# EM Image Formation and Single-Particle Reconstruction

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Phase contrast and the contrast transfer function

#### Phase contrast and the contrast transfer function, Part I

- 1. Complex numbers: review
- 2. Defocus contrast (the simple version)
- 3. Image delocalization

#### Why complex numbers?

- They make the equations simpler
- Natural for Fourier transforms
- Give us the magnitude and phase of structure factors

#### i, the imaginary unit

$$i = \sqrt{-1}$$

A complex number Real part 
$$z = a + ib$$
 Imaginary part

#### You can do arithmetic with complex numbers

$$z = a + ib$$

$$w = c + id$$

Add 
$$z + w = (a + c) + i(b + d)$$

Multiply 
$$zw = (ab - bd) + i(ad + bc)$$

Real part 
$$Re(z) = a$$

Imaginary part 
$$Im(z) = b$$

Absolute value 
$$|z| = \sqrt{a^2 + b^2}$$

Conjugate 
$$z^* = a - ib$$

#### The exponential function $e^x$

$$e = 2.718...$$

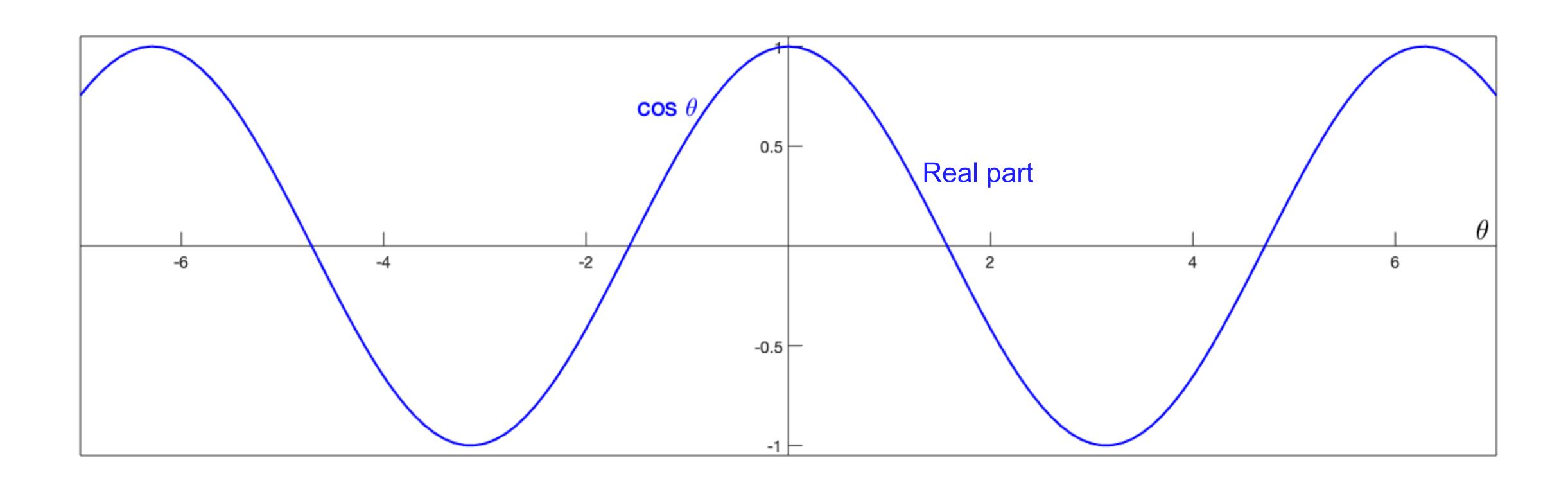
$$e^x = 1 + x + \frac{x^2}{2} + \frac{x^3}{2 \cdot 3} + \dots$$

A very important approximation, valid when  $x \ll 1$ , is

$$e^x \approx 1 + x$$

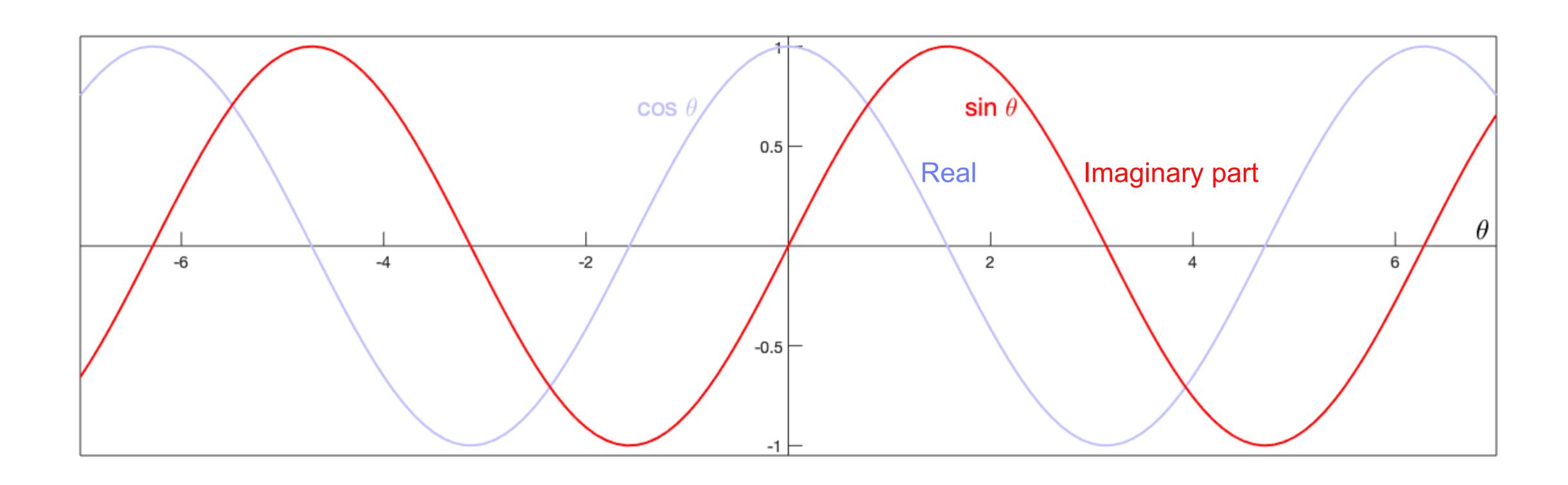
## The complex exponential $e^{i\theta}$

$$e^{i\theta} = \cos\theta + i\sin\theta$$

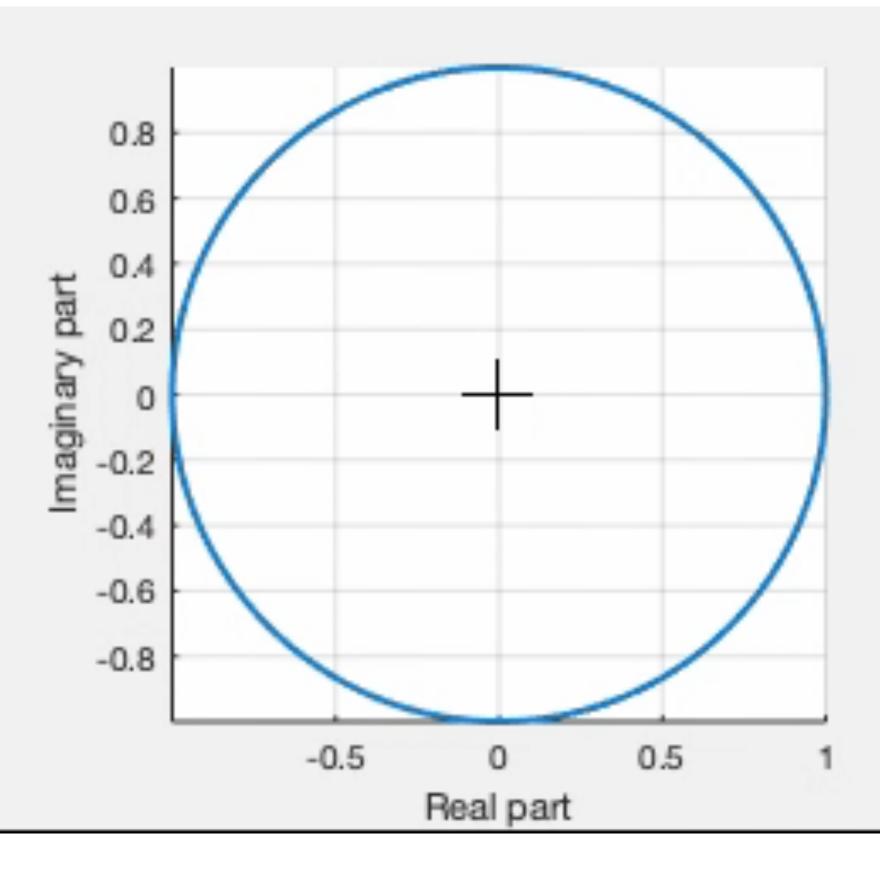


### The complex exponential $e^{i\theta}$

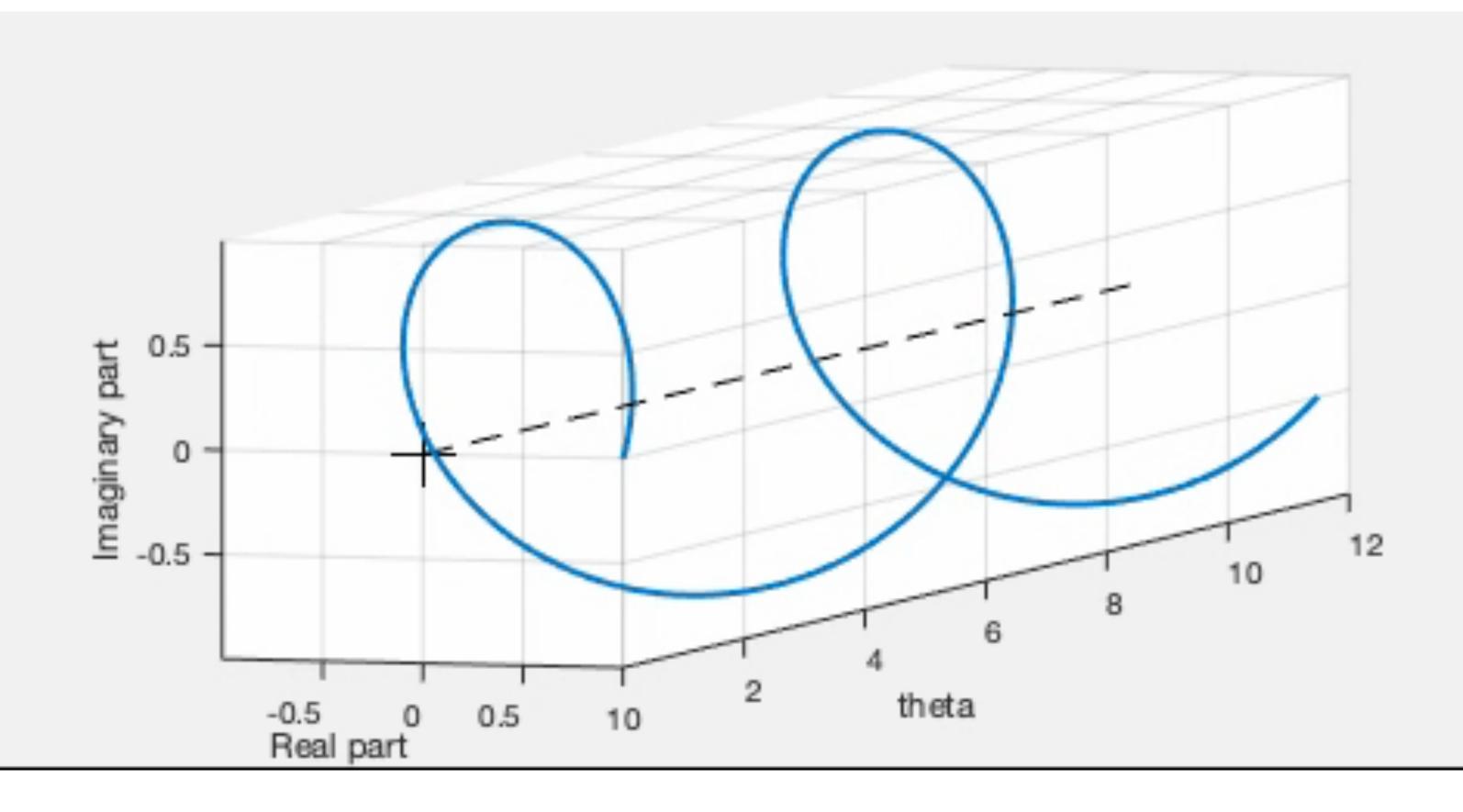
$$e^{i\theta} = \cos\theta + i\sin\theta$$



# A plot of $e^{i\theta}$



# A plot of $e^{i\theta}$



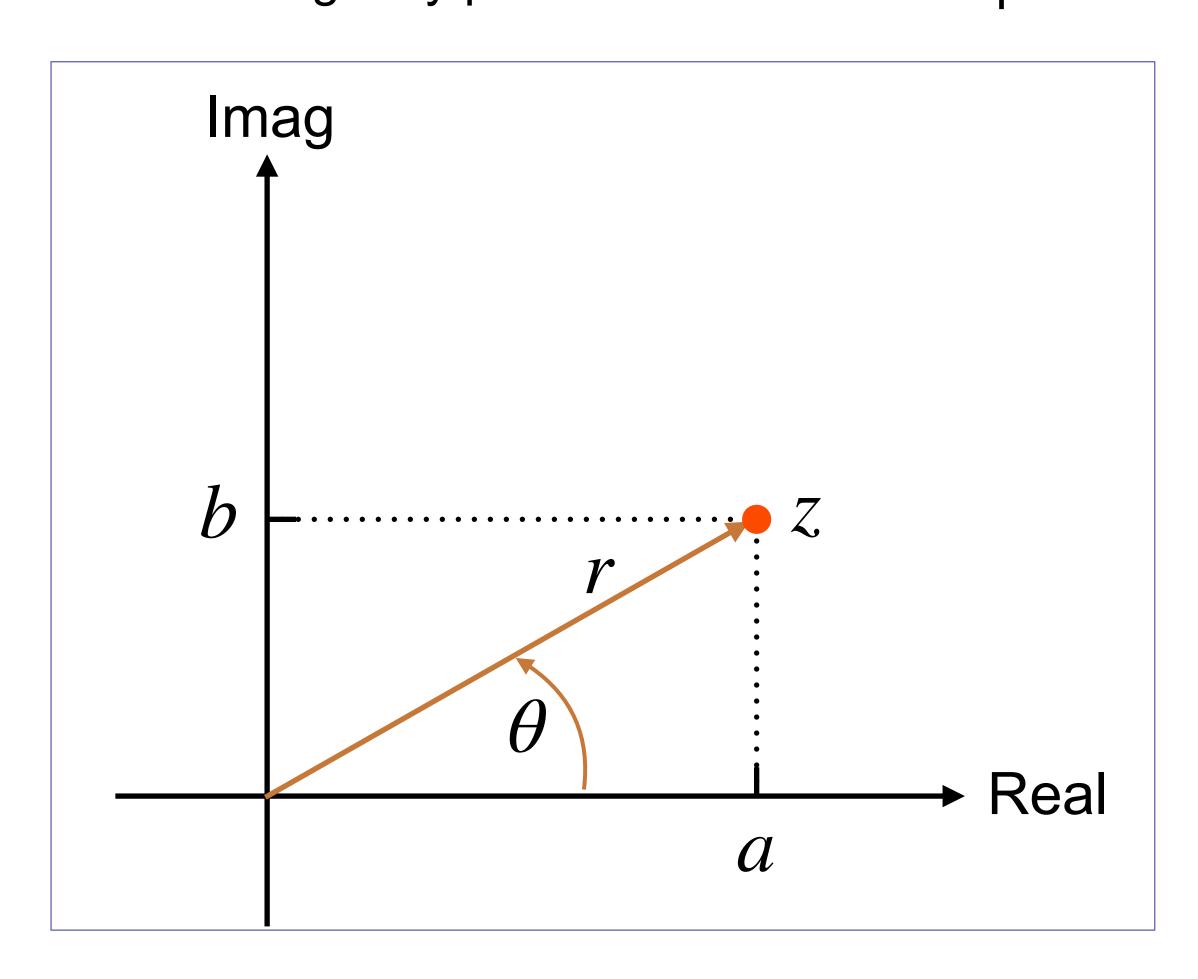
### Any z can be represented as (a, b) or as $(r, \theta)$

$$z = a + ib$$

a is the real part b is the imaginary part

$$z = re^{i\theta}$$

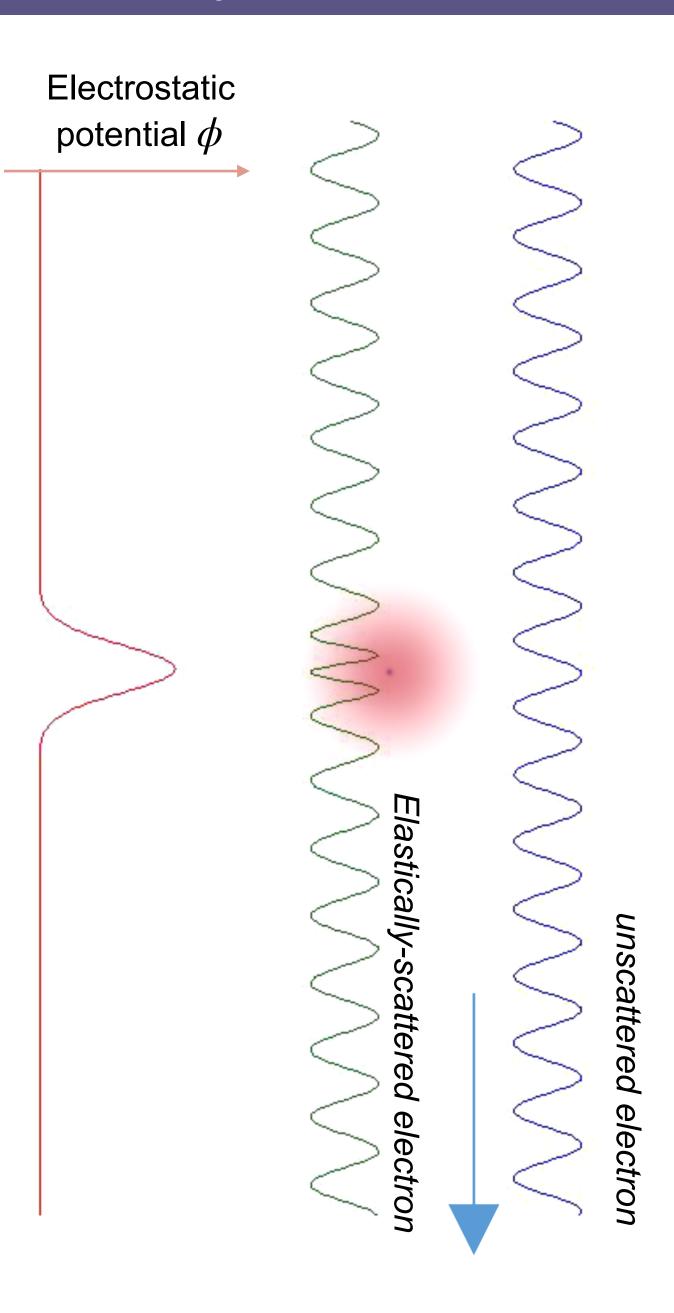
r is the magnitude  $\theta$  is the phase



#### Phase contrast and the contrast transfer function, Part I

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#### Cryo-EM specimens are imaged by phase contrast

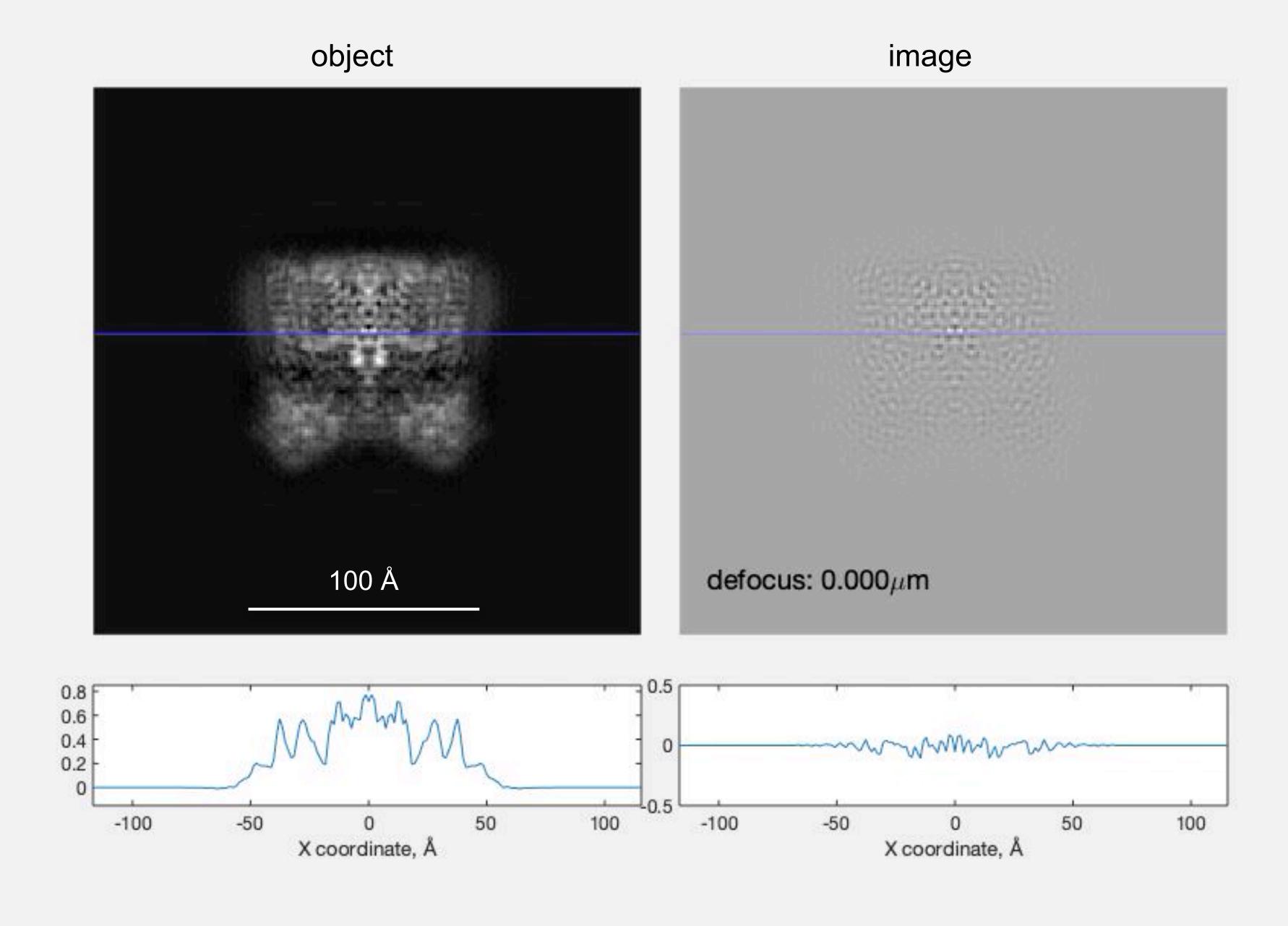


The imaging electrons are phase-shifted when passing near atomic nuclei or fixed charges.

The phase shift coefficient  $\sigma$  is about 0.5 milliradian per volt-angstrom of integrated potential.

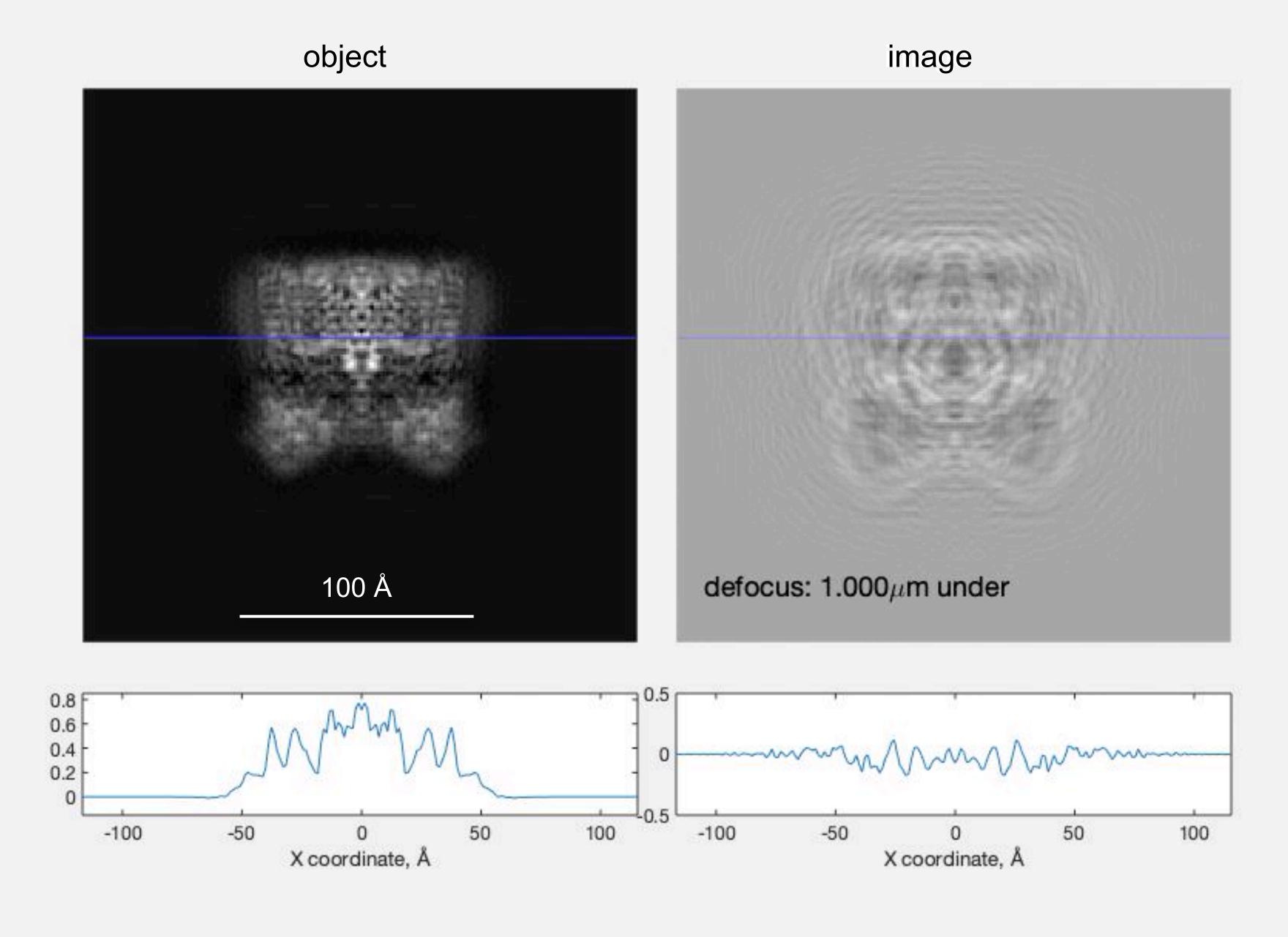
The phase shift near a single atom is ~1 milliradian.

#### Most cryo-EM data are acquired using defocus contrast



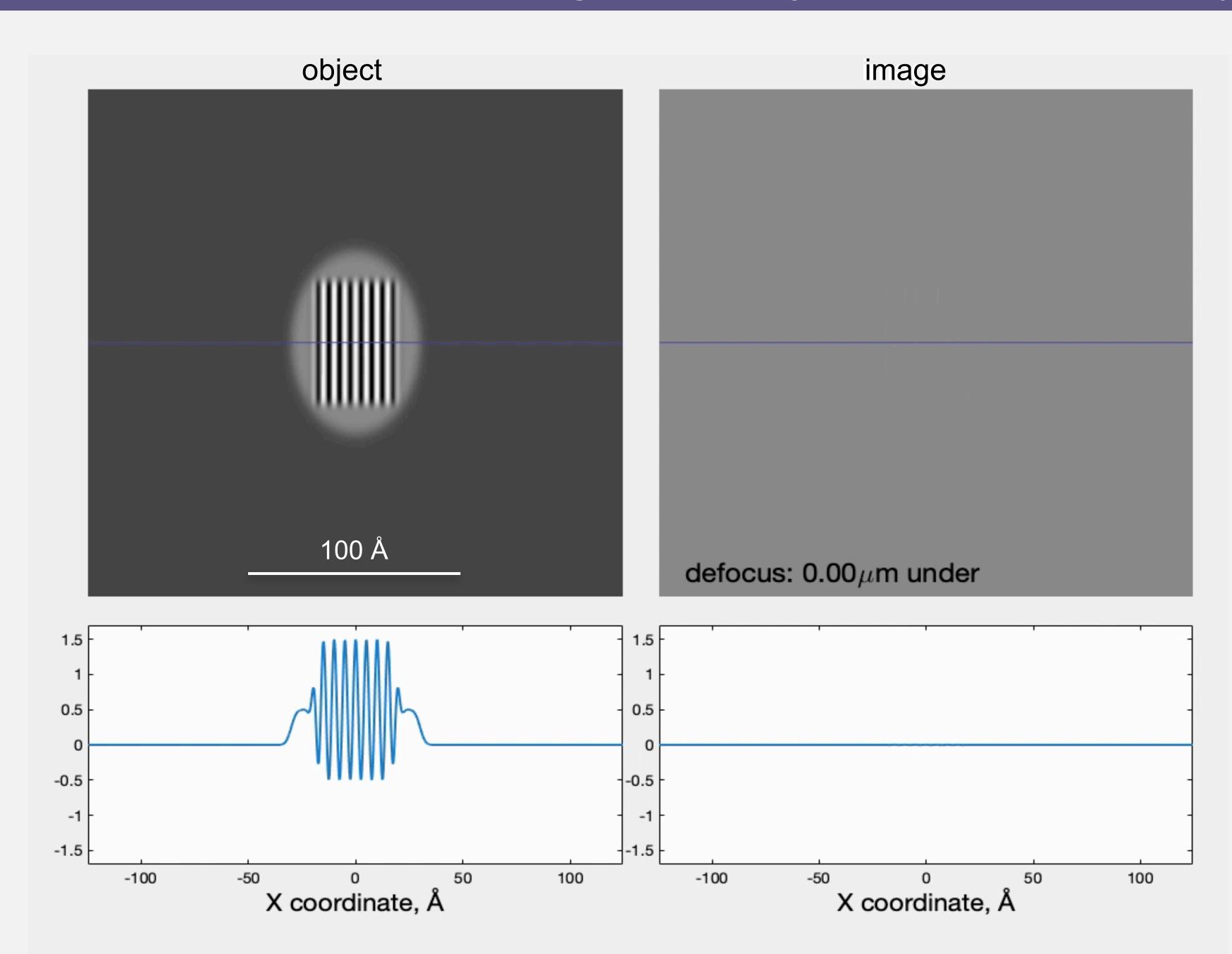
- At high defocus, highresolution information in the image is strongly delocalized
- Image processing can relocalize the signals, but at most only about half of the theoretical contrast is preserved by defocusing.

#### Most cryo-EM data are acquired using defocus contrast



- At high defocus, highresolution information in the image is strongly delocalized
- Image processing can relocalize the signals, but at most only about half of the theoretical contrast is preserved by defocusing.

#### Image of an object with 5Å periodicity

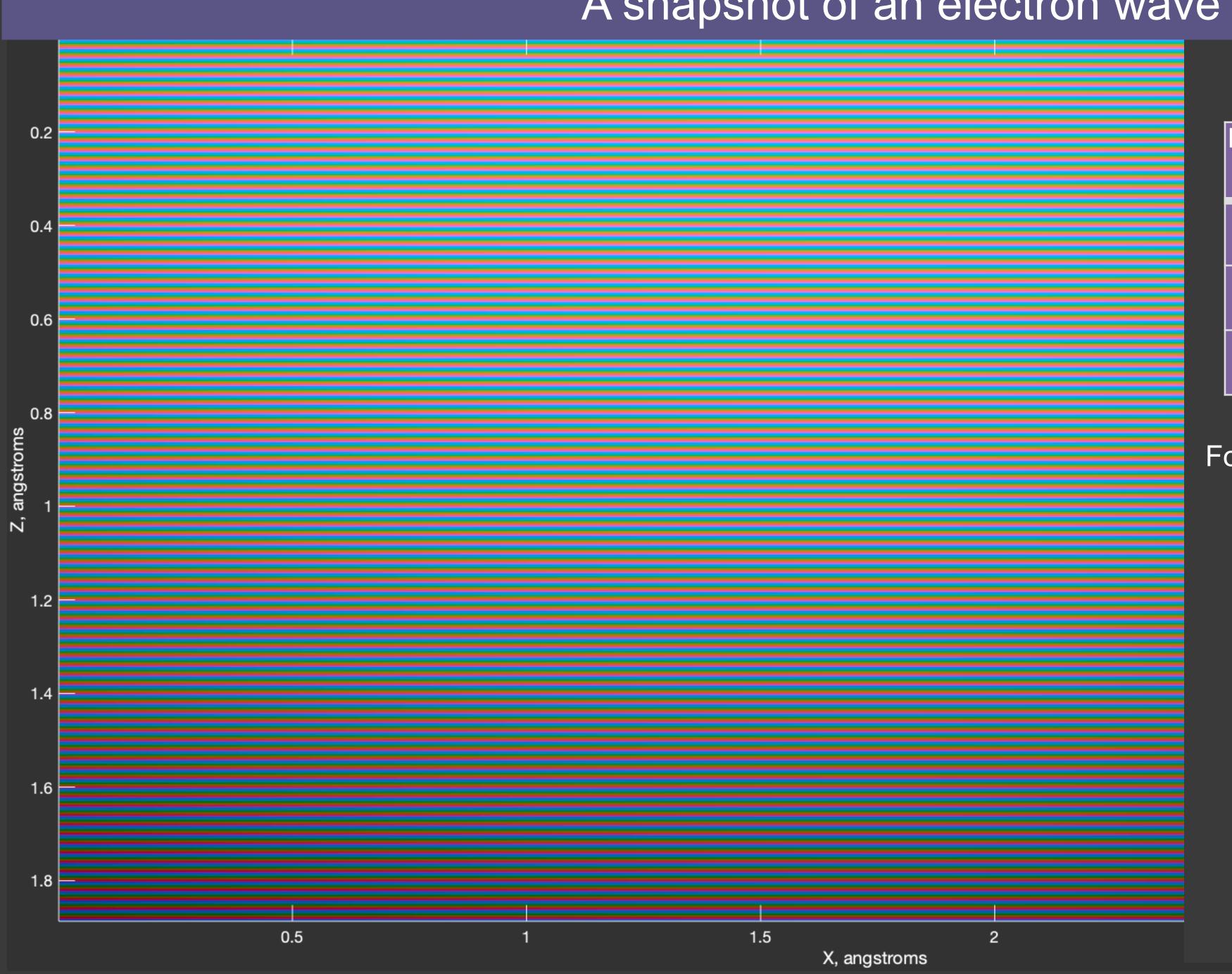


- At high defocus, highresolution information in the image is strongly delocalized
- Image processing can relocalize the signals, but at most only about half of the theoretical contrast is preserved by defocusing.

#### Defocus contrast in a nutshell

- 1. The contrast in the image of a grating object varies with the amount of defocus.
- 2. The grating object produces diffracted waves with shifting phase.
- 3. When the diffracted waves interfere with the undiffracted waves, we have contrast.

#### A snapshot of an electron wave



Energy (keV)	Wavelength (Å)	Velocity (fraction of c)
120	0.033	0.59
200	0.025	0.70
300	0.020	0.78

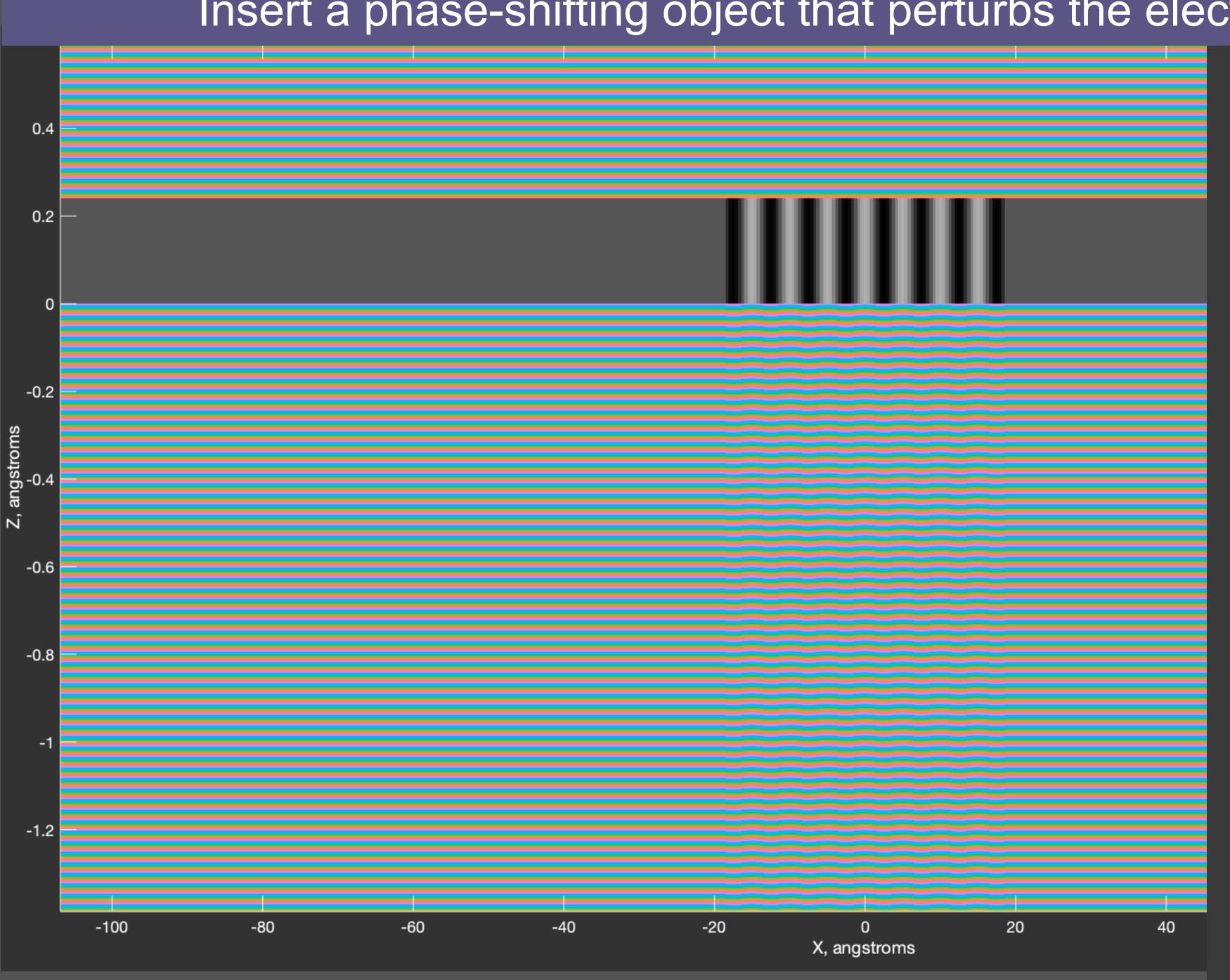
For an electron propagating in the z direction, the time-independent wave function is

$$\Psi_0 = e^{ikz}$$

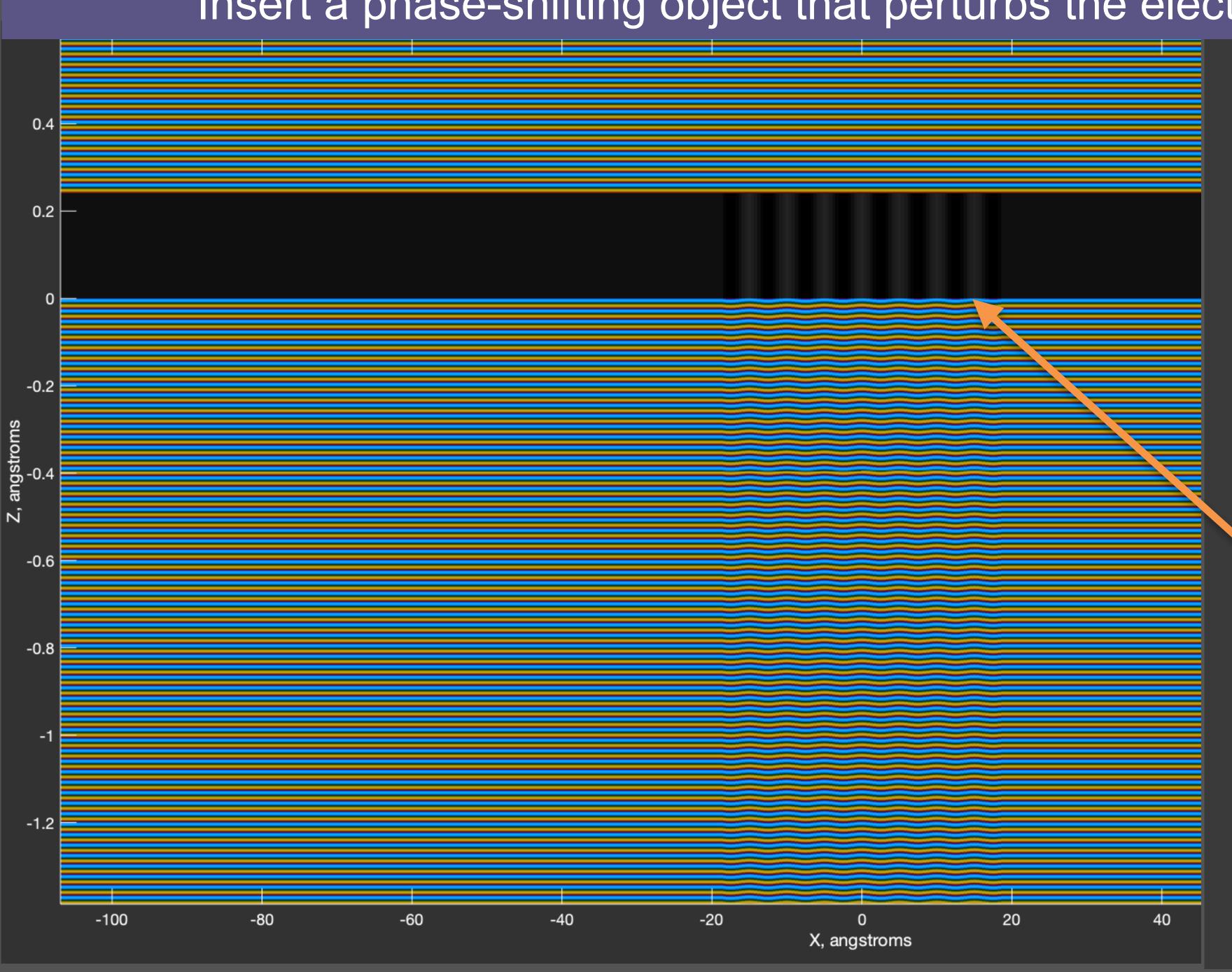
with

$$k = 2\pi/\lambda$$

Insert a phase-shifting object that perturbs the electron wave function



Insert a phase-shifting object that perturbs the electron wave function



The object is a grating,  $\epsilon \phi(x) = \epsilon \cos(2\pi x/d).$ 

Example:

 $d=5\mathring{\mathrm{A}}$  and  $\epsilon\ll1$ .

At 
$$z = 0$$
,
$$\Psi = e^{i\epsilon\phi(x)}$$

#### The weak-phase approximation

- Just below the specimen, at z=0 , the electron wave function is  $\Psi=e^{i\epsilon\phi(x)}$ .
- Then, by the approximation  $e^x \approx 1 + x$  we have just after the specimen

$$\Psi \approx 1 + i\epsilon\phi(x)$$

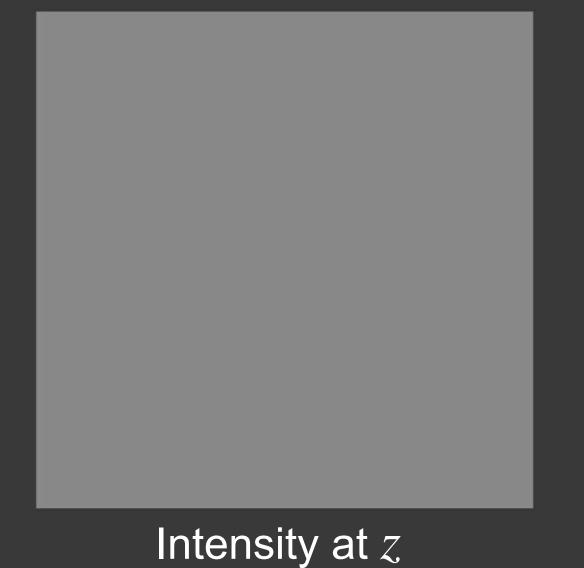
This is the weak phase approximation.

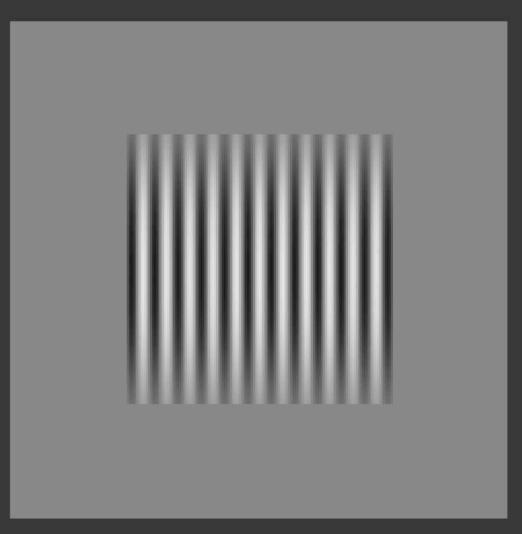
What are the two terms in the approximation?

- There is an undiffracted wave —essentially the same as the incident wave—of amplitude 1. We'll call this  $\,\Psi_0\,$
- And there is a new wave combination of amplitude  $\epsilon$ . In this example of a grating there are actually two diffracted waves,  $\Psi_+$  and  $\Psi_-$ 
  - The full wavefunction is

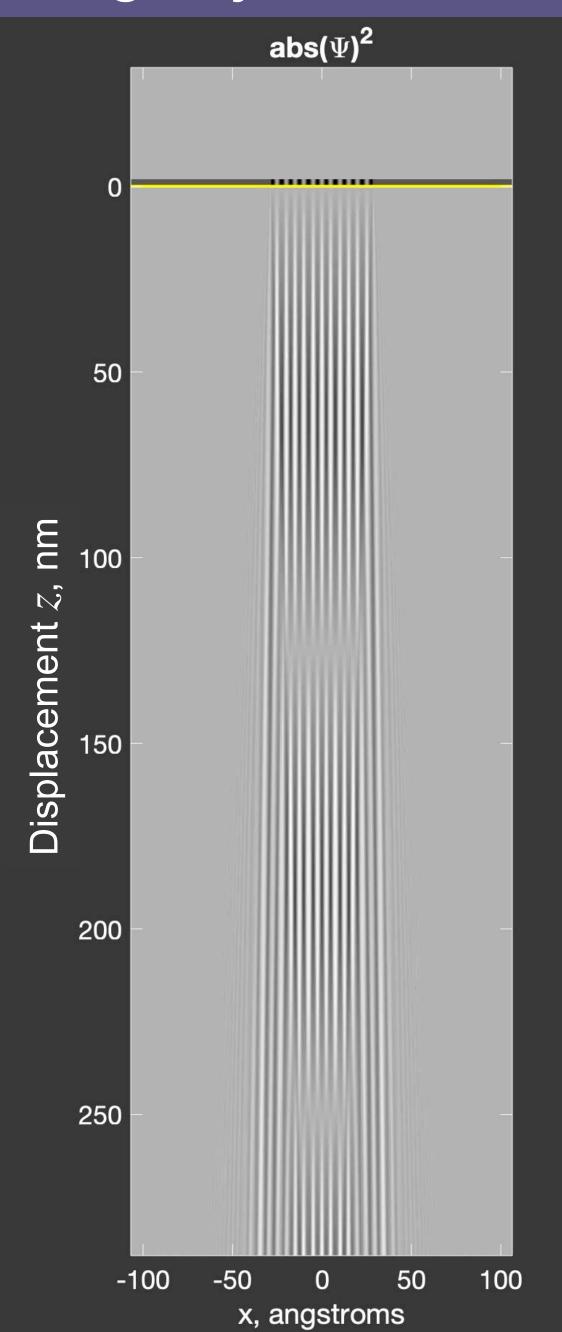
$$\Psi = \Psi_0 + \Psi_+ + \Psi_-$$

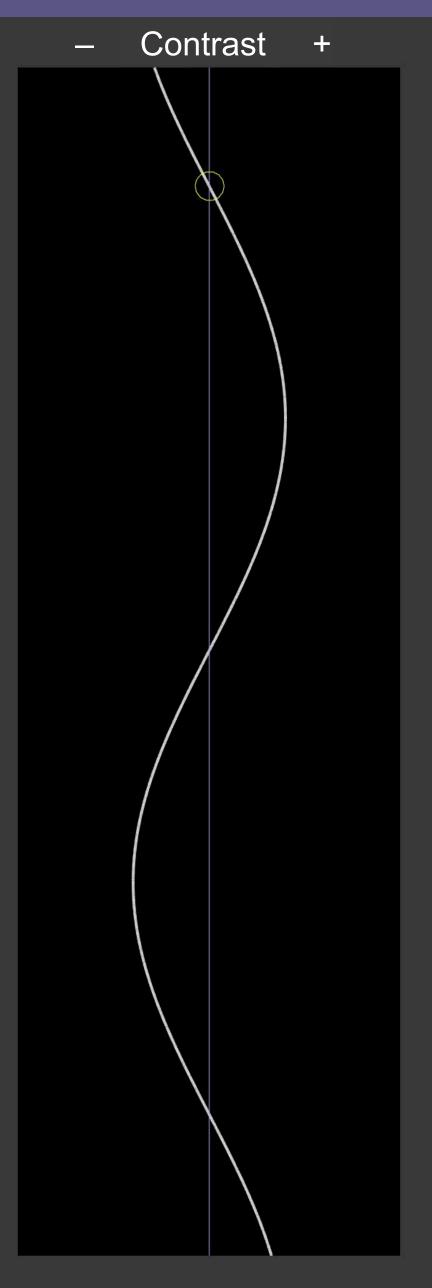
#### The contrast of a grating object varies with the distance below the object





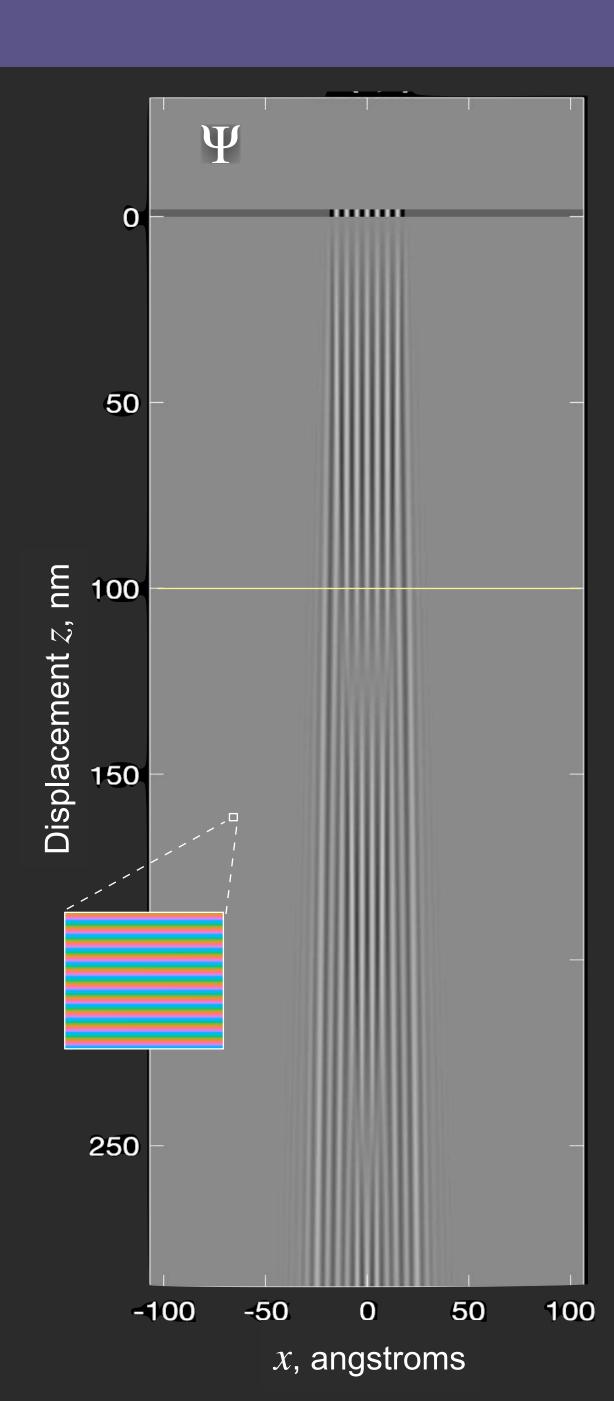
The grating  $\phi(x)$ 

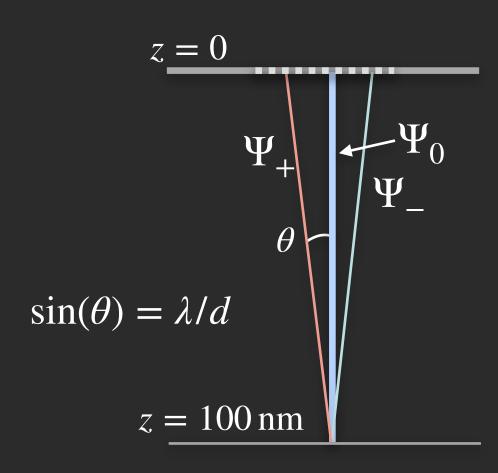




Interference between the undiffracted wave and diffracted waves produces contrast.

#### Waves interfere to make contrast





- The two diffracted waves  $\Psi_+$  and  $\Psi_-$  travel at very small angles  $+\theta$  and  $-\theta$  to the undiffracted wave.
- To reach a distance z below the specimen, they take a path longer than  $\Psi_0$  does. Let  $\zeta$  = the path length difference.

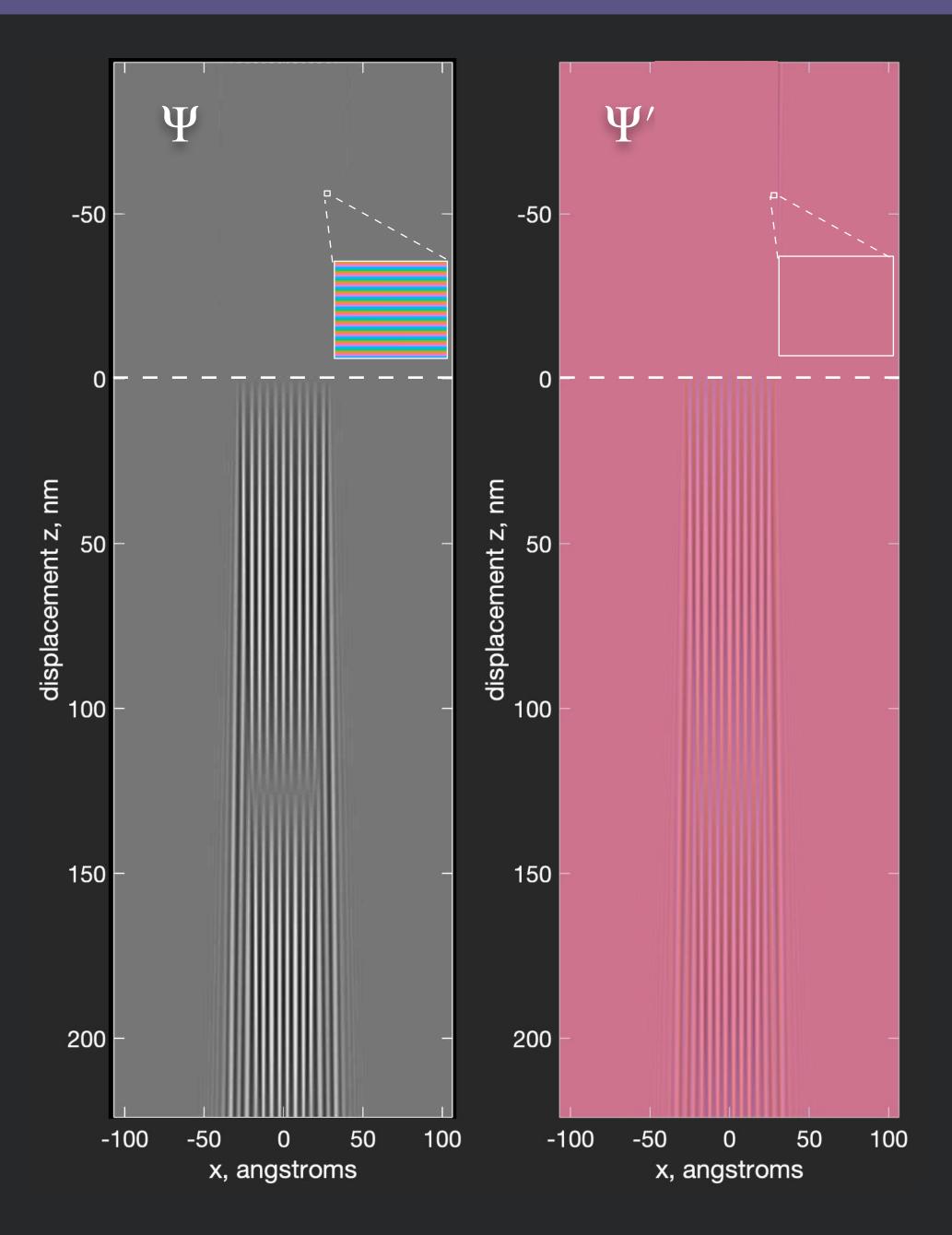
$$\zeta = \frac{z}{\cos \theta} - z \approx z \lambda^2 / 2d^2.$$

- In our example  $\lambda=.02 {\rm \AA}$  and the grating  $d=5 {\rm \AA}$ . At the level  $z=100\,{\rm nm}$ ,  $\zeta=.008 {\rm \AA}$ , about half a wavelength.
- Define  $\chi$  = the phase difference between the undiffracted and diffracted waves.

$$\chi = 2\pi \zeta / \lambda$$
$$= \pi \lambda z / d^2$$

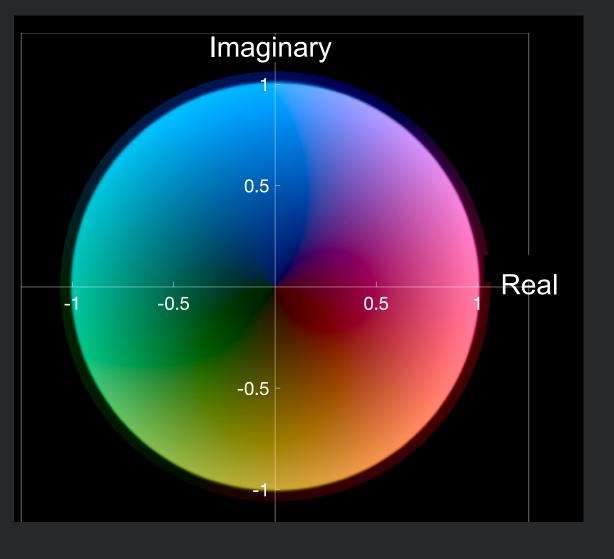
• In this example  $\chi=0.8\pi$ 

#### Where the phase of the diffracted waves is right, we have contrast.

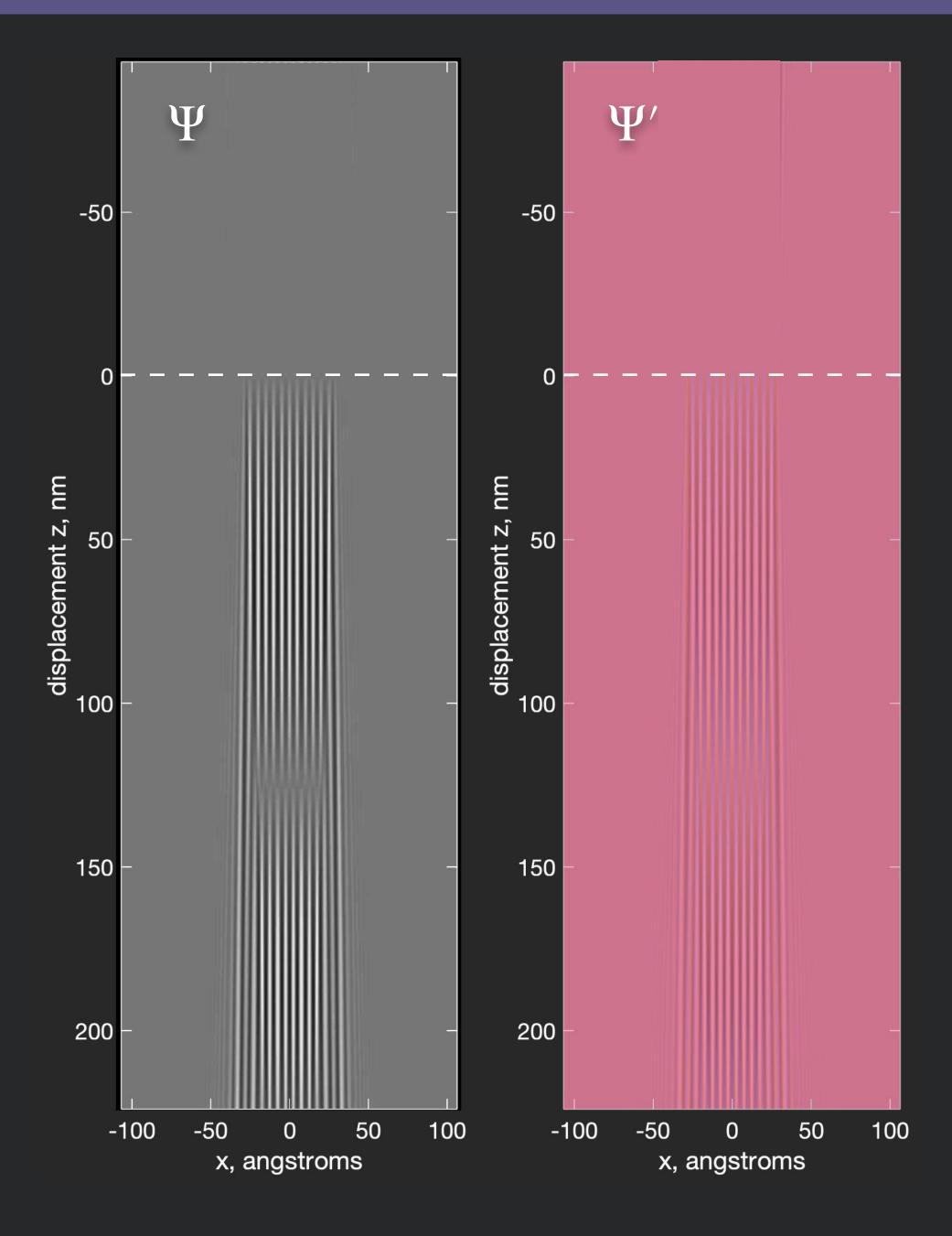


Let's unwrap the oscillations in  $\Psi$ : We'll define  $\Psi' = \Psi/\Psi_0$ 





#### Where the phase of the diffracted waves is right, we have contrast.

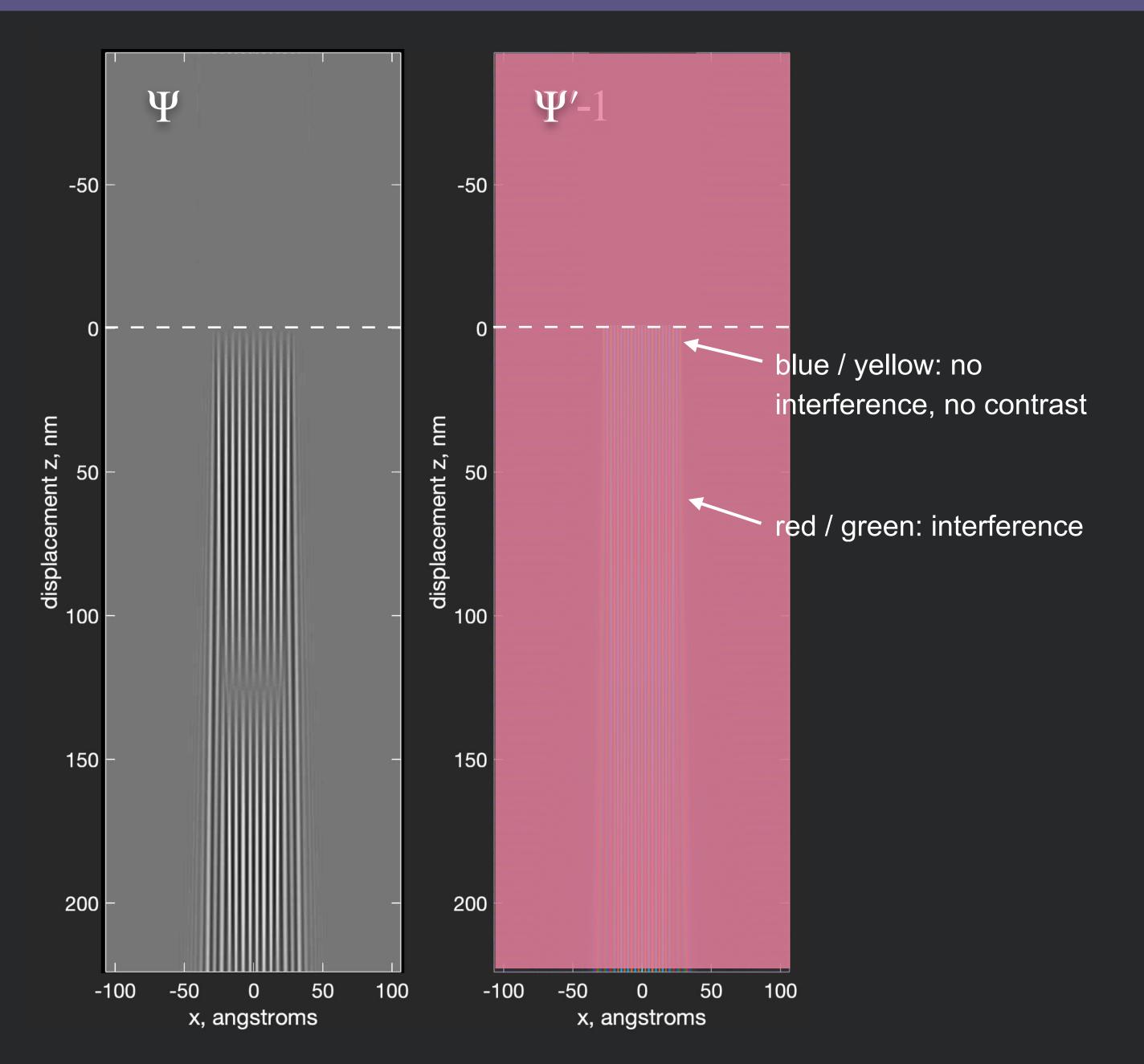


Let's unwrap the oscillations in  $\Psi$ : We'll define  $\Psi' = \Psi/\Psi_0$ 

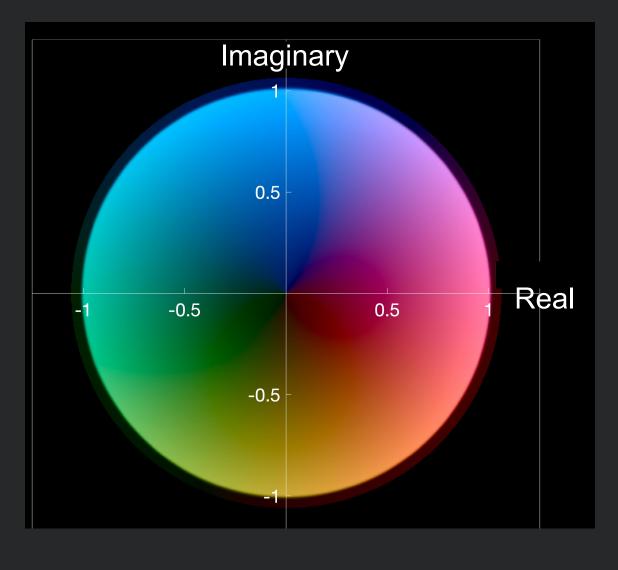
Let's remove the undiffracted wave, so we have just the diffracted waves,

$$(\Psi' - 1) = \Psi_+ + \Psi_-$$

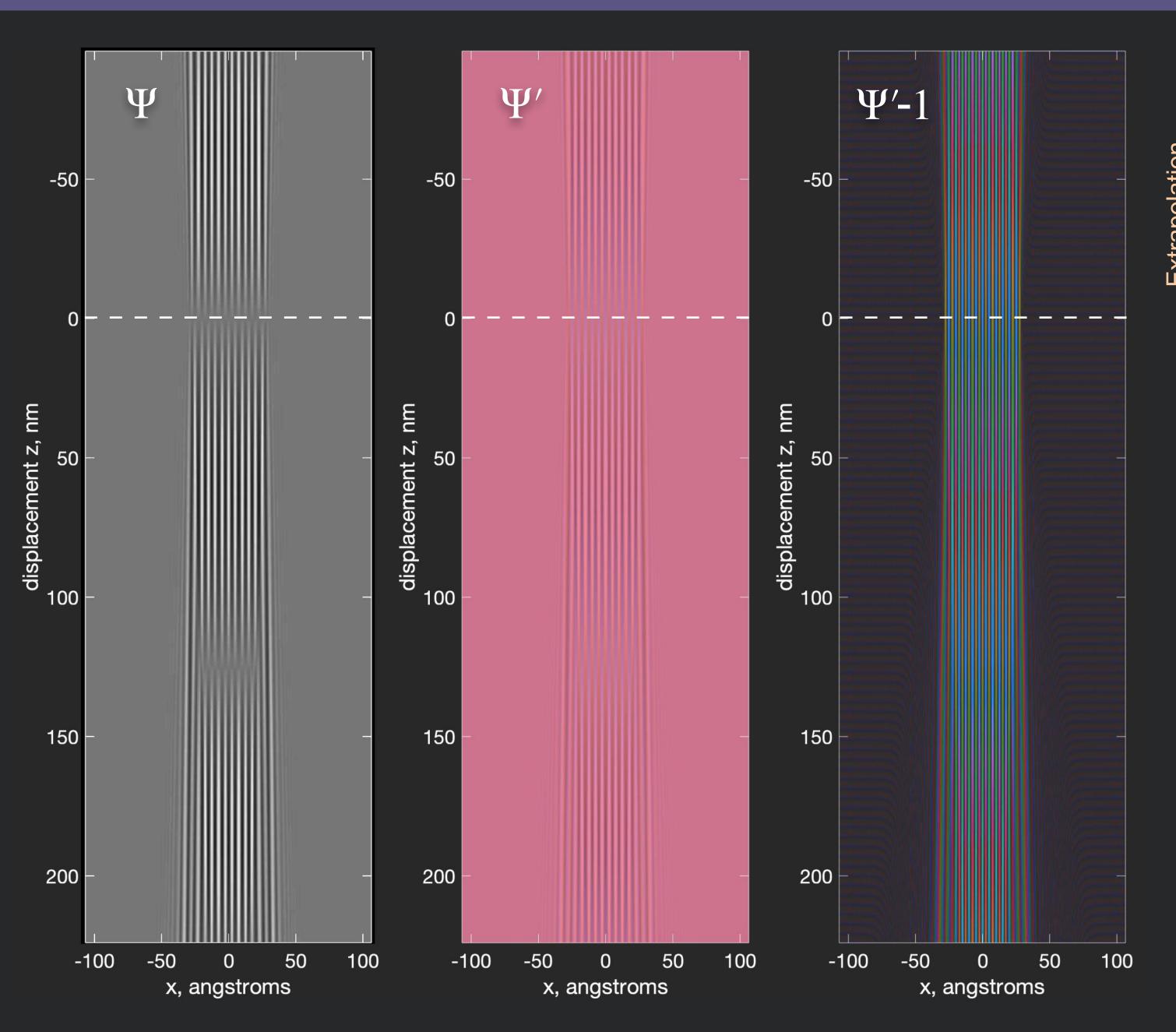
#### When the phase of the diffracted waves is right, we have contrast.



#### Complex number color scheme



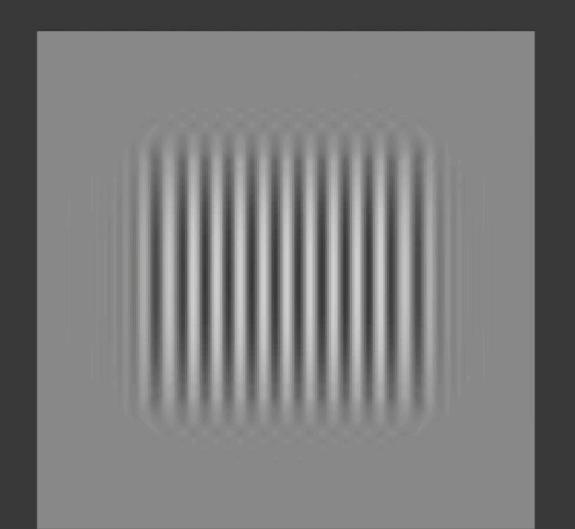
#### What happens when the objective lens is focused above the specimen?



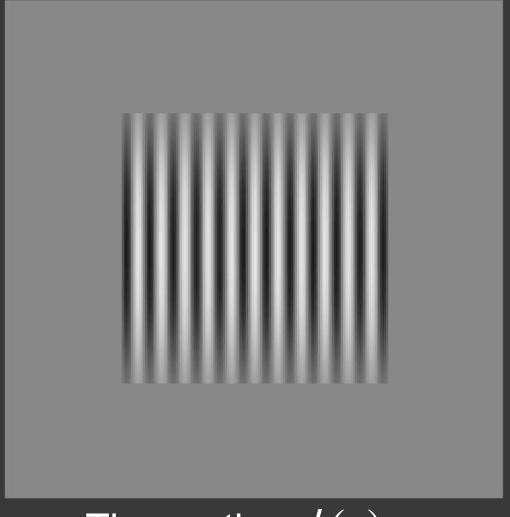
What wavefunction above the specimen would give rise to what we see below it?

We can back-propagate  $\Psi\mbox{:}$  this is what the objective lens "sees"

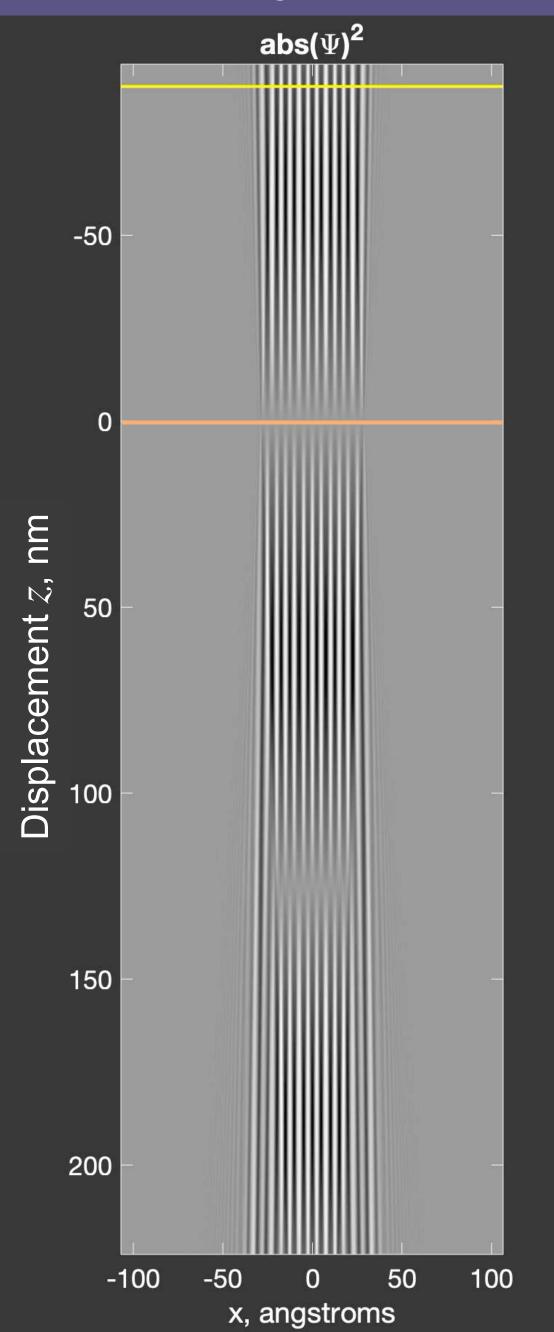
#### What happens when the objective lens is focused above the specimen?

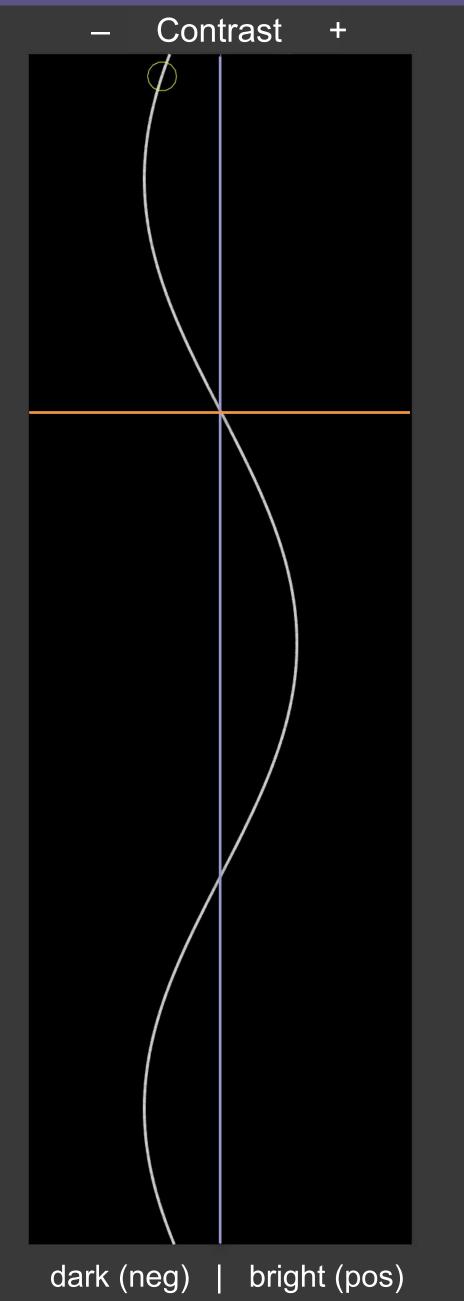


Intensity at z



The grating  $\phi(x)$ 





#### <u>Terminology</u>

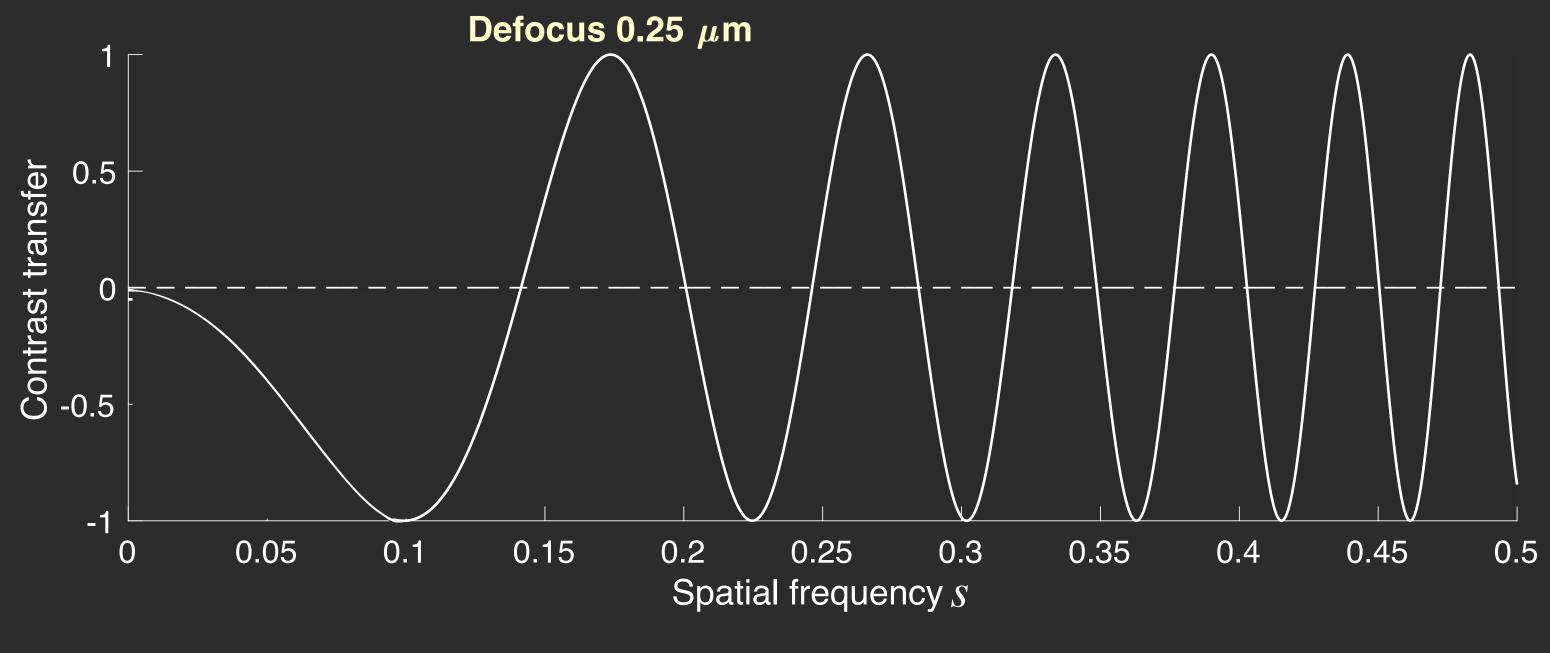
- "Underfocus" is focusing the objective lens above the specimen.
- $\bullet$  By convention, defocus values  $\delta$  are positive for underfocus:

$$\delta = -z$$

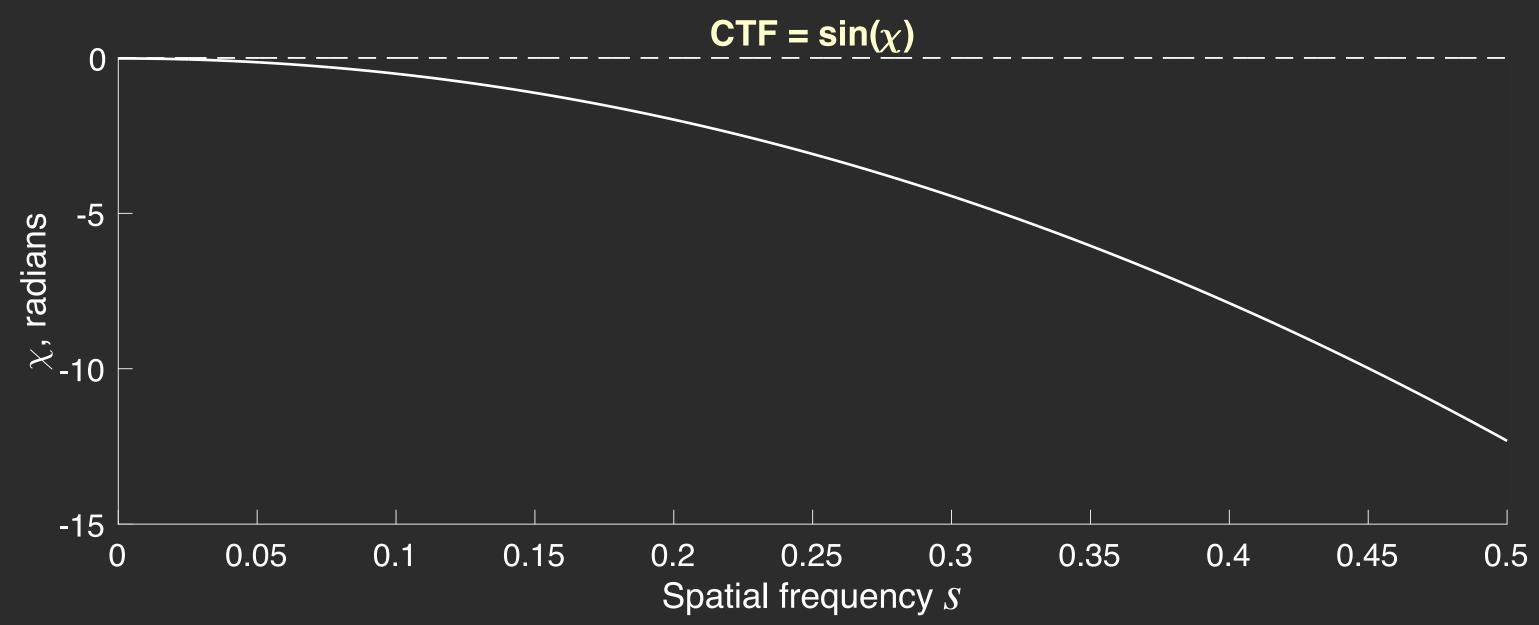
- Spatial frequency is s = 1/d
- The phase shift  $\chi$  is proportional to  $\delta$ .
- The contrast transfer function is given by

CTF = 
$$sin(\chi)$$
  
=  $sin(-\pi\lambda\delta s^2)$ 

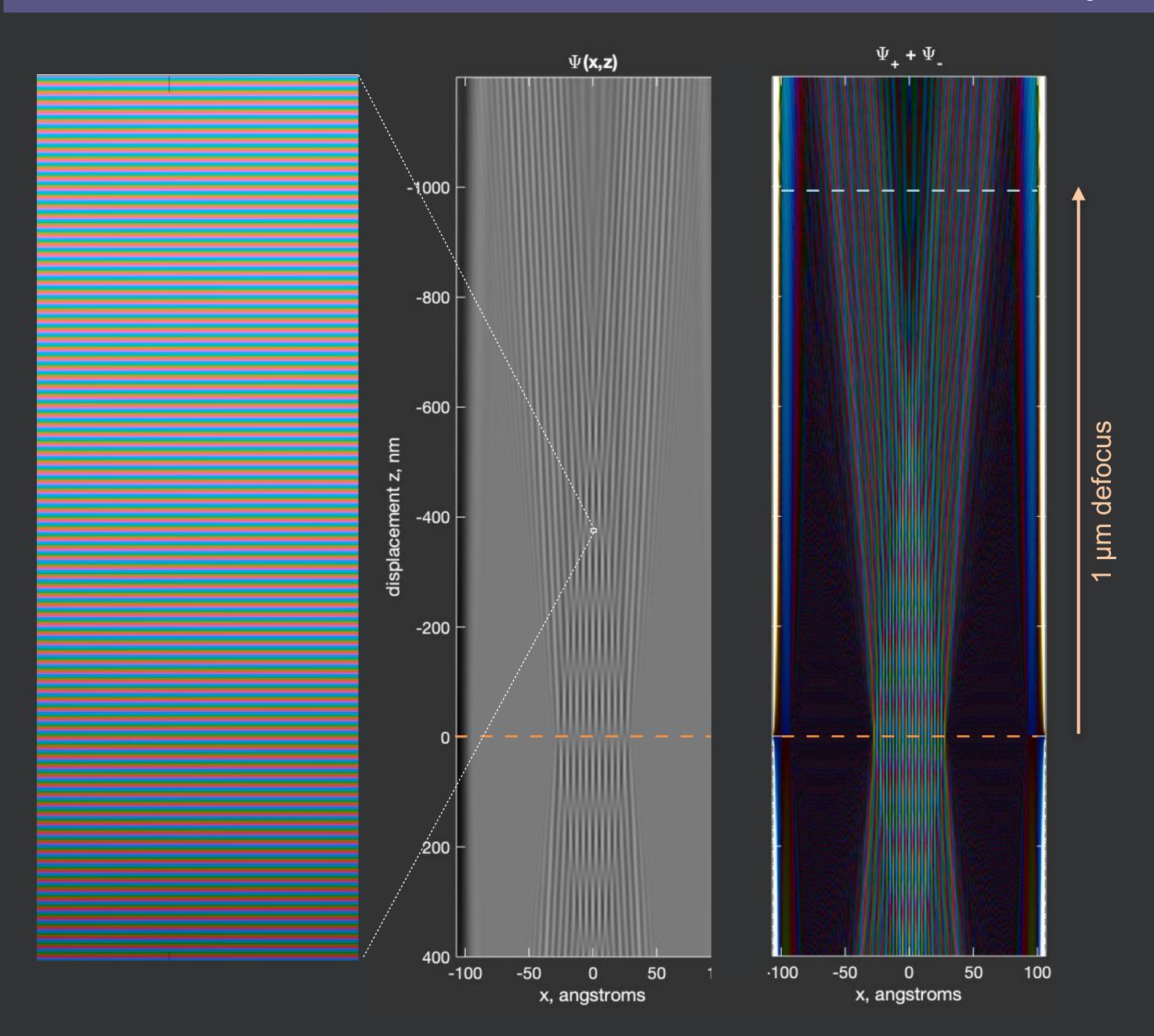
#### The basic contrast-transfer function as a function of s







#### A little defocus is actually a long distance

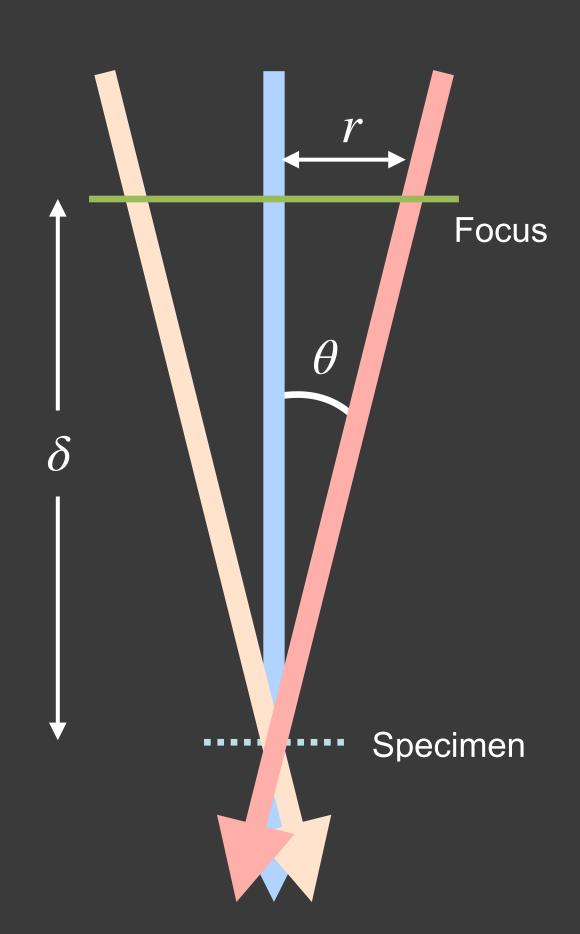


1 μm—a moderate defocus for cryo-EM imaging—is 500,000 wavelengths!

This has ramifications regarding

- beam coherence
- specimen charging
- delocalization

#### With large defocus, how bad is the image delocalization?



The dispersion radius is given by

$$r = \delta \tan \theta$$
  
=  $\delta \lambda s$  (small angle approx\*)

For example at 3µm defocus and 3Å resolution

$$\delta = 3 \times 10^4 \text{Å}$$

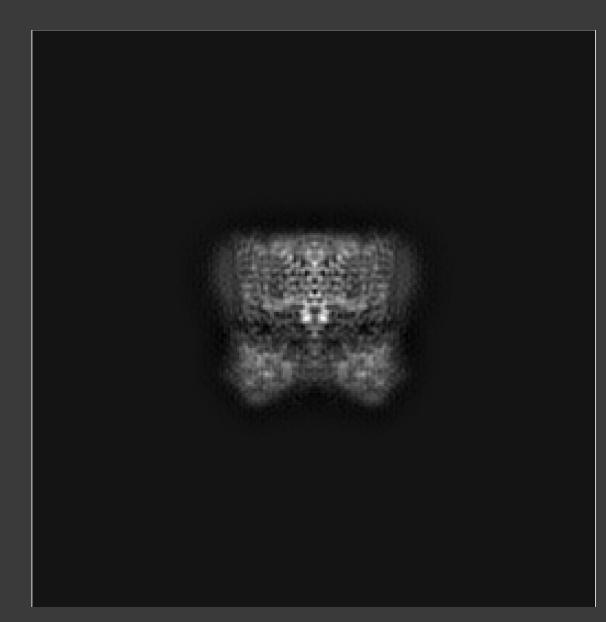
$$\lambda = .02 \mathring{A}$$

$$s = 0.33 \text{Å}^{-1}$$

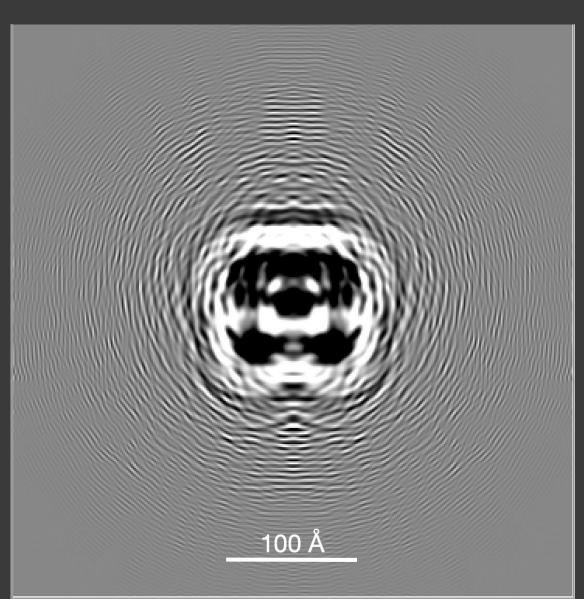
then

$$r = 200 \text{Å}$$

In this case one would want 200Å of space in the box around each particle image.



Object



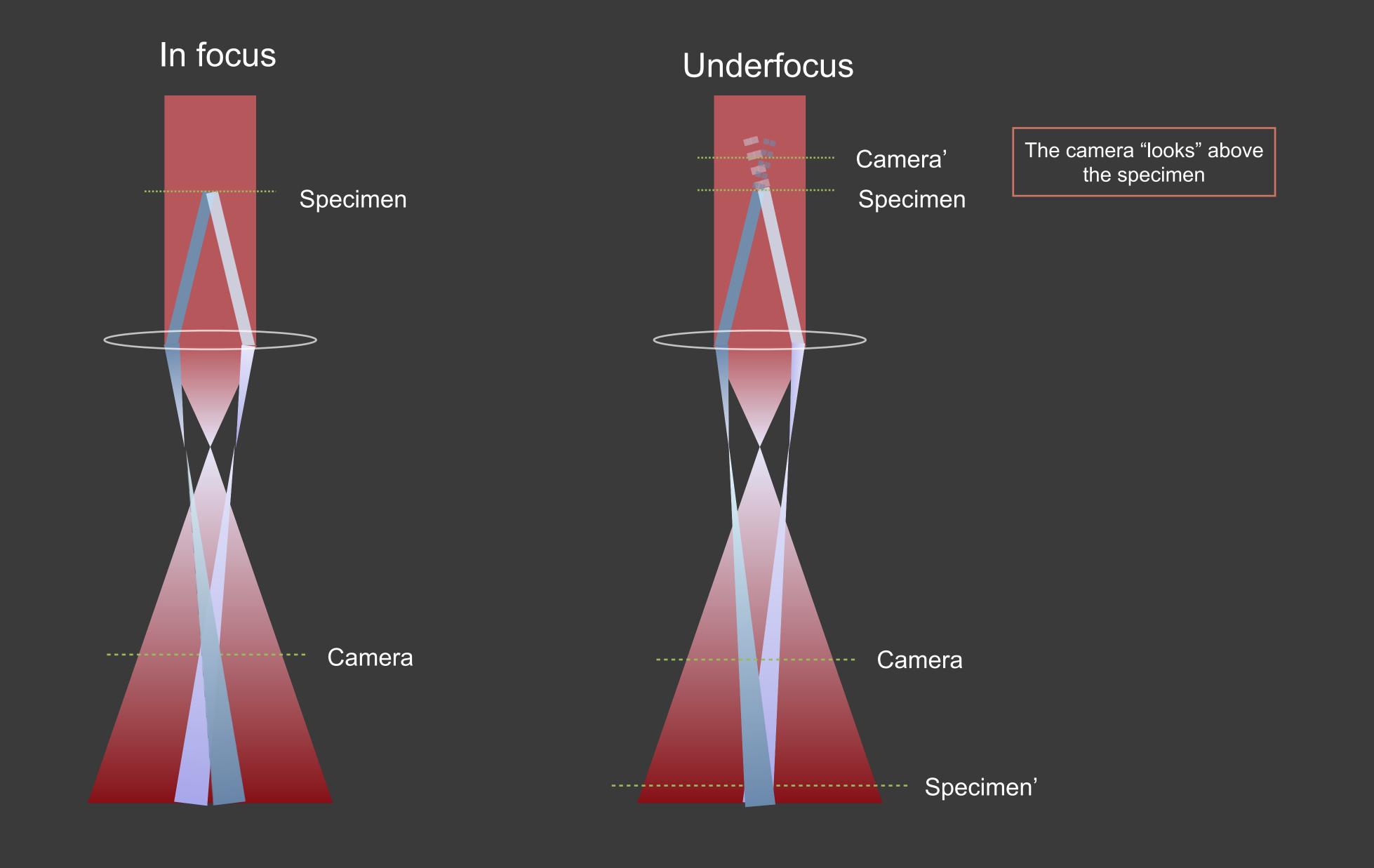
3 µm defocus

<sup>\*</sup>Note: beyond about 3Å, spherical aberration needs to be taken into account too.

#### Contrast Transfer Function, Part 2: Advanced Topics

- Lens aberrations and the image plane
- Why use underfocus?
- The diffraction plane and phase plate

#### An objective lens reproduces interference patterns at the camera



#### With spherical aberration a lens bends high-angle rays more strongly

Spherical aberration changes the defocus by

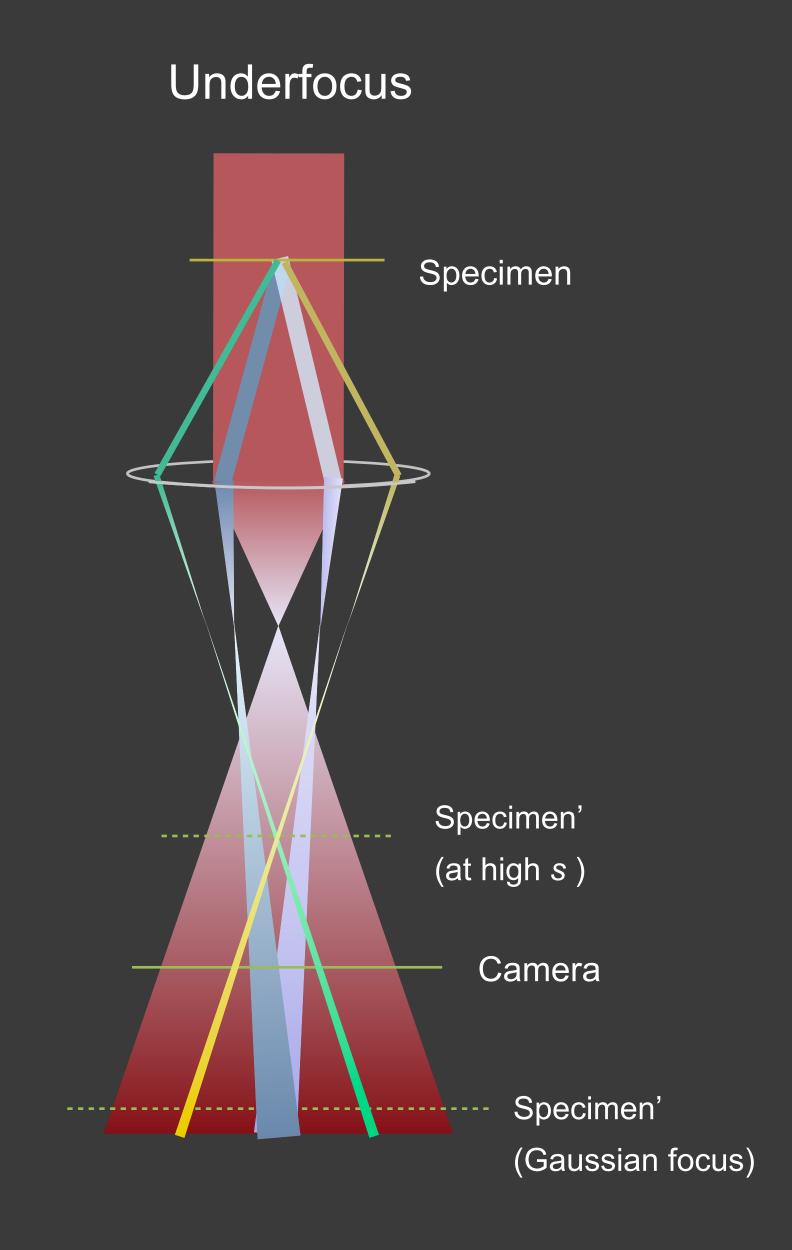
$$\delta' = -C_s \lambda^2 s^2 / 2.$$

The contrast transfer function now includes  $\delta'$ ,

CTF = 
$$\sin(-\pi\lambda (\delta + \delta') s^2)$$
 or, expanded,

$$CTF = \sin(-\pi\lambda\delta s^2 + \frac{\pi}{2}C_s\lambda^3 s^4)$$

The coefficient  $C_s$  is typically ~2mm. Spherical aberration typically becomes important for  $s \gtrsim 0.25 \text{Å}^{-1}$ , or about 4 Å resolution.



#### Very high-angle scattering yields amplitude contrast

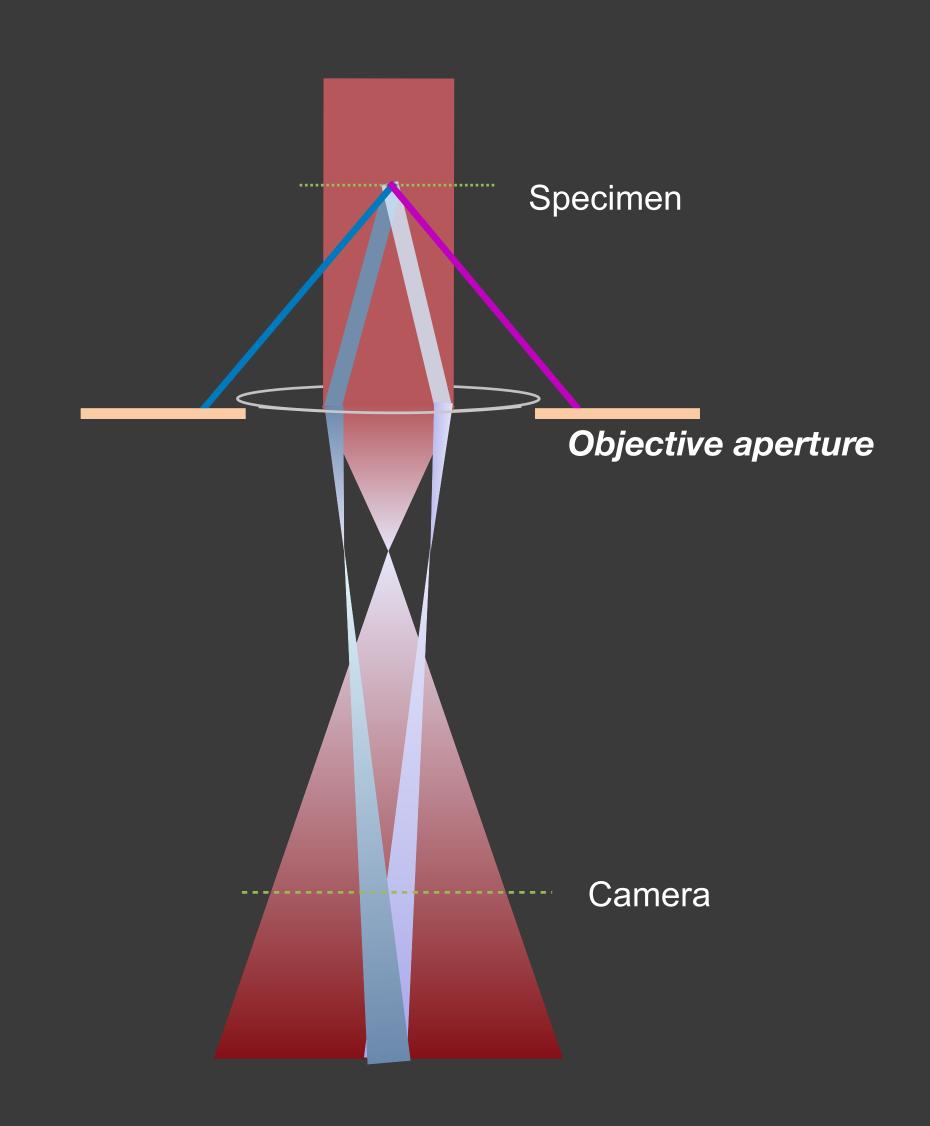
Electrons that pass very close to an atomic nucleus are scattered at high angles, and are caught by the objective aperture.

- The loss of these electrons results in a small amount of negative amplitude contrast.
- For proteins  $\alpha$  is typically around -0.05.
- The amplitude contrast term allows the CTF to have a small negative value even at zero spatial frequency.

Combining all these terms, the contrast transfer function is given by

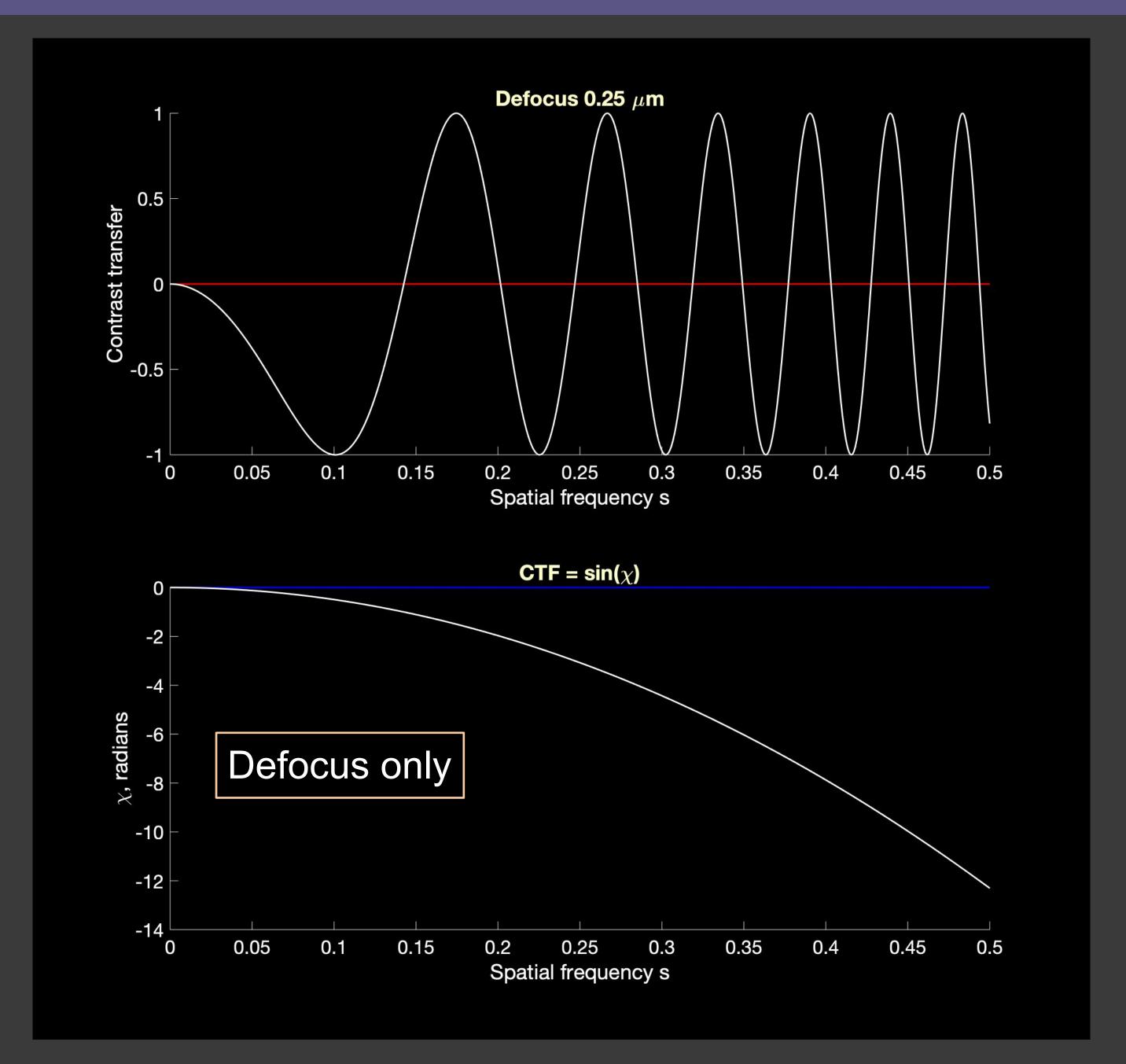
$$\text{CTF} = \sin(-\pi\lambda \delta s^2 + \frac{\pi}{2} C_s \lambda^3 s^4 - \alpha)$$

$$\text{defocus sphere abb. amplitude}$$



#### The simple defocus contrast is what we've seen before

CTF = 
$$\sin(-\pi\lambda\delta f^2 + \frac{\pi}{2}C_s\lambda^3 f^4 - \alpha)$$
defocus sphere abb. amplitude



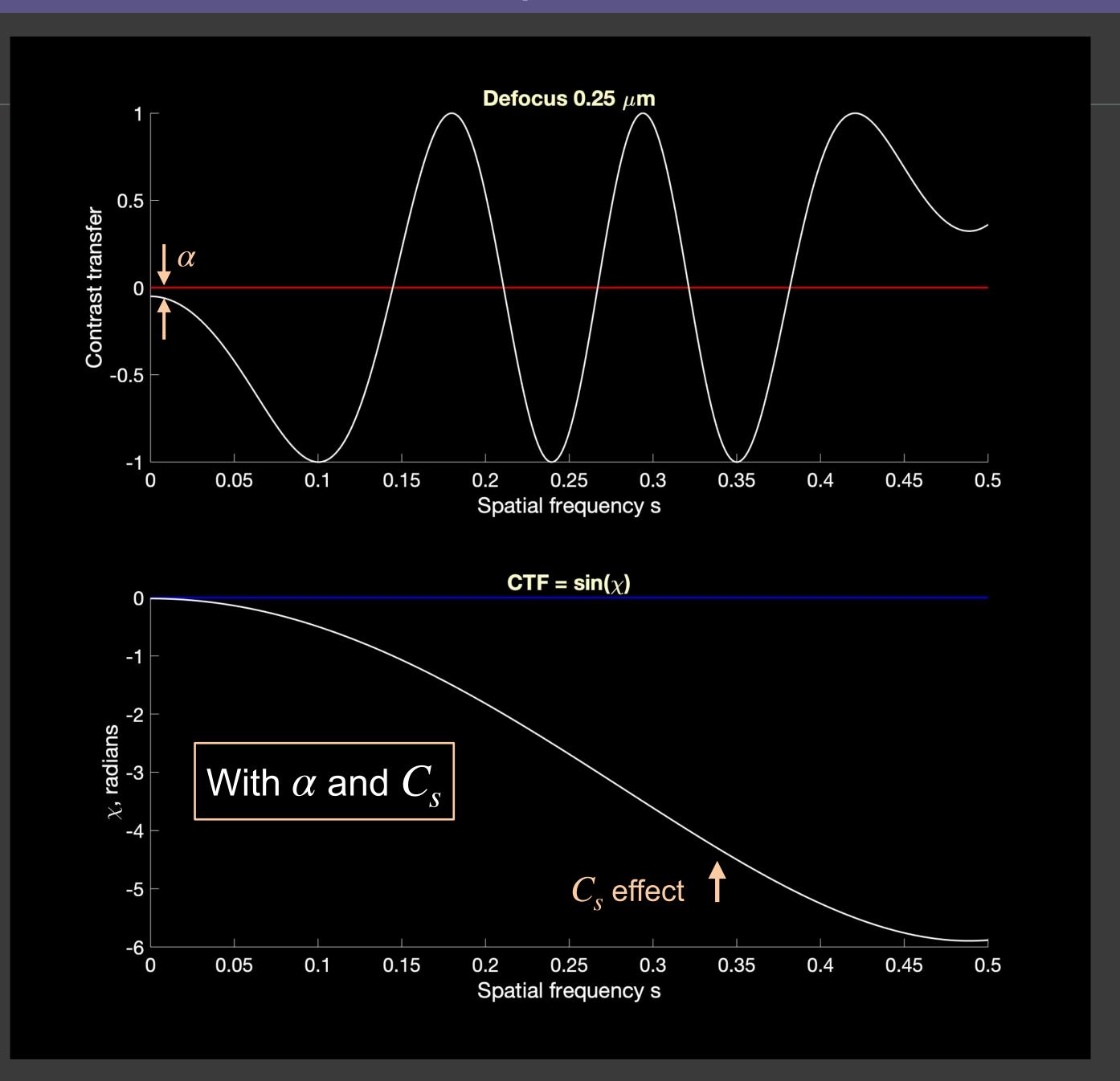
#### Now adding in spherical aberration and amplitude contrast

Here you can see why everyone uses underfocus: the amplitude contrast and defocus contrast are additive in this case.

Also, Cs has the effect of reversing some of the oscillations in the CTF.

Combining all these terms, the contrast transfer function is given by

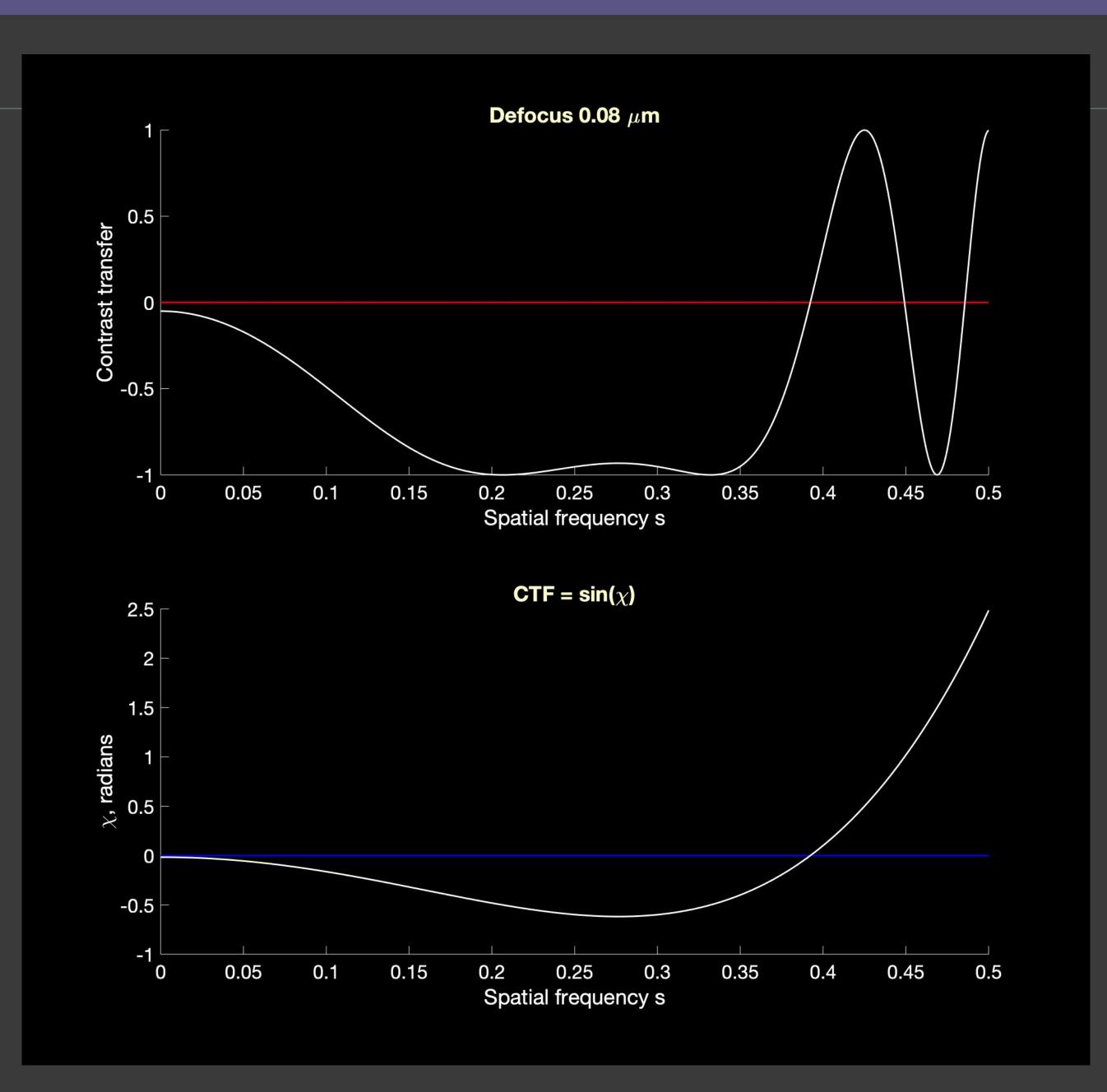
$$\text{CTF} = \sin(-\pi\lambda\delta f^2 + \frac{\pi}{2}C_s\lambda^3 f^4 - \alpha)$$
 defocus sphere abb. amplitude



#### Spherical aberration can be our friend

If we're not using image processing to remove CTF effects, Scherzer defocus is a good solution: just enough defocus to give signal over a broad range of spatial frequencies.

It's popular in materials science but not much for cryoEM: the signal transfer at low frequencies is poor.



- 1.Complex numbers: review
- 2. Defocus contrast (the simple version)
- 3. Contrast at the camera plane
- 4. Defocus contrast (formal version)
- 5.Phase plate

### Formal derivation of the CTF for a grating of spacing d

The non-oscillating wavefunction

$$\Psi' = 1 + ie^{-ik\zeta} \cdot \epsilon \cos(2\pi x/d)$$

can be written as

$$\Psi' = 1 + ie^{-i\chi} \epsilon \phi(x).$$

The measured intensity is

$$|\Psi|^2 = |\Psi'|^2 = (\text{real part})^2 + (\text{imag part})^2$$

$$= \left[1 + \sin(\chi) \epsilon \phi(x)\right]^2 + \left[\cos(-\chi) \epsilon \phi(x)\right]^2$$

$$= \left[1 + 2\sin(\chi) \epsilon \phi(x)\right] + \mathcal{O}\epsilon^2 + \left[\mathcal{O}\epsilon^2\right].$$

#### In practice

- We ignore the constant background intensity.
- Everyone ignores the factor of 2 also.
- So we say the transfer from phase shift to intensity change is

$$CTF = \frac{\Delta Intensity}{\Delta Electron phase} = sin(\chi)$$

#### Grating object:

$$\epsilon \phi(x) = \epsilon \cos(2\pi x/d)$$

Electron propagation:

$$k = 2\pi/\lambda$$

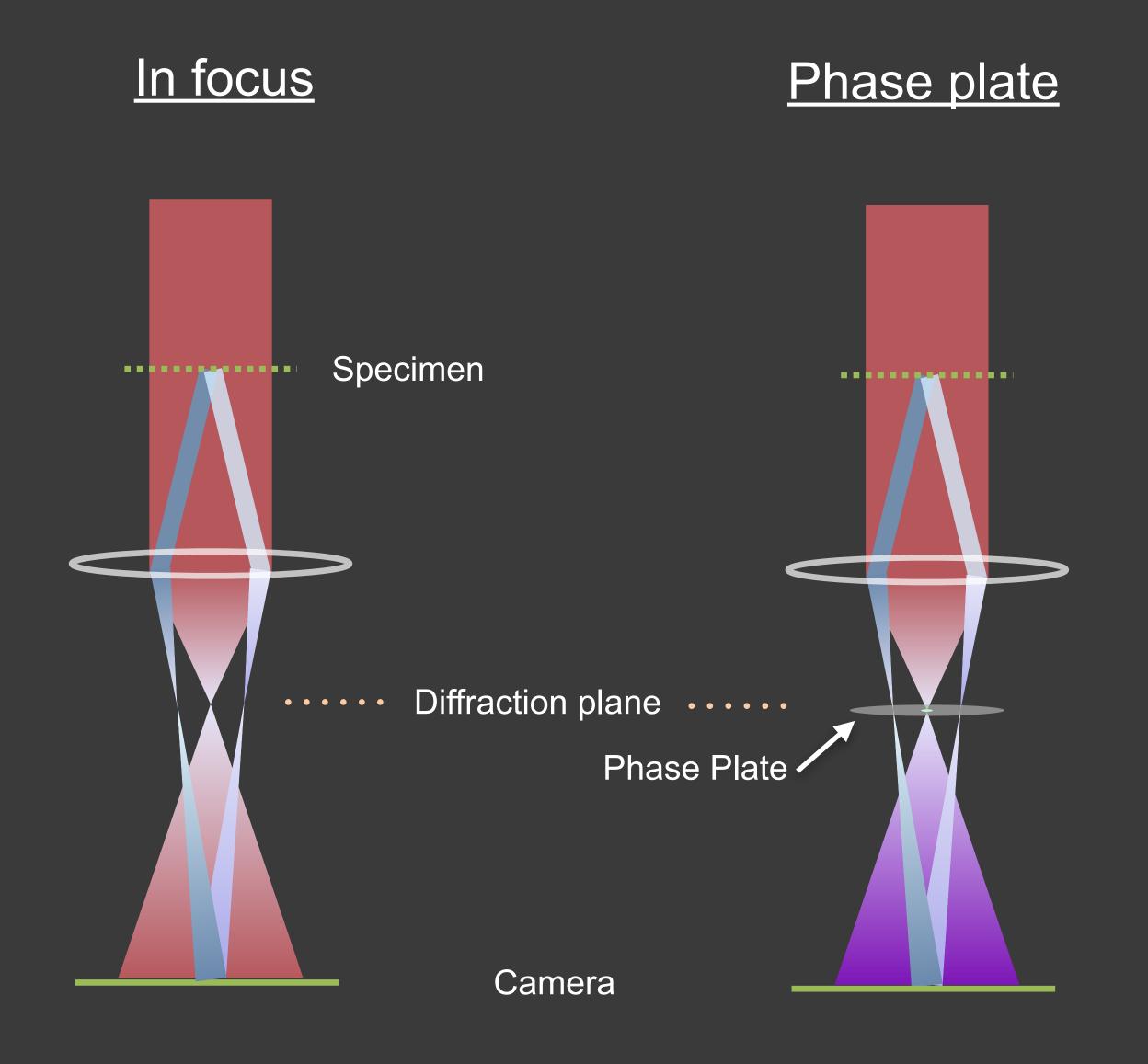
Diffracted wave path difference:

$$\zeta \approx z\lambda^2/2d^2$$

Wave aberration function:

$$\chi = k\zeta \approx \pi \lambda z / d^2$$

#### A phase plate modifies the interference of electron waves at the camera



The phase plate shifts the phase of the undiffracted beam  $\Psi_0$  by some angle  $\phi$ .

Then CTF = 
$$\sin(\chi - \phi)$$
.

If 
$$\phi = 90^{\circ}$$
 then
$$CTF = -\cos(\chi)$$

### The phase plate allows in-focus imaging, given precise focusing.

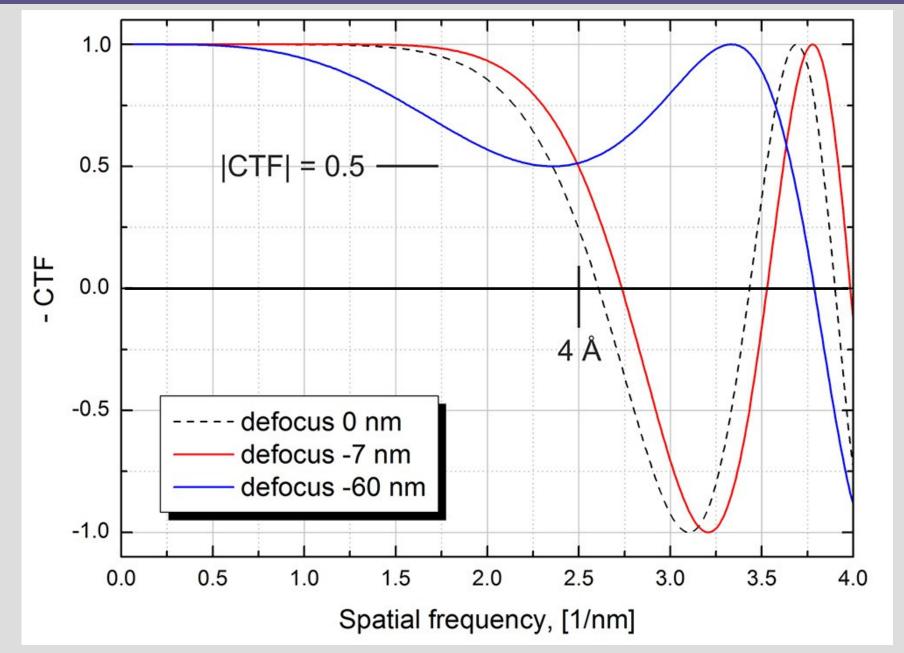
# Cryo-EM single particle analysis with the Volta phase plate

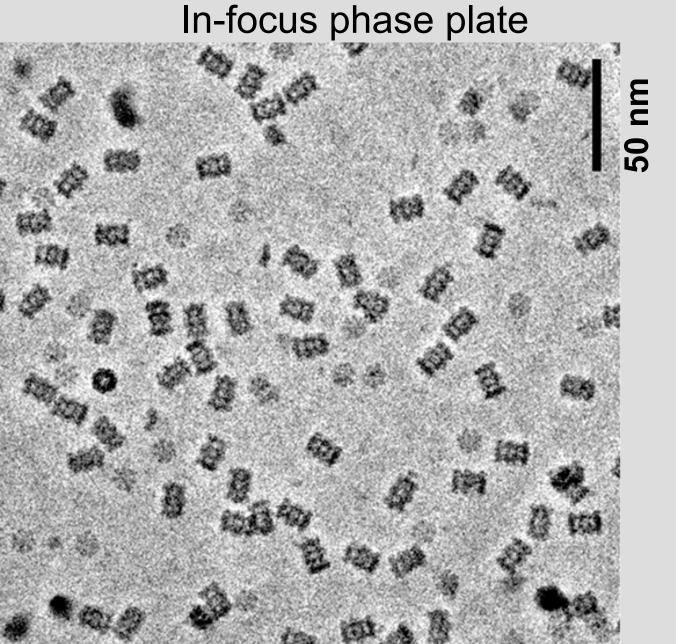
Radostin Danev\*, Wolfgang Baumeister

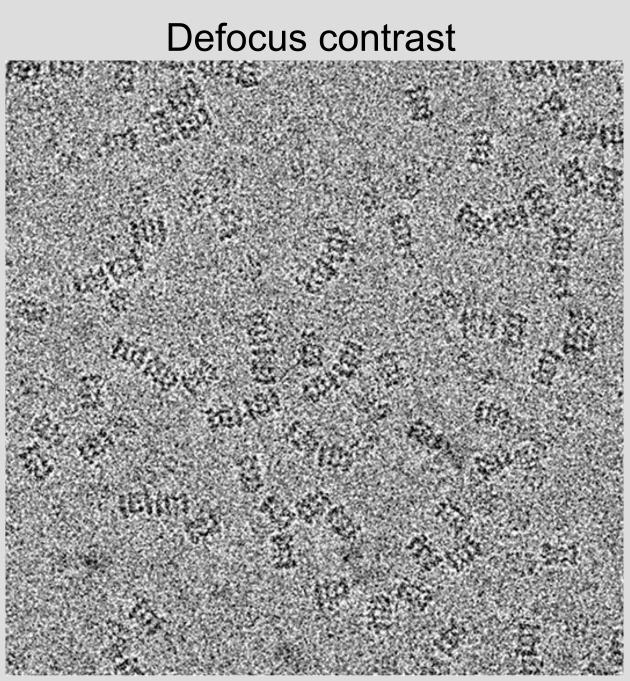
Department of Molecular Structural Biology, Max Planck Institute of Biochemistry, Martinsried, Germany

*eLife* 2016

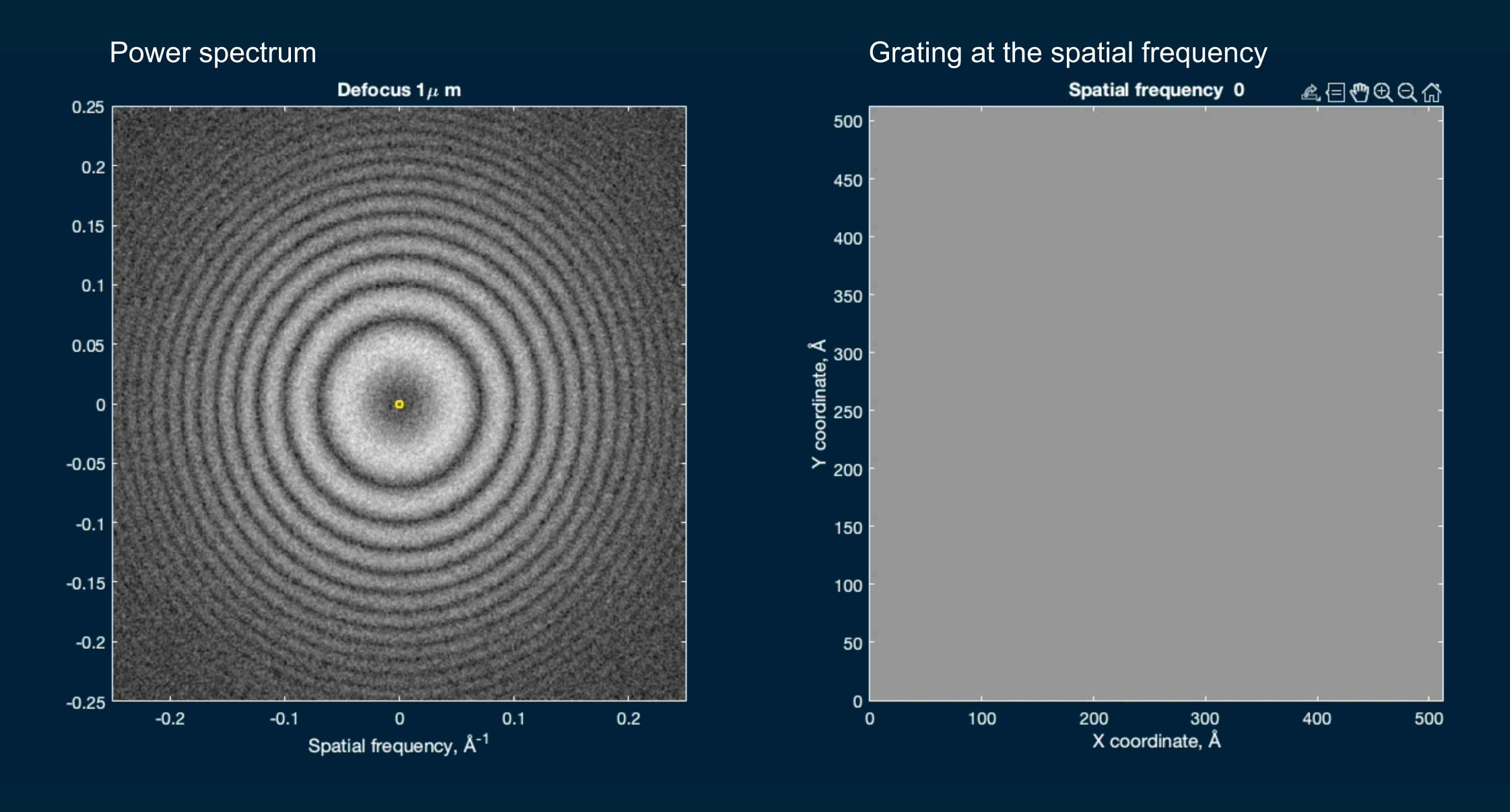
- The better low-frequency contrast makes particles much more visible.
- The defocus value must be precise within 60 nm in order to get 4 Å resolution.







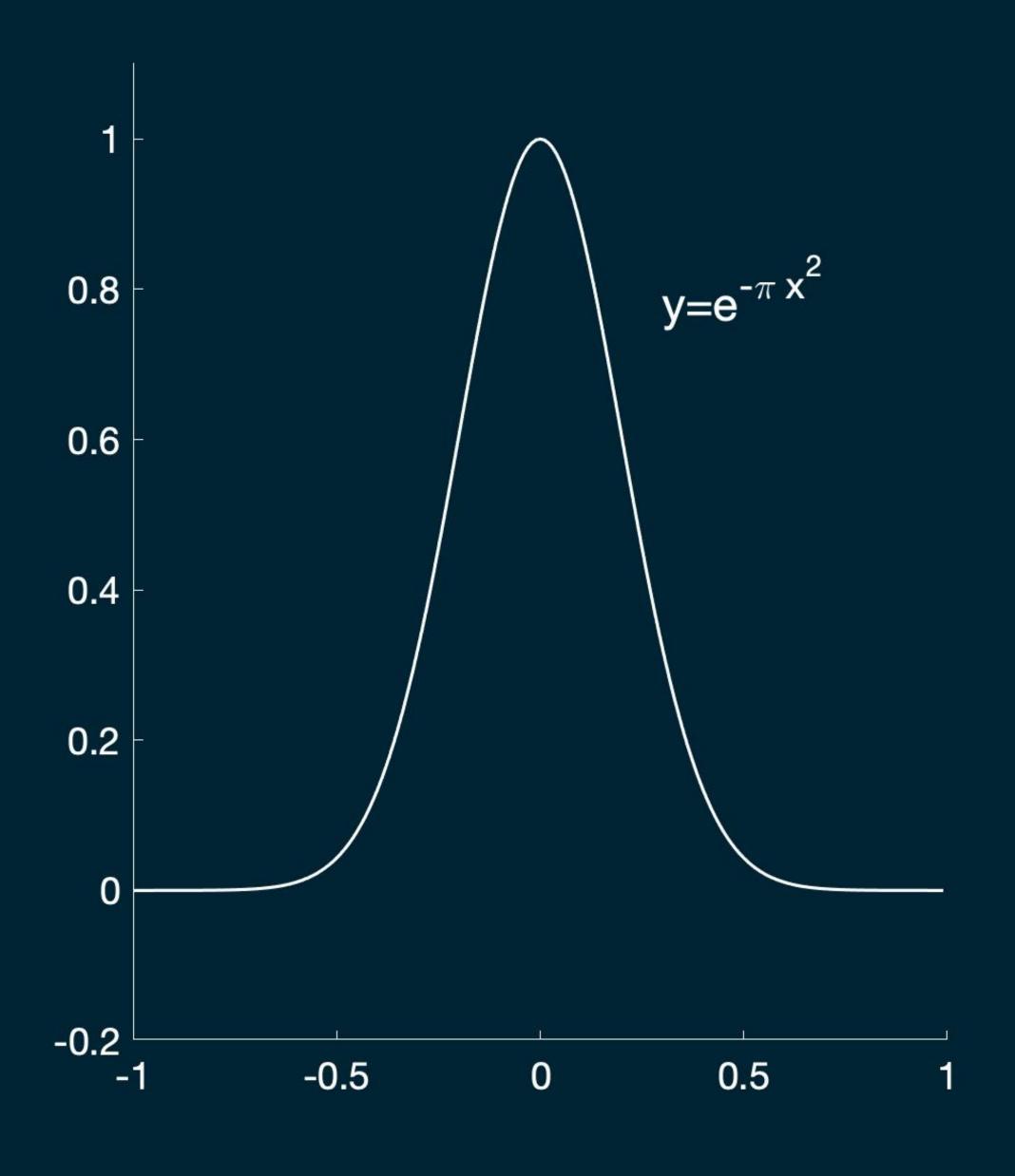
### Thon rings in the power spectrum show zeros in the CTF

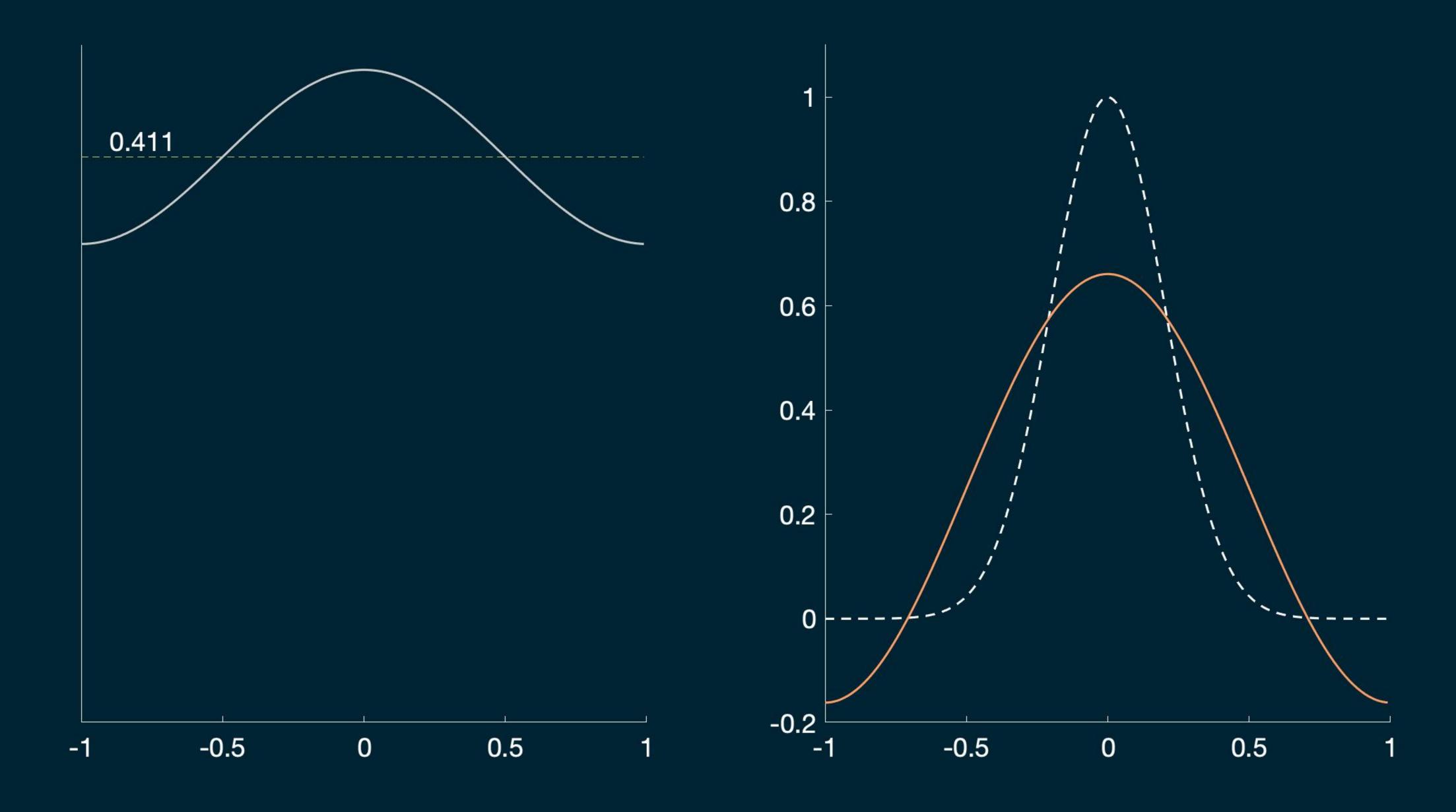


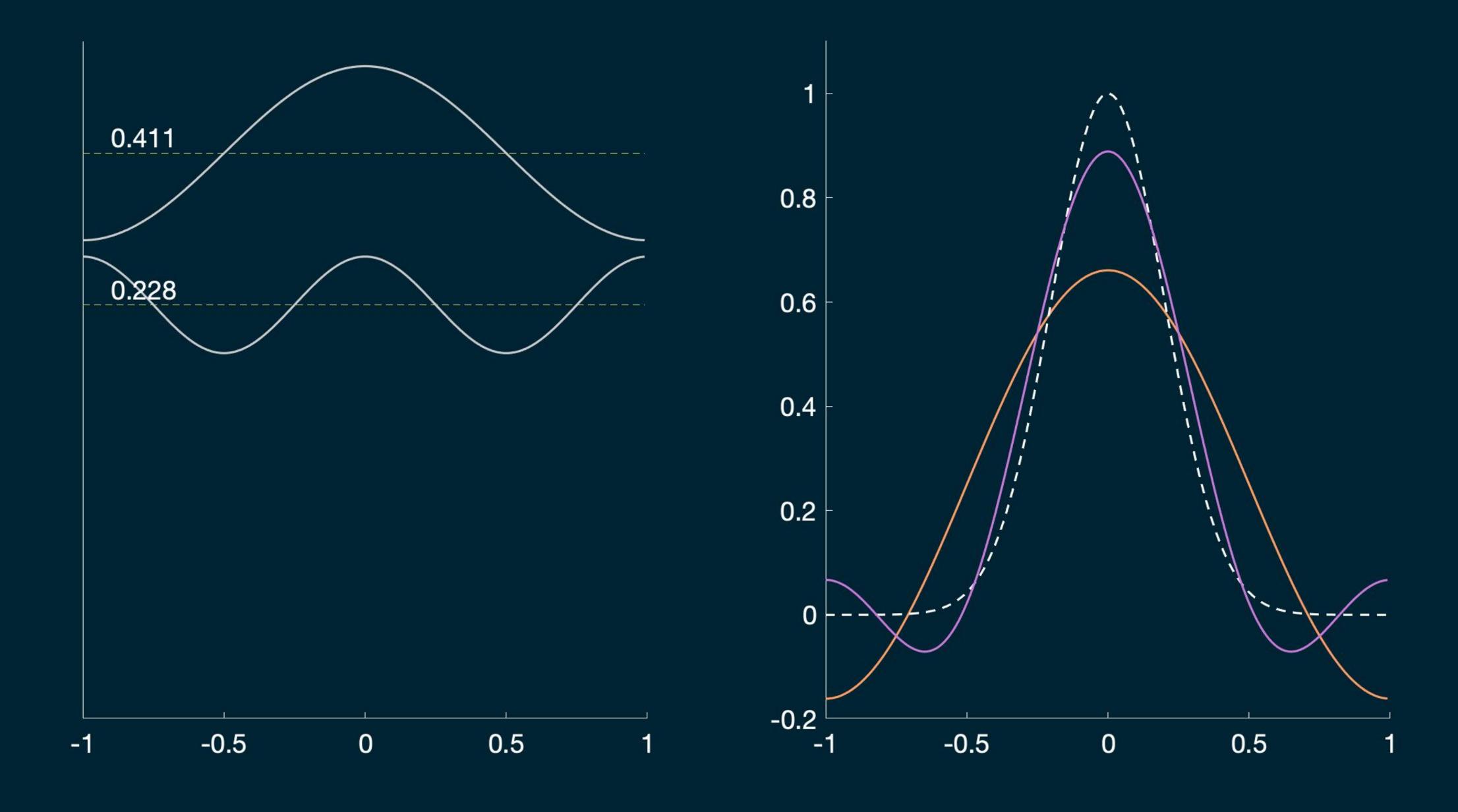
(Optional)

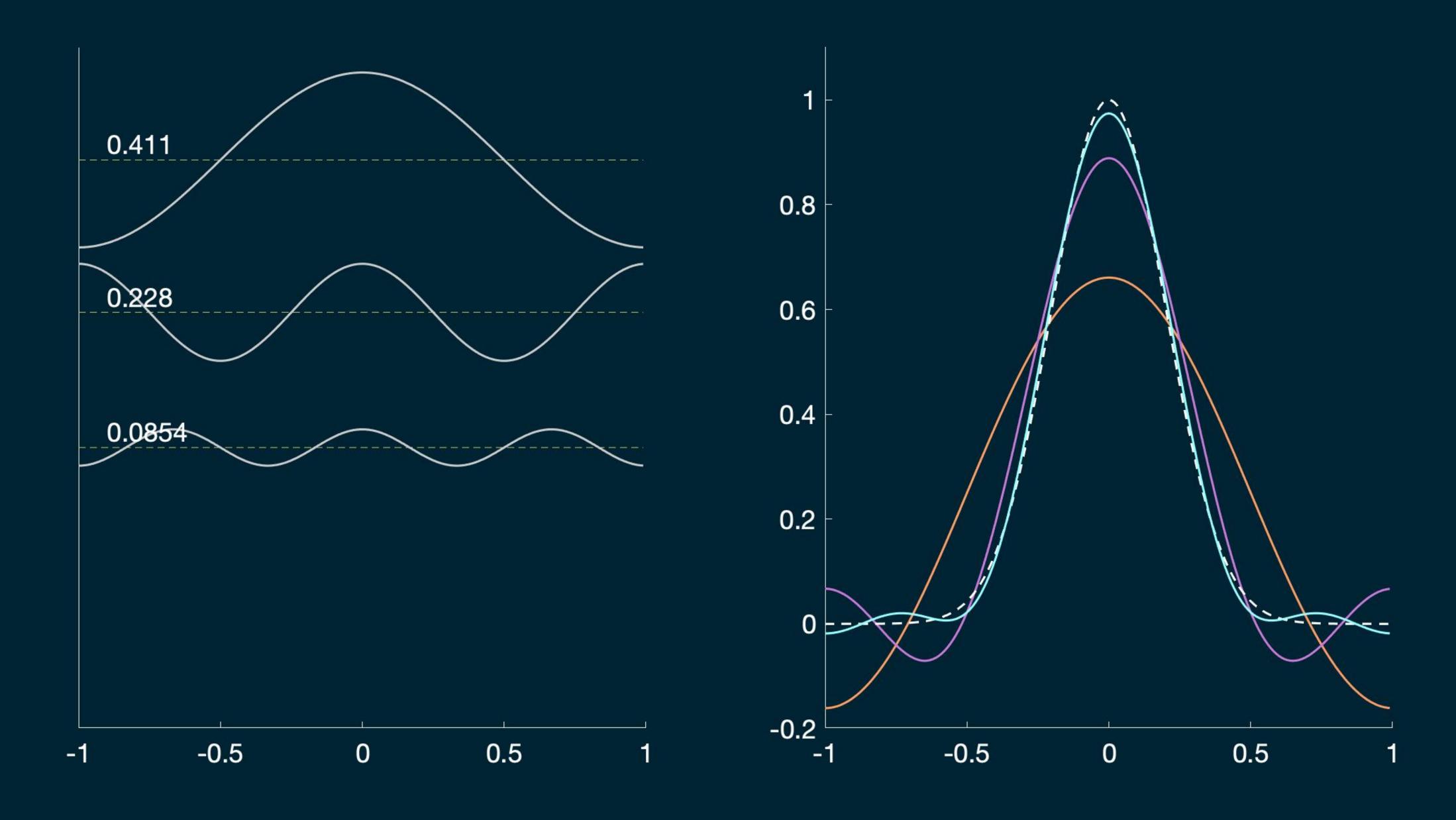
A quick introduction to Fourier transforms

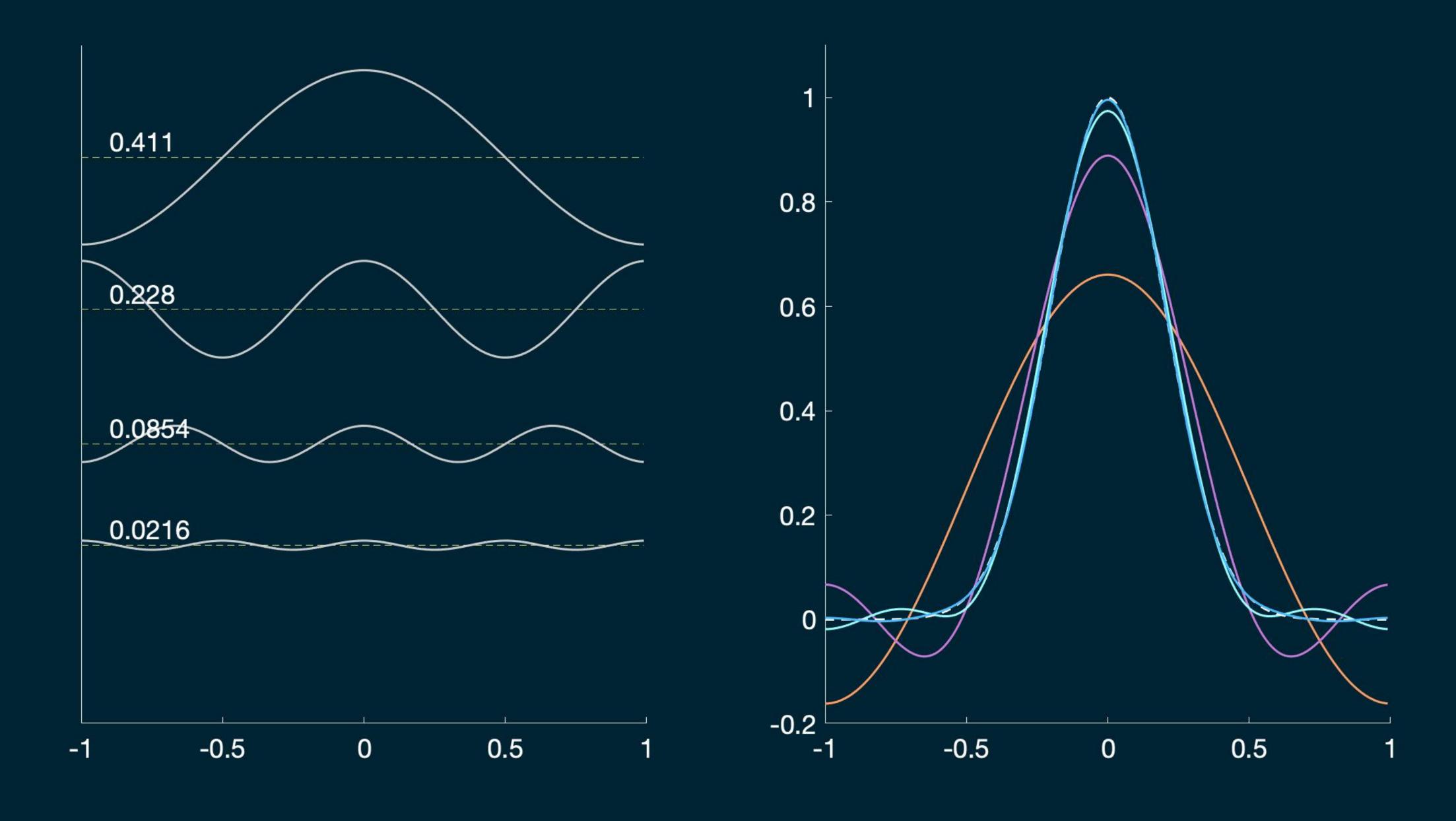
The Fourier transform in one dimension

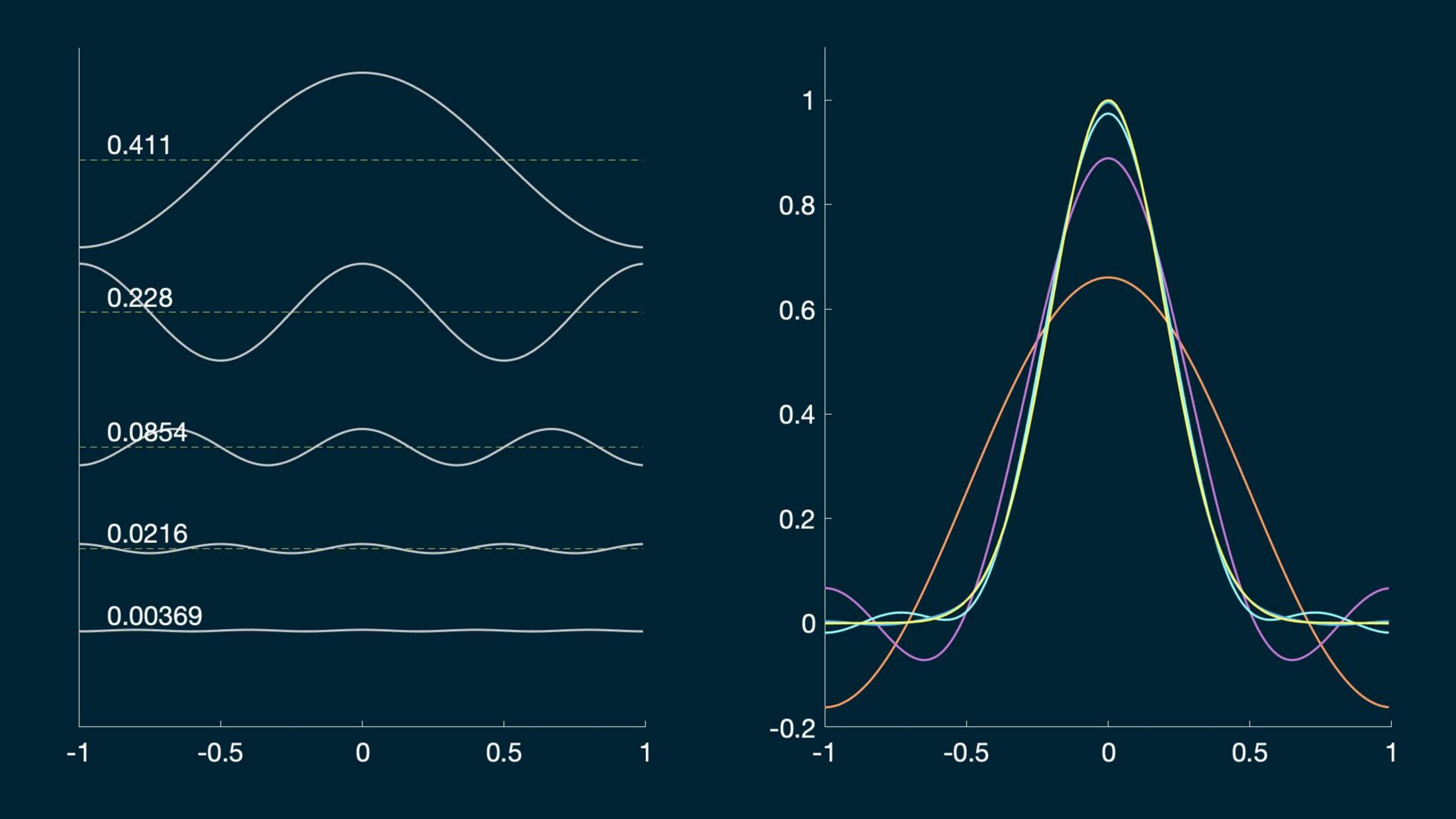




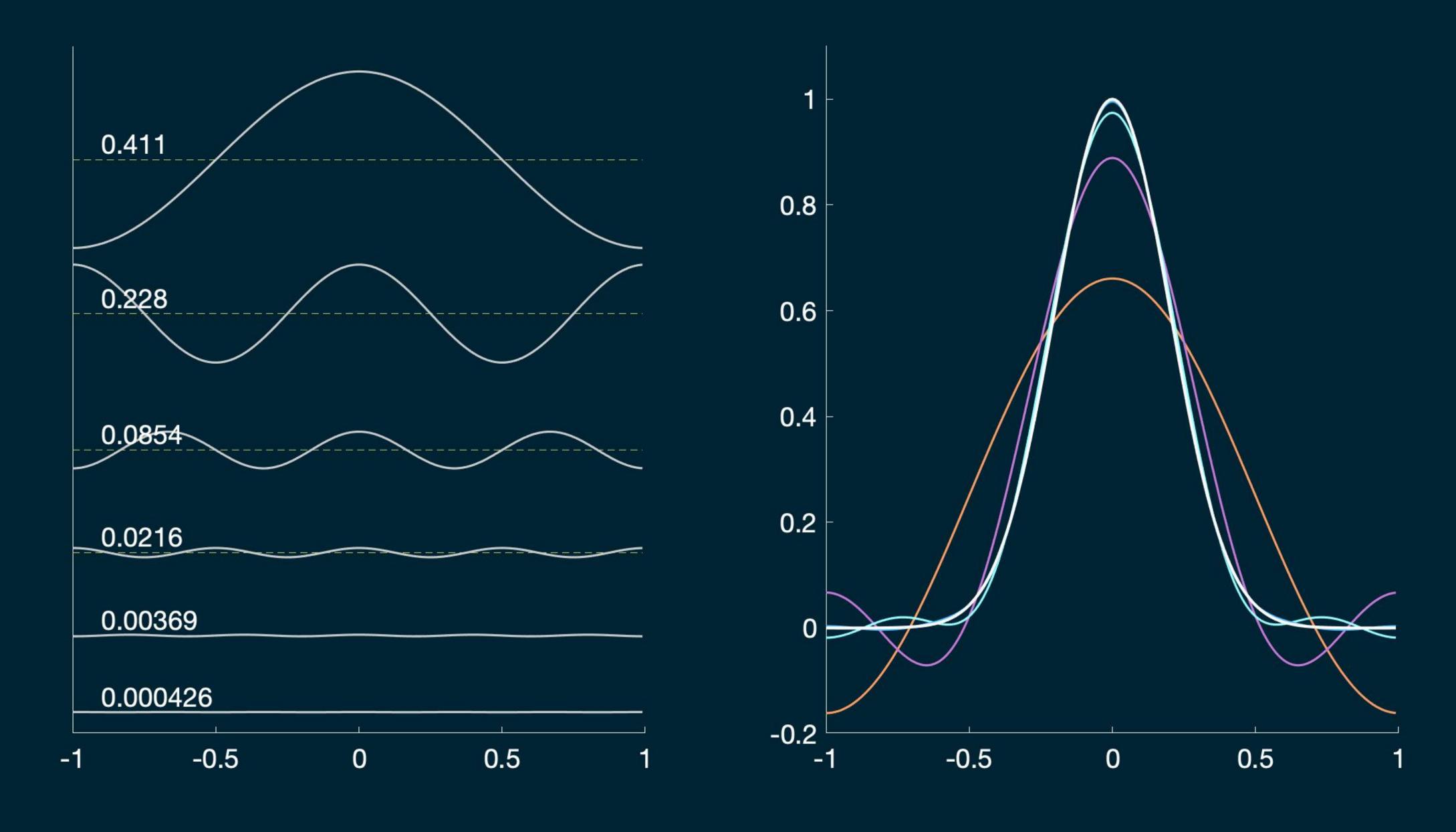




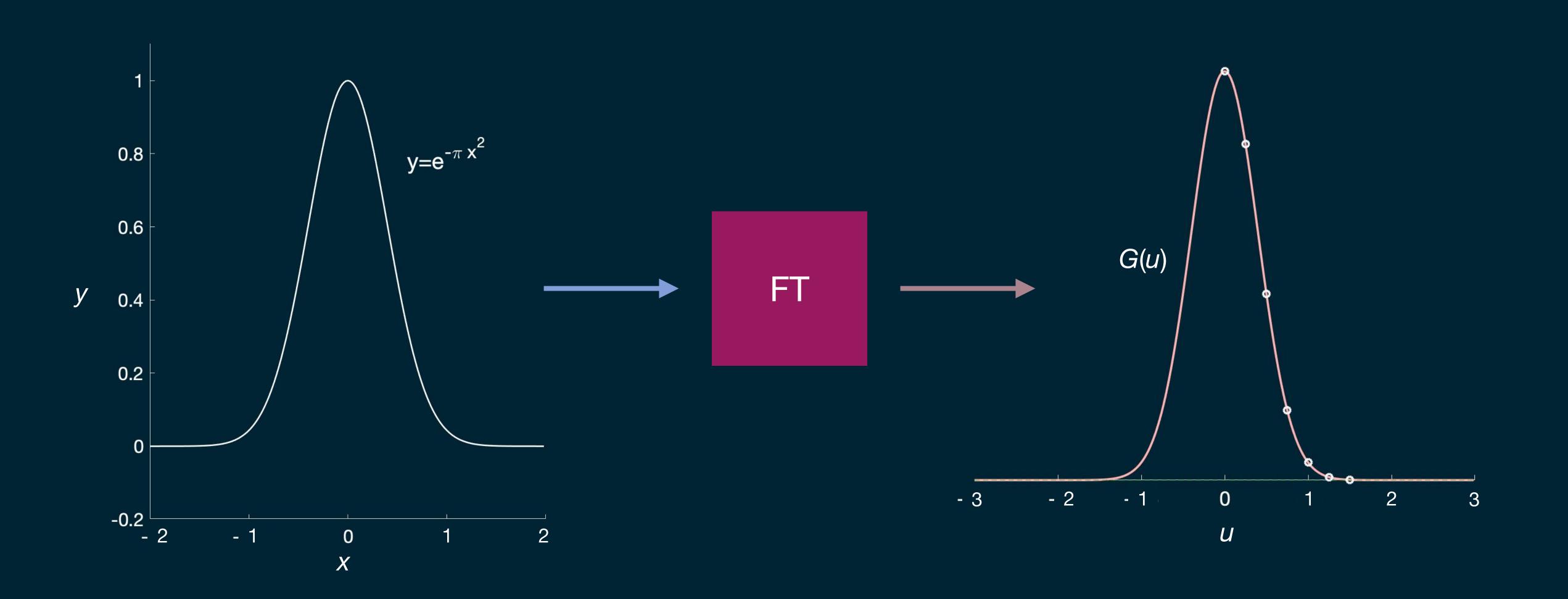




### "Converged" at 6 terms



### The Fourier Transform gives us the coefficients



#### The formulas

#### Fourier transform

$$G(u) = \int g(x)e^{-i2\pi ux}dx$$

#### Inverse Fourier transform

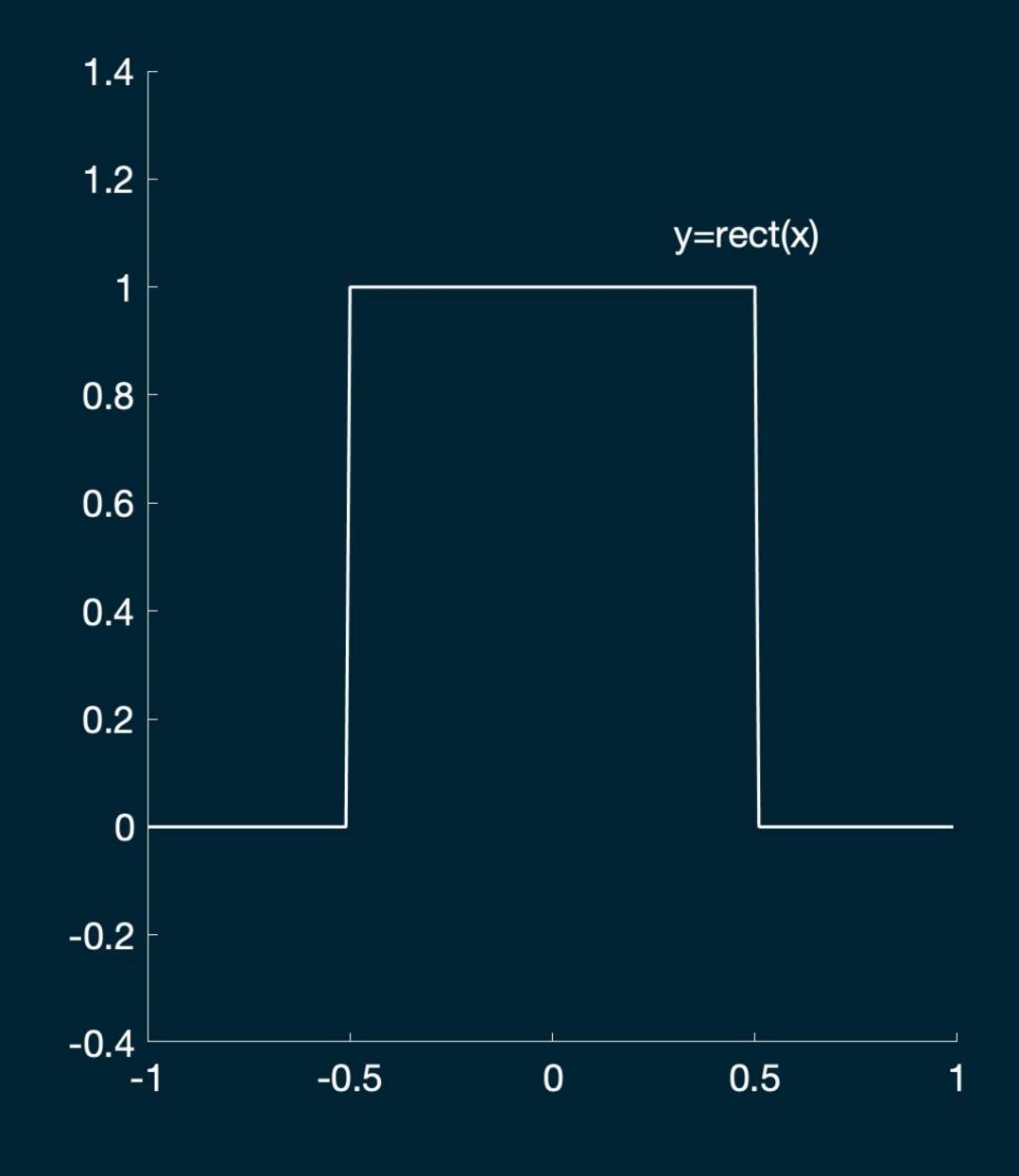
$$g(x) = \int G(u)e^{+i2\pi ux}du$$

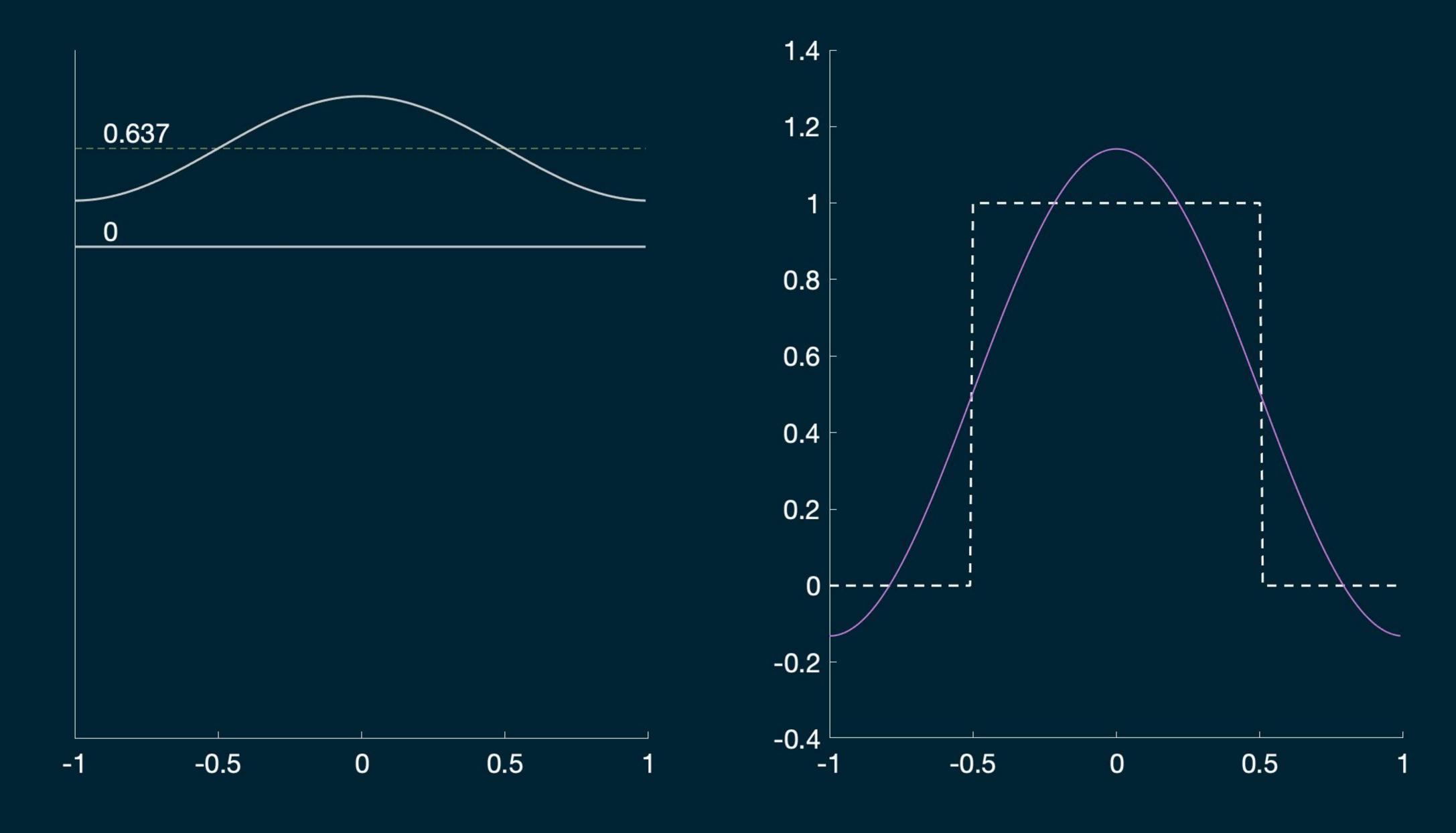
#### Example:

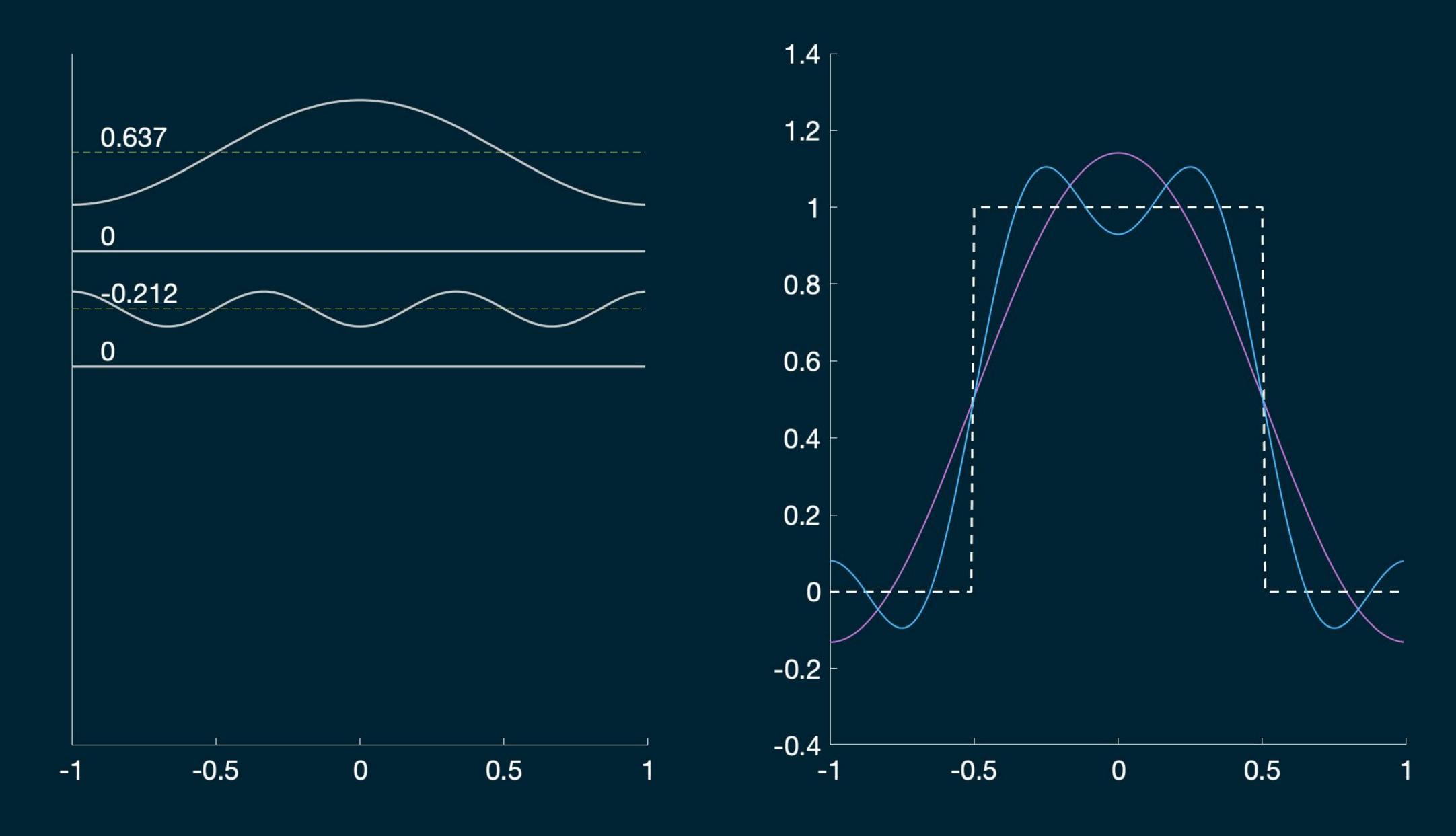
$$g(x) = e^{-\pi x^2}$$

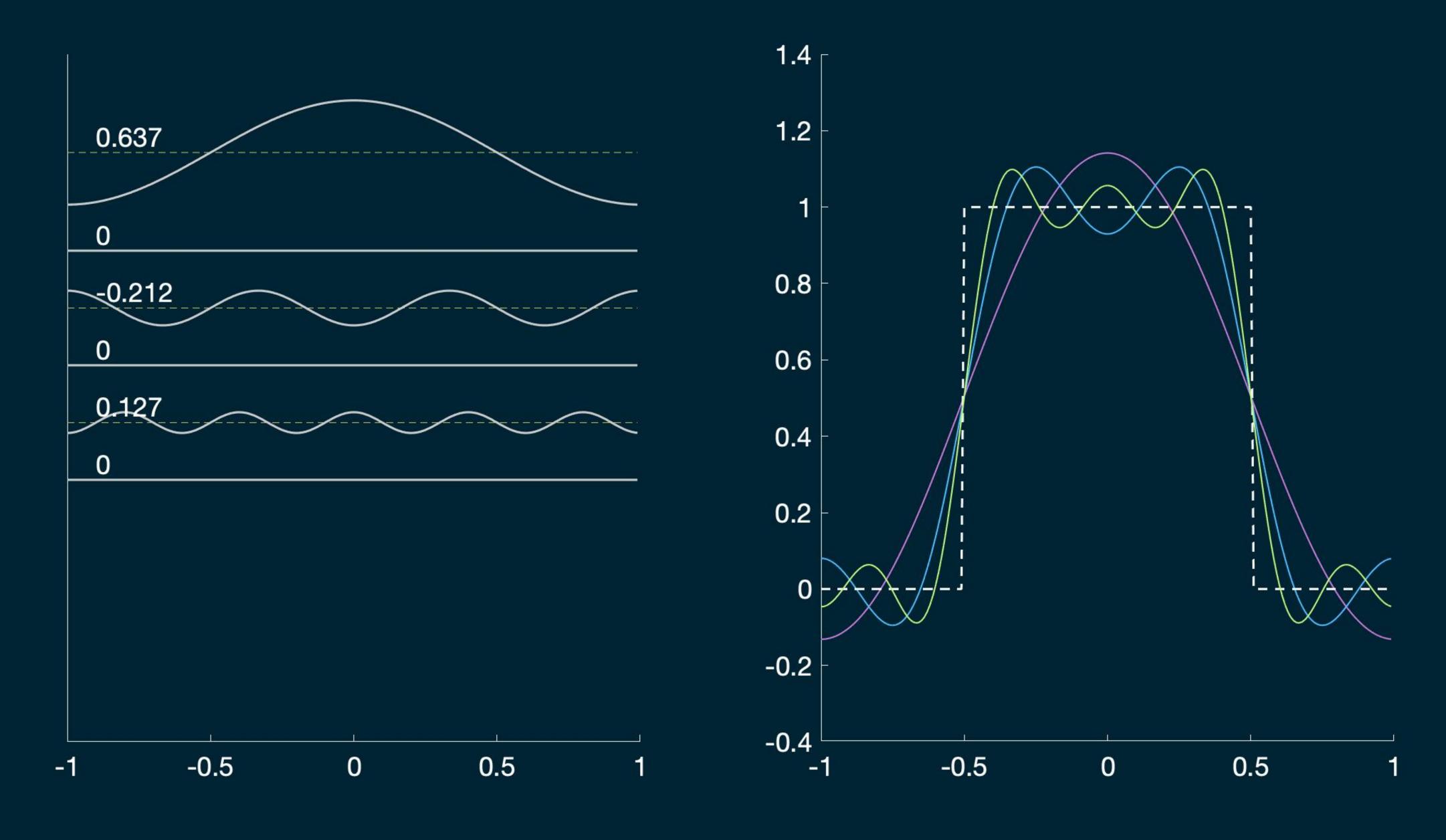
$$G(u) = e^{-\pi u^2}$$

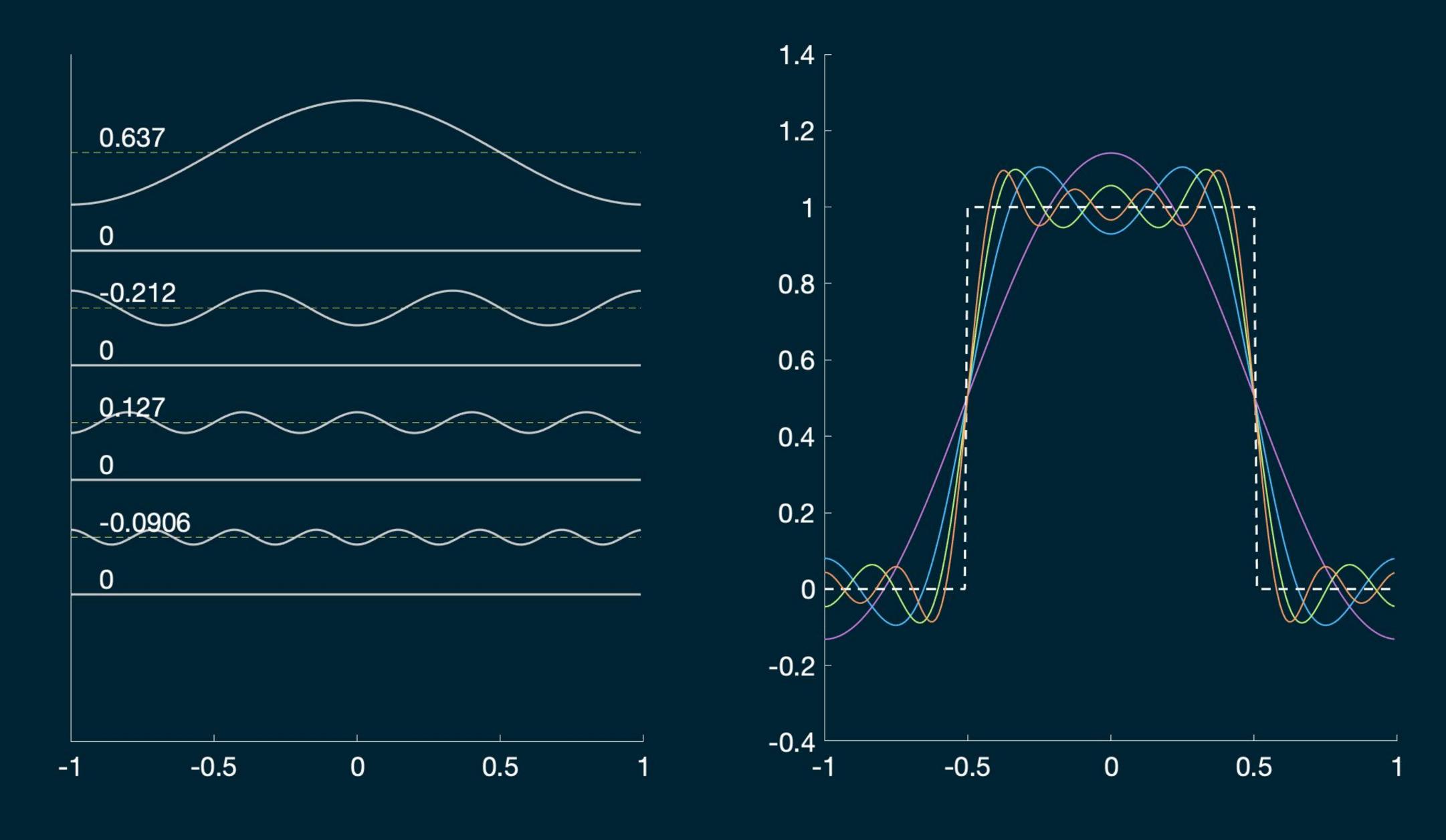
### Fourier reconstruction of a rectangular function



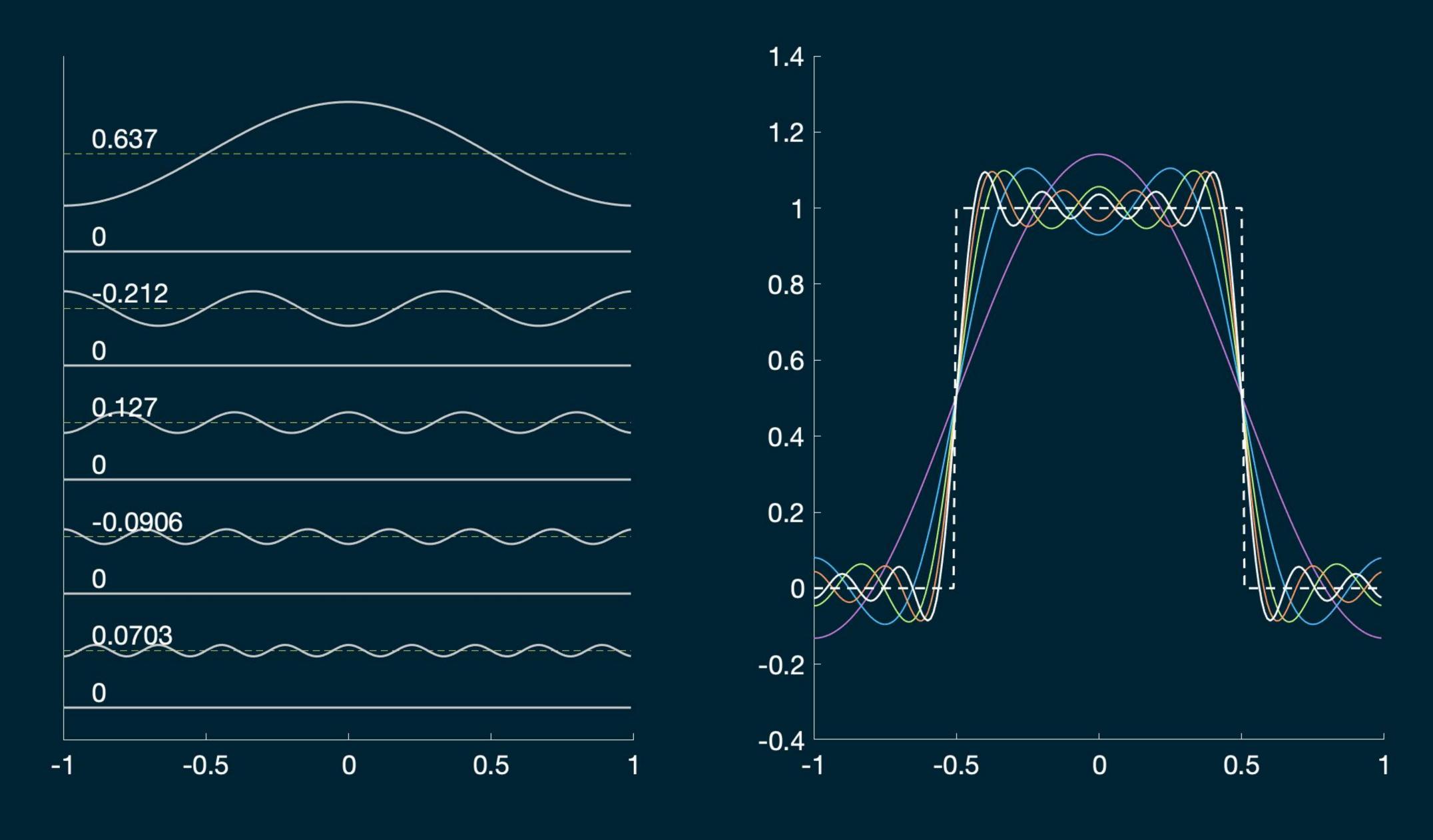




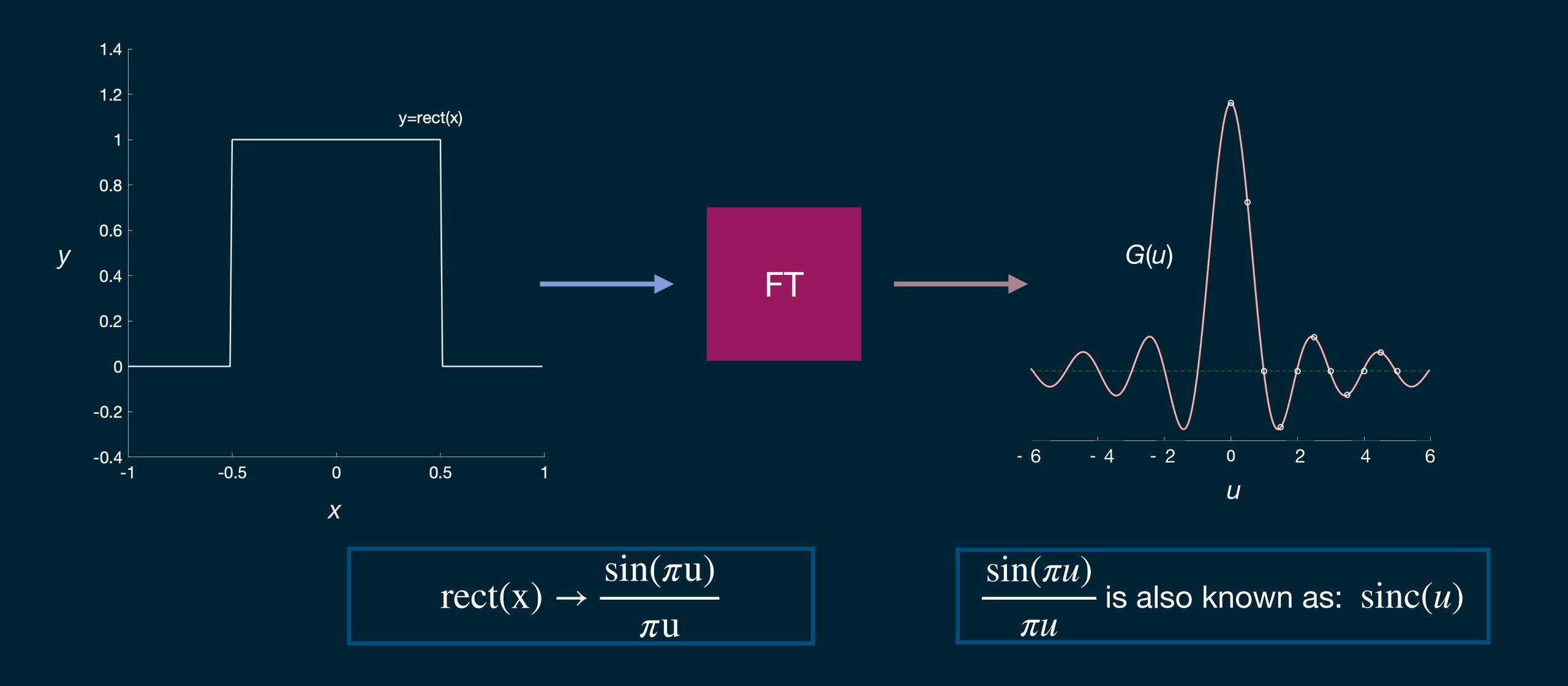




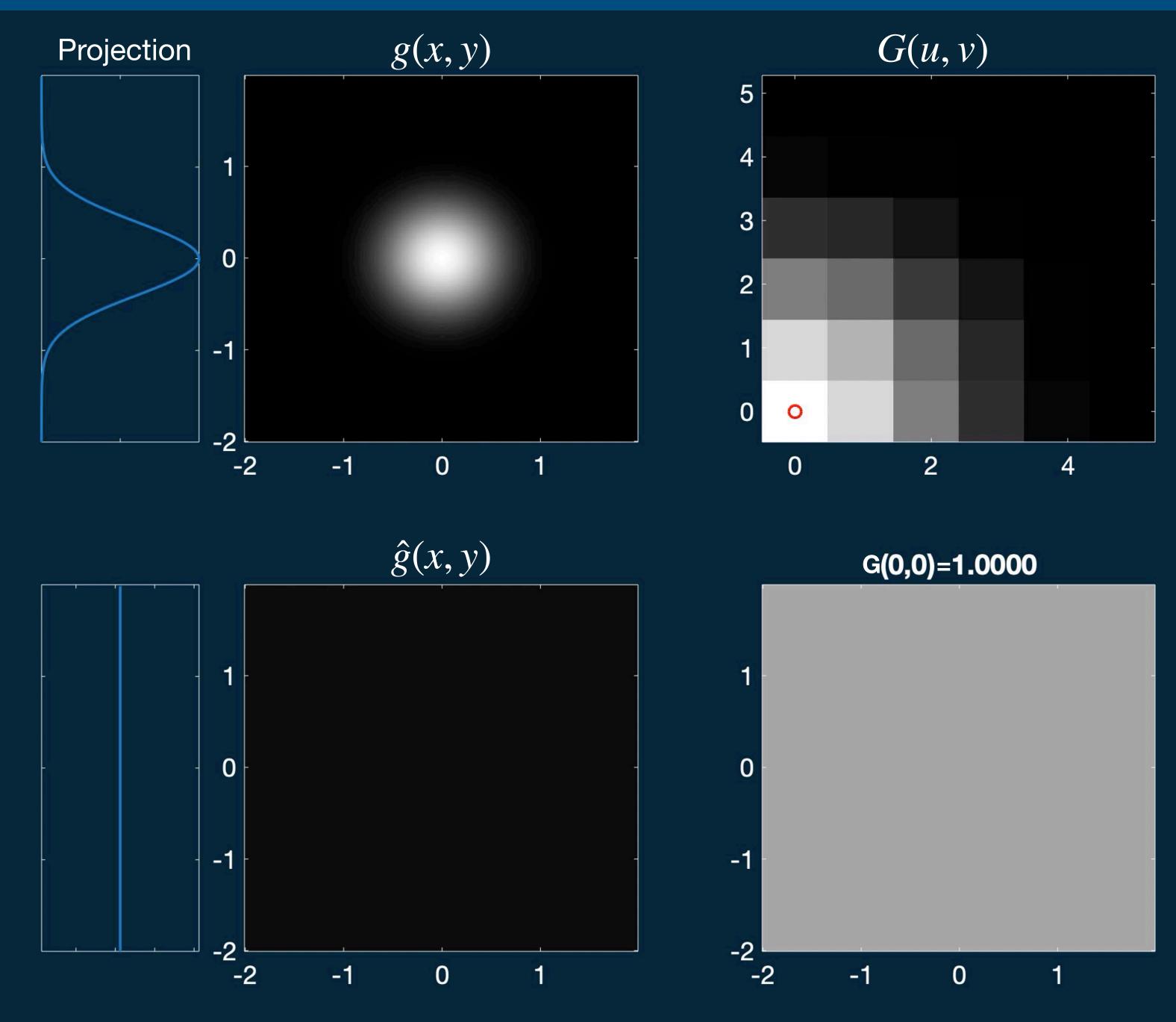
### Nowhere near convergence at 10 terms

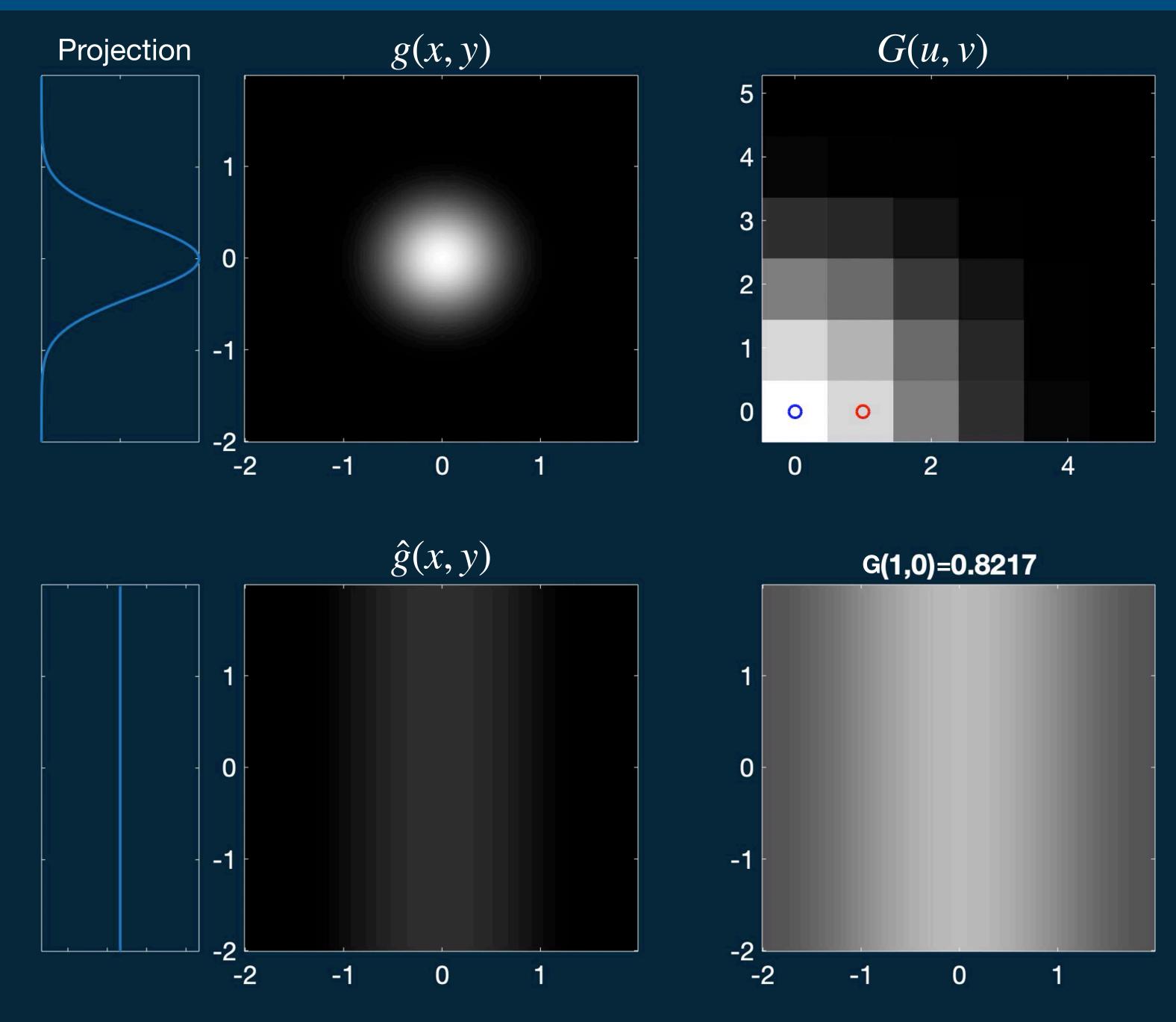


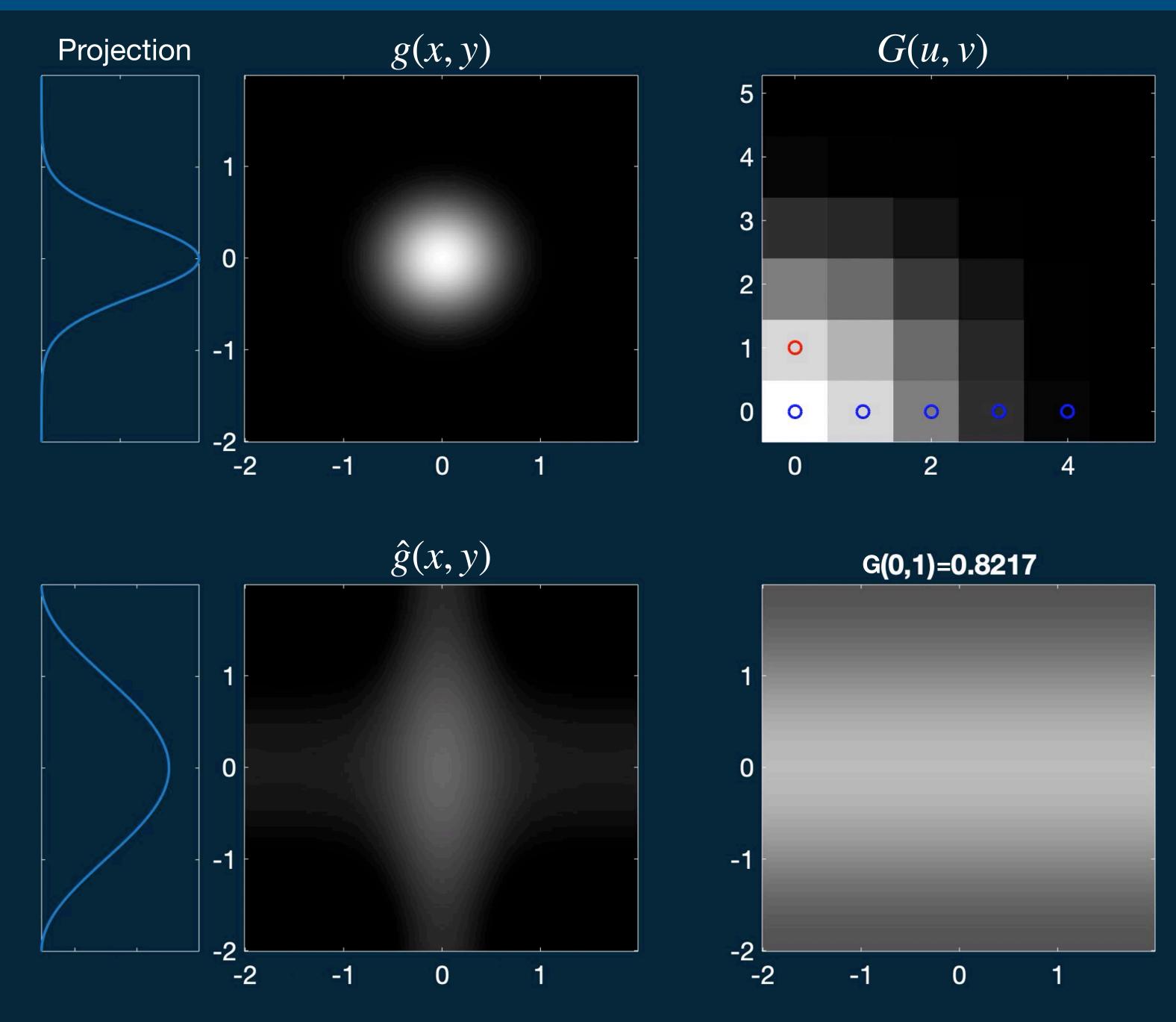
### The Fourier Transform of rect(x) is sinc(u)

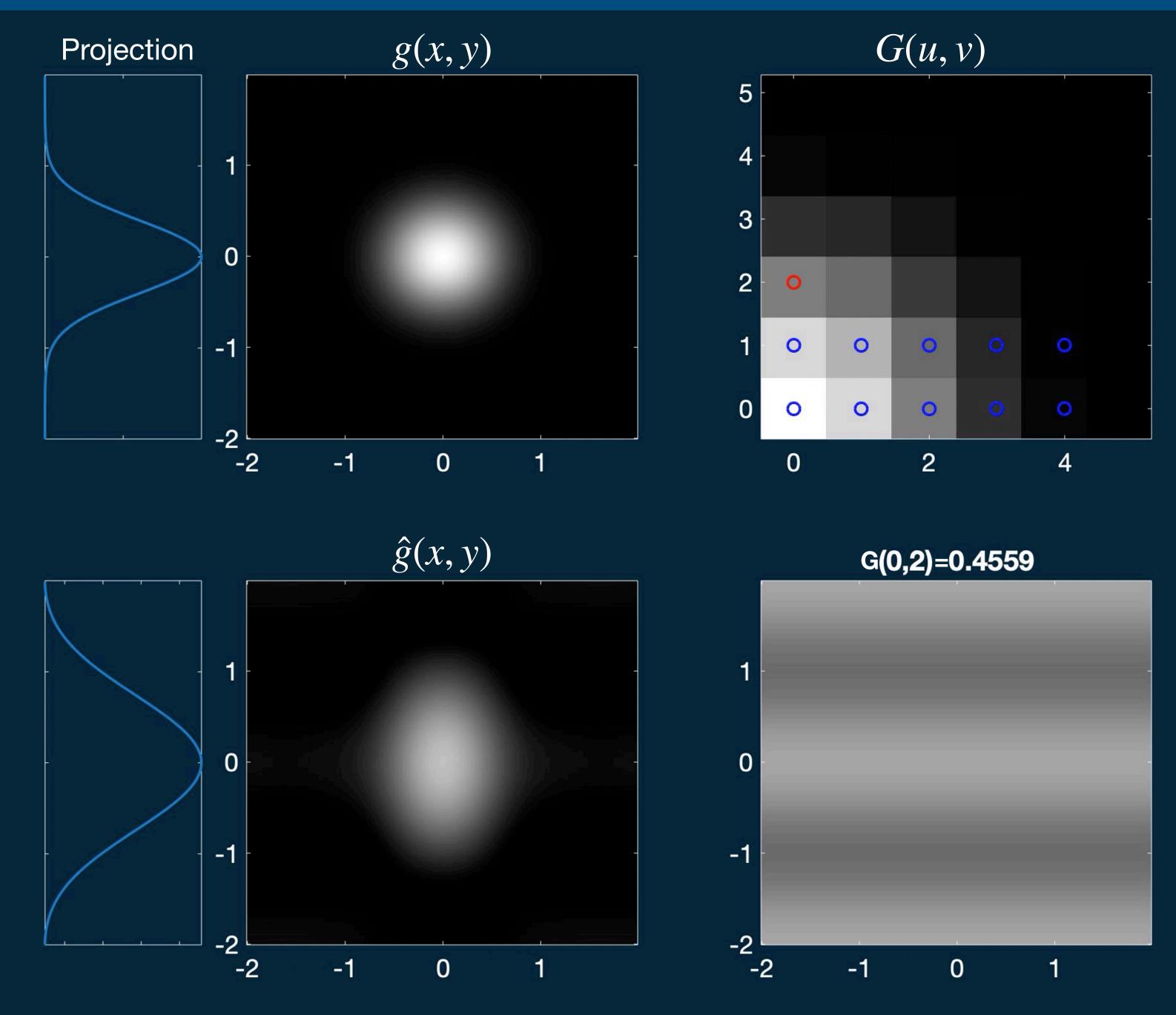


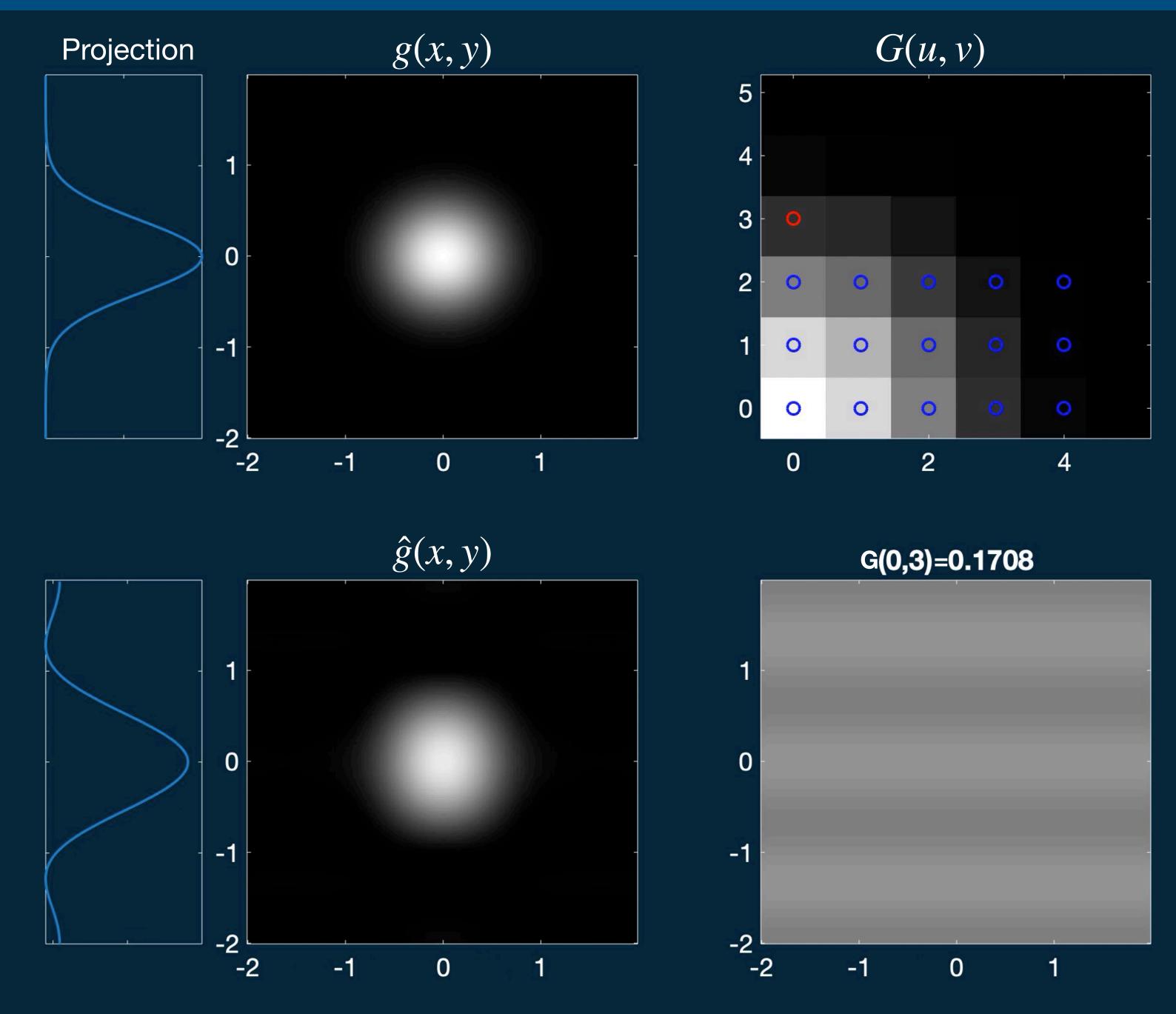
The Fourier transform in two dimensions











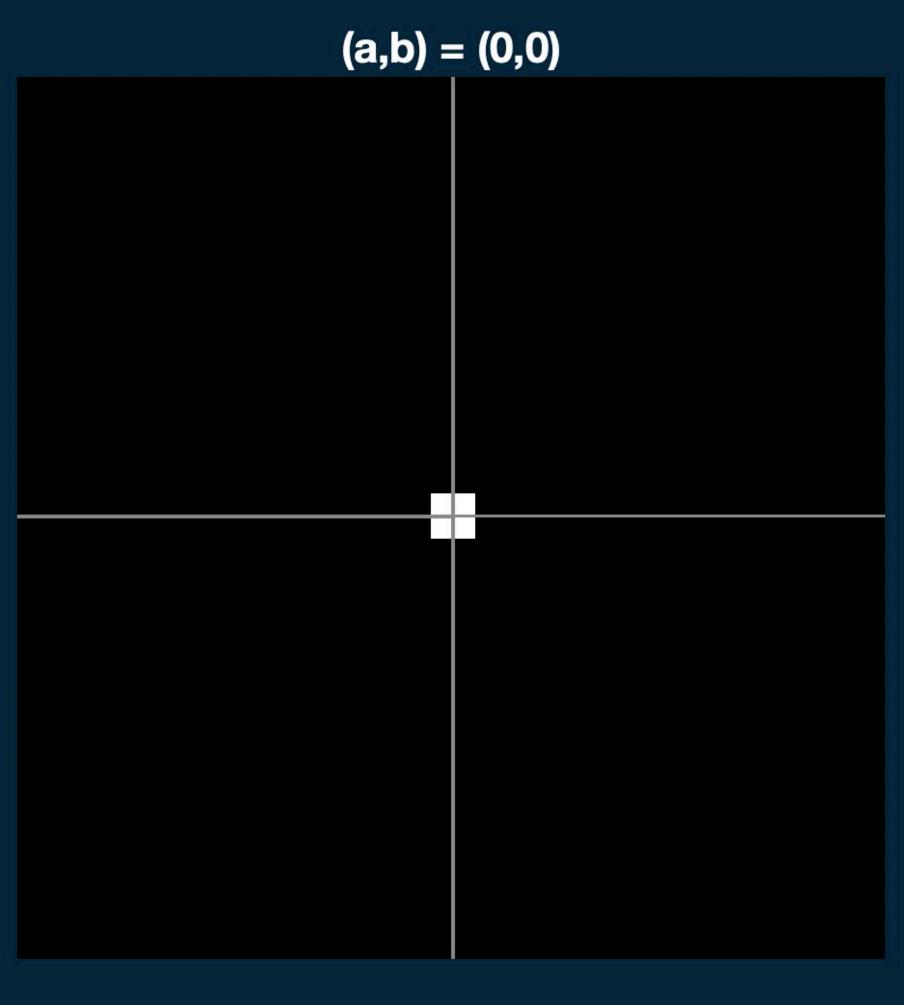
#### 2D Fourier transform

$$G(u, v) = \iint g(x, y) e^{-i2\pi(ux + vy)} dx dy$$

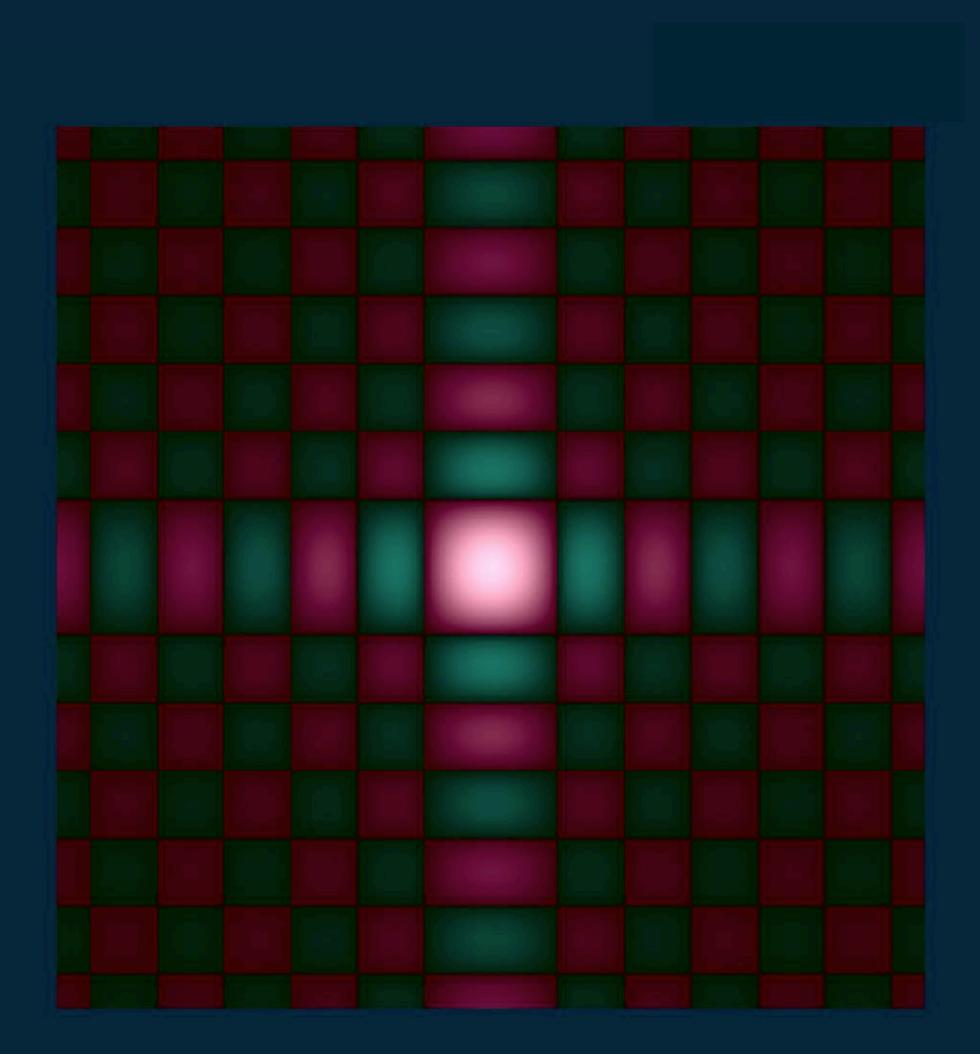
#### 2D inverse Fourier transform

$$g(x,y) = \iint G(u,v) e^{i2\pi(ux+vy)} du \, dv$$

## FT of a square

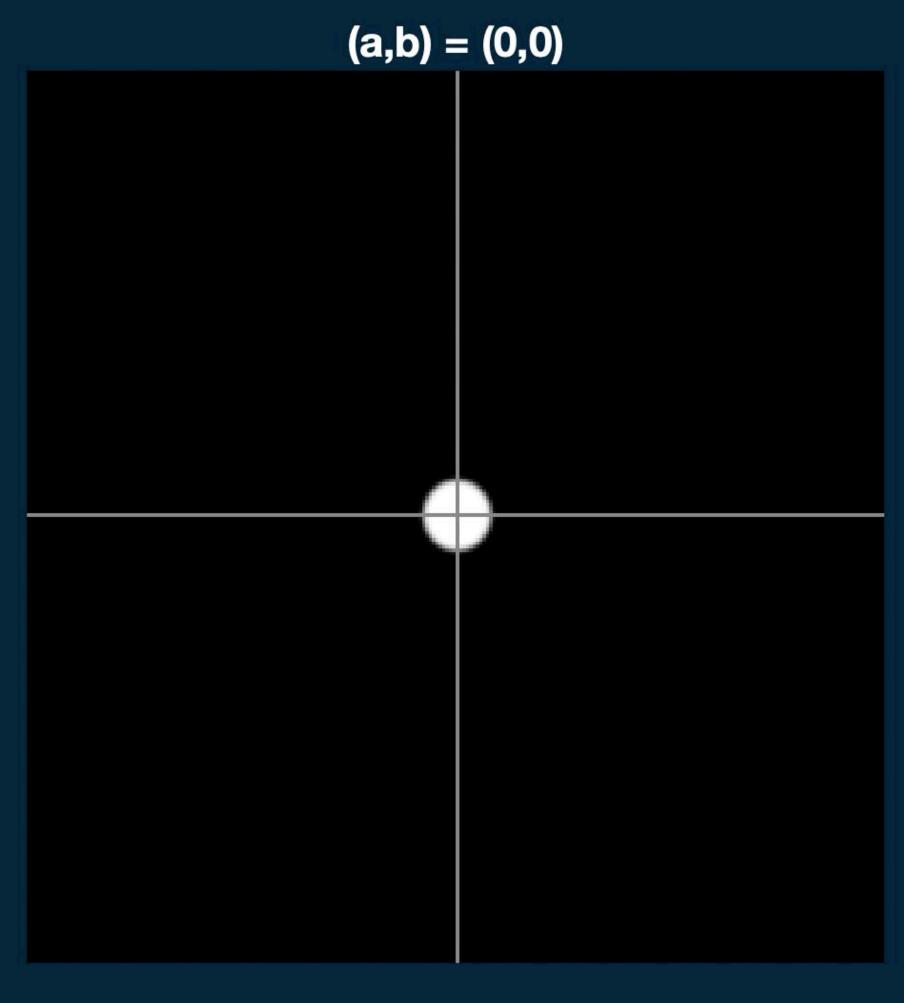


$$g = rect(x) rect(y)$$

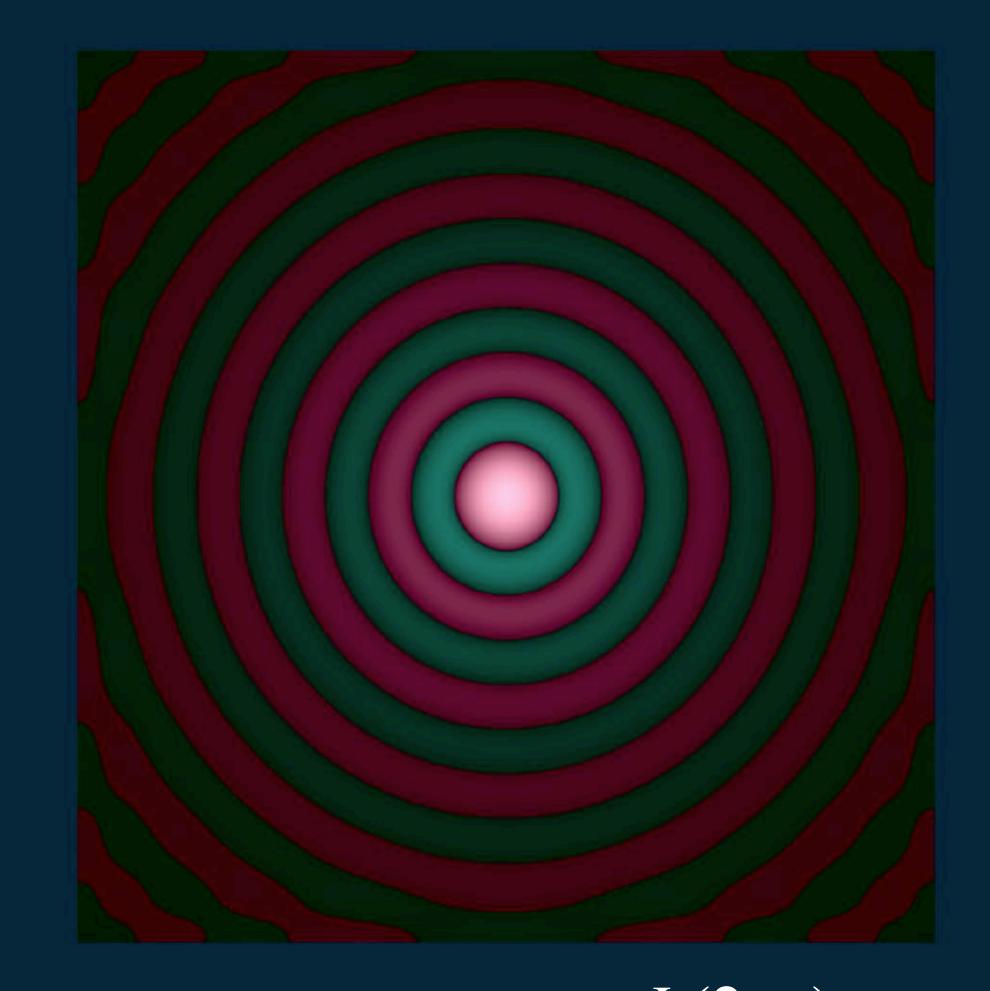


$$G = \operatorname{sinc}(u) \operatorname{sinc}(v)$$

### FT of a disc



$$g(x, y) = \operatorname{circ}(r)$$



$$G(u,v) = \frac{J_1(2\pi\rho)}{\rho}$$

#### 2D Fourier transform properties

$$ab g(ax, by) \rightarrow G(u/a, v/b)$$
 Scale

$$g(x-a,y-b) \rightarrow G(u,v)e^{-i2\pi(au+bv)}$$
 Shift

$$g * h \rightarrow GH$$
 Convolution

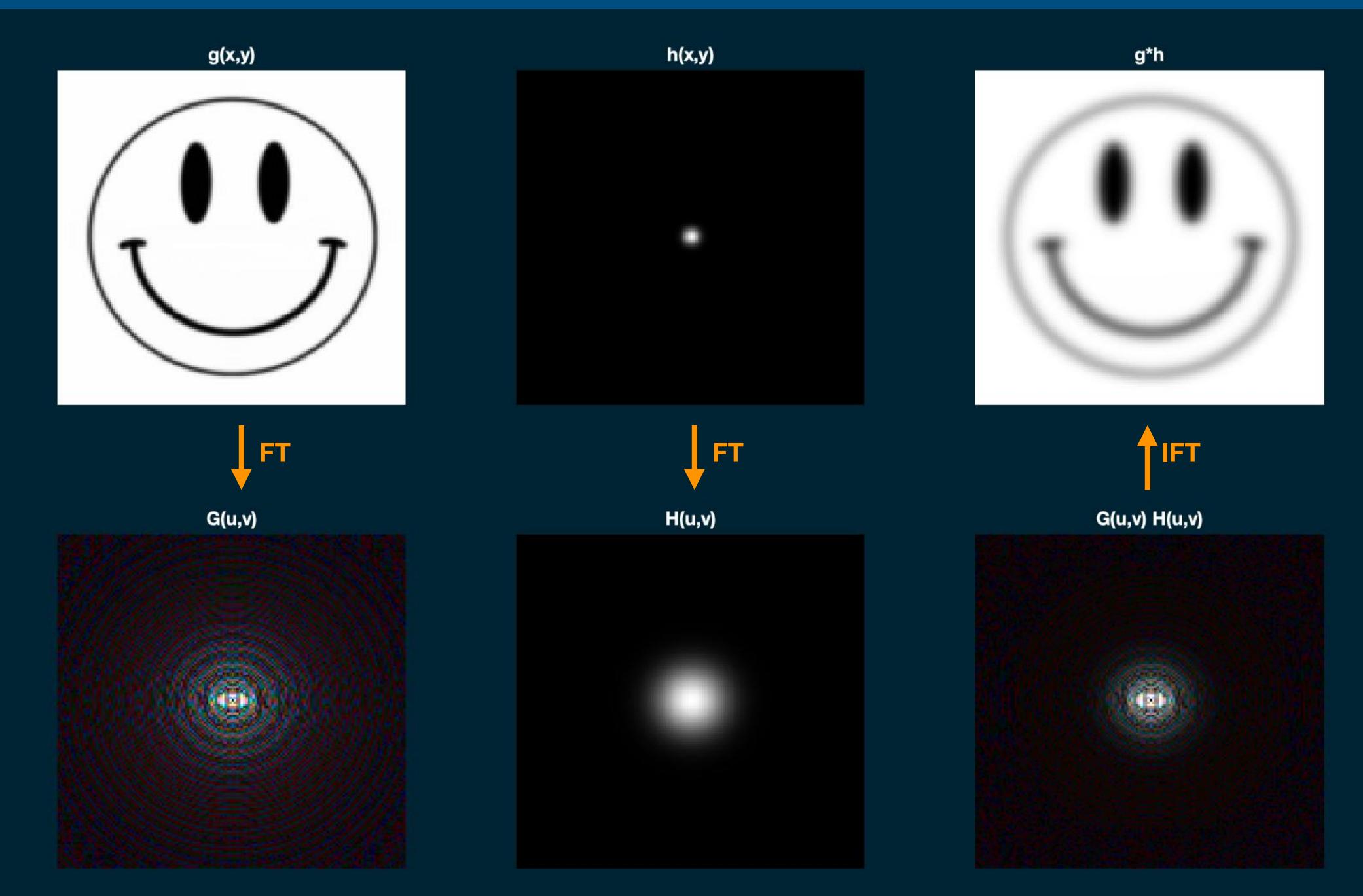
$$g(x', y') \rightarrow G(u', v')$$
 Rotation

$$P_y g(x, y) \rightarrow G(u, 0)$$
 Projection

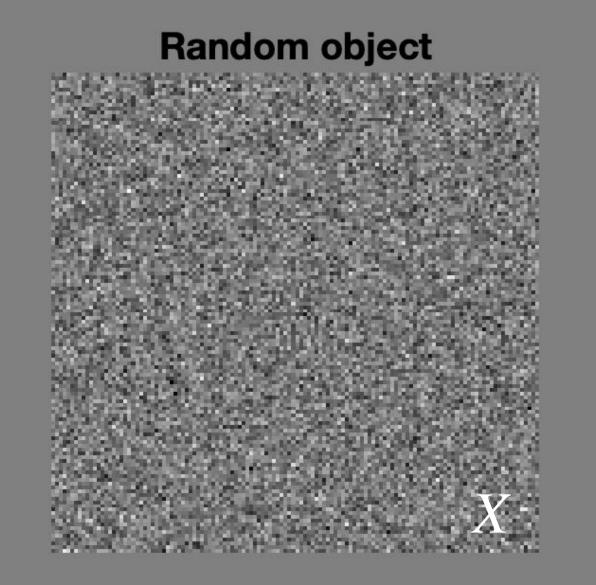
### Convolution in 2D

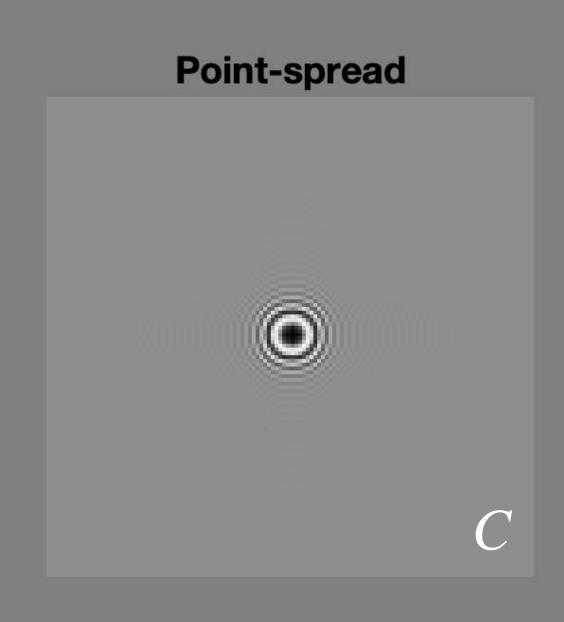
$$G \star H = \iint g(x - s, y - t) h(s, t) ds dt$$

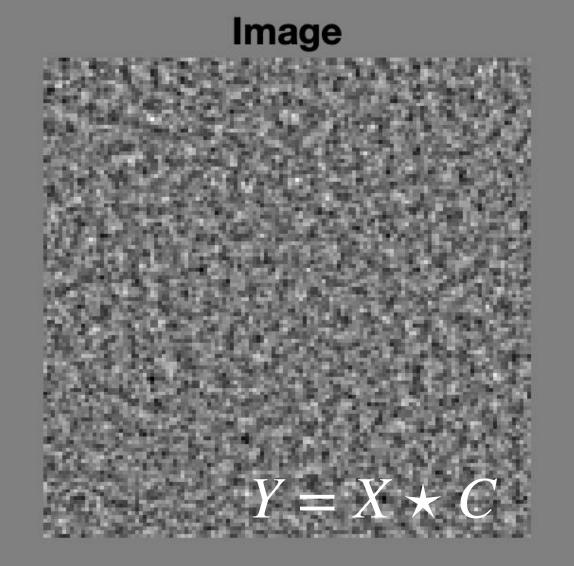
## Convolution with a Gaussian

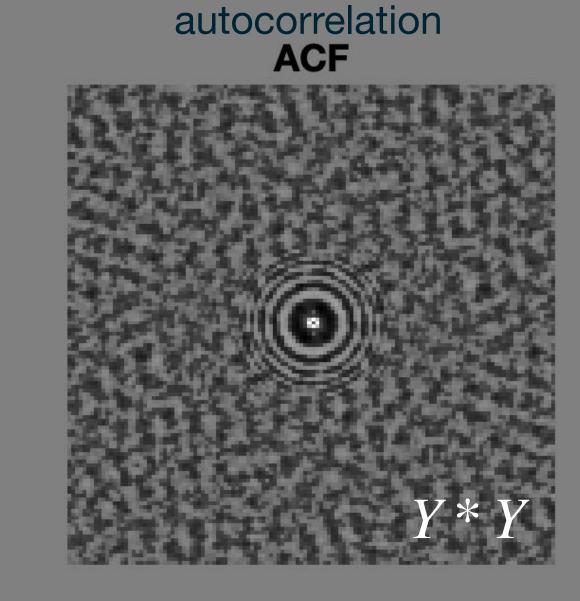


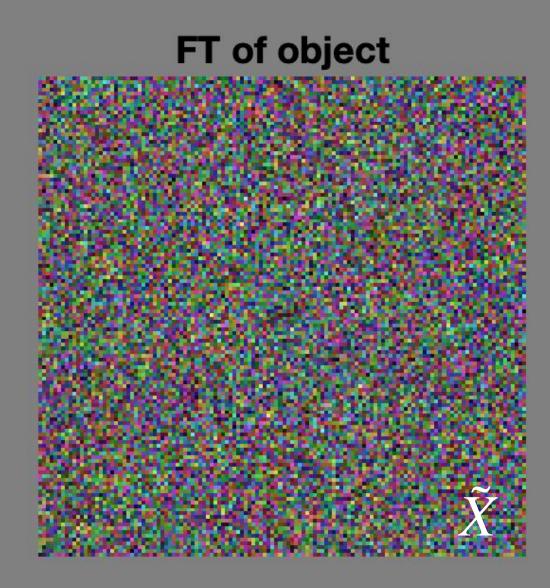
# Visualizing the contrast transfer function

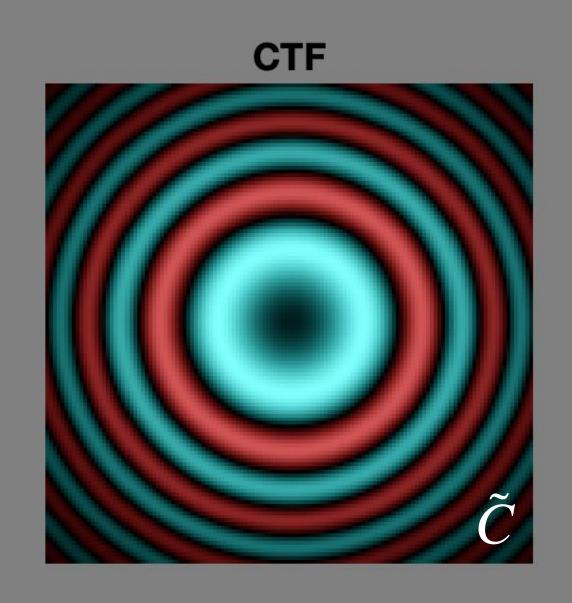


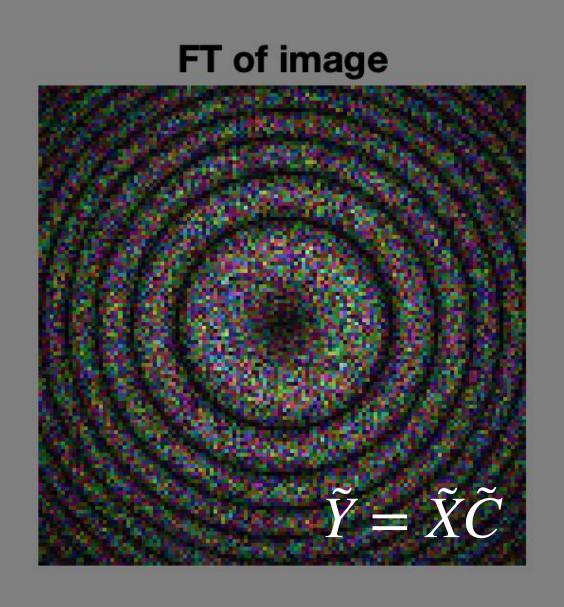


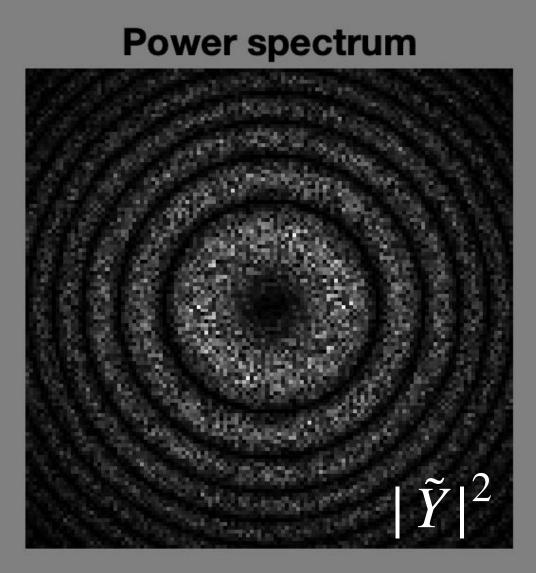












### The rotation property

2D Fourier Transform

$$G(u, v) = \iint g(x, y)e^{-i2\pi(ux+vy)}dxdy$$

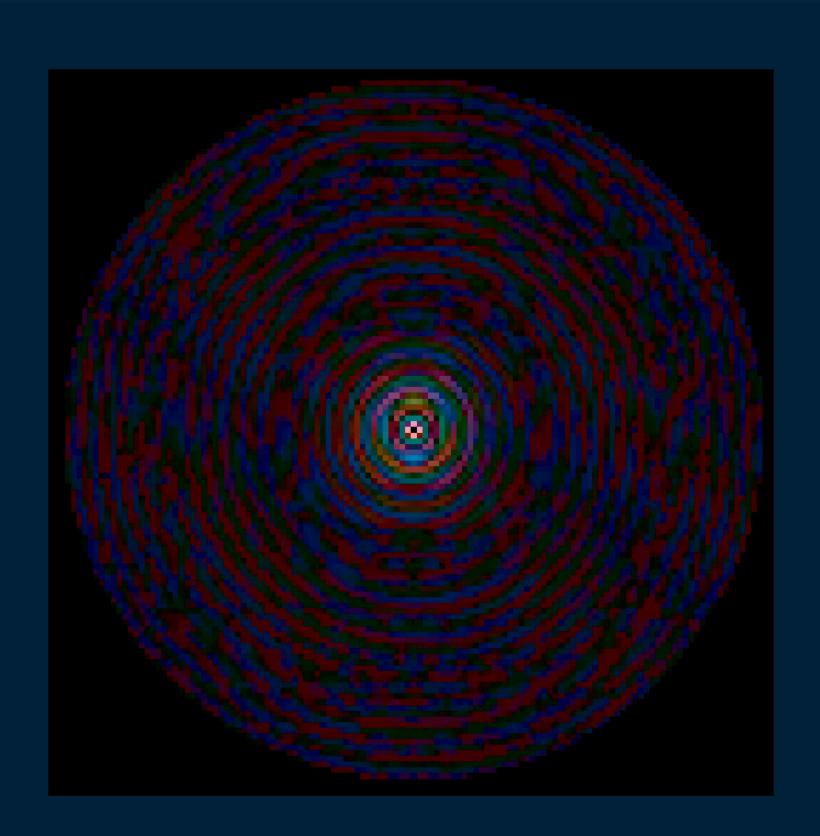
FT using 2D vectors

$$G(\mathbf{u}) = \iint g(\mathbf{x})e^{-i2\pi(\mathbf{u}\cdot\mathbf{x})}d^2\mathbf{x}$$

The dot-product is invariant under rotations!







Let  $R_{\theta}$  signify a rotation, and

$$(x', y') = R_{\theta}(x, y)$$
$$(u', v') = R_{\theta}(u, v)$$

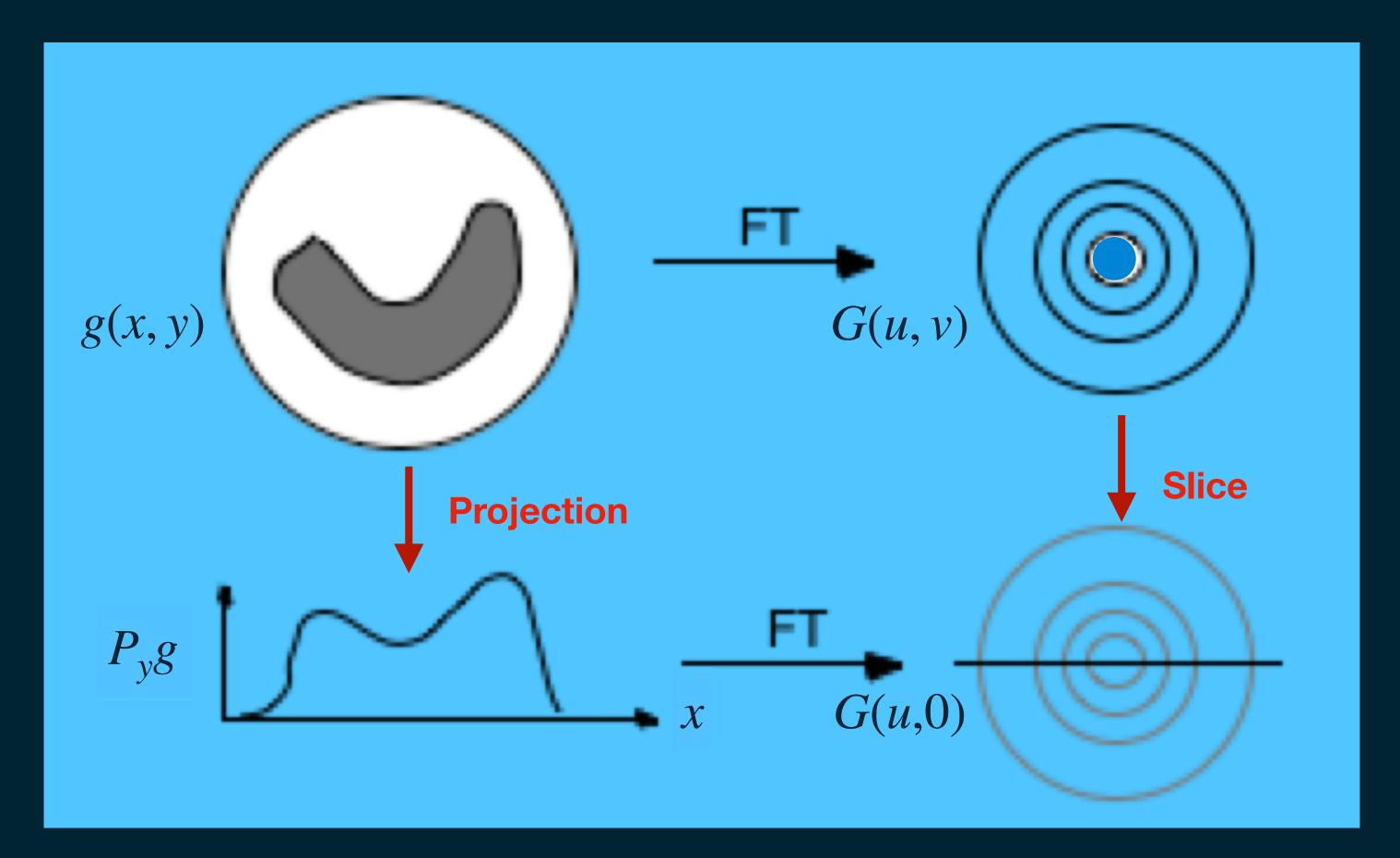
then

$$g(x',y') \rightarrow G(u',v')$$

or alternatively,

$$g(R_{\theta}\mathbf{x}) \to G(R_{\theta}\mathbf{u})$$

#### The Fourier Slice Theorem

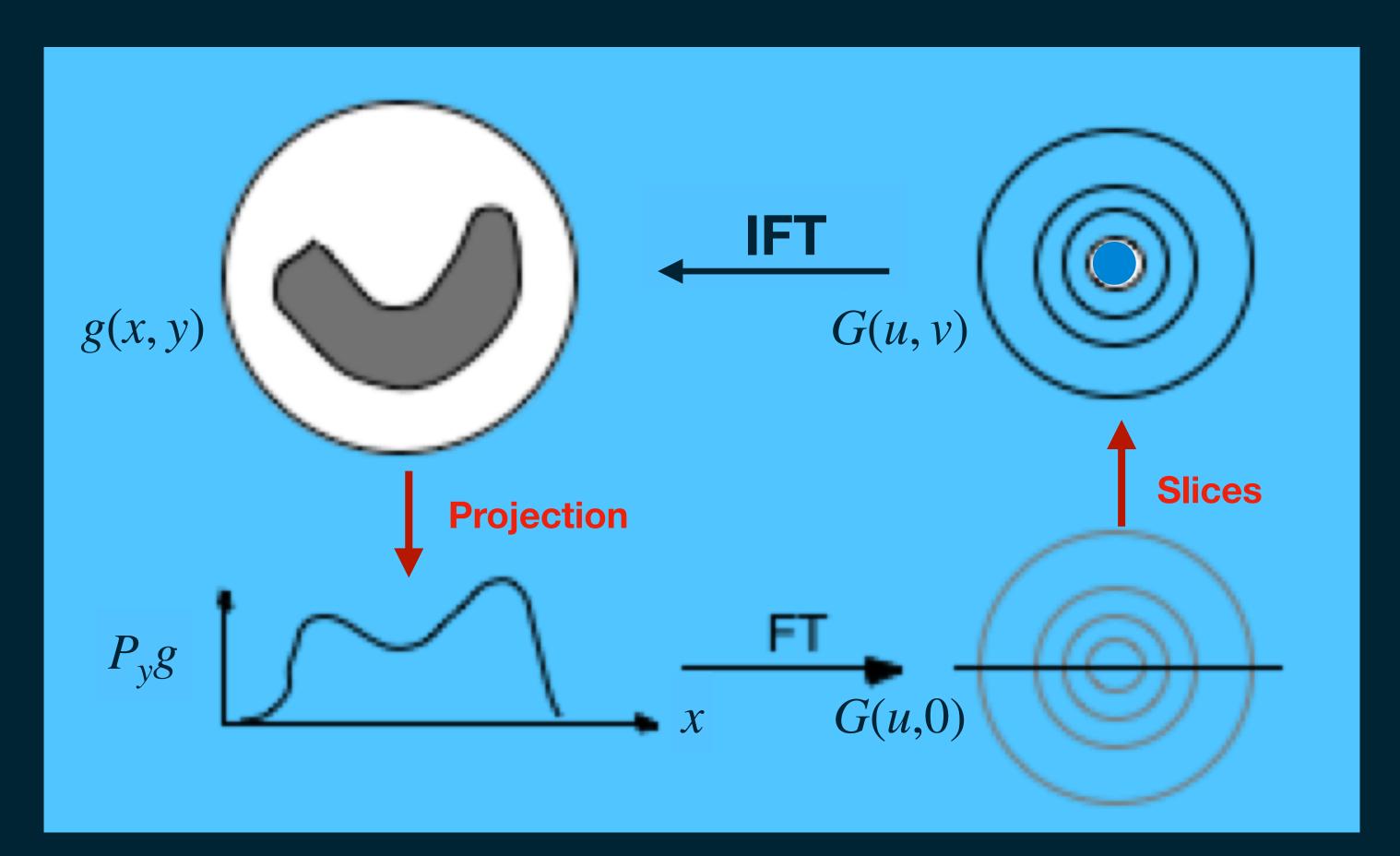


$$G(u, v) = \iint g(x, y)e^{-i2\pi(ux+vy)}dxdy$$

$$G(u,0) = \iint g(x,y)dy e^{-i2\pi(ux)}dx$$
$$= \mathcal{F}\{P_yg\}$$

$$P_{y}g(x,y) = \int g(x,y)dy$$

## Reconstruction using the Fourier Slice Theorem



$$G(u, v) = \iint g(x, y)e^{-i2\pi(ux+vy)}dxdy$$

$$G(u,0) = \iint g(x,y)dy e^{-i2\pi(ux)}dx$$
$$= \mathcal{F}\{P_yg\}$$

$$P_{y}g(x,y) = \int g(x,y)dy$$

The rotation property says: If we can collect projections from all directions, we can construct all of G(u, v)

### The discrete FT is what is calculated on a computer

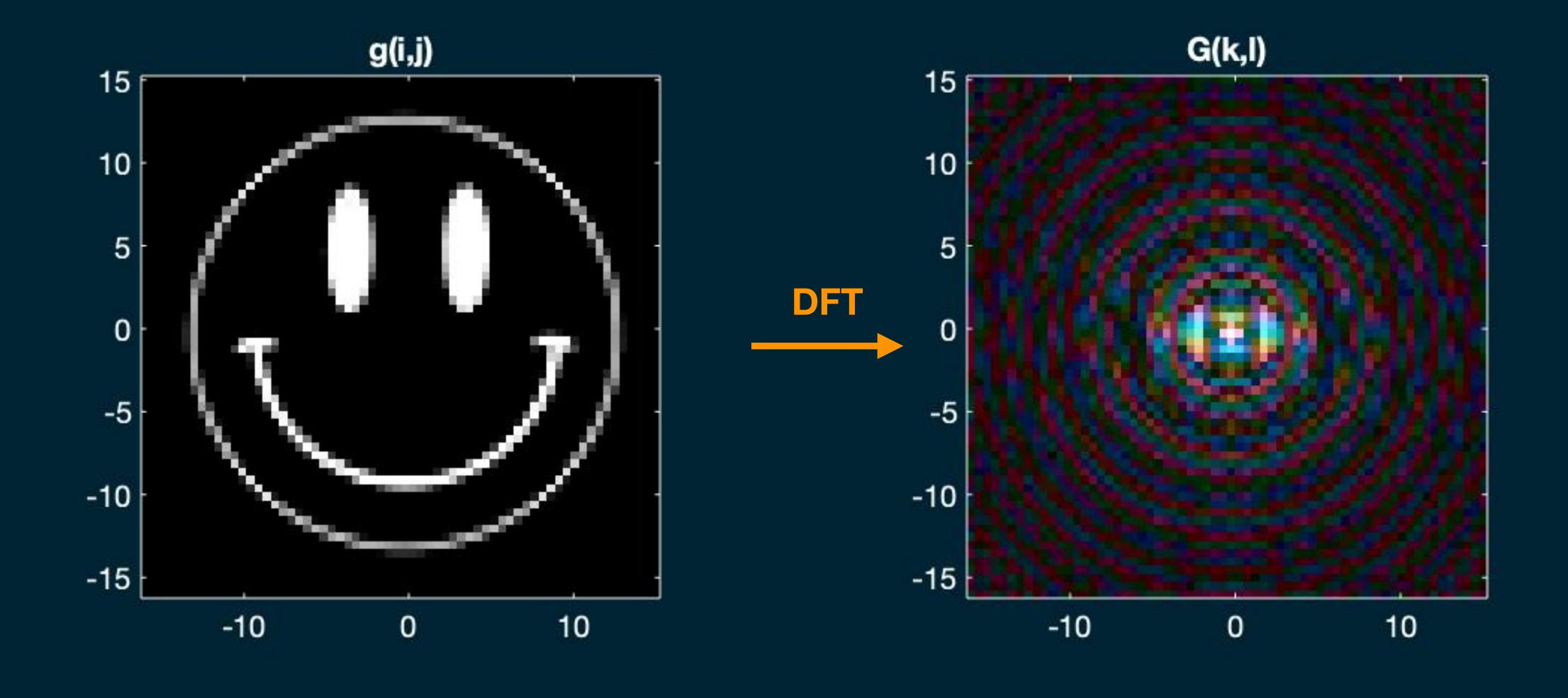
#### 2D Fourier transform

$$G(u, v) = \iint g(x, y) e^{-i2\pi(ux+vy)} dx dy$$

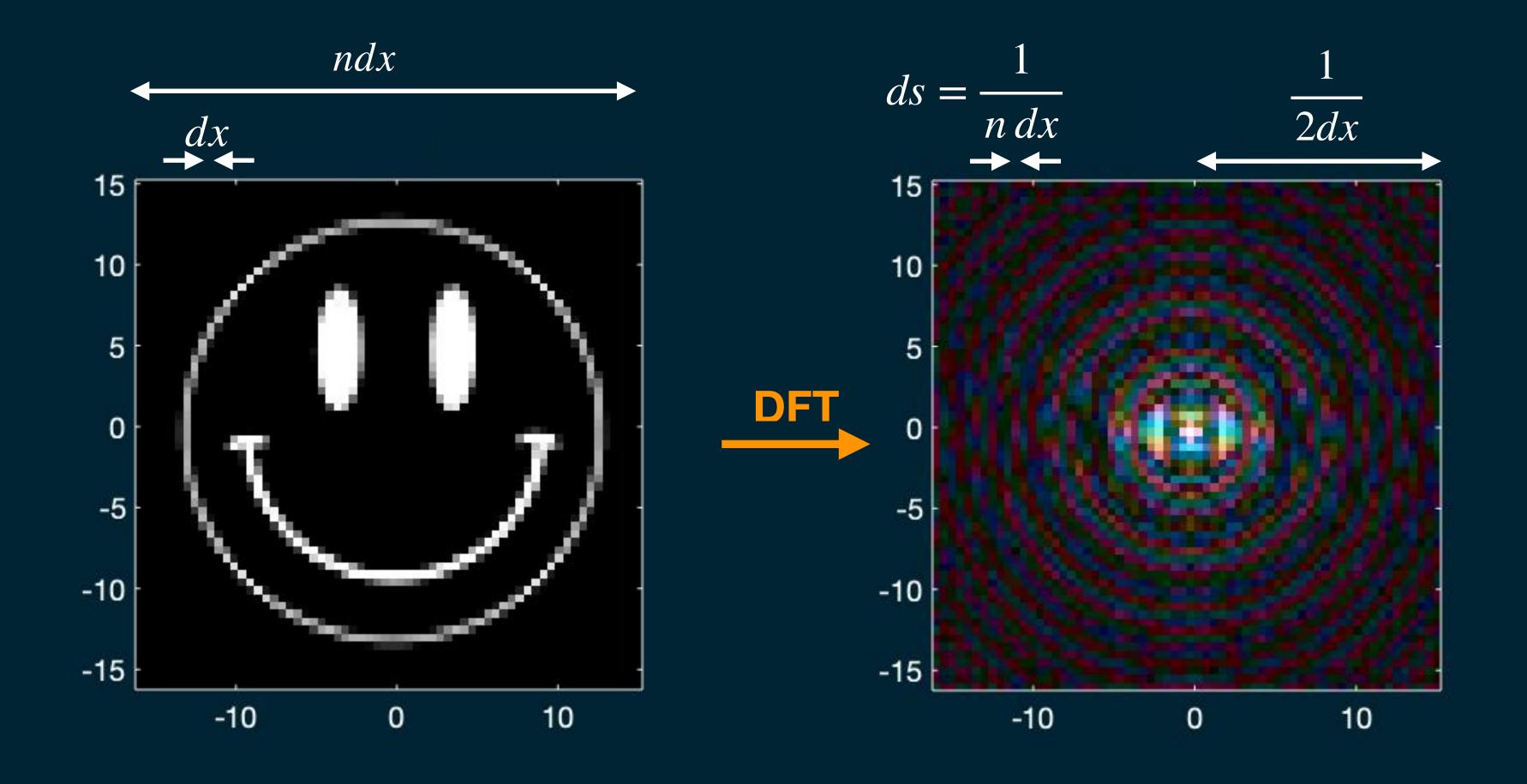
#### 2D discrete Fourier transform

$$G(k, l) = \frac{1}{N} \sum_{i,j=-N/2}^{N/2-1} g(i,j) e^{-i2\pi(ik+jl)N}$$

# The DFT of a 32 x 32 pixel image has 32 x 32 complex pixel values



# What are the dimensions of the transformed image?



Note that the sampling frequency 1/dx ..... corresponds to twice the maximum accessible frequency nds/2.

#### The 3D transform

#### 3D Fourier transform

$$G(u, v, w) = \iiint g(x, y, z)e^{-i2\pi(ux+vy+wz)}dx dy dz$$

#### 3D Inverse Fourier transform

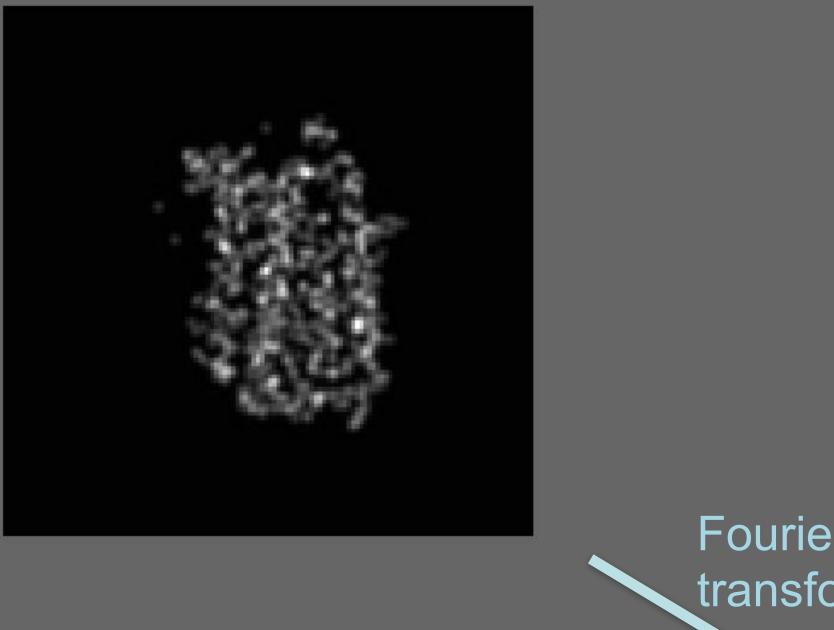
$$g(x, y, z) = \iiint G(u, v, w)e^{+i2\pi(ux+vy+wz)}du\,dv\,dw$$

Theoretical basis of single-particle reconstruction

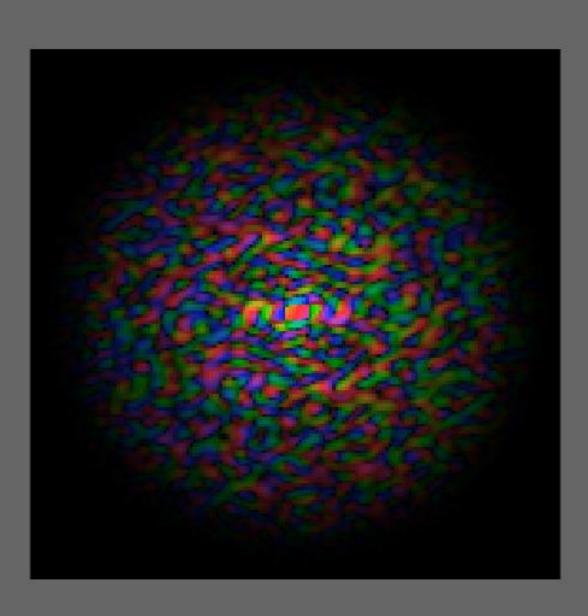
### Image processing with Fourier transforms

$$g(x,y) o G(u,v)$$
 Fourier Transform  $g \star h o GH$  Convolution  $g \otimes h o GH^*$  Correlation  $g(x',y') o G(u',v')$  Rotation  $P_y g(x,y) o G(u,0)$  Projection

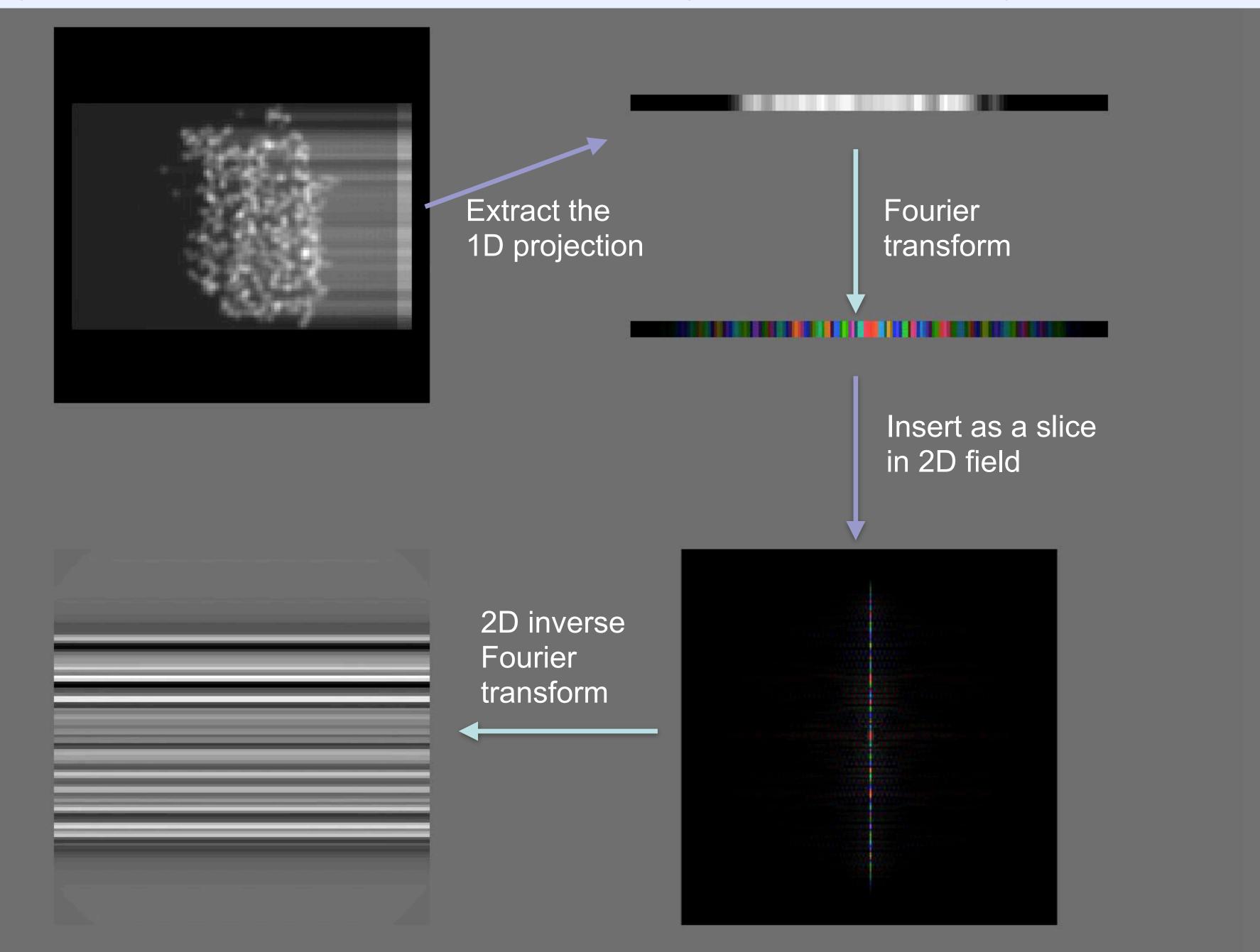
# How to get 3D structures from 2D images? The Fourier slice theorem



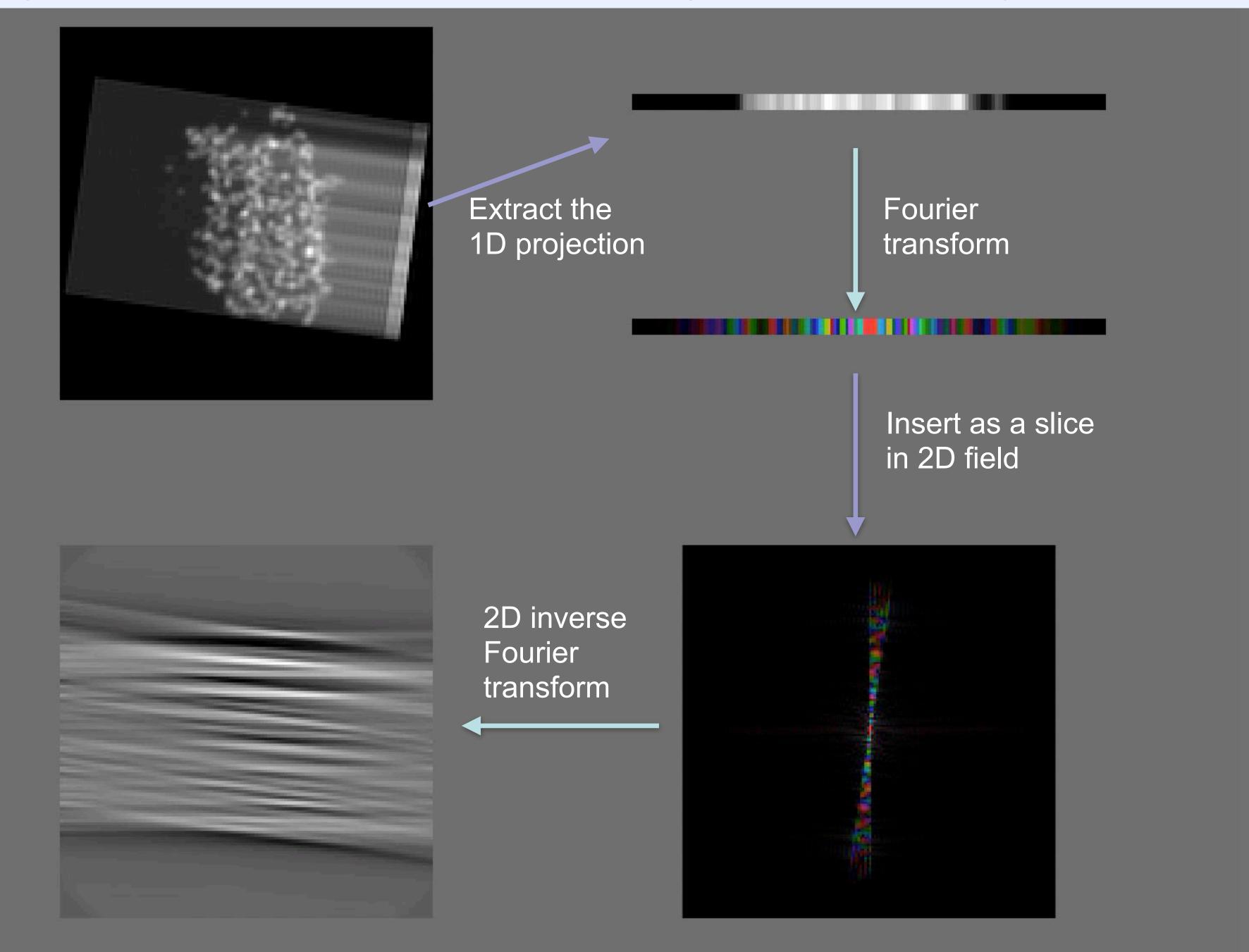




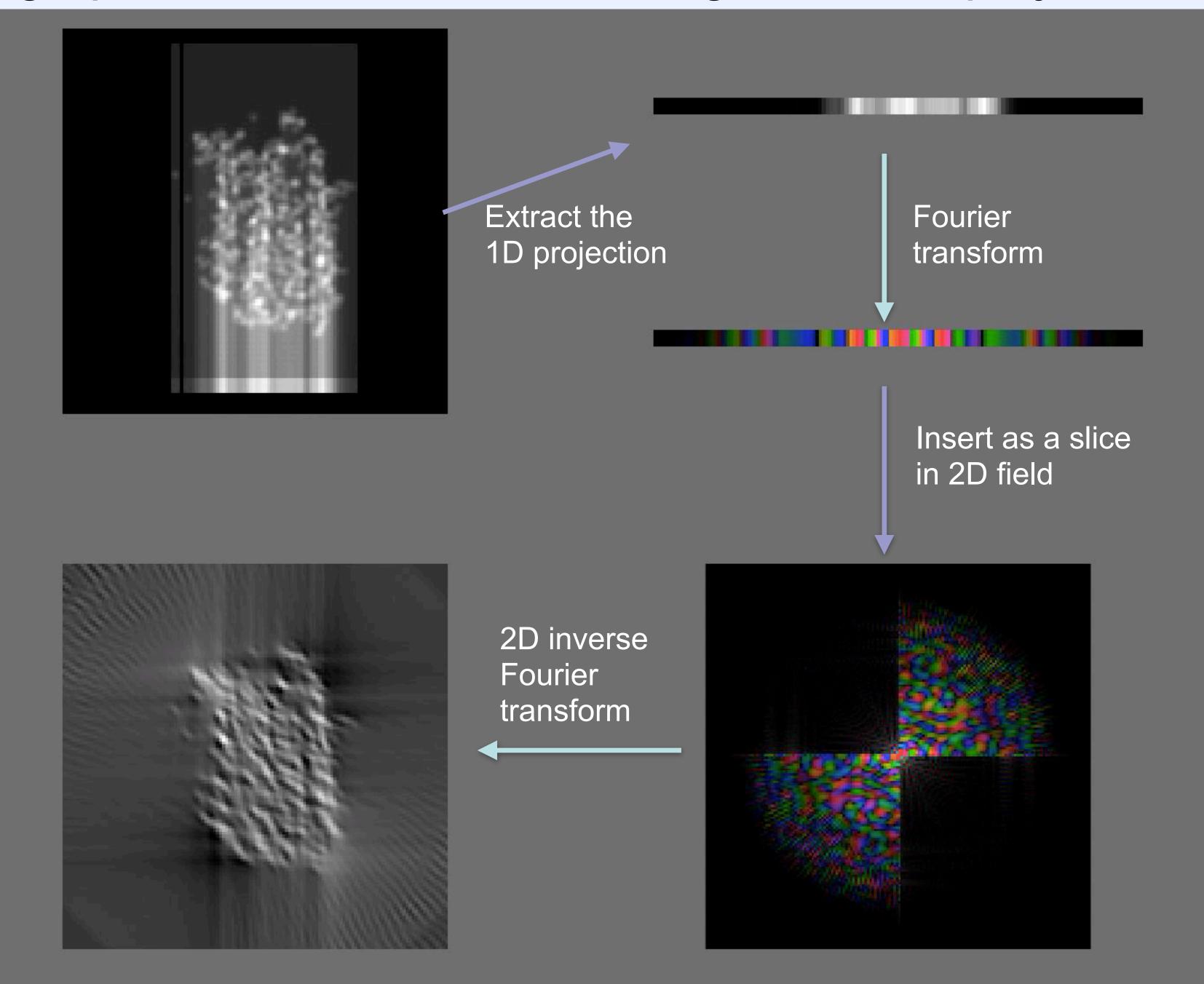
# Tomographic reconstruction: 2D image from 1D projections



# Tomographic reconstruction: 2D image from 1D projections



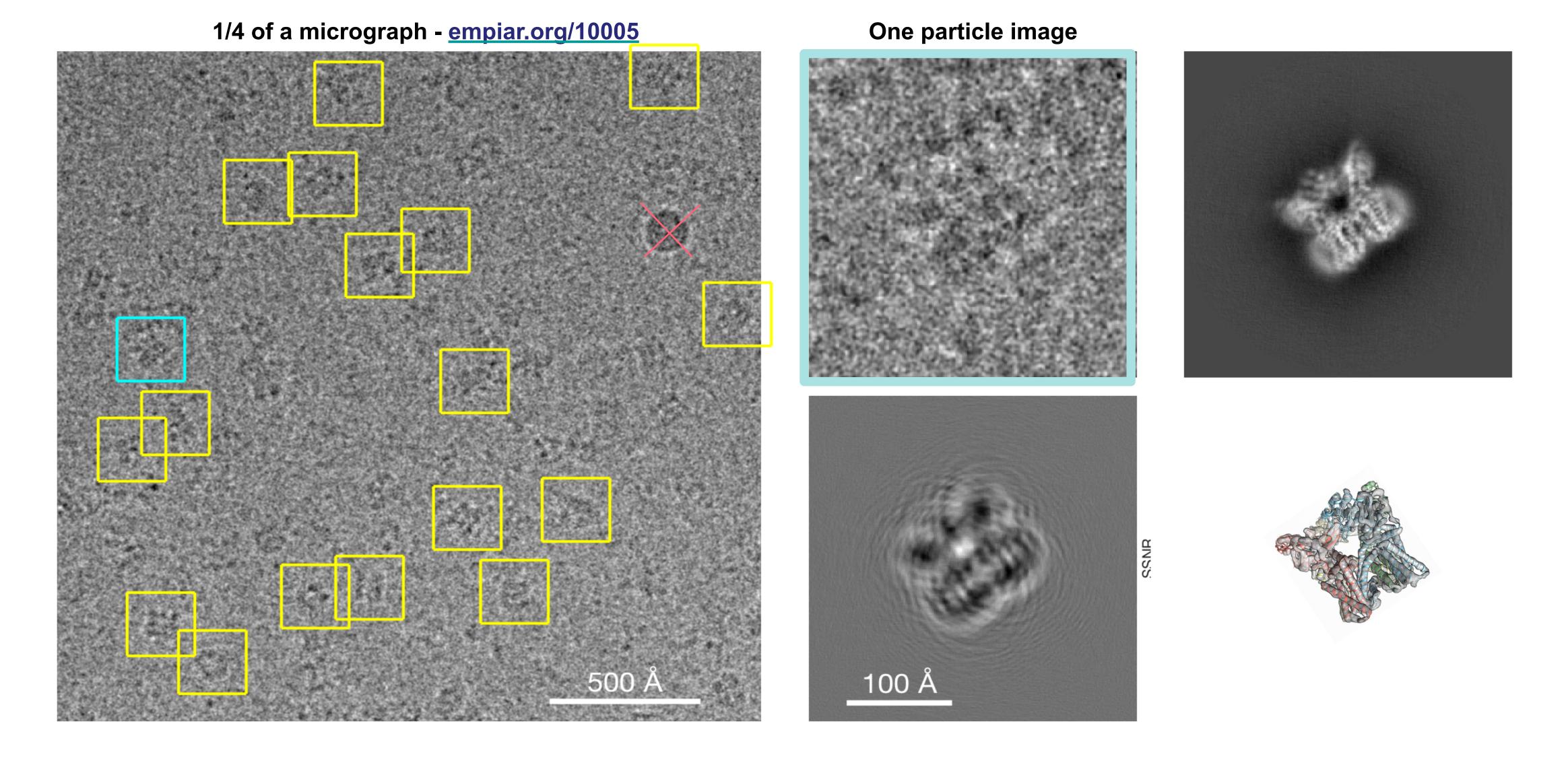
# Tomographic reconstruction: 2D image from 1D projections



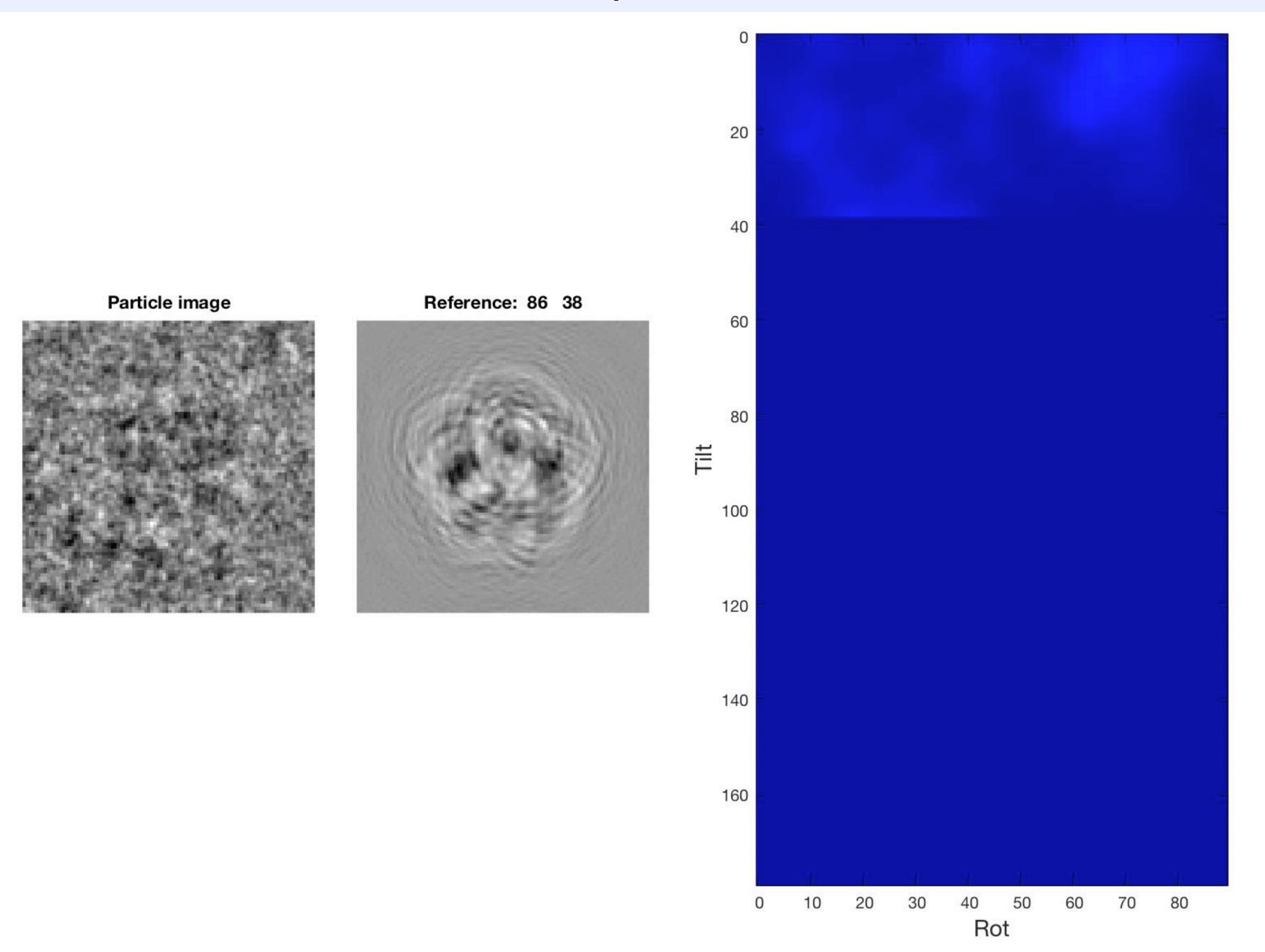
#### Determining the orientation angles: example from the TRPV1 dataset

# Structure of the TRPV1 ion channel determined by electron cryo-microscopy

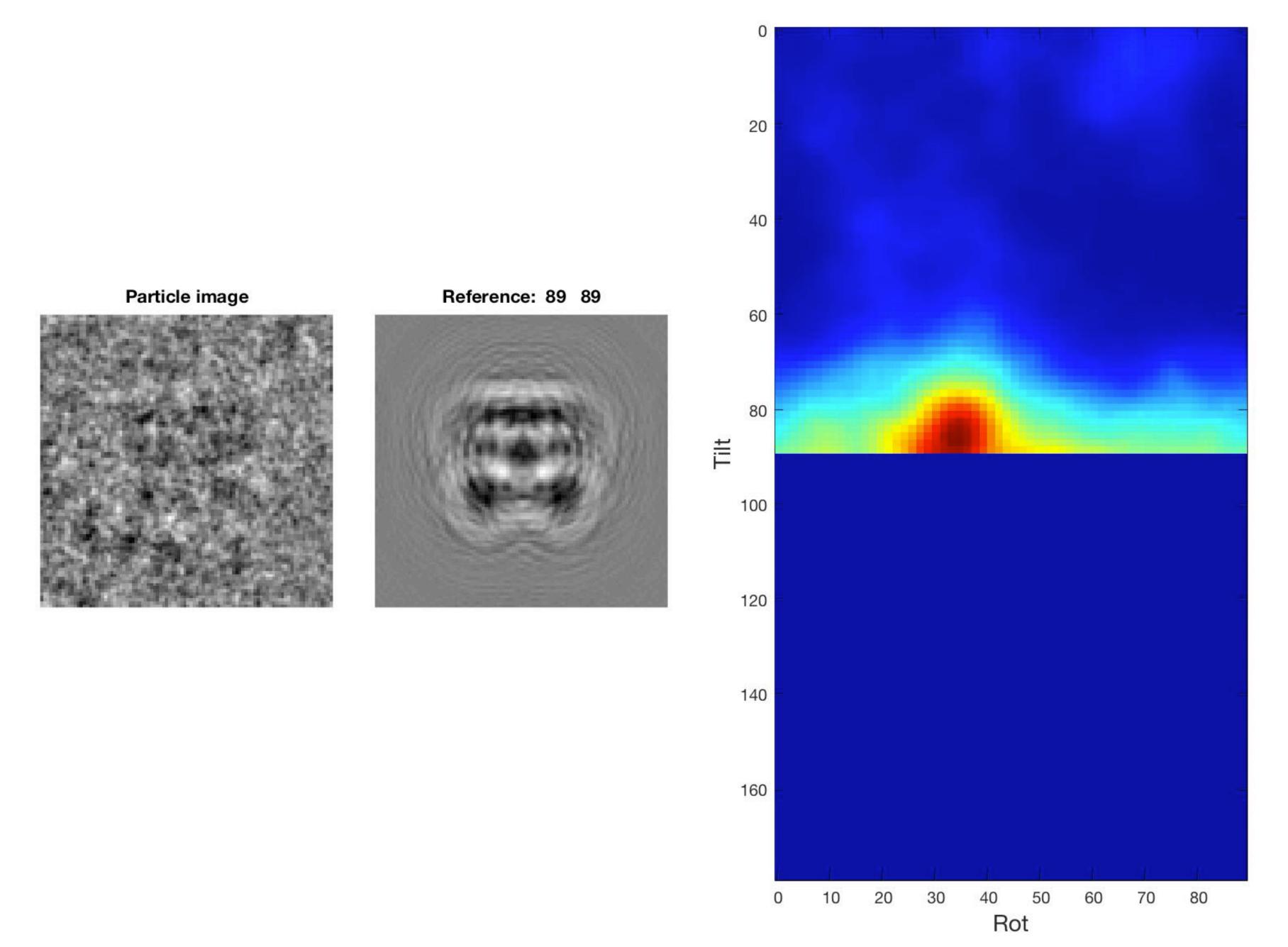
Maofu Liao¹\*, Erhu Cao²\*, David Julius² & Yifan Cheng¹



# The probability of orientations $P(\phi \mid X, V)$ is remarkably sharp



# The probability of orientations $P(\phi | X, V)$ is remarkably sharp

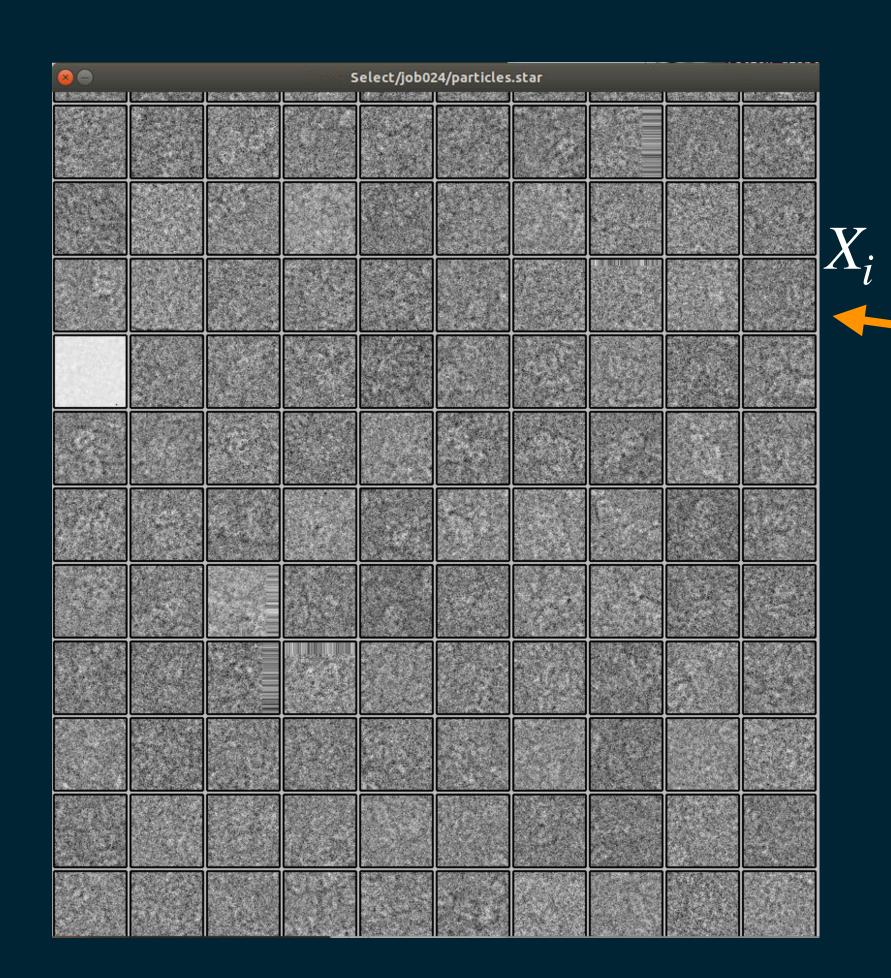


### Single-particle reconstruction

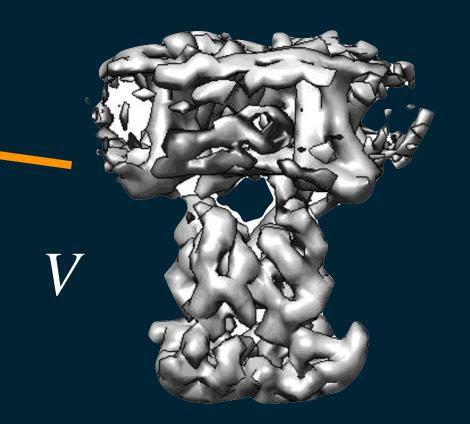
We assume that image  $X_i$  comes from a projection in direction  $\phi_i$  of volume V according to

$$X_i = C_i \mathbf{P}_{\phi_i} V + N_i$$

The goal is to discover the volume  ${\cal V}$ 



Project along  $\phi_i$ 



The first step is to compare images to determine orientations...

## There are various ways to compare images

Define the "reference" as the true image A modified by the CTF C:

$$R = CA$$

We wish to compare a data image X with it.

#### **Squared difference**

$$||X - R||^2 = \sum_{j} (X_j - R_j)^2$$
$$= ||X||^2 - 2X \cdot R + ||R||^2$$

#### Correlation

$$Cor = X \cdot R$$

$$= \sum_{j} X_{j}R_{j}$$

#### **Correlation coefficient**

$$CC = \frac{X \cdot R}{|X| |R|}$$

Notation used here:

A single pixel in the image X:

$$X_i$$
 —the  $j^{th}$  pixel (out of  $J$  pixels total)

The  $i^{th}$  image in the dataset X:

# First the 2D problem: reconstruct an image

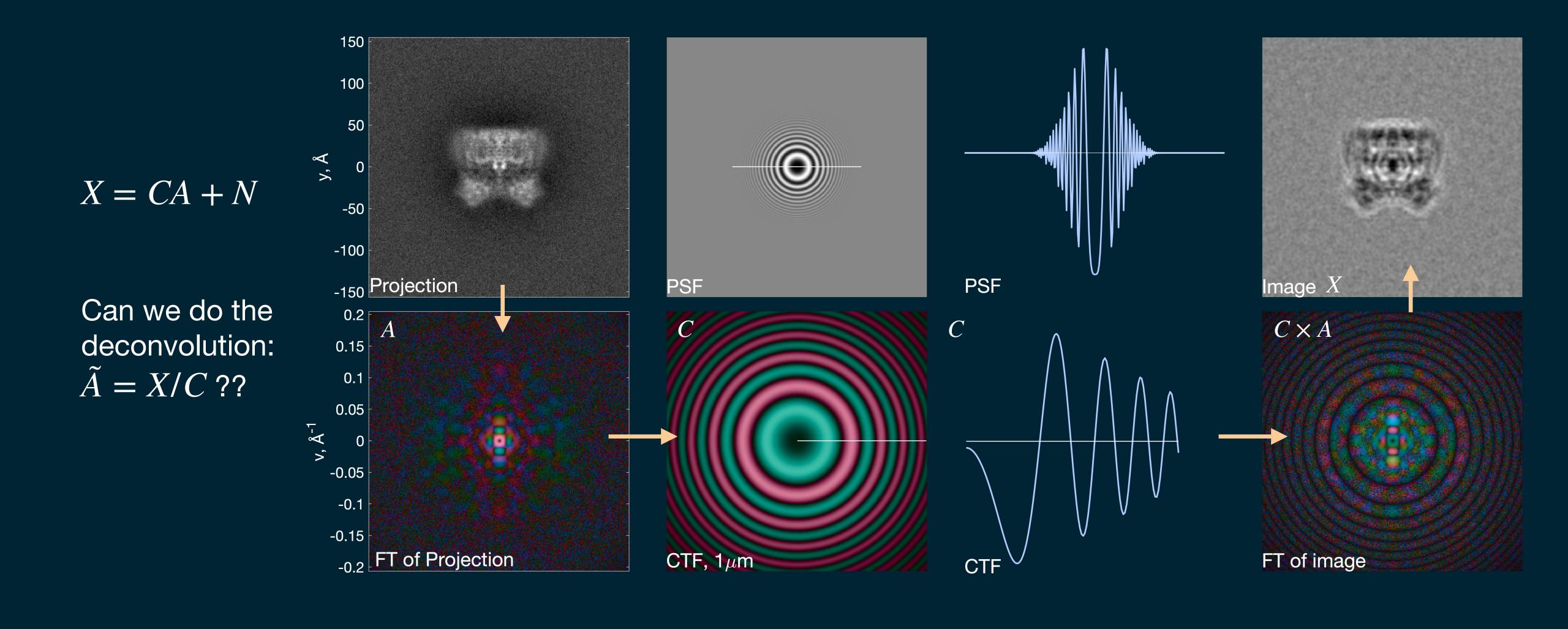
Model of an image

$$X = CA + N$$

- A "true" image
- C contrast-transfer function
- N noise image

We can interpret *C* as either the CTF operator (*x*,*y* space), or just the multiplicative CTF factor (*u*,*v* space)

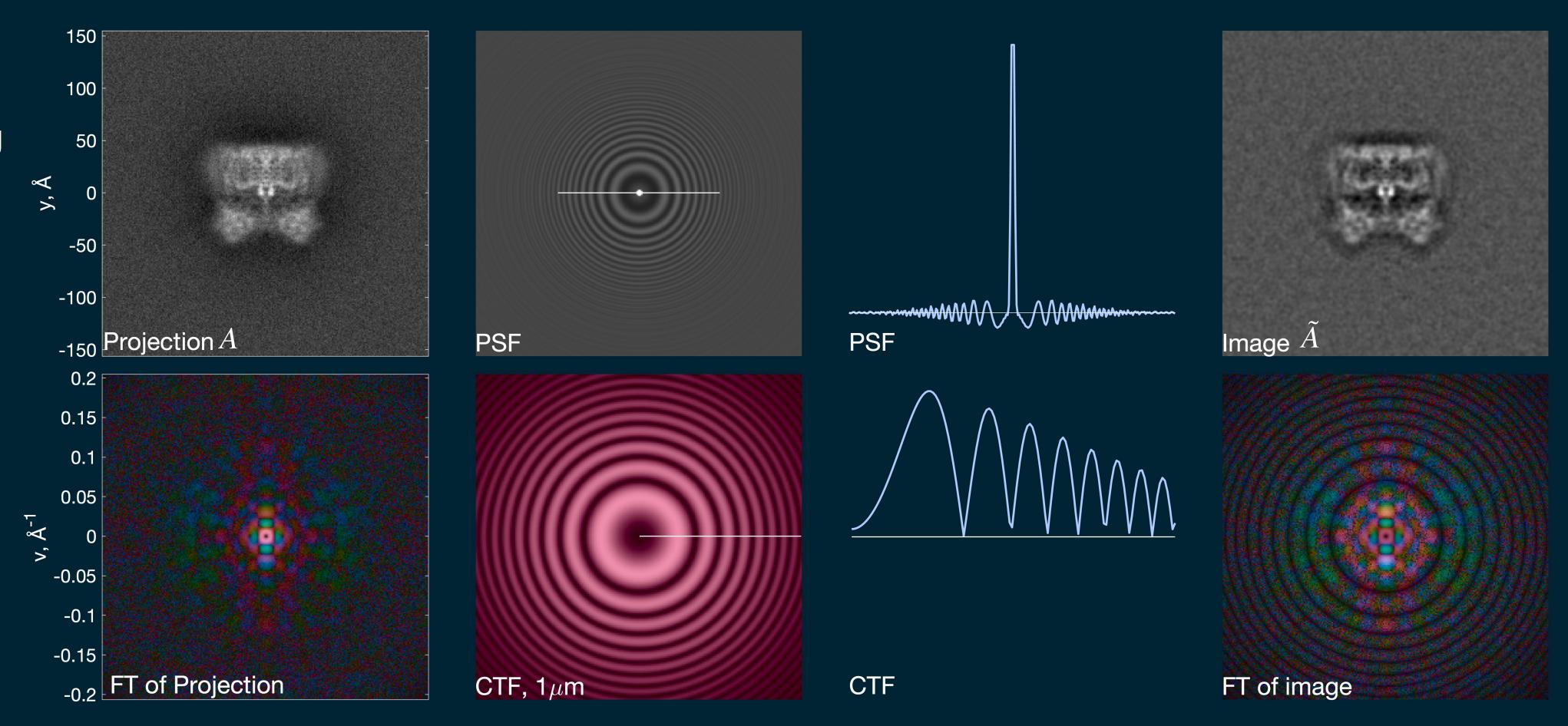
# Modeling the CTF effect on an image



# How to undo the CTF effects?

#### 1. Phase flipping

$$\tilde{A} = \operatorname{sgn}(C)X$$



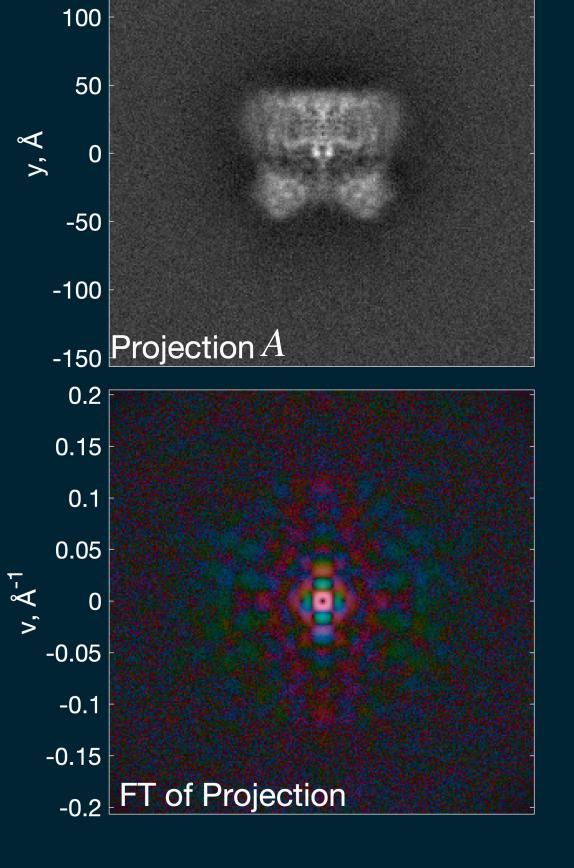
### How to undo the CTF effects?

#### 1. Phase flipping

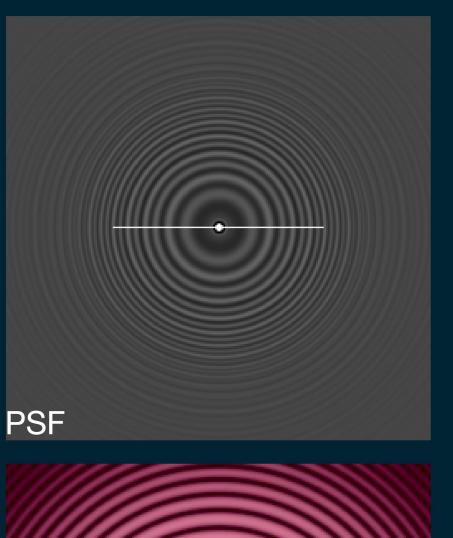
$$\tilde{A} = \operatorname{sgn}(C)X$$

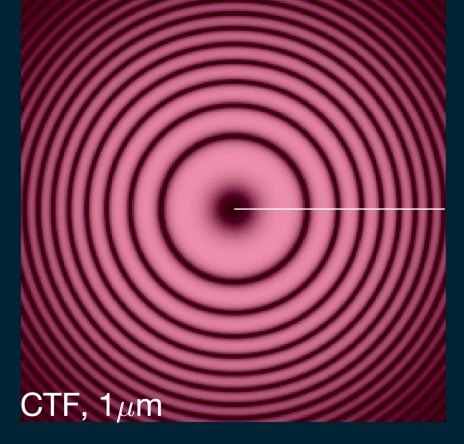
#### 2. Wiener filter

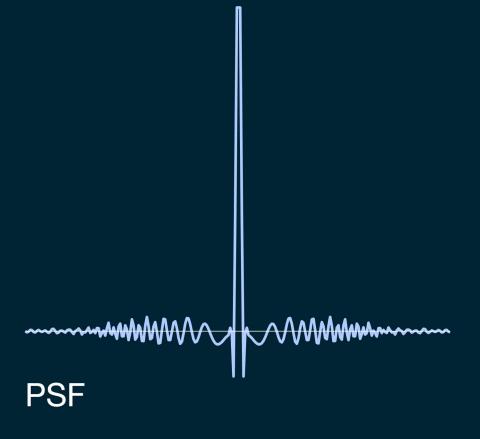
$$\tilde{A} = \frac{CX}{C^2 + k}$$

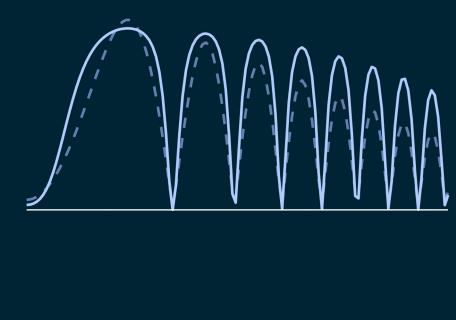


150

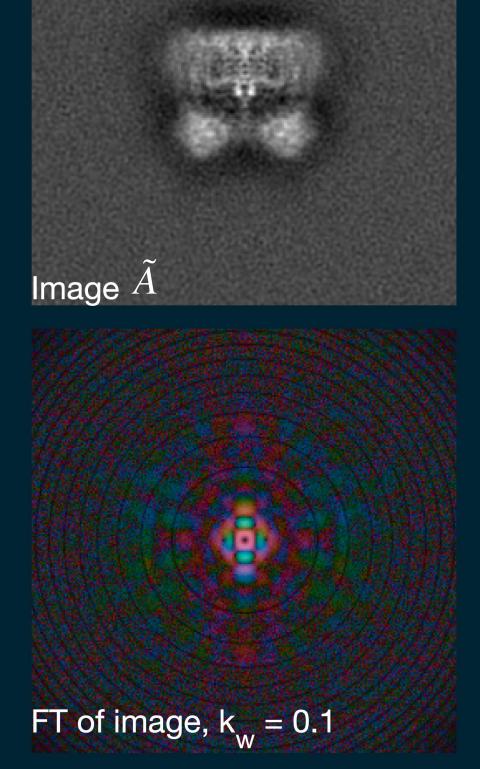








CTF



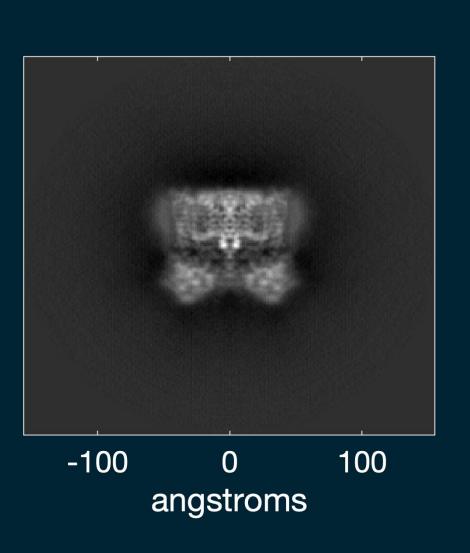
# How to undo the CTF effects in noisy images?

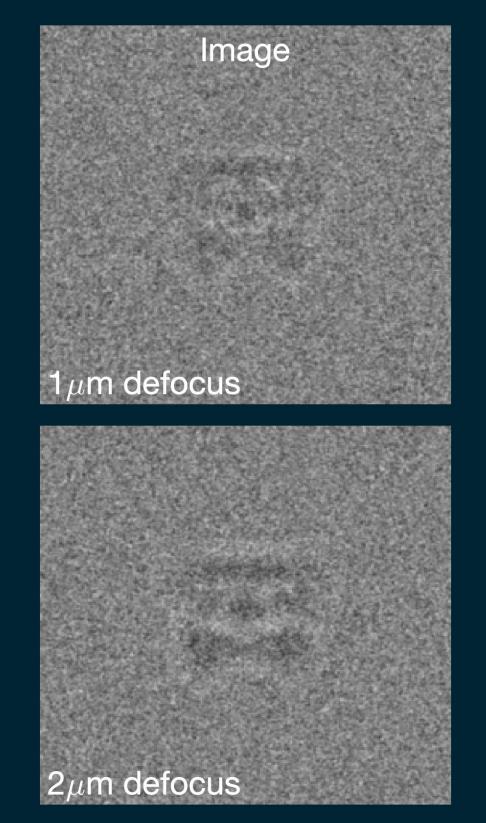
#### 1. Phase flipping

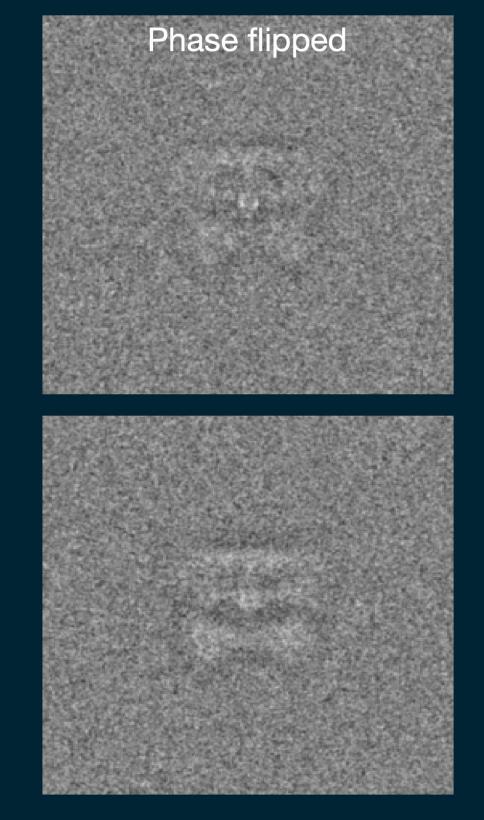
$$\tilde{A} = \operatorname{sgn}(C)X$$

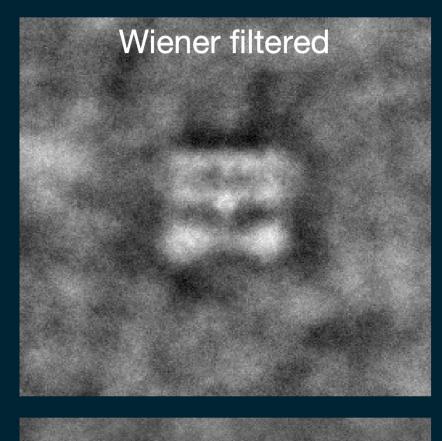
#### 2. Wiener filter

$$\tilde{A} = \frac{CX}{C^2 + k}$$



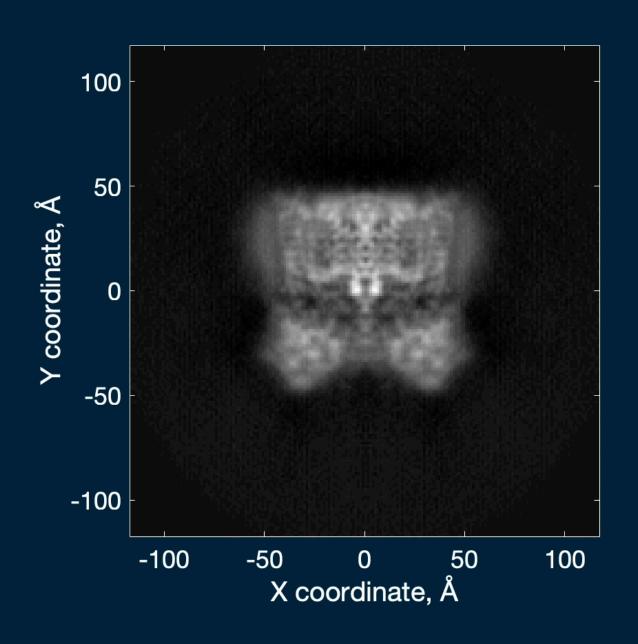


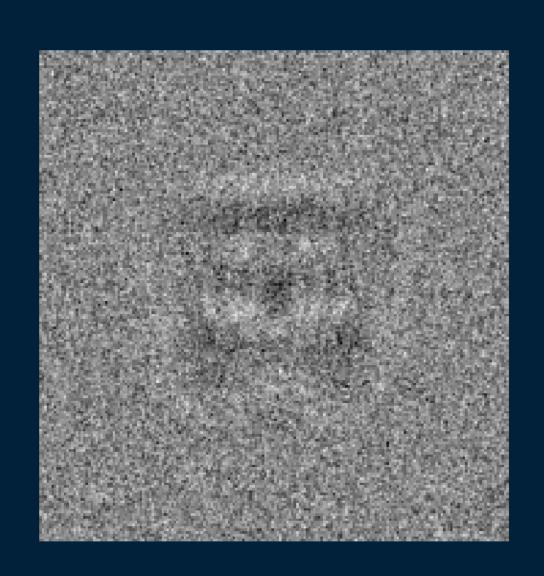


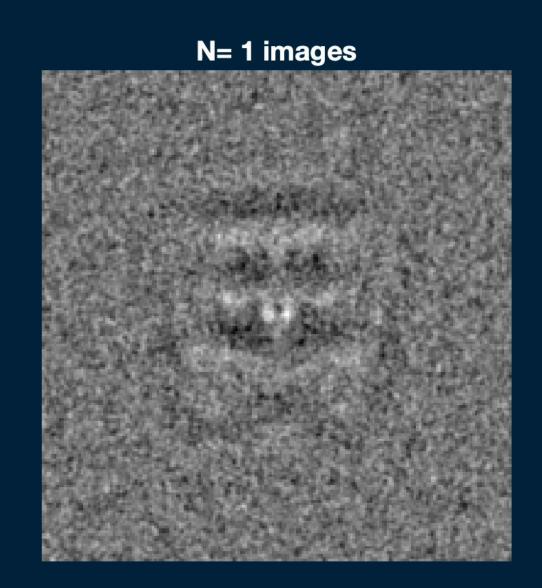




# How to undo the CTF effects in noisy images?



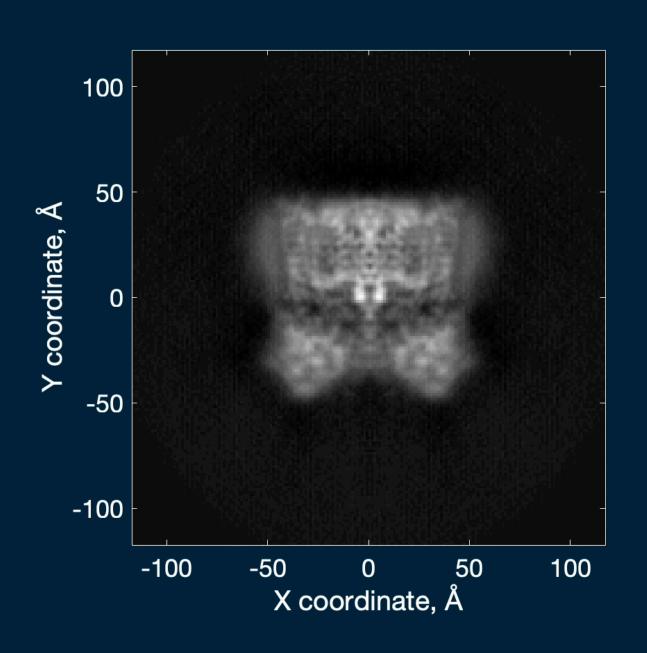


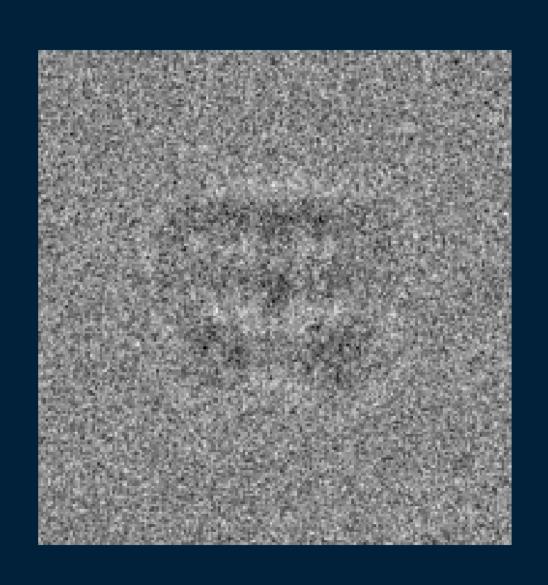


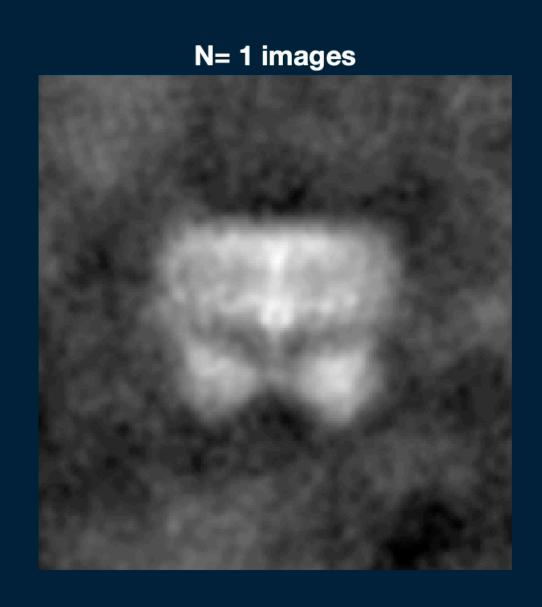
#### 3. Wiener from multiple images

$$\tilde{A} = \frac{\sum_{i}^{N} C_{i} X_{i}}{k + \sum_{i}^{N} C_{i}^{2}}$$

# How to undo the CTF effects in noisy images?







#### 3. Wiener from multiple images

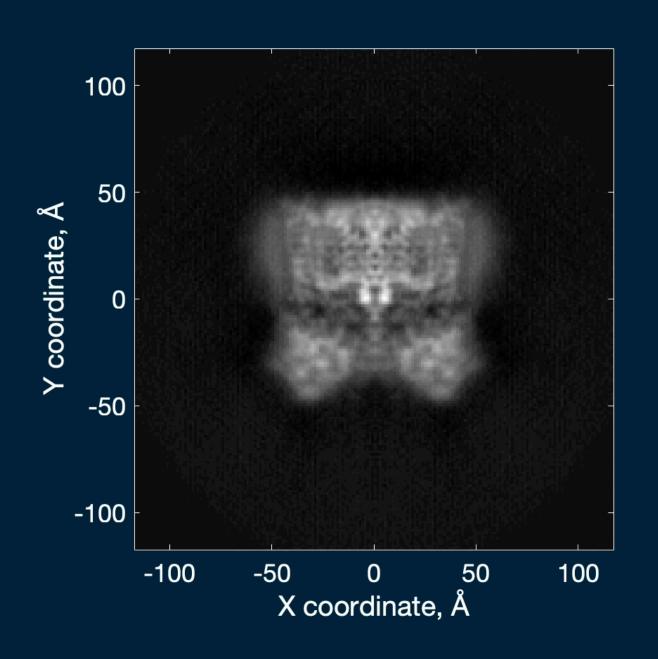
$$\tilde{A} = \frac{\sum_{i}^{N} C_{i} X_{i}}{k(s) + \sum_{i}^{N} C_{i}^{2}} \qquad k(s) = 1/\text{SNR}$$
$$= \frac{|N|^{2}}{|A|^{2}}$$

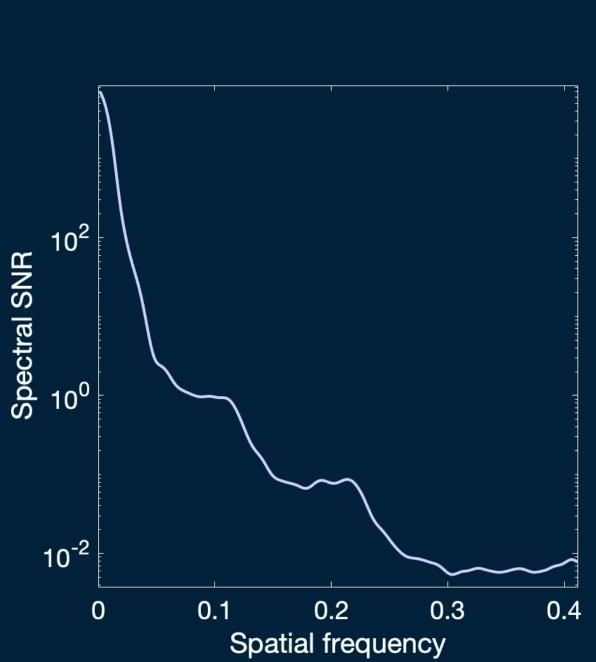
# Image restoration when spectral SNR is known

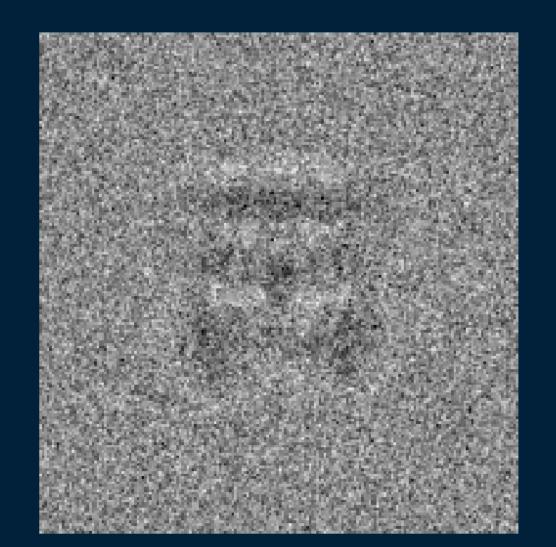
Restoration from multiple images

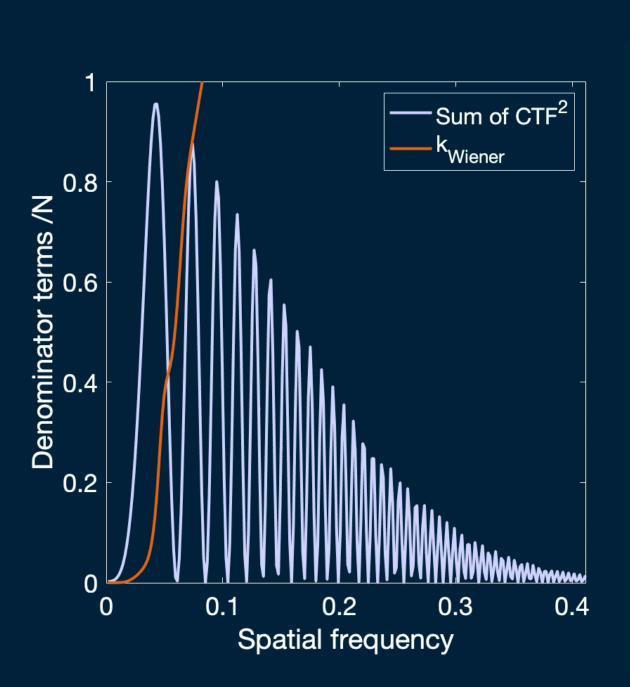
$$\tilde{A} = \frac{\sum_{i}^{N} C_{i} X_{i}}{\frac{1}{\text{SSNR}} + \sum_{i}^{N} C_{i}^{2}}$$

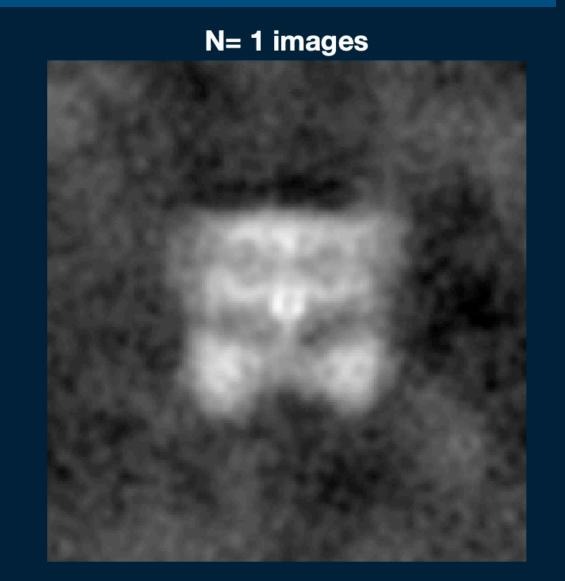
The defocus varies to fill in CTF zeros









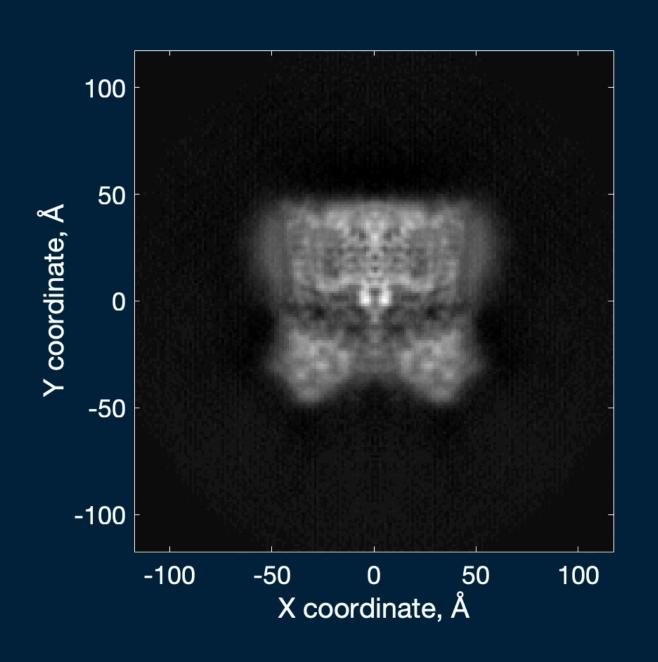


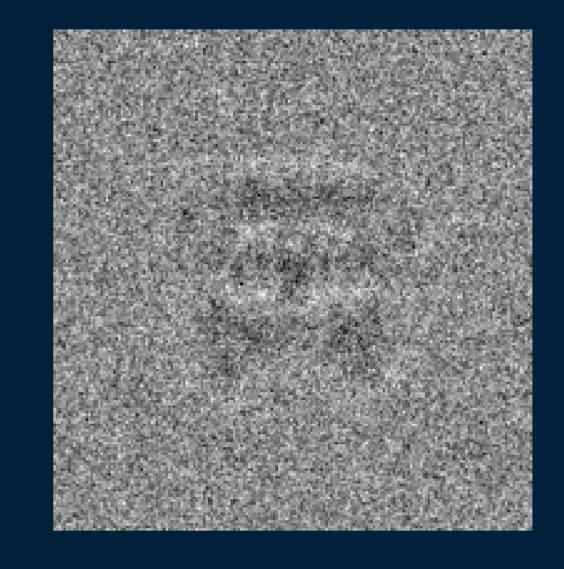
# Image restoration when spectral SNR is known

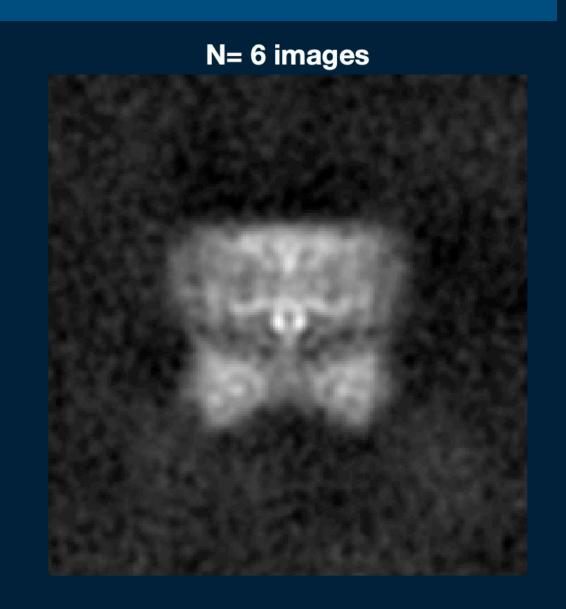
# Restoration from multiple images

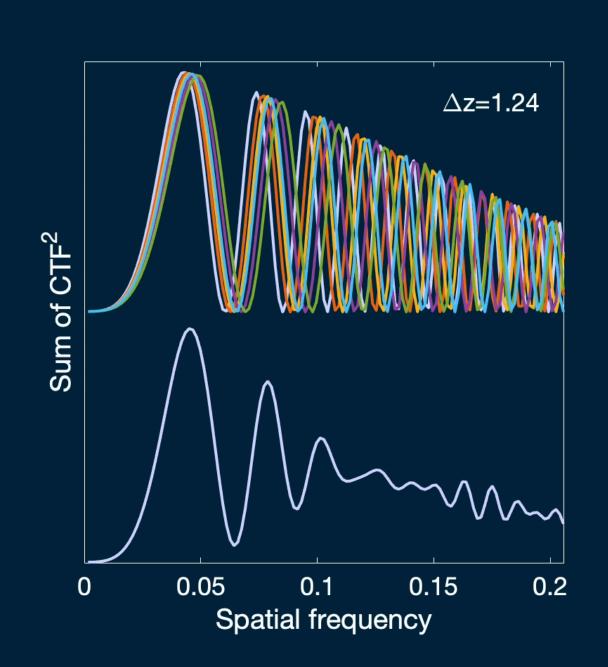
$$\tilde{A} = \frac{\sum_{i}^{N} C_{i} X_{i}}{\frac{1}{\text{SSNR}} + \sum_{i}^{N} C_{i}^{2}}$$

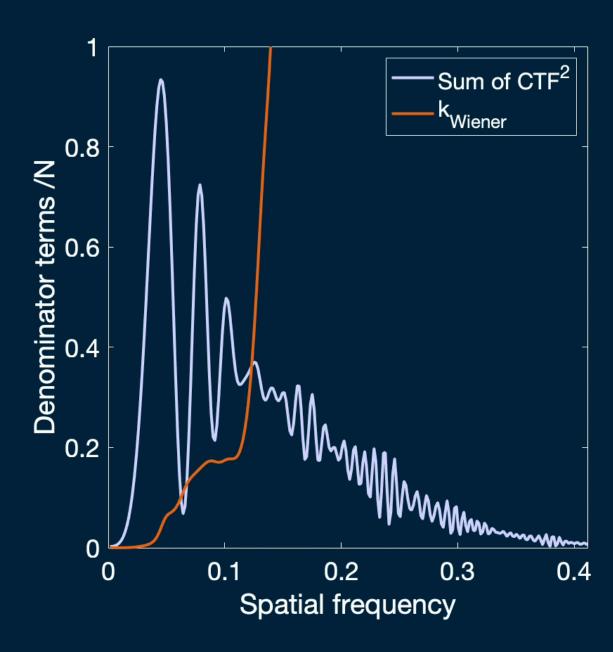
The defocus varies to fill in CTF zeros









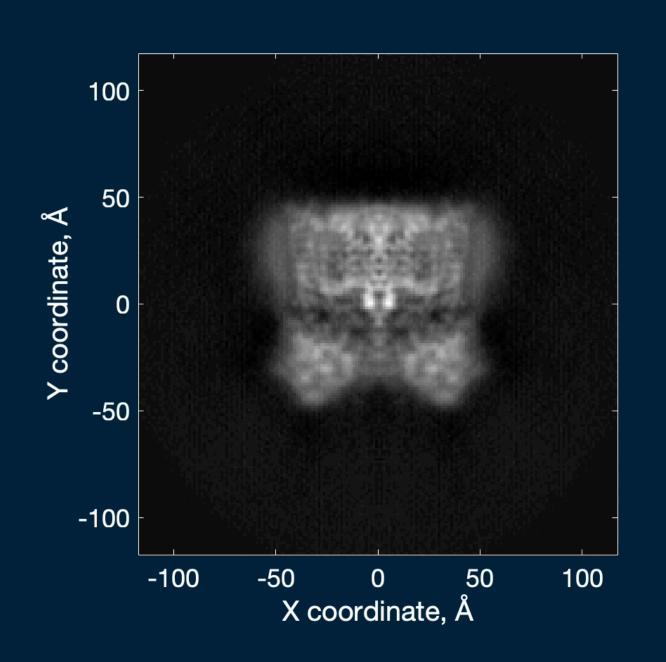


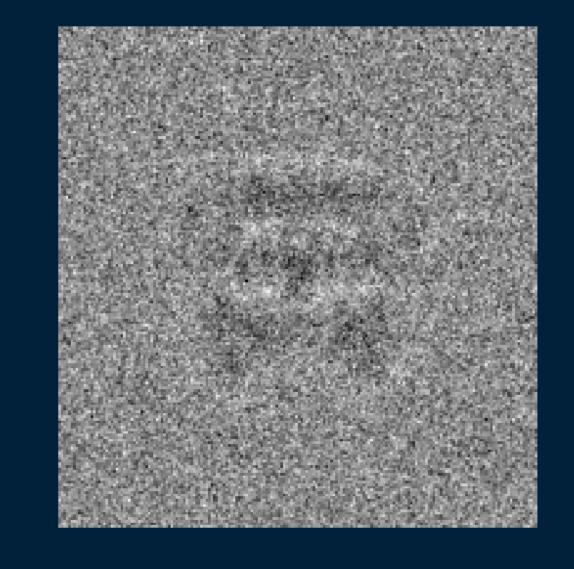
### Image restoration when spectral SNR is known

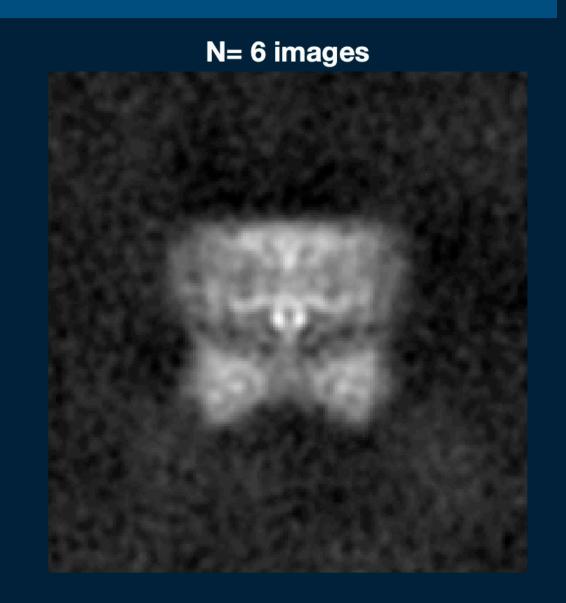
# Restoration from multiple images

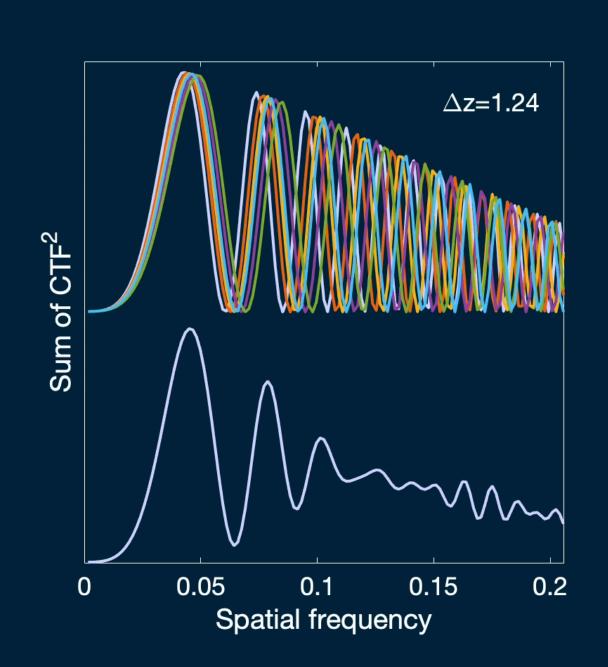
$$\tilde{A} = \frac{\sum_{i}^{N} C_{i} X_{i}}{\frac{1}{\text{SSNR}} + \sum_{i}^{N} C_{i}^{2}}$$

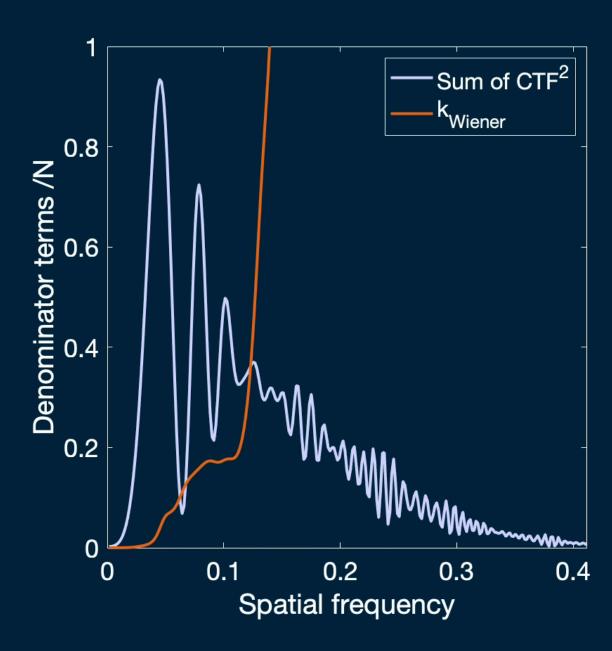
The defocus varies to fill in CTF zeros

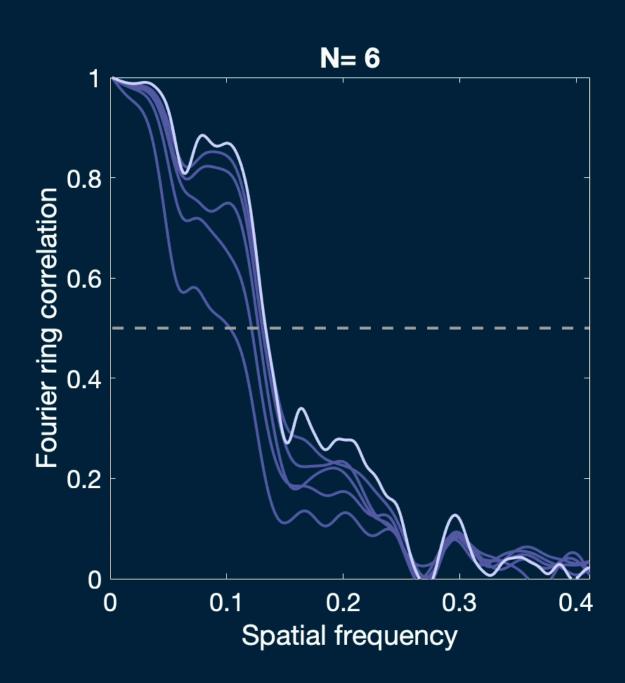












### 3D reconstruction in FREALIGN: correlation and Wiener filtering

A Frealign iteration, refining  $V^{(n)}$  to  $V^{(n+1)}$ , consists of two steps:

1. Vary the projection direction  $\phi_i$  to find the projection image  $R_i = C_i \mathbf{P}_{\phi_i} V^{(n)}$  that maximizes the correlation coefficient for each image  $X_i$ ,

$$CC = \frac{X_i \cdot R_i}{|X_i| |R_i|}.$$

2. Knowing the best projection direction  $\phi_i$  for each image  $X_i$ , update the volume according to

$$V^{(n+1)} = \frac{\sum_{i}^{N} \mathbf{P}_{\phi_{i}}^{\mathsf{T}} C_{i} X_{i}}{k + \sum_{i}^{N} \mathbf{P}_{\phi_{i}}^{\mathsf{T}} C_{i}^{2}}$$

#### <u>Notes</u>

- 1.  $C_i$  is the CTF corresponding to the image  $X_i$ .
- 2. The projection operator  $\mathbf{P}_{\phi}$  also includes translations. So  $\phi$  consists of five variables:  $\phi = \{\alpha, \beta, \gamma, t_x, t_y\}$ .
- 3.  $\mathbf{P}_{\phi_i}^{\mathbf{T}}$  is the corresponding <u>back</u> <u>projection</u> operator. In Fourier space it yields a volume that is all zeros except for values along a slice.
- 4. The sum

$$\sum_{i}^{N} \mathbf{P}_{\phi_{i}}^{\mathbf{T}} C_{i} X_{i}$$

is therefore the insertion of N slices.

### 3D reconstruction in FREALIGN—iterations

1. Start with a preliminary structure  $V^{(n)}$ , n=1

Iterate

2.For each particle image  $X_i$  find the projection angles  $\phi_i$  that gives the best match, so  $X_i \approx C_i \mathbf{P}_{\phi_i} V^{(n)}$ 

3.Use the Frealign iteration to produce a new 3D volume  $V^{(n+1)}$ 

#### 3D Classification in FREALIGN

Suppose our model is that an image X can come from any of K different particle types  $V_1, V_2, \ldots V_K$  and our images are selected randomly from these volumes, projected with noise added.

#### 1. The references are

$$R_{ik} = C_i \mathbf{P}_{\phi_i} V_k$$
.

We assign  $k_i$  such that  $V_{k_i}$  yields the projection (with direction  $\phi_i$ ) that gives the highest correlation coefficient with  $X_i$ .

2. Update the volume according to

$$V_k^{(n+1)} = \frac{\sum_{i \in k} \mathbf{P}_{\phi_i}^{\mathsf{T}} C_i X_i}{k_w + \sum_{i \in k} \mathbf{P}_{\phi_i}^{\mathsf{T}} C_i^2}$$

Maximum-likelihood methods

### Probabilities, another way to compare images

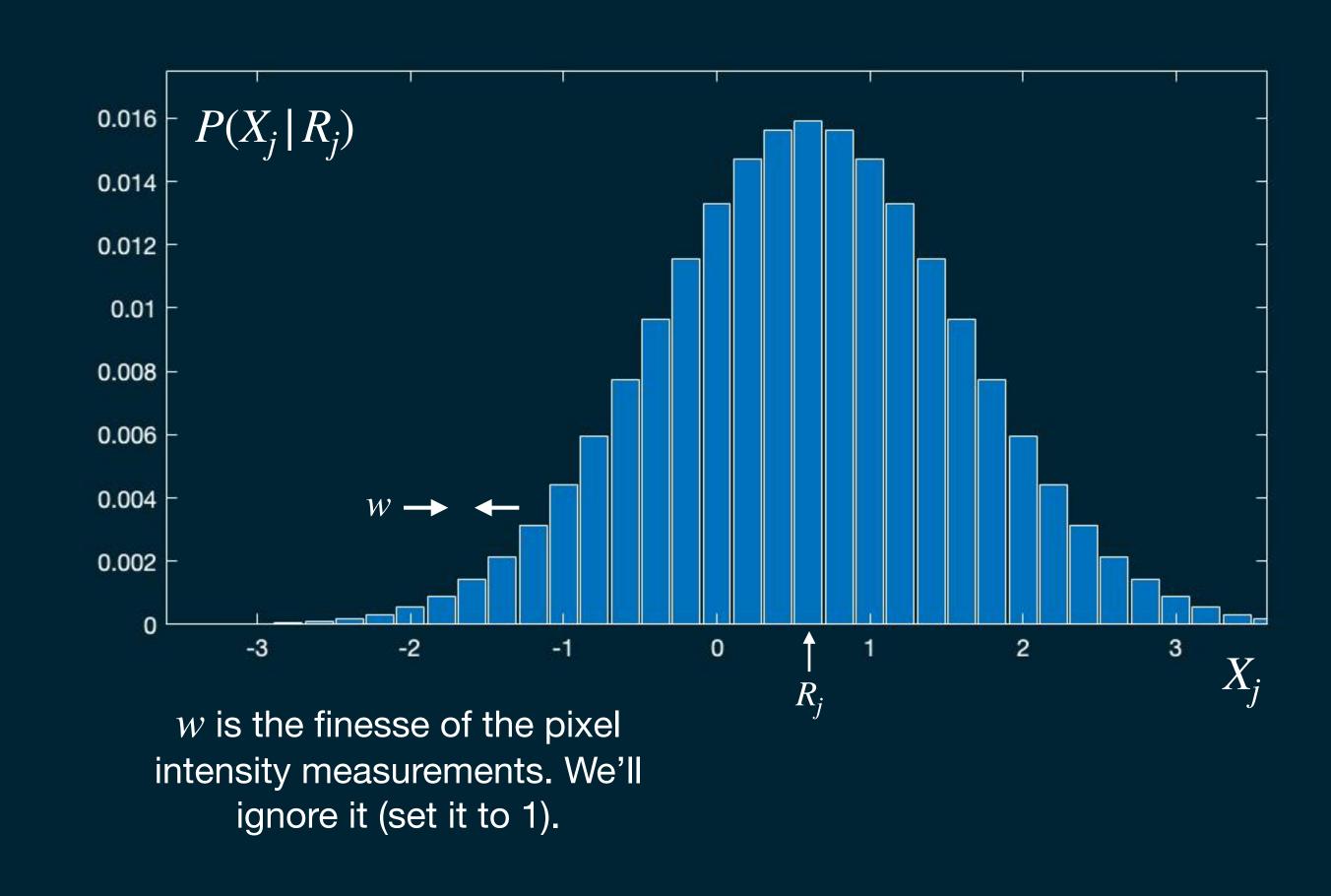
Image model: X = R + N

Probability of the jth pixel value:

$$P(X_j | R_j) = \frac{\sqrt{1}}{\sqrt{2\pi\sigma^2}} e^{-(X_j - R_j)^2/2\sigma^2}$$

Probability of observing an entire image that comes from *R*:

$$P(X|R) = \frac{\sqrt{1}}{(2\pi\sigma^2)^{J/2}} e^{-||X-R||^2/2\sigma^2}$$

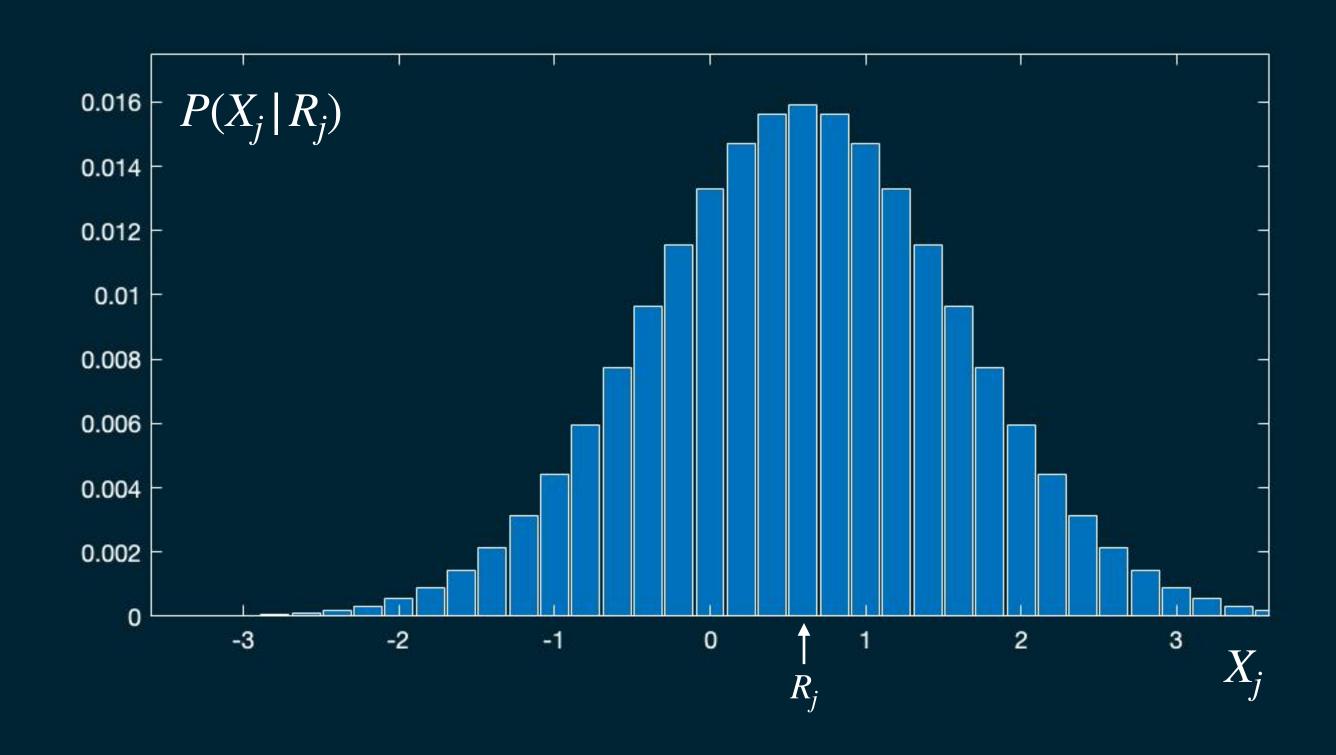


## Simplified image probability

$$X = R + N$$

Probability of observing an image that comes from *R*:

$$P(X|R) = c e^{-||X-R||^2/2\sigma^2}$$



#### The Likelihood

Let  $\mathbf{X} = \{X_1 ... X_N\}$  be our "stack" of particle images. We'd like to find the best 3D volume V consistent with these data, say maximizing the posterior probability

$$P(V|\mathbf{X})$$
.

According to Bayes' theorem,

$$P(V|\mathbf{X}) = P(\mathbf{X}|V) \frac{P(V)}{P(\mathbf{X})}.$$

*prior* → Experiment → *posterior likelihood* 

- $\cdot P(\mathbf{X})$  doesn't depend on V so we can ignore it.
- P(V) is called the <u>prior probability</u>. It reflects any knowledge about V that we have before considering the data set.
- $P(X \mid V)$  is something we can calculate. It's called the likelihood of V.

$$Lik(V) = P(X | V)$$

## We know how to compute the likelihood

We know that

$$P(X|V,\phi) = c e^{-\|X - \mathbf{CP}_{\phi}V\|^2/2\sigma^2}$$

To get the likelihood for one image we just integrate over all the  $\phi$ 's:

$$P(X|V) = \int P(X|V,\phi) P(\phi) d\phi,$$

assuming  $P(\phi)$  is uniform.

To get the likelihood for the whole dataset we compute the product over all the images,

$$P(\mathbf{X} | V) = \prod_{i}^{N} \int P(X_i | V, \phi) d\phi,$$

For numerical sanity, we compute the log likelihood,

$$L = \sum_{i}^{N} \ln \left( \int P(X_i | V, \phi) d\phi \right).$$

Maximum-likelihood reconstruction is finding V that maximizes L.

## Maximum-likelihood estimation is asymptotically unbiased

If the size of the dataset grows without bounds (and the number of parameters to be estimated does not) Maximum Likelihood converges to the right answer.

# To maximize the likelihood, we'll need a probability function $\Gamma(\phi)$

A projection

$$A = \mathbf{P}_{\phi}V$$

Probability of observing an image  $X_i$  if we know  $\phi$ :

$$P(X_i | V, \phi) = c e^{-||X_i - \mathbf{CP}_{\phi}V||^2/2\sigma^2}$$

Probability of a projection direction for  $X_i$ :

$$\Gamma_i(\phi) = P(\phi \mid X_i, V) = \frac{P(X_i \mid V, \phi)}{\int P(X_i \mid V, \phi) d\phi}$$

## The E-M algorithm finds a local maximum of the likelihood

The Expectation-Maximization (E-M) algorithm has this iteration, guaranteed to increase the likelihood:

$$V^{(n+1)} = \frac{\sum_{i} \int \Gamma_{i}^{(n)}(\phi) \mathbf{P}_{\phi}^{\mathsf{T}} C_{i} X_{i} d\phi}{\frac{\sigma^{2}}{T\tau^{2}} + \sum_{i} \int \Gamma_{i}^{(n)}(\phi) \mathbf{P}_{\phi}^{\mathsf{T}} C_{i}^{2} d\phi}$$

...Relion's compute-intensive "Expectation" step is basically the evaluation of  $\Gamma_i(\phi)$  for each image  $X_i$ , based on the current estimate  $V^{(n)}$  of the volume.

For comparison, here is Frealign's iteration:

- 1. Find the best orientation  $\phi_i$  for each particle image  $X_i$
- 2. Update the volume according to

$$V^{(n+1)} = \frac{\sum_{i} \mathbf{P}_{\phi_{i}}^{\mathbf{T}} C_{i} X_{i}}{k + \sum_{i} \mathbf{P}_{\phi_{i}}^{\mathbf{T}} C_{i}^{2}}$$

#### 3D Classification

We can use Expectation-Maximization to optimize K different volumes  $V_1, V_2, \ldots V_K$  simultaneously. The formula is essential the same except that the function  $\Gamma$  depends also on k:

$$\Gamma^{(n)}_{\phi_i,k}$$

The iteration, guaranteed to increase the likelihood:

$$V_k^{(n+1)} = \frac{\sum_i \int \Gamma_{i,k}^{(n)}(\phi) \mathbf{P}_{\phi}^{\mathsf{T}} C_i X_i \, d\phi}{\frac{\sigma^2}{T\tau^2} + \sum_i \int \Gamma_{i,k}^{(n)}(\phi) \mathbf{P}_{\phi}^{\mathsf{T}} C_i^2 \, d\phi}$$

...Relion's compute-intensive "Expectation" step is basically the evaluation of  $\Gamma_{i,k}(\phi)$  for each image  $X_i$  and volume  $V_k$ 

For comparison, here is Frealign's iteration:

- **1.** Find the best orientation  $\phi_i$  for each particle image  $X_i$
- 2. Update the volume according to

$$V^{(n+1)} = \frac{\sum_{i} \mathbf{P}_{\phi_i}^{\mathsf{T}} C_i X_i}{k + \sum_{i} \mathbf{P}_{\phi_i}^{\mathsf{T}} C_i^2}$$

#### The orientation determination is the most expensive step

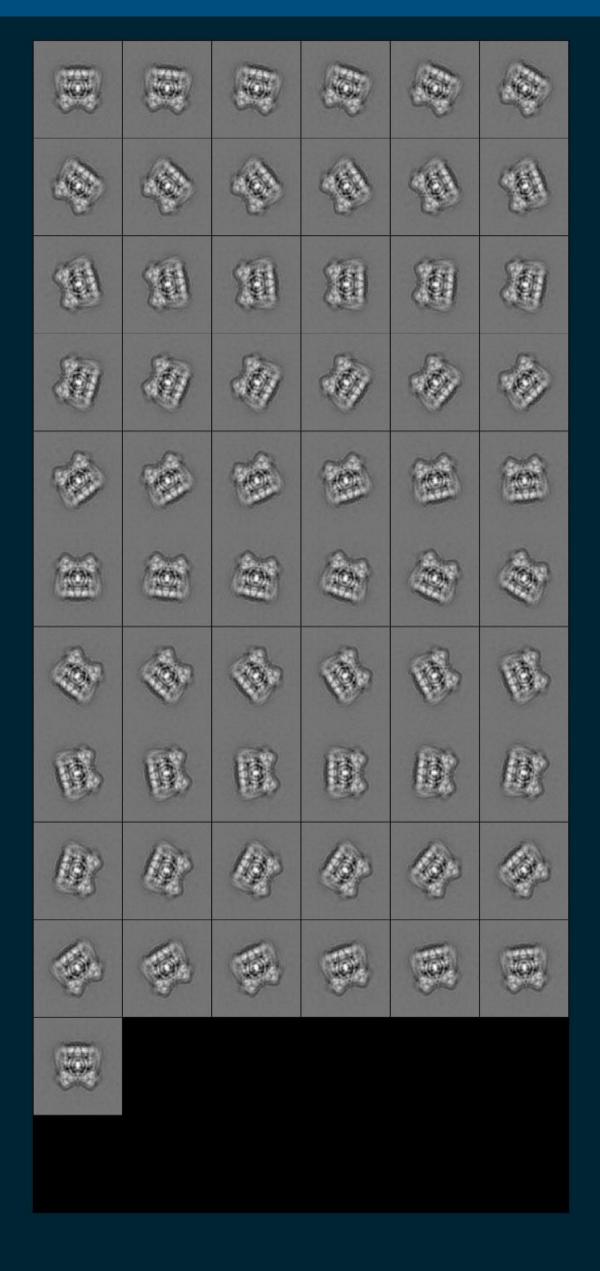
No. operations 
$$\approx \frac{\pi^3}{8} t^2 n^5 N + \pi n^4 + N n^2$$
Solution finding orientations and struction orientations

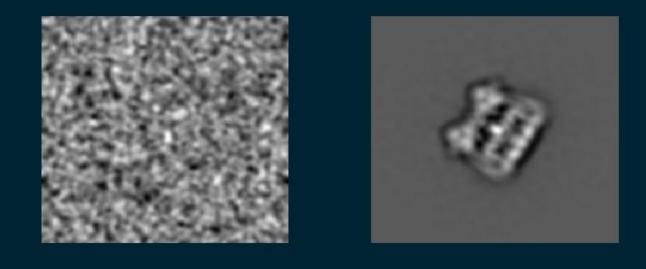
#### The orientation determination is the most expensive step

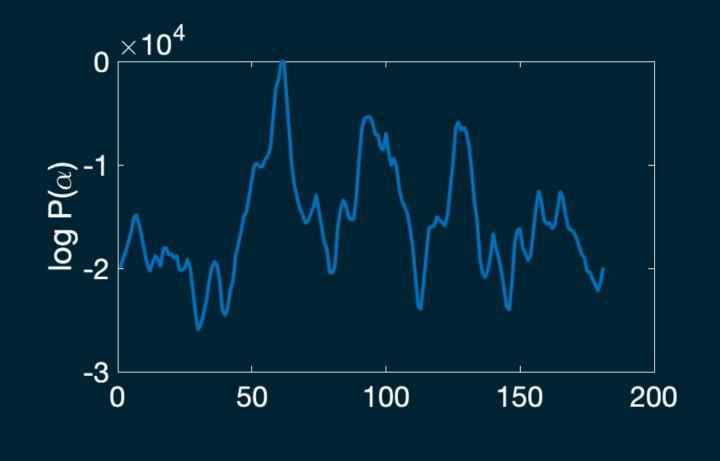
No. operations 
$$\approx \frac{\pi^3}{8} t^2 n^5 N + \pi n^4 + N n^2$$
Solution orientations and struction orientations  $\approx \frac{\pi^3}{8} t^2 n^5 N + \pi n^4 + N n^2$ 

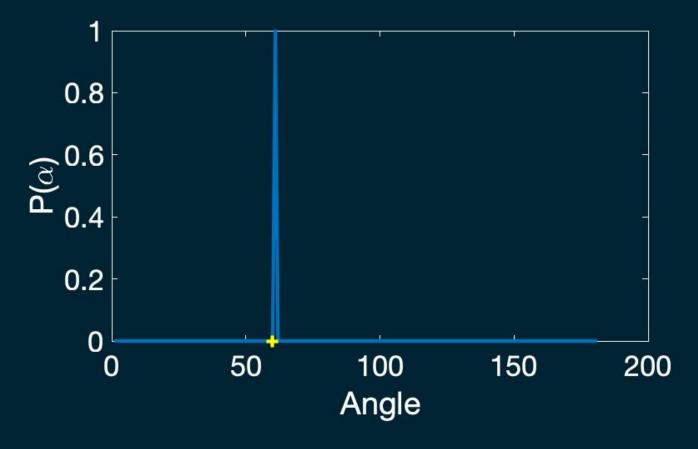
No. operations  $\approx 6 \times 10^{17} \approx 19$  CPU-years

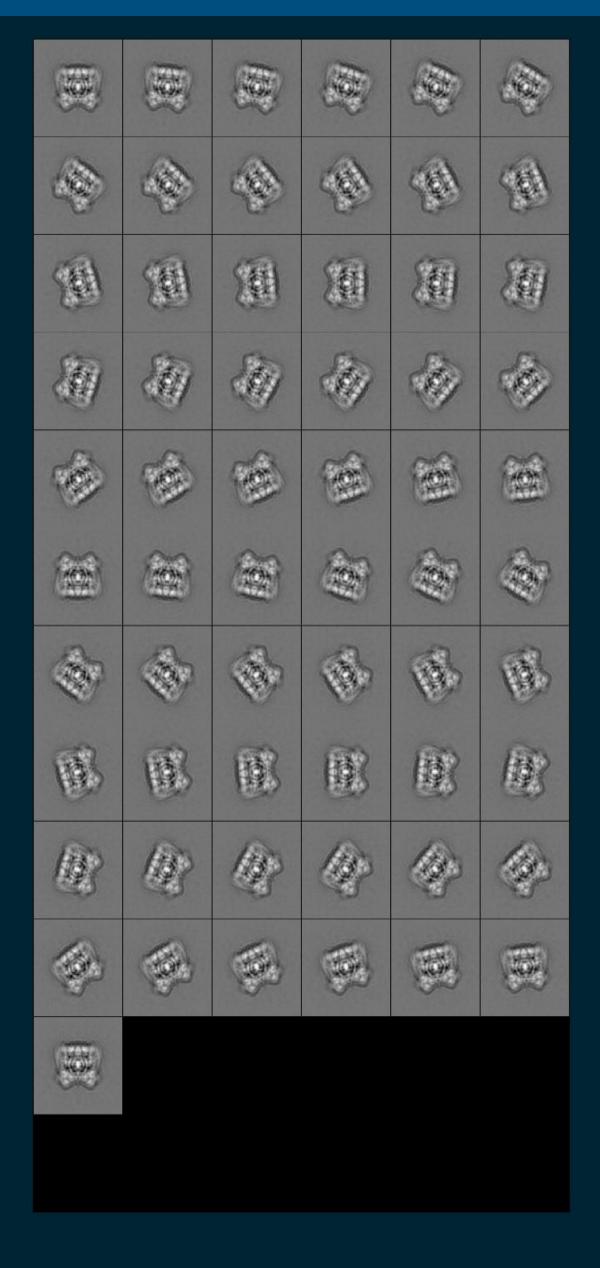
With efficient programs, ~ 1 CPU-day

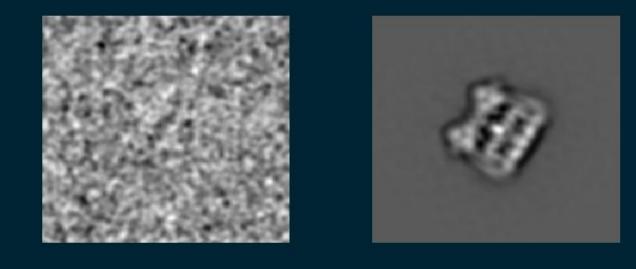


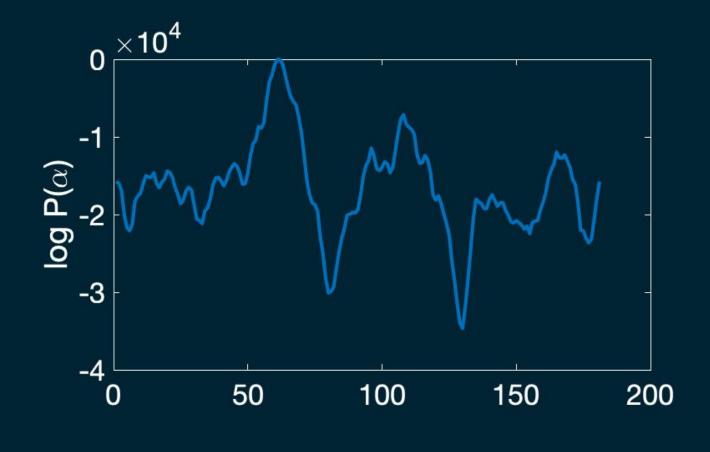


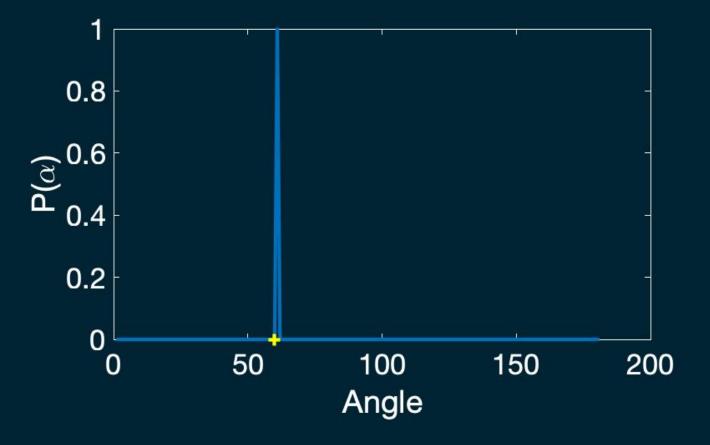


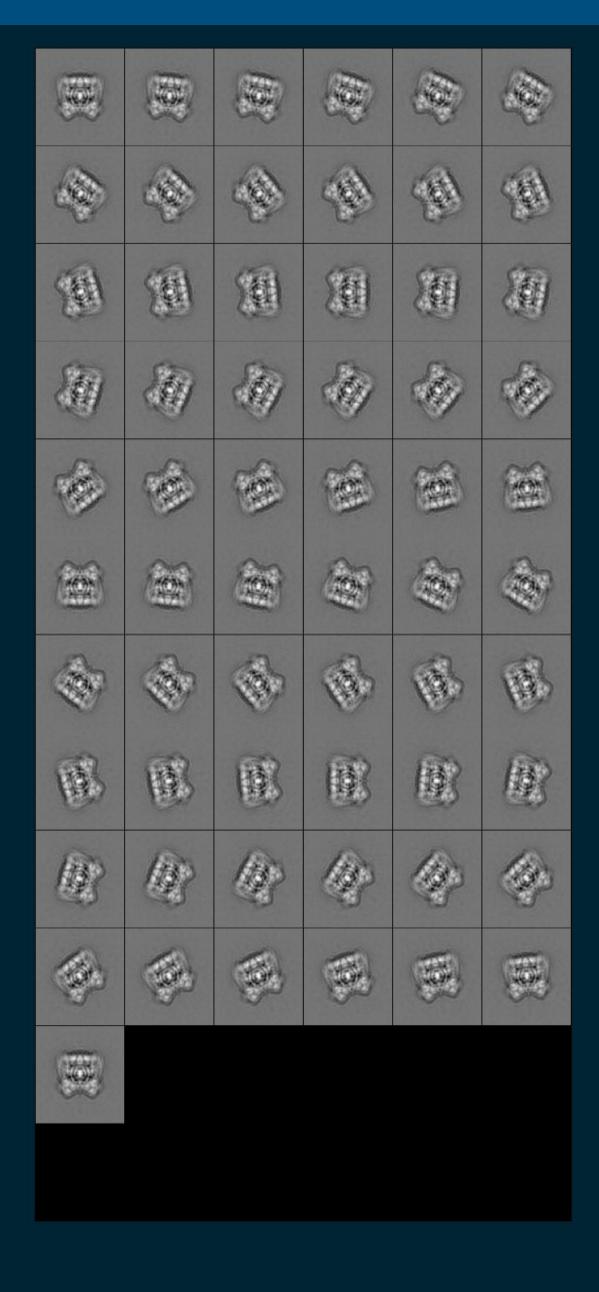


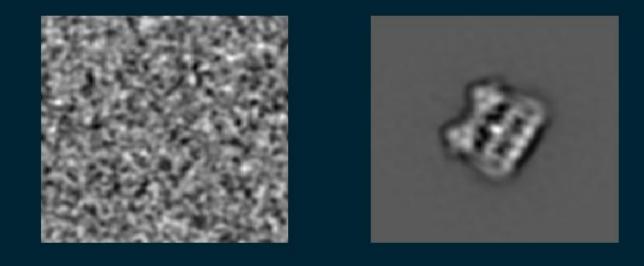


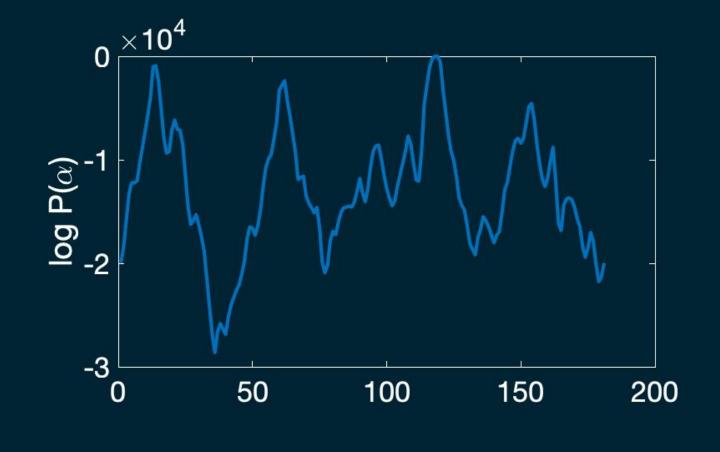


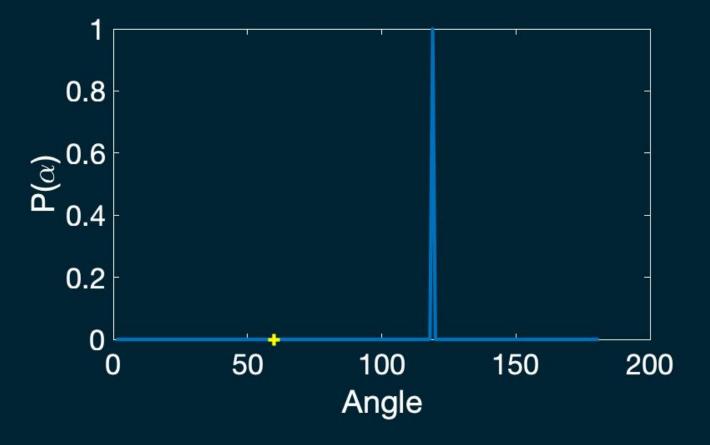




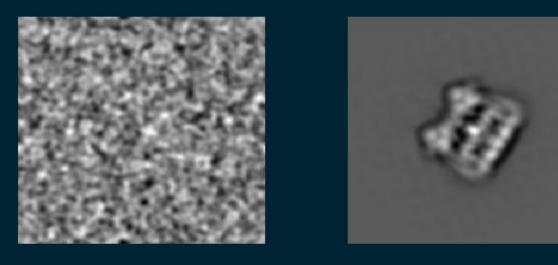


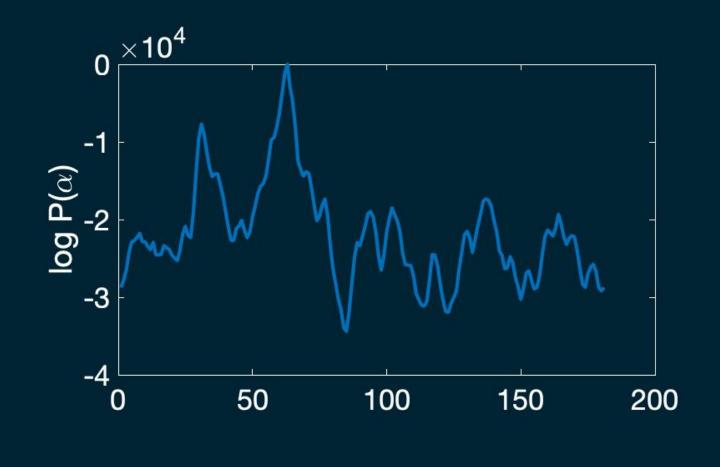


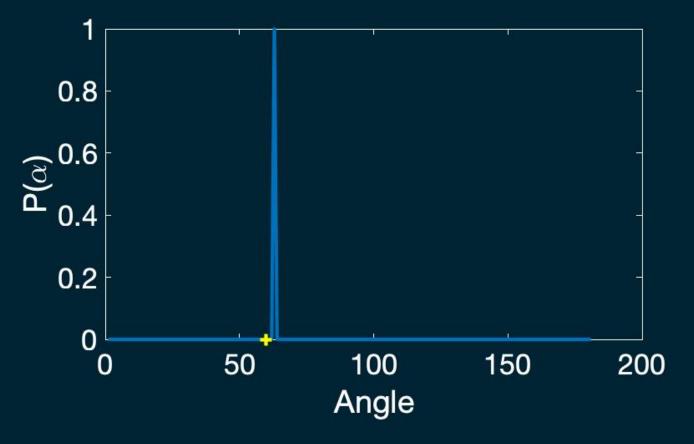




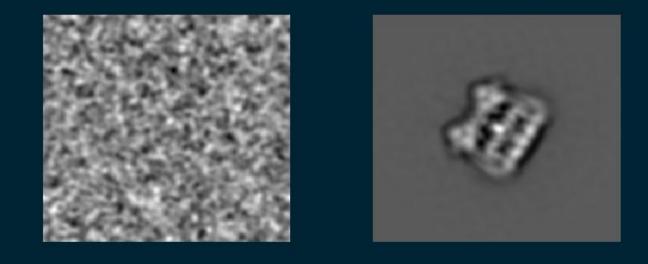


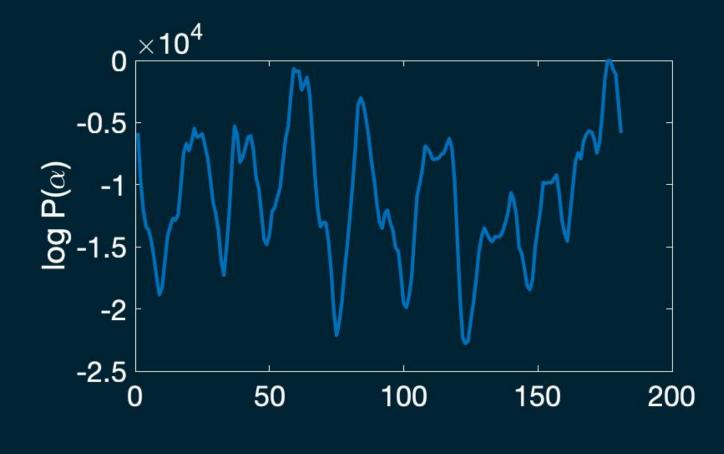


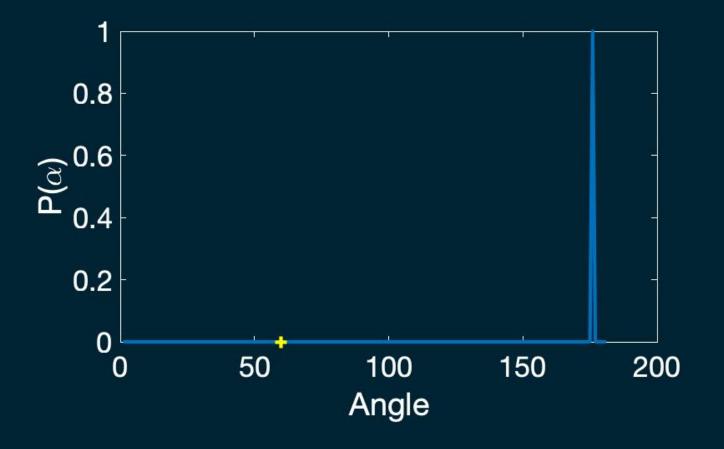






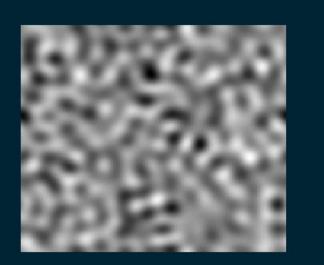






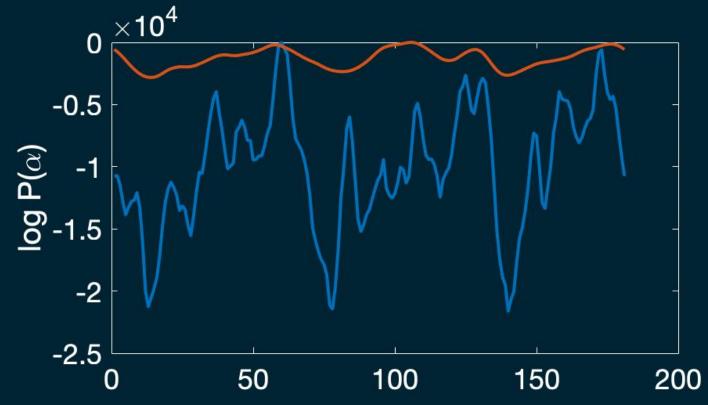
How to decrease the effort?

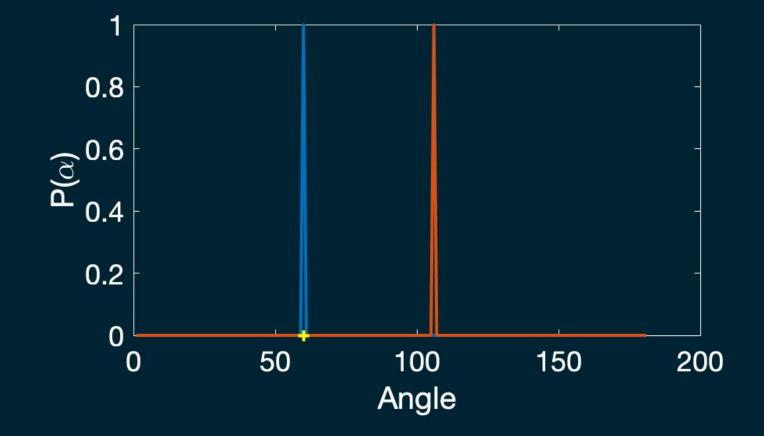
## Domain reduction: branch and bound, illustrated for 1D





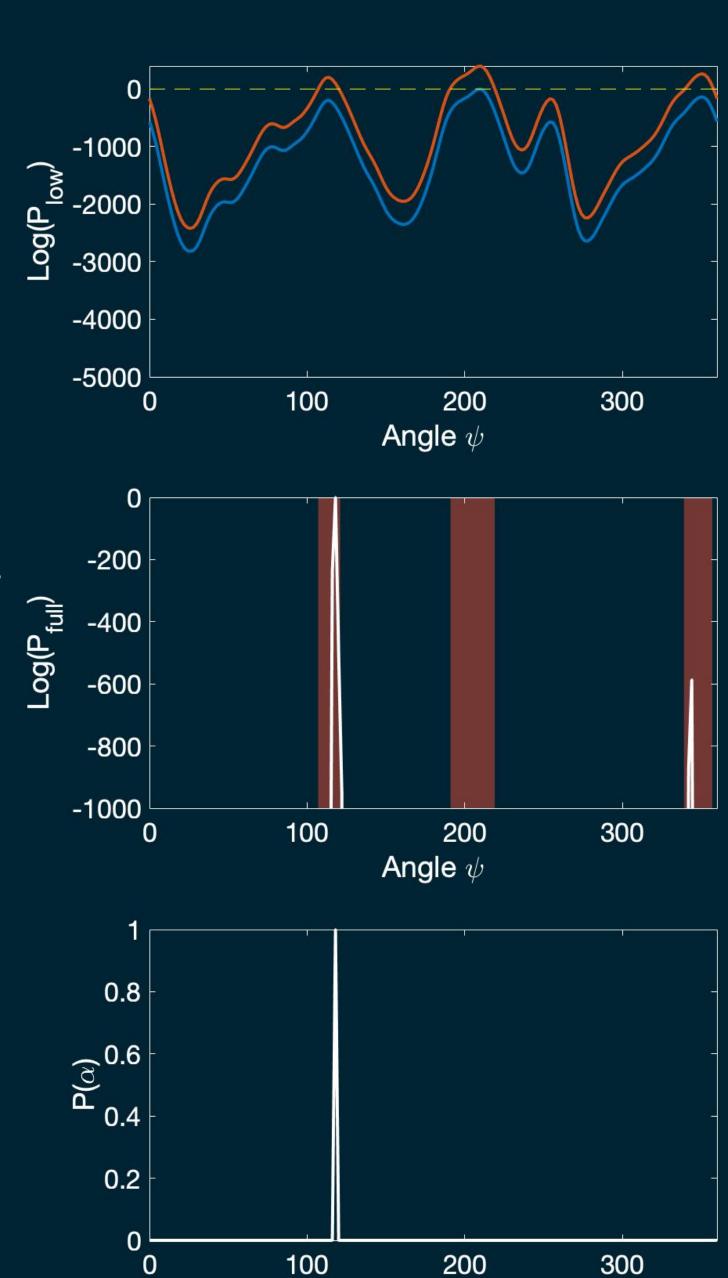
1. To save time, we compute probabilities of orientations at low resolution.





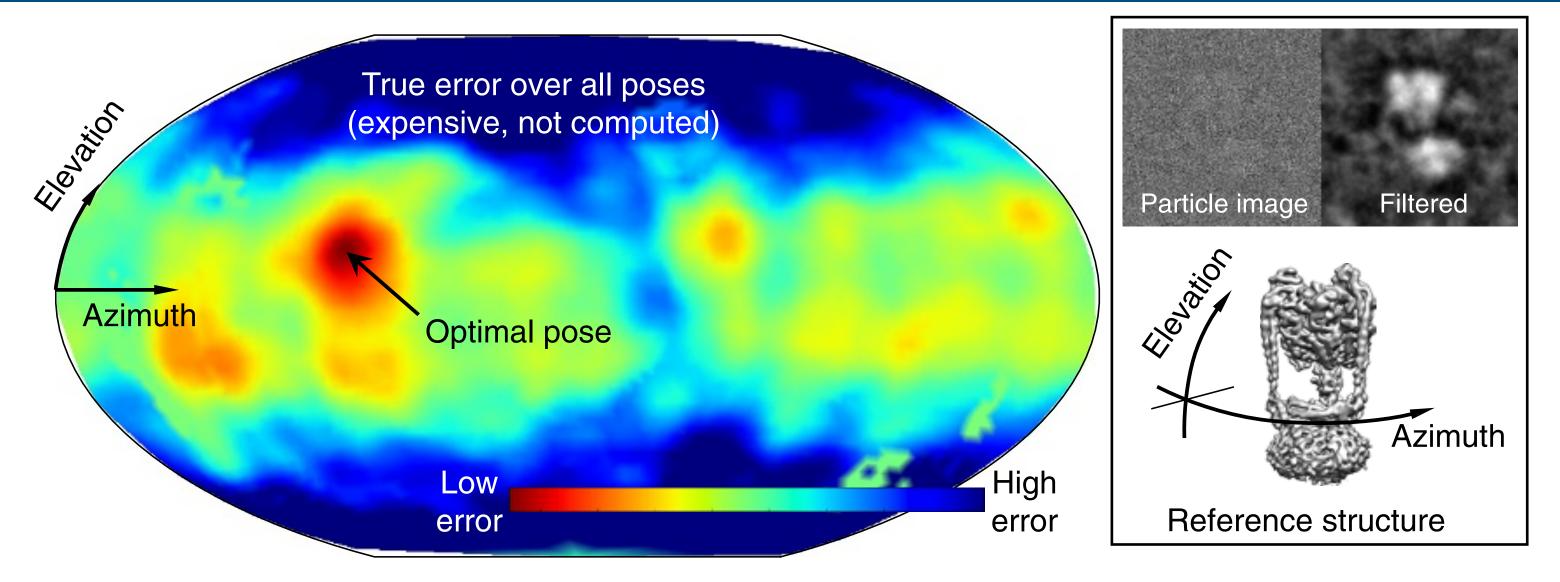
2. We place bounds on how much higher the probabilities could be at full resolution.

Given a cutoff value, we evaluate over a fraction of the domain.

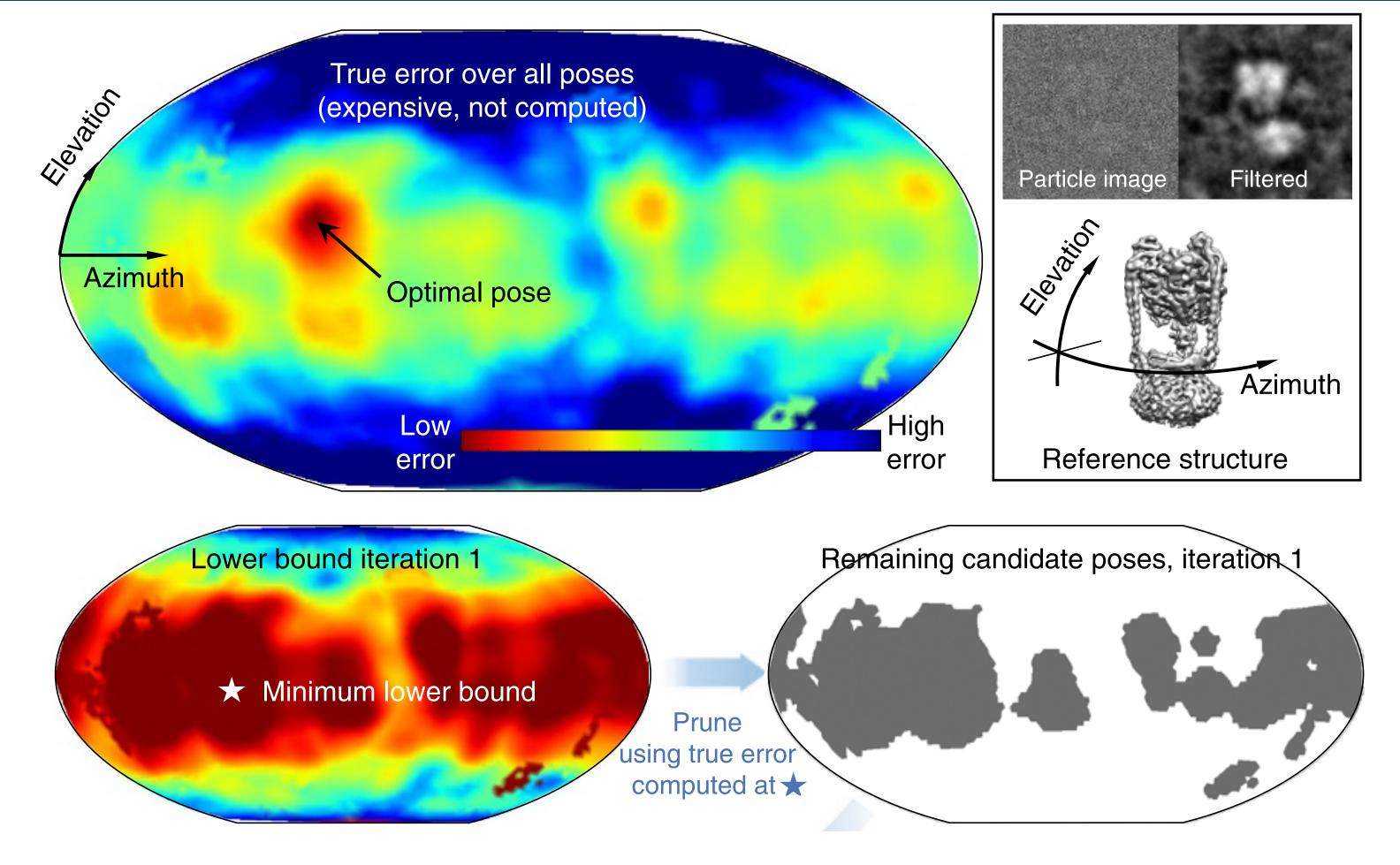


Angle  $\psi$ 

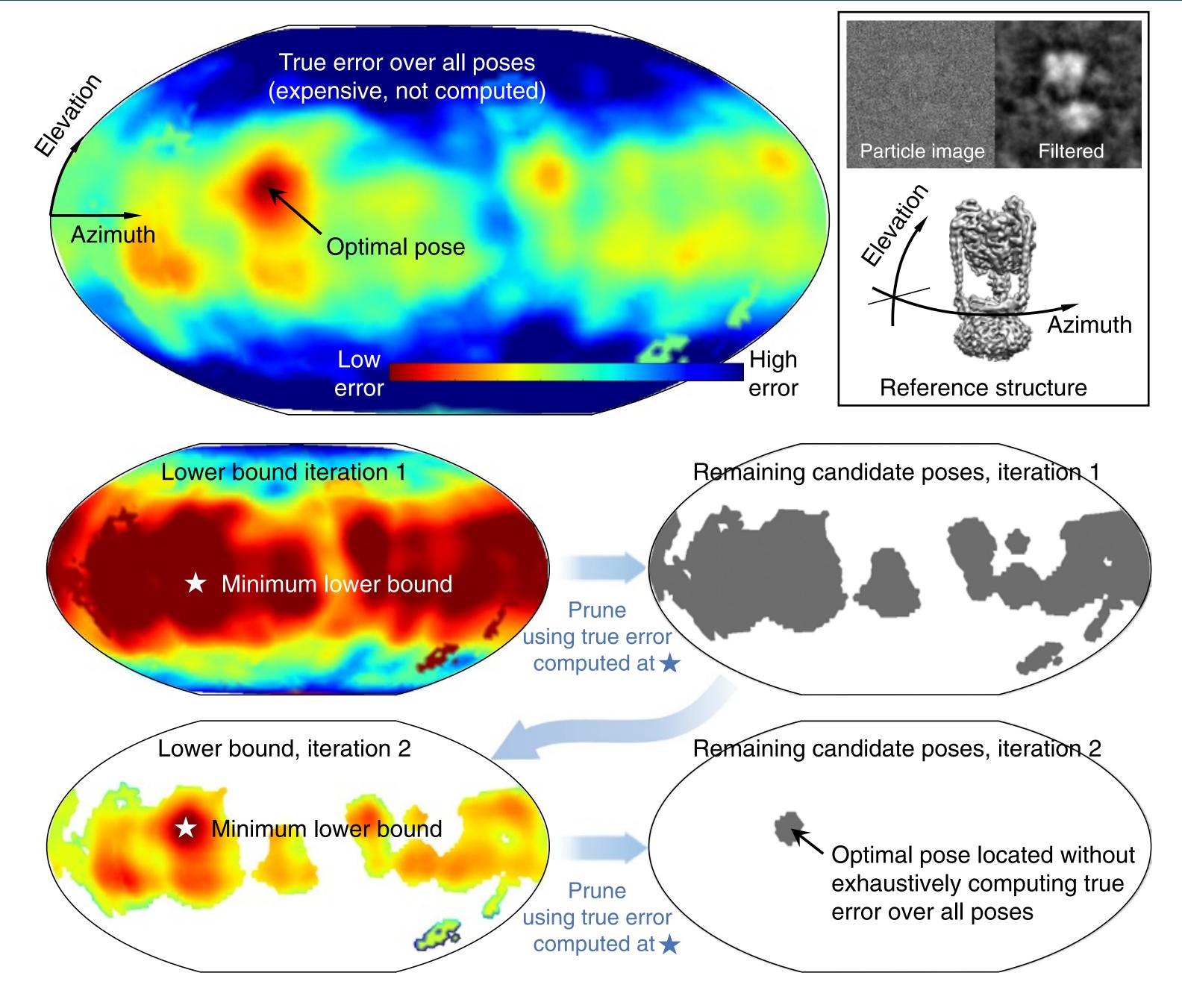
# Branch-and-bound in cryoSPARC for integrating over orientations



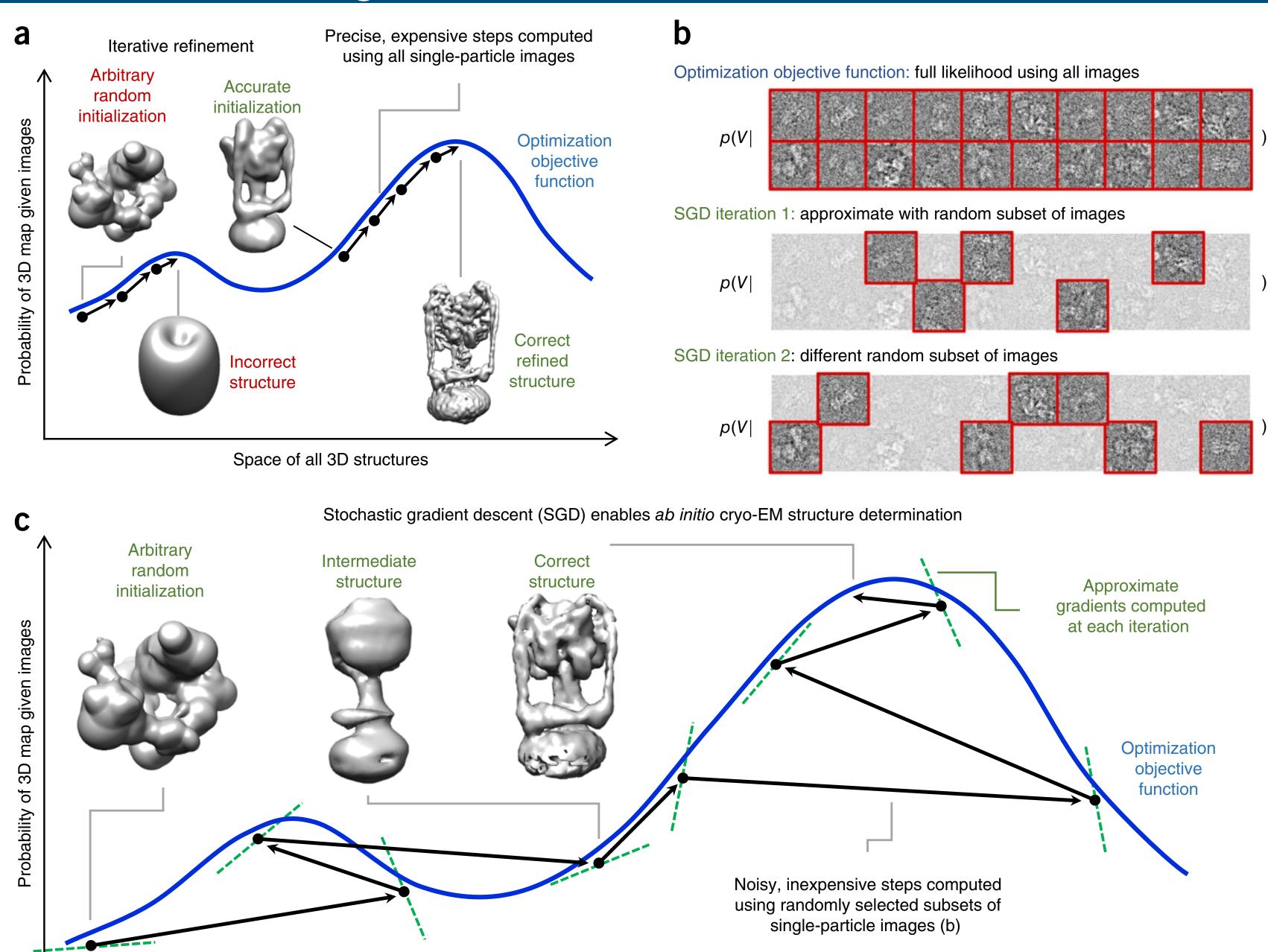
# Branch-and-bound in cryoSPARC for integrating over orientations

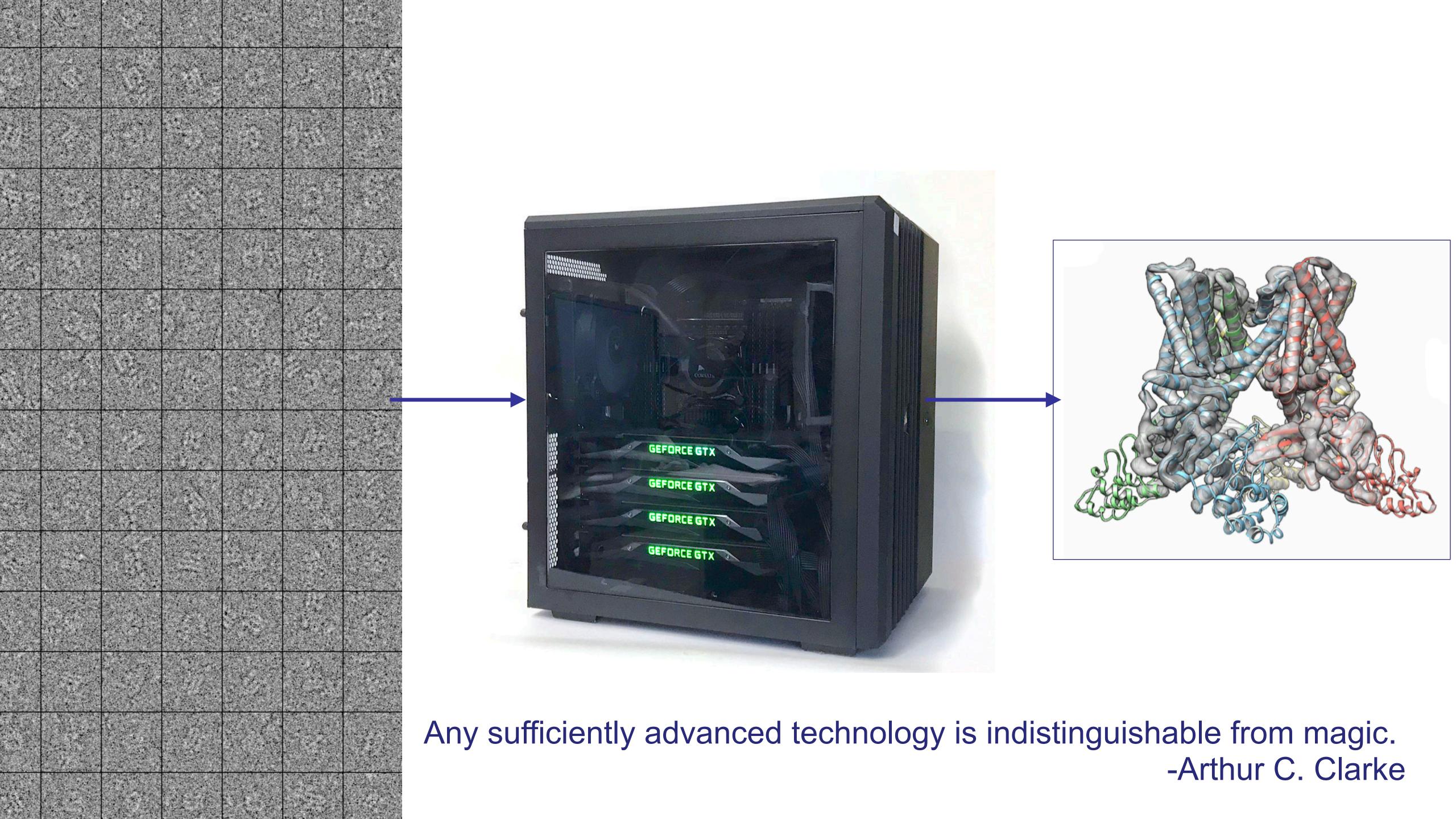


# Branch-and-bound in cryoSPARC for integrating over orientations



## Stochastic gradient descent to avoid model bias





#### **Resources:**

"I don't know of good textbooks. Here's a review that we wrote for a mathematical audience, but might be useful in understanding some details. It has a big appendix that may be worth looking at too." – Fred Sigworth

Computational Methods for Single-Particle Electron Cryomicroscopy

PMCID: PMC8412055

DOI: 10.1146/annurev-biodatasci-021020-093826

Lecture notes at <a href="https://cryoemprinciples.yale.edu/">https://cryoemprinciples.yale.edu/</a> Select the link to All Files.