

Time-resolved Cryo-EM

Jack Fu

Joachim Frank's lab

Columbia University

Questions to address?

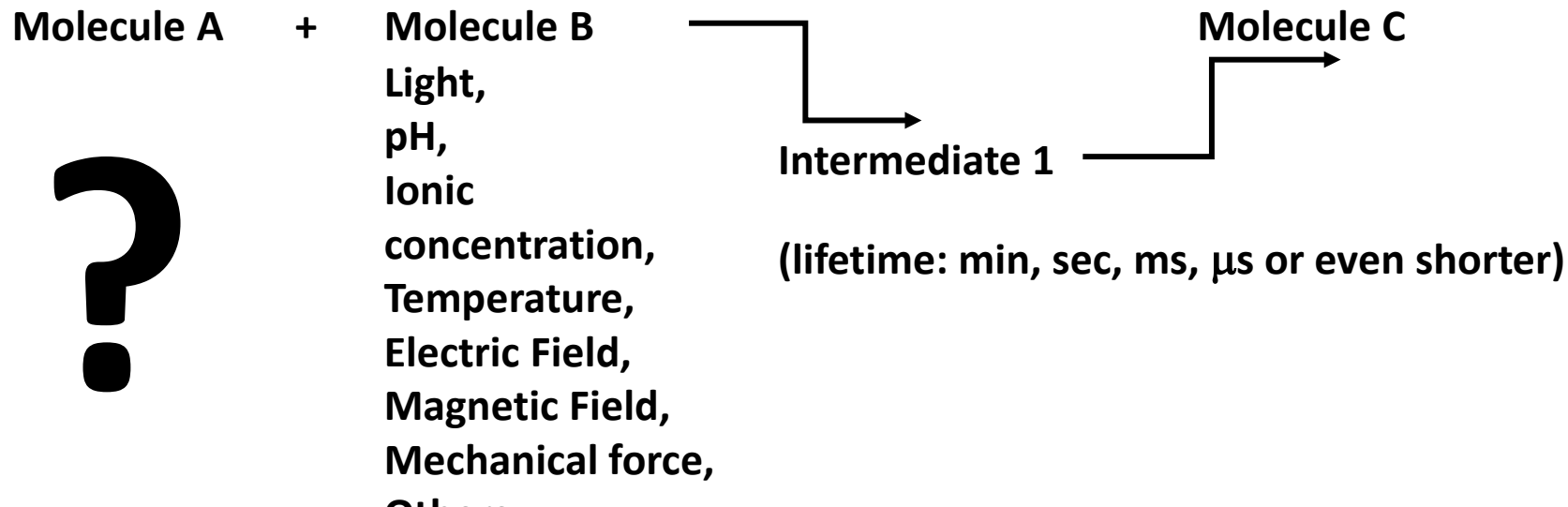
- ~~How can time-resolved cryo-EM help you in your research?~~
- **We need your help.**
- ~~What are the obstacles to success?~~
- **There are a lot of issues in time-resolved Cryo-EM method.**

Time-resolved cryo-electron microscopy

- Time-resolved cryo-electron microscopy (cryo-EM) combines the known advantages of **single-particle cryo-EM in visualizing molecular structure** with the ability **to dissect the time progress of a reaction between molecules in vitro.**

Time-resolved cryo-electron microscopy

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What has been tested?

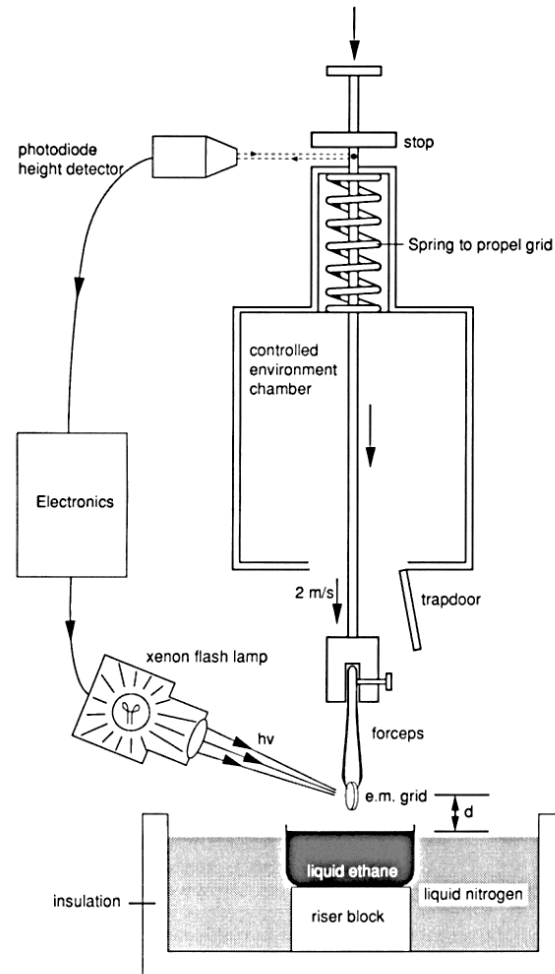
The EMBO Journal vol.12 no.1 pp.1 – 8, 1993

Electron diffraction analysis of structural changes in the photocycle of bacteriorhodopsin

**Sriram Subramaniam^{1,2}, Mark Gerstein¹,
Dieter Oesterhelt³ and Richard Henderson¹**

¹MRC Laboratory of Molecular Biology, Hills Road, Cambridge CB2 2QH, UK, ²Department of Biological Chemistry, Johns Hopkins University School of Medicine, Baltimore, MD 21205, USA and ³Max-Planck Institut für Biochemie, D-8033, Martinsried, Germany

Communicated by R.Henderson

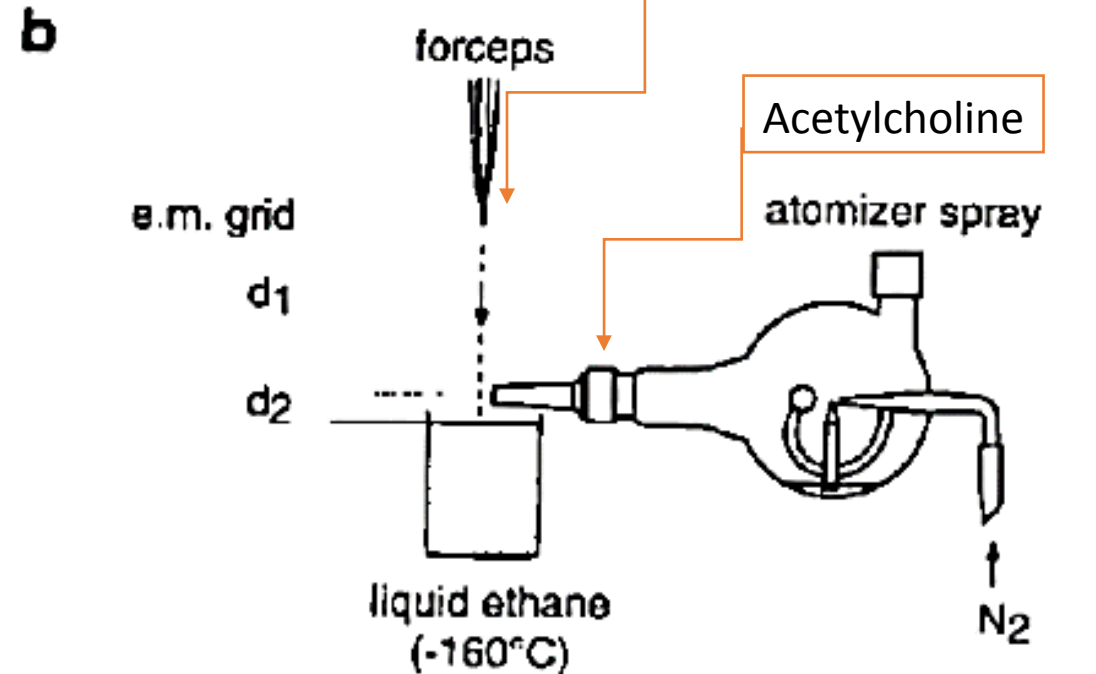
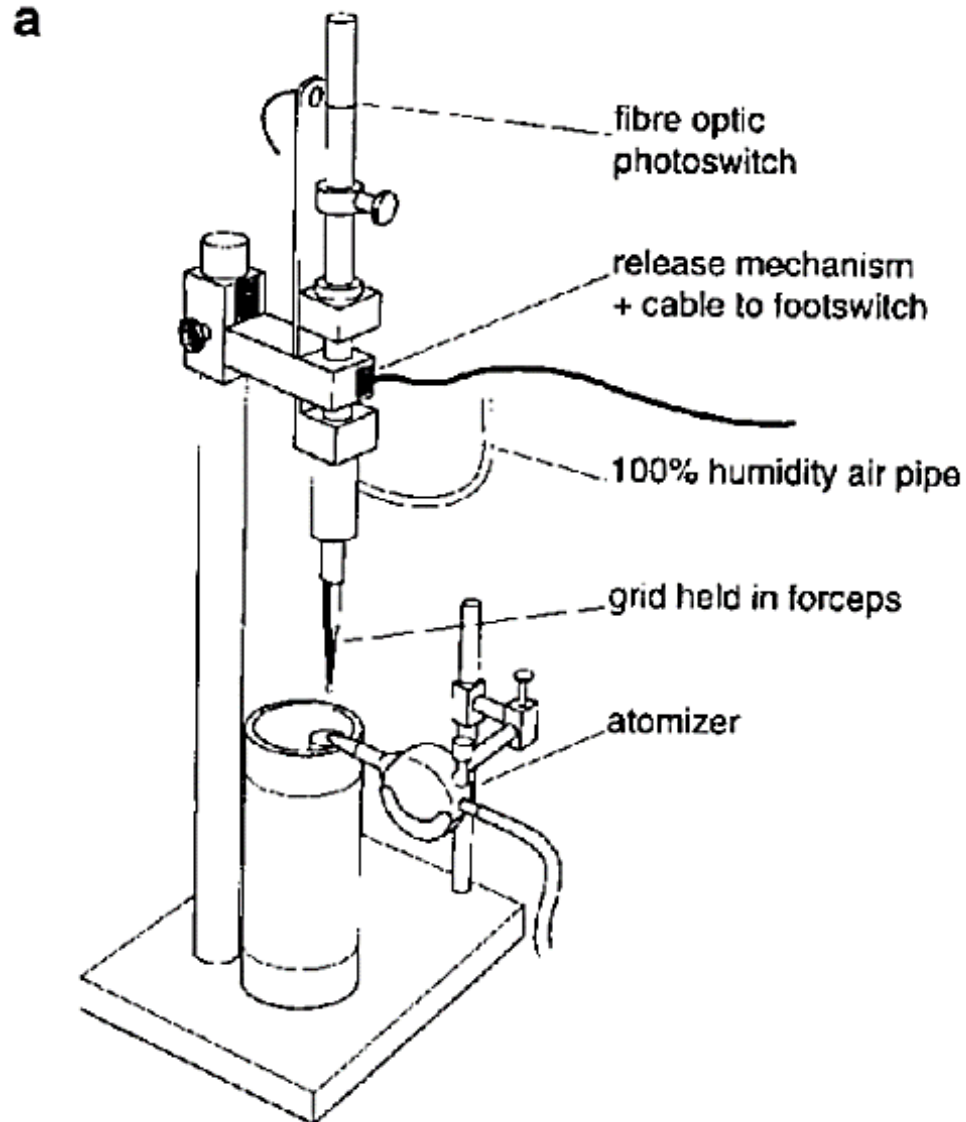


Analysis of transient structures by cryo-microscopy combined with rapid mixing of spray droplets

John Berriman, Nigel Unwin

MRC Laboratory of Molecular Biology, Hills Road, Cambridge, CB2 2QH, UK

Received 15 June 1994



Millisecond Time Resolution Electron Cryo-microscopy of the M-ATP Transient Kinetic State of the Acto-Myosin ATPase

M. Walker,* J. Trinick,* and H. White[‡]

***Muscle and Collagen Group, Bristol University Veterinary School, Langford, Bristol BS18 7DY United Kingdom, and [‡]Biochemistry Department, Eastern Virginia Medical School, Norfolk, Virginia 23507 USA**

JOURNAL OF STRUCTURAL BIOLOGY **121**, 306–313 (1998)
ARTICLE NO. SB983968

A Computer-Controlled Spraying-Freezing Apparatus for Millisecond Time-Resolution Electron Cryomicroscopy

H. D. White

Department of Biochemistry, Eastern Virginia Medical School, Norfolk, Virginia 23507

M. L. Walker and J. Trinick

Department of Human Biology, University of Leeds, Leeds LS2 9JT, United Kingdom

Received November 18, 1997, and in revised form January 26, 1998

Limitations in the spraying-freezing method

Molecule A	+	Molecule B
On Cryo-EM grid	+	Small molecule Acetylcholine/ ATP Marker to identify the droplets

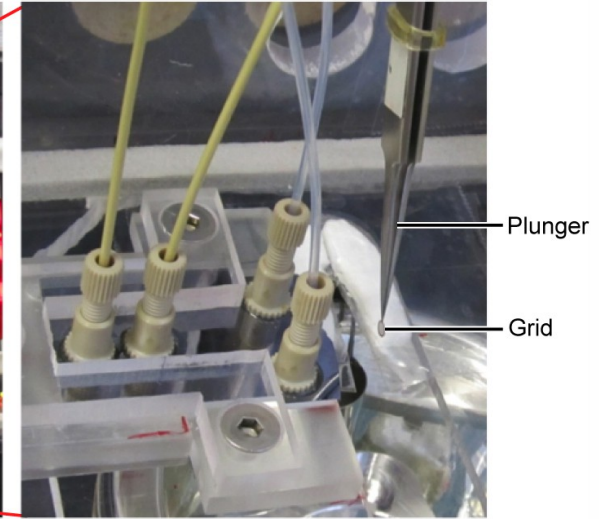
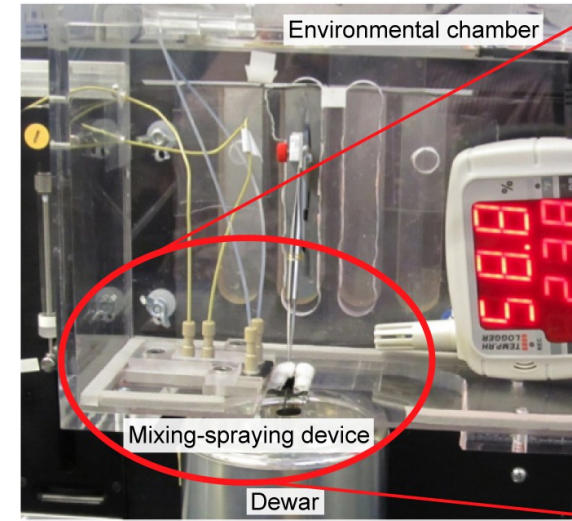
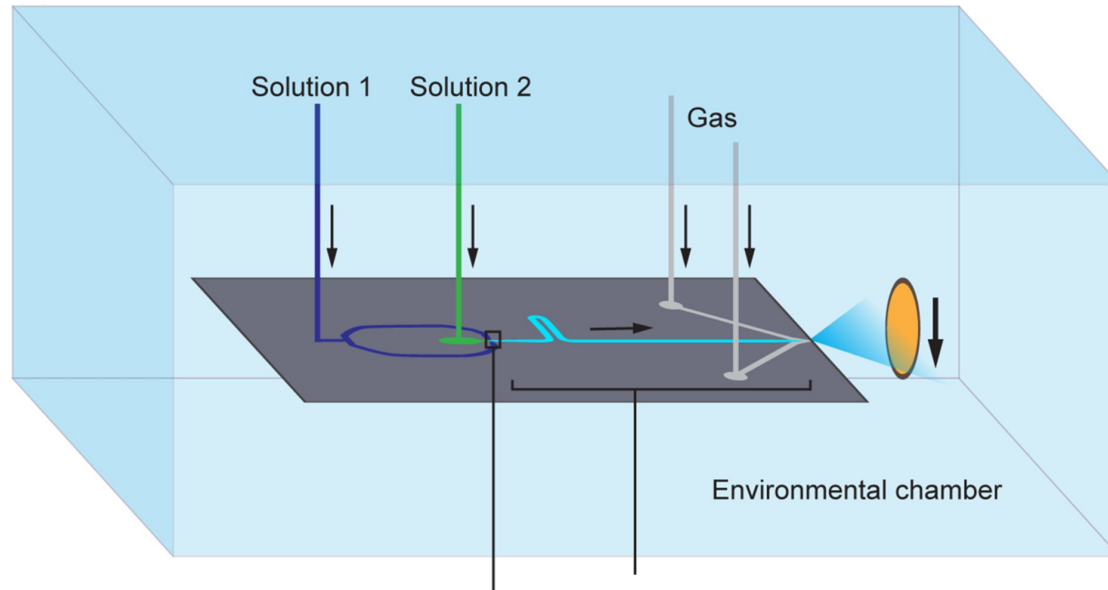
The mixing is dependent on diffusion

1. slow

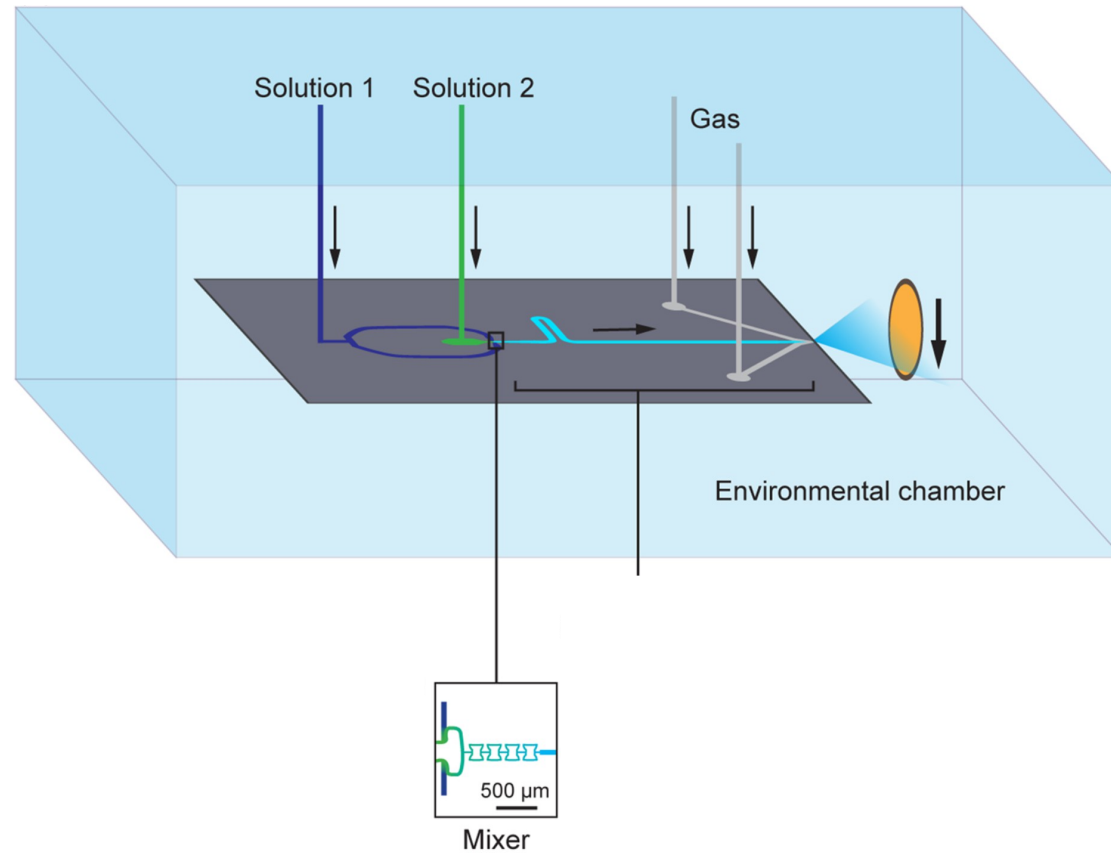
2. dependence on molecular weight

Our solution is the mixing-spraying-freezing method

Experimental setup – Microfluidic chip



Experimental setup – Mixer performance

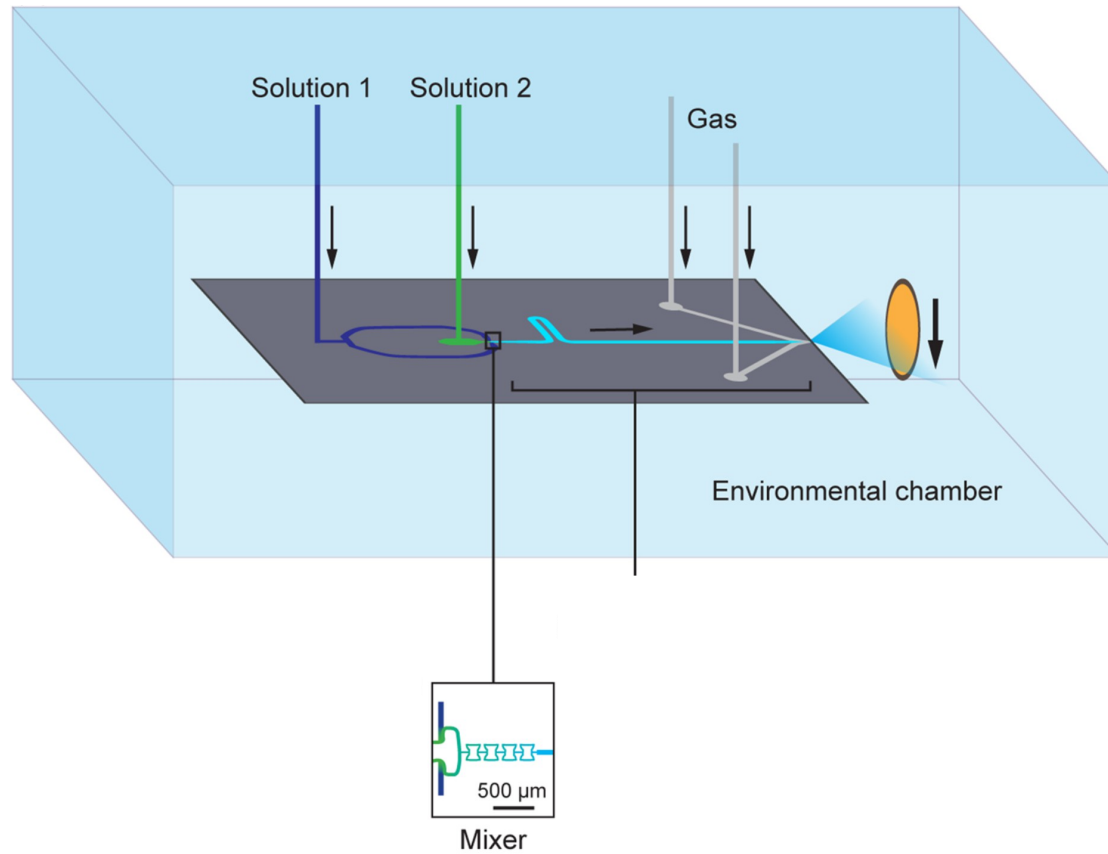


Not mixed well Flow rate: 1 $\mu\text{L/s}$

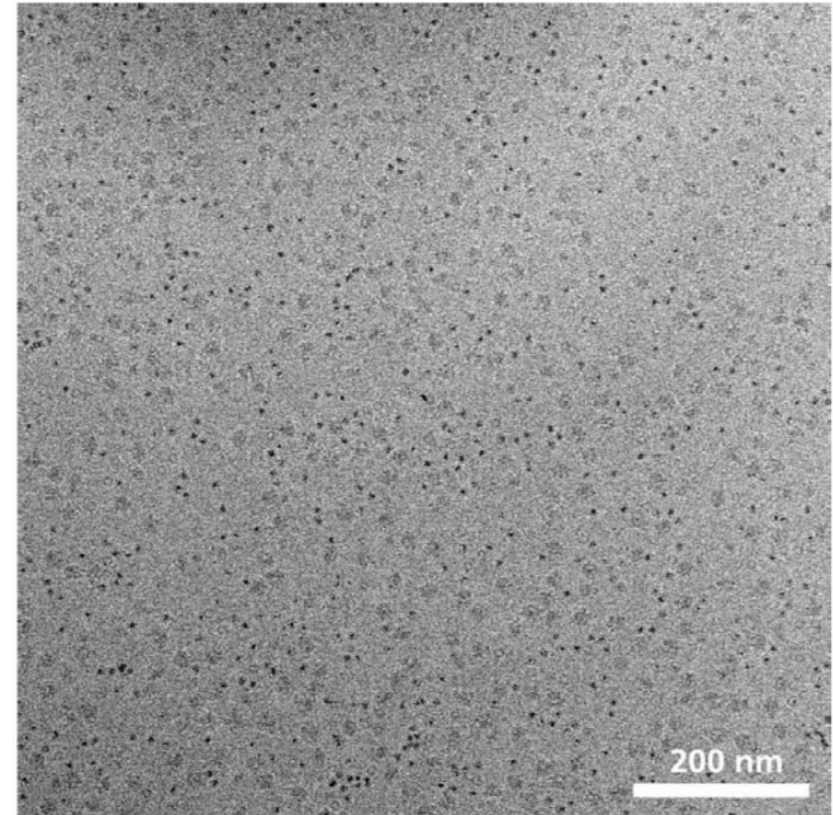


Very well mixed Flow rate: 6 $\mu\text{L/s}$

Experimental setup – Mixer performance

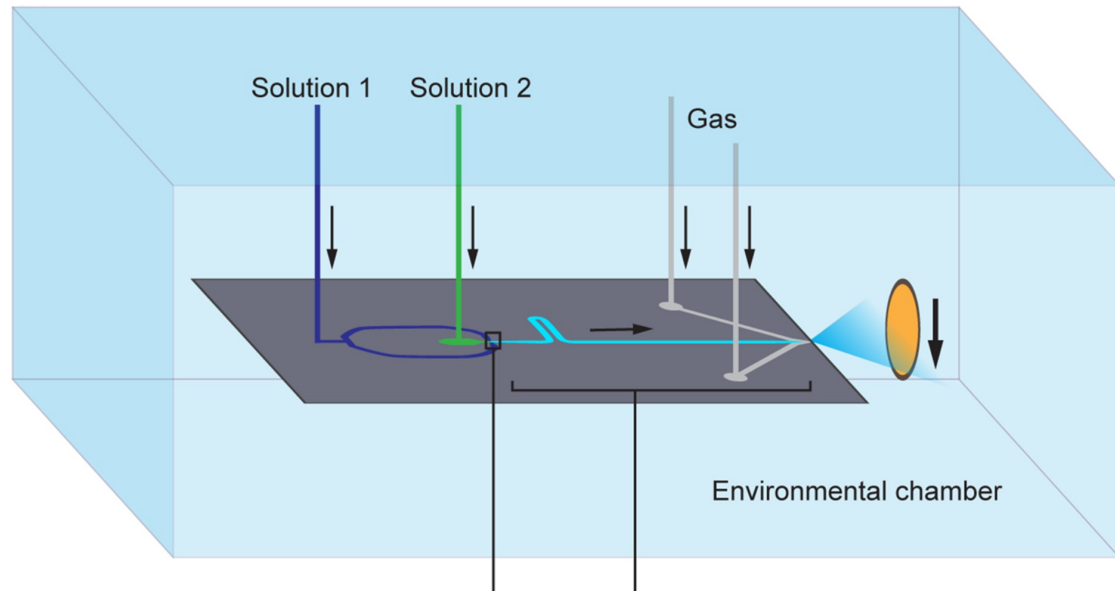


Mixing time 0.5 ms

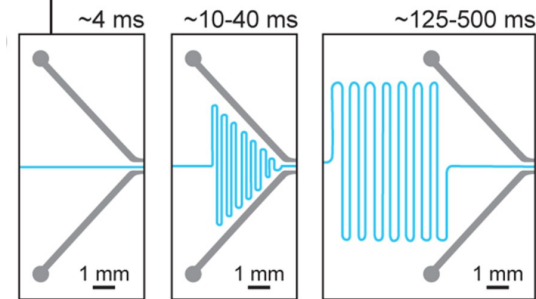


Mixing 70S ribosomes and Ferritin molecules

Experimental setup – Reaction time



Length of the reaction channels and the flow rate determine the reaction time in the microfluidic chips.

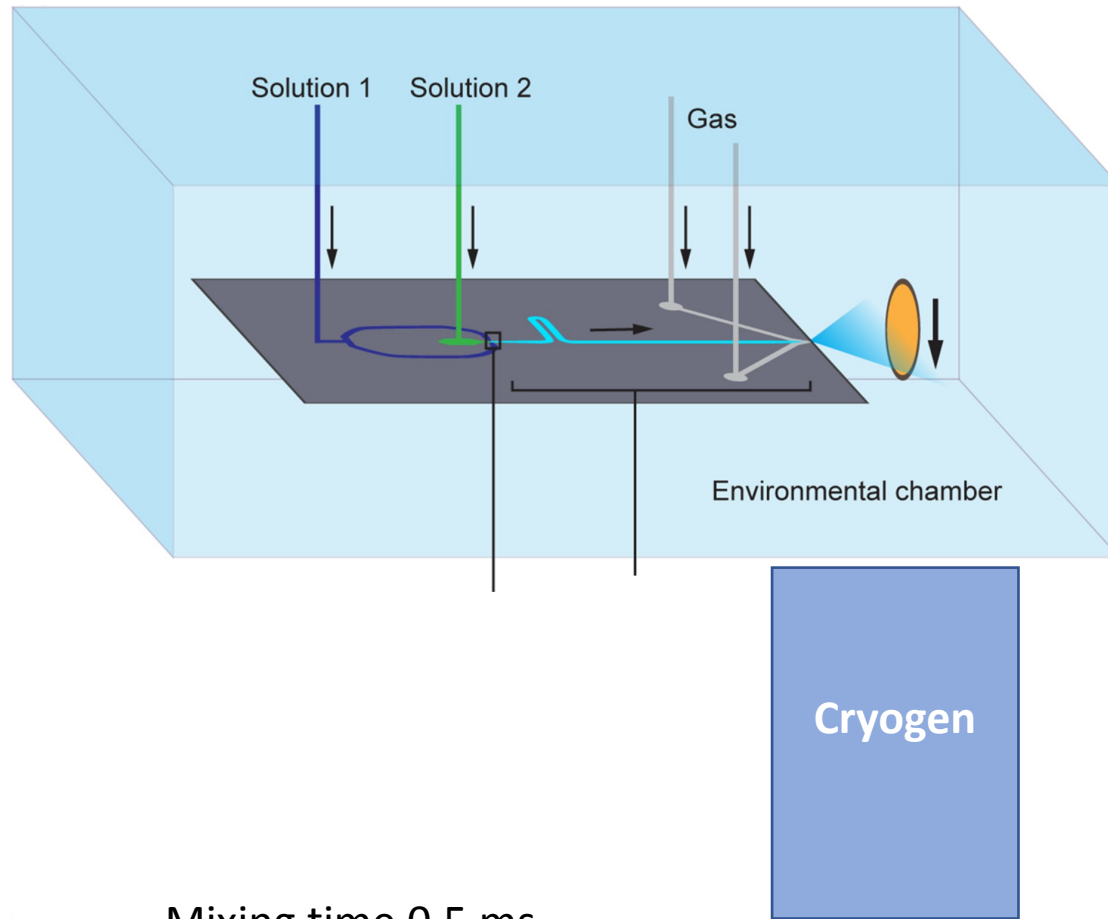


Different Reaction Channels

Mixing time 0.5 ms

Reaction time 4-500 ms

Experimental setup – Plunging and freezing



Reaction is stopped by plunging into cryogen.

Mixing time 0.5 ms

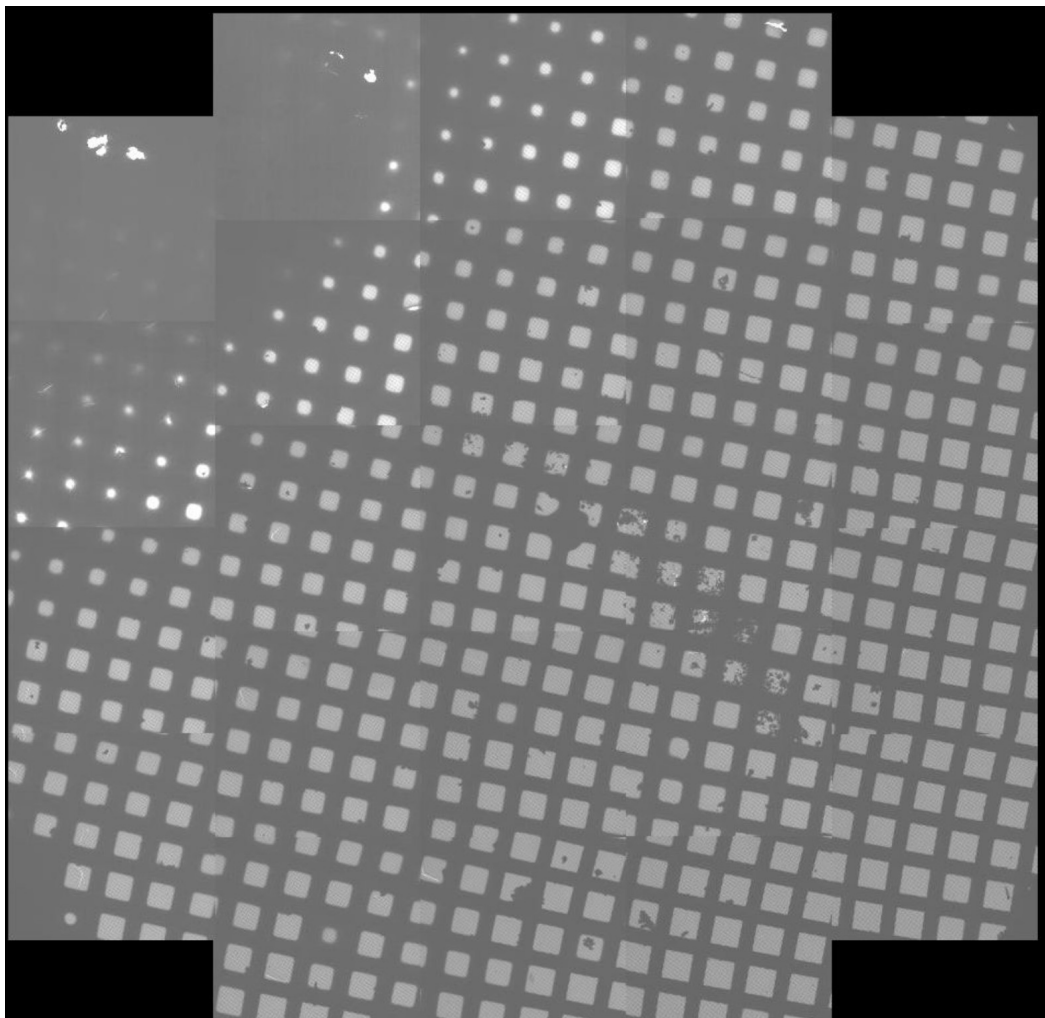
Reaction time 4-500 ms

Plunging time 18 ms

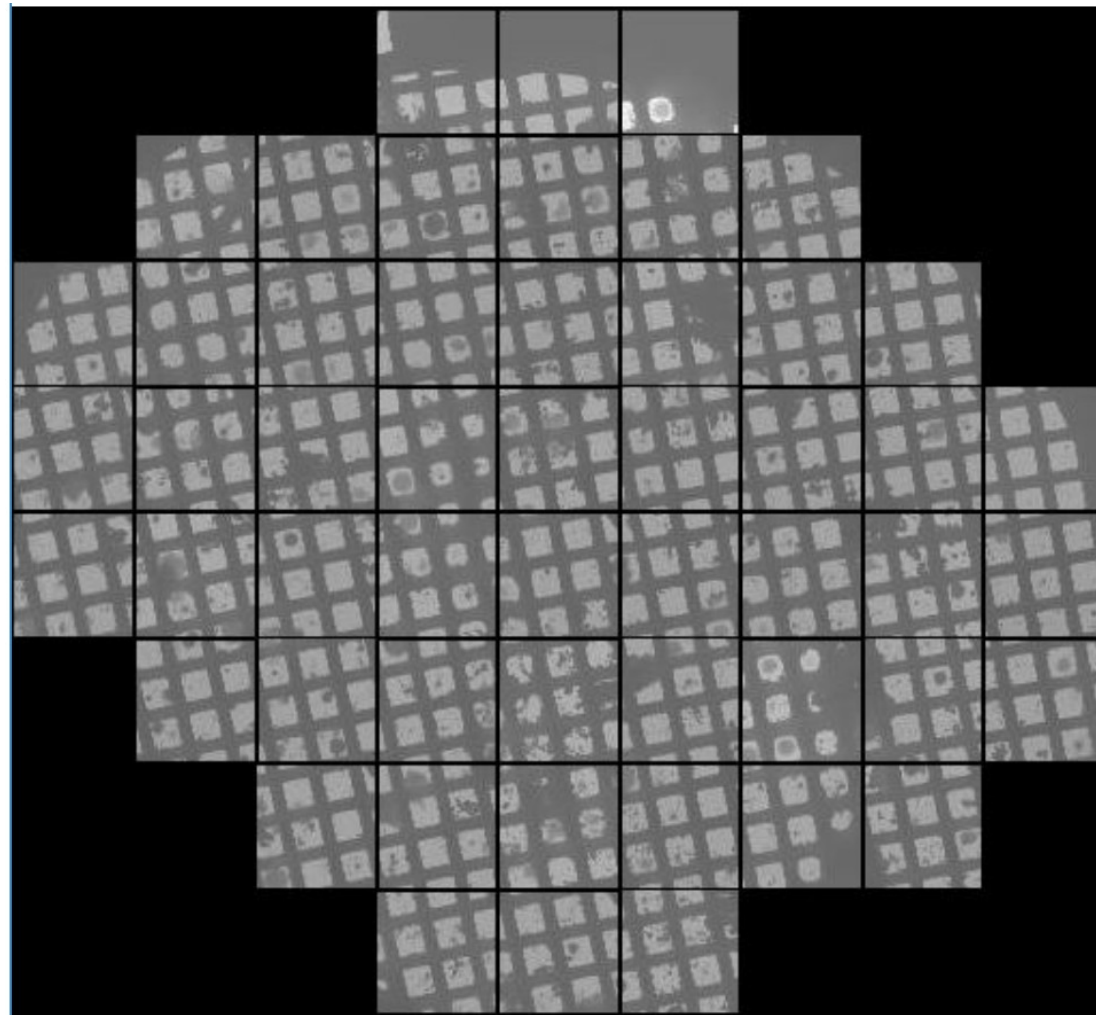
Limitation

- 1. How to get right ice thickness?

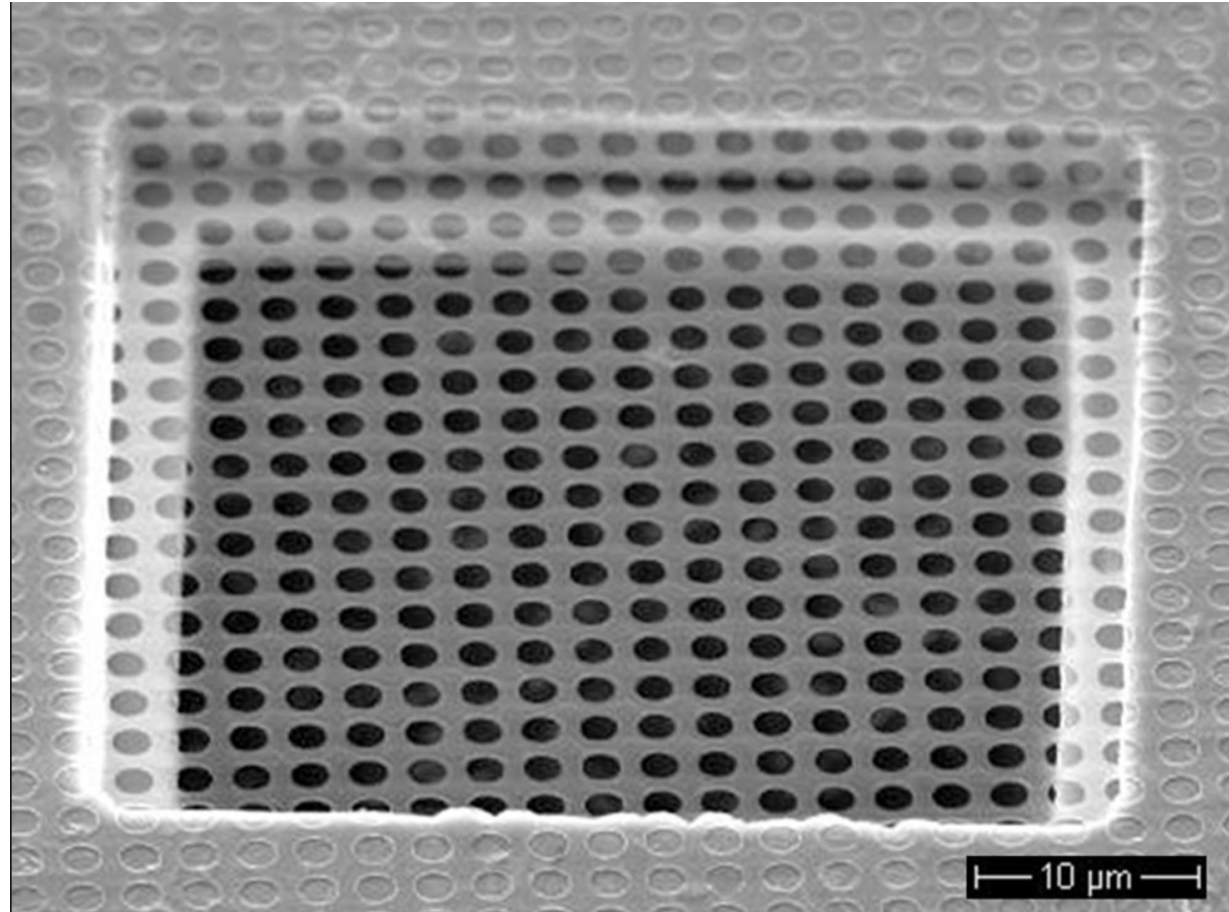
Blotting grid



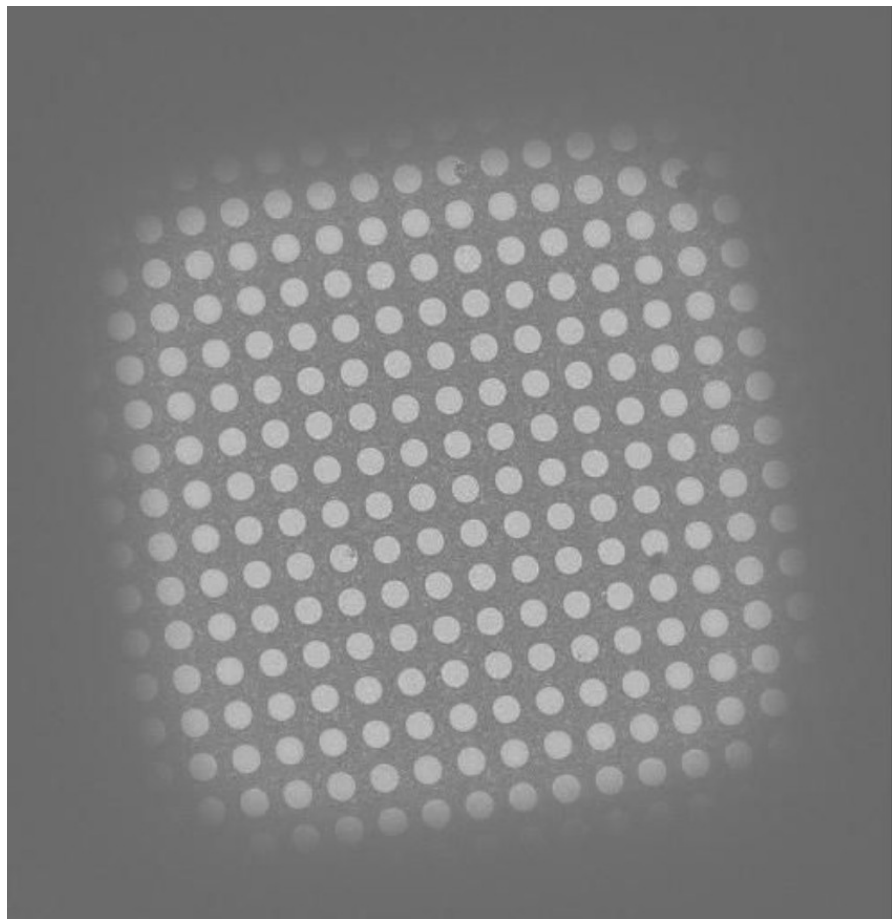
Spraying grid



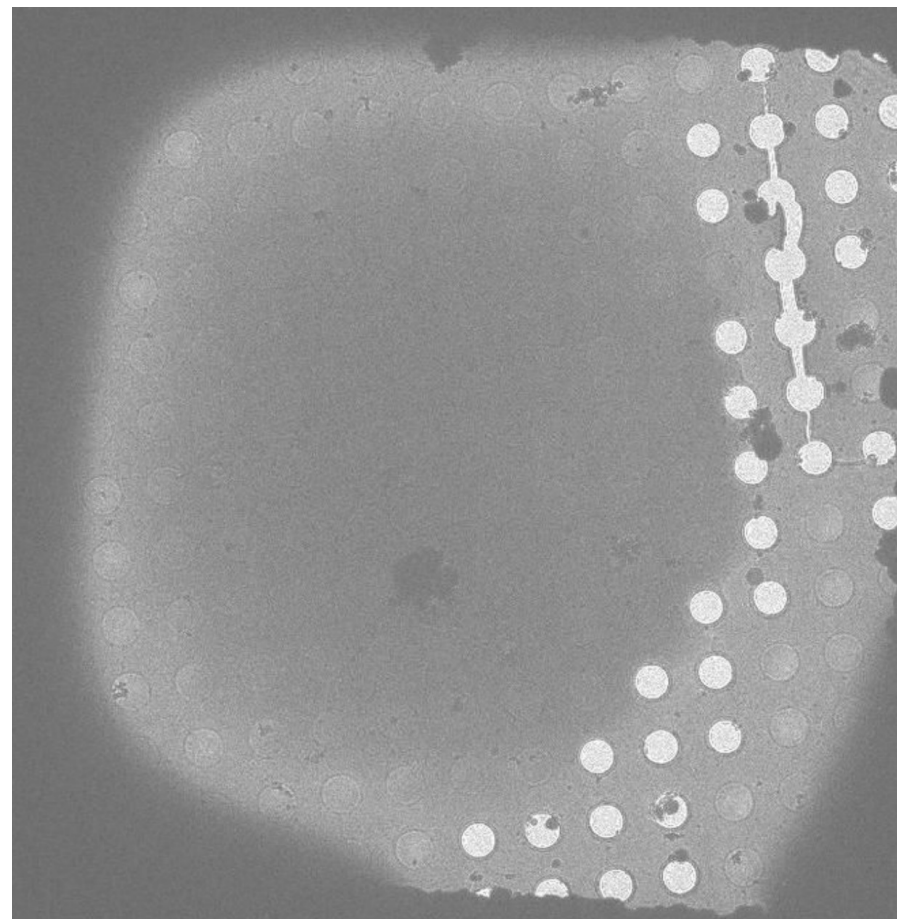
Quantifoil R1.2/1.3 400 mesh



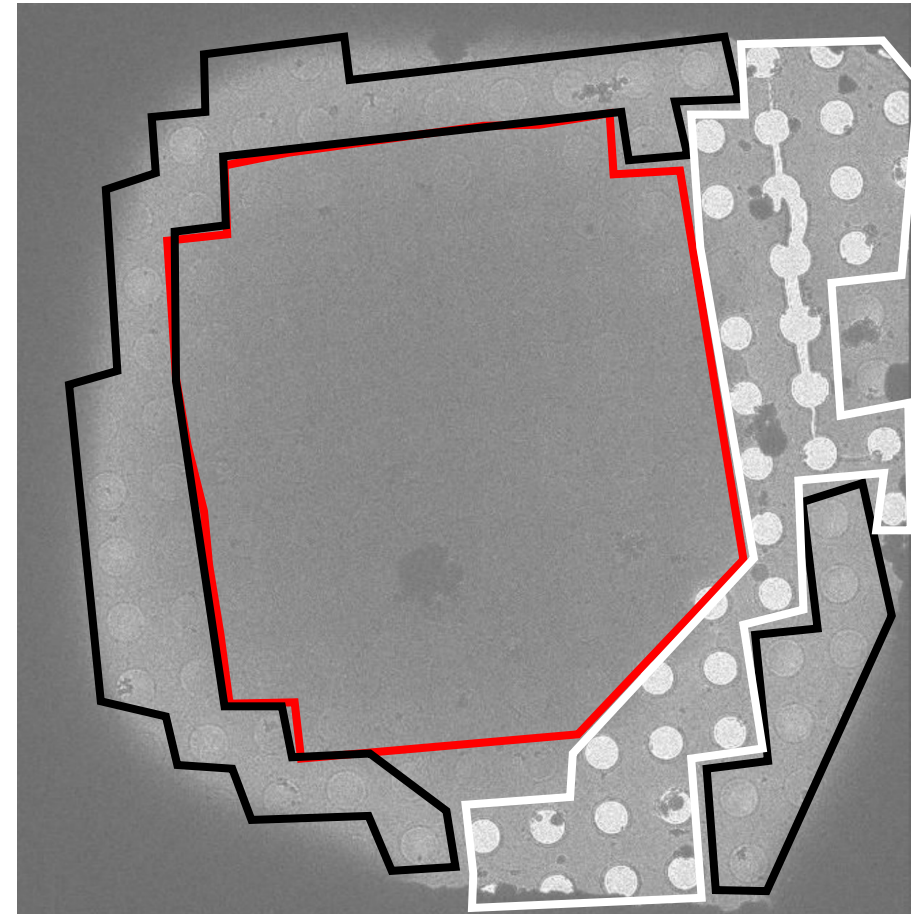
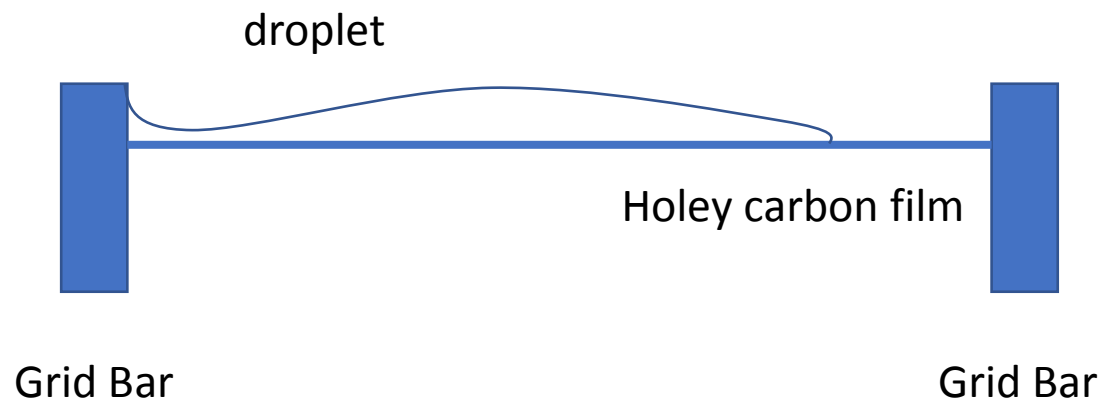
Blotting grid



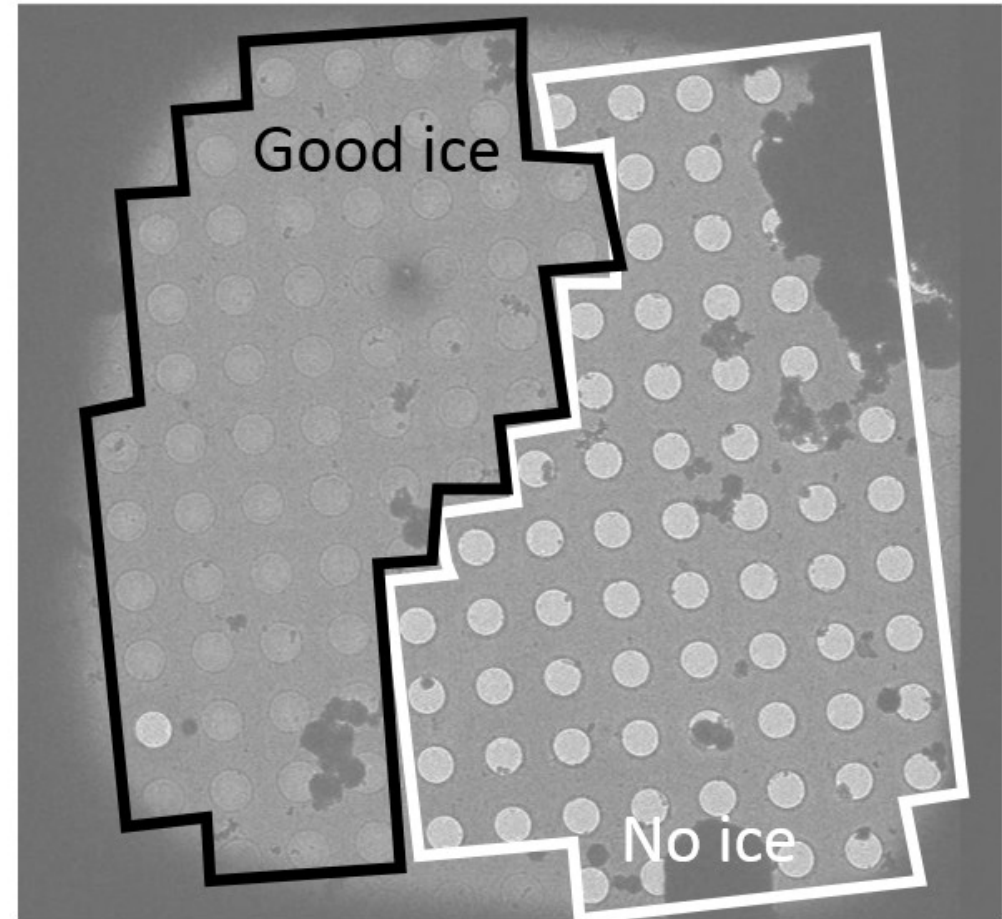
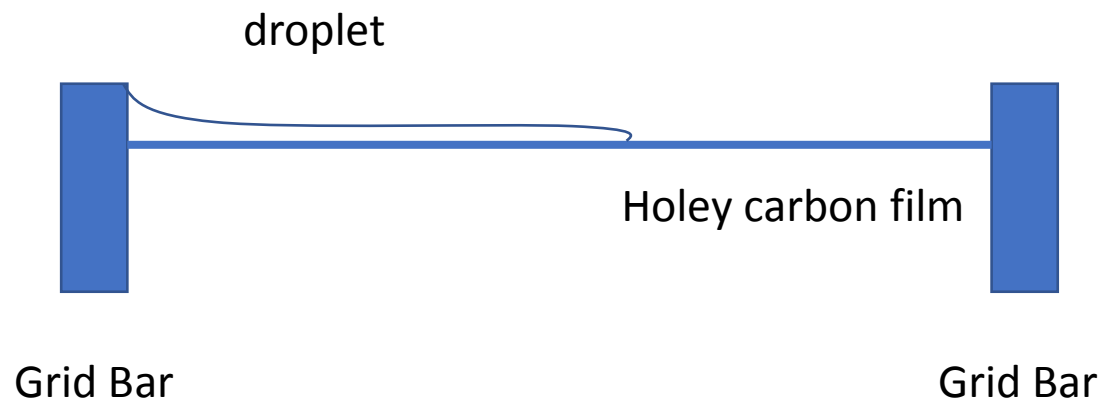
Spraying grid



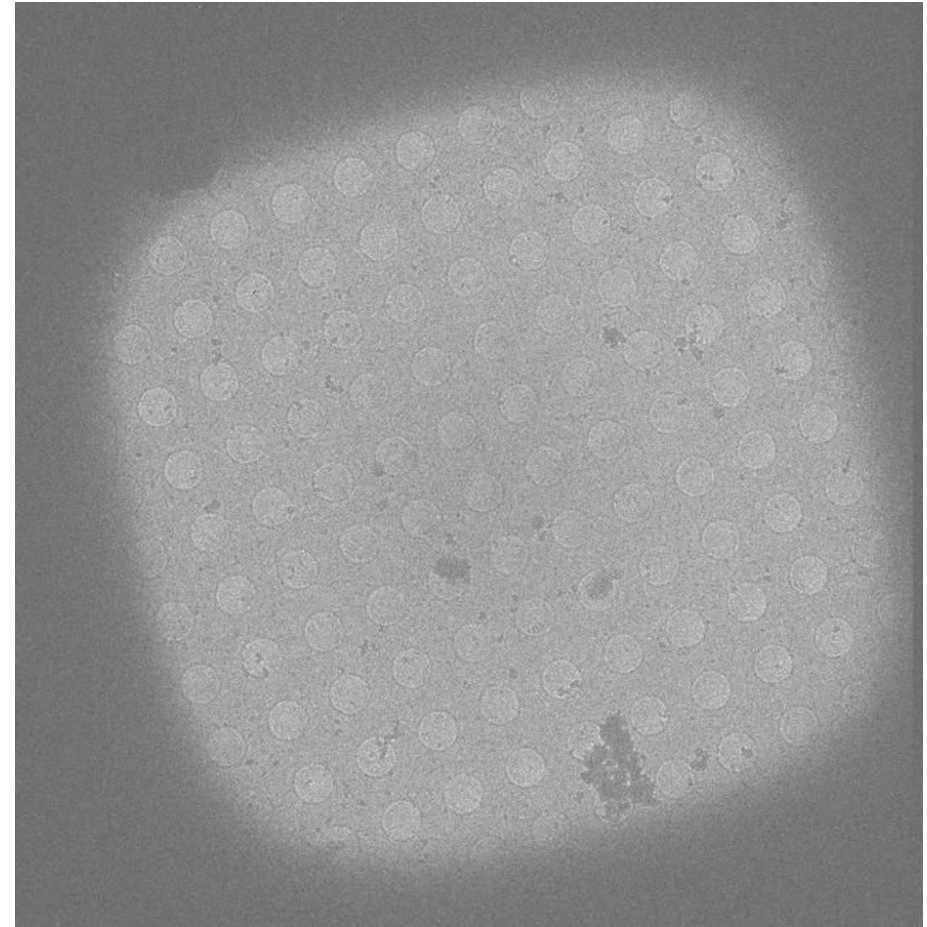
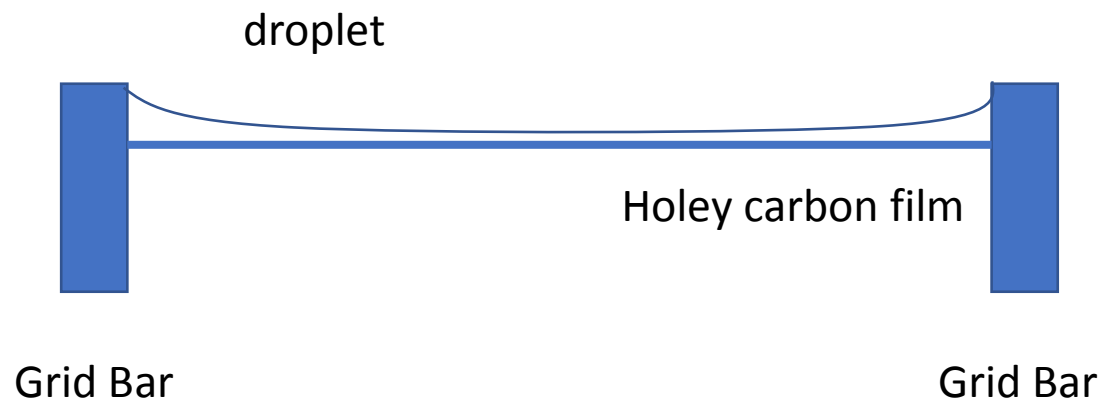
Spraying grid



Spraying grid

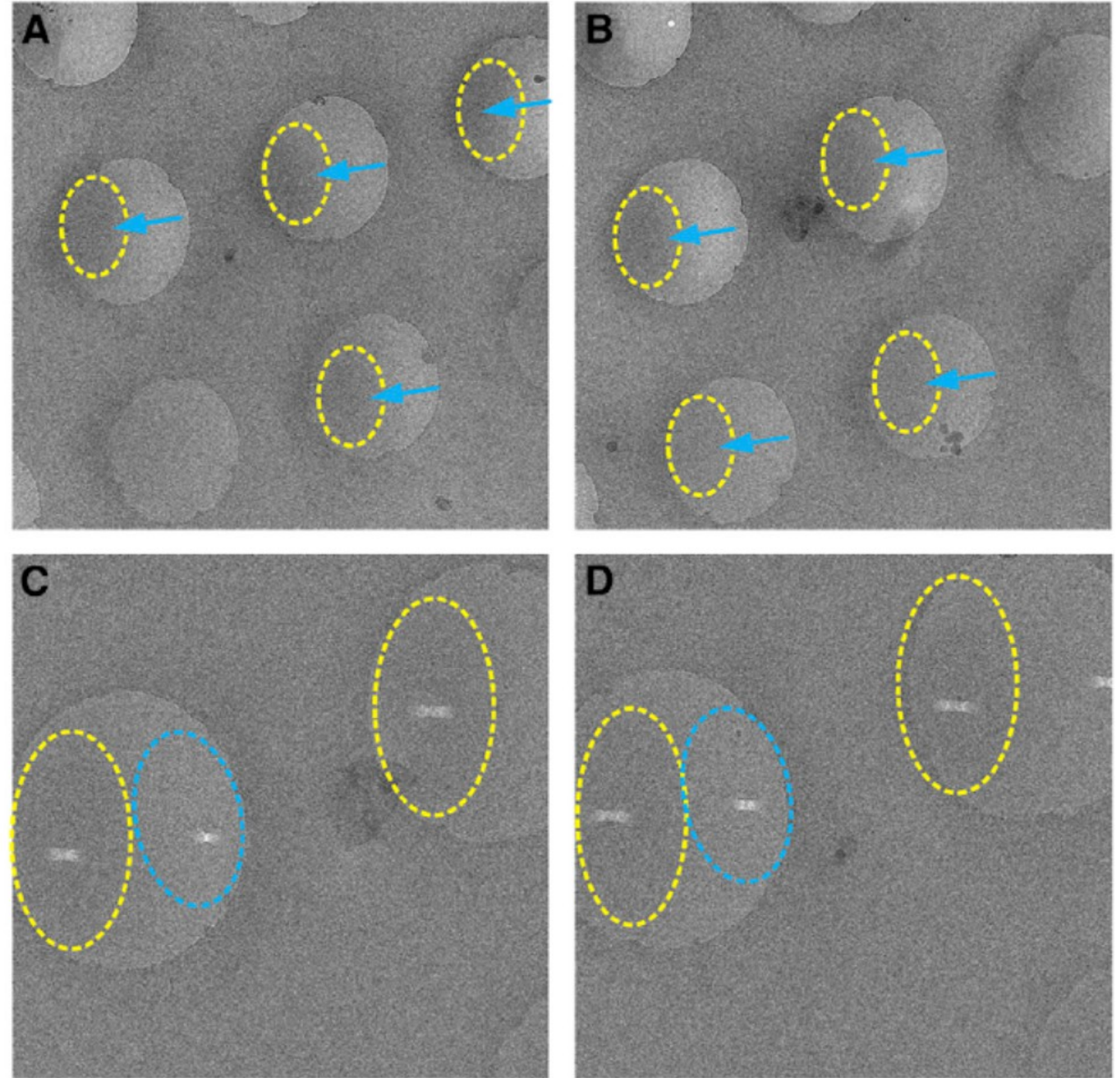


Spraying grid

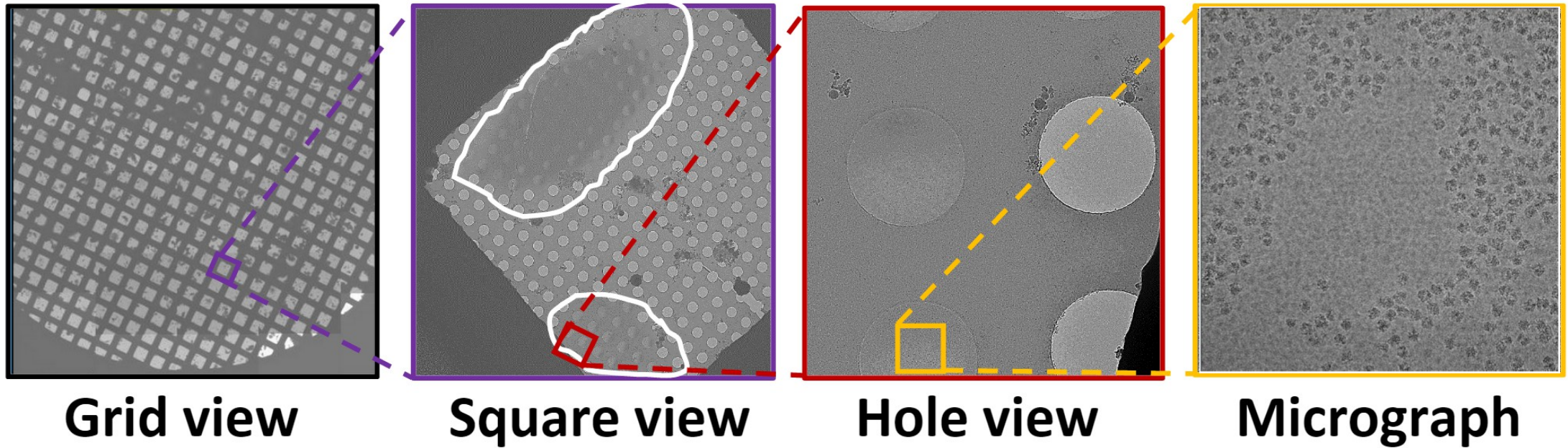


(A and B) The ice thickness is different from the leading to the trailing side of each hole (blue arrows), which is different from grids obtained by the blotting method.

(C and D) The ice is thinner on one side than on the other side as indicated by the different lengths of the tunnels drilled on the two sides. The thicker



Data collection



Mean droplet size – flow rate ratio between liquid and gas

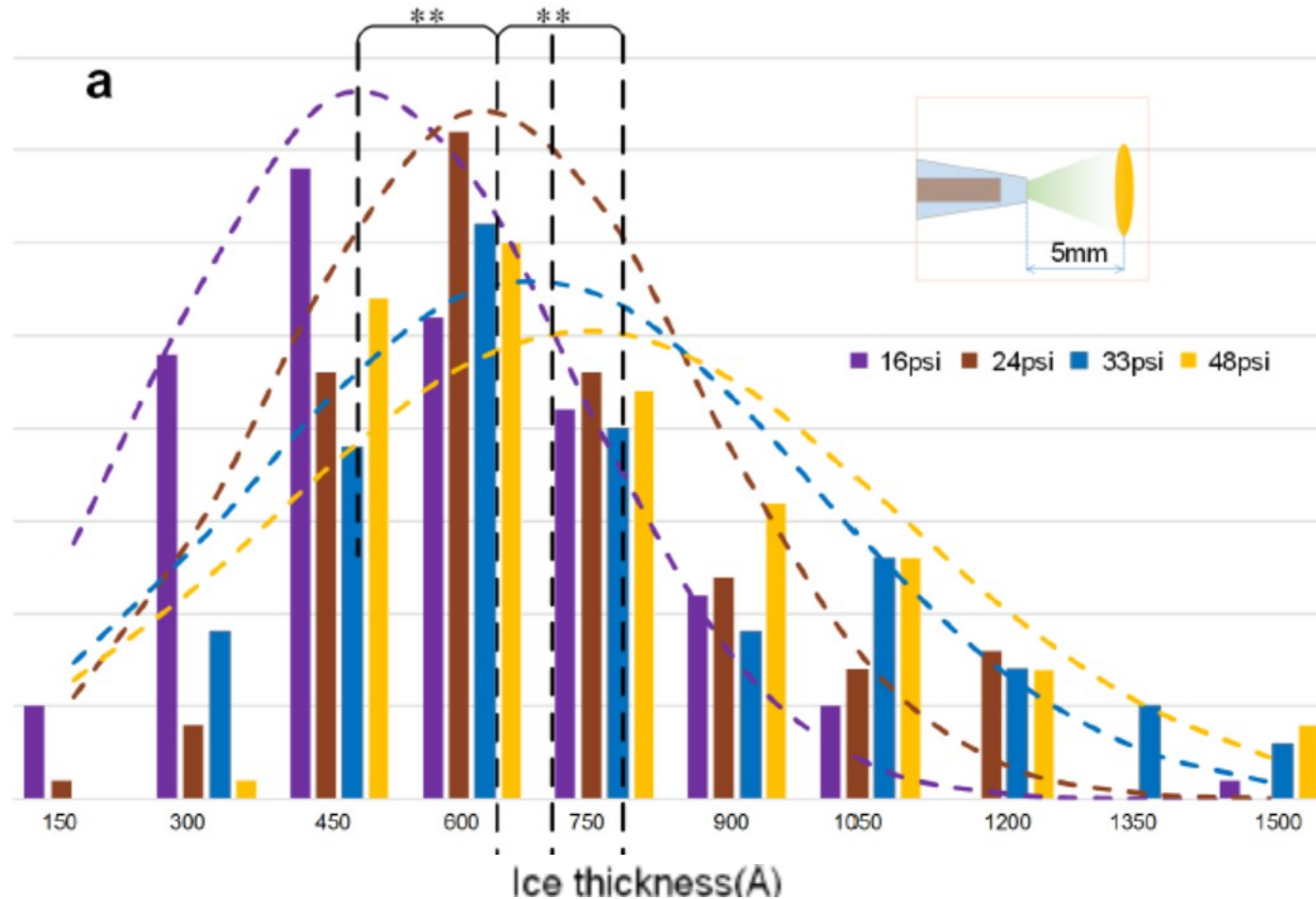
- Diameter: 36.2 to 4.4 μm (Volume : 24.4 pL - 0.044 pL)

Gas pressure: 16 psi to 48 psi

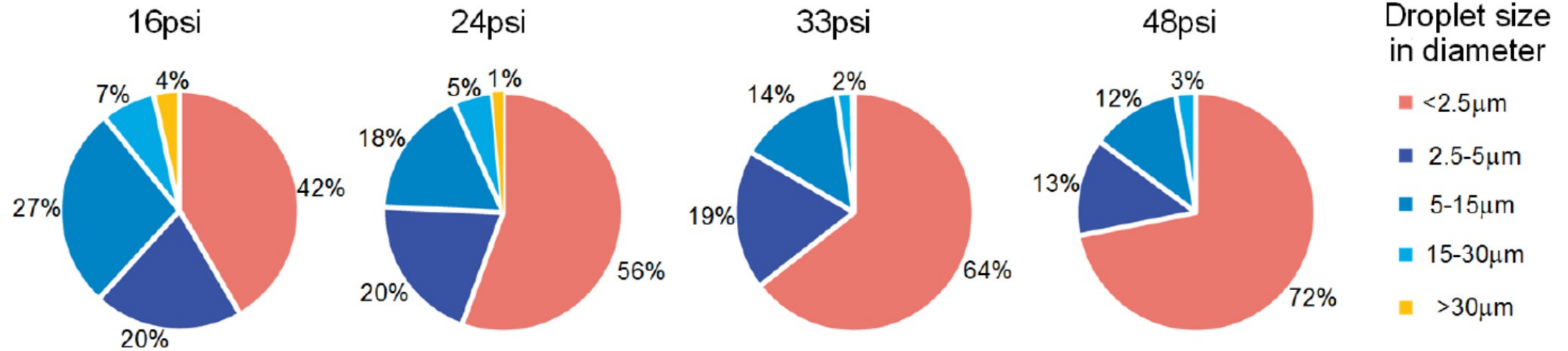
$$\text{SMD} = 0.95 \left[\frac{(\sigma_l \dot{m}_l)}{V_r \rho_l^{11/30} \rho_g^{3/10}} \right] \left[1 + \frac{\dot{m}_l}{\dot{m}_g} \right]^{17/10} + 0.13 \mu_l \left[\frac{D}{\sigma_l \rho_l} \right]^{1/2} \left[1 + \frac{\dot{m}_l}{\dot{m}_g} \right]^{17/10}$$

Where m is the mass flow rate, and subscripts g and l denote gas and liquid. Suppose that the solution sprayed is water, the viscosity μ , surface tension σ , density ρ are $0.89 \times 10^{-3} \text{ Pa}\cdot\text{s}$, 0.072 N/m , $1 \times 10^3 \text{ kg/m}^3$, respectively.

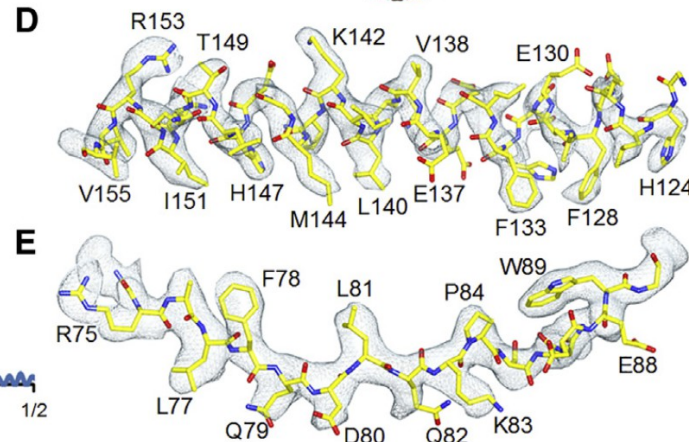
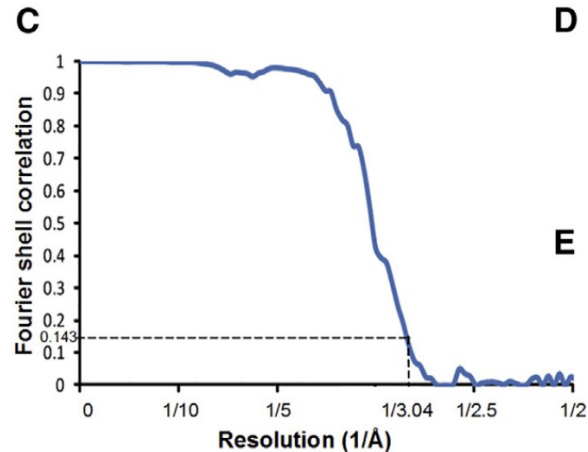
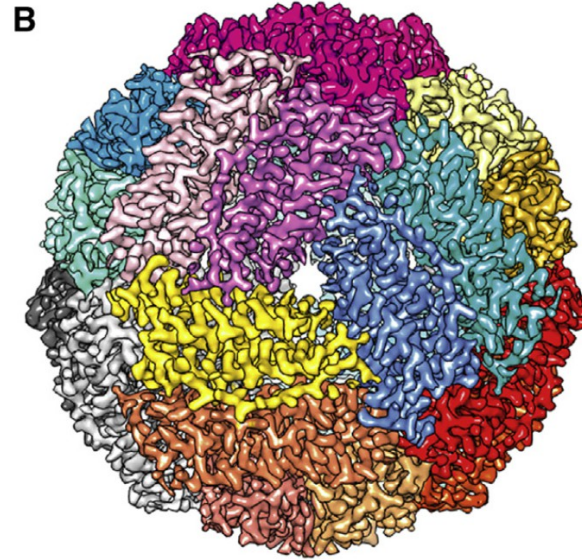
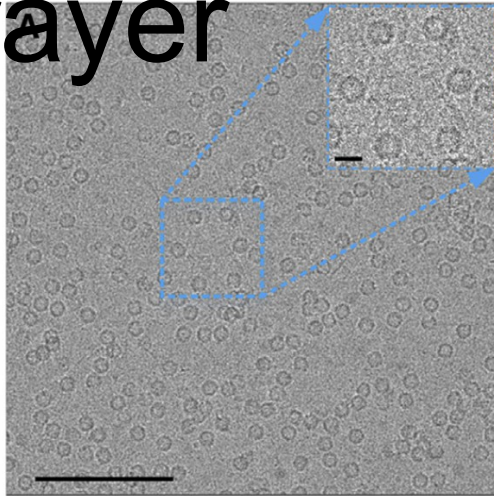
Measurements of Ice Thickness of Droplets Sprayed on the EM Grid



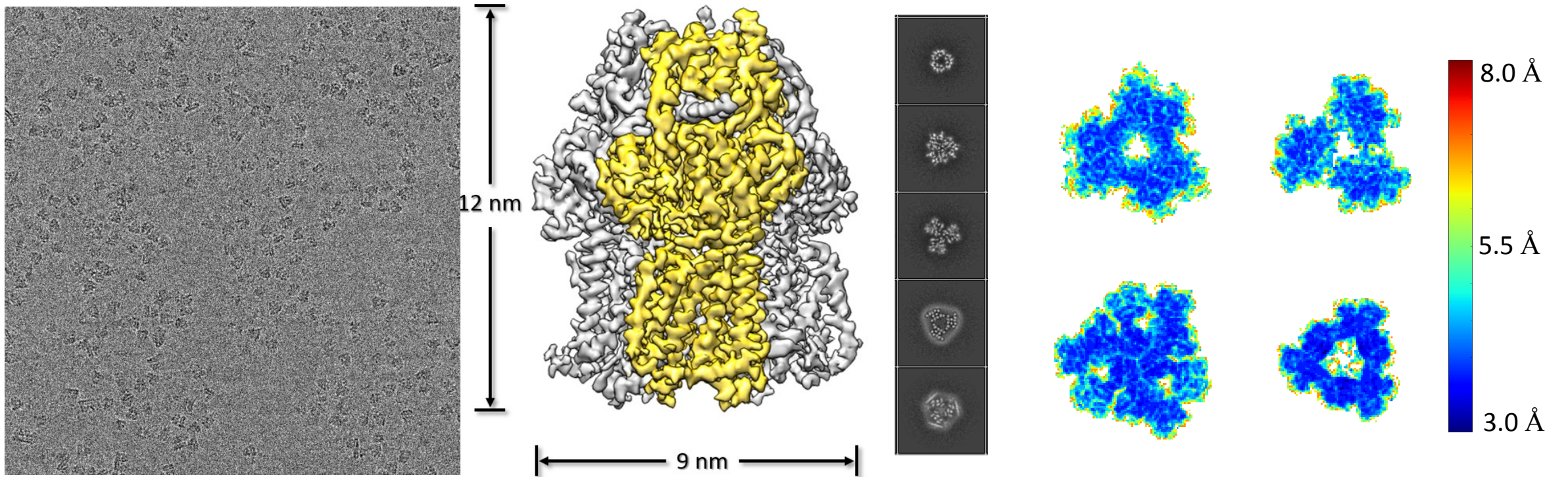
Pie charts illustrating the droplet size distribution under four different spraying conditions.



3.0-Å Resolution Structure of Apoferritin Obtained by Spraying with the Microsprayer

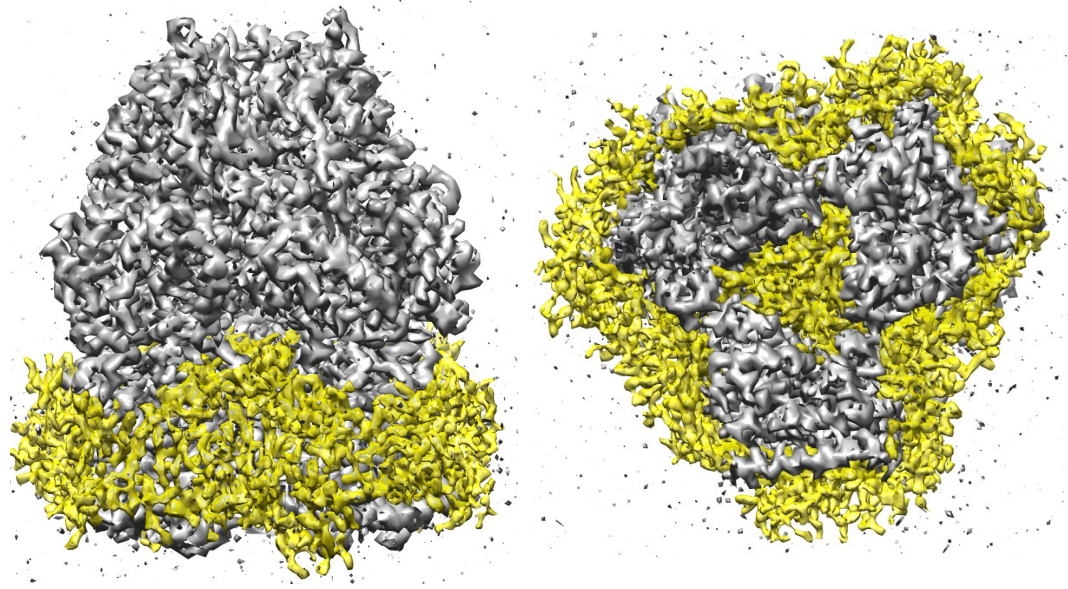


360-kDa membrane protein, AcrB 3.7 Å

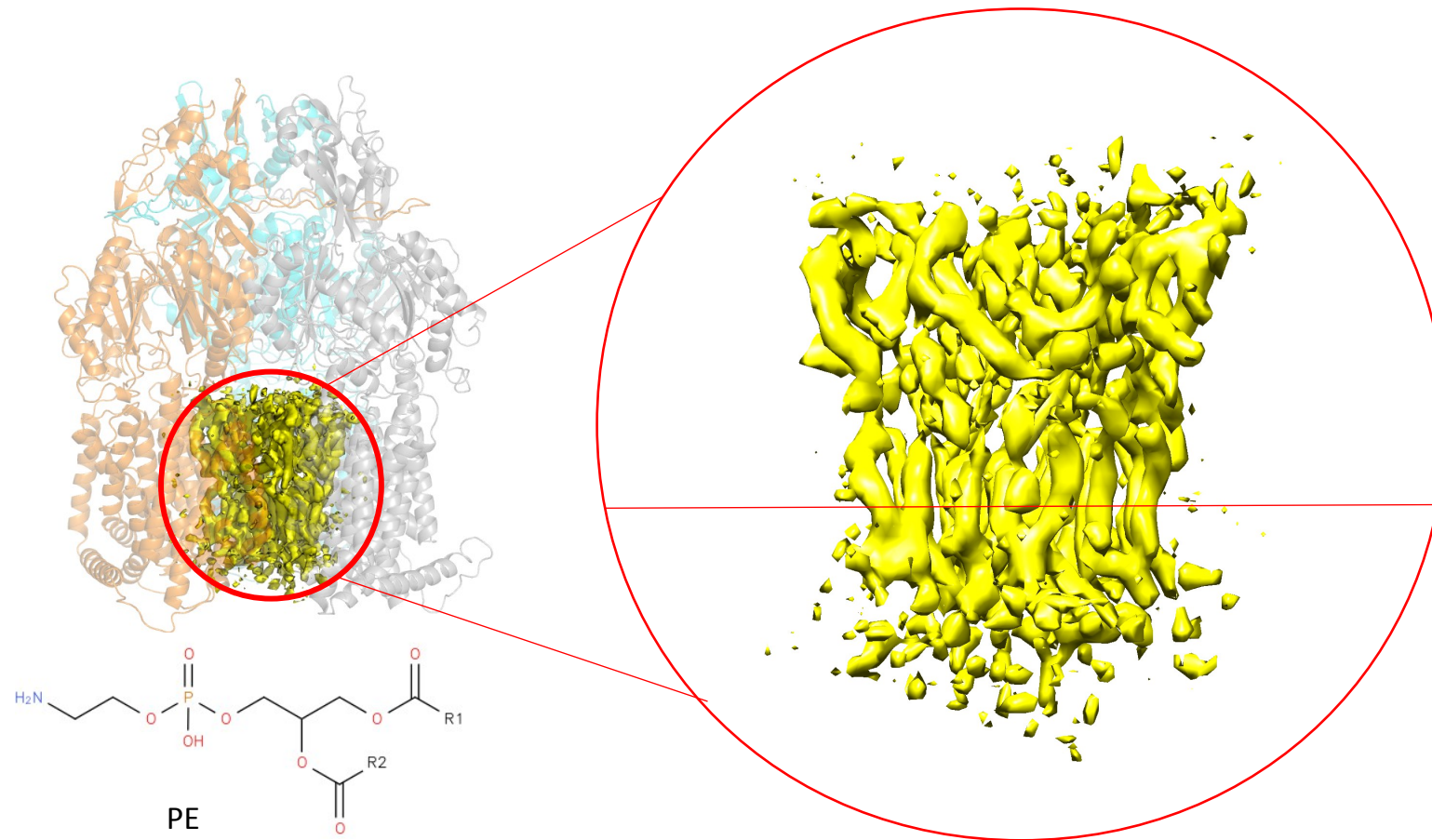


In preparation

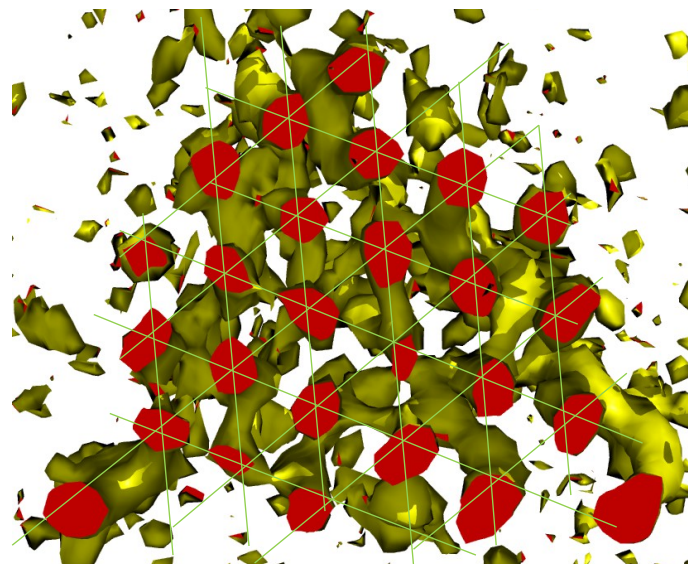
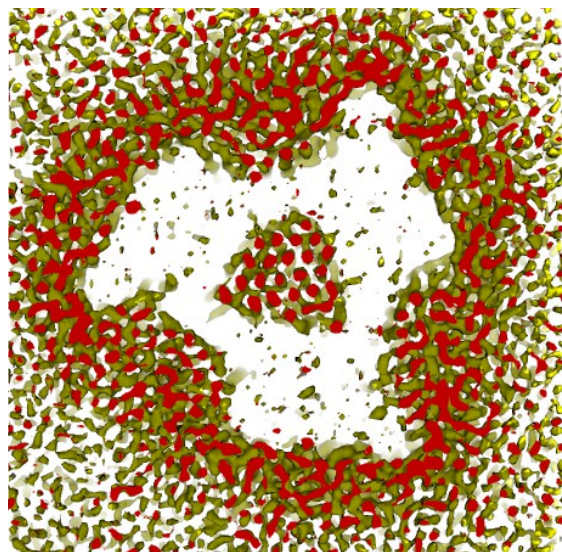
3D EM-density (3.2 Å) of AcrB in Native Cell Membrane Nanoparticle



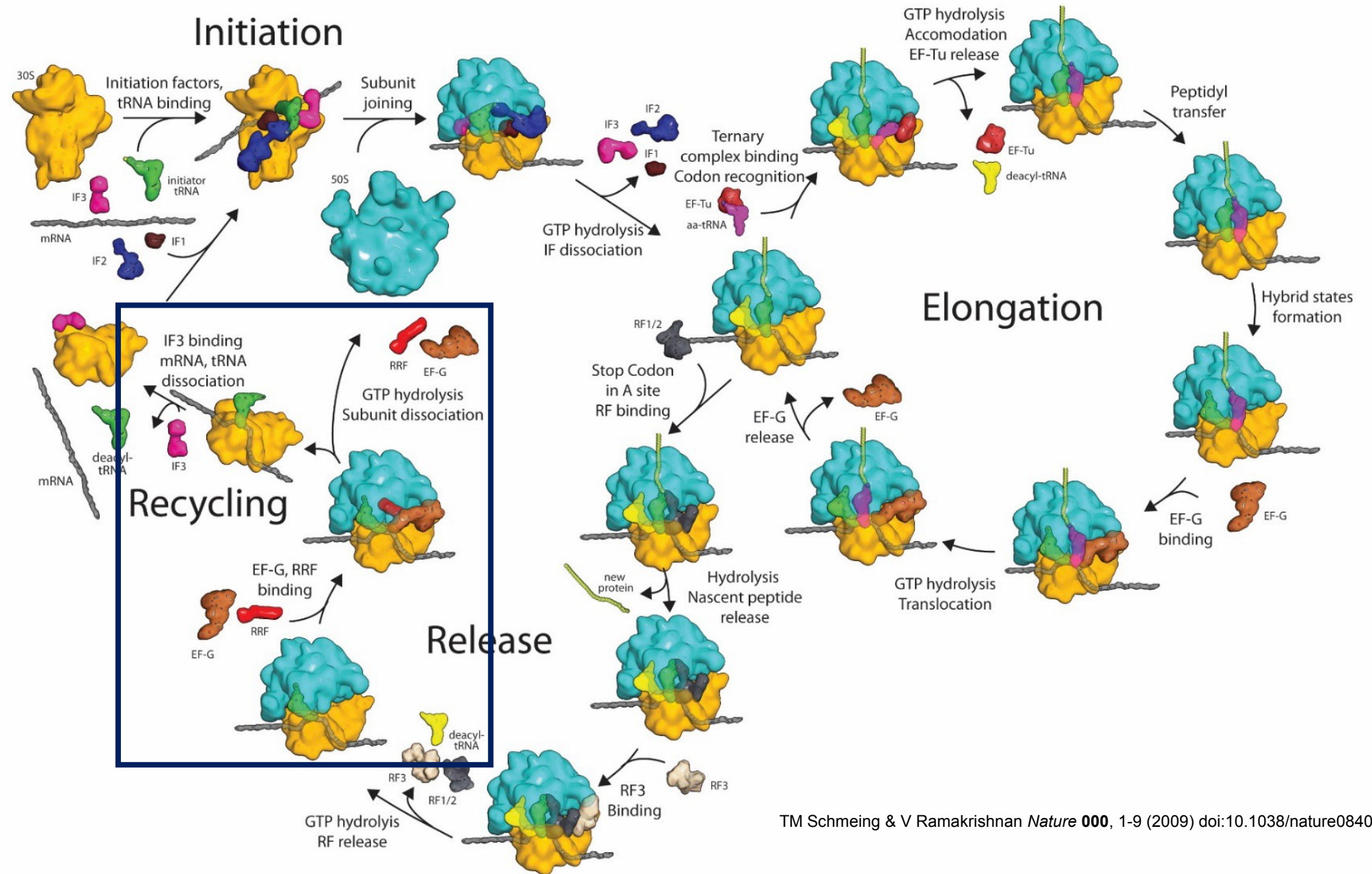
3D EM-density: Native Cell Membrane Bilayer



Lipid Belt in Sliced View and Hexagonal Pattern of Lipid Arrangement



Overview of translation



The recycling process

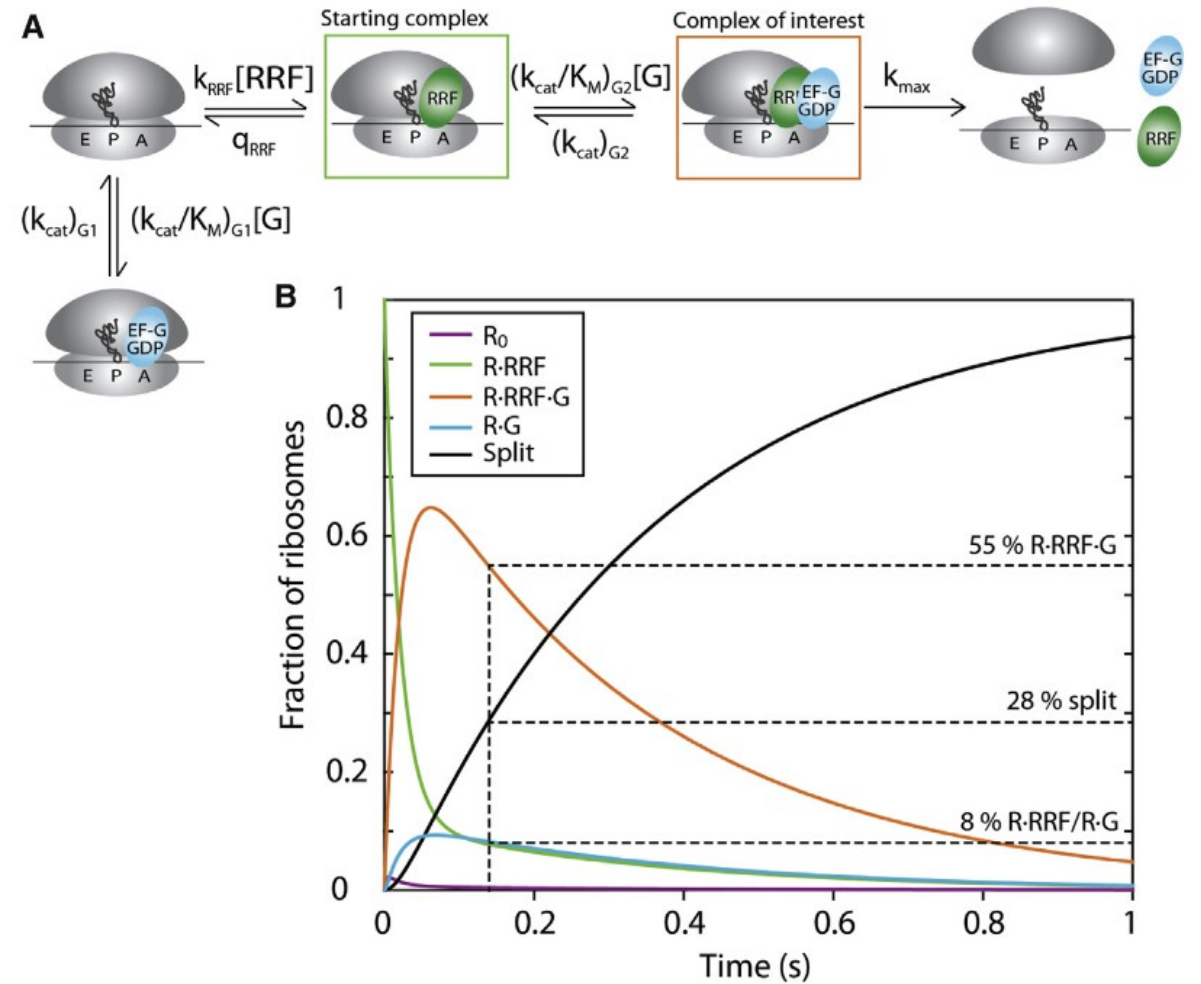
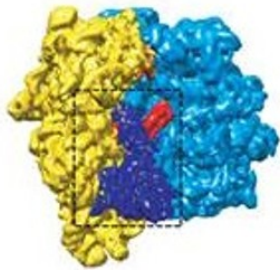
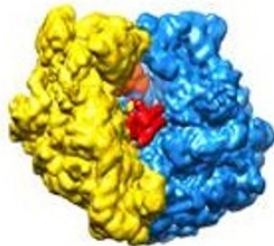
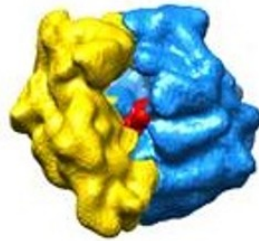
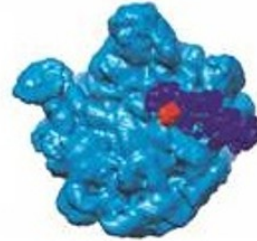


Table 1. List of the Cryo-EM Structures Obtained in this Study

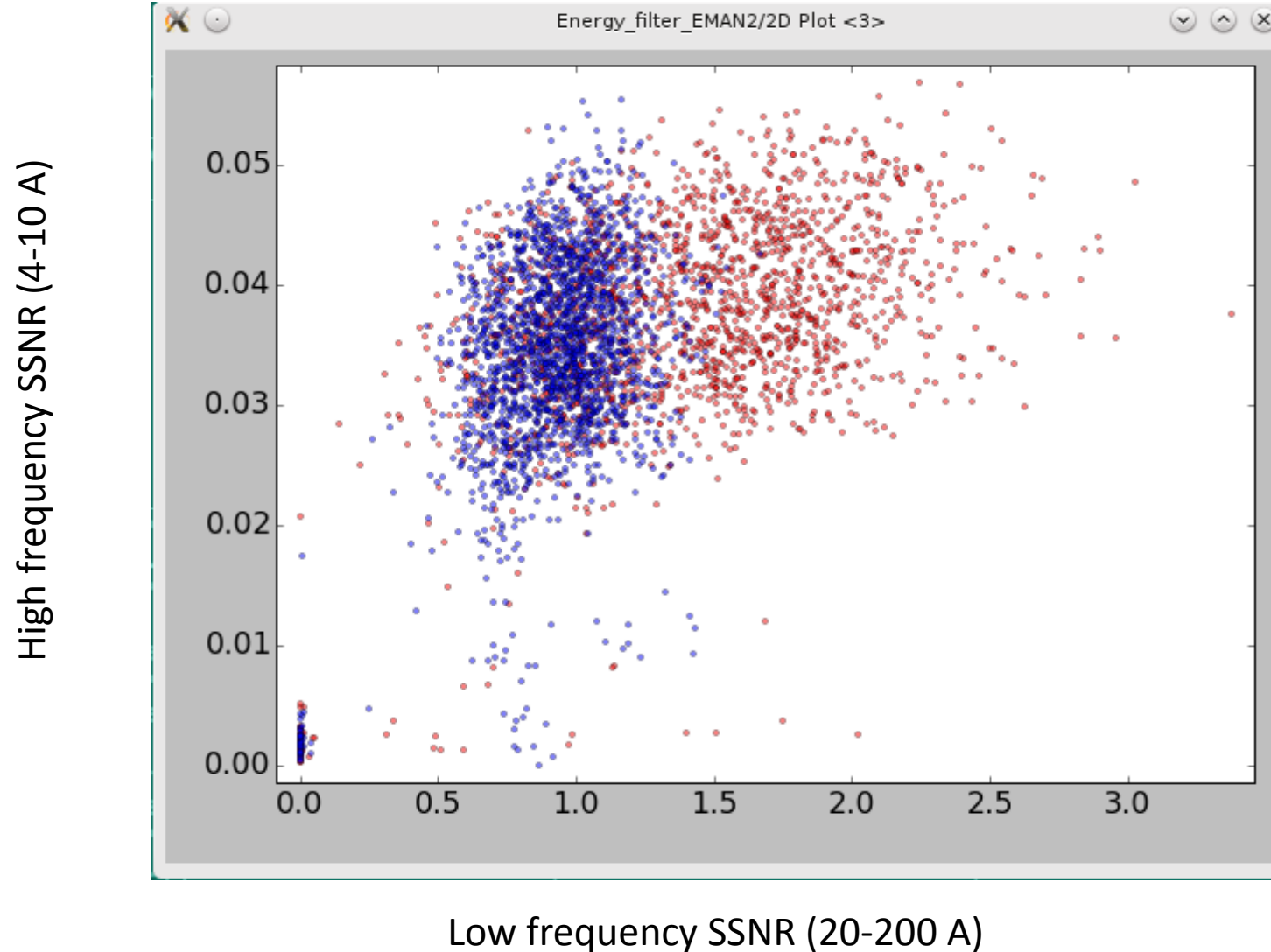
Experiment	Ribosome and Other Components (Rot or Non-rot)	Abbreviation (Resolution, Å)
Control	70S ribosome, RRF, P/E tRNA, mRNA (Rot)	PostTC·RRF _{control} (10)
140 ms	70S ribosome, RRF, P/E tRNA, mRNA (Rot)	PostTC·RRF ₁₄₀ (15)
	70S ribosome, RRF, mRNA (Non-rot)	NR-PostTC·RRF ₁₄₀ (16)
	70S ribosome, RRF, P/E tRNA, EF-G, mRNA (Rot)	PostTC·RRF·EF-G ₁₄₀ (7.4)
	50S subunit, RRF, EF-G	50S·RRF·EF-G ₁₄₀ (12)
	50S subunit, RRF, EF-G, E tRNA	50S·RRF·EF-G·tRNA ₁₄₀ (16)
	30S subunit, P/I tRNA, mRNA	30S·tRNA ₁₄₀ (10)
	30S subunit, IF3, mRNA	30S·IF3 ₁₄₀ (22)
Long incubation	50S subunit, RRF, EF-G	50S·RRF·EF-G _{long} (14)
	50S subunit, RRF, EF-G, E tRNA	50S·RRF·EF-G·tRNA _{long} (12)
	50S subunit, RRF, E tRNA	50S·RRF·tRNA _{long} (16)
	50S subunit, E tRNA	50S·tRNA _{long} (16)
	30S subunit, IF3, mRNA	30S·IF3 _{long} (10)

**PostTC·RRF·EF-G₁₄₀ (7.4)****PostTC·RRF_{control} (10)****NR-PostTC·RRF₁₄₀ (16)****50S·RRF·EF-G₁₄₀ (12)****50S·RRF·EF-G·tRNA₁₄₀ (16)****30S·tRNA₁₄₀ (10)****30S·IF3_{long} (10)**

Next steps

- 1. General application
- 2. Nano-fluidic system (less sample consumption)
- 3. Sub-millisecond system
 - (mixing time < 50 μ s, freezing time < 100 μ s)

Energy filter 20 eV (red) vs no slit (blue)



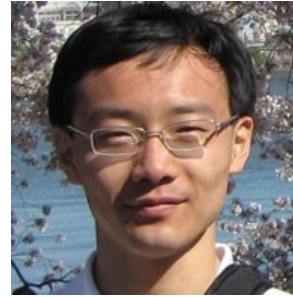
Frank Lab Team



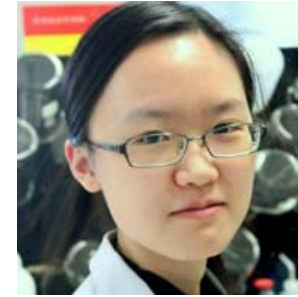
Prof. Joachim Frank



Sandip Kaledhonkar



Bo Chen

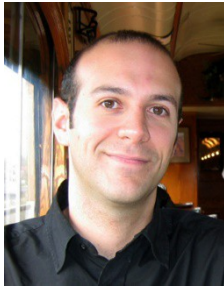


Ming Sun



Bob Grassucci

Gonzalez Lab



Ruben



Kelvin

Ehrenberg Lab



Mans



Anneli

3D sprayer Microfluidic device with PDMS



Dr. Qiao Lin



Xiangsong Feng



Yuan Jia

Dept. of Mechanical Engineering
Columbia University

New Time-Resolved Machine



Prof. Howard D. White
Eastern Virginia Medical School

References

Key intermediates in ribosome recycling visualized by time-resolved cryoelectron microscopy

Z Fu, S Kaledhonkar, A Borg, M Sun, B Chen... - Structure, 2016

A Fast and Effective Microfluidic Spraying-Plunging Method for High-Resolution Single-Particle Cryo-EM

X Feng*, Z Fu*, S Kaledhonkar, Y Jia, B Shah, A Jin... - Structure, 2017

Lipid Bilayer Structure in Native Cell Membrane Nanoparticles of Multidrug Exporter AcrB

Qiu W*, Z Fu*, Xu G, ... - Submitted