



THE UNIVERSITY *of* TEXAS

SCHOOL OF HEALTH INFORMATION
SCIENCES AT HOUSTON

X-Ray Crystallography Pt. I

For students of HI 6001-125

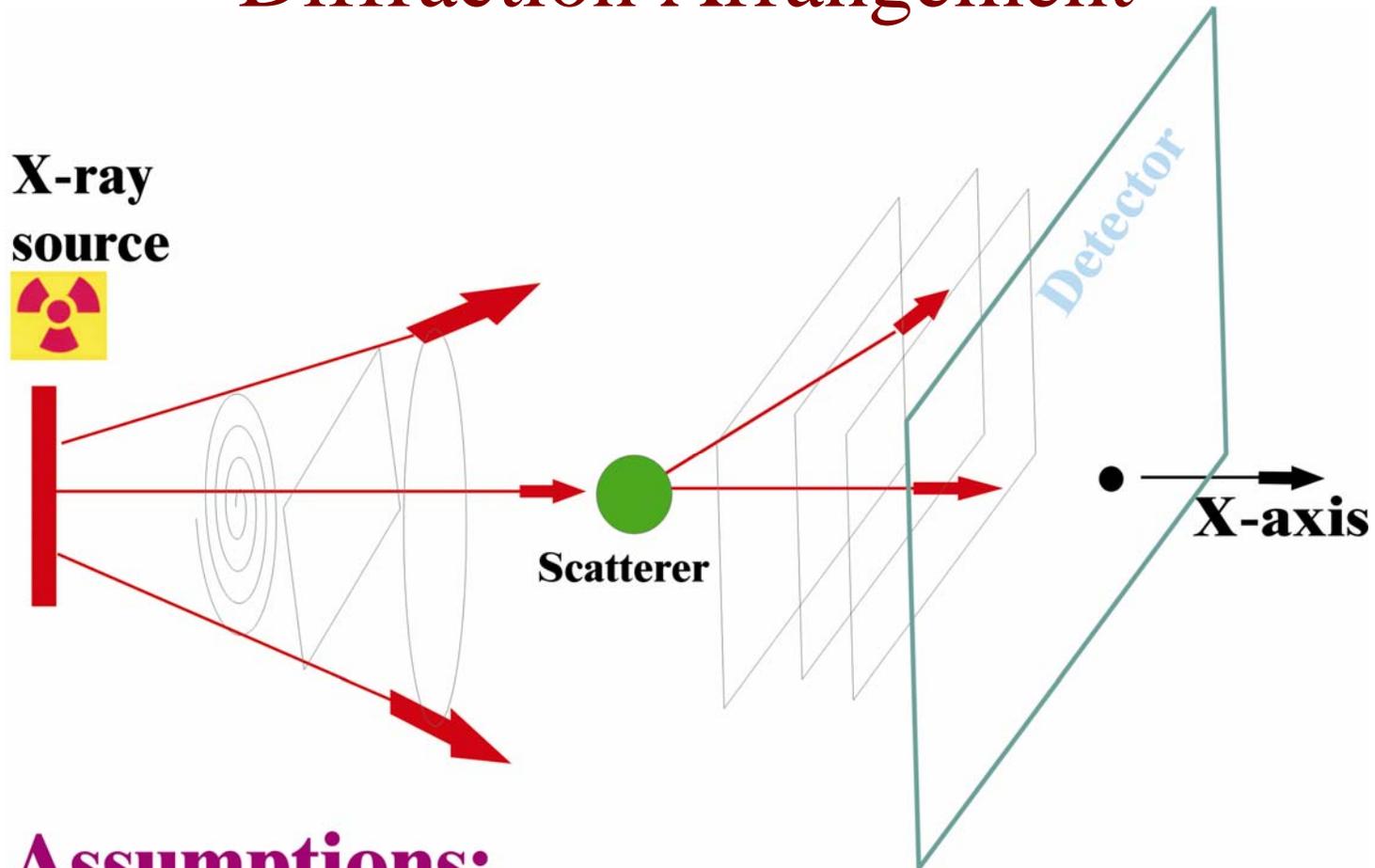
“Computational Structural Biology”

Sugoto Chakravarty, Ph.D.

Baylor College of Medicine

<http://biomachina.org/courses/structures/02.html>

Diffraction Arrangement



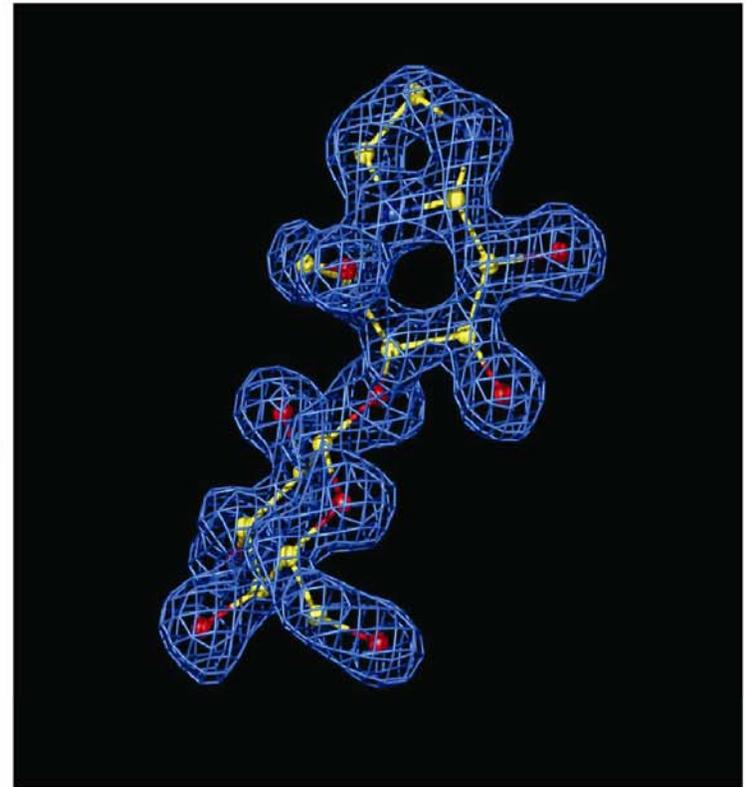
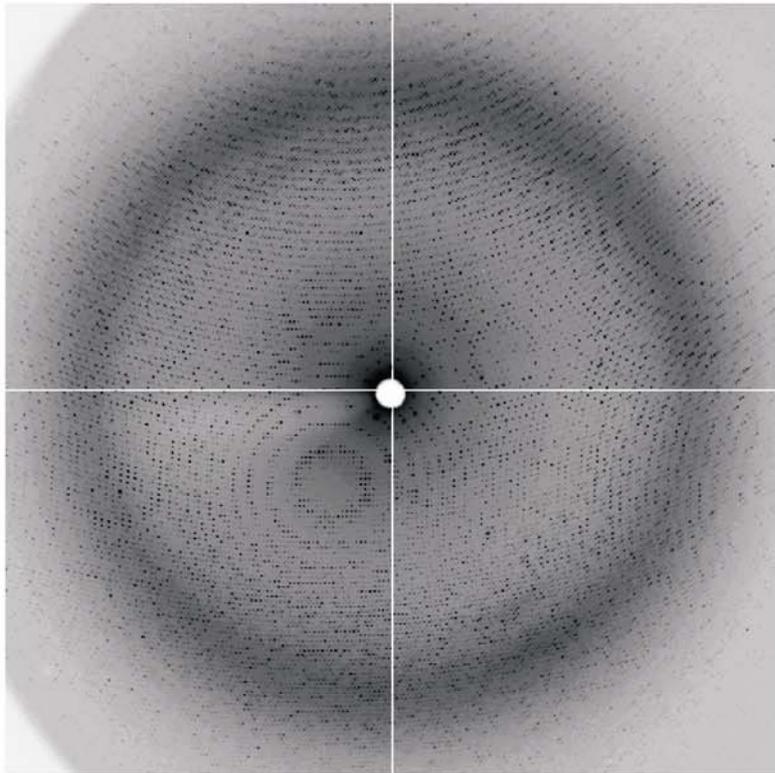
Assumptions:

- **Incident and scattered plane waves**
- **Elastic scattering: No change in wavelength**

Probing Macromolecular Atomic Structures

- Crystalline sample
- Probe should be chosen such that its wavelength is of atomic dimensions: X-rays
- Suitable detection systems

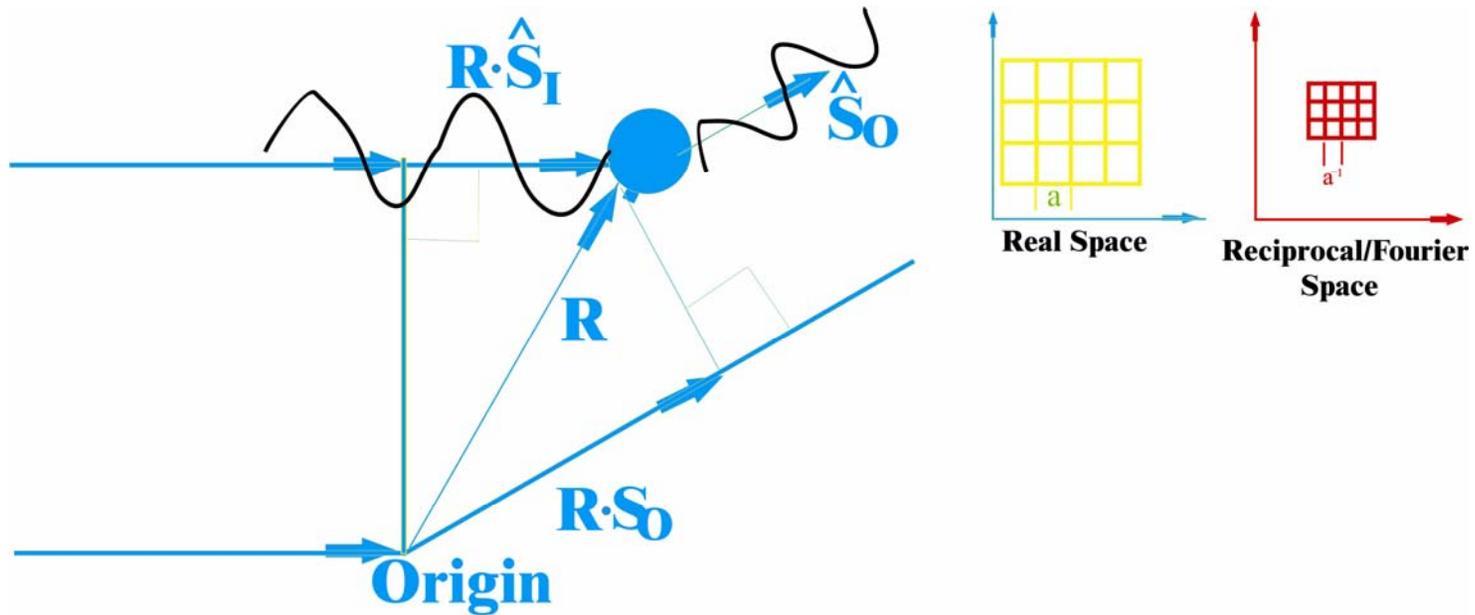
From Diffraction Spots to Atomic Models



Scattering from Regular Atomic Arrays

- Diffraction pattern may be understood from principles of optical diffraction on gratings
- Scattering from single atom
- Scattering from an atomic array

Path and Phase Difference



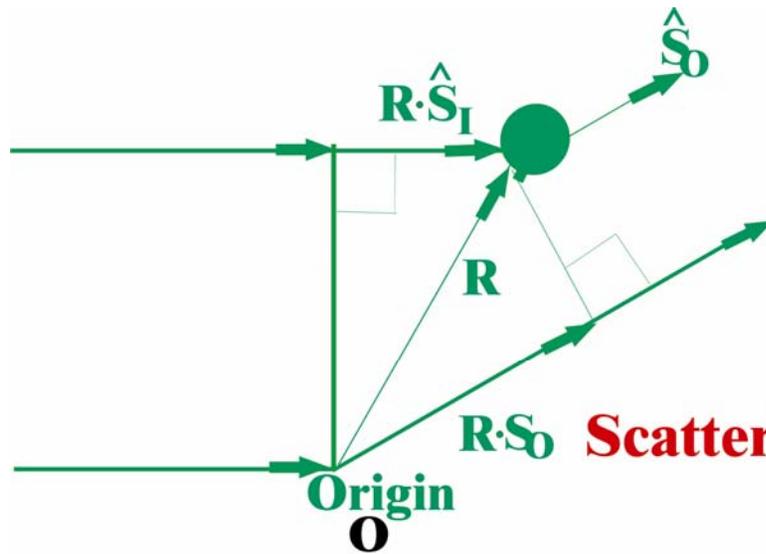
Path difference

$$\mathbf{R} \cdot \mathbf{S}_O - \mathbf{R} \cdot \mathbf{S}_I$$

Phase difference

$$\frac{2\pi}{\lambda} [\mathbf{R} \cdot \mathbf{S}_O - \mathbf{R} \cdot \mathbf{S}_I]$$

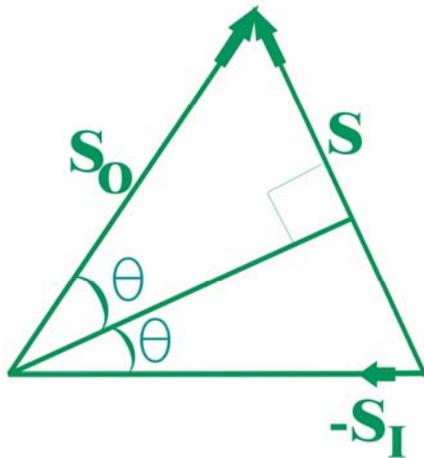
Scattering from a Single Atom



Path difference

$$\mathbf{R} \cdot \hat{S}_0 - \mathbf{R} \cdot \hat{S}_I$$

Scattering vector, $\hat{\mathbf{S}}: \hat{S}_0/\lambda - \hat{S}_I/\lambda$

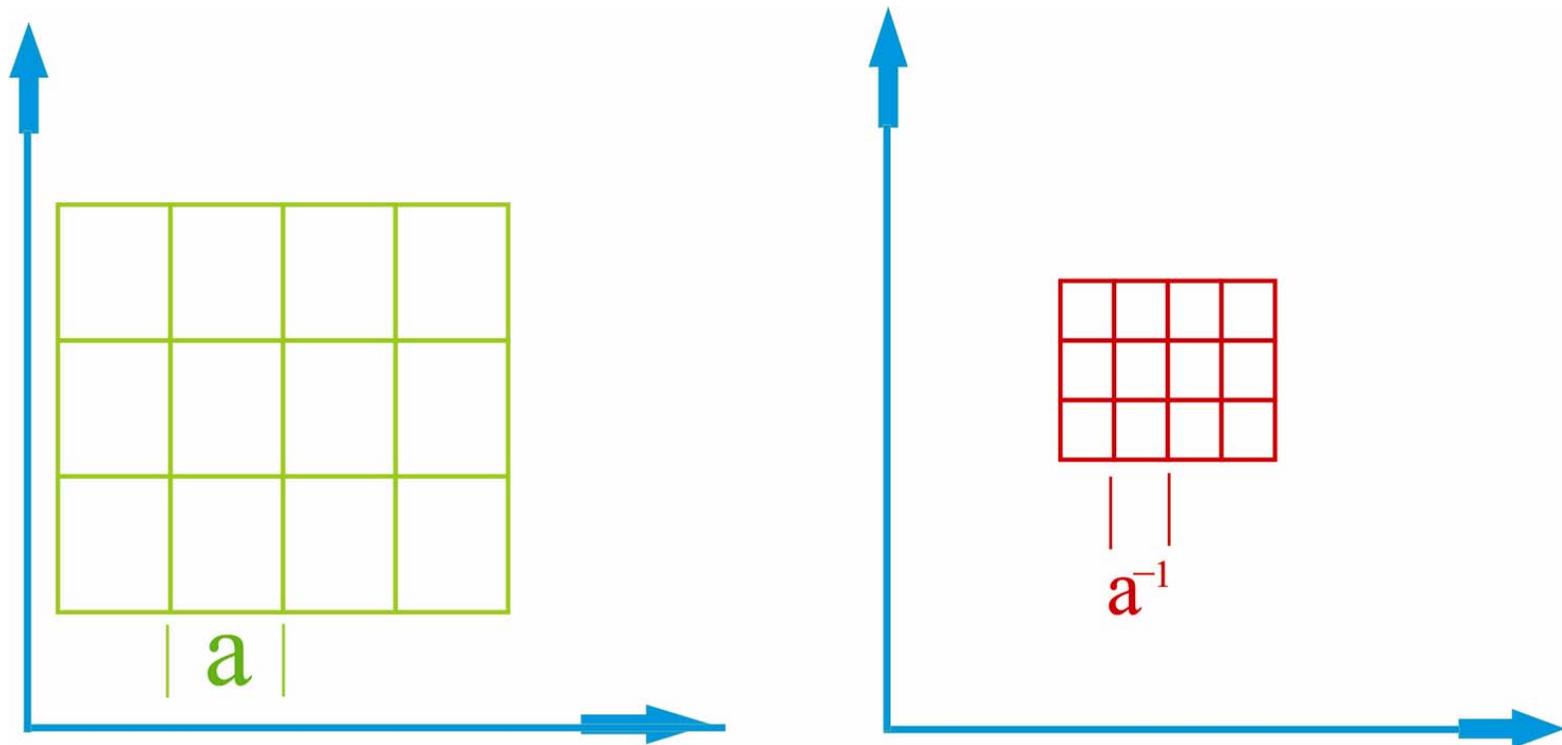


$$\mathbf{S} \cdot \mathbf{S} = 4 \sin^2 \theta / \lambda^2$$

$$|\mathbf{S}| = 2 |\sin \theta| / \lambda$$

$$0 \leq |\mathbf{S}| \leq 2 / \lambda$$

Real and Reciprocal Space



real space/
direct space/
Cartesian space

reciprocal space/
frequency space/
Fourier space

Phase Difference and Scattering Power

Real Space

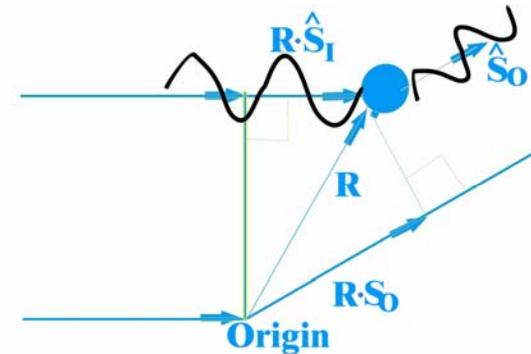
Reciprocal space

Path difference

$$\mathbf{R} \cdot \mathbf{S}_0 - \mathbf{R} \cdot \mathbf{S}_1$$

Phase difference

$$\frac{2\pi}{\lambda} [\mathbf{R} \cdot \mathbf{S}_0 - \mathbf{R} \cdot \mathbf{S}_1]$$



**If electron at origin has a scattering power $E(\mathbf{S})$ along direction \mathbf{S} ,
Scattering power of an electron at**

\mathbf{R} is

$$E(\mathbf{S}) e^{2\pi i \mathbf{S} \cdot \mathbf{R}}$$

Structure factor: $e^{2\pi i \mathbf{S} \cdot \mathbf{R}}$
 (Geometrical part
 of the scattering)

SF for Electron Density Distribution

Structure factor for a unit volume: $e^{2\pi i\mathbf{S}\cdot\mathbf{r}}$

For a volume element $d\mathbf{r}$

Structure factor: $\rho(\mathbf{r})e^{2\pi i\mathbf{S}\cdot\mathbf{r}} d\mathbf{r}$

For a continuous electron density distribution,

Structure factor:

$$\mathbf{F}(\mathbf{S}) = \int \rho(\mathbf{r}) e^{2\pi i\mathbf{S}\cdot\mathbf{r}} d\mathbf{r}$$

Plane Waves and Fourier Transforms

$$\mathbf{F}(\mathbf{S}) = \int \rho(\mathbf{r}) e^{2\pi i \mathbf{S} \cdot \mathbf{r}} d\mathbf{r}$$

Structure factor is the Fourier Transform of the electron density distribution

Conversely, the electron density distribution is the inverse Fourier transform of the structure factor

$$\rho(\mathbf{r}) = \frac{1}{V} \int \mathbf{F}(\mathbf{S}) e^{-2\pi i \mathbf{S} \cdot \mathbf{r}} d\mathbf{S}$$

Phase Problem in Crystallography

The structure factor $F(\mathbf{S})$ for a given direction \mathbf{S} is a complex quantity.

$$F(\mathbf{S}) = |F(\mathbf{S})| e^{i\phi}$$

$|F(\mathbf{S})|$: Structure factor amplitudes

ϕ : Phases

The measured intensity $I(\mathbf{S})$ is

$$I(\mathbf{S}) = F(\mathbf{S}) * F(\mathbf{S}) = |F(\mathbf{S})|^2$$

Thus the phases ϕ are lost during measurement.

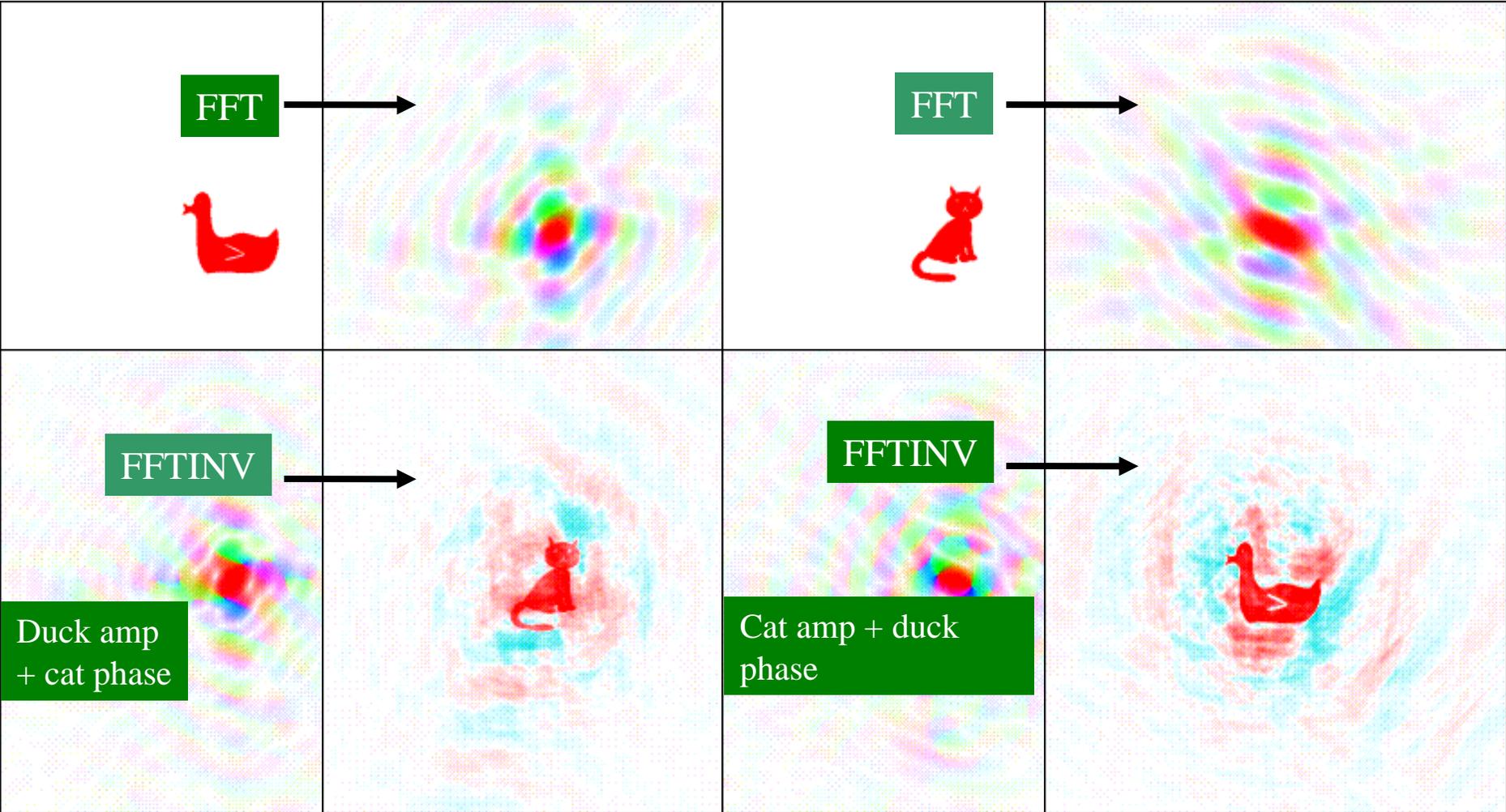
Phases Needed to Reconstruct Images

Scattered X-rays carry phase information

**But no X-ray lens to recombine the amplitudes
and the phases to reconstruct the original image!**

**Which is more important:
Amplitudes or phases?**

Phases more Important than Amplitudes



Phase Problem in Crystallography

How to estimate the phases that give the best electron density $\rho(r)$ from the structure factors $F(S)$

Uniform Sample Cannot Deflect X-rays

Atoms must have heterogenous electron density distribution to scatter X-rays.

Assume a uniform electron density distribution:

$$\begin{aligned}\rho(\mathbf{r}) &= \rho \\ \mathbf{F}(\mathbf{S}) &= \int \rho(\mathbf{r}) e^{2\pi i \mathbf{S} \cdot \mathbf{r}} d\mathbf{r} \\ &= \rho \int e^{2\pi i \mathbf{S} \cdot \mathbf{r}} d\mathbf{r} \\ &= \rho \delta(\mathbf{S} - \mathbf{0}) \\ &= \mathbf{F}(\mathbf{0})\end{aligned}$$

Uniform Sample Cannot Deflect X-rays

$$\mathbf{F}(\mathbf{S}) = \mathbf{F}(\mathbf{0}) \quad \text{if } \rho(r) = \rho$$

Thus, a uniform electron density gets scattered only in the forward direction.

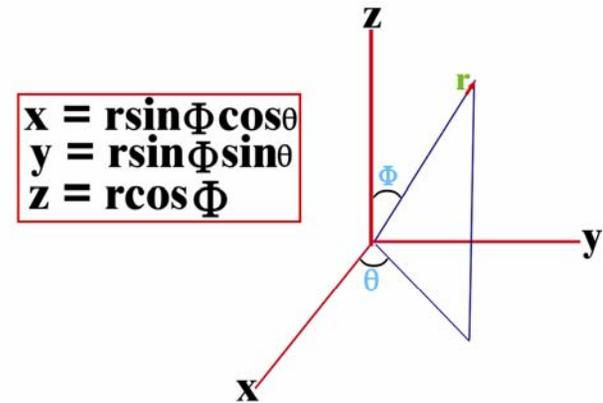
Spatial or temporal heterogeneity that creates a contrast between the electrons and the background, is a must to scatter X-rays in all directions.

X-ray Scattering from a Single Atom

Assume a spherically symmetric electron density distribution

$$\mathbf{F}(\mathbf{S}) = \int \rho(\mathbf{r}) e^{2\pi i \mathbf{S} \cdot \mathbf{r}} d\mathbf{r}$$

$$\begin{aligned} \mathbf{F}(\mathbf{S}) &= \int_0^{2\pi} d\phi \int_0^{\pi} \sin\theta d\theta \int_0^{\infty} dr \rho(r) r^2 e^{2\pi i \mathbf{S} \cdot \mathbf{r}} \\ &= 2\pi \int_0^{\infty} dr \rho(r) r^2 \int_0^{\pi} d\theta \sin\theta e^{2\pi i S r \cos\theta} \end{aligned}$$



X-ray Scattering from a Single Atom

$$F(\mathbf{S}) = 2\pi \int_0^{\infty} dr \rho(r) r^2 \int_0^{\pi} d\theta \sin\theta e^{2\pi i \mathbf{S} r \cos\theta}$$

Substitute $x = r \cos\theta$

$$F(\mathbf{S}) = 4\pi \int_0^{\infty} dr \rho(r) r^2 \left[\frac{\sin 2\pi S r}{2\pi S r} \right]$$

$f(\mathbf{S})$: Atomic form factor

Finite Atomic Size & Form Factor

Assuming a tailed electronic distribution

$$\rho(r) = Zne^{-kr^2} \quad \text{where } Z \text{ is the atomic number}$$

it can be shown that $f(S) = Ze^{\left(\frac{-\pi^2}{k}\right)S^2}$ 

Thus, finite size of atom introduces a resolution dependent fall-off of the atomic form factor.

This is due to the destructive interference of the scattered radiation from electrons.

Resources

- Cantor, R.C. & Schimmel, P.R. (1980). *Biophysical Chemistry Part II: Techniques for the study of biological structure and function*. New York; W.H. Freeman & Co, Chapter 13.
- Blundell, T.L. & Johnson, L.N. (1976). *Protein Crystallography*. London; Academic Press.
- Stout, G.H. & Jensen, L.M. (1968). *X-ray Structure Determination: A Practical Guide*. New York: Macmillan