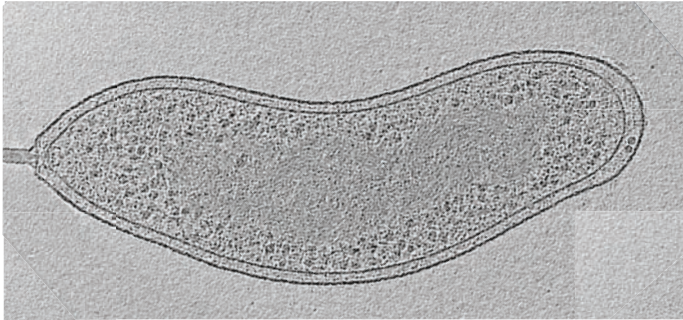


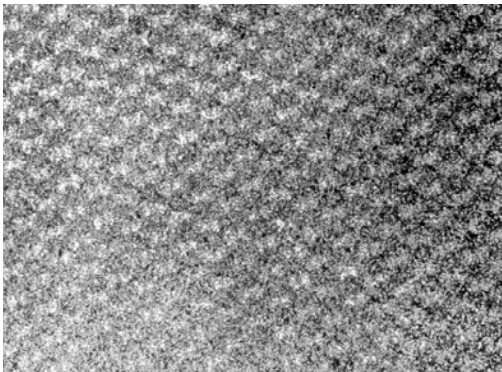
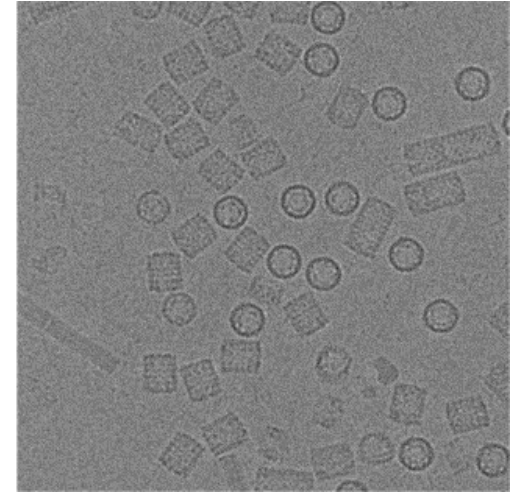
Part 7: Electron crystallography

Basic approaches in cryo-EM



Tomography

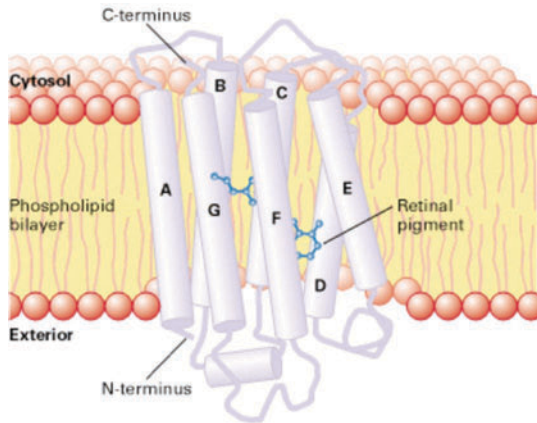
Single particle analysis



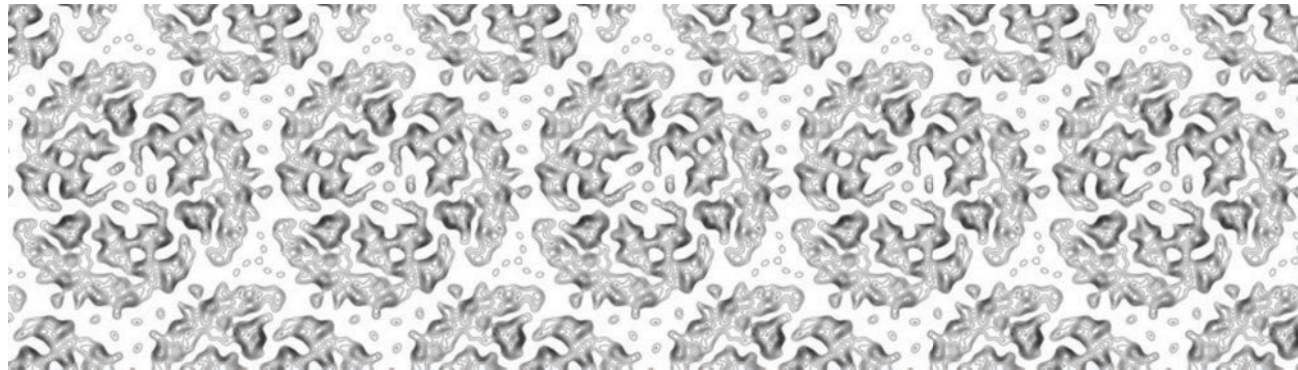
2D crystallography

Some proteins naturally assemble into 2D arrays

Example:
bacteriorhodopsin

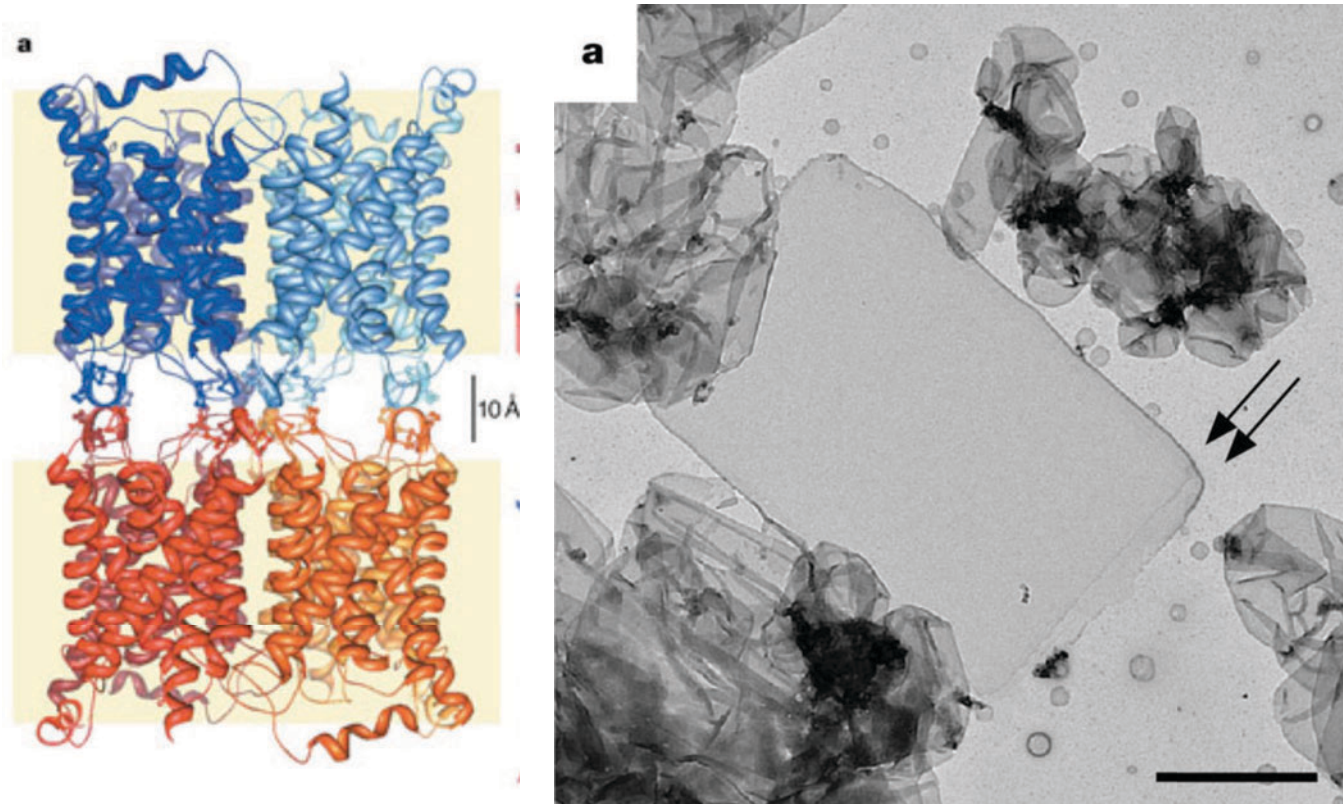


[http://www.unitus.it/
scienze/
corsonew/
lezione11.html](http://www.unitus.it/scienze/corsonew/lezione11.html)



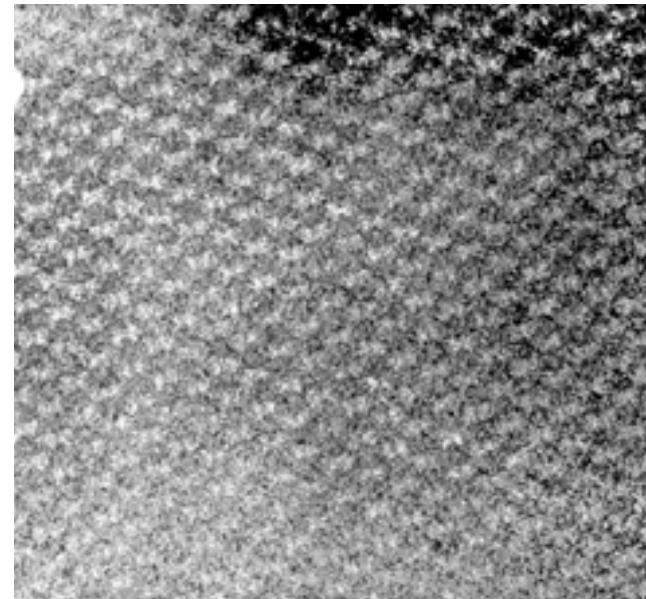
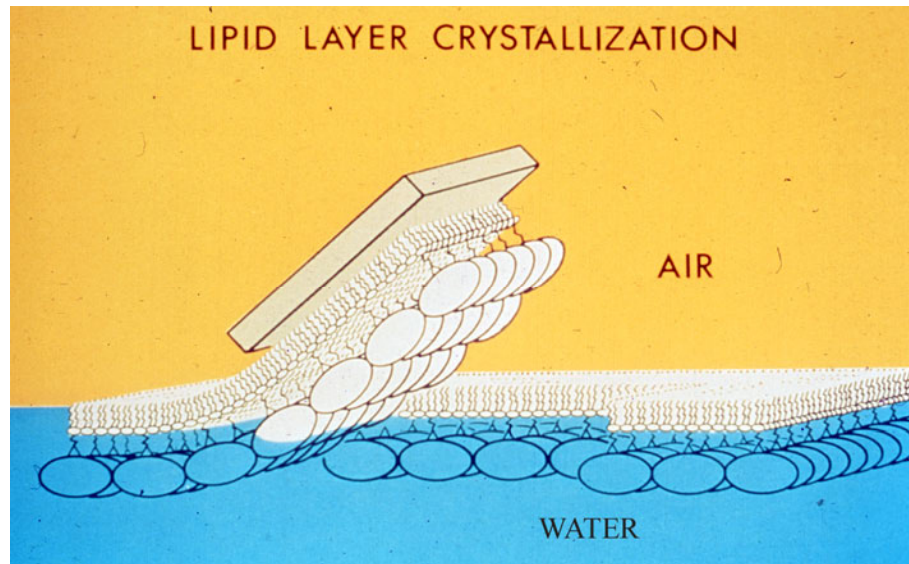
Grigorieff et al., JMB 1995

Second example: aquaporin



Gonen et al., Nature 2004

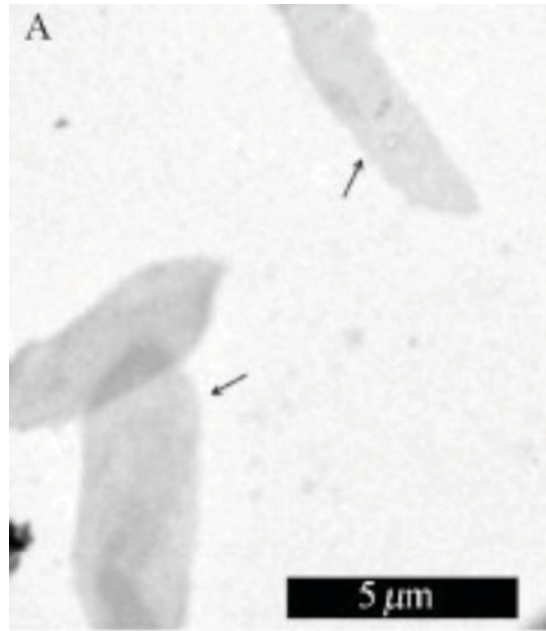
Others can be crystallized in-vitro



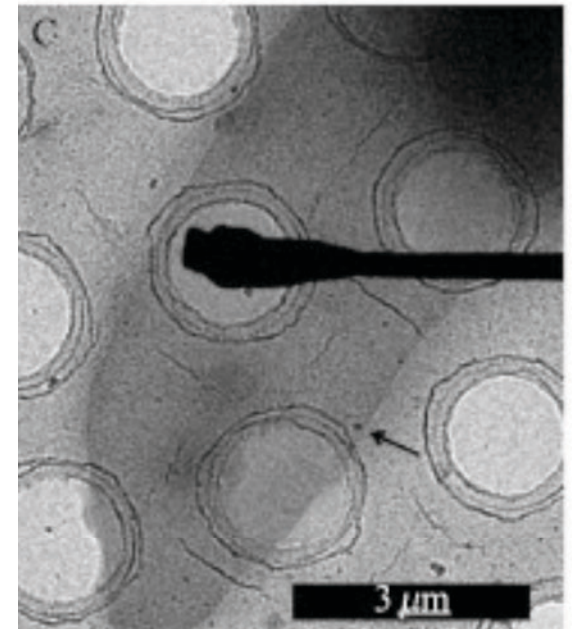
RNA polymerase

Courtesy Roger Kornberg

After formation, crystals can be embedded in a sugar like trehalose and dried, or plunge-frozen



embedded in trehalose
between two layers of
continuous carbon film



adsorbed onto holey
carbon film and
plunge-frozen

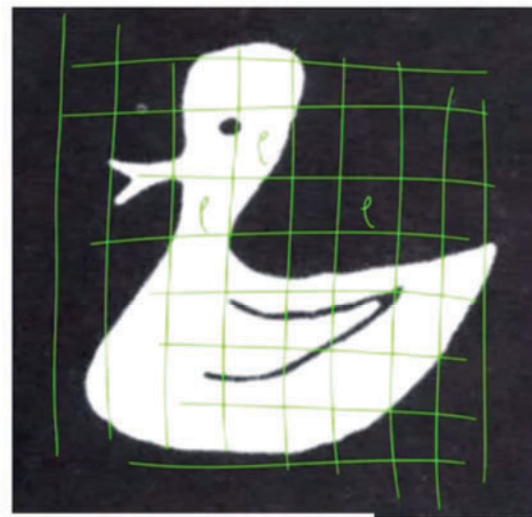
Abeyrathne et al., MIE 2010

2-D crystallography - Intro and sample prep

Concept check questions:

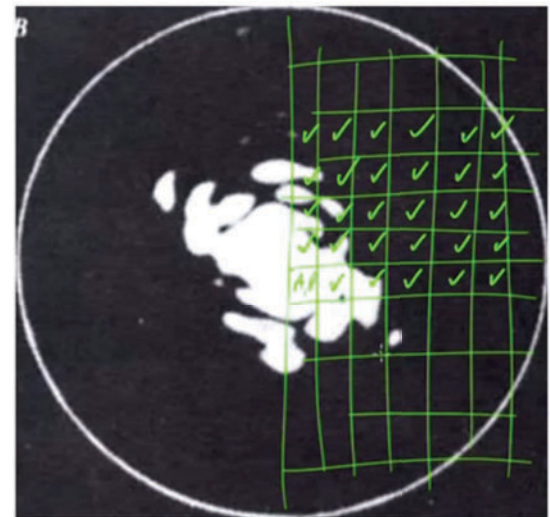
- What is a “2-D crystal”?
- When is 2-D crystallography the cryo-EM approach of choice?
- Describe a method for inducing a protein of interest to form a 2-D crystal.
- In addition to plunge-freezing, what other way have 2-D crystals been stabilized for EM imaging?

The Fourier transform of an asymmetric object



10x10

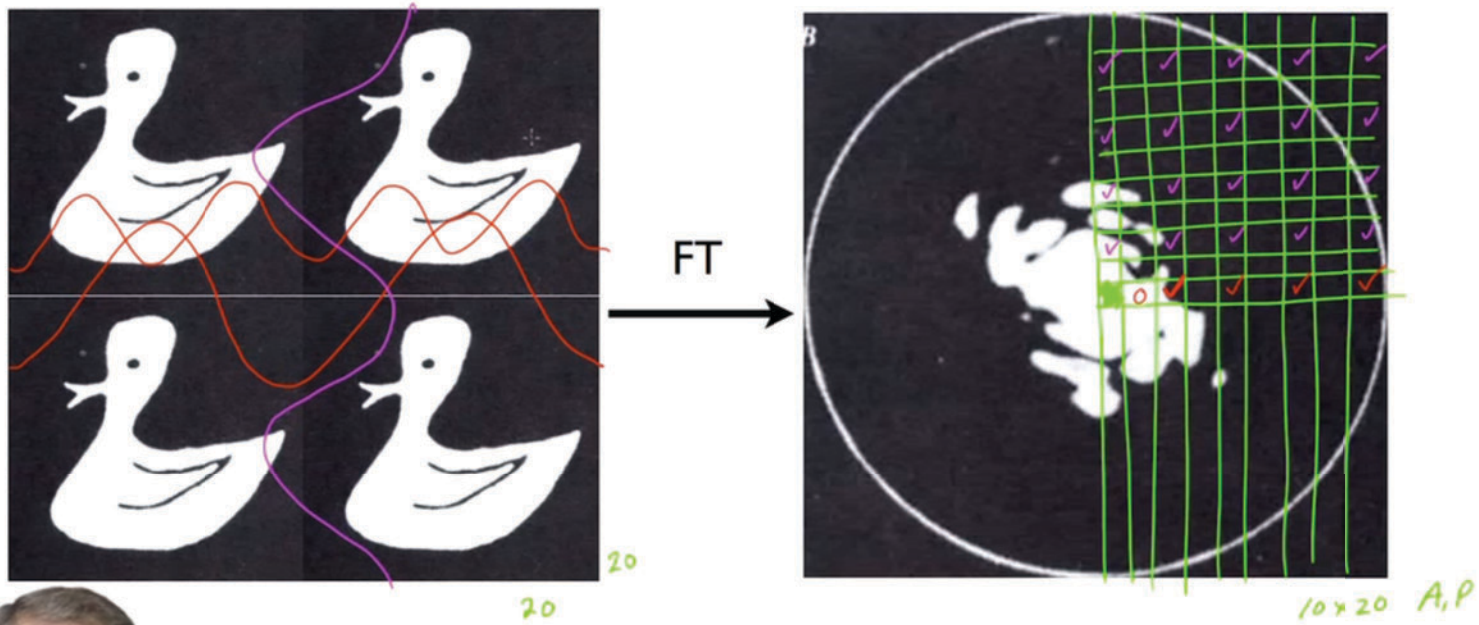
FT
→



5x10 48P

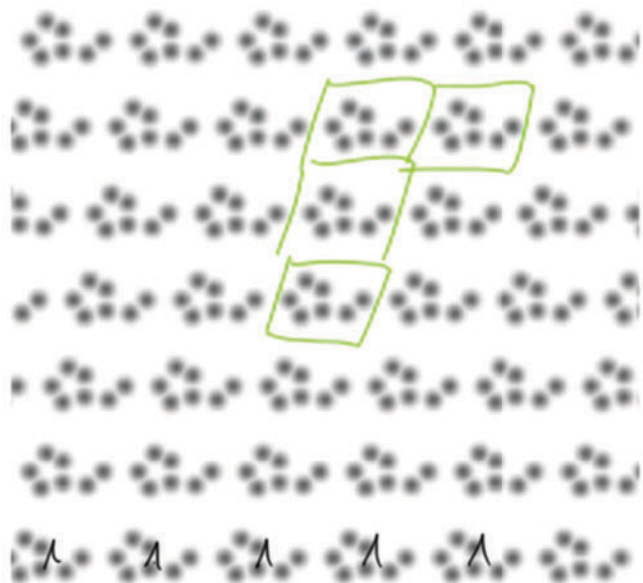
all pixels are significant



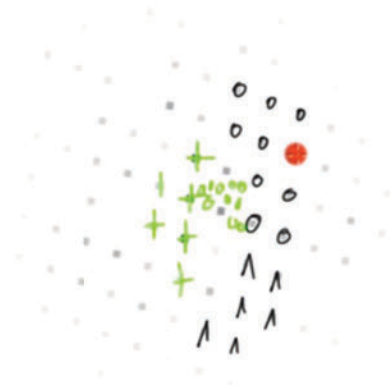


In the Fourier transform of a crystal, some pixels have significant non-zero values and others do not





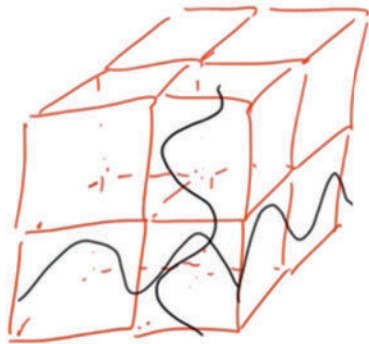
FT
→



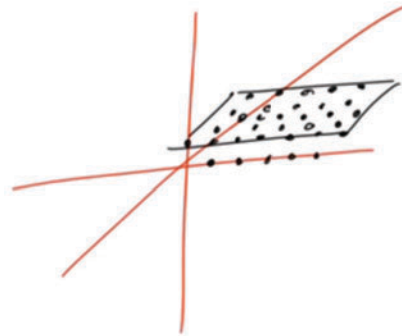
Object \otimes lattice

$$FT\{obj \otimes \mathcal{L}\} = FT\{obj\} \cdot \underbrace{FT\{\mathcal{L}\}}_L$$

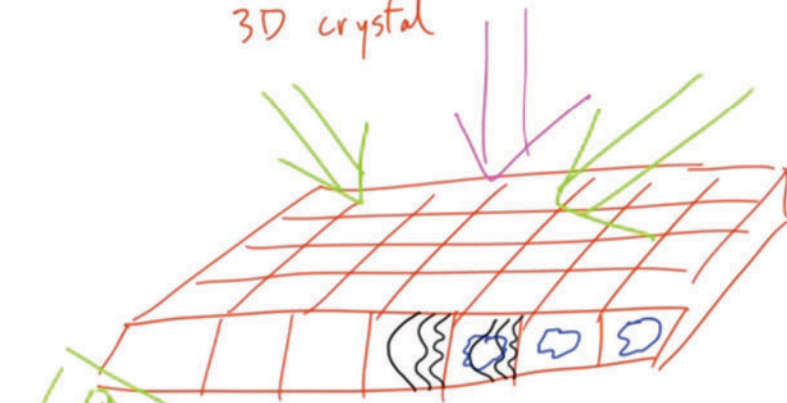




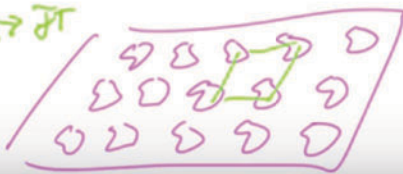
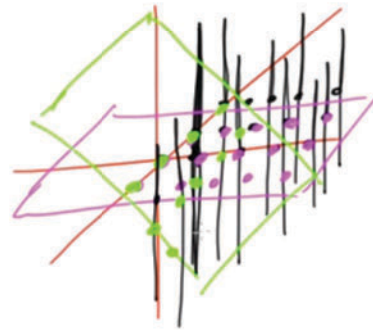
$\tilde{F}T$



3D crystal



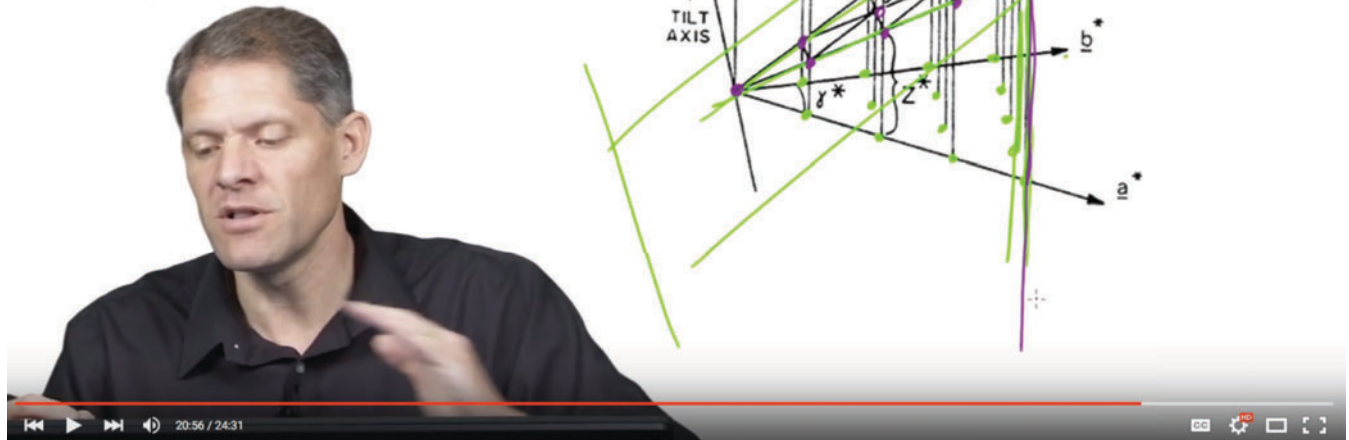
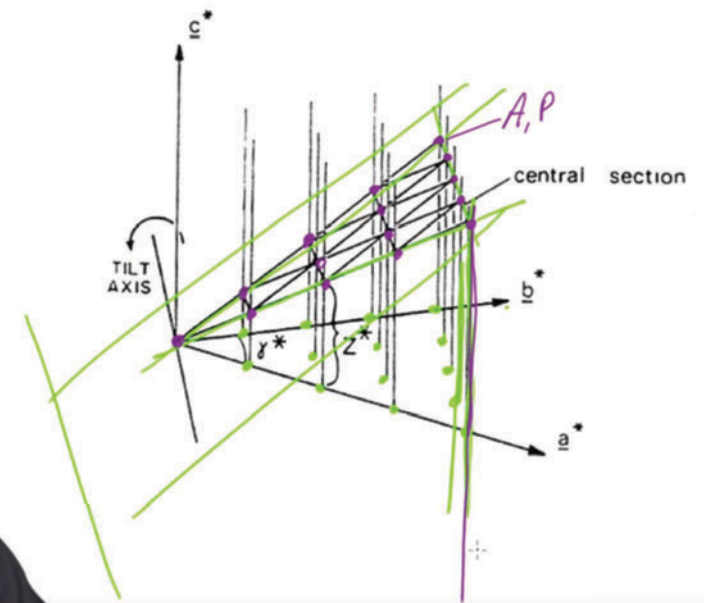
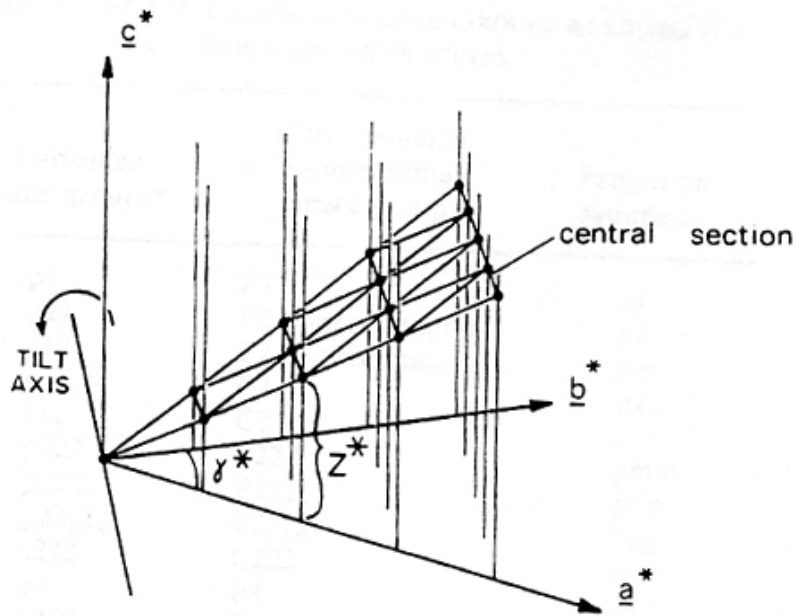
$\tilde{F}T$

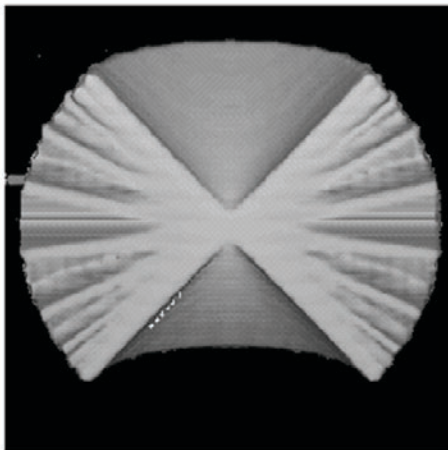
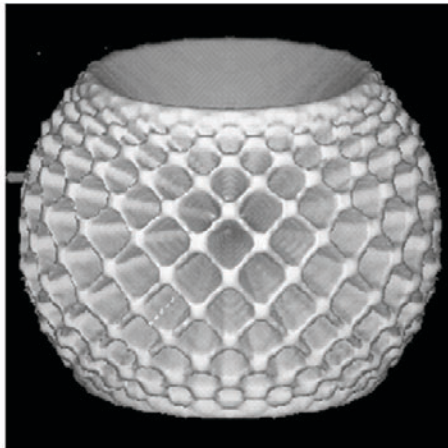


$\tilde{F}T$

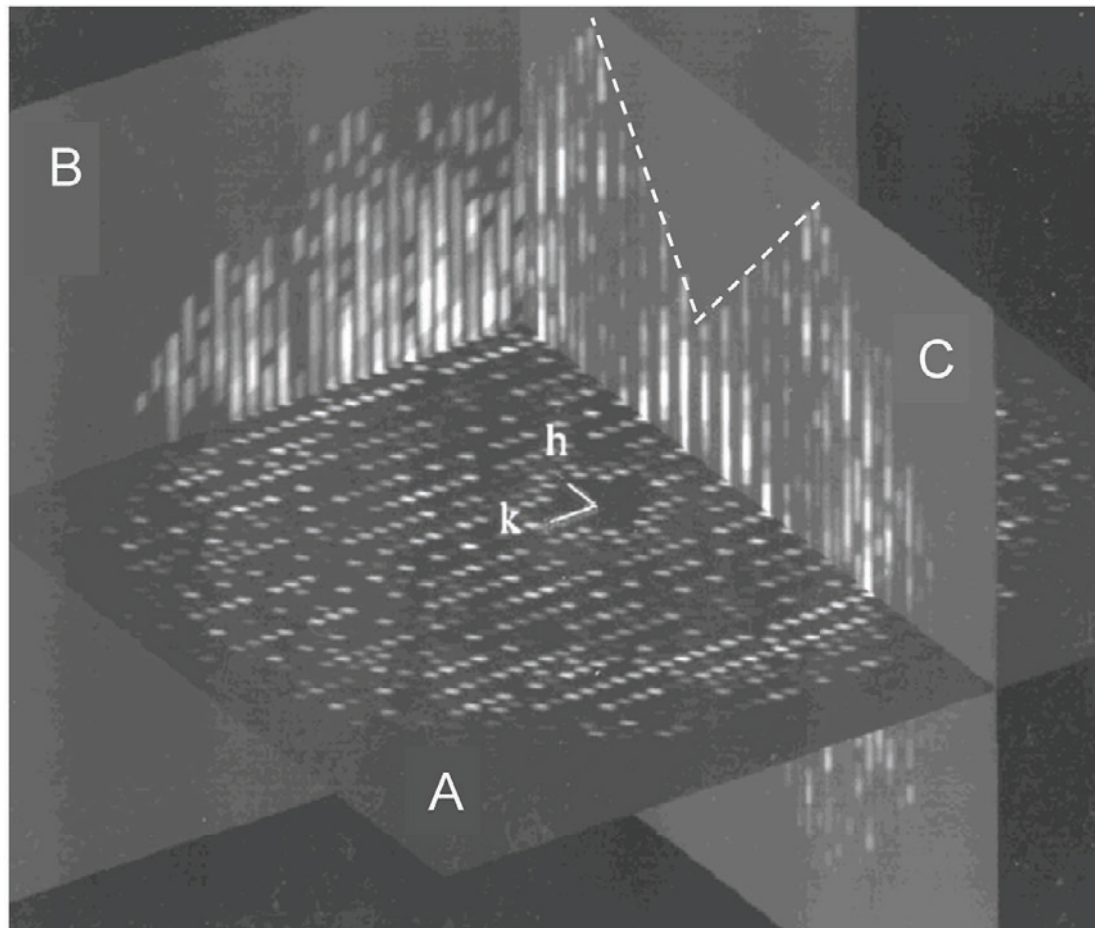


How the Fourier transforms of images of 2-D crystals sample the 3-D Fourier transform of the object

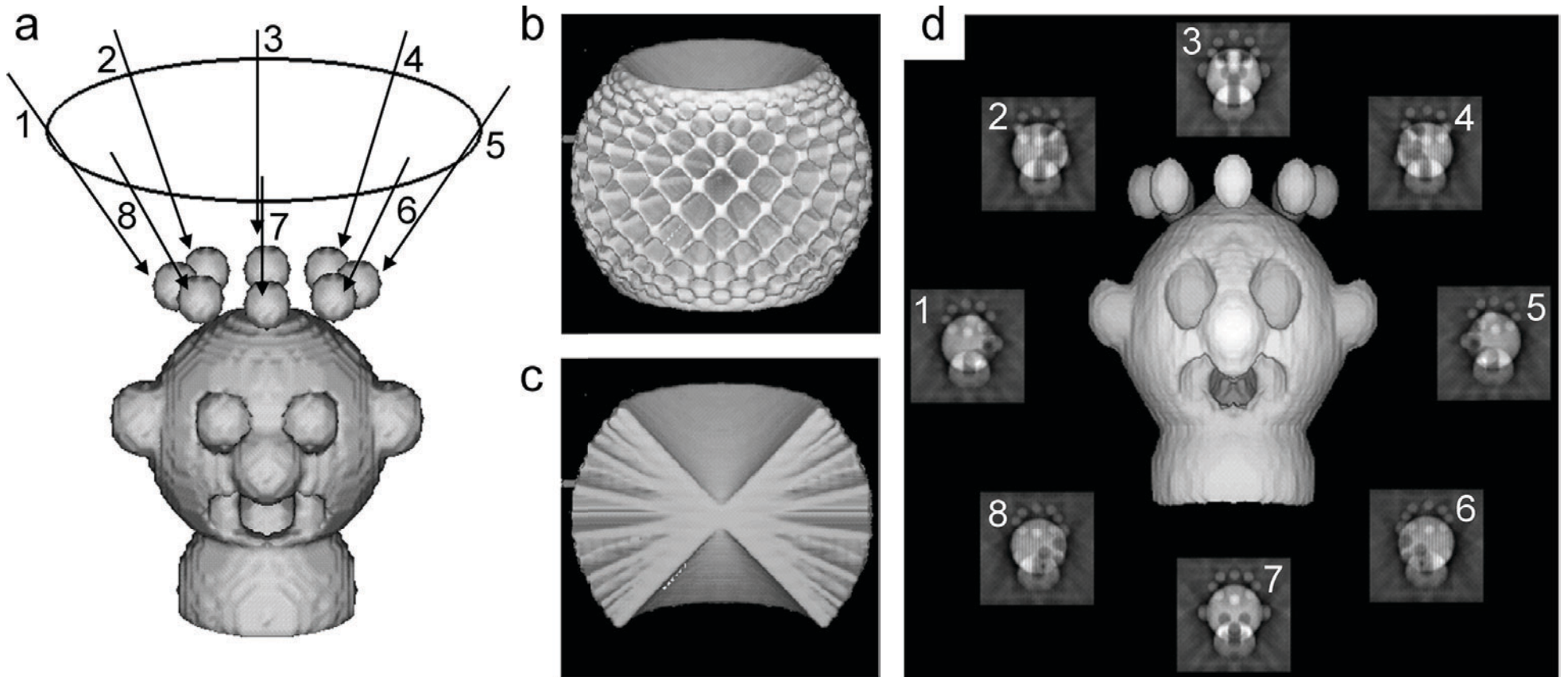




Boisset et al.,
Ultramicroscopy 1998



Orlova and Saibil,
Chemical Reviews 2011

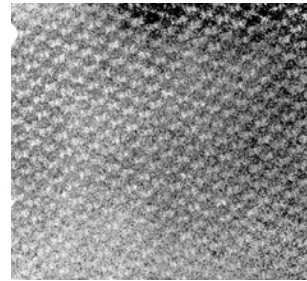
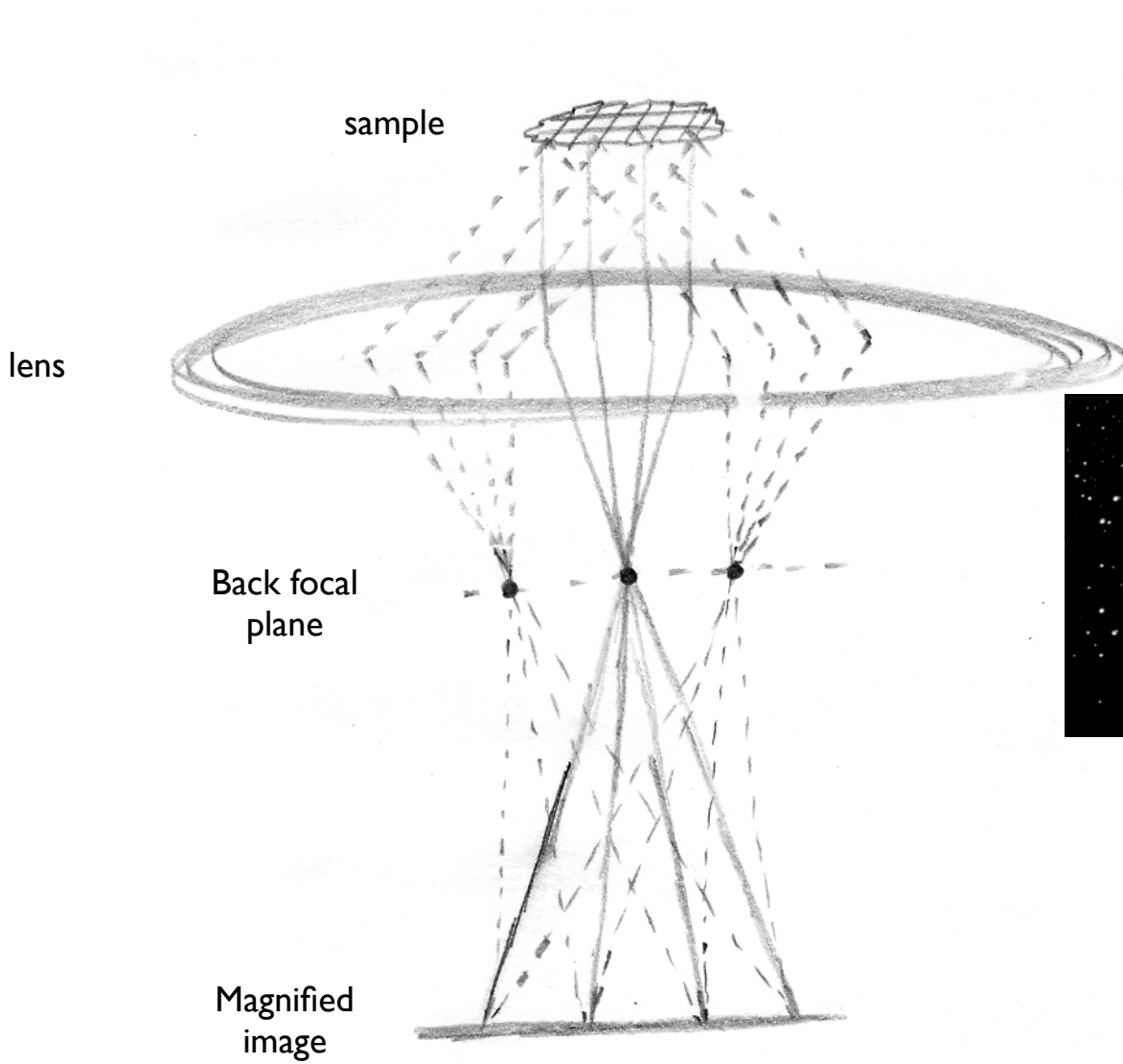


Boisset et al., Ultramicroscopy 1998

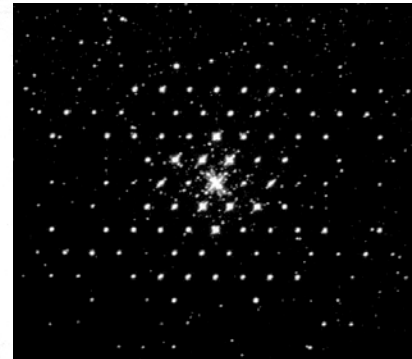
Fourier transform of a 2-D crystal

Concept check questions:

- Why does the Fourier transform of a crystalline object have discrete spots separated by pixels with near-zero amplitudes?
- What is the convolution theorem, and what does it have to do with crystallography?
- What does the Fourier transform of a 2-D crystal look like?
- What is the “missing cone,” why is it “missing,” and what effect does it have on 2-D crystallographic reconstructions?

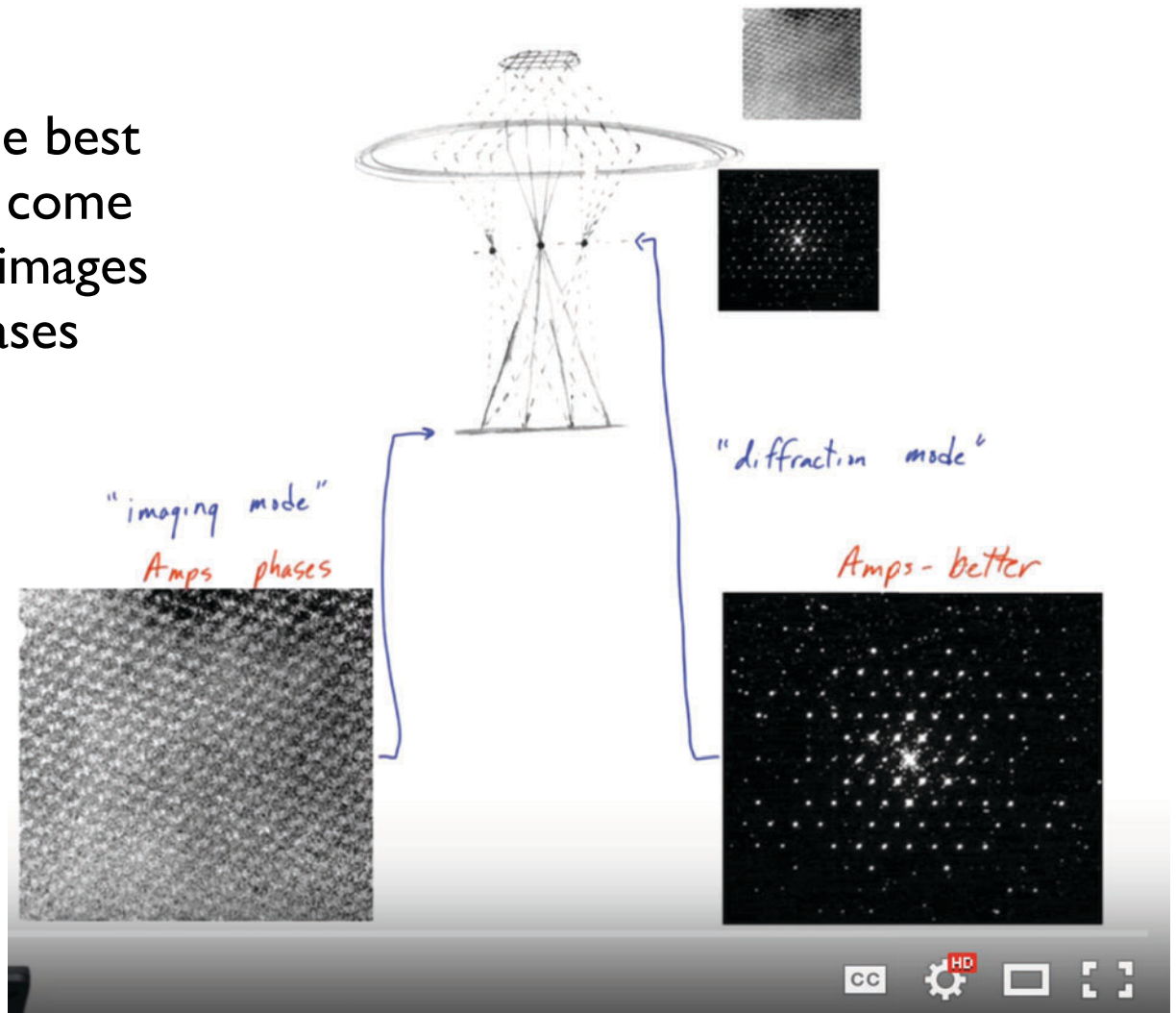


2D crystal

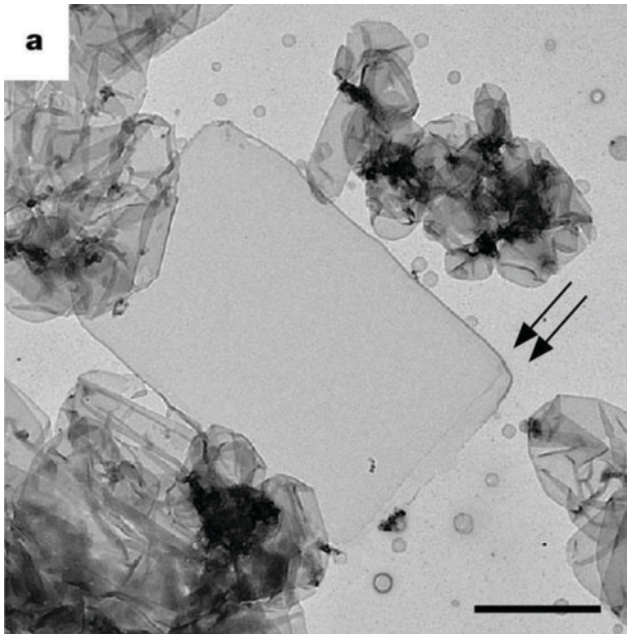


diffraction pattern on back focal plane

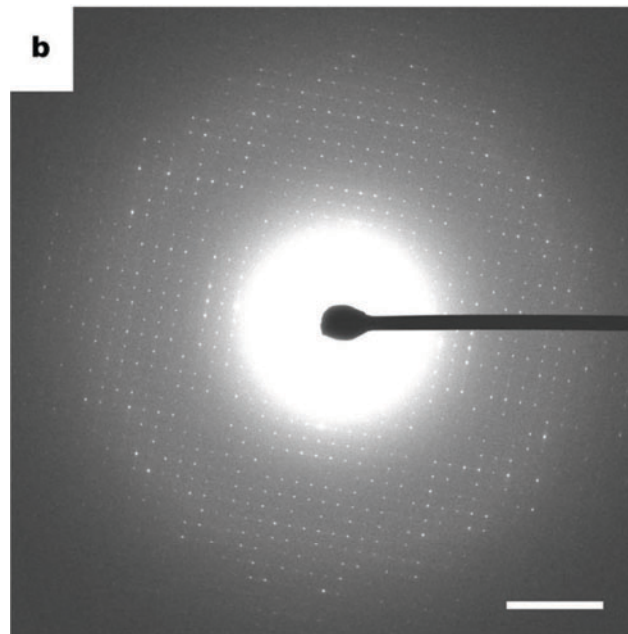
In electron crystallography, the best measurements of amplitudes come from diffraction patterns, but images are recorded to obtain phases



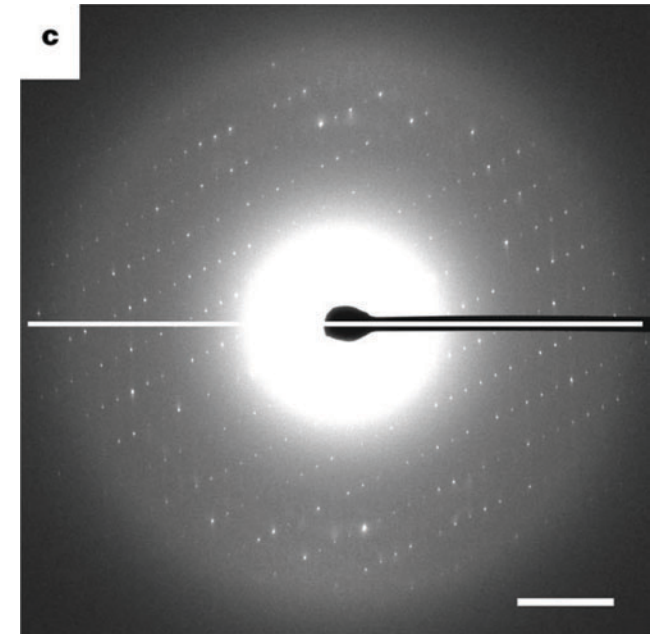
Example images and diffraction patterns from aquaporin crystals



Aquaporin crystal



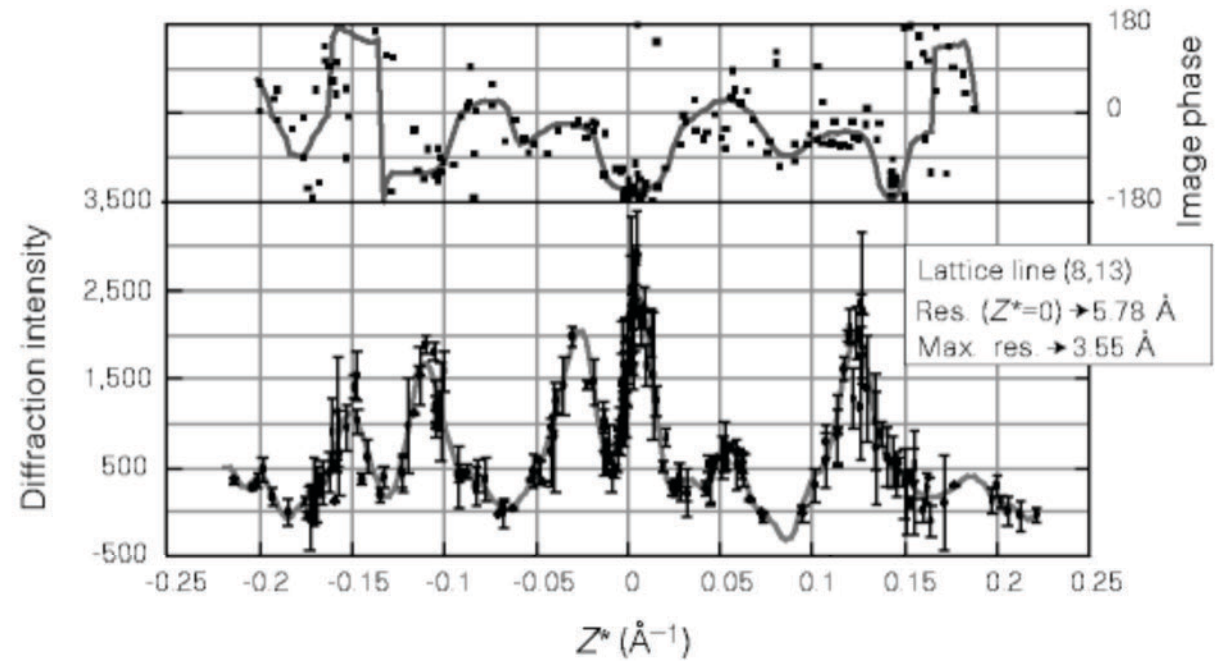
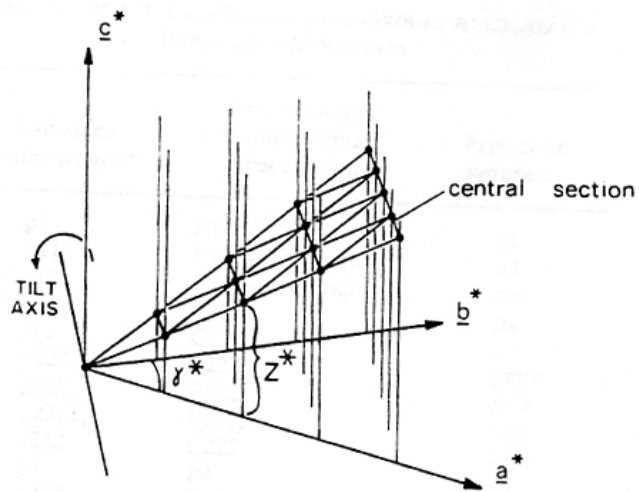
Electron diffraction pattern
of an untilted crystal



Electron diffraction pattern
of a crystal tilted to 70°

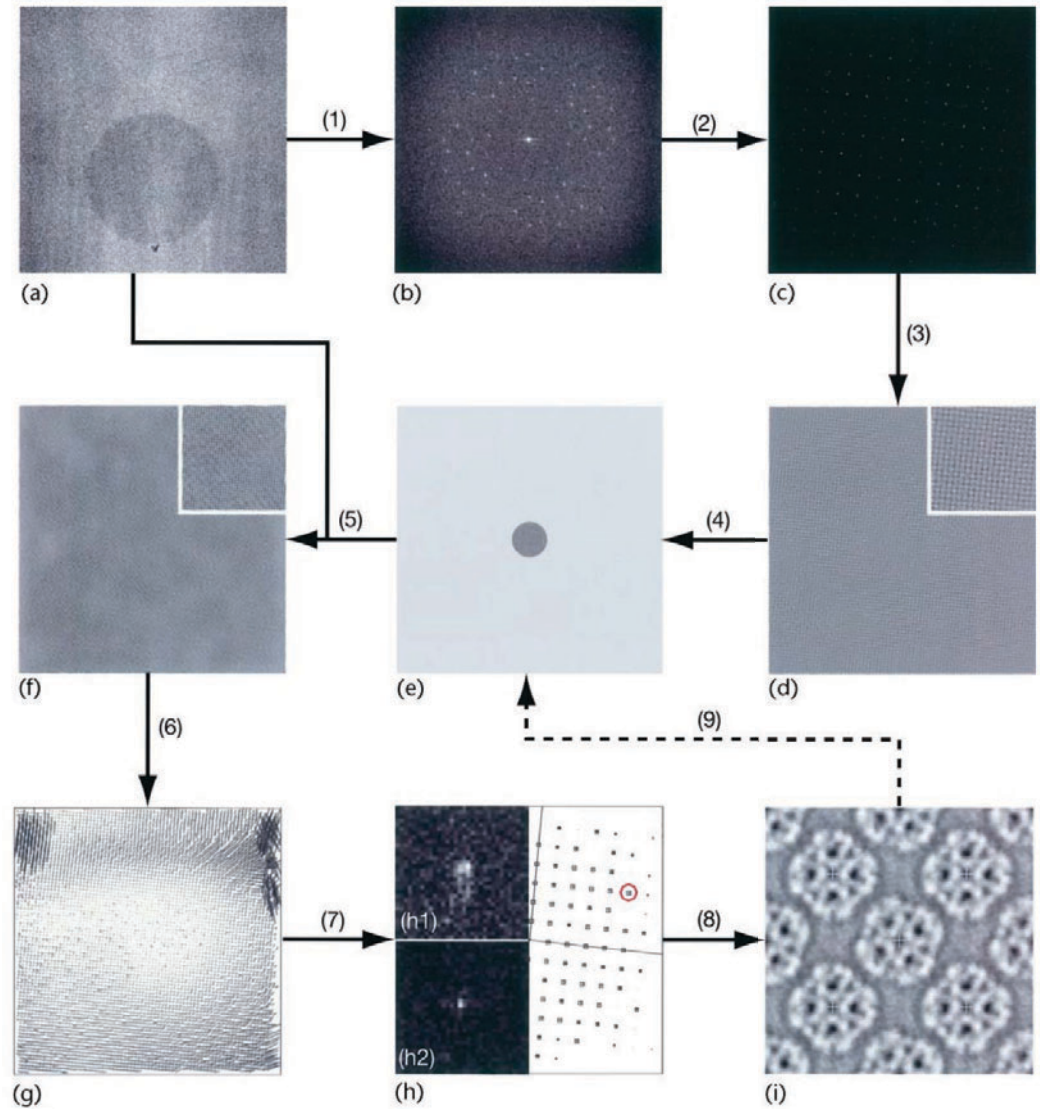
Gonen et al., Nature 2004

Example lattice line data (amplitude and phase) and curve fitting



Nogales et al., Nature 1998

Crystal “unbending”



Braun and Engel,
Encyclopedia of Life Sciences 2005

Challenges in 2D crystallography

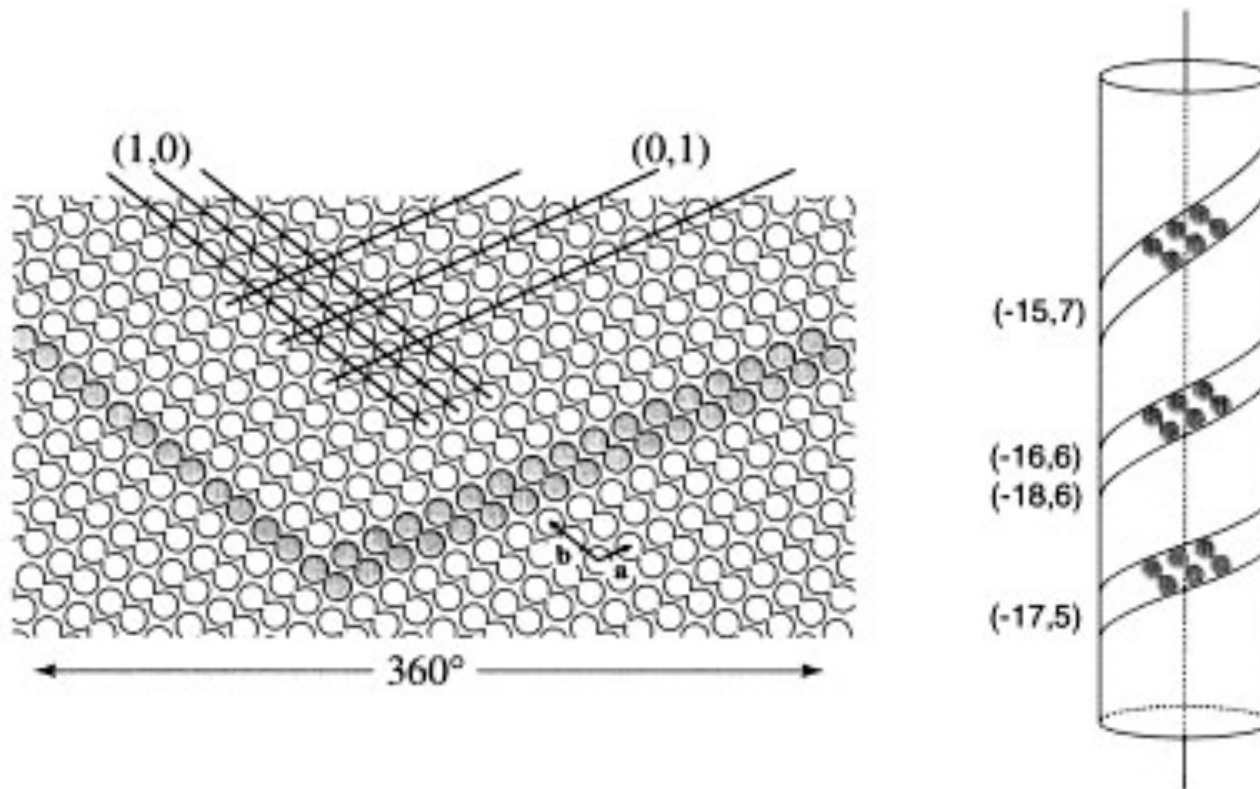
- Hard to get well-ordered crystals
- Hard to get flat crystals
- Charging, beam-induced movement can blur images
- “Missing cone”

2-D crystallography - Data collection and reconstruction

Concept check questions:

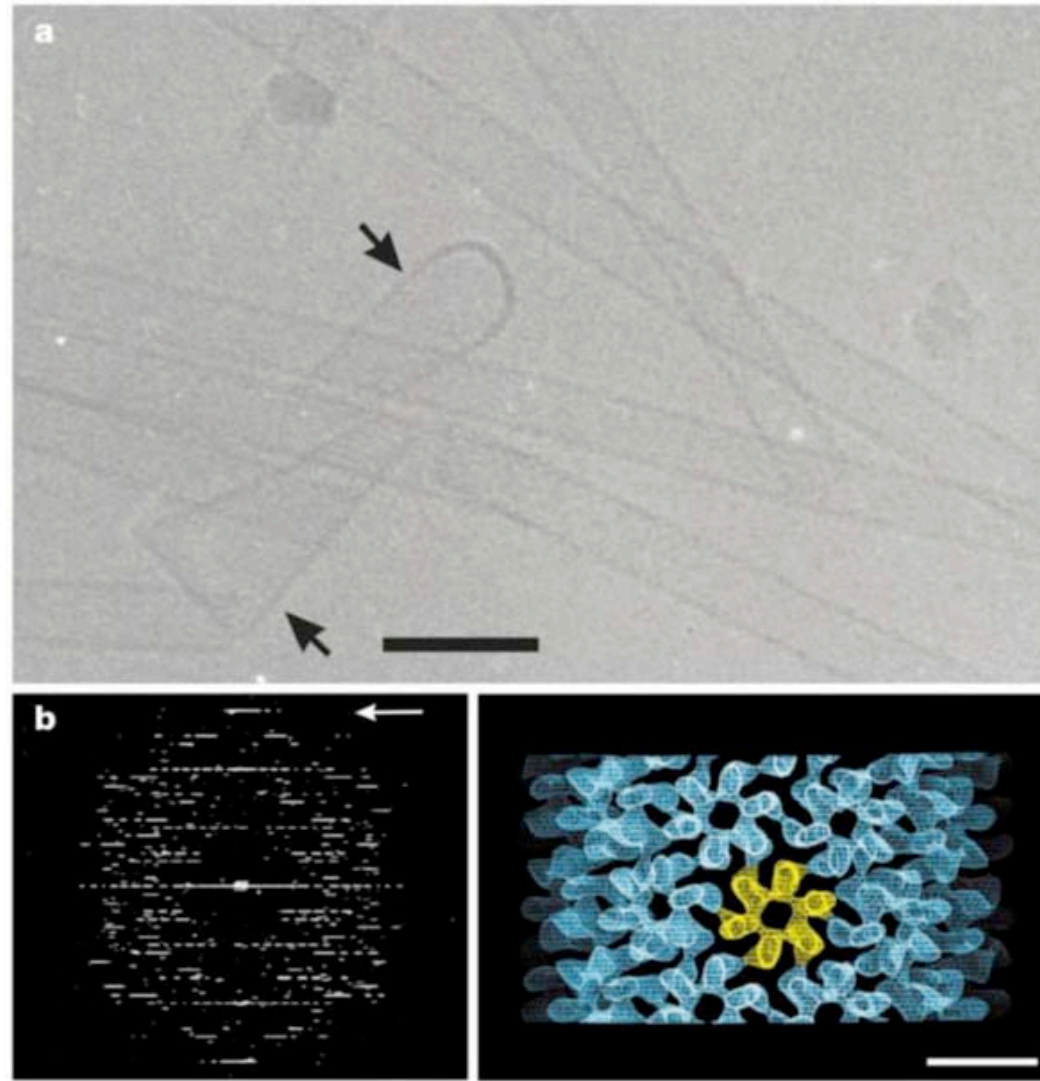
- What is the difference between “imaging” and “diffraction” modes on an EM?
- Why are both images and diffraction patterns of 2-D crystals recorded in a 2-D crystallography project?
- Why are images of both untilted and tilted samples recorded?
- How is all the data from all these images and diffraction patterns merged to produce the reconstruction?
- What is crystal “unbending”? How and why is it done?
- Describe four common challenges in 2-D crystallography projects.

Helical tubes are “rolled up” versions of 2-D crystals, can be rolled up into different families of tubes with different pitches



Miyazawa et al., JMB (1999)

Cryo-EM projection
image of a helical
tube of purified HIV
CA protein



Su Li et al.
Nature 2000

Power spectrum
shows “layer lines”

3D
reconstruction

Helical tubes

Concept check questions:

- How are helical tubes related to 2-D crystals?
- Why are helical tubes particularly good samples for cryo-EM reconstruction?
- What does the diffraction pattern of a helical tube look like?