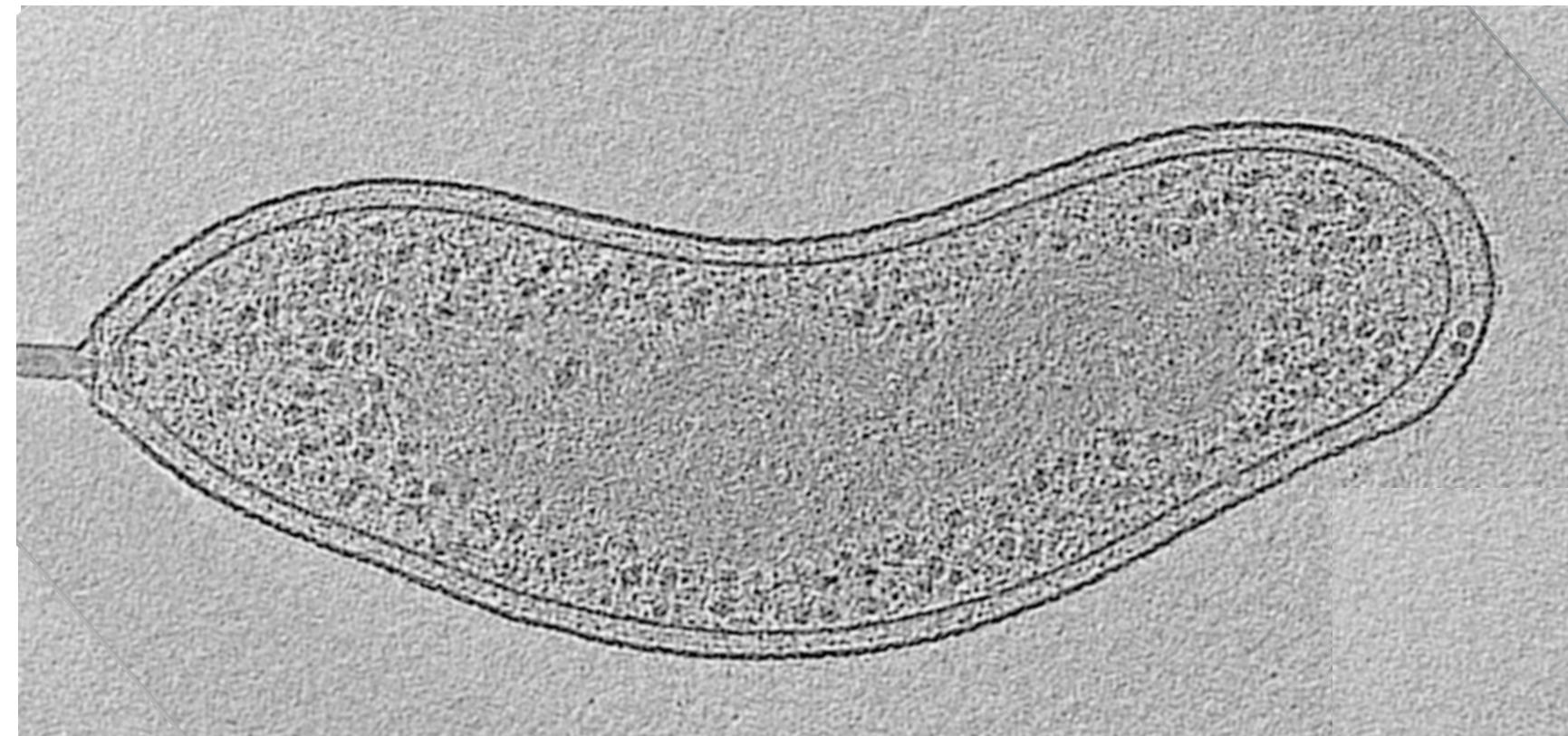


Electron cryotomography

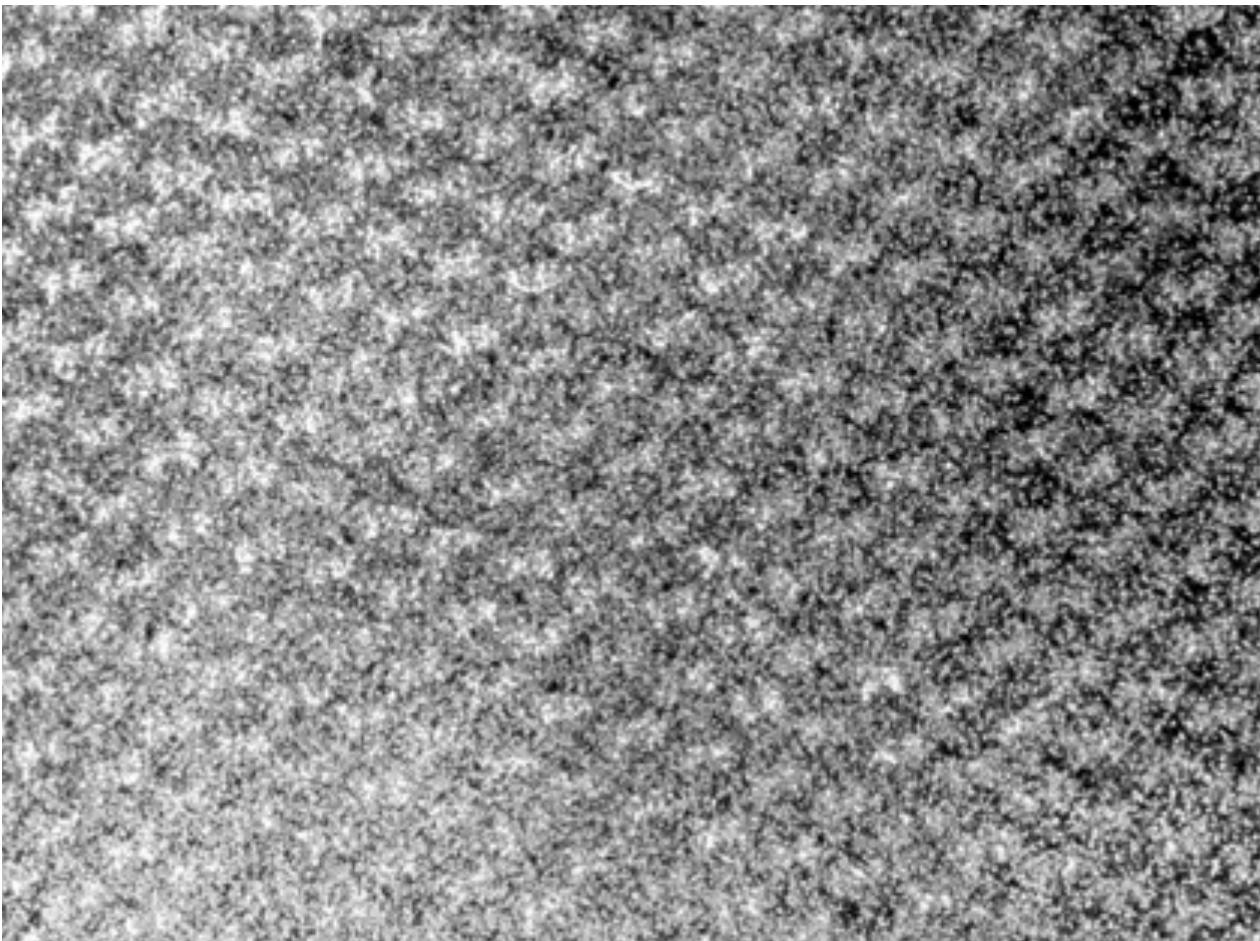
Grant Jensen
BYU

Basic approaches in cryo-EM



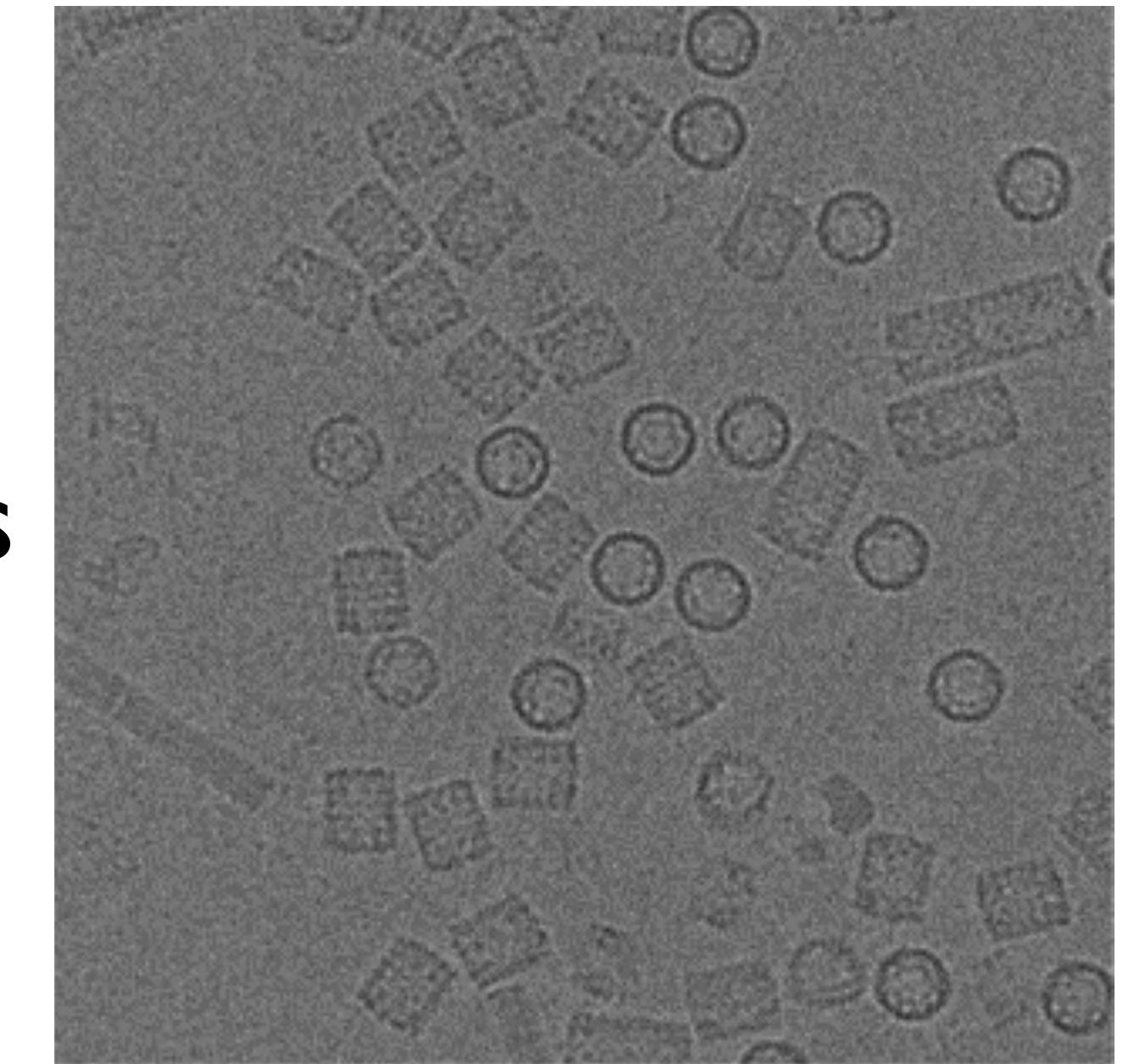
Tomography

*1 object
~4 nm resolution*



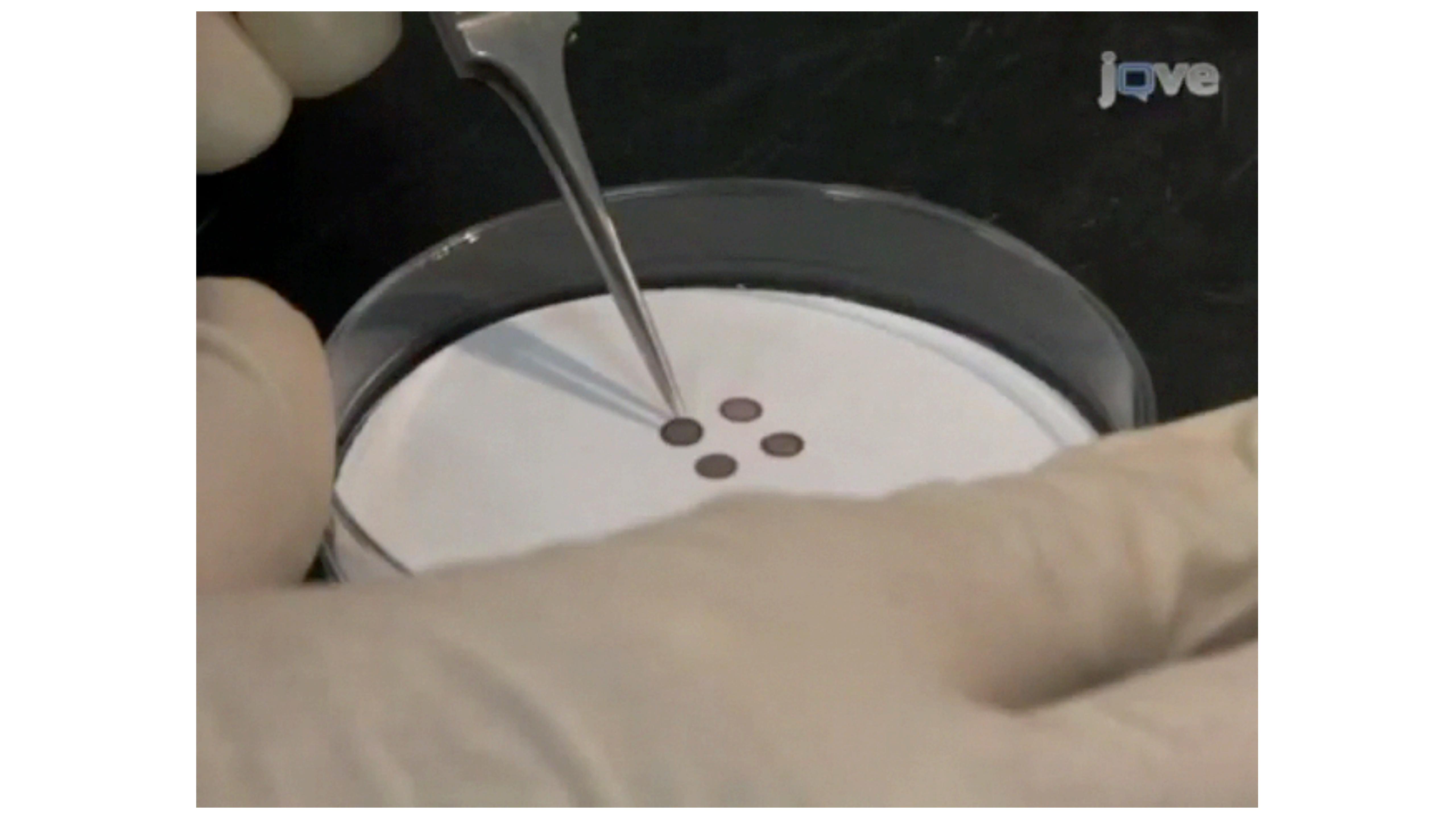
Single particle analysis

*Average 10^4 objects
1-4 Å resolution*

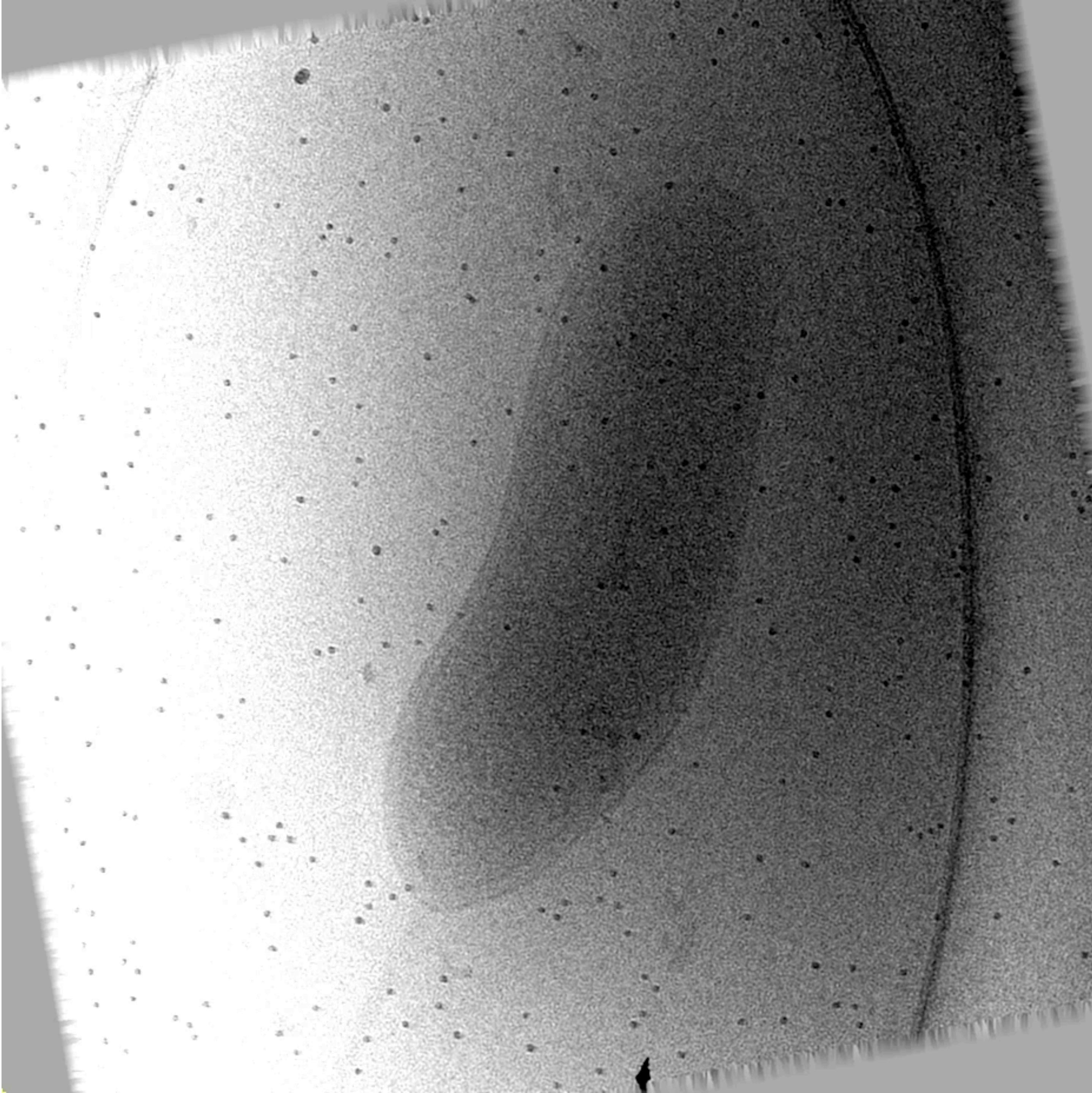


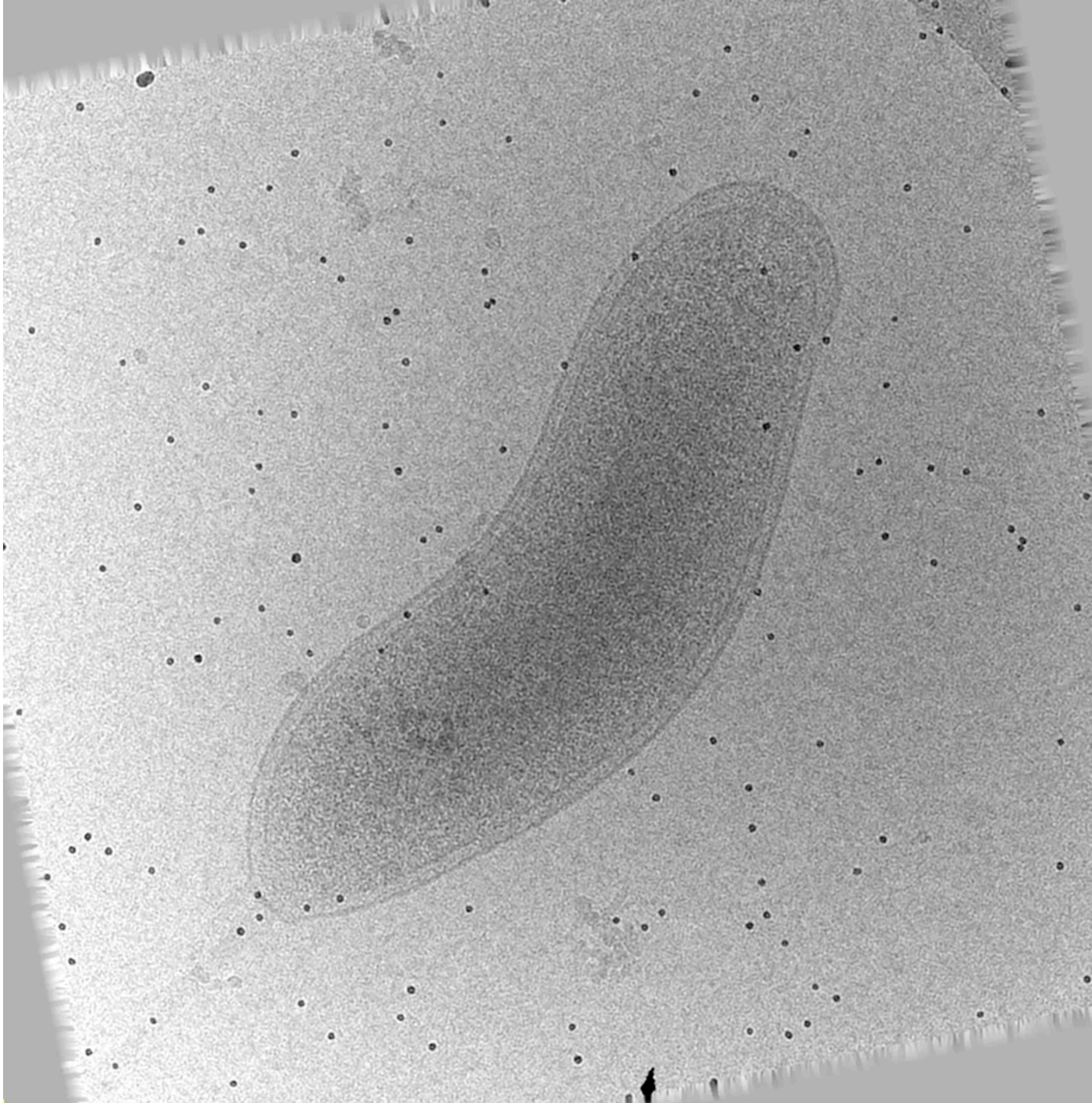
Electron crystallography

*Average 10^7 objects
0.5-3 Å resolution*



jove





Aligning tilt series

For each image, determine

x,y shifts

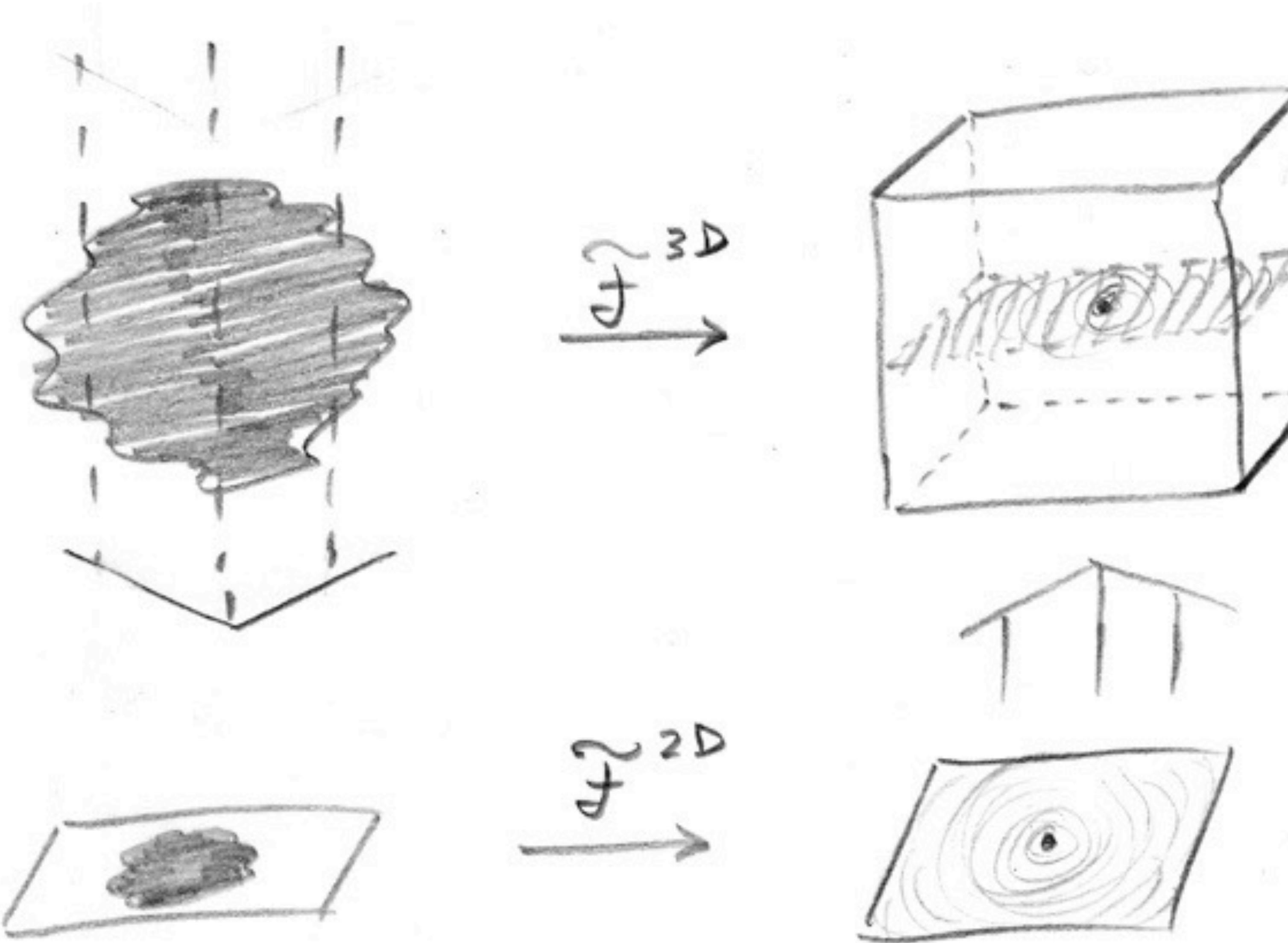
rotation (position of tilt axis)

tilt angle

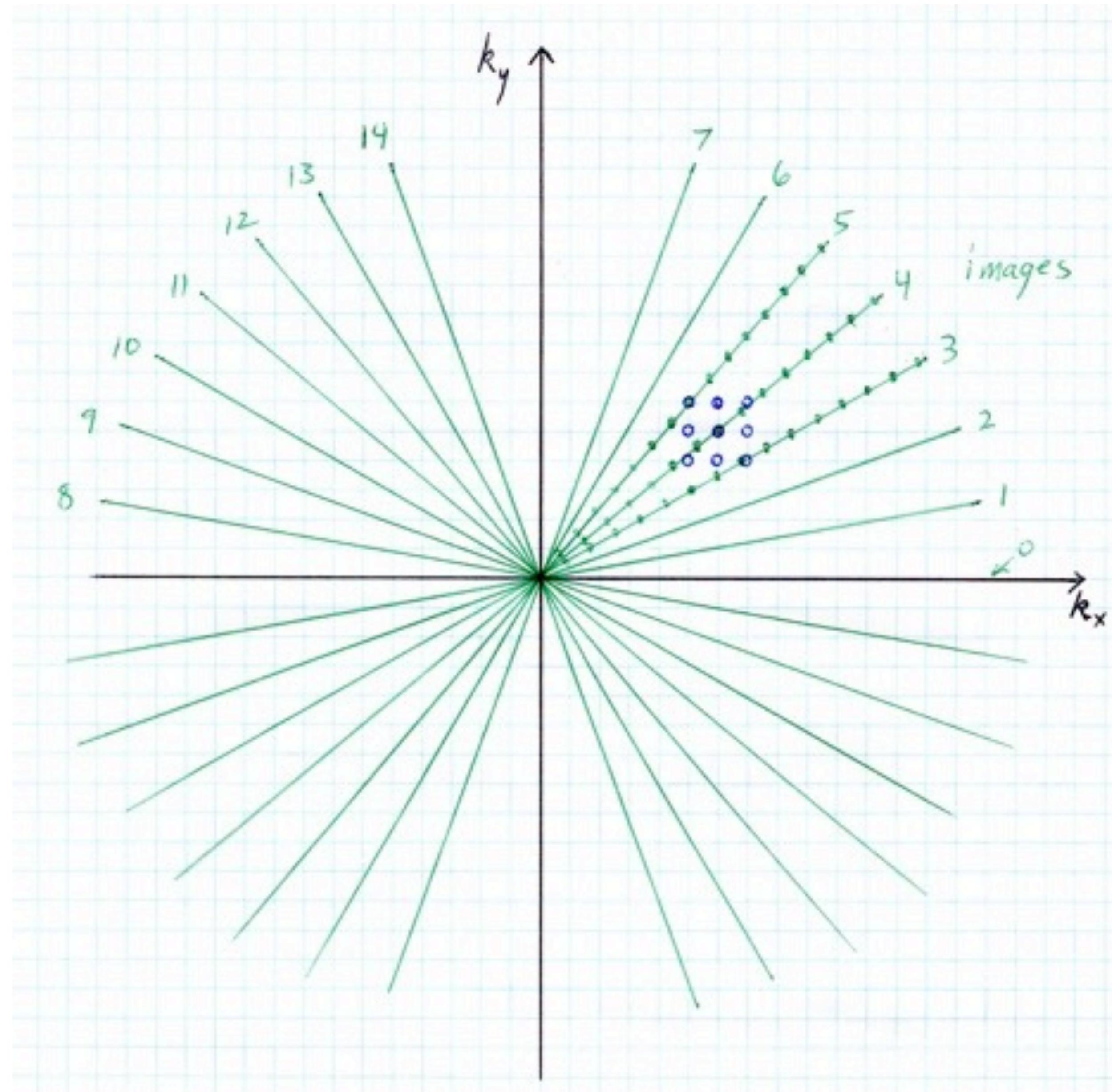
magnification

defocus

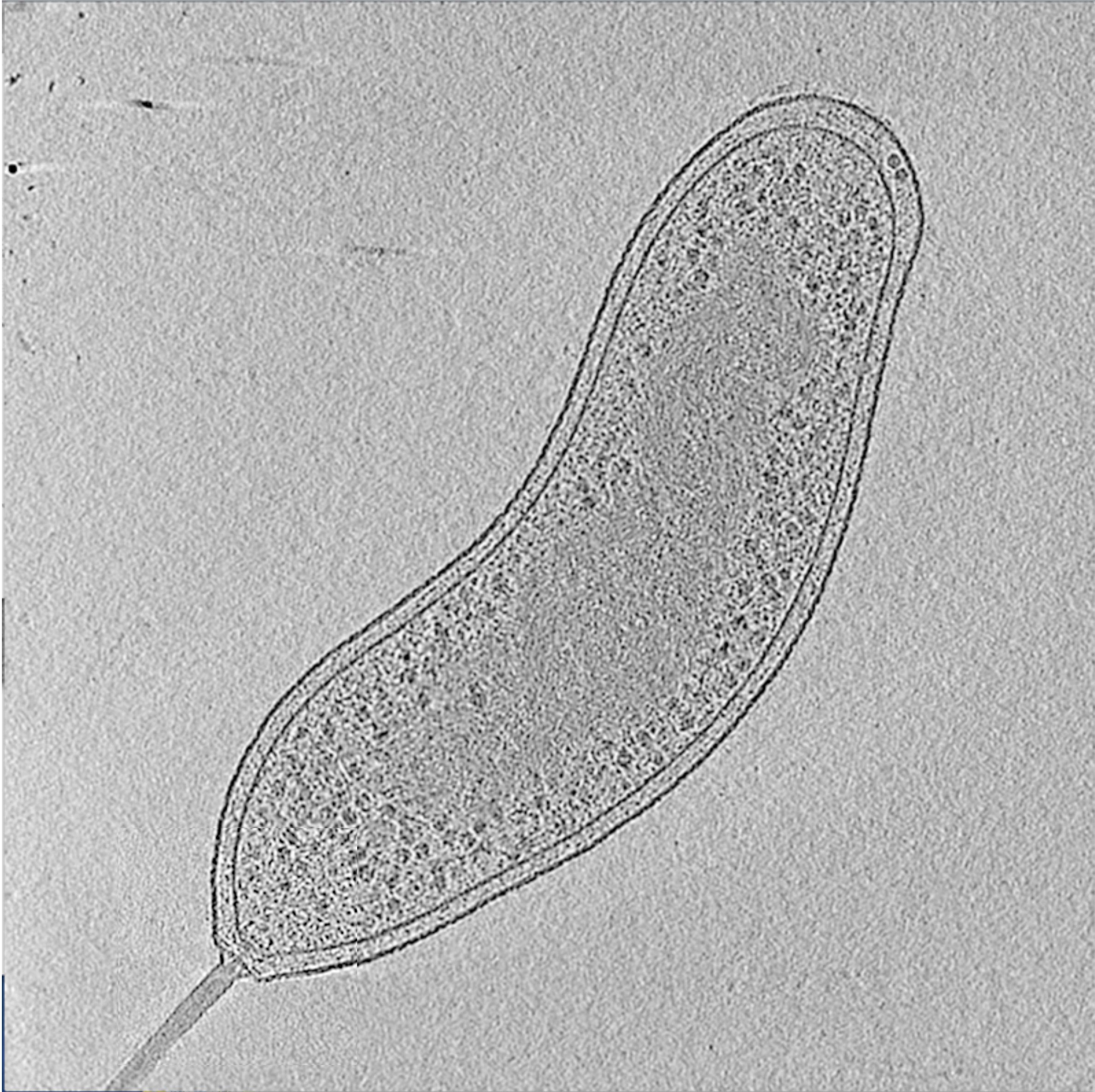
The projection theorem



Tilt series are typically merged in reciprocal space

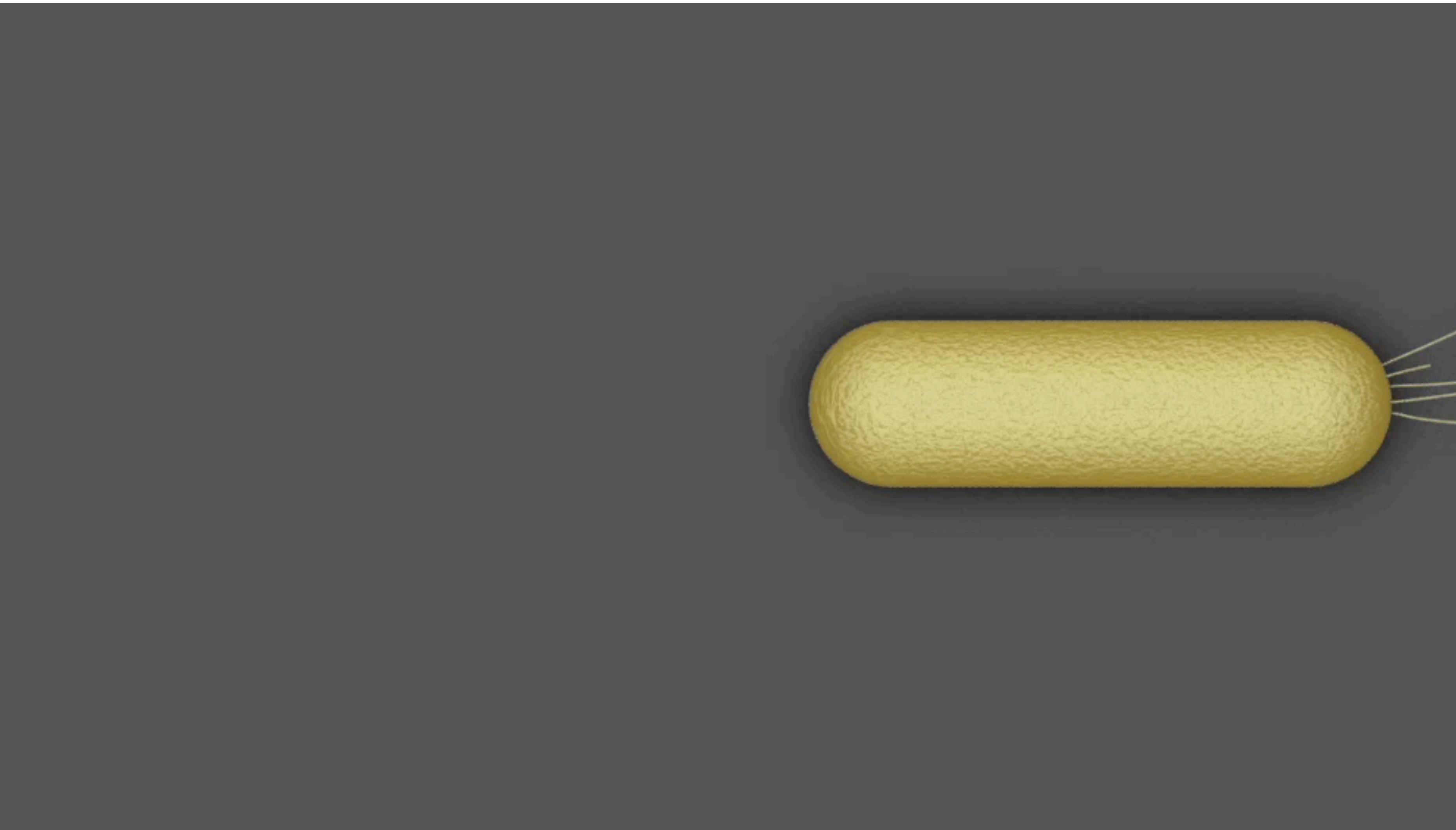


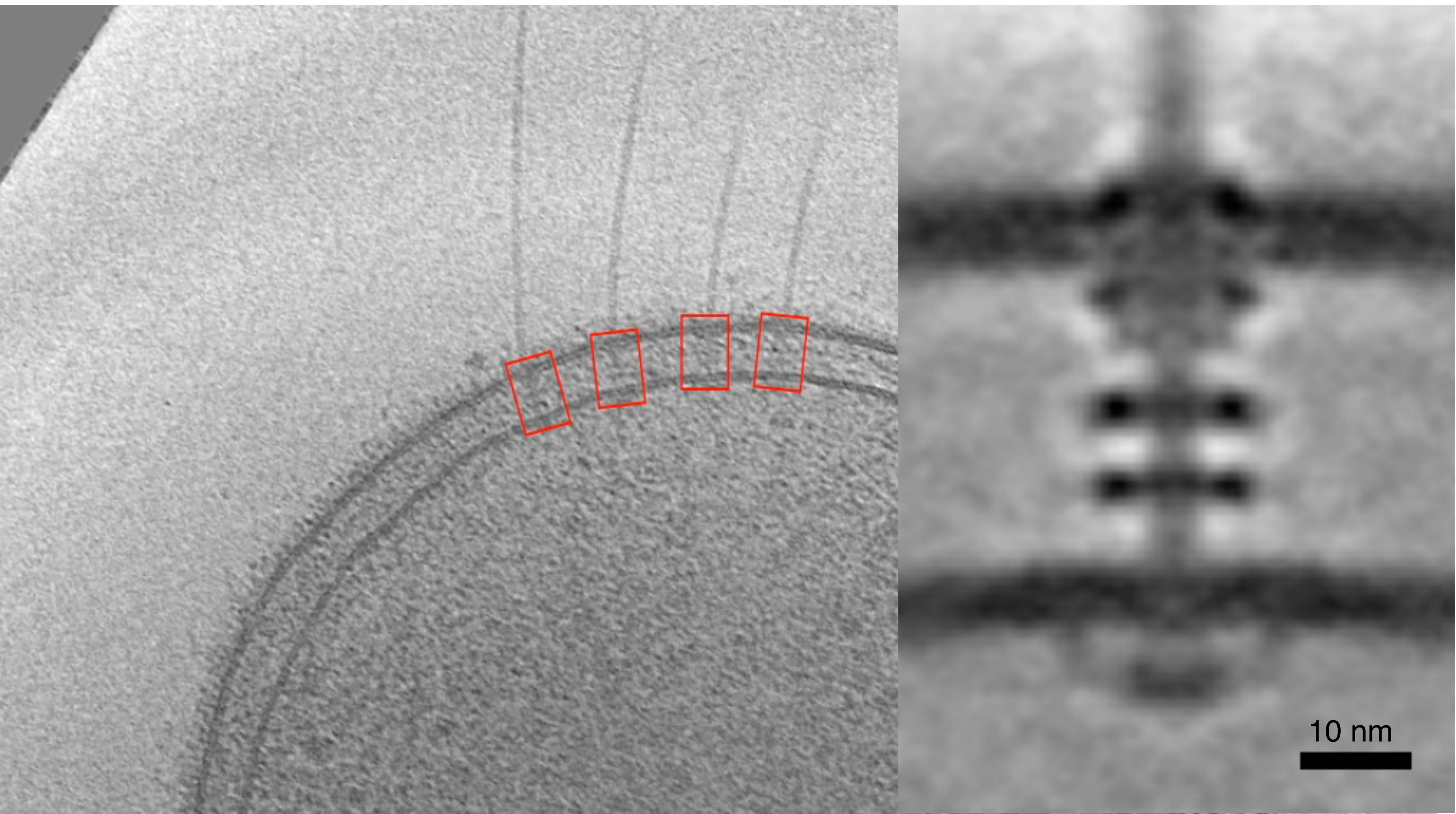


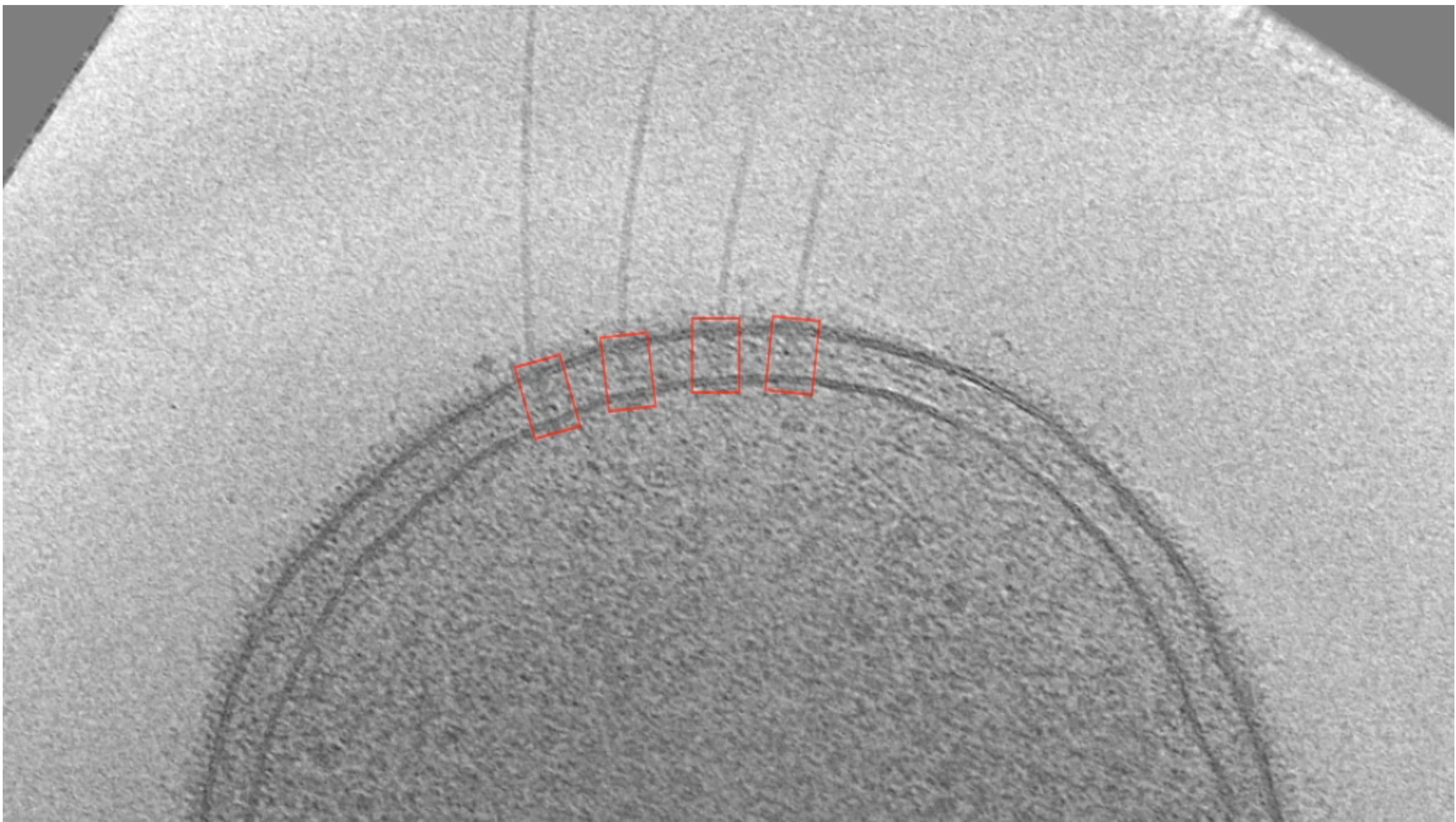


Myxococcus xanthus



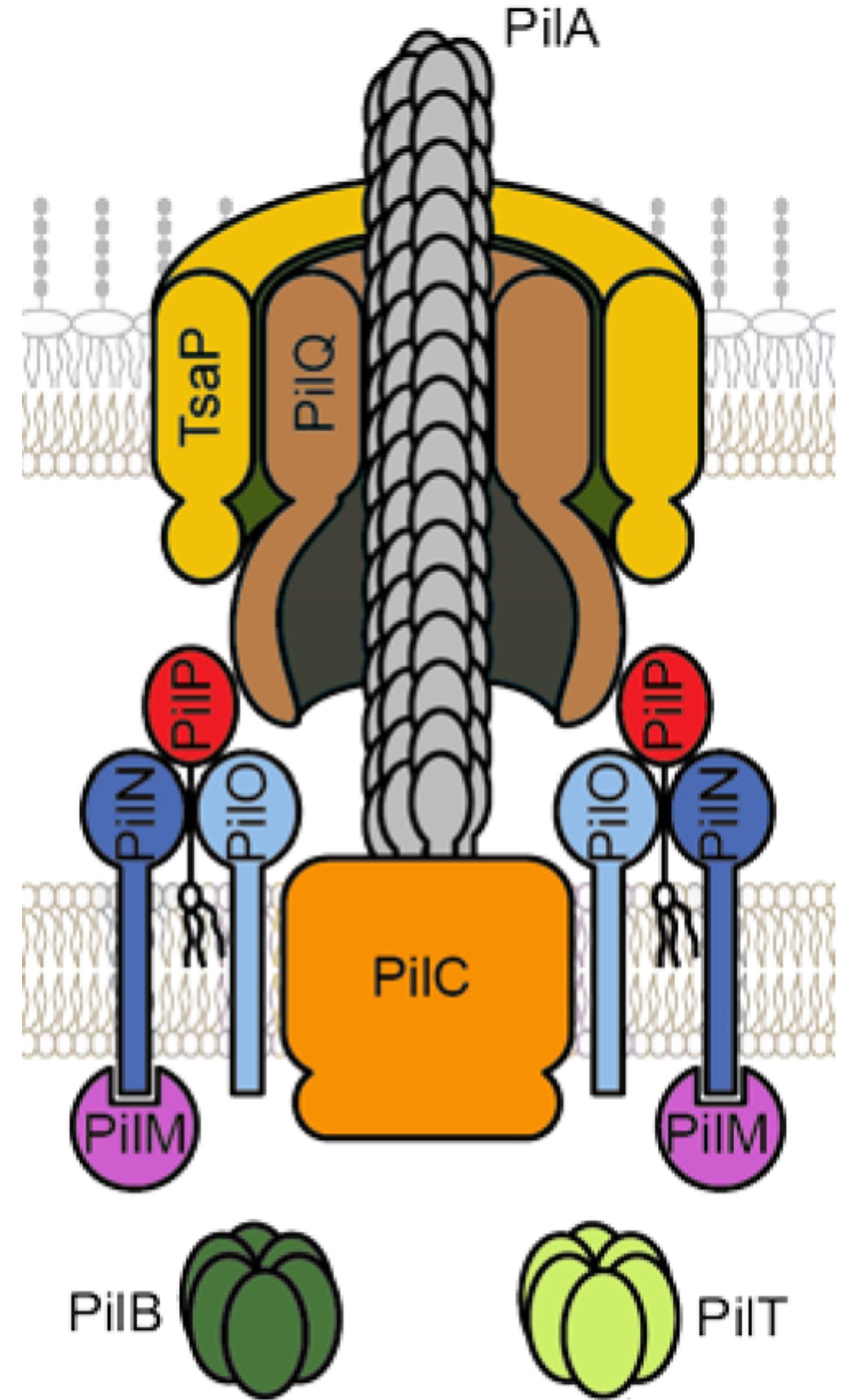






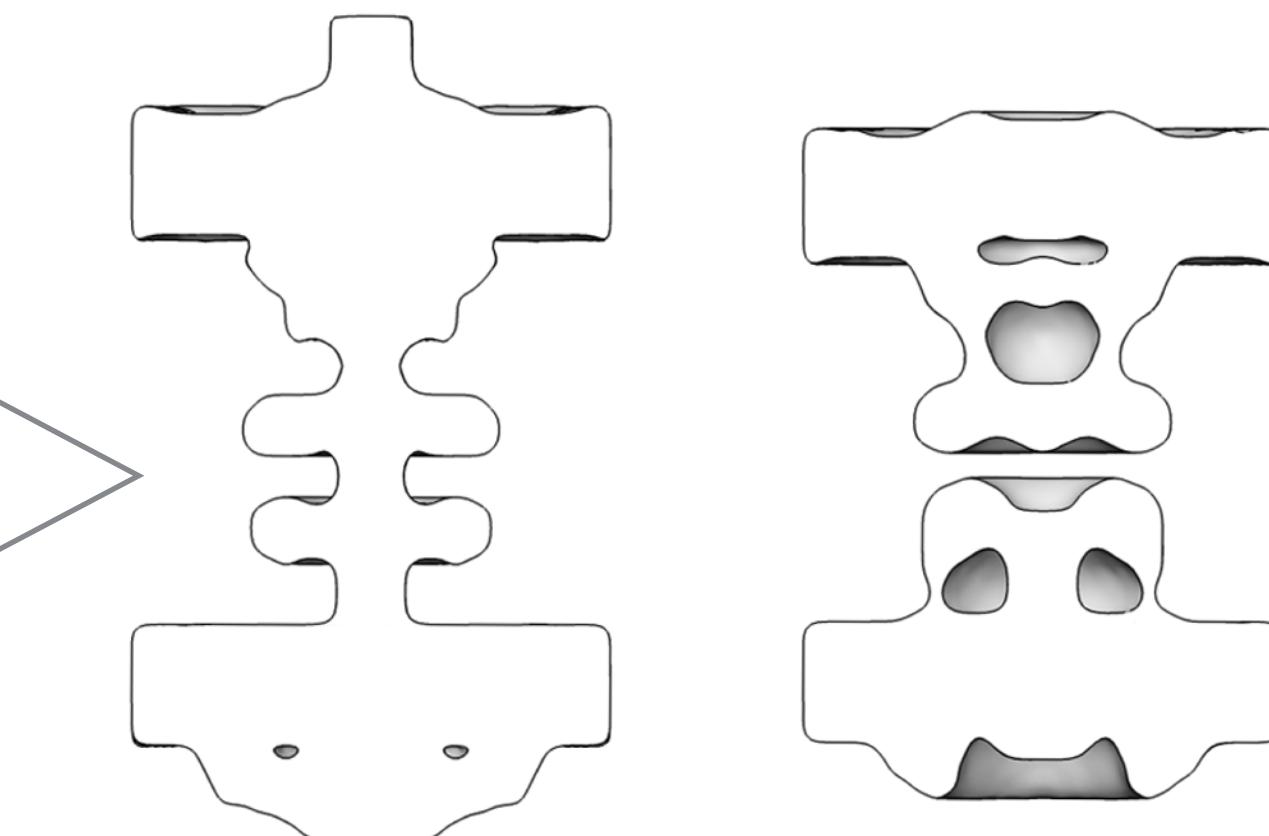
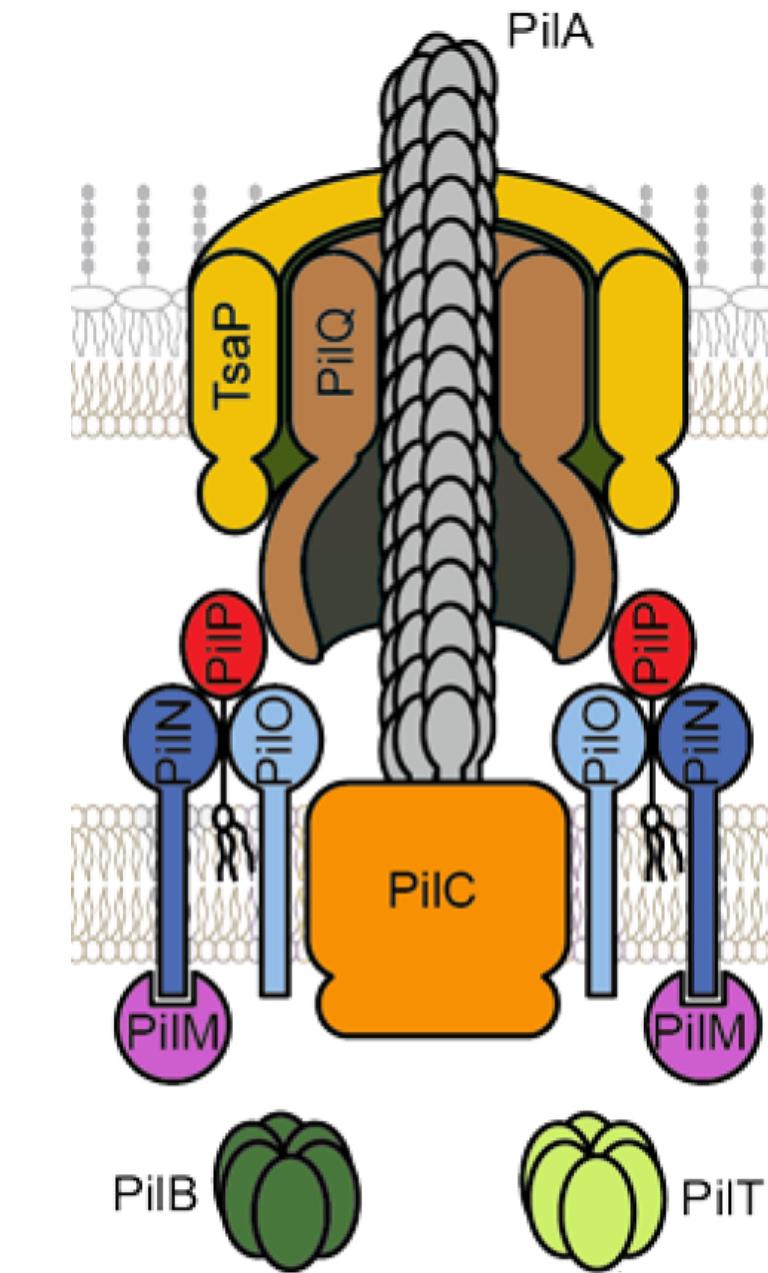


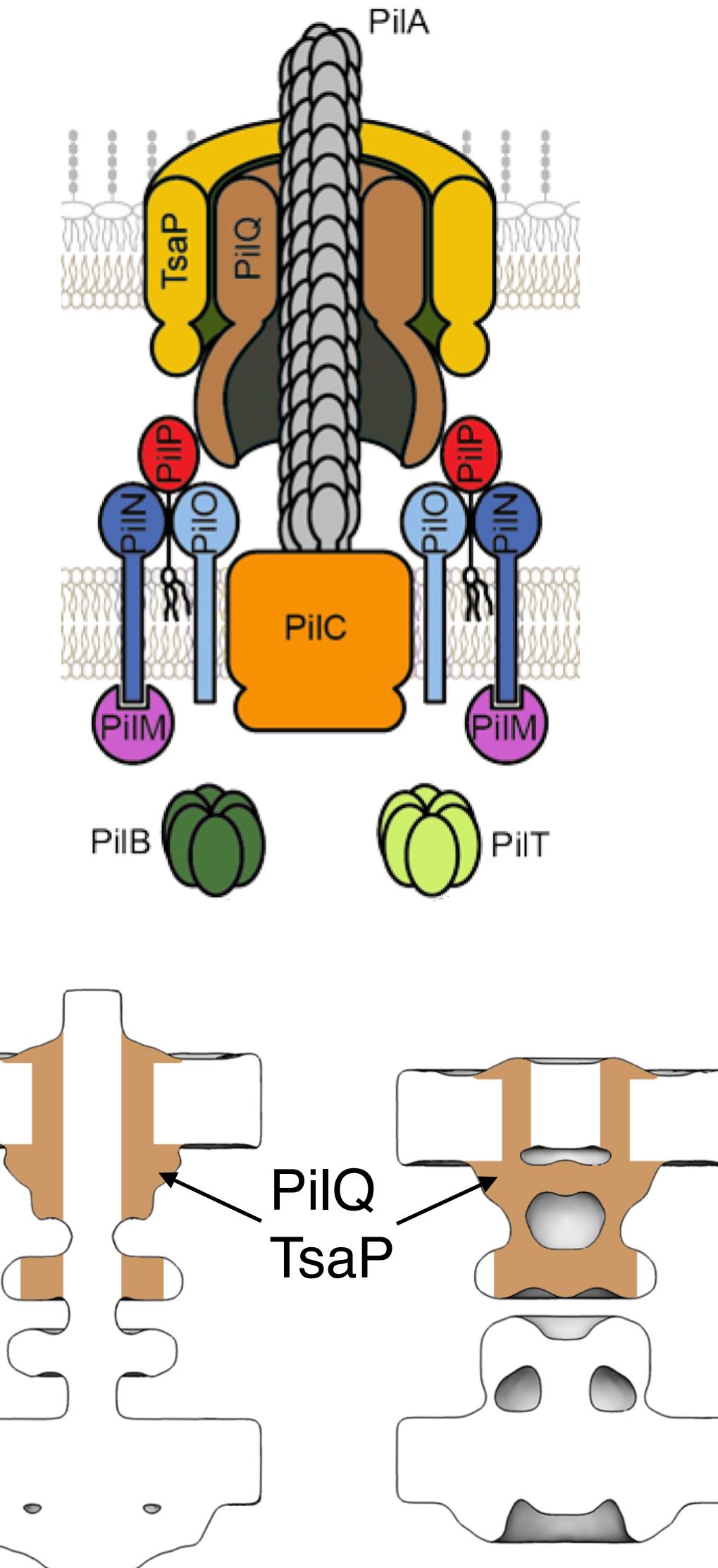
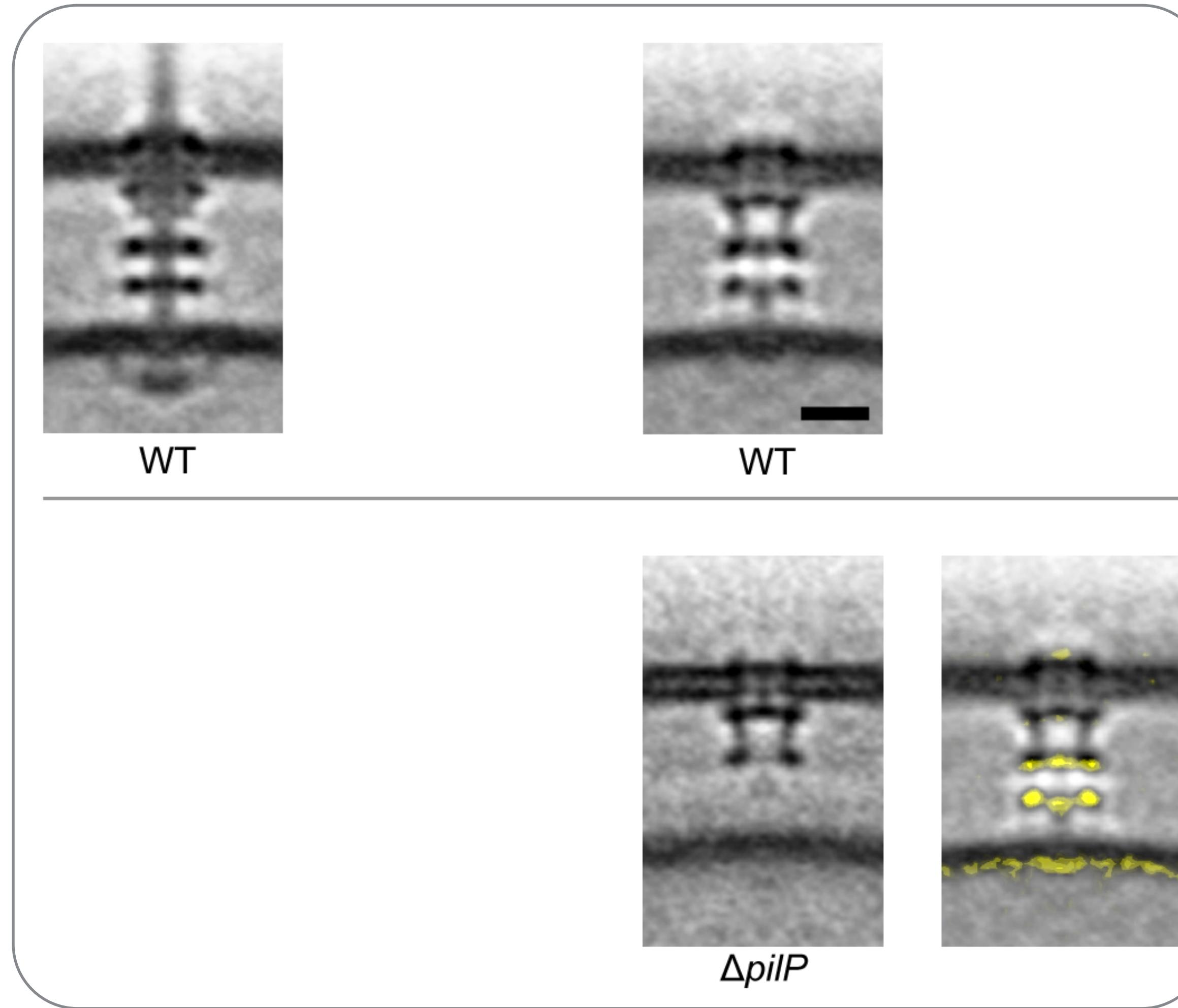
Non-piliated

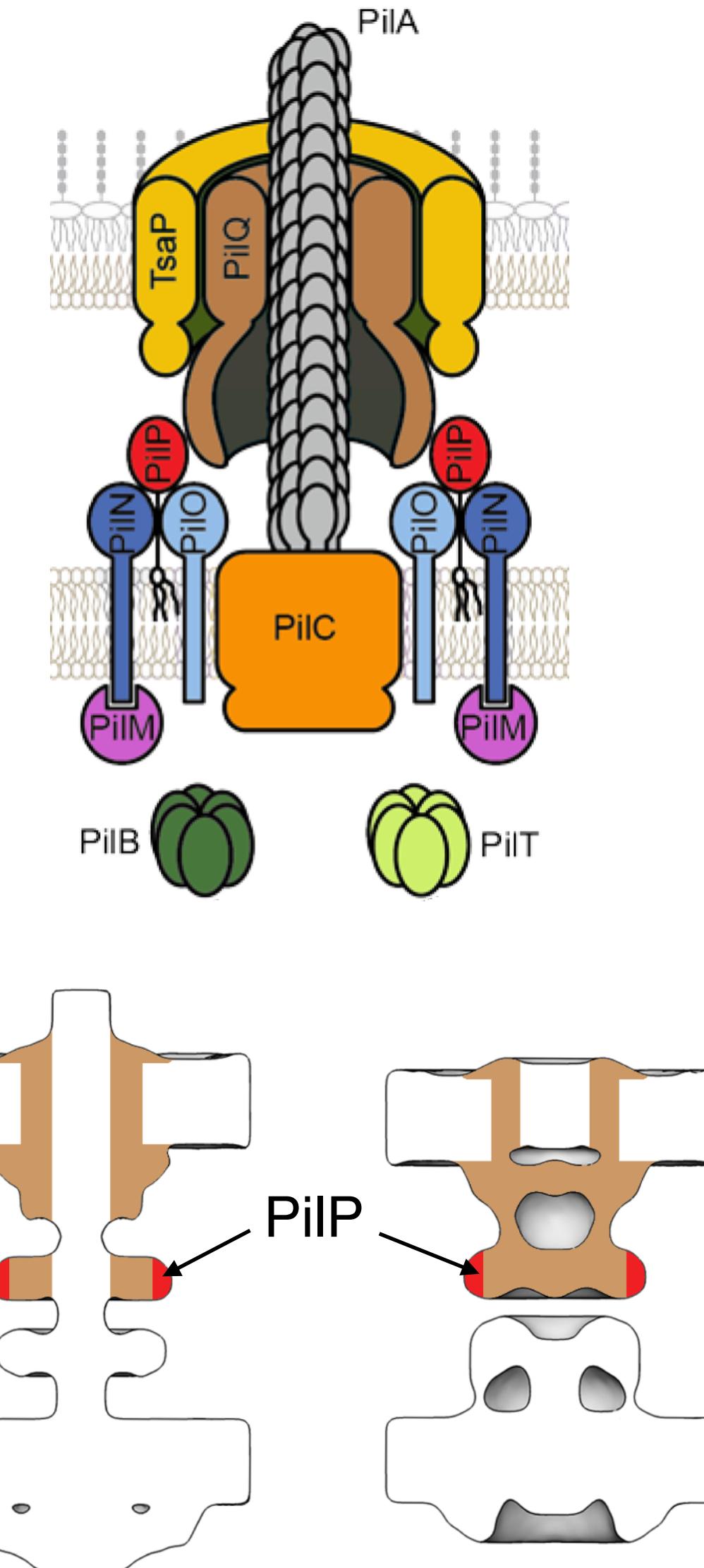
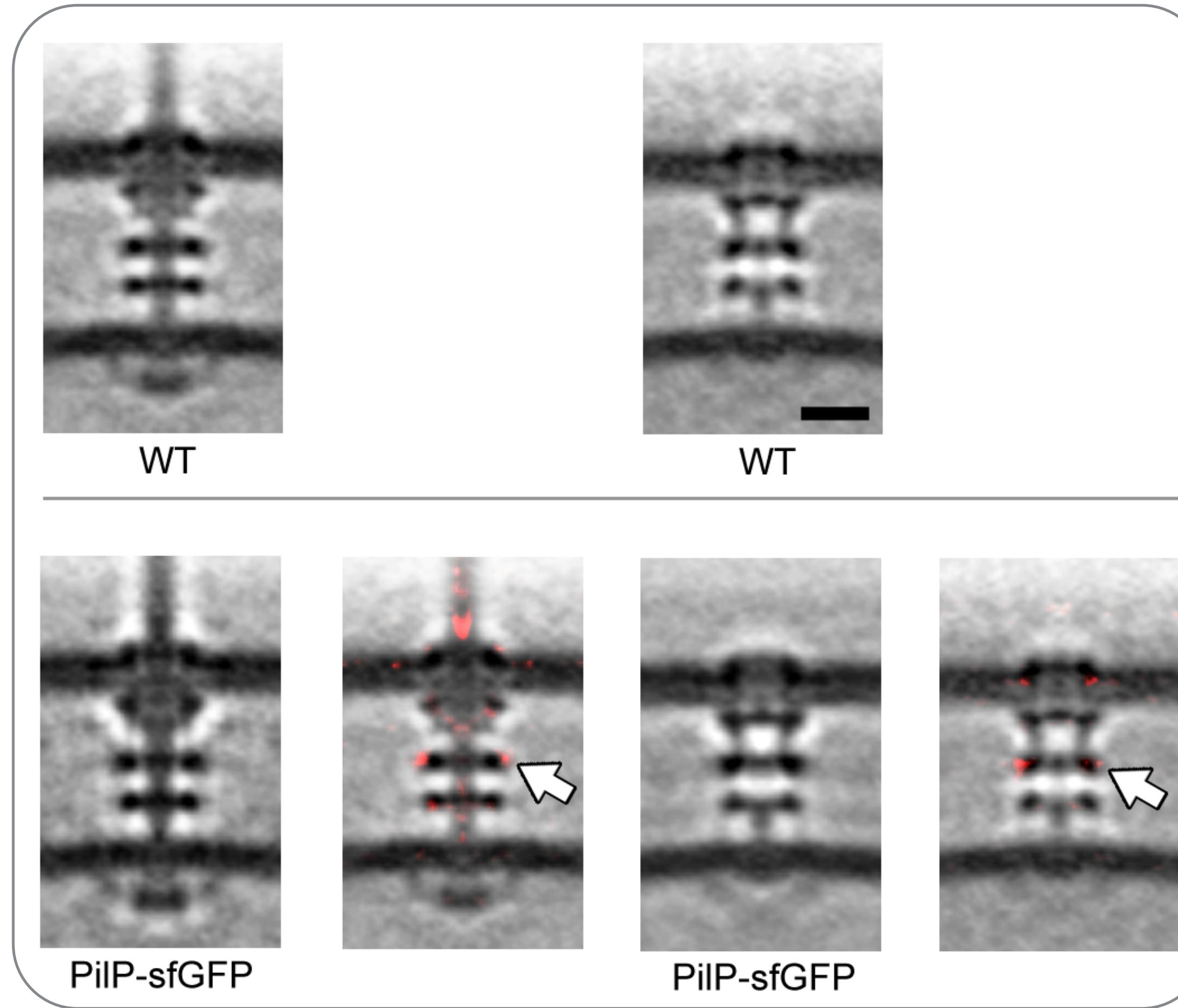


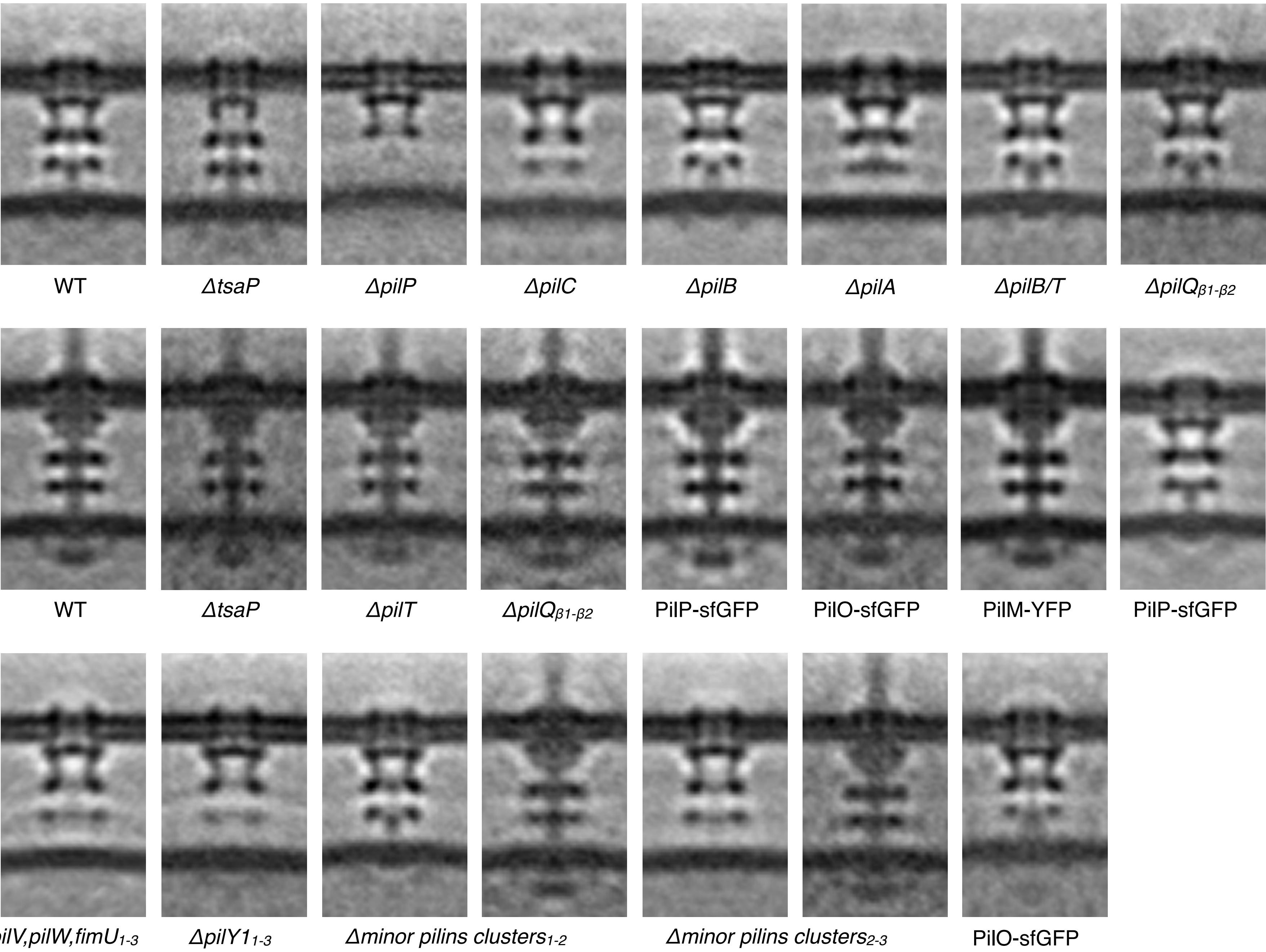


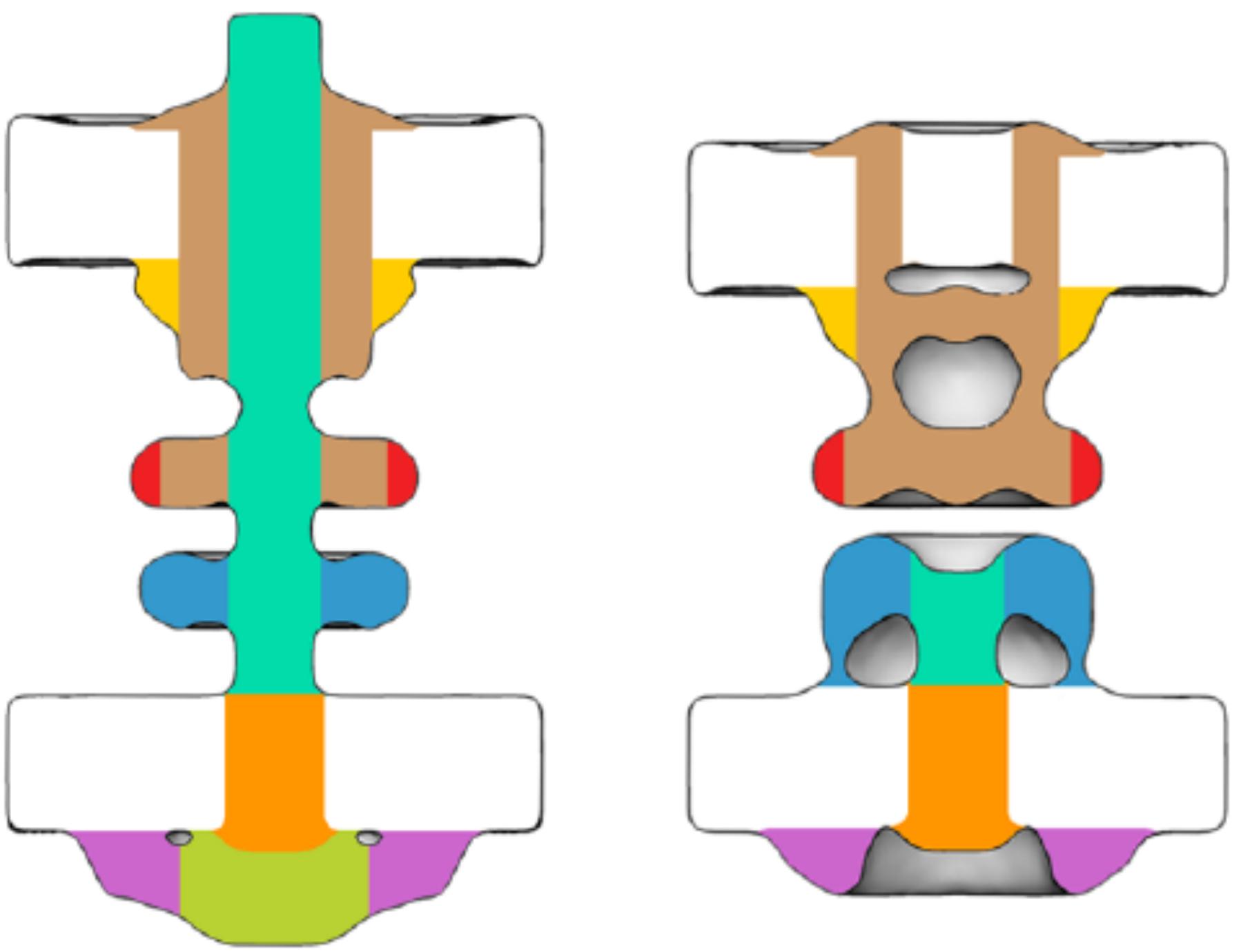
strategy: image knockouts and tagged strains



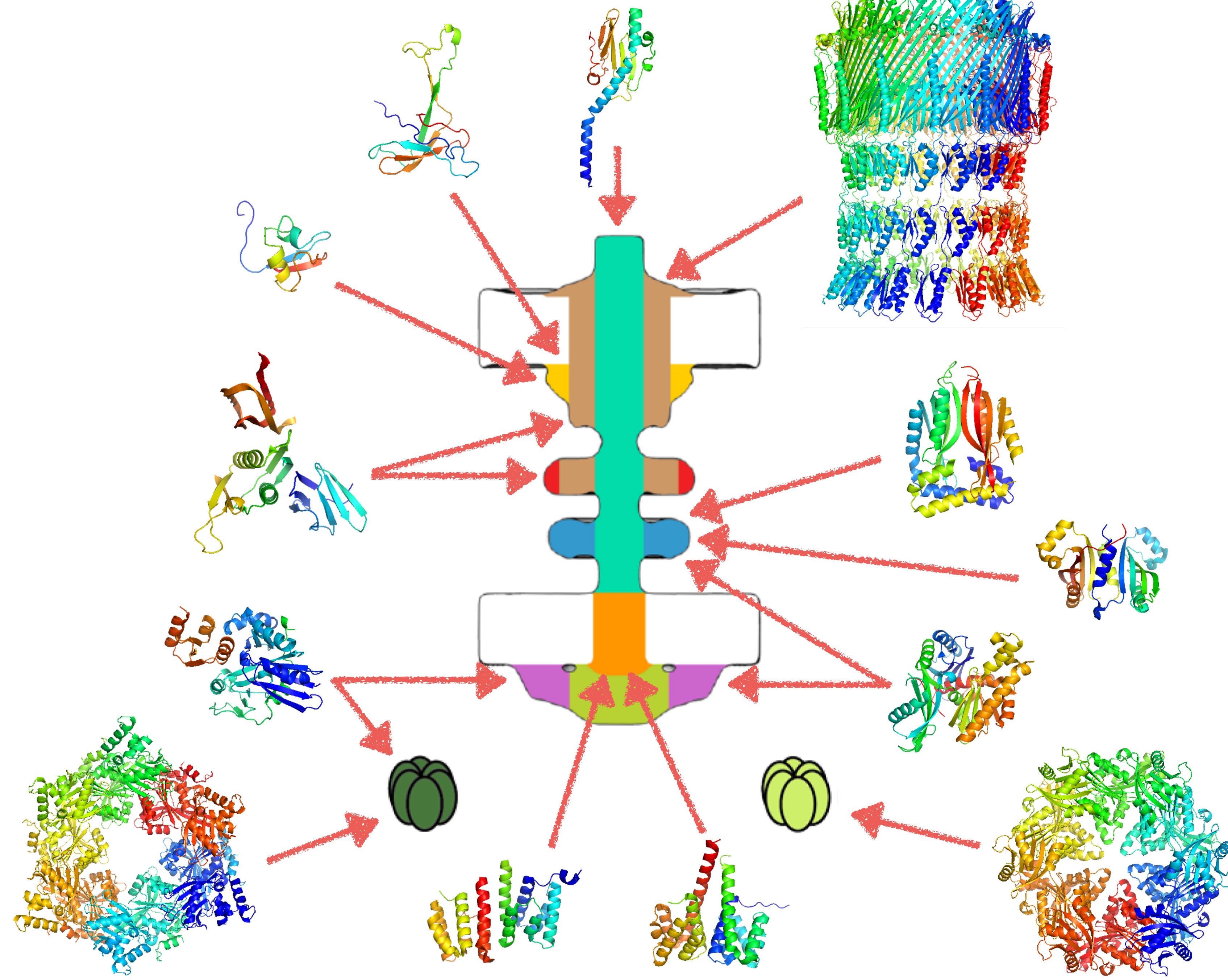


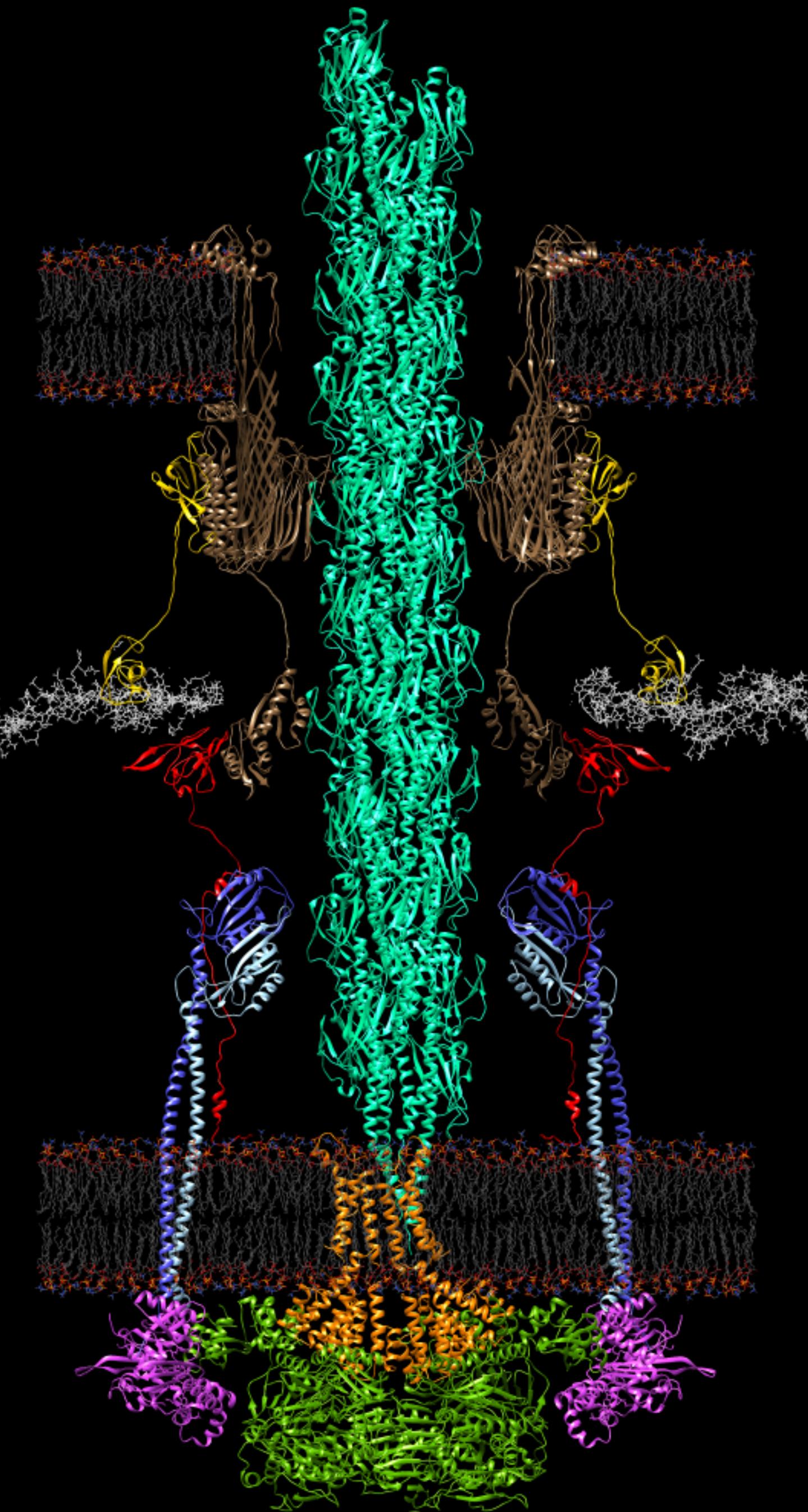
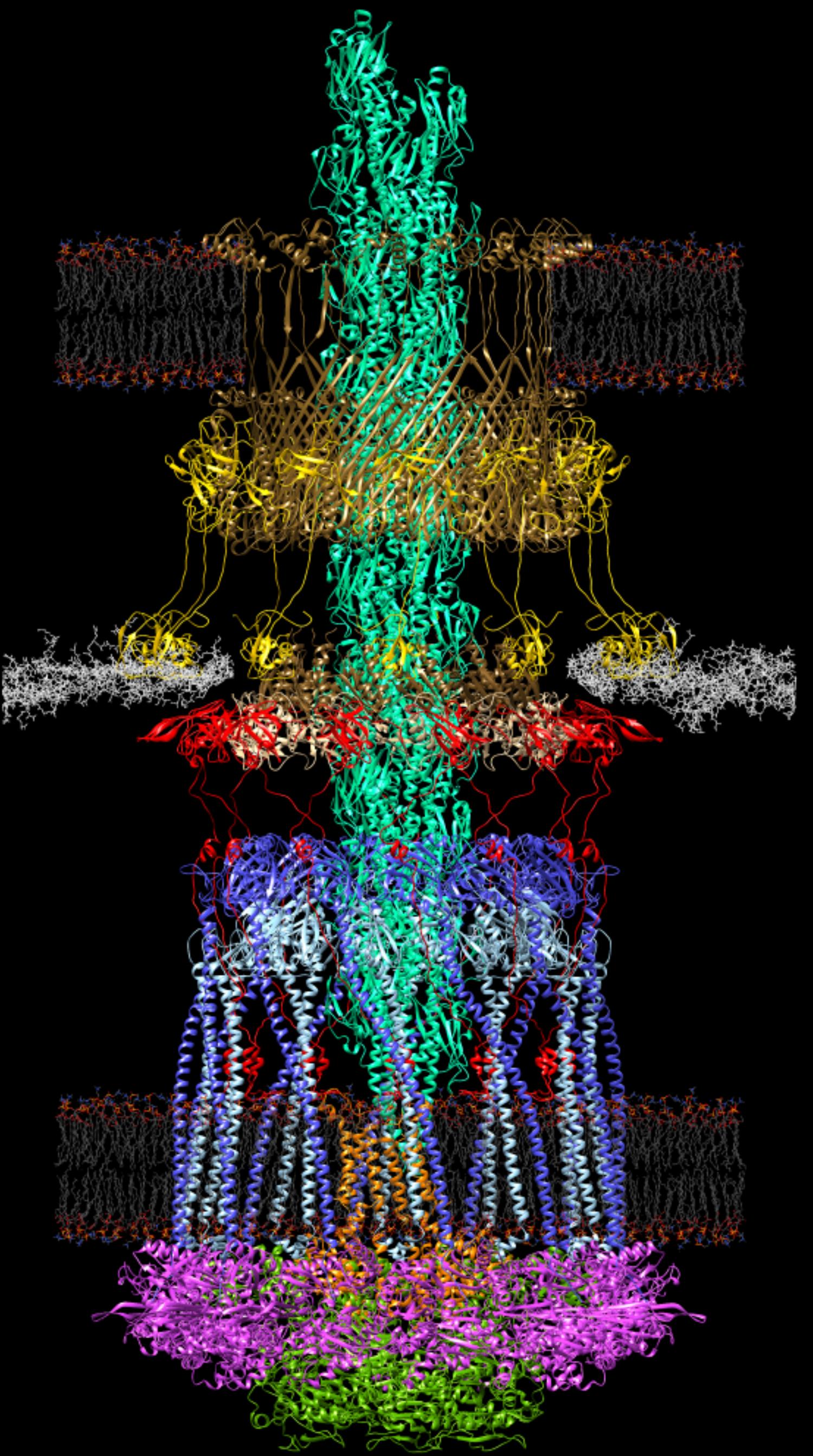




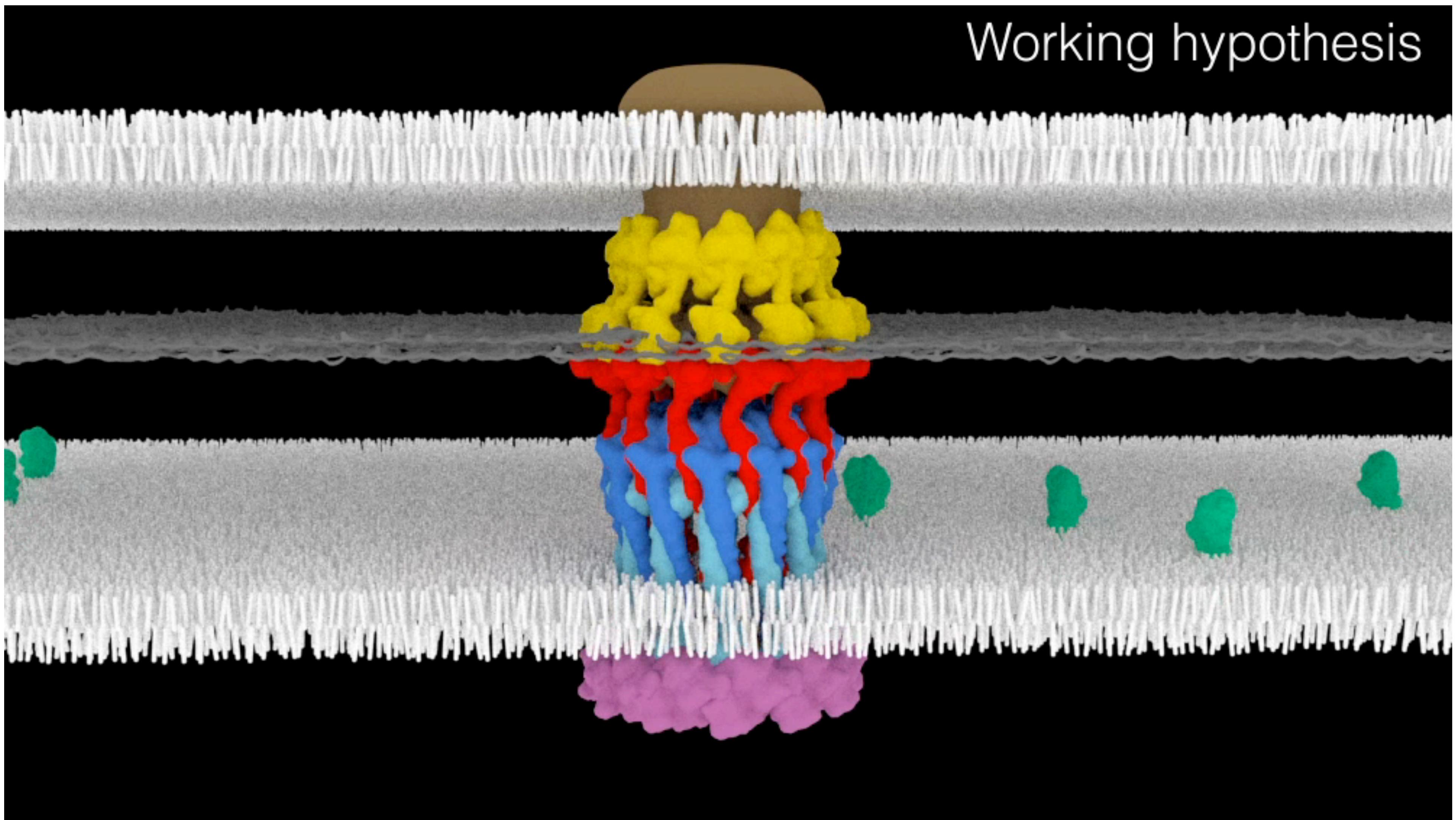


| | | |
|------|------|-----------|
| PilQ | PilA | PilN&PilO |
| TsaP | PilC | PilB/PilT |
| PilP | PilM | Membrane |





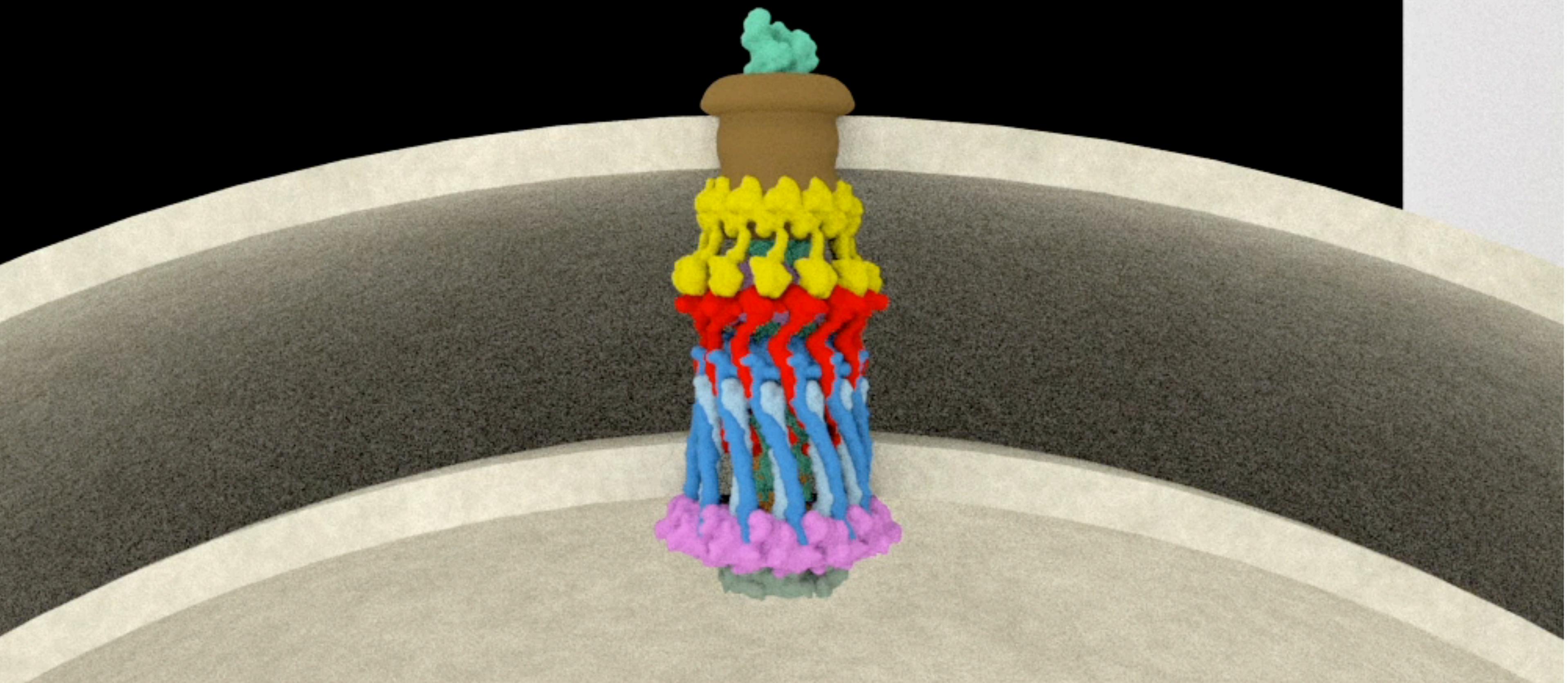
Working hypothesis



Animation by Janet Iwasa



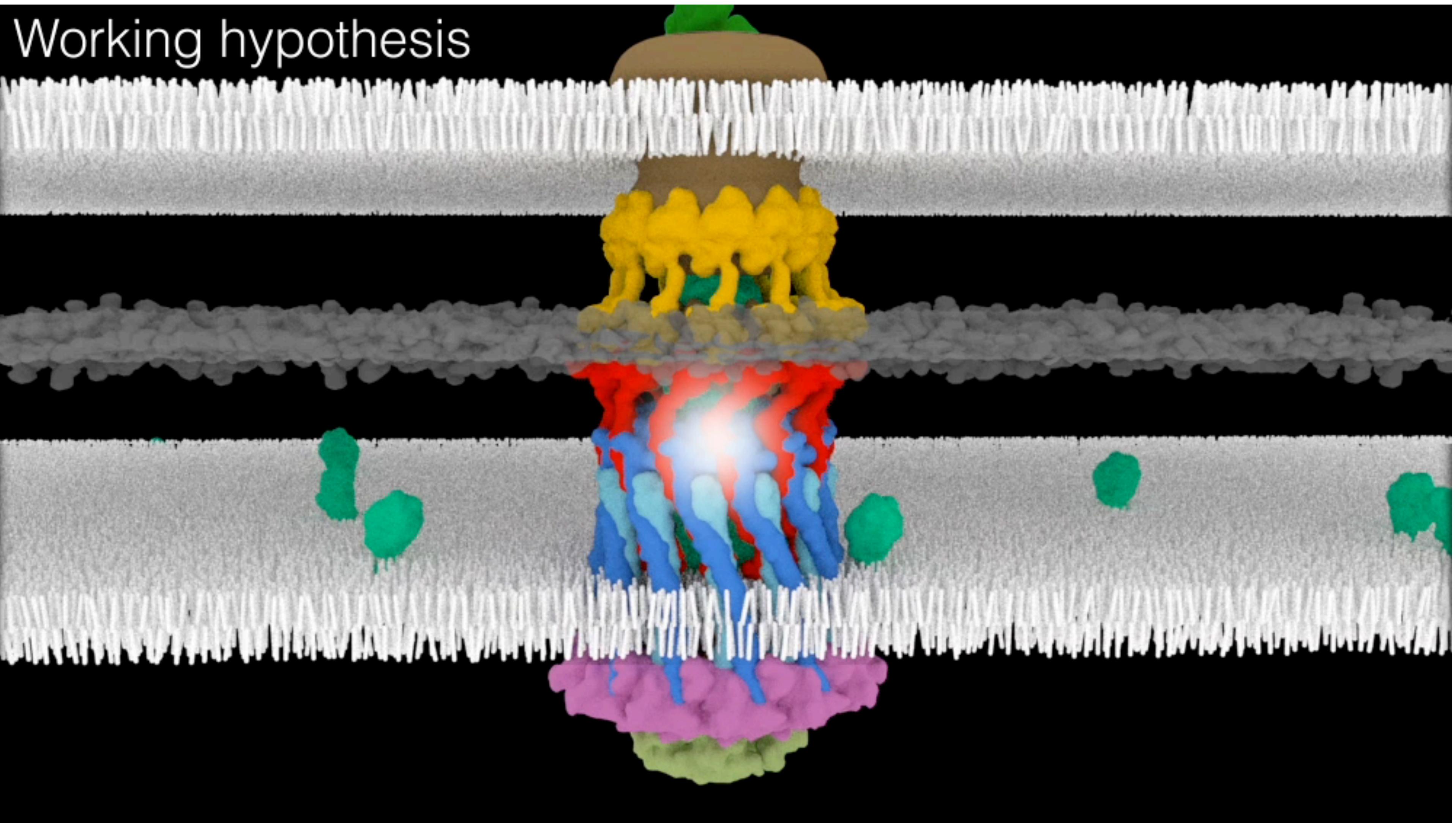
Working hypothesis

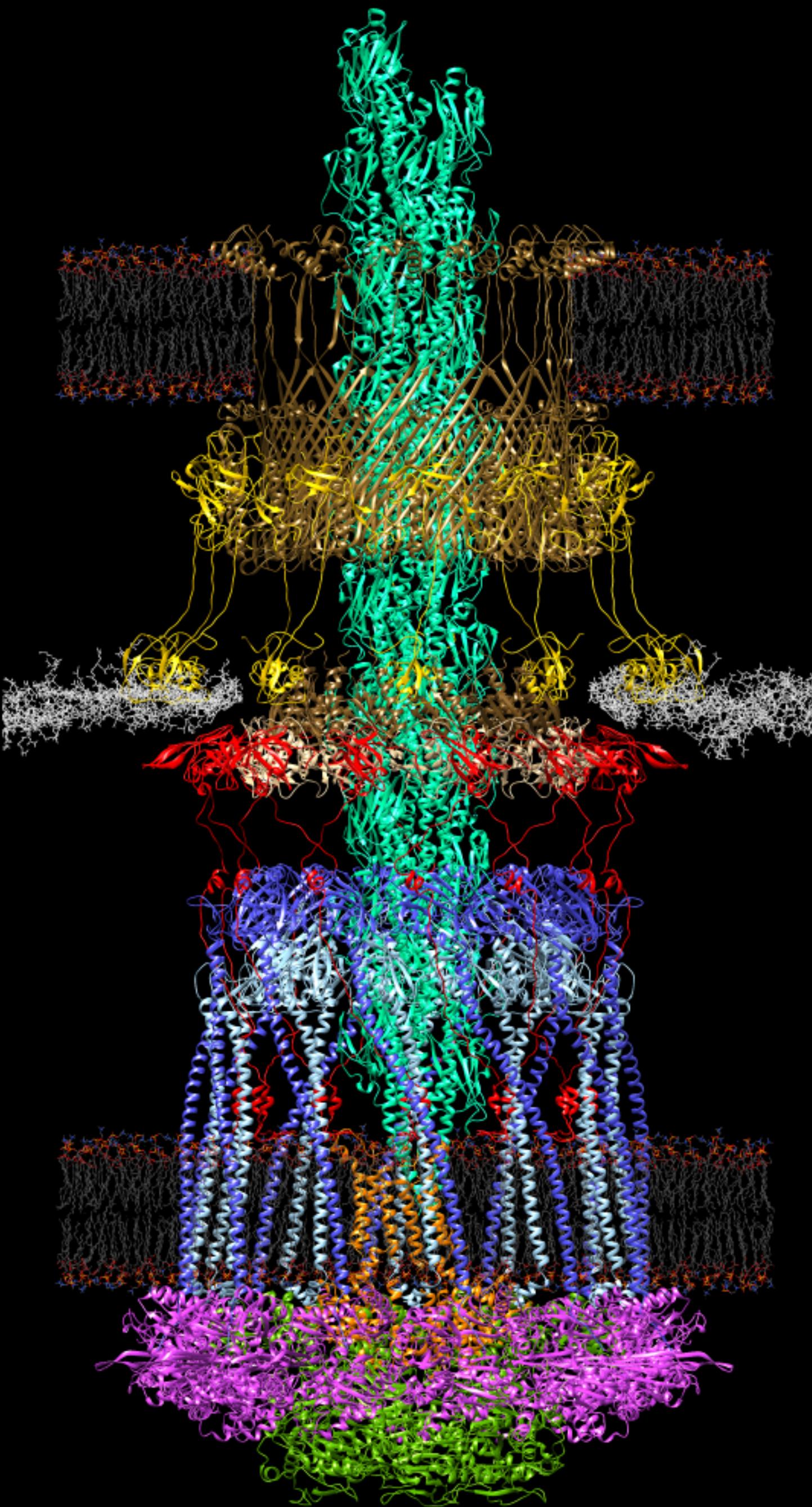


Working hypothesis



Working hypothesis





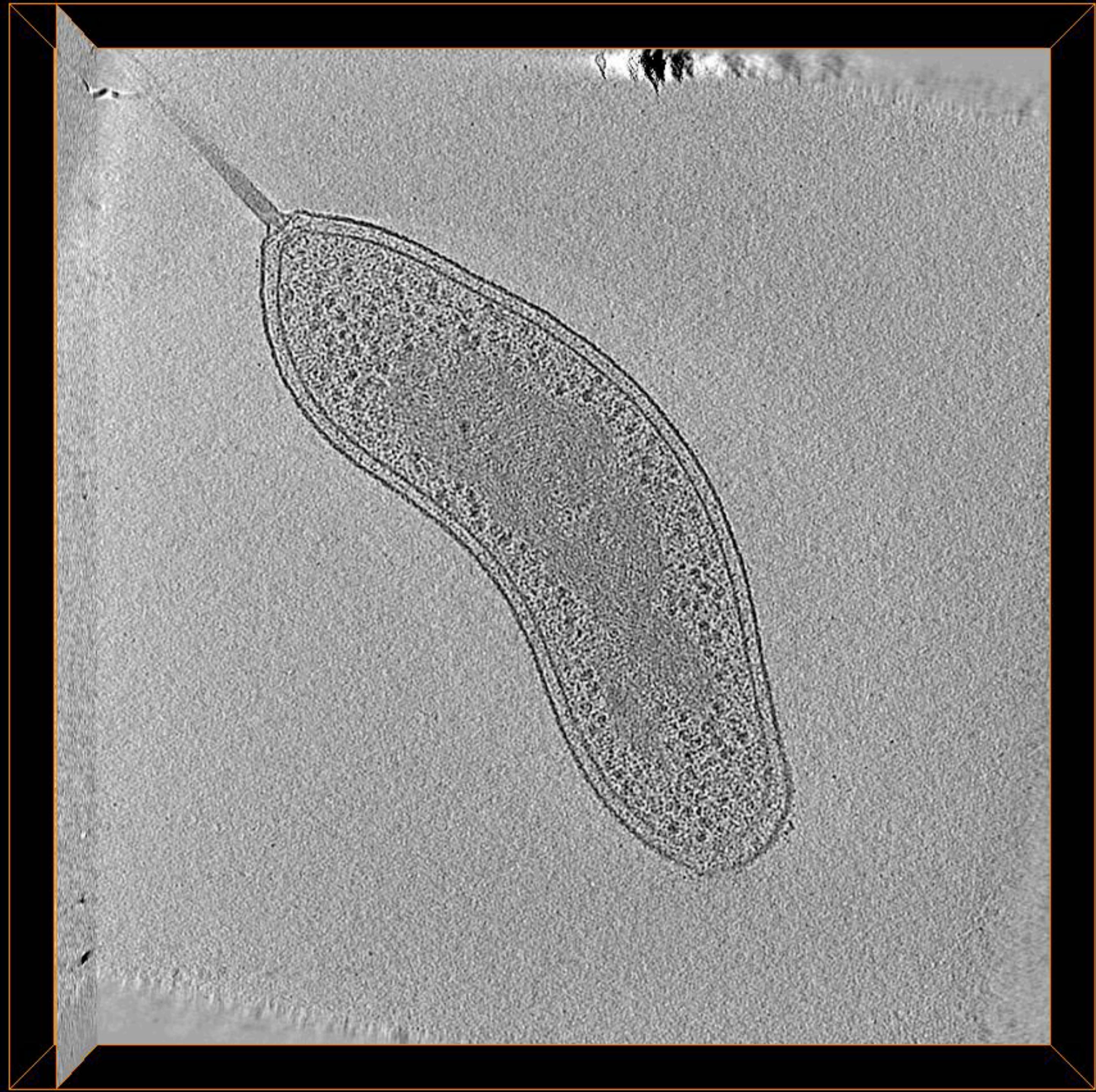
Yi-Wei
Chang

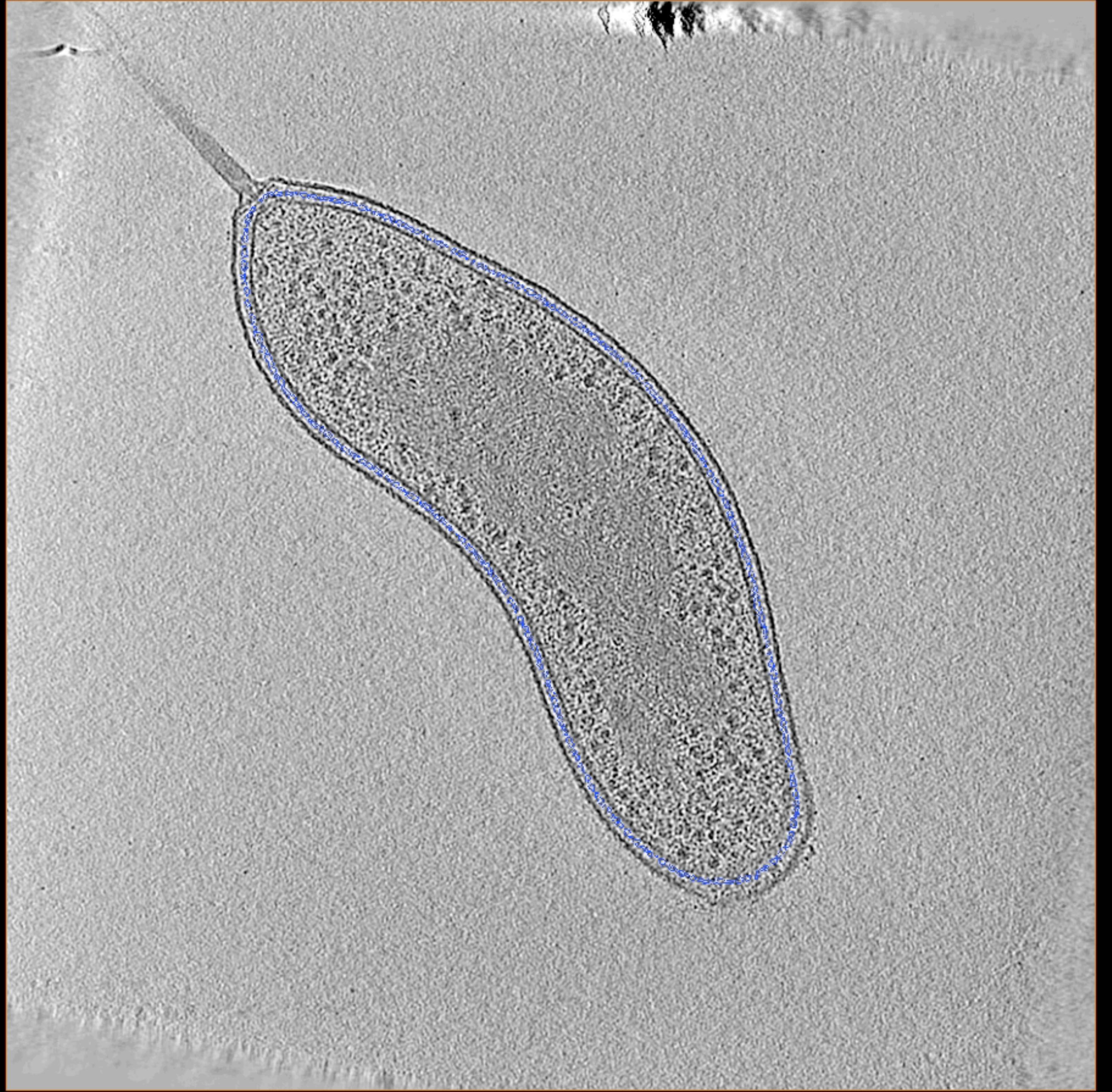


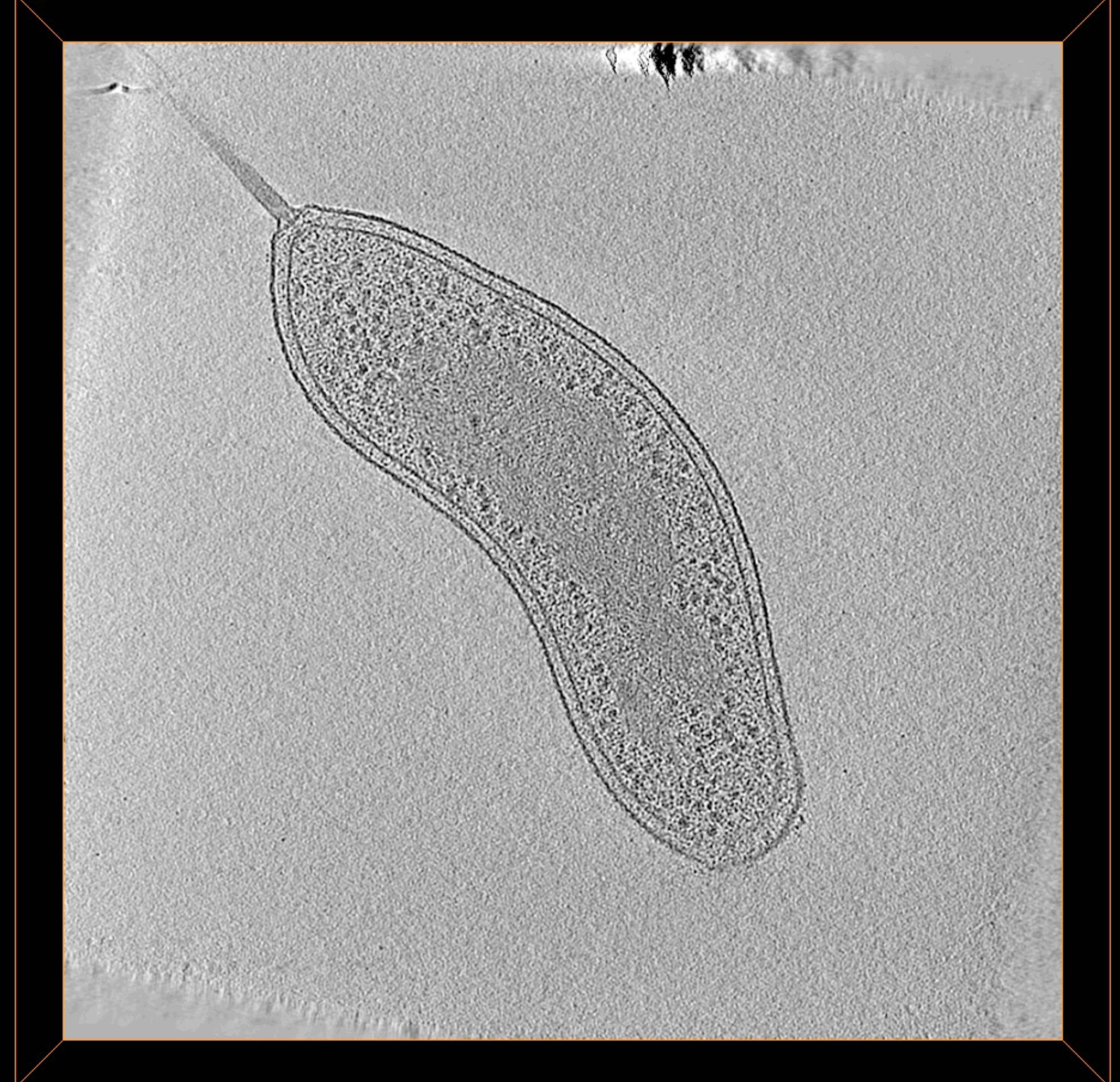
Sara
Weaver

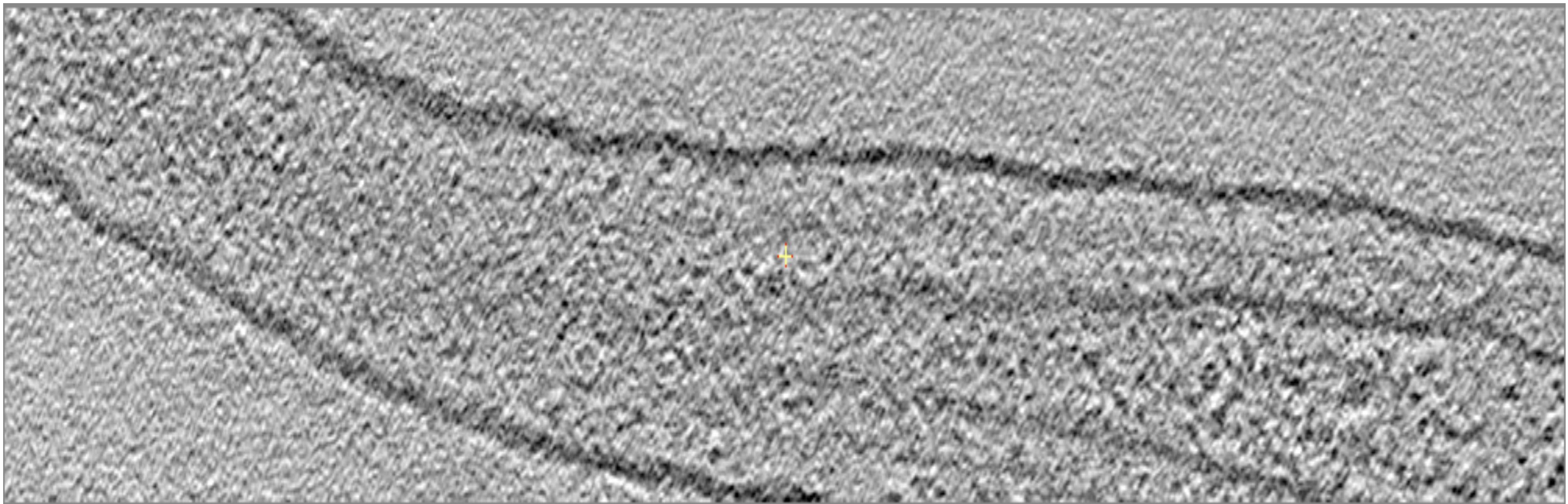


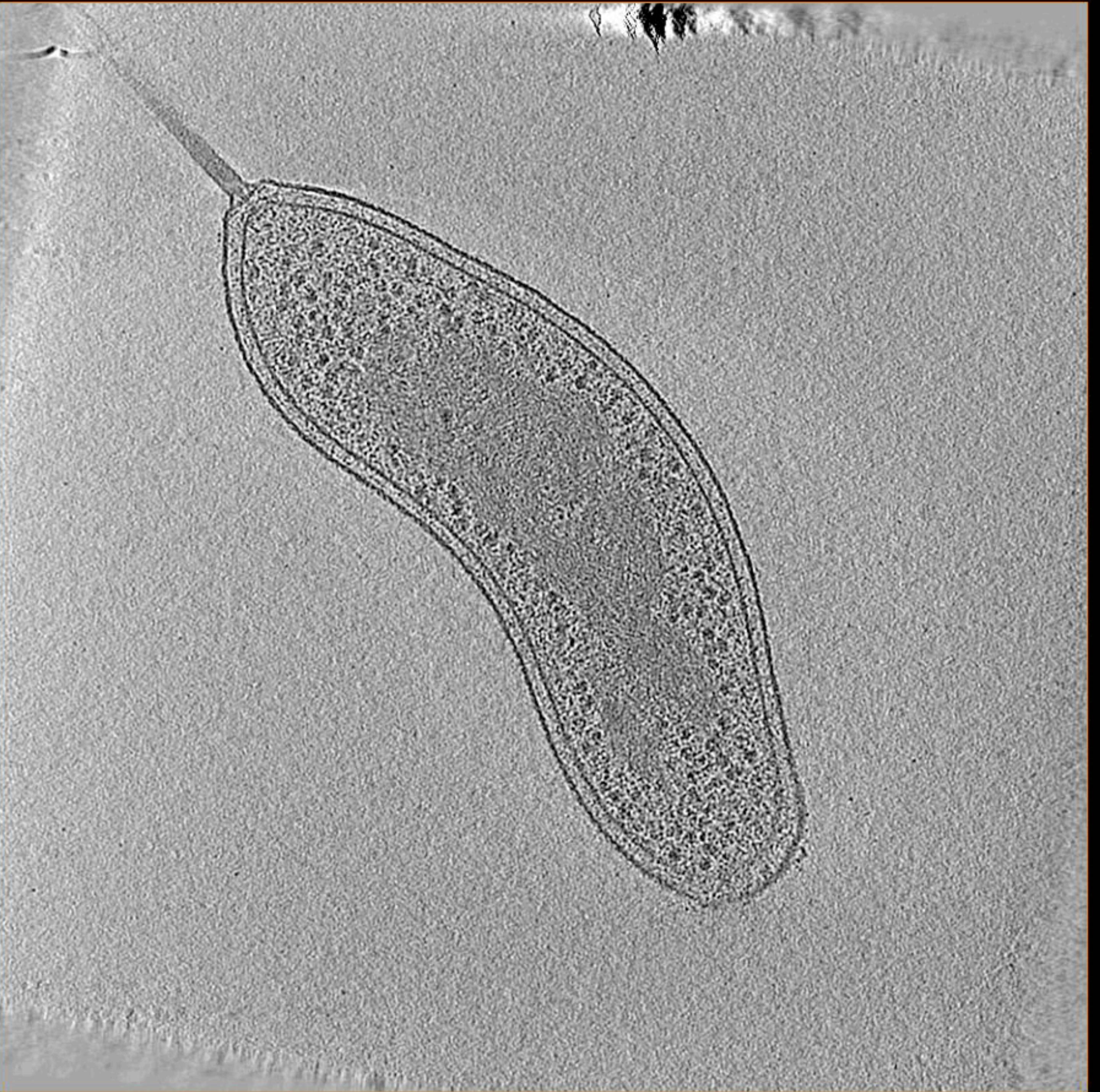
Lotte
Søgaard-Andersen



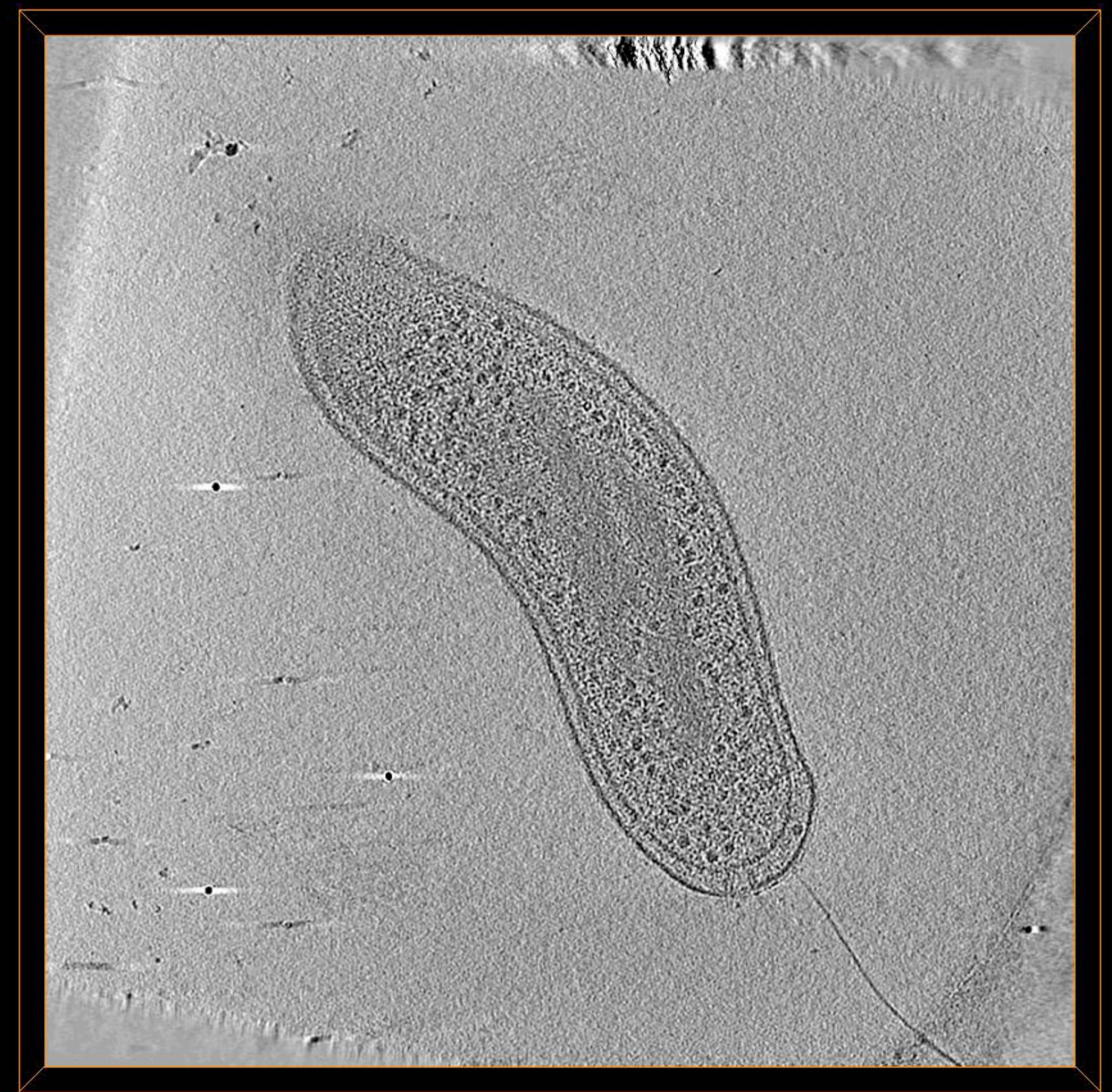


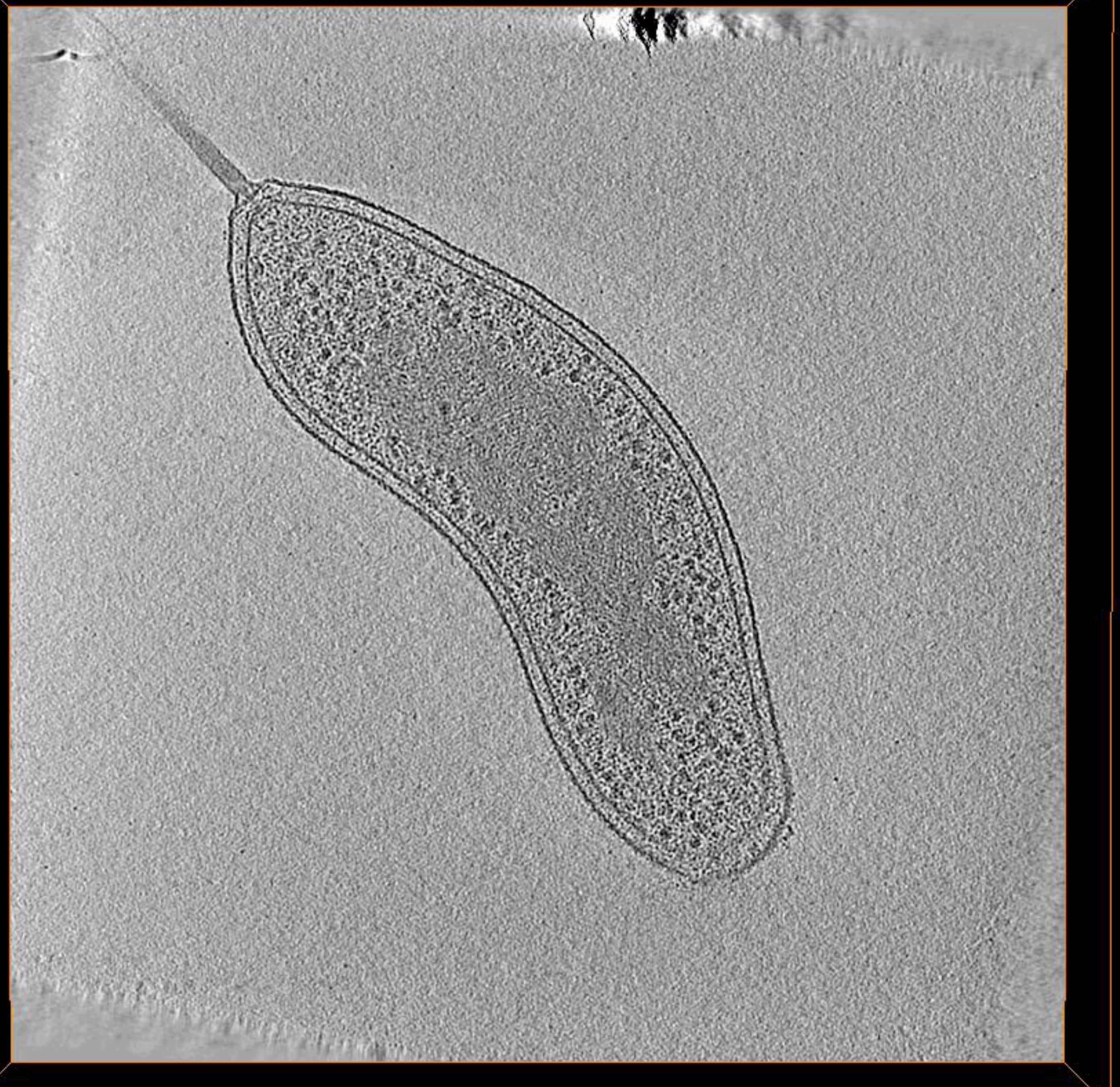


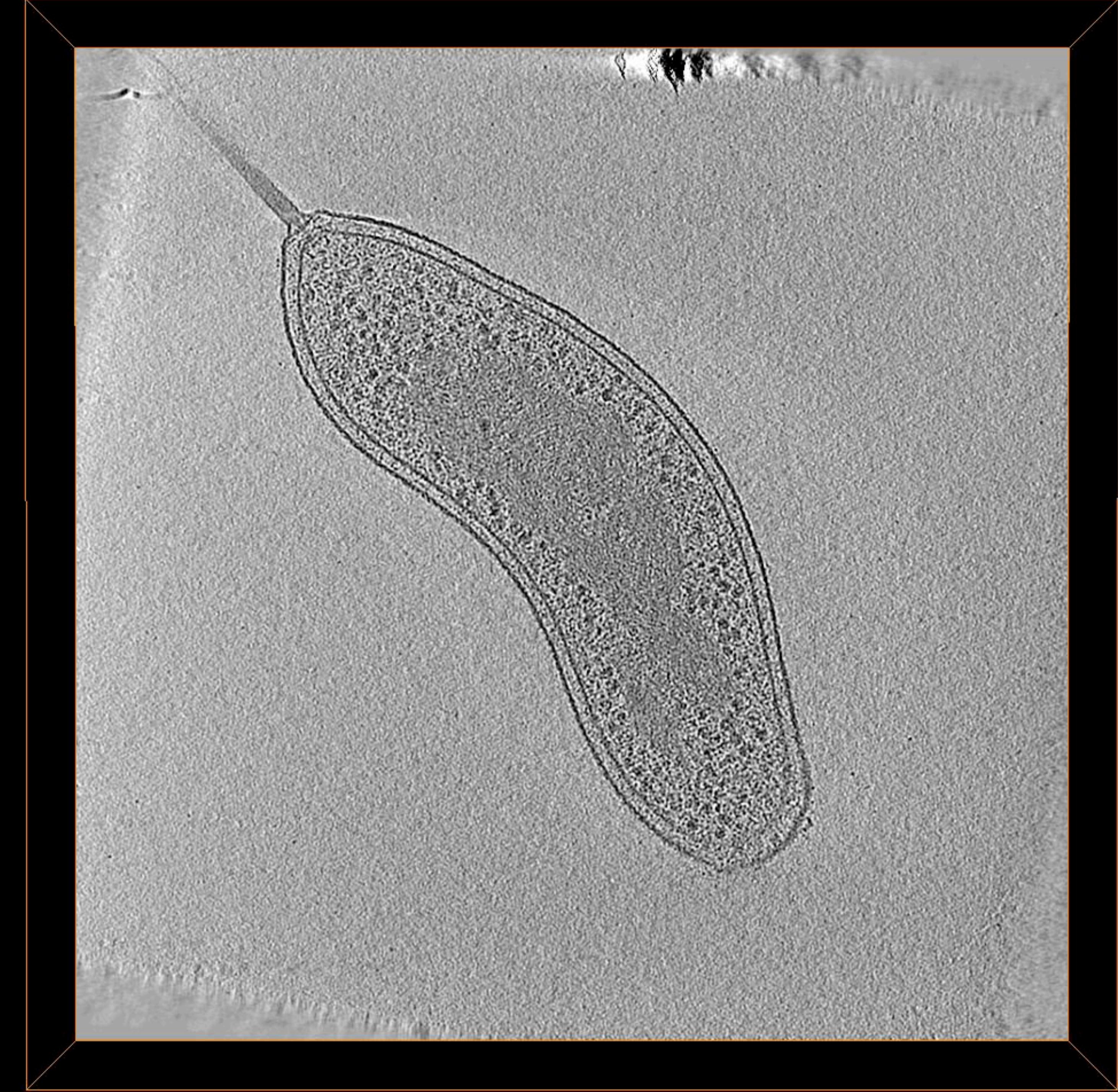












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by Catherine M. Oikonomou & Grant J. Jensen

This open access digital textbook offers a tour of microbial cells guided by cutting-edge 3D electron microscopy.

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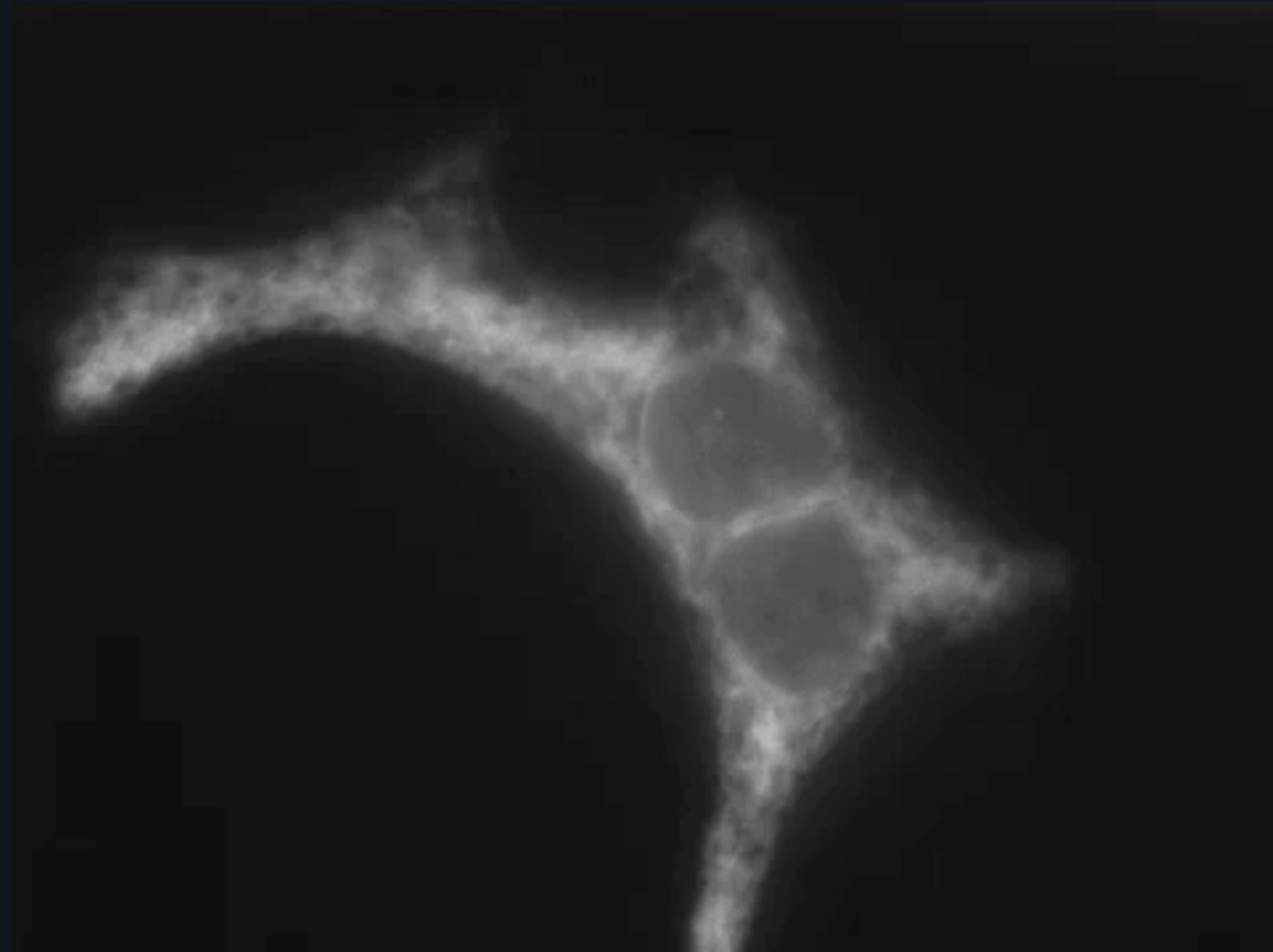
by Catherine M. Oikonomou & Grant J. Jensen

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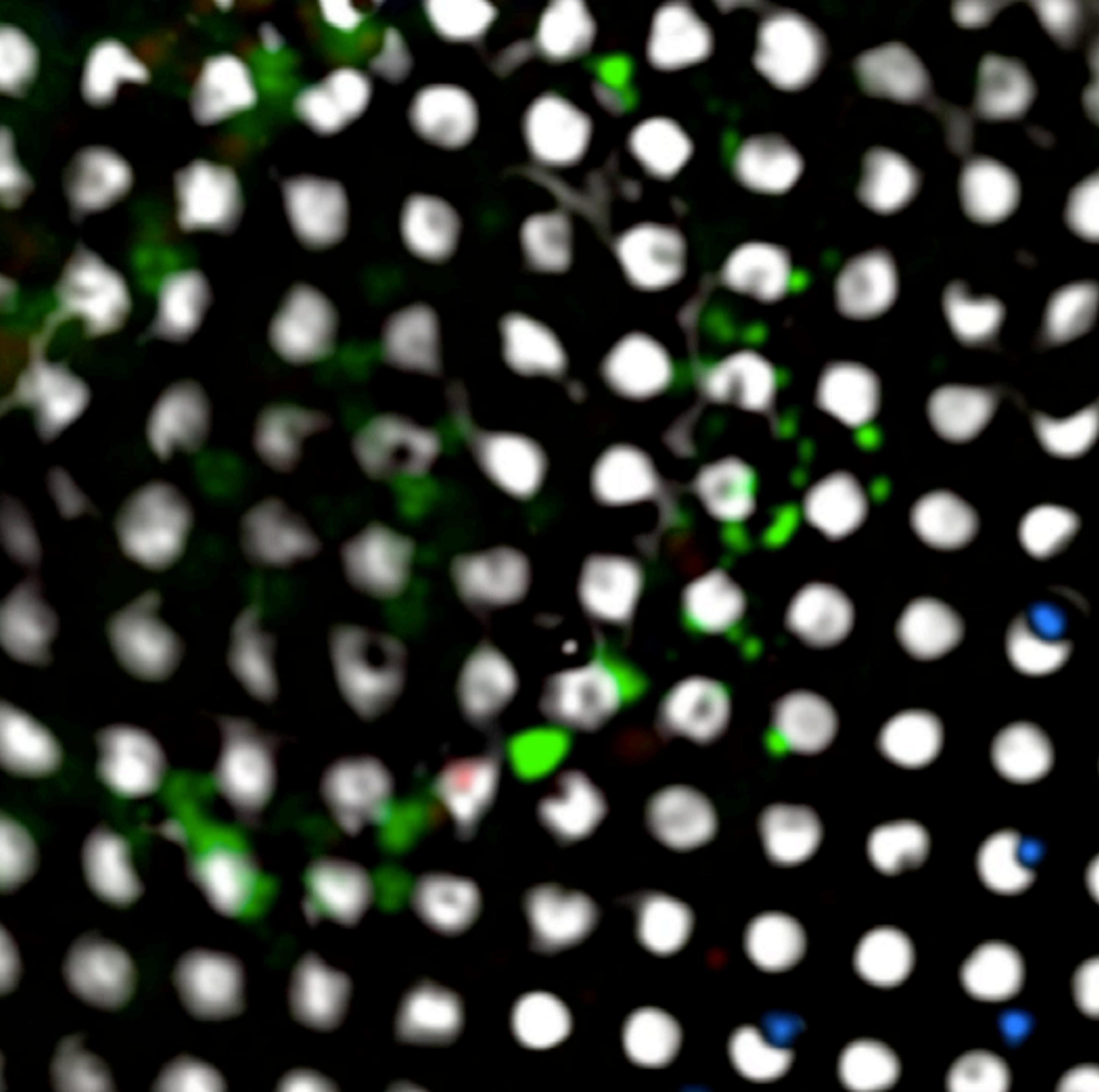
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<https://www.cellstructureatlas.org/>

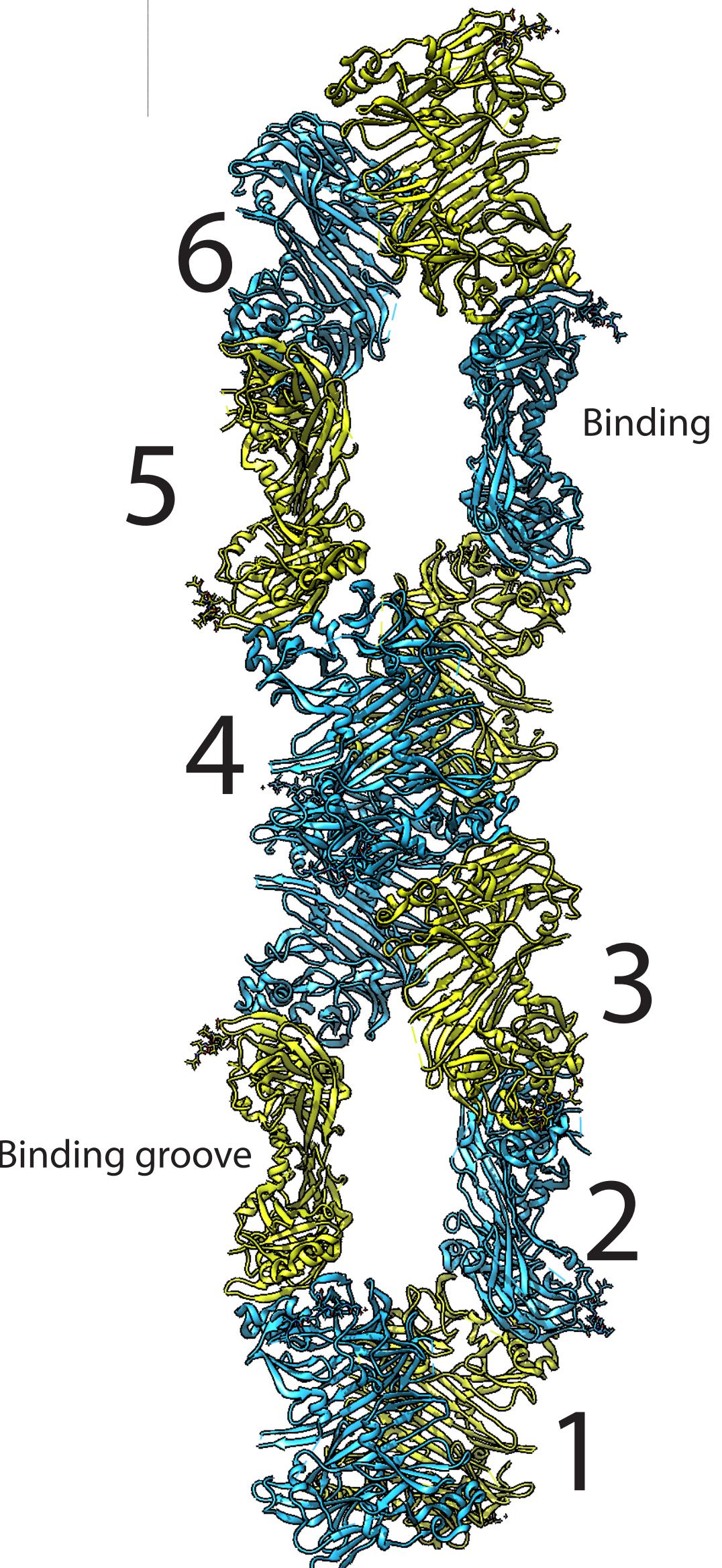
Evidence of IRE1 α oligomerization *in vivo*



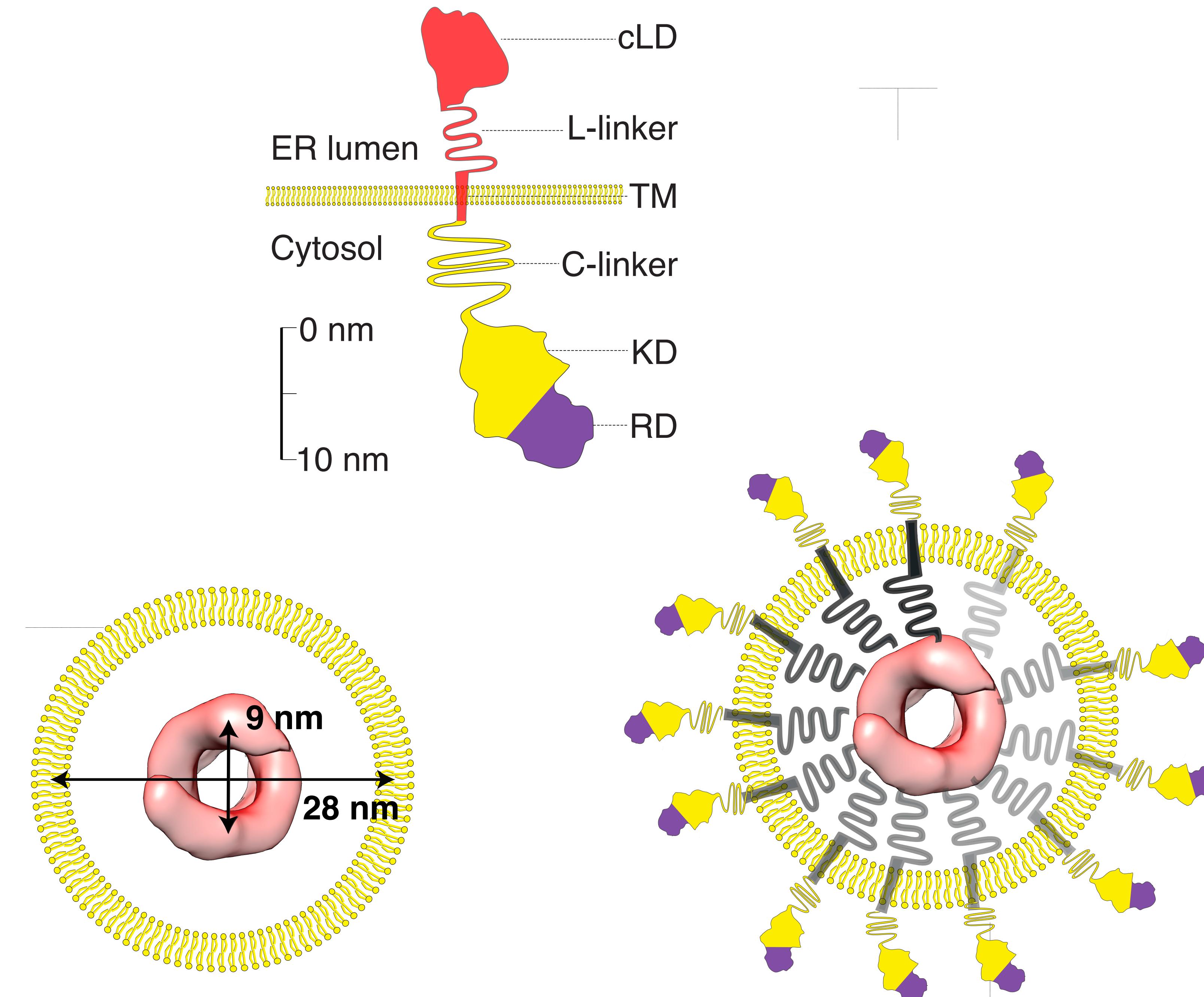
Walter and Ron, 2011; Li *et al*, 2010; Aragón *et al*, 2009







Yeast
crystal
structure





Stephen
Carter
Caltech

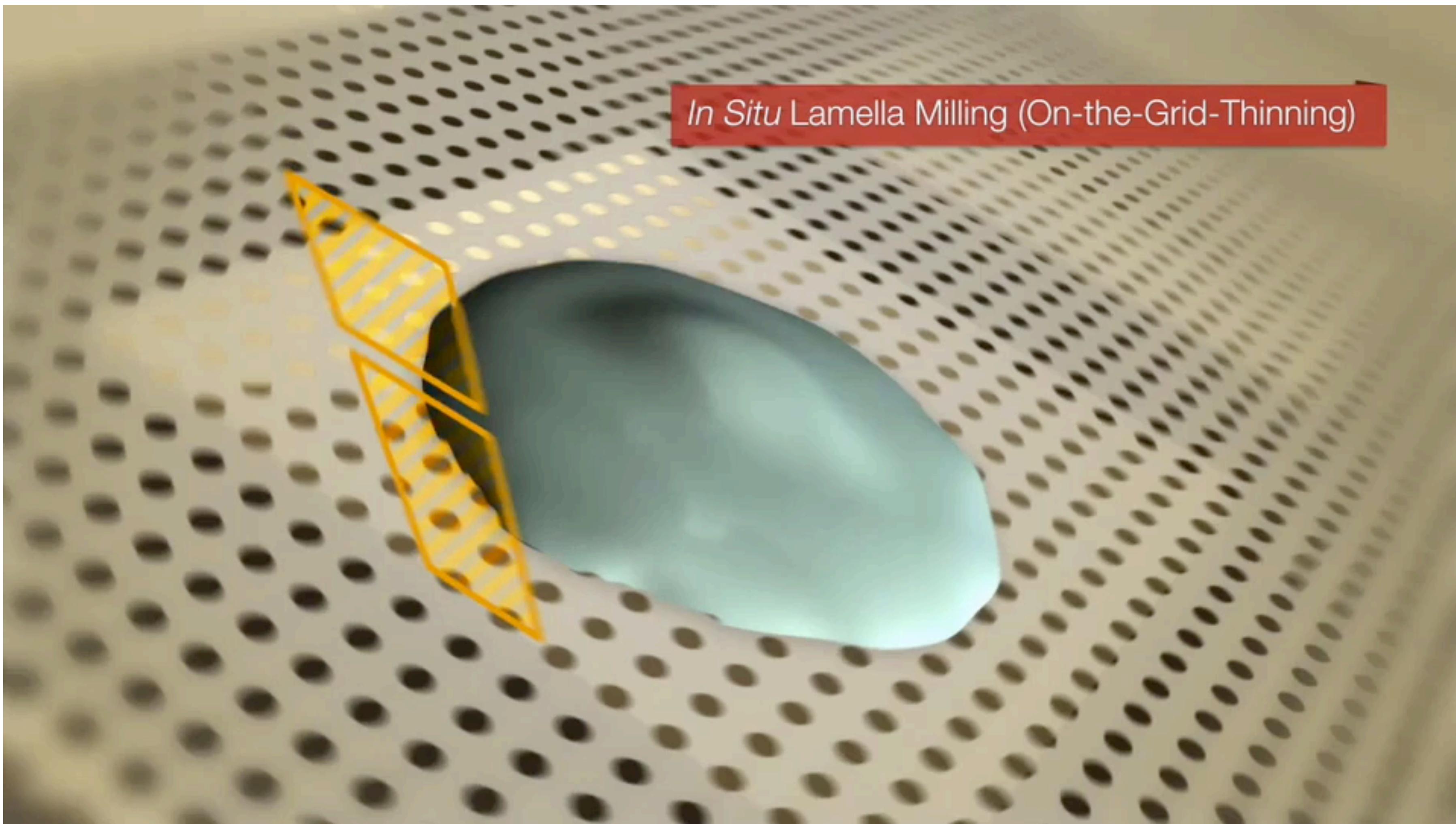


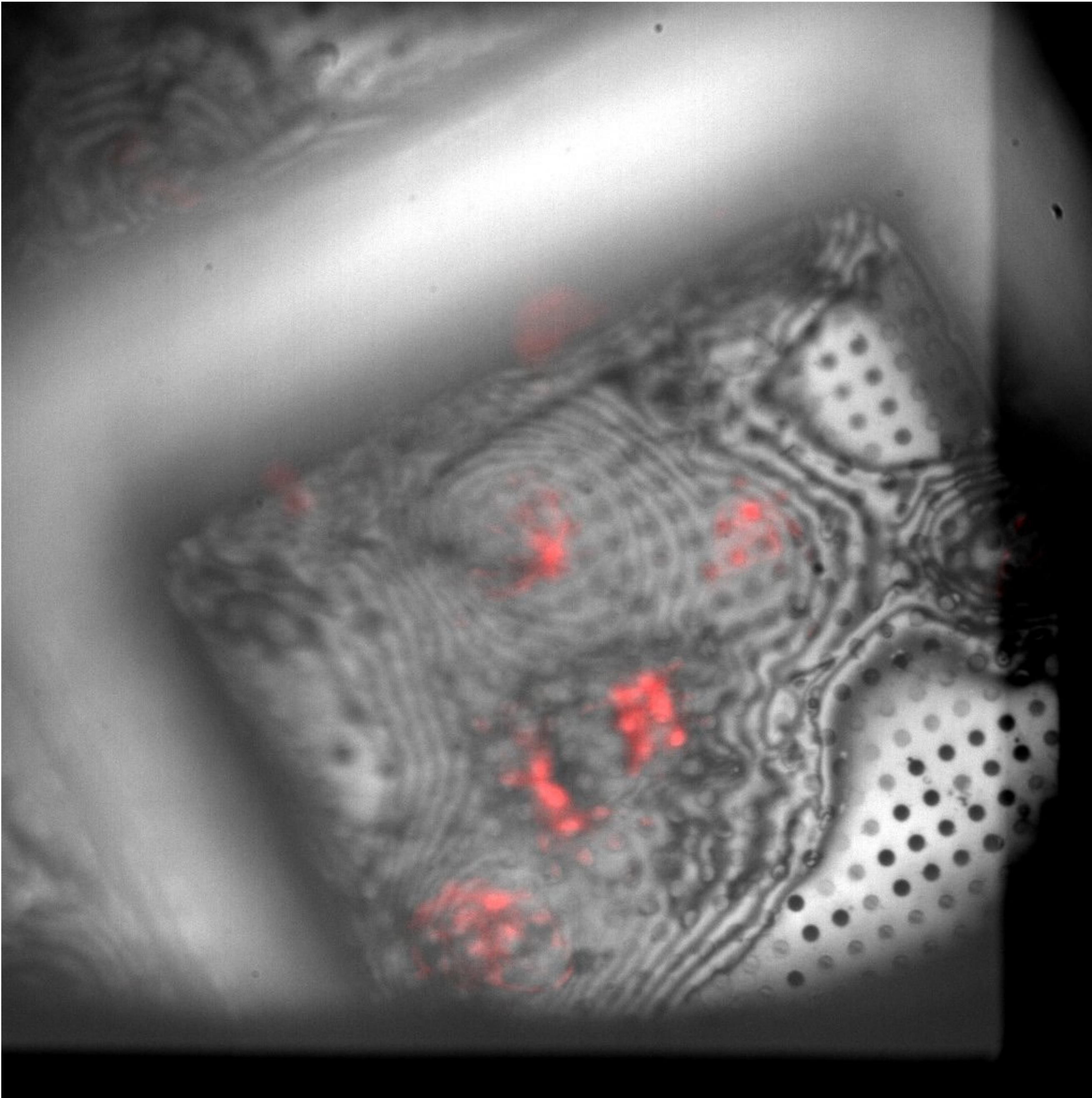
Han
Tran
UCSF



Peter
Walter
UCSF

Focused ion beam (FIB) milling

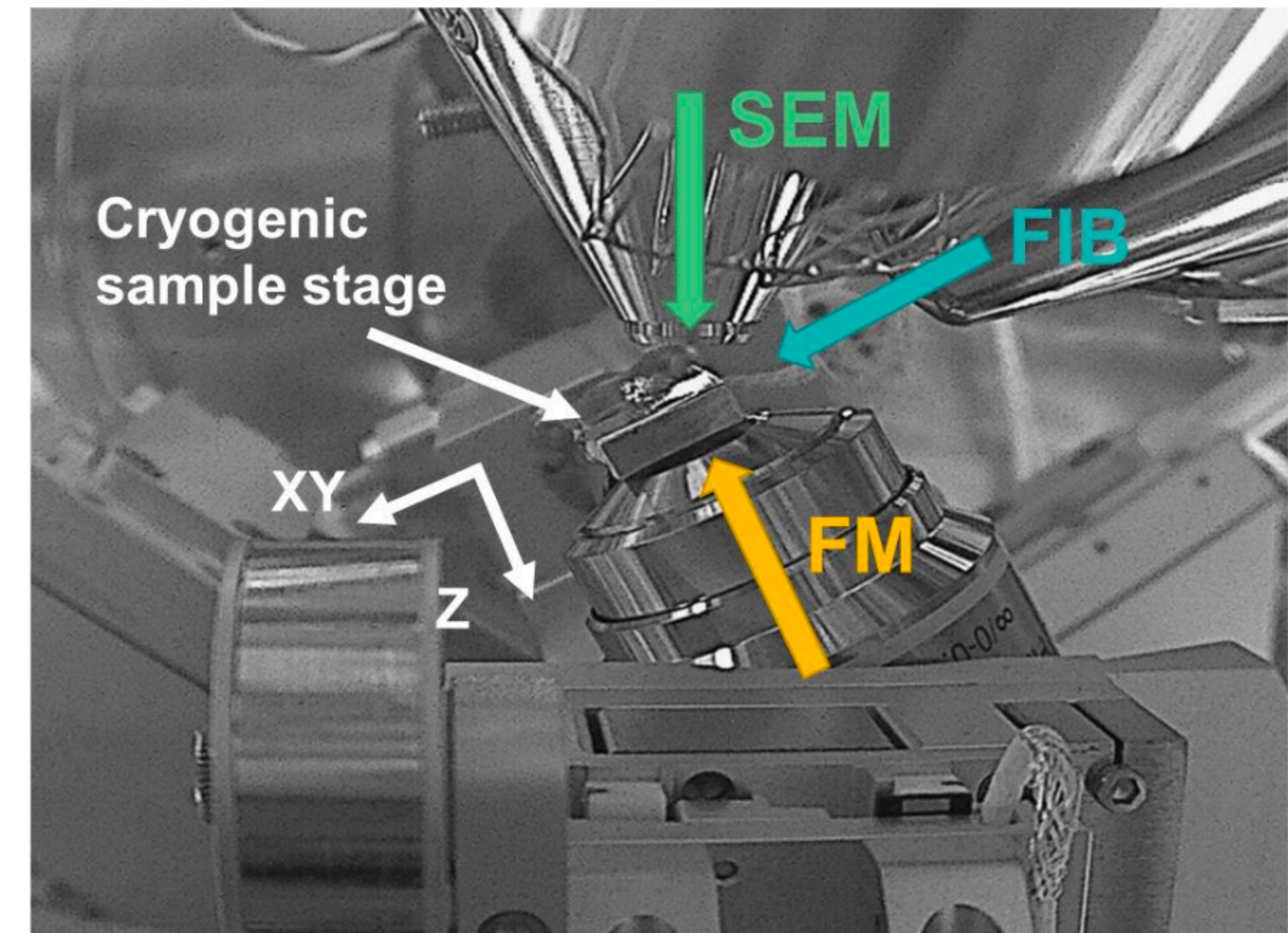
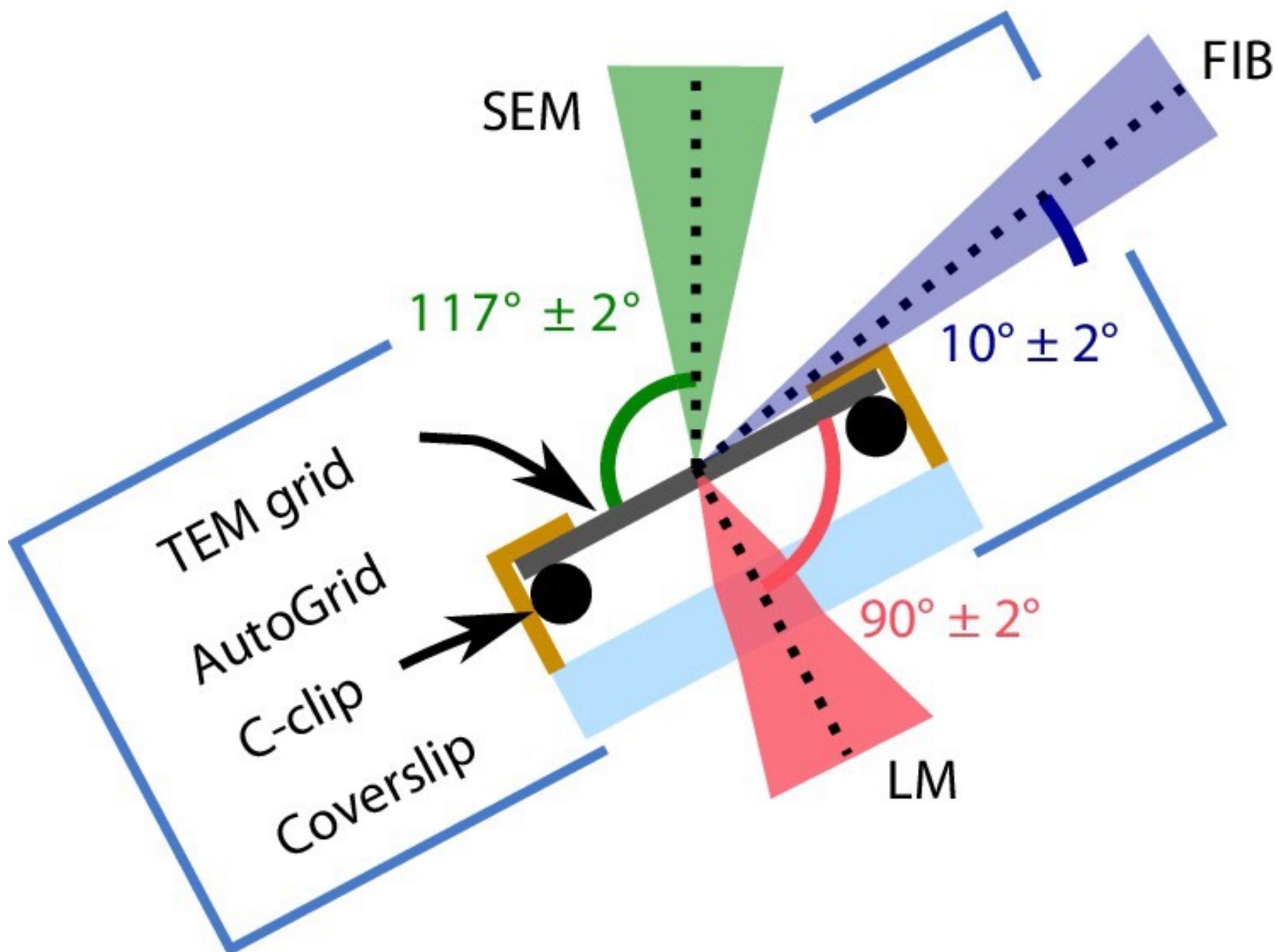


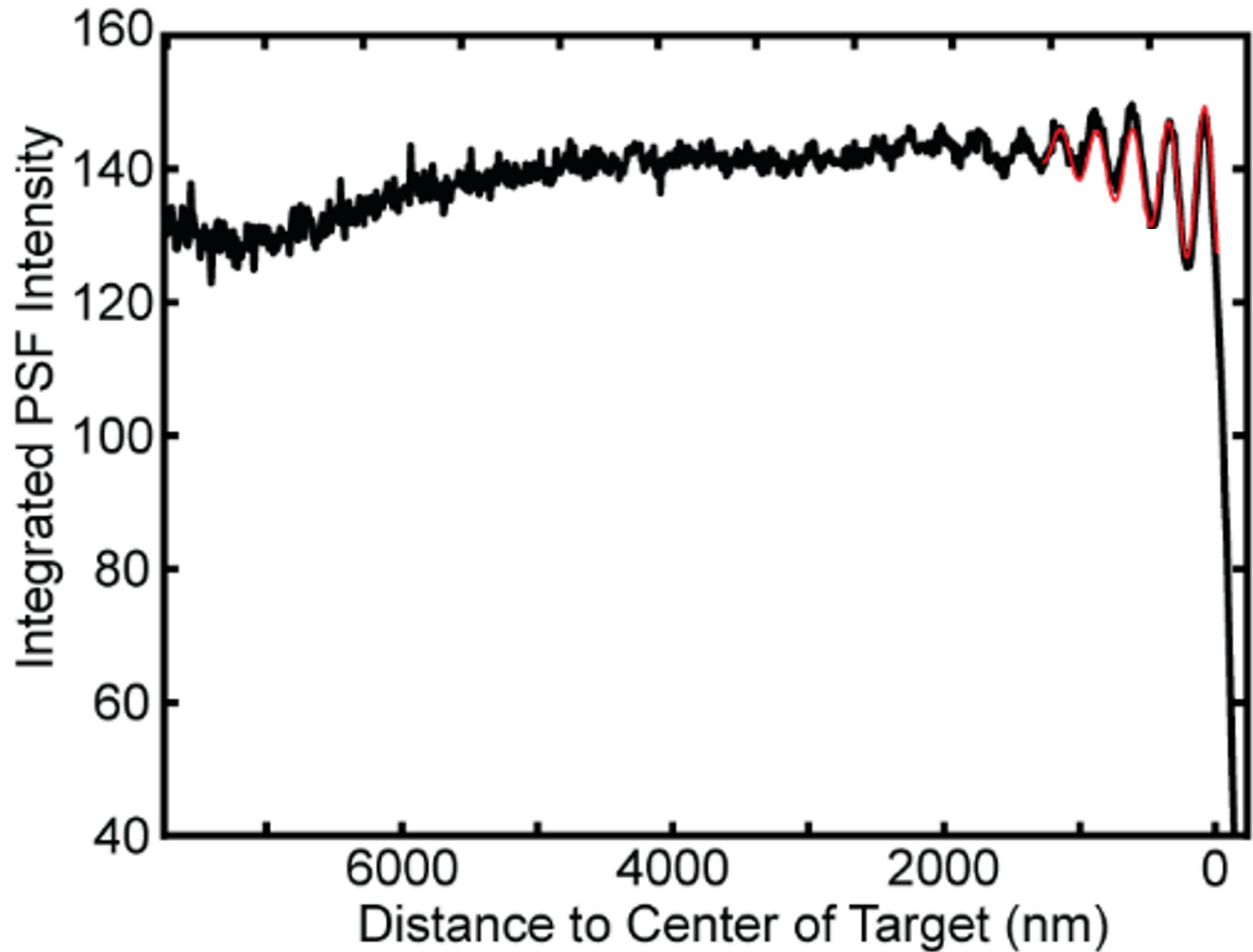
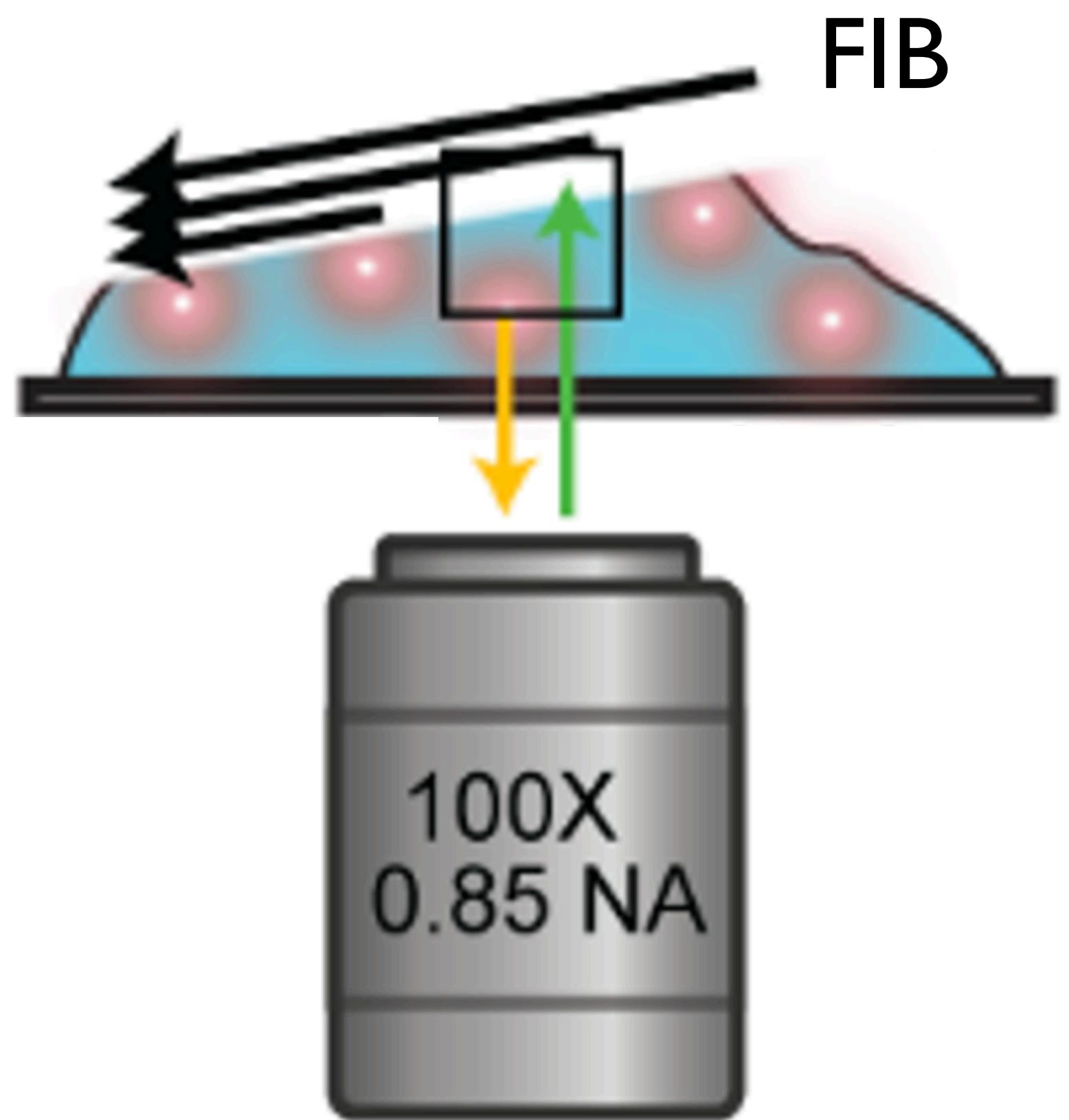


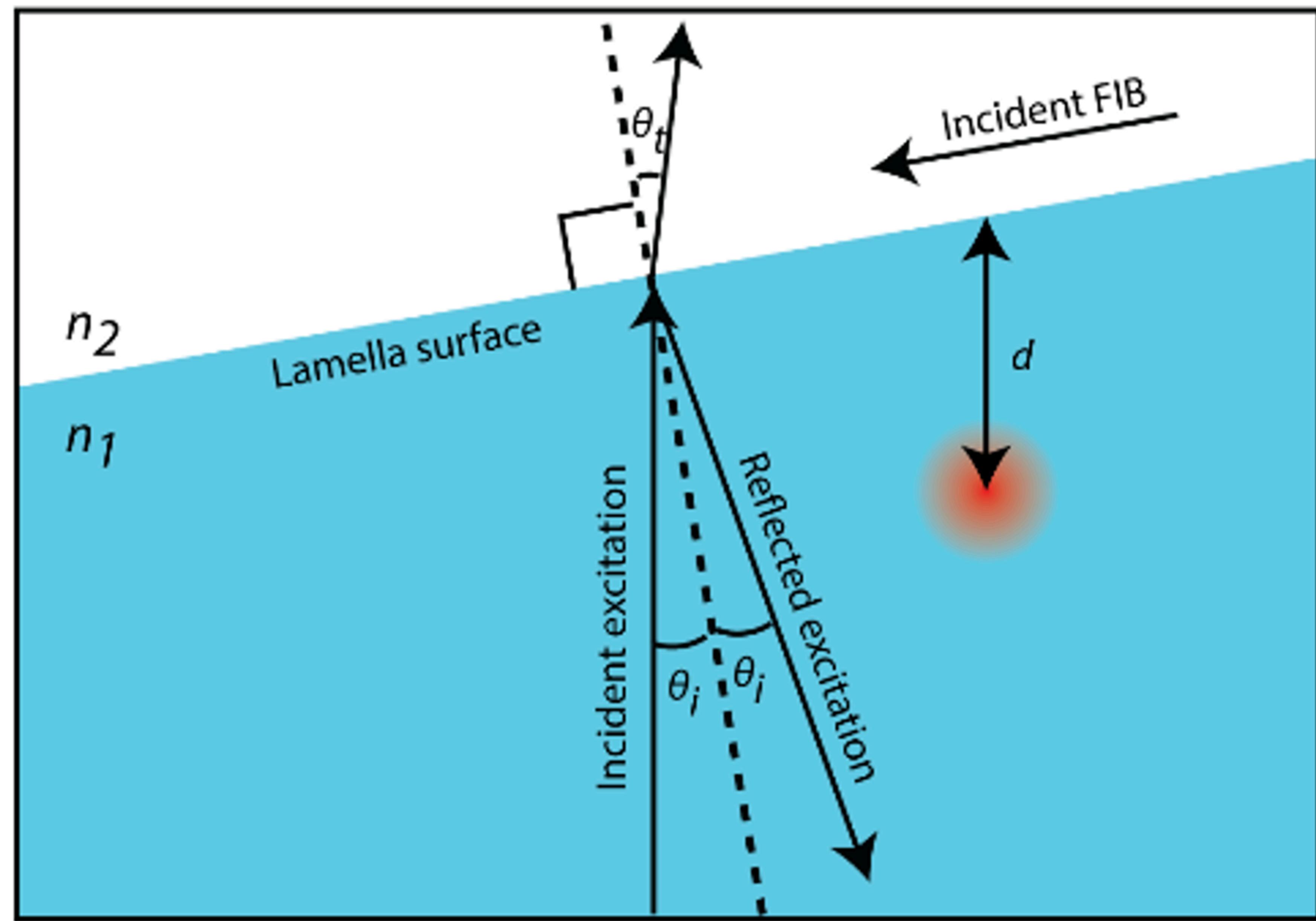
It's easy enough to target
milling in x and y,
but z is hard

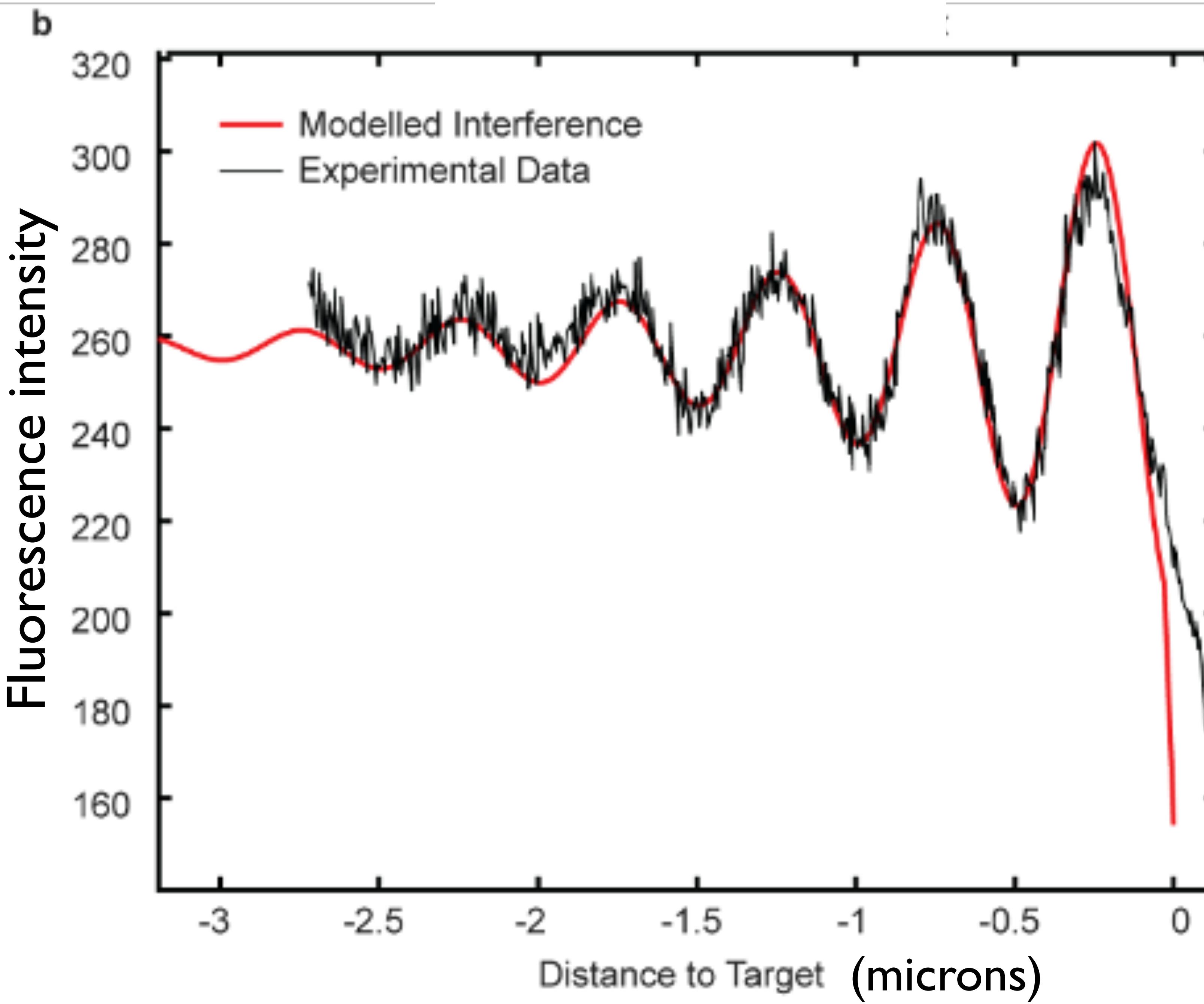
ENZEL

Cryo-FIB stage









Strategy:
Mill from top until close
Finish milling from below



Peter Dahlberg
Stanford

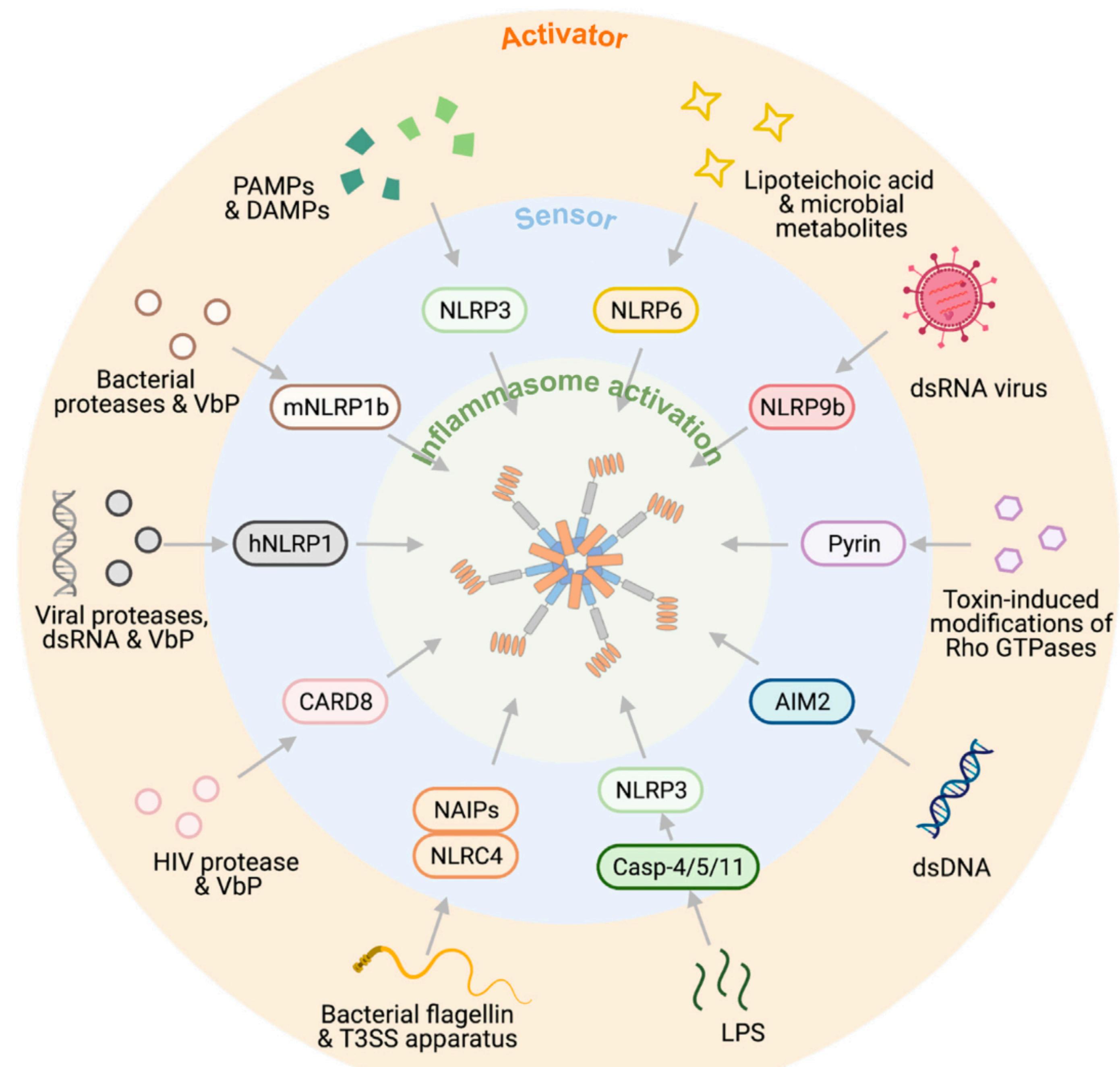


Phyllis Wang
Caltech/BYU/Stanford



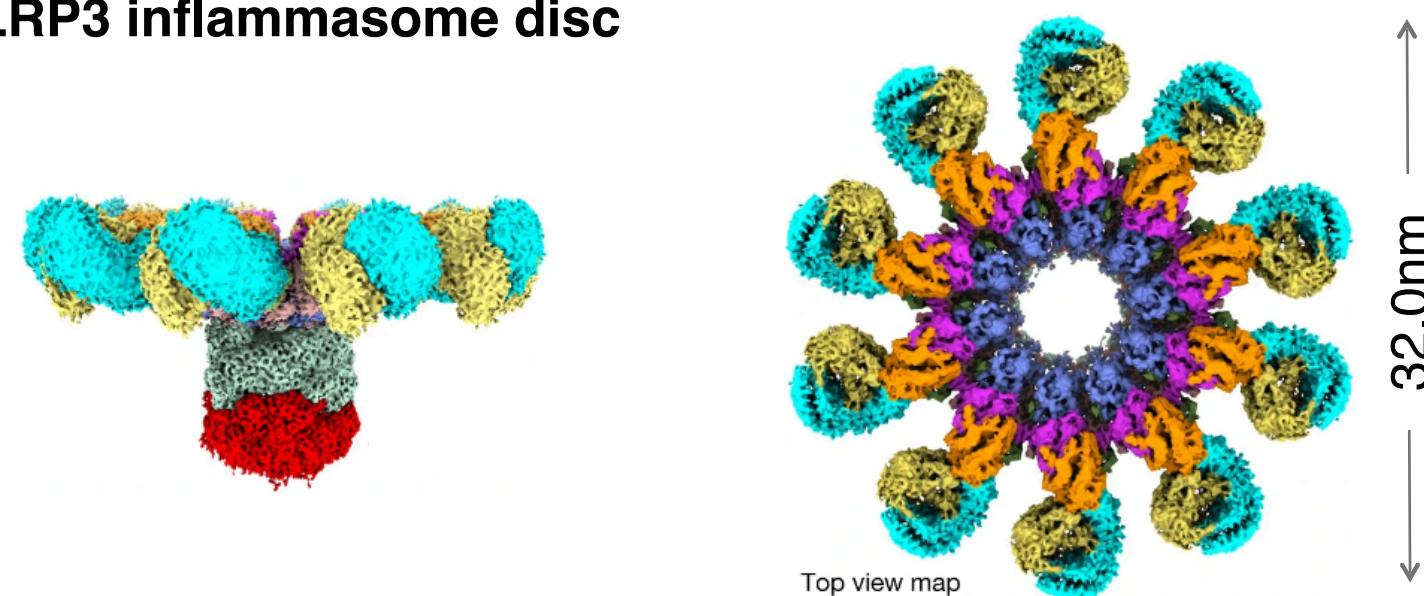
Anthony Sica
Stanford

NLRP3 inflammasome

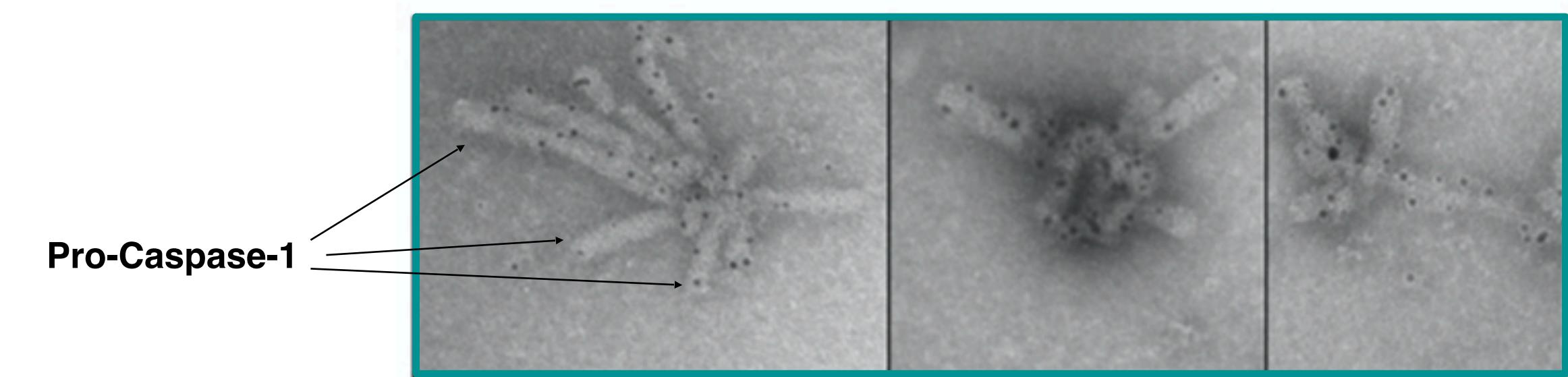
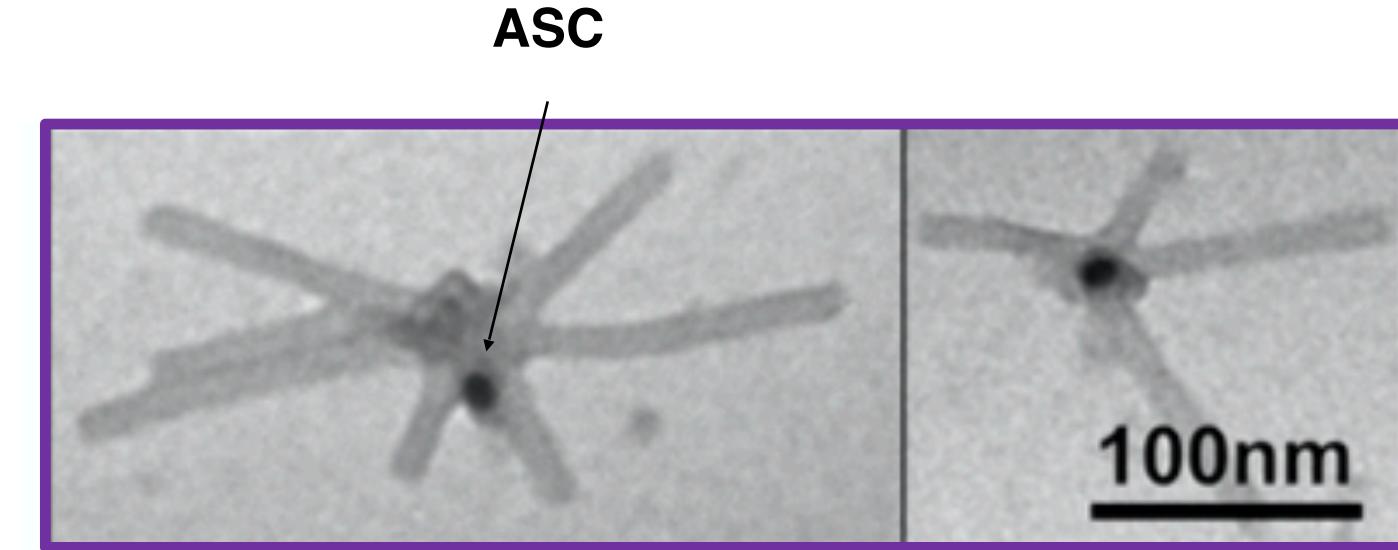
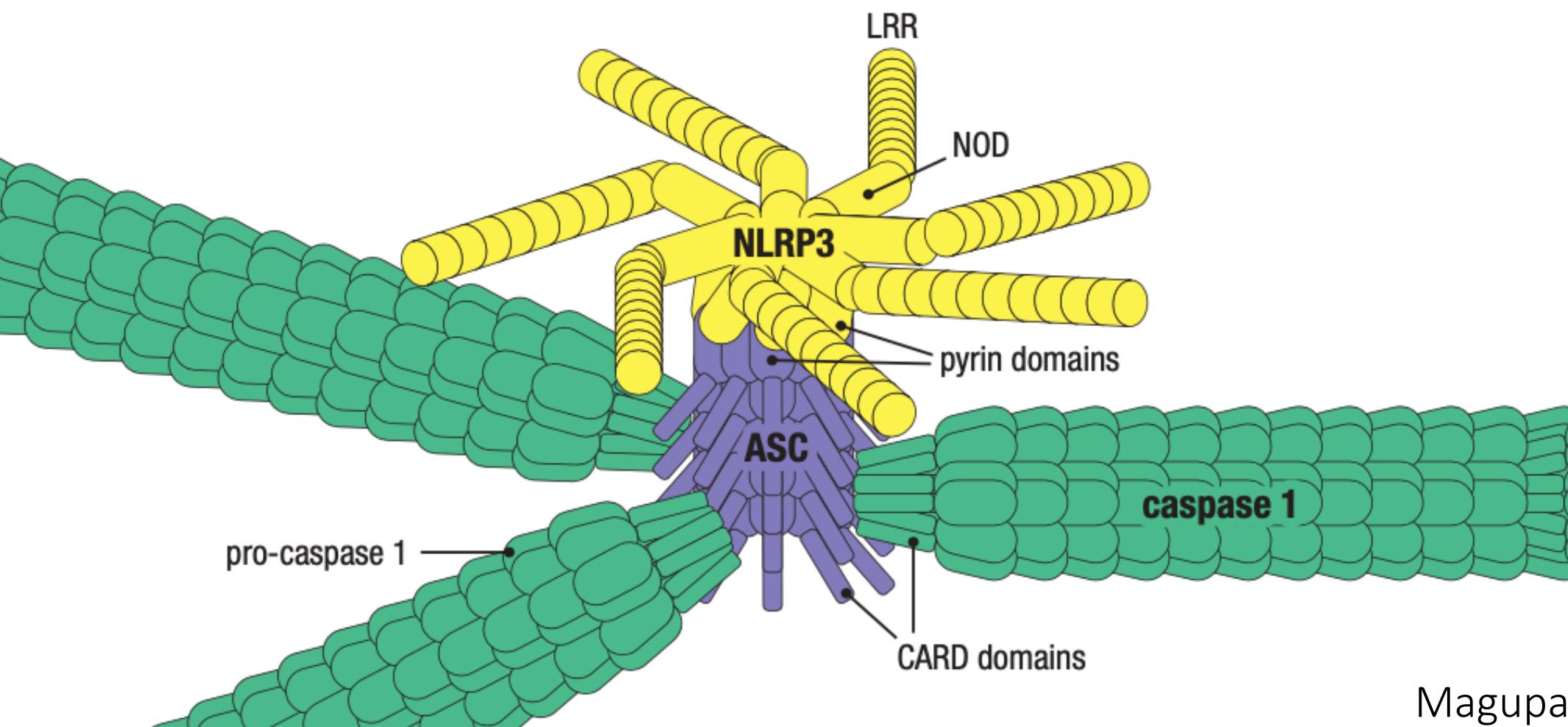
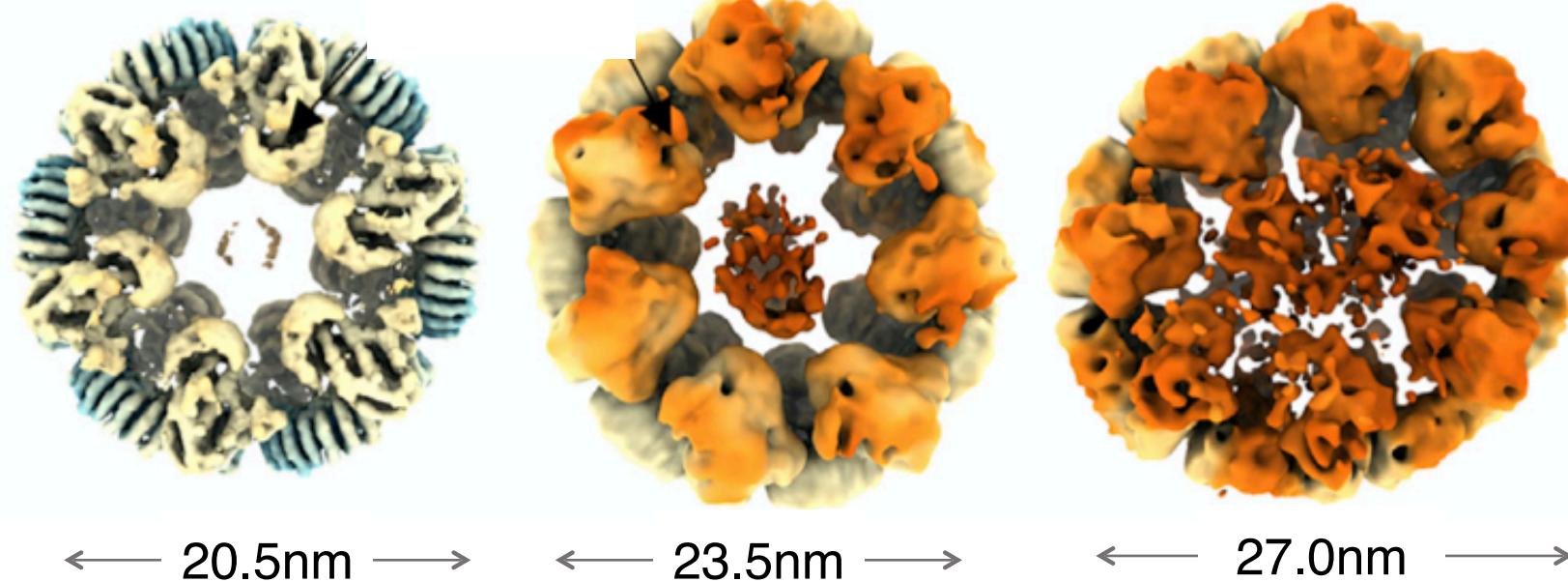


Structures of the NLRP3 inflammasome

NLRP3 inflammasome disc

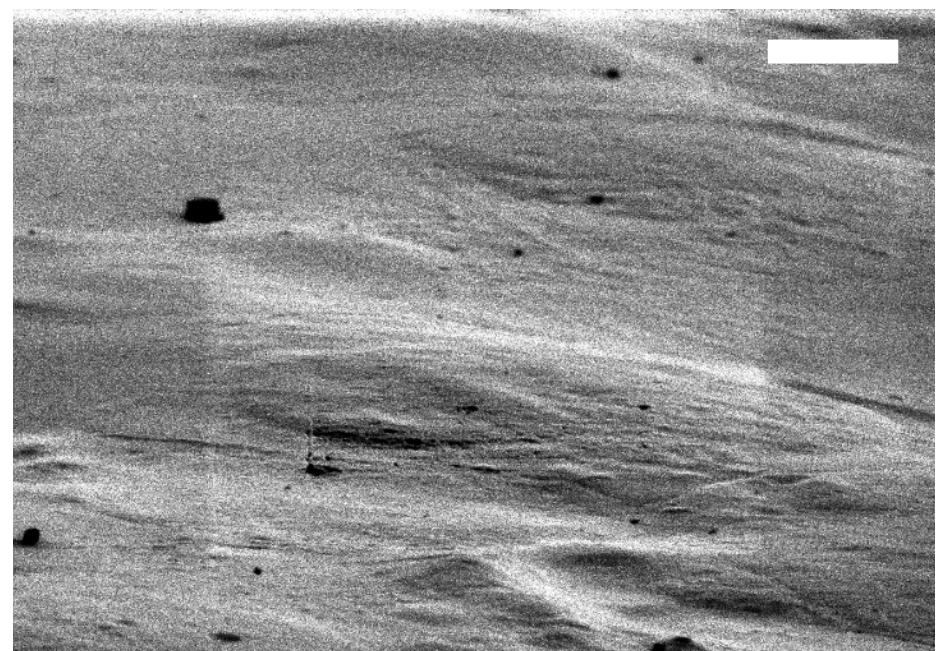


NLRP3 double-ring cage (autoinhibited conformation)

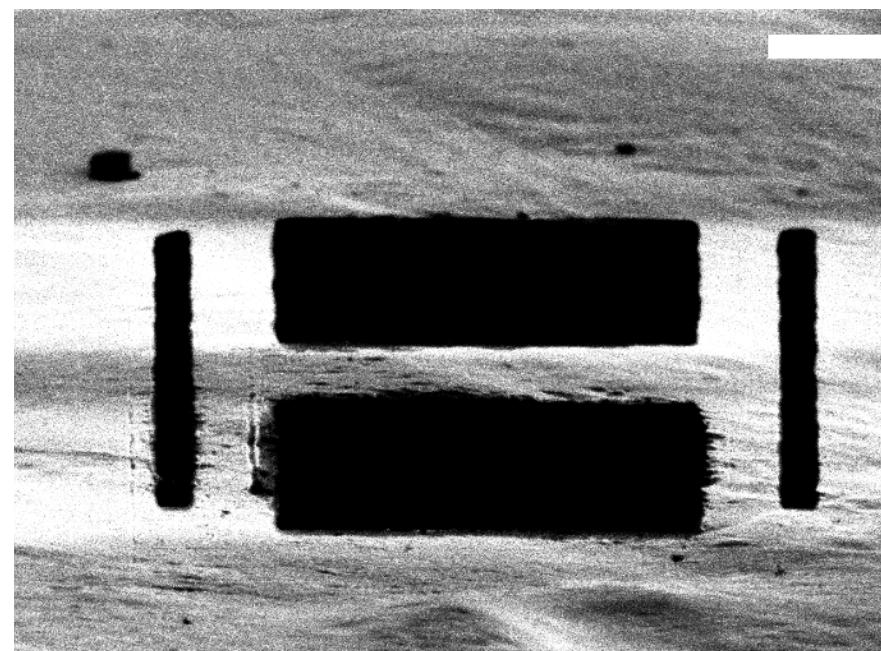


Magupalli, V.G. et al. *Science*. 2020
Lu, A. et al. *Cell*. 2014
Zhang, L. et al. *Science*. 2015
Andreeva, L. et al. *Cell*. 2021
Xiao, L. et al. *Nature*. 2022

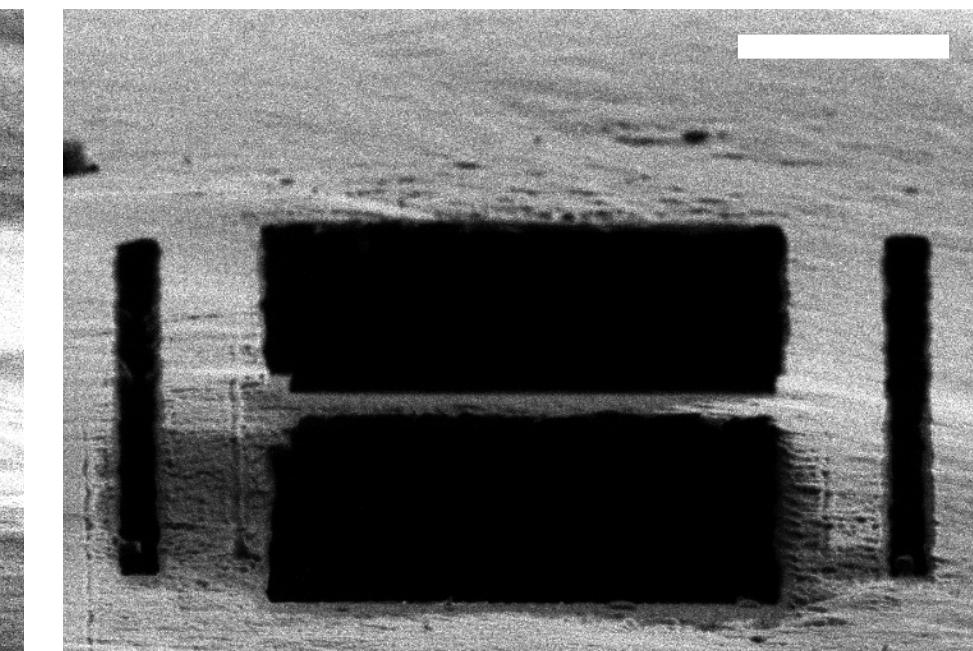
Pre-milling



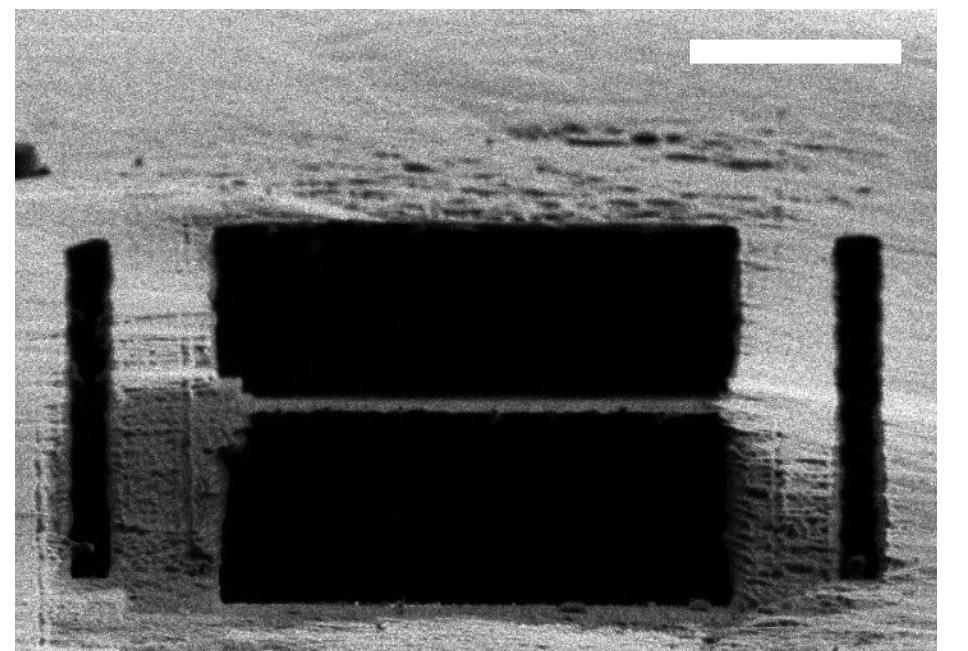
2μm



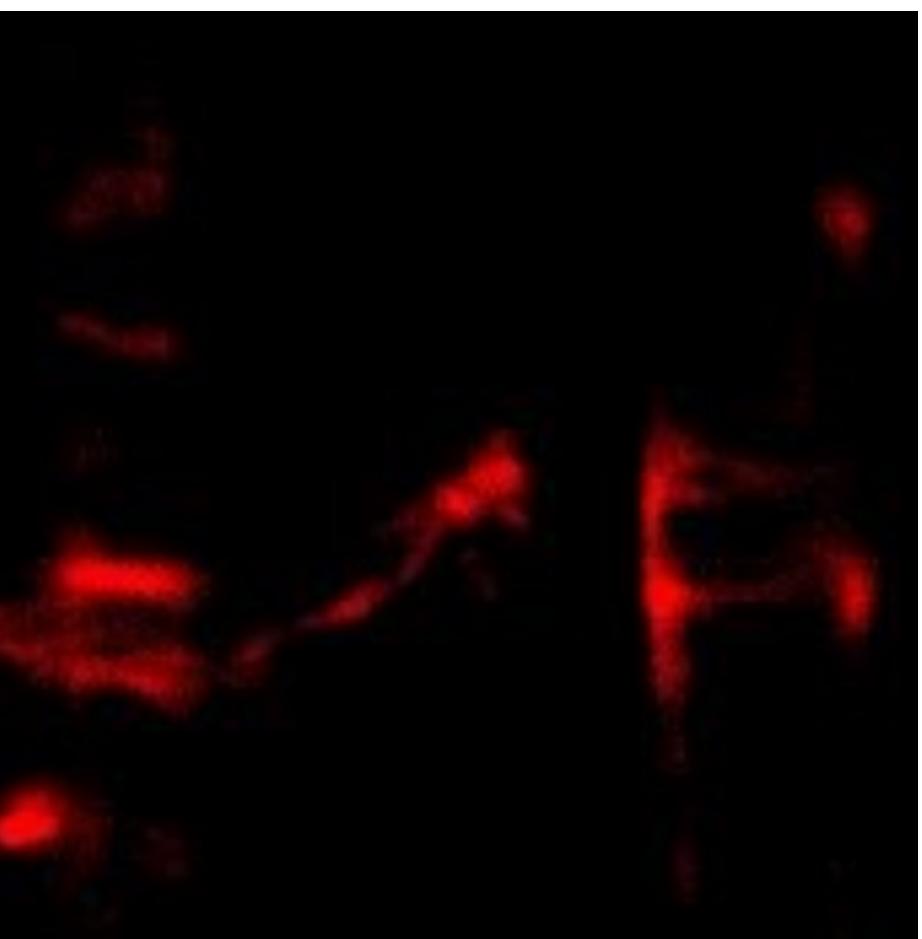
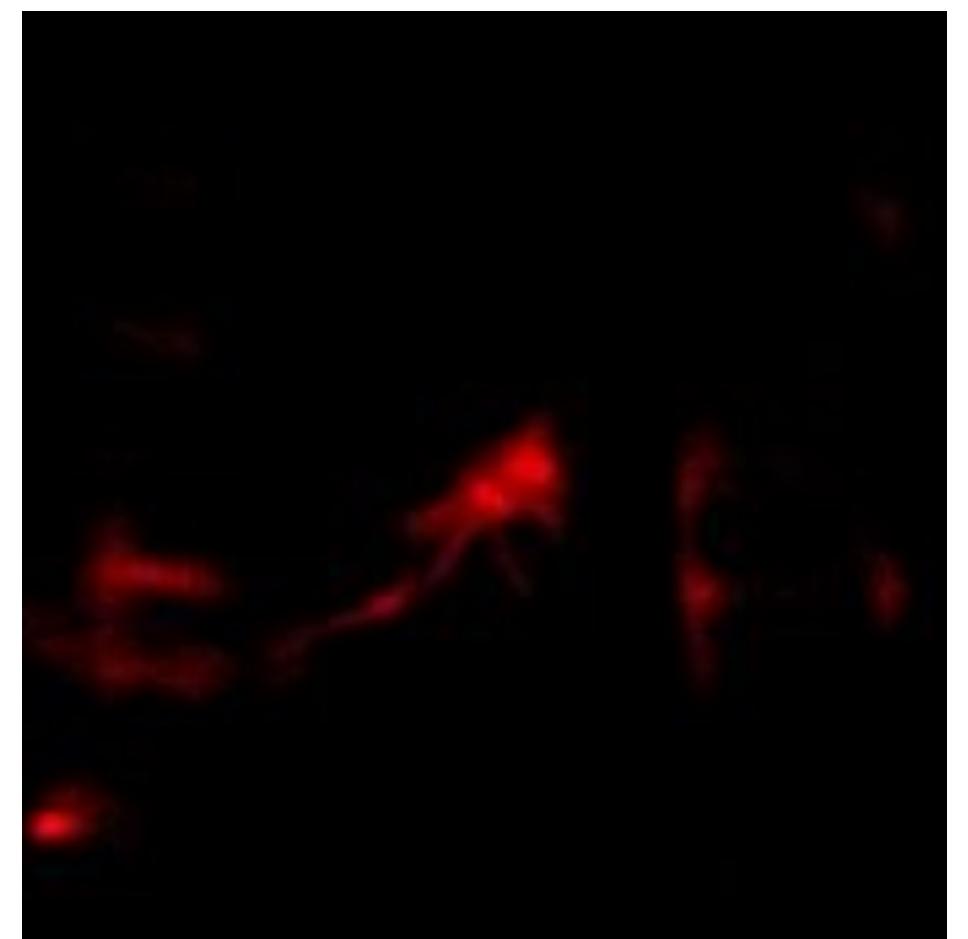
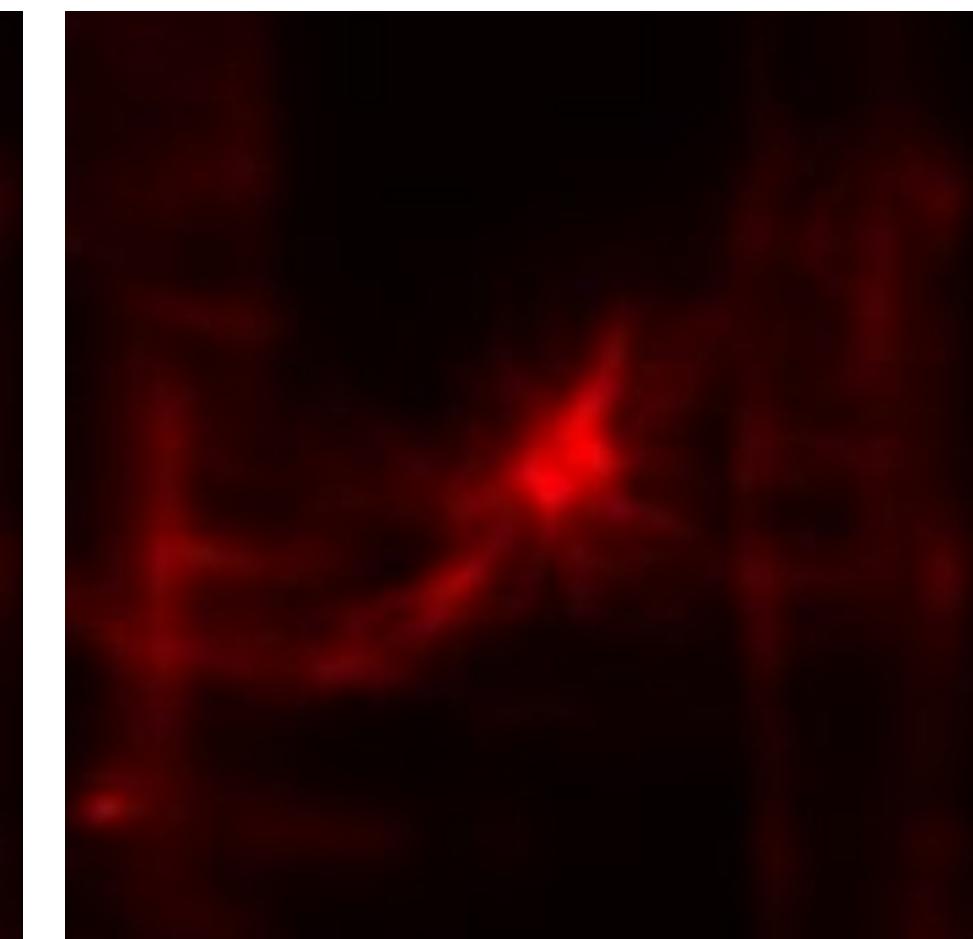
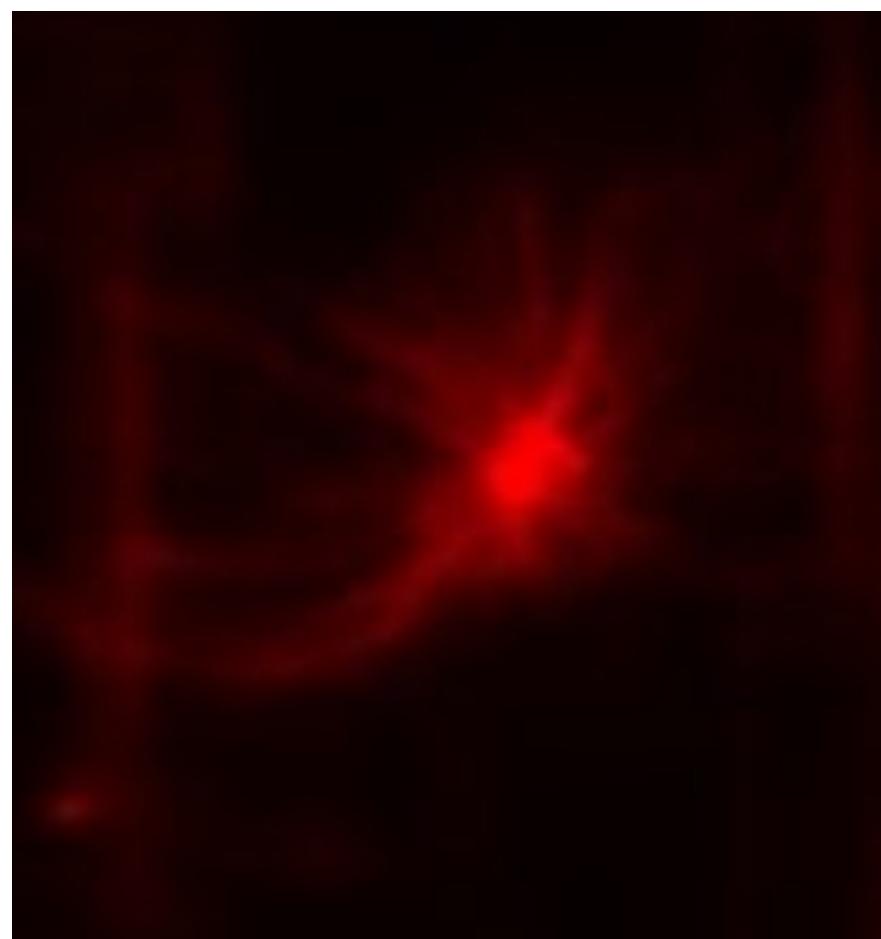
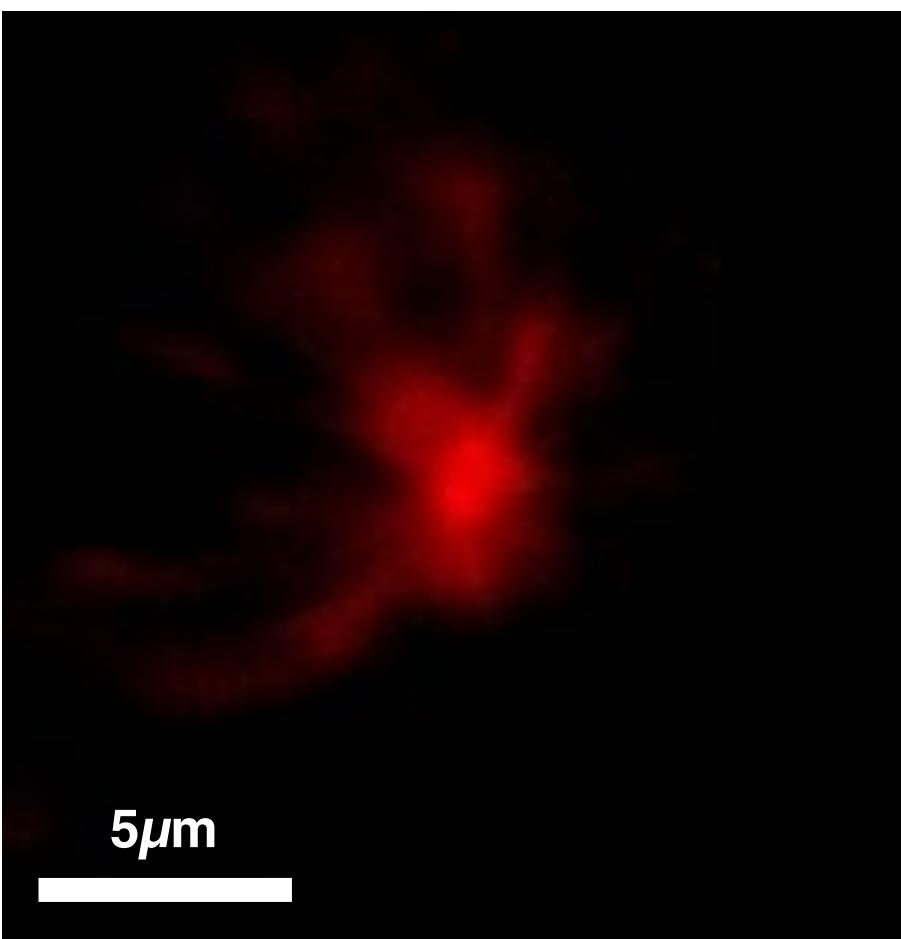
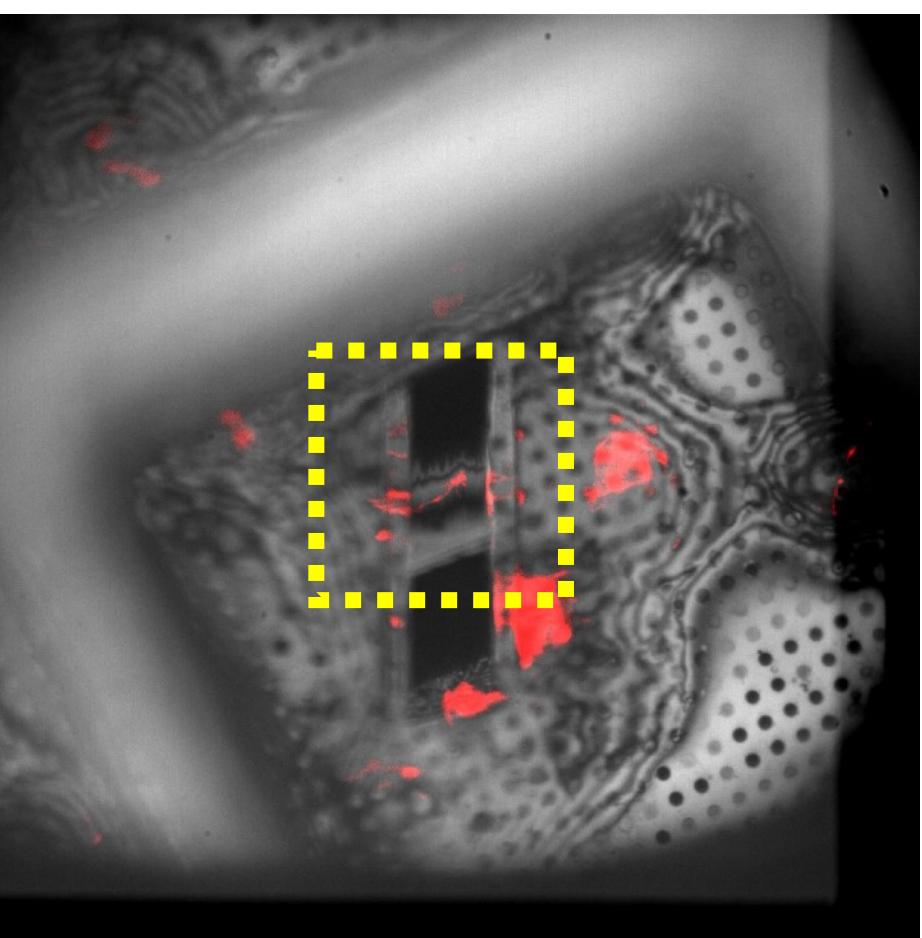
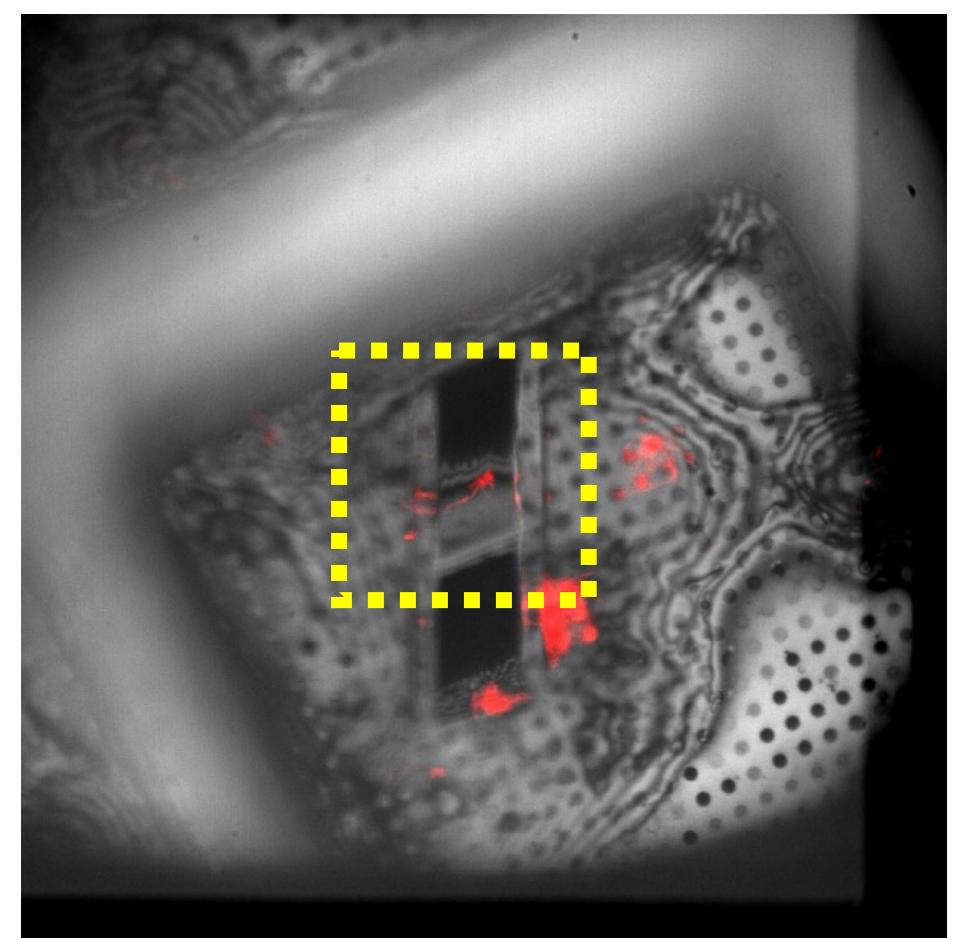
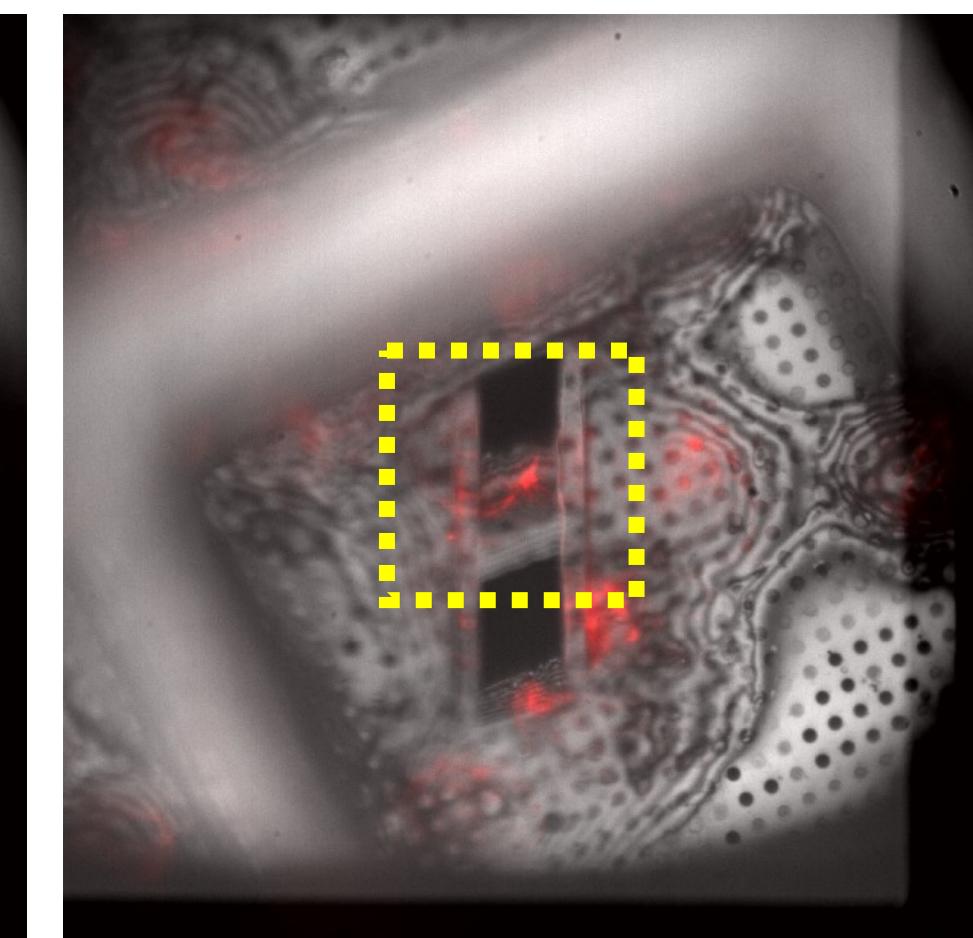
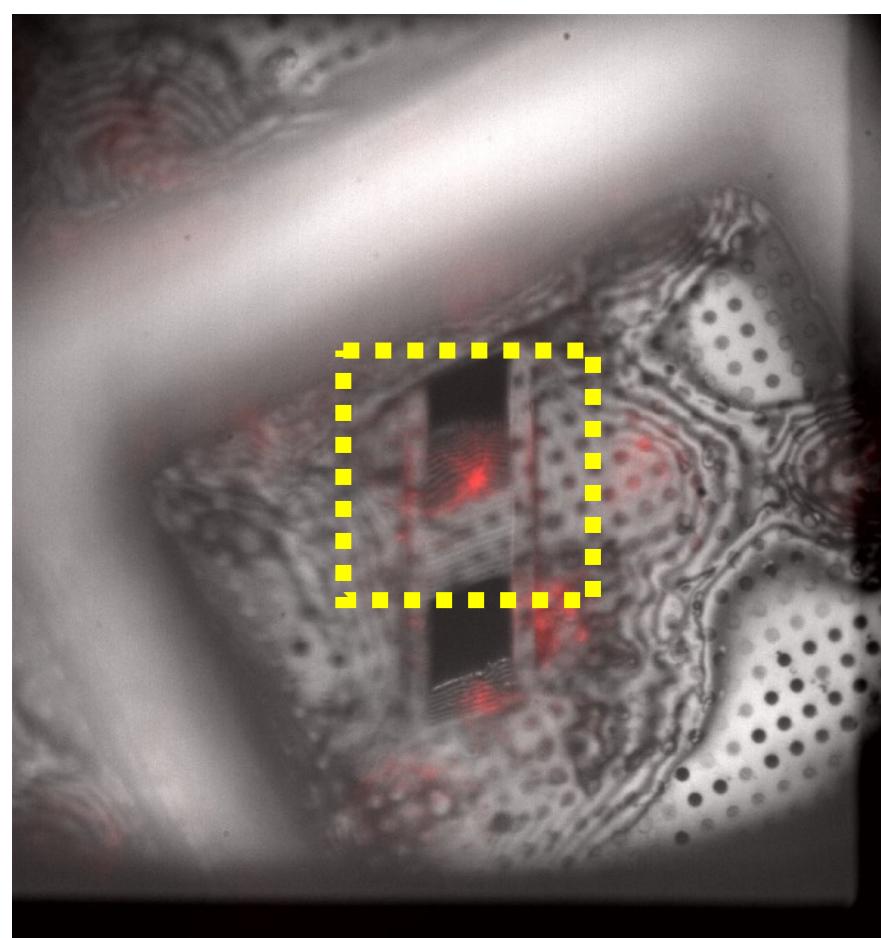
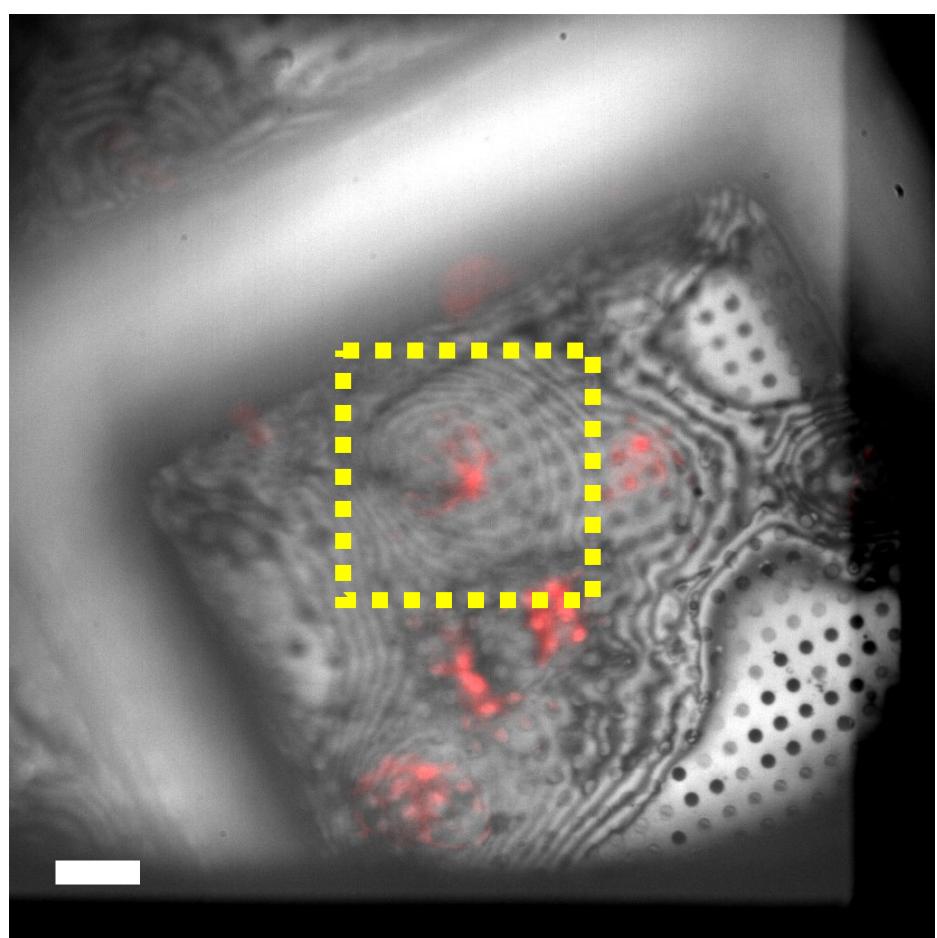
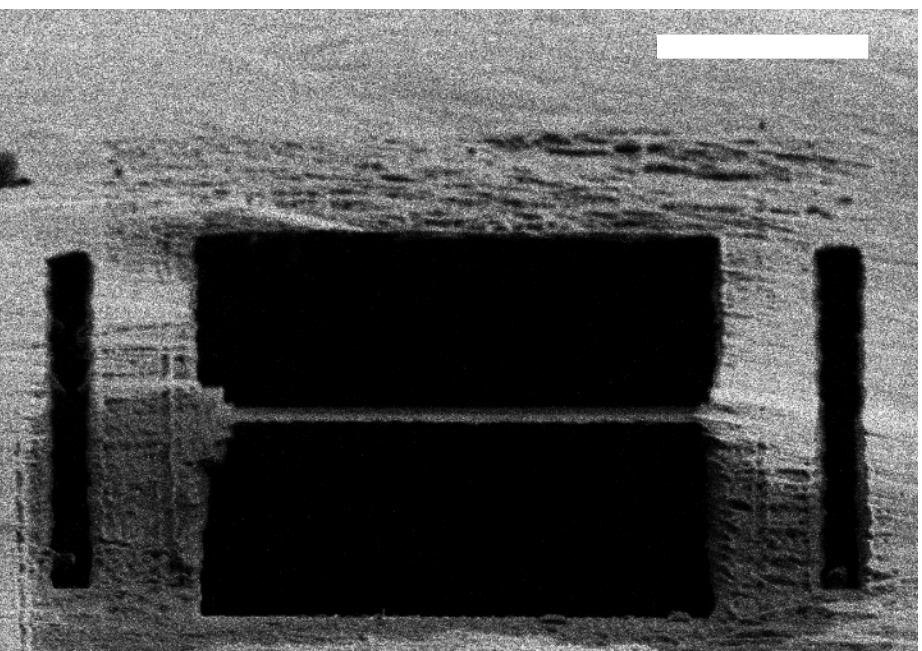
800nm



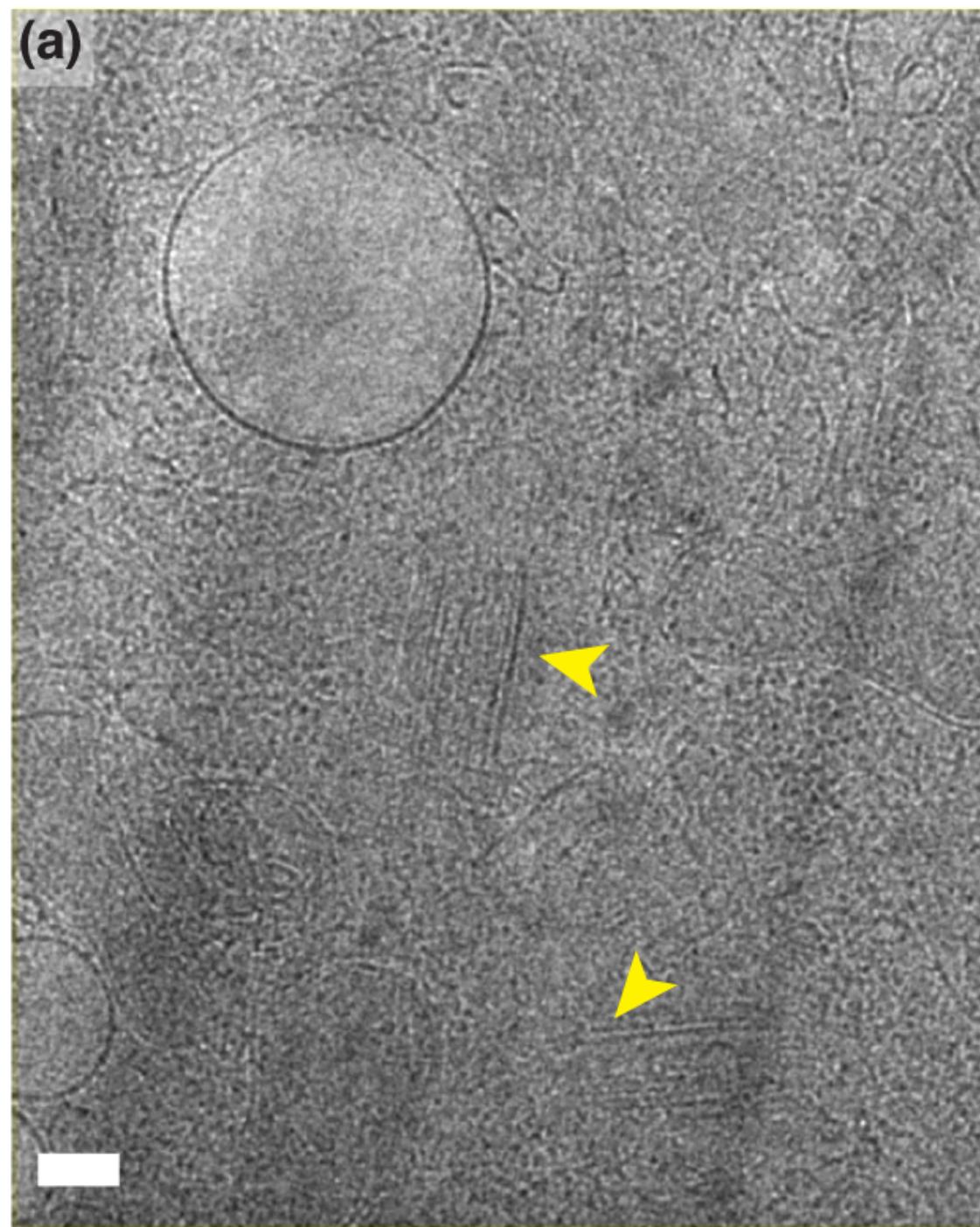
500nm



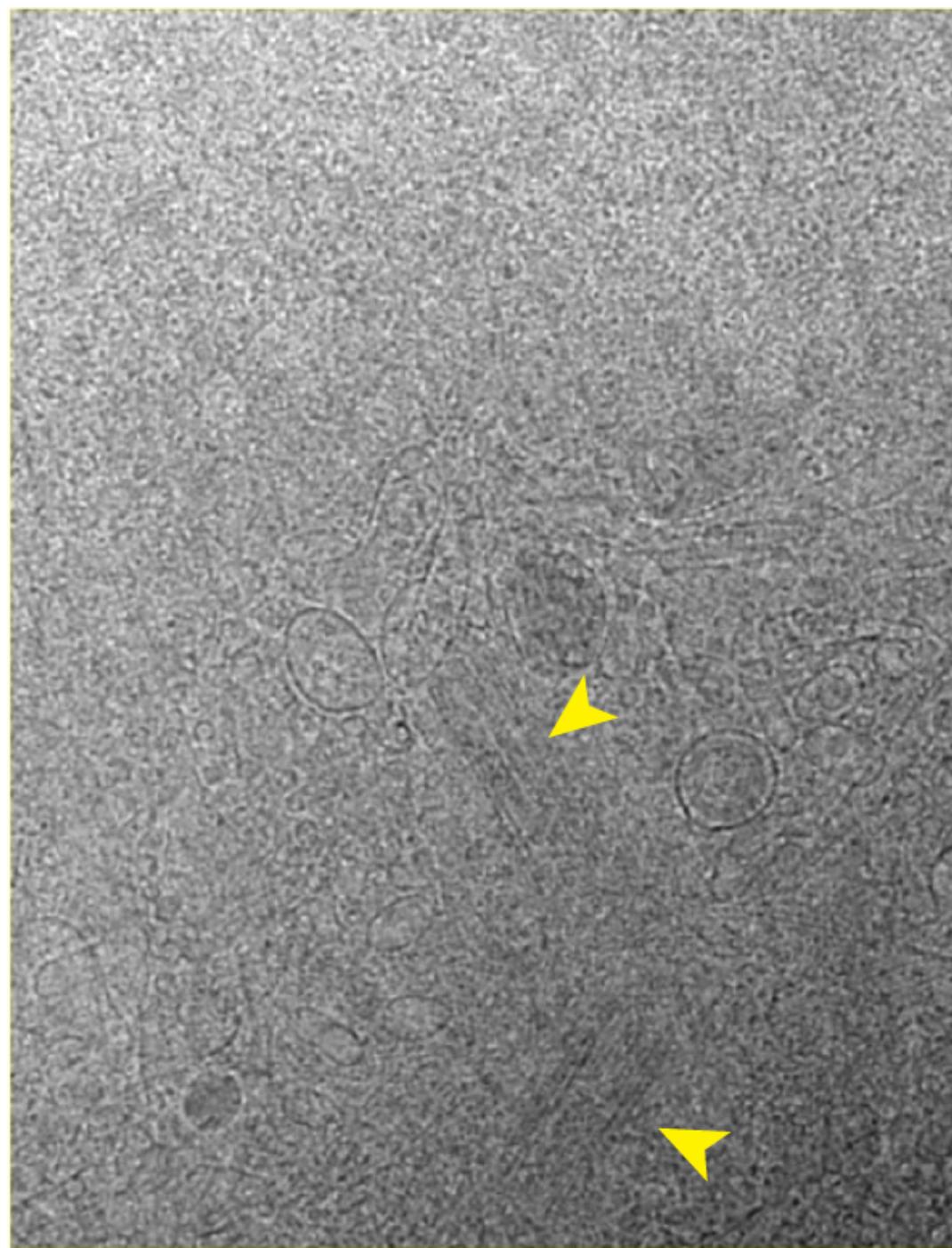
180nm



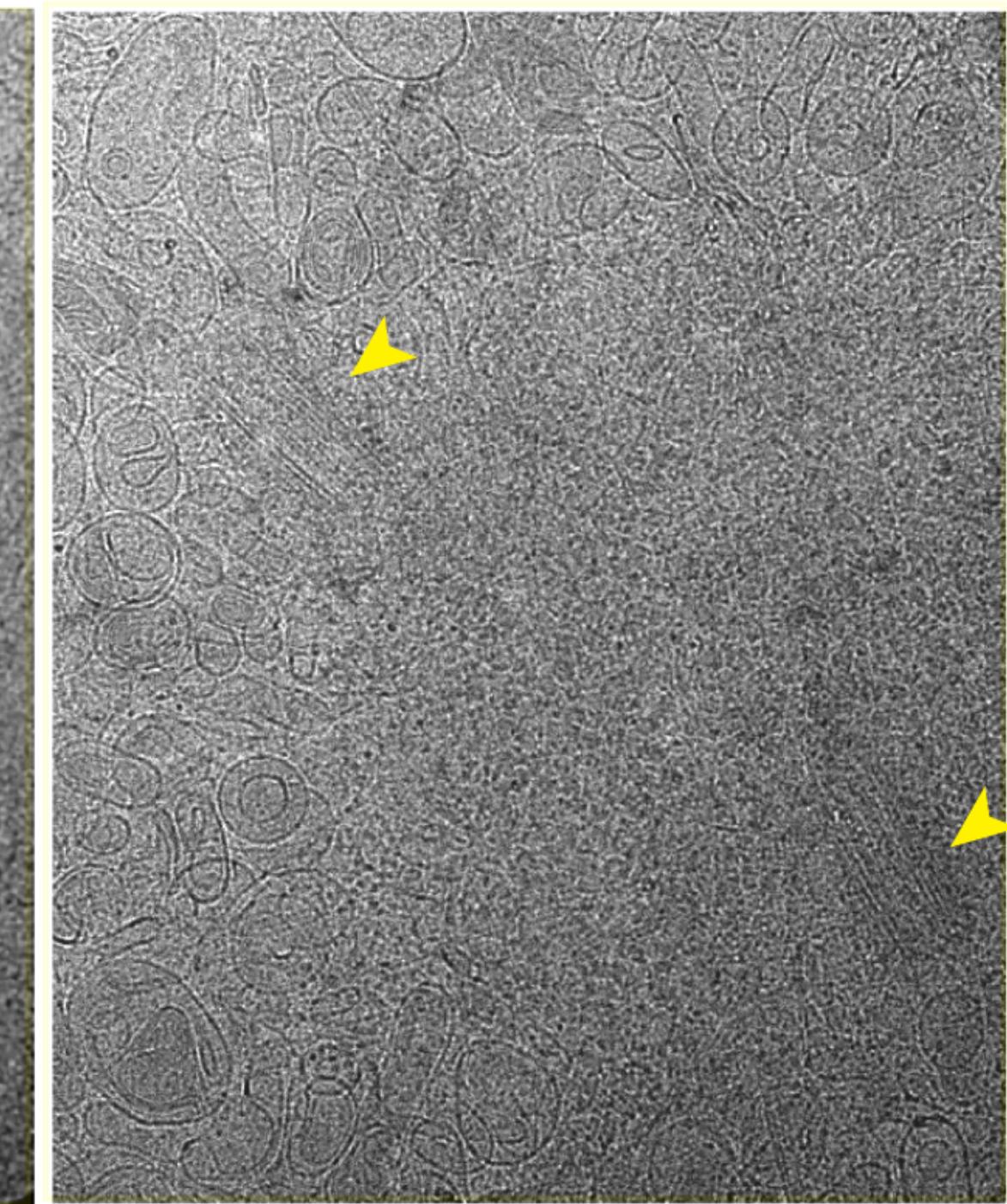
Untreated



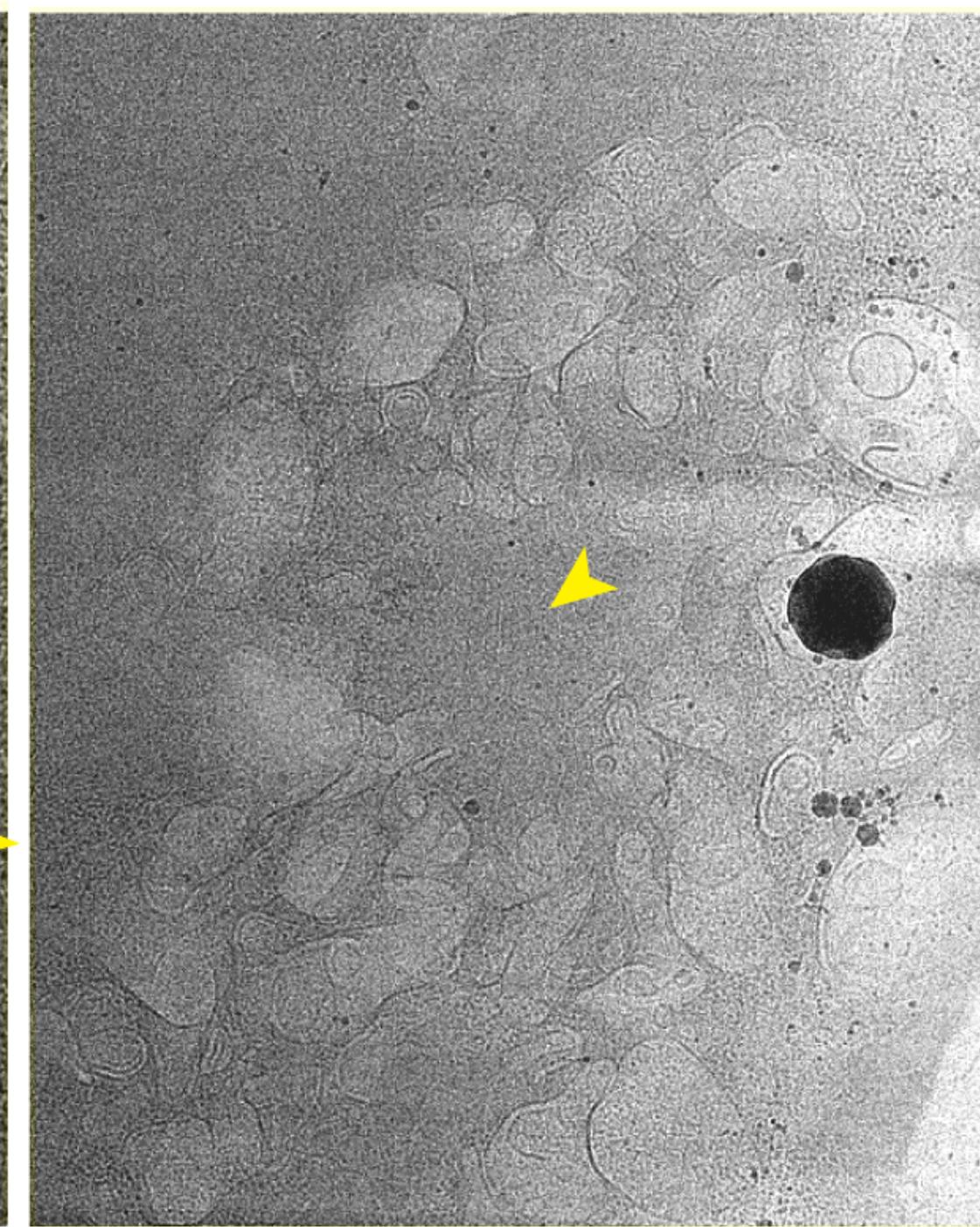
LPS



LPS+Nig 10 min



LPS+Nig > 30min



MTOC

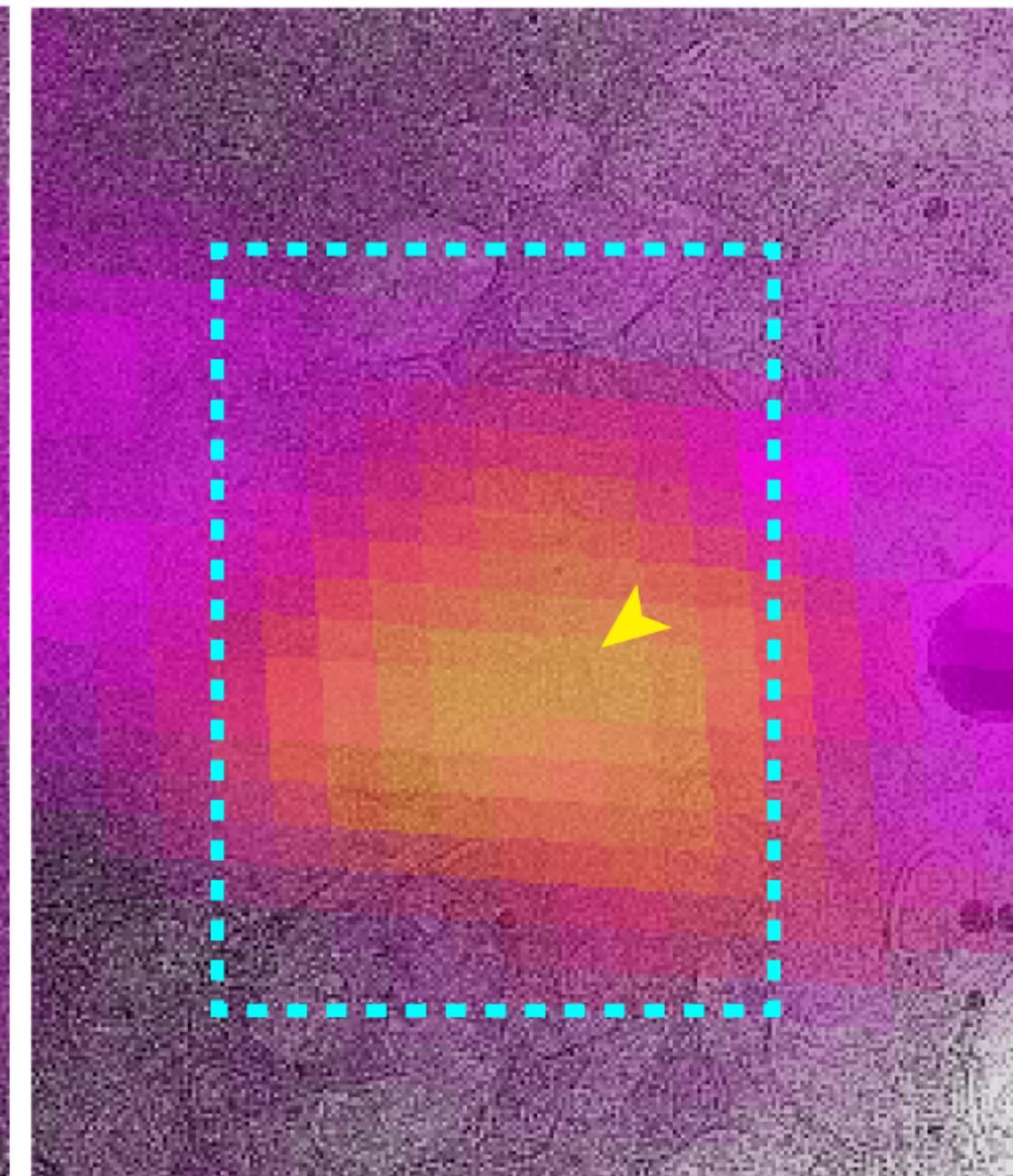
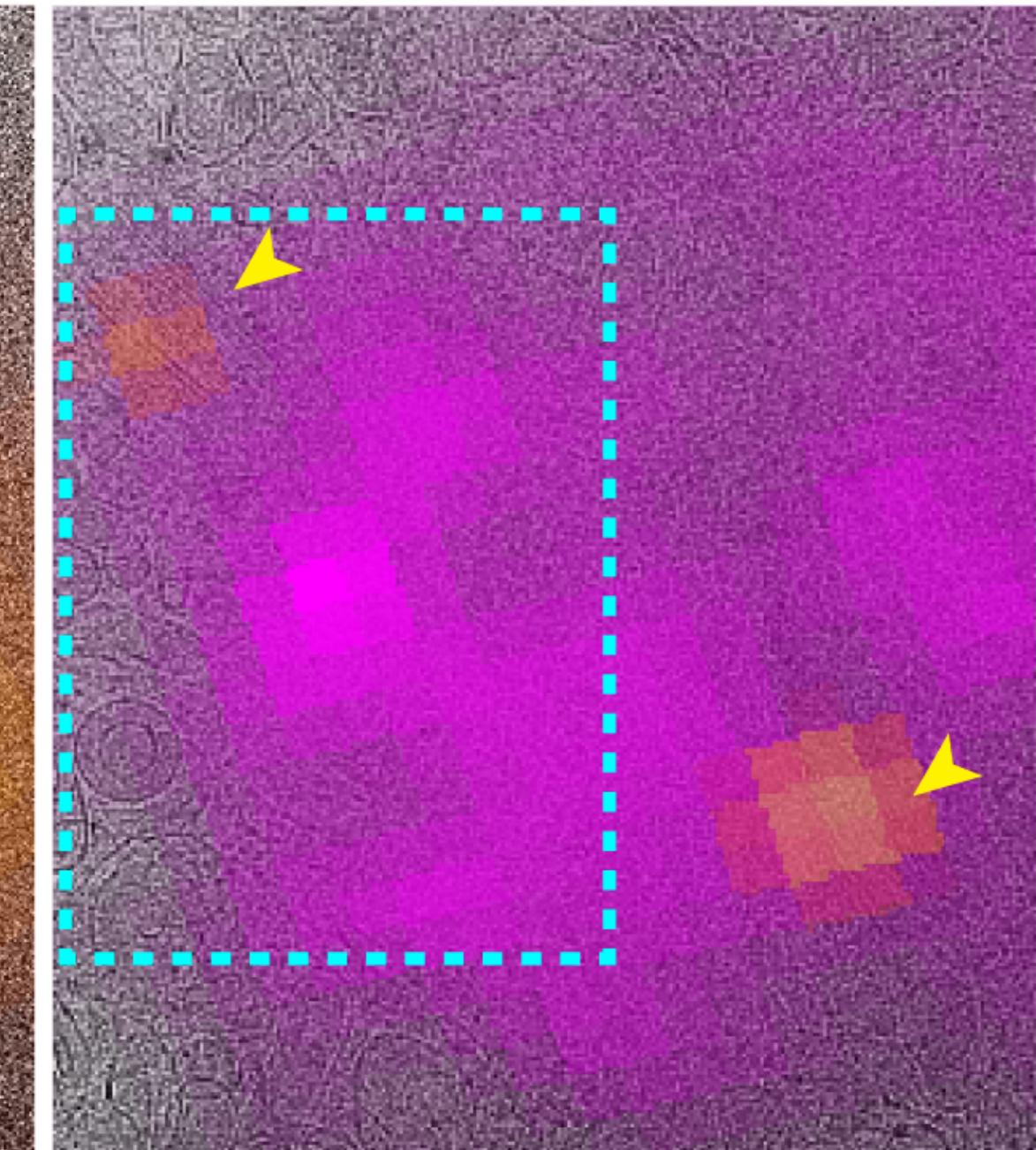
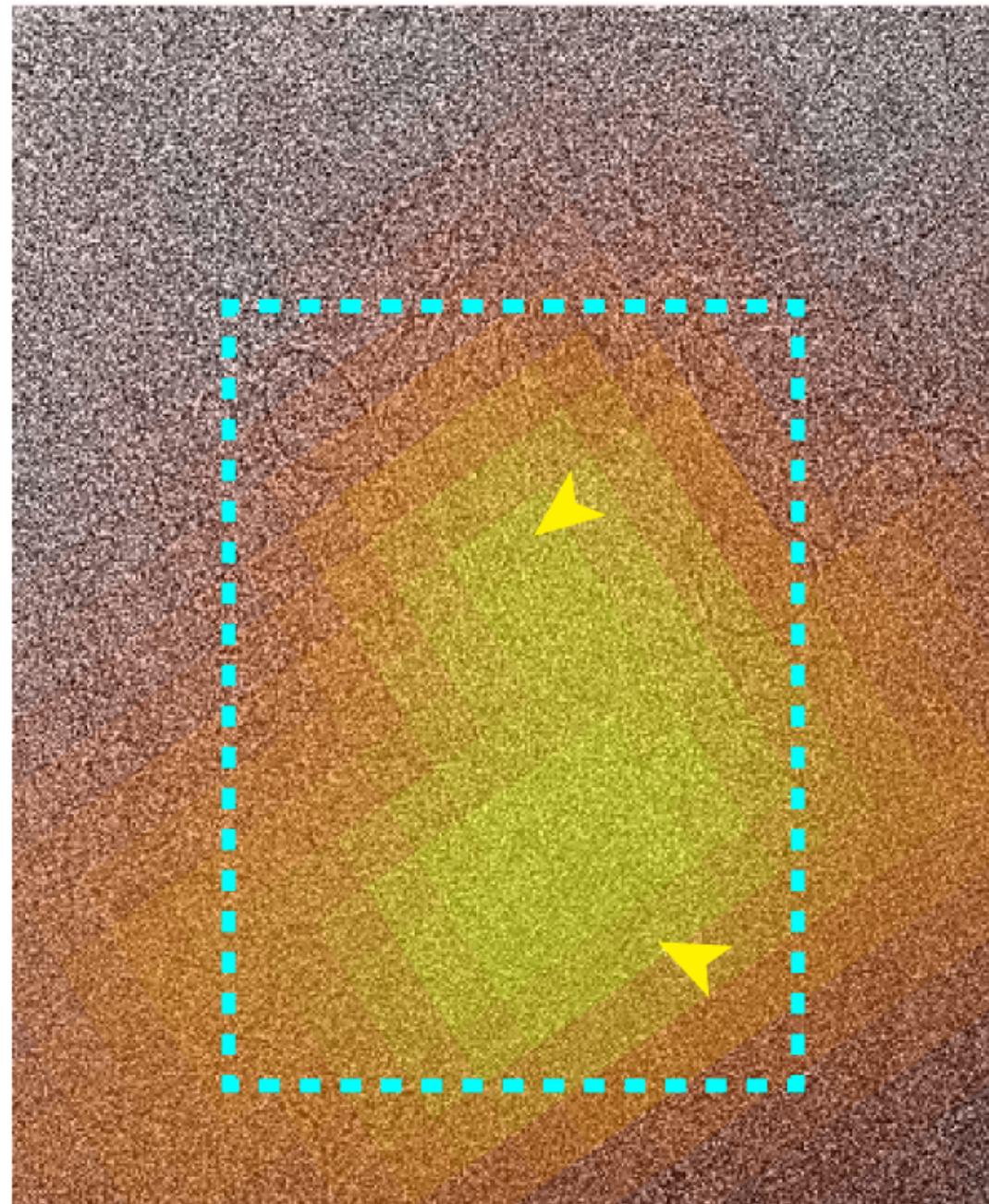
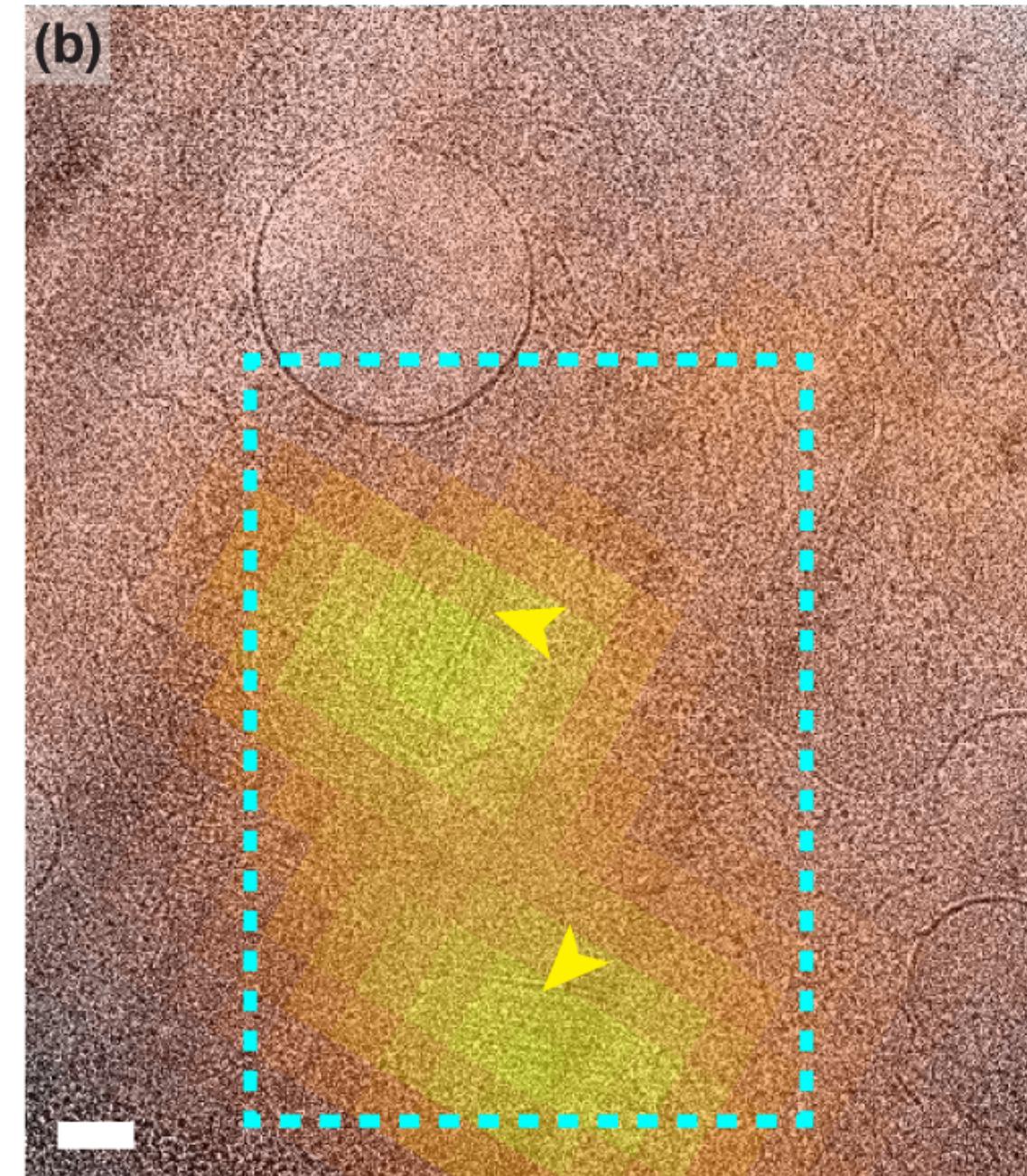
Low

High

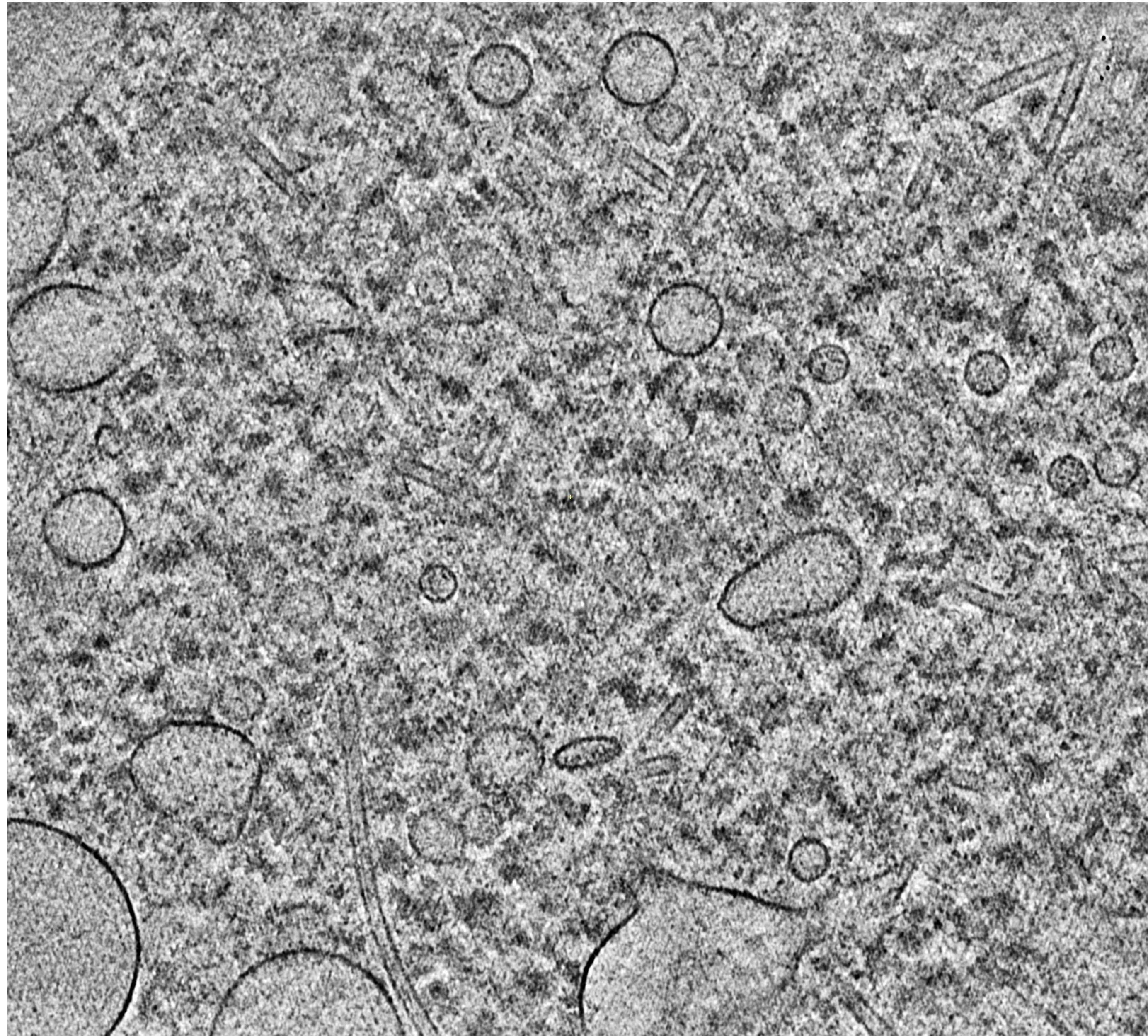
NLRP3

Low

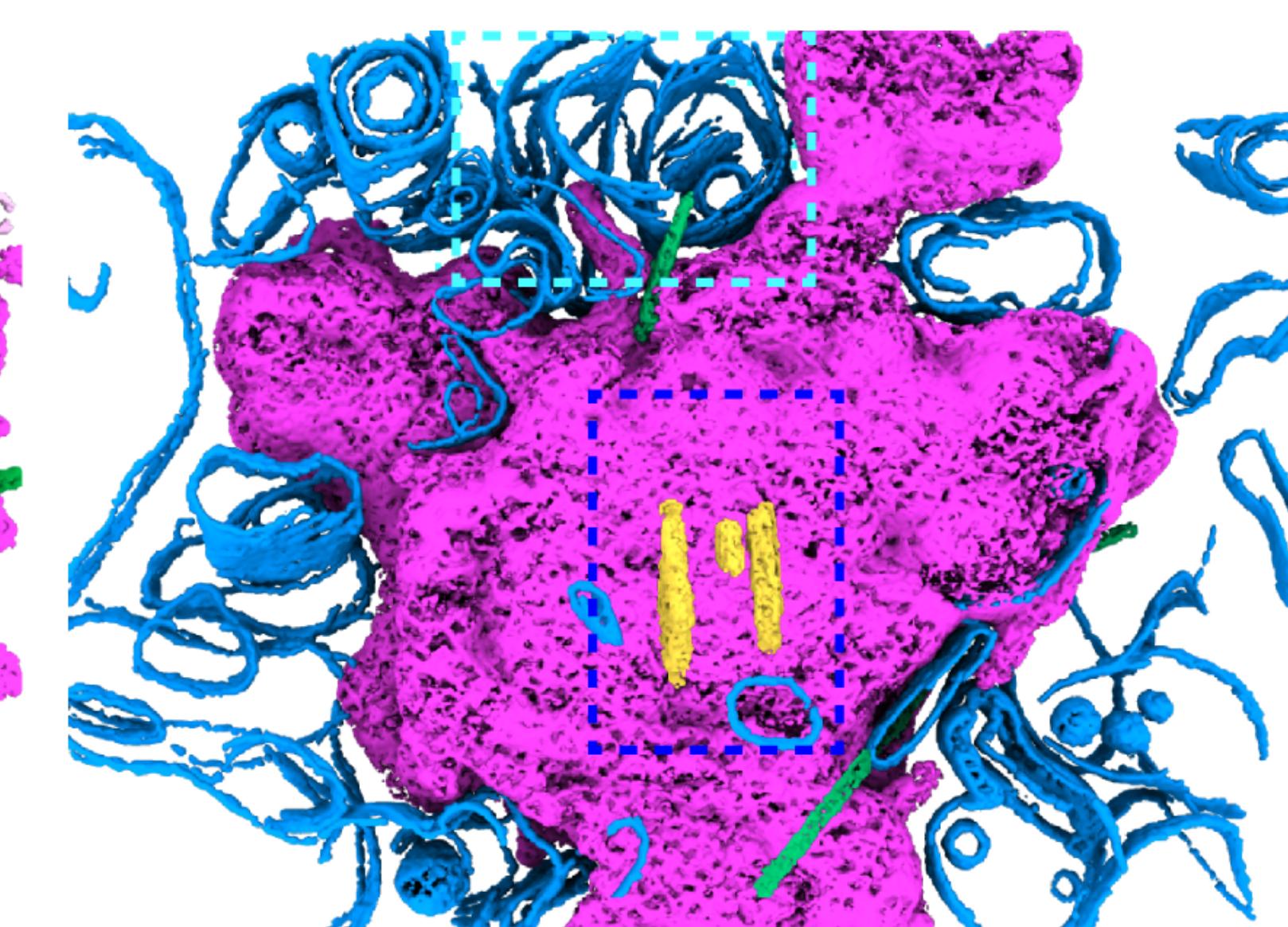
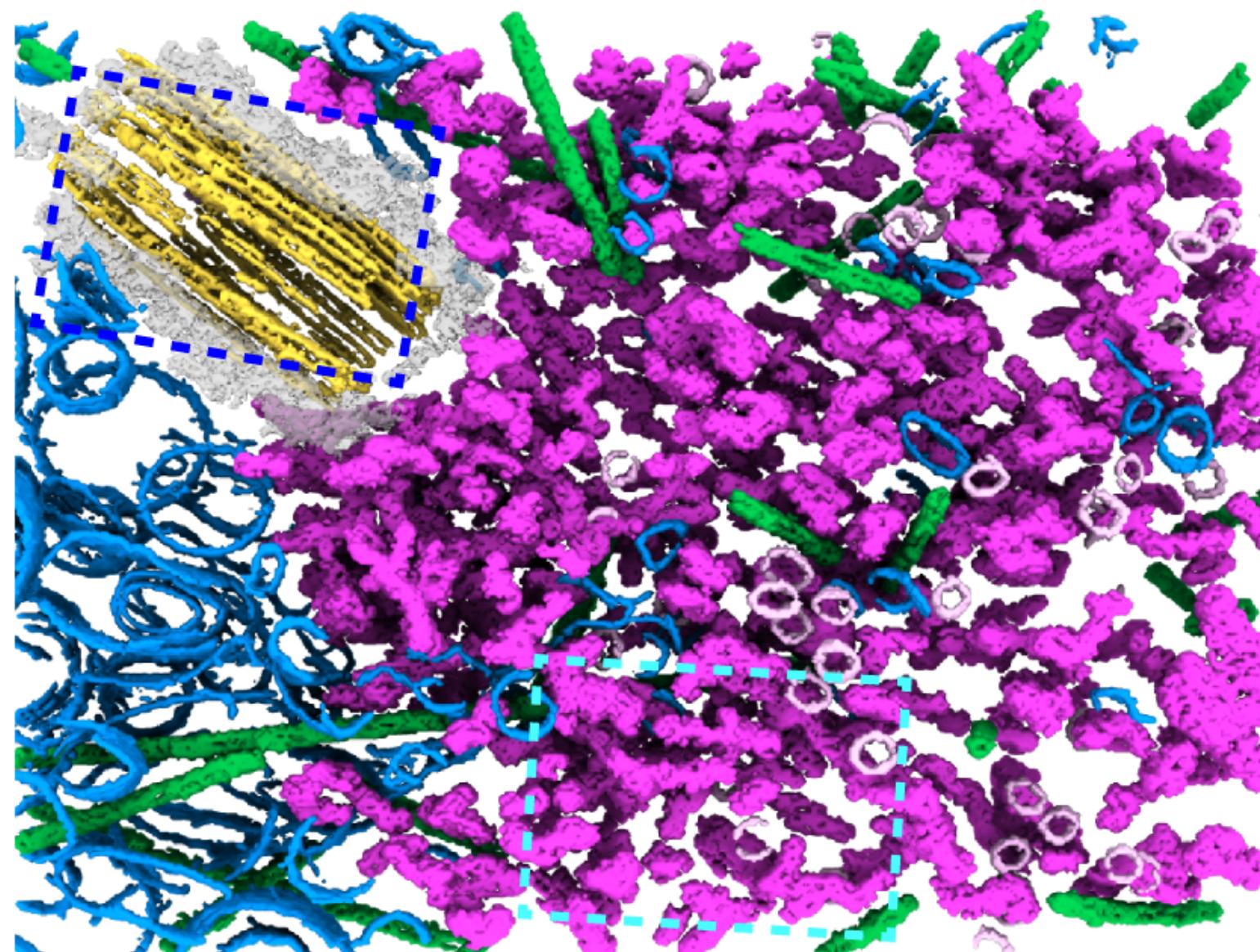
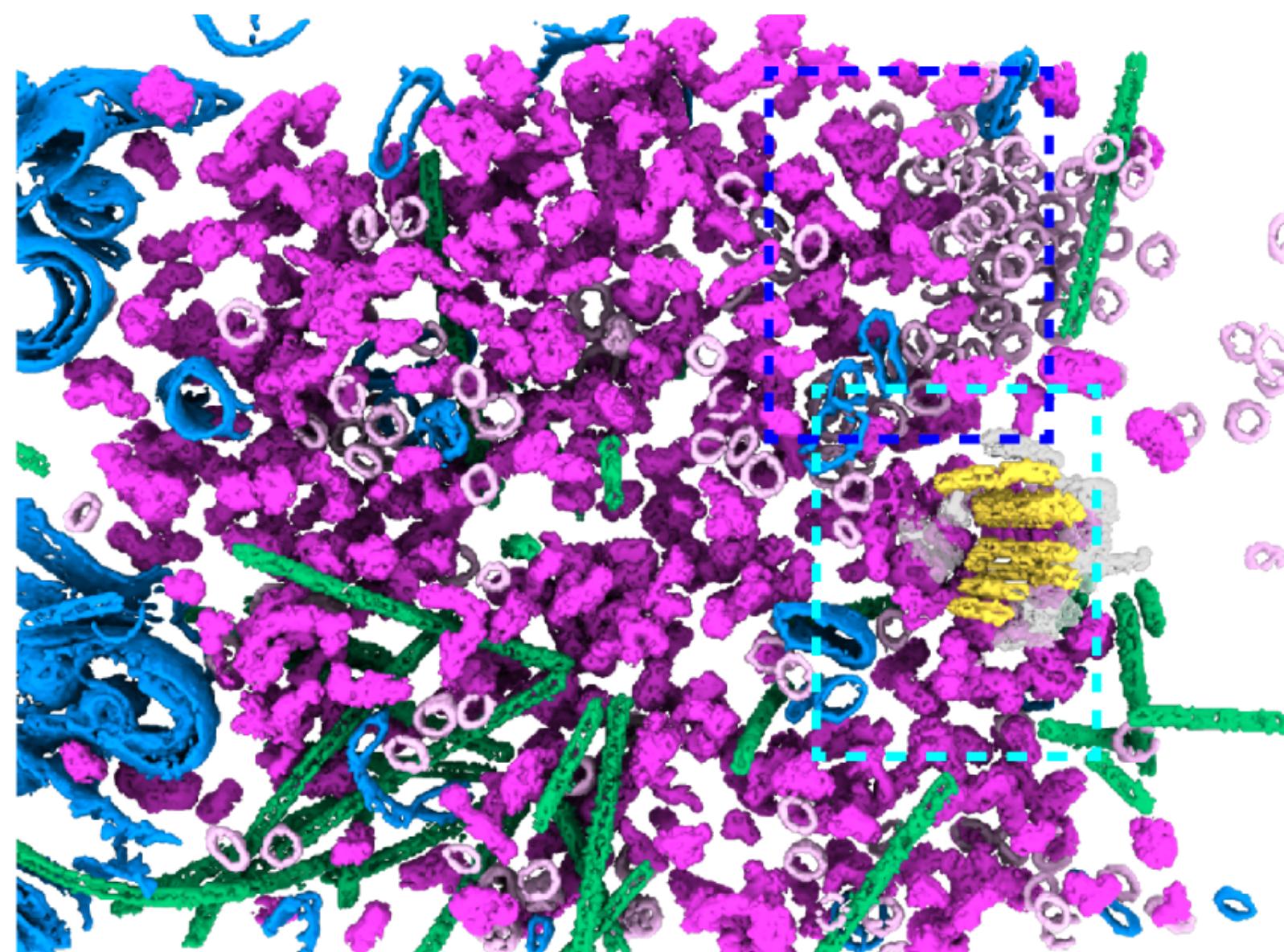
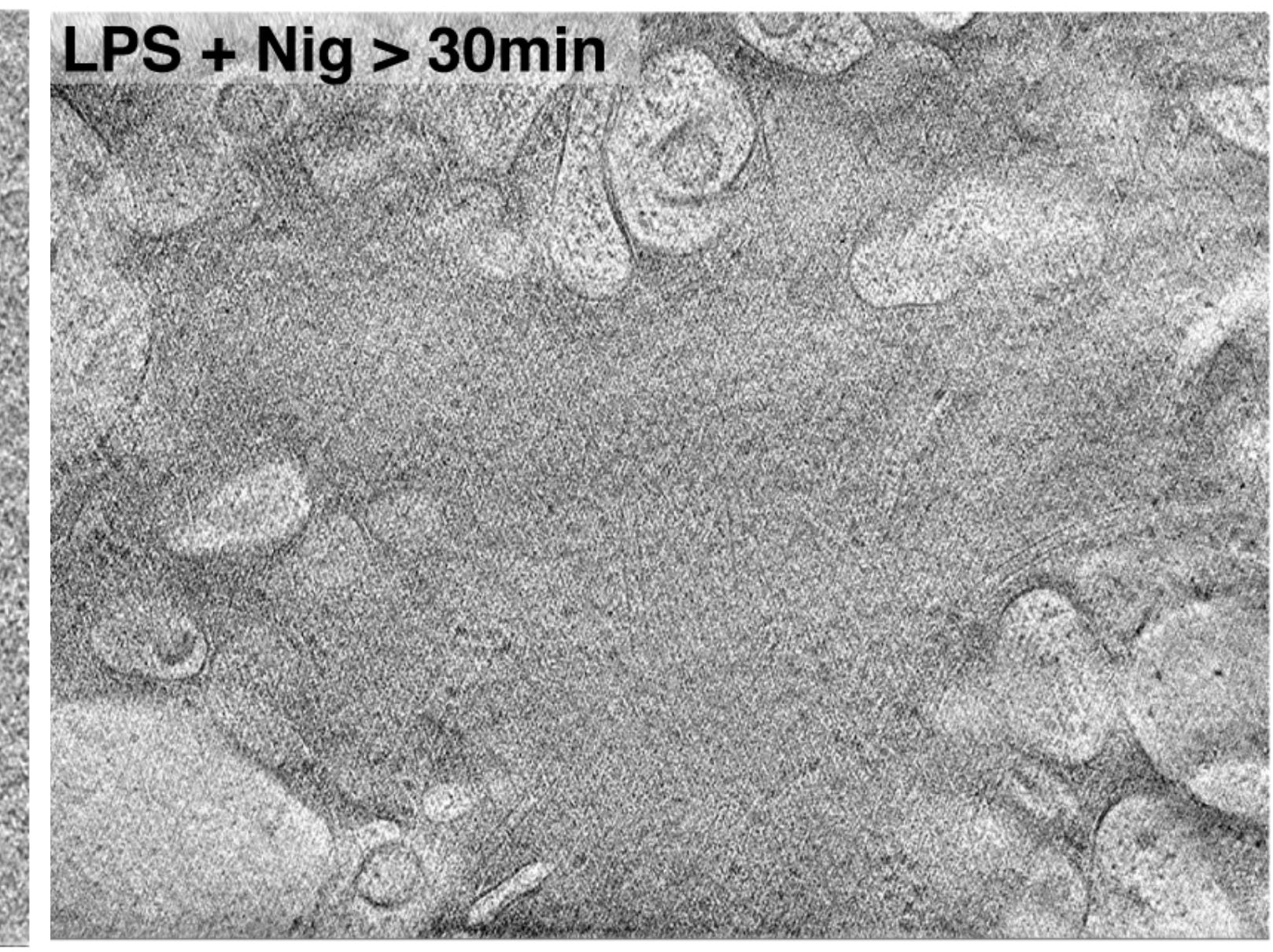
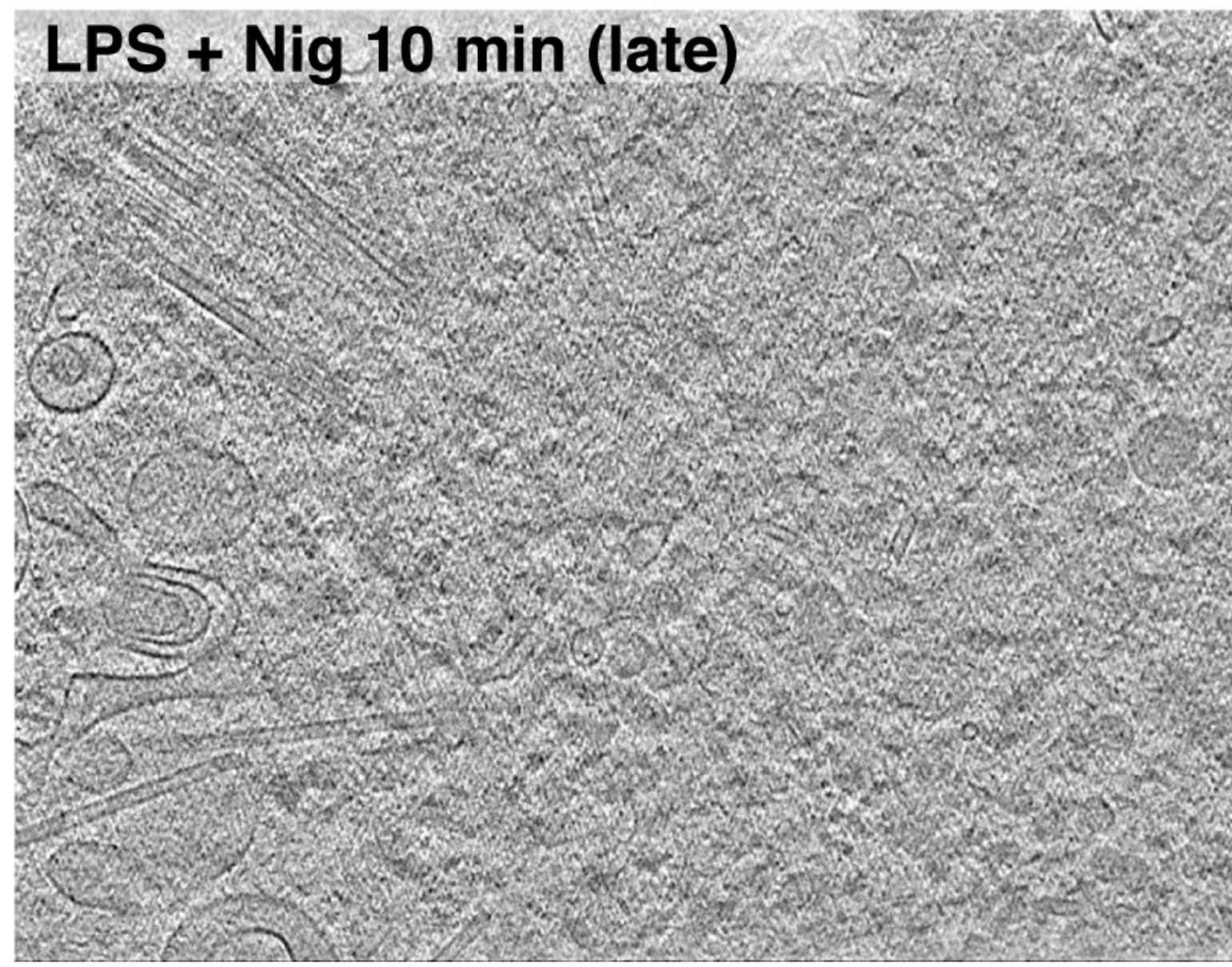
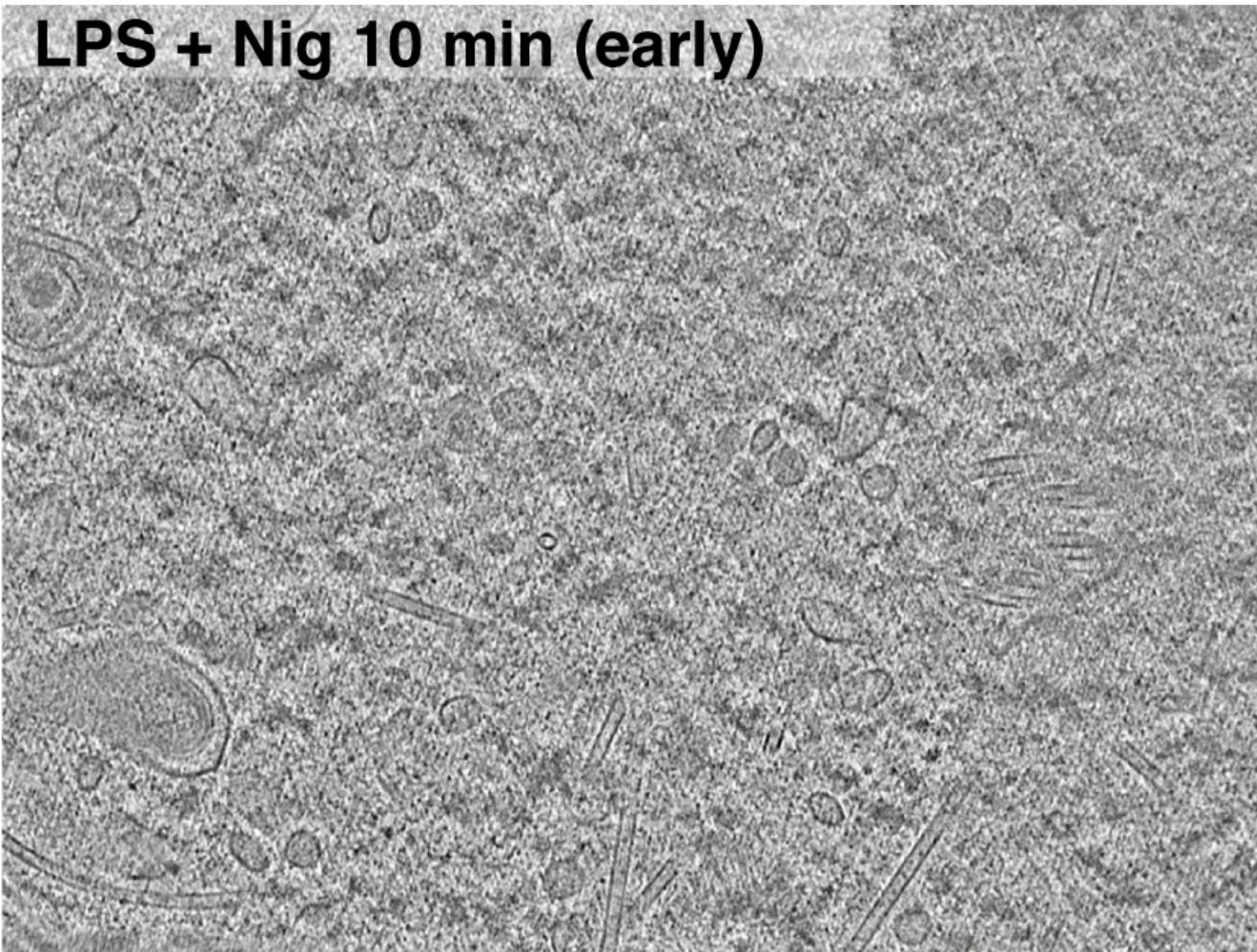
High

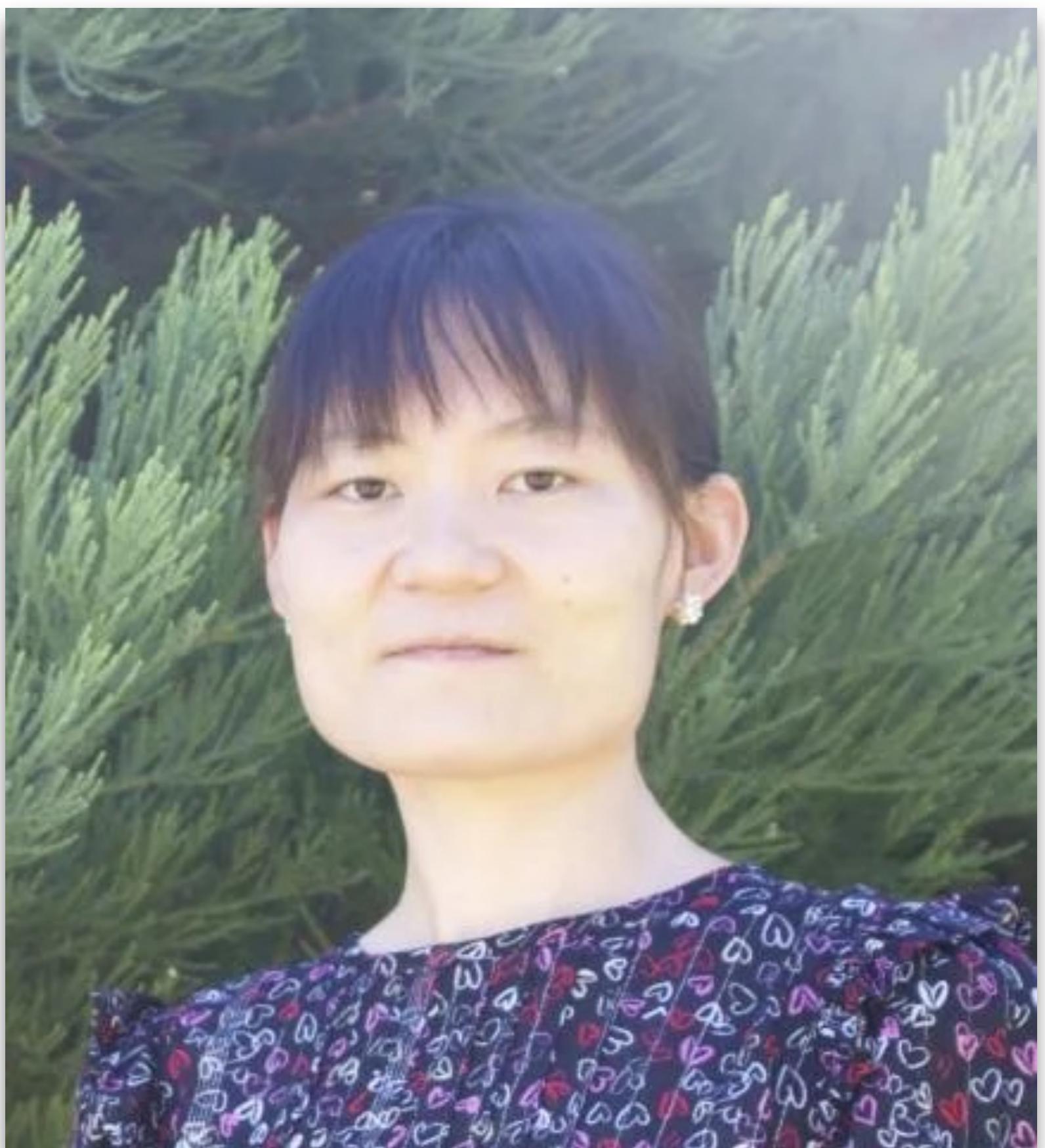


Activated human NLRP3 inflammasomes are unstructured condensates



The NLRP3 condensate grows denser





Phyllis Wang
Caltech/Stanford



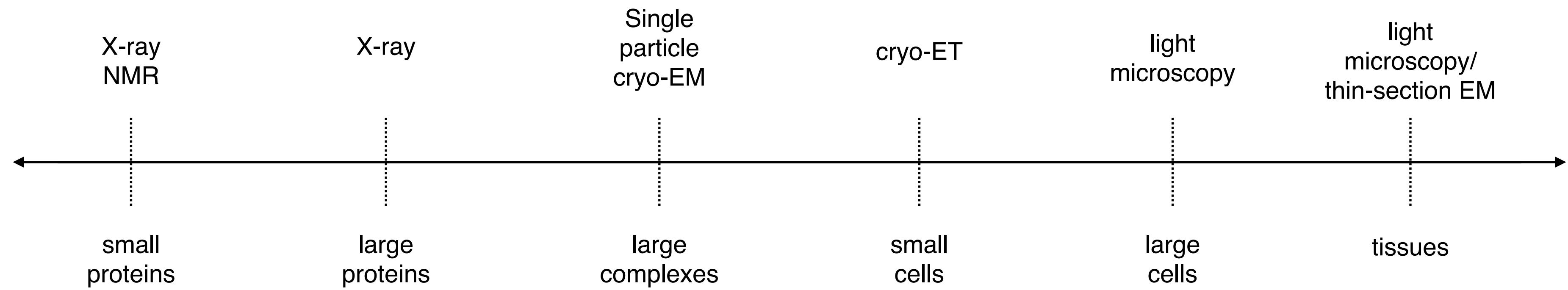
Hao Wu
Harvard

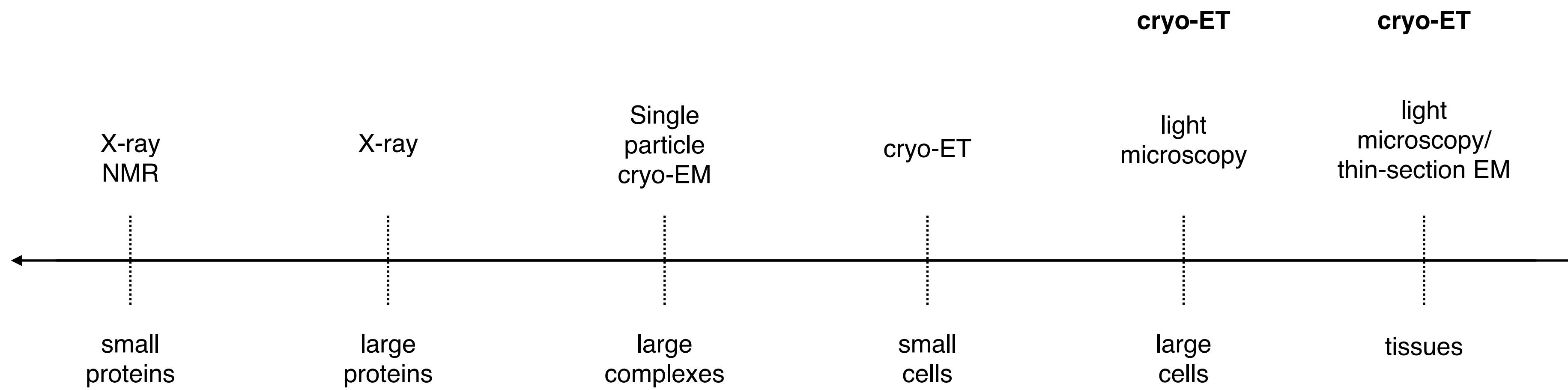


Peter Dahlberg
Stanford

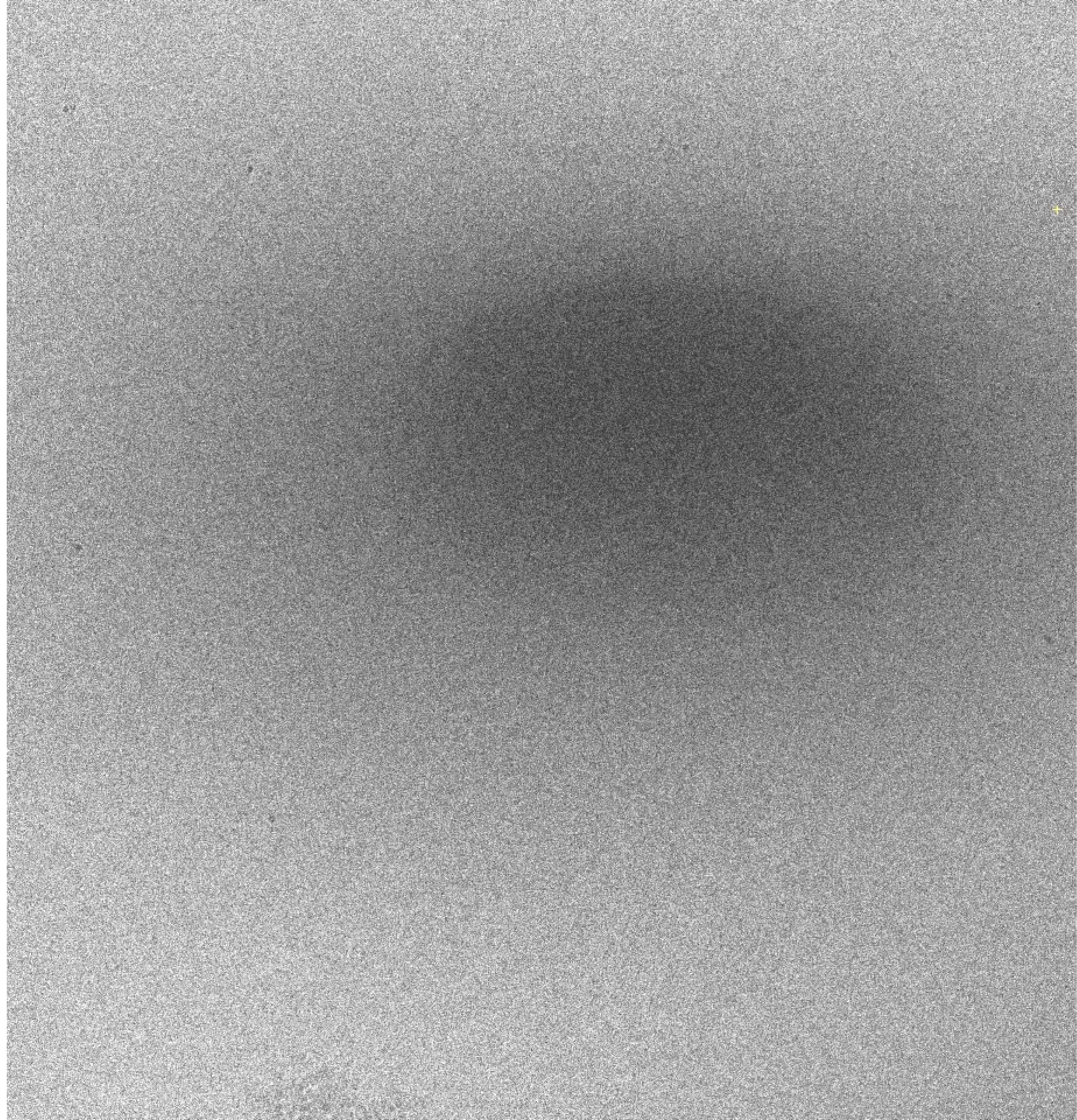
Future technical advances

- Direct detectors: quantum efficiency/MTF, size, speed
- Phase plates
- Lower temperatures with liquid helium-cooling
- Better images at high tilt - what is the problem?
- Larger areas: montage imaging
- Faster - continuous tilting?
- Automation (cryo-FIB milling, data collection, image processing)
- Super-resolution CLEM
- Routine serial-section lift-out

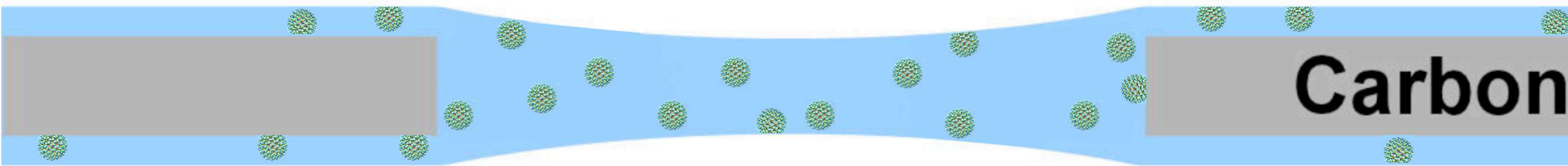




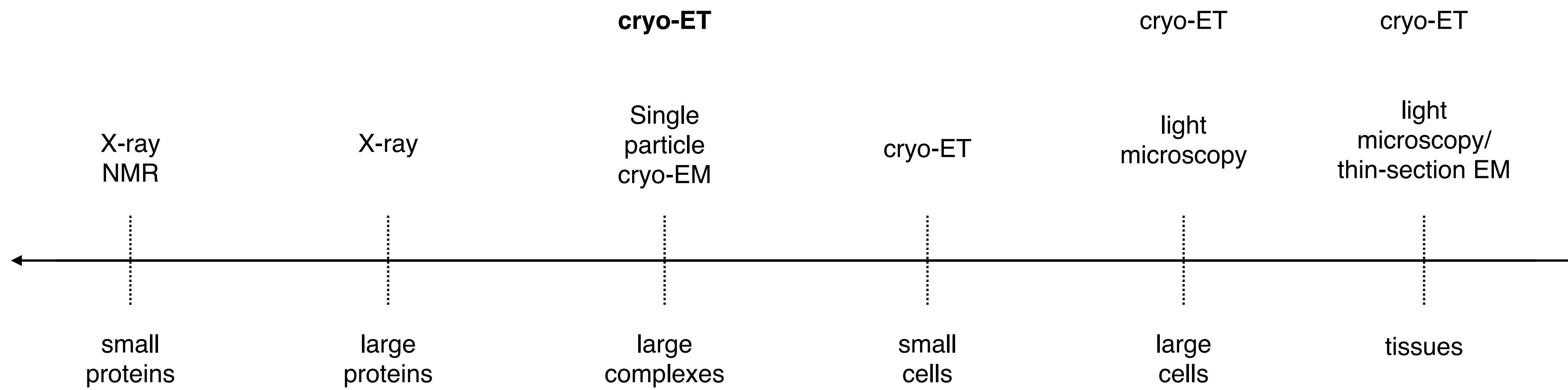
1. Fluorescence-guided FIB-milling



How fast tilt-series could supplement single particle projections

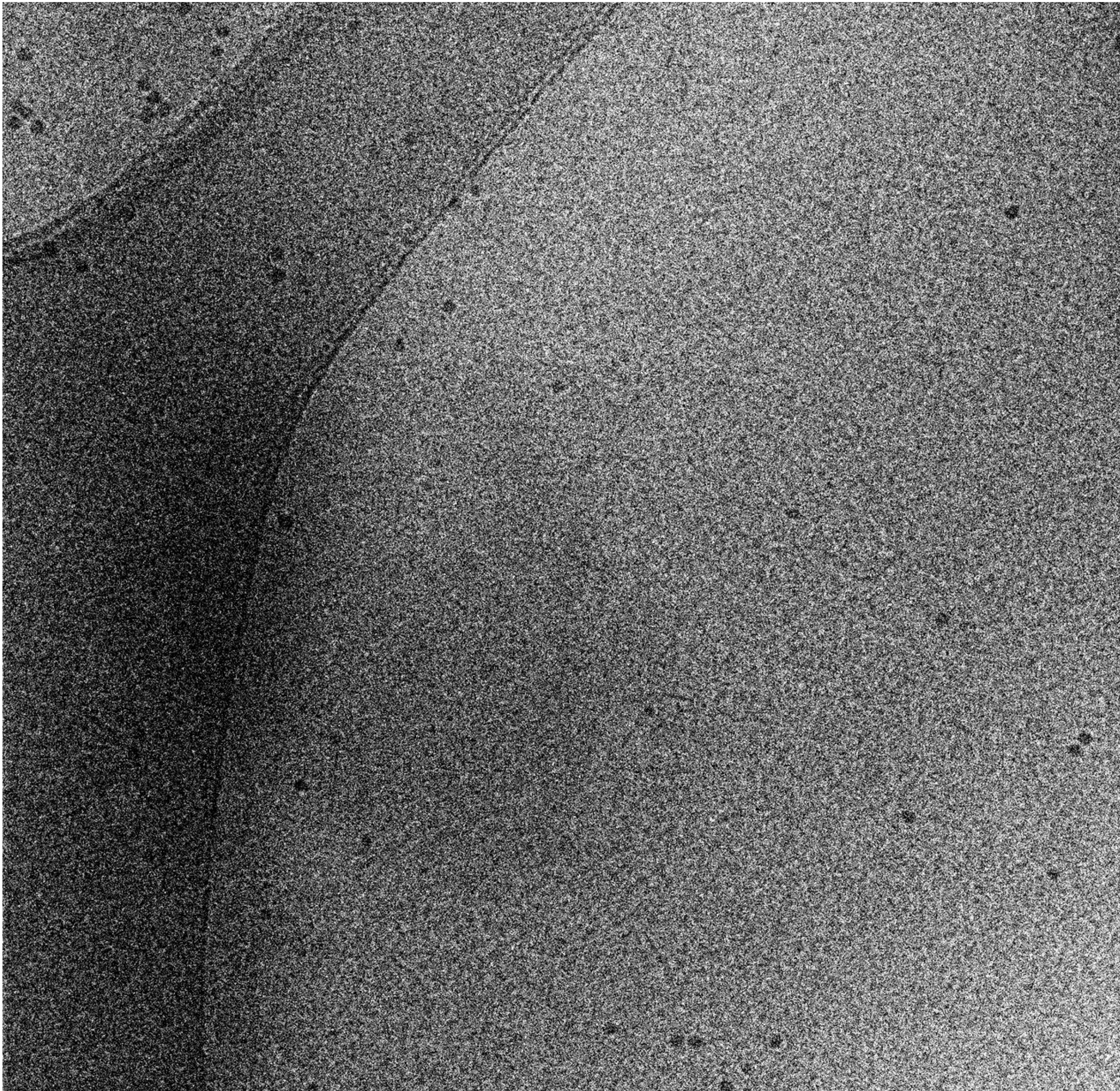


- detect and exclude particles on the air/water interface (which are likely damaged)
- improve per-particle CTF-refinement by determining each particle's z-coordinate within the ice
- disambiguate conformational changes from differences in orientation
- start alignment searches very near the true 3-D orientation



1. Fluorescence-guided FIB-milling
2. Rapid tilt-series

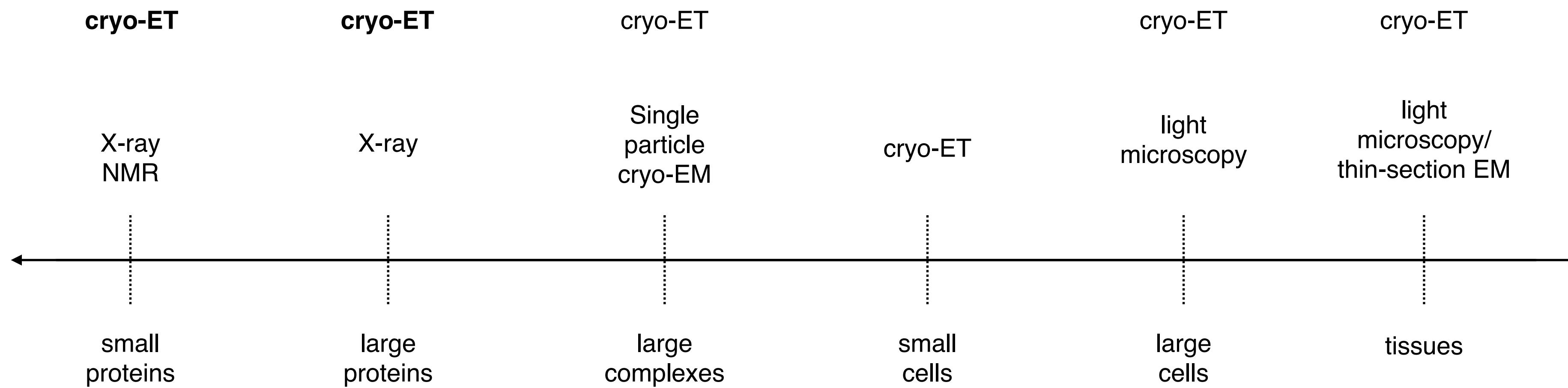
Tomography of nanocrystals





Electron tomography of nanocrystals

- Uses crystals only a few hundred nanometers thick
- Could use poorly-ordered crystals
- Could resolve twinning and joints
- Fast, accurate phases with no need for heavy atom derivatives or a molecular replacement model
- Cheaper (no synchrotron needed)



1. Fluorescence-guided FIB-milling
2. Rapid tilt series
3. **Tomography of nanocrystals**

cryo-ET



- Forecast rests on two principles:**
1. Electrons are better than X-rays
 - they can be focused
 - give more info for the same damage
 2. 3D is better than 2D

The physics is clear, if the technical challenges can be solved!