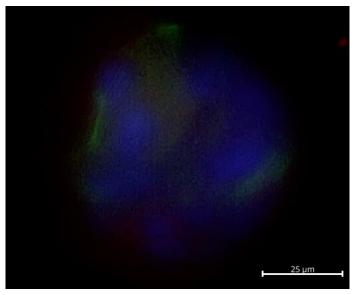
- Used high-pressure freezing to freeze a sheet of EPH4 cells
- Patterned a "button" to confine the cells within the spacer center
- Lamellae showed very little non-vitreous ice
- Tilt-series also had small to no areas non-vitreous ice

# Test Case: Confined Cysts for High Pressure Freezing

- Used the "fill in the grid wells" method to grow cysts of epithelial cells (MDCK)
- PRIMO patterned 1 mm circle, added PLL and 2% matrigel
- Use larger hexagon shaped grids (100 mesh)
- Fixed some grids to make sure they grew properly
- High-pressure froze the rest
- Currently imaging on CLEM in preparation for FIB-milling







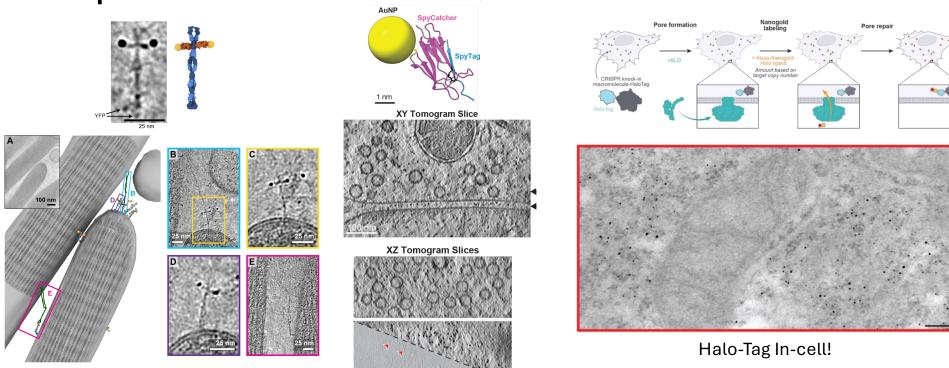
#### Labeling targets for CLEM and TEM

- Labeling with a fluorescent tag allows the localization of the target molecules within the sample
  - Bleaching is not much of an issue at cryogenic temperatures
  - Membranes (especially multimembrane vesicles) auto-fluoresce at cryogenic temperatures, strongest in GFP channel
  - Can't localize more than ~0.1 um area
- Labeling protein of interest with a small (~3-5 nm) gold nanoparticle (AuNP) allows precise localization within the tomogram
  - Not easy to get nanoparticles inside of cells, so normally this is limited to extracellular targets
  - Need to conjugate the AuNP to a molecule that would bind to the target
- Other methods include genetically encoded multimeric tags (GEMS) or Ferritags to label inside of cells

AuNP labeling for Localization in Tomograms:

Examples

FAB-conjugated AuNP



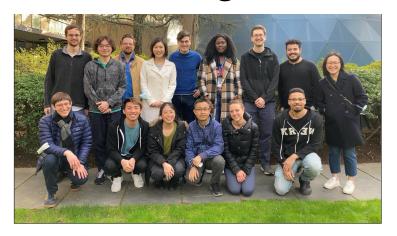
Elferich, J., Clark, S., ..., Gouaux, E. (2021). Molecular structures and conformations of protocadherin-15 and its complexes on stereocilia elucidated by cryo-electron tomography. eLife, 10, e74512.

Held, R. G., Liang, J., ..., Azubel, M., & Brunger, A. T. (2024). In-Situ Structure and Topography of AMPA Receptor Scaffolding Complexes Visualized by CryoET. bioRxiv 2024.10.19.619226.

Young, L. N., ..., Rosen, M. K., Giraldez, A. J., & Villa, E. (2024). ExoSloNano: Multi-Modal Nanogold Tags for identification of Macromolecules in Live Cells & Cryo-Electron Tomograms. bioRxiv 2024.10.12.617288

Spycatcher

#### **Acknowledgements**





SCIENCE FOR THE BENEFIT OF HUMANITY



#### **Alushin Lab:**

Gregory Alushin
Ayala Carl
Alfred Chin
Rui Gong
Joe Levin
Donovan Phua
Matt Reynolds
Ariel Pan
Xiaoyu Sun
John Watters
Opeoluwa Olaniyan
Michael Barrett
Lily Sutton
lan Windham

#### **CEMRC:**

Mark Ebrahim Johanna Sotiris Honkit Ng

#### PIT:

Peer Strogies
James Petrillo

#### **EMRC**:

Hilda Amalia Pasolli

#### CFC:

Ruby Room Teachers Blue Room Teachers General Staff

#### **Funding:**





ΙΔΡΥΜΑ ΣΤΑΥΡΟΣ ΝΙΑΡΧΟΣ STAVROS NIARCHOS FOUNDATION