

# Handout 8: Data Collection

Sources, detectors and methods

Chem 6850/8850

X-ray Crystallography

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THE UNIVERSITY OF  
TOLEDO

# Measurement of diffracted beams

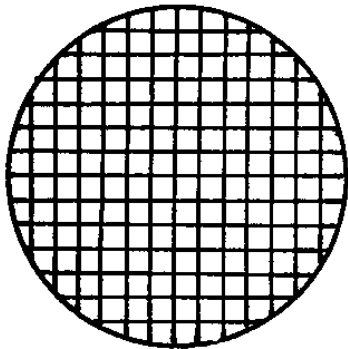
- **Position of diffracted beams gives information about unit cell dimensions and translational symmetry elements**
  - systematic absences
- **Intensity of beams arises from electron density distribution inside the unit cell**
  - type and position of atoms
- **There are many different ways of recording the diffracted beams**

# Important parts of the experimental setup

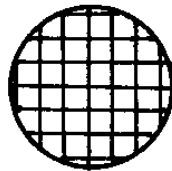
- **Radiation source**
  - X-ray tubes, synchrotrons, neutrons
- **Specimen mount**
  - single crystal or powder
- **Detector setup**
  - film, image plate, point detector, area detector
- **Experimental method**
  - how do you sample reciprocal space, how much of it

# How many reflections are needed?

- For each parameter to be determined, you should measure ~10 reflections
- The number of reflections that can be measured depends on the X-ray wavelength
  - $N = (4\pi/3)(8V/n\lambda^3)$ 
    - $n$  = number of lattice points in the unit cell



MoK $\alpha$   
0.71 Å



CuK $\alpha$   
1.54 Å



CrK $\alpha$   
2.29 Å

“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.



# Radiation sources

- **X-ray tube**
  - lab X-ray source
  - line source with white background
  - low flux
- **Synchrotron radiation**
  - high flux
  - white, tunable radiation
- **Neutrons from nuclear reactor or spallation source**
  - most expensive method
  - useful for light elements or distinction of elements that are close in Z

# X-ray tubes

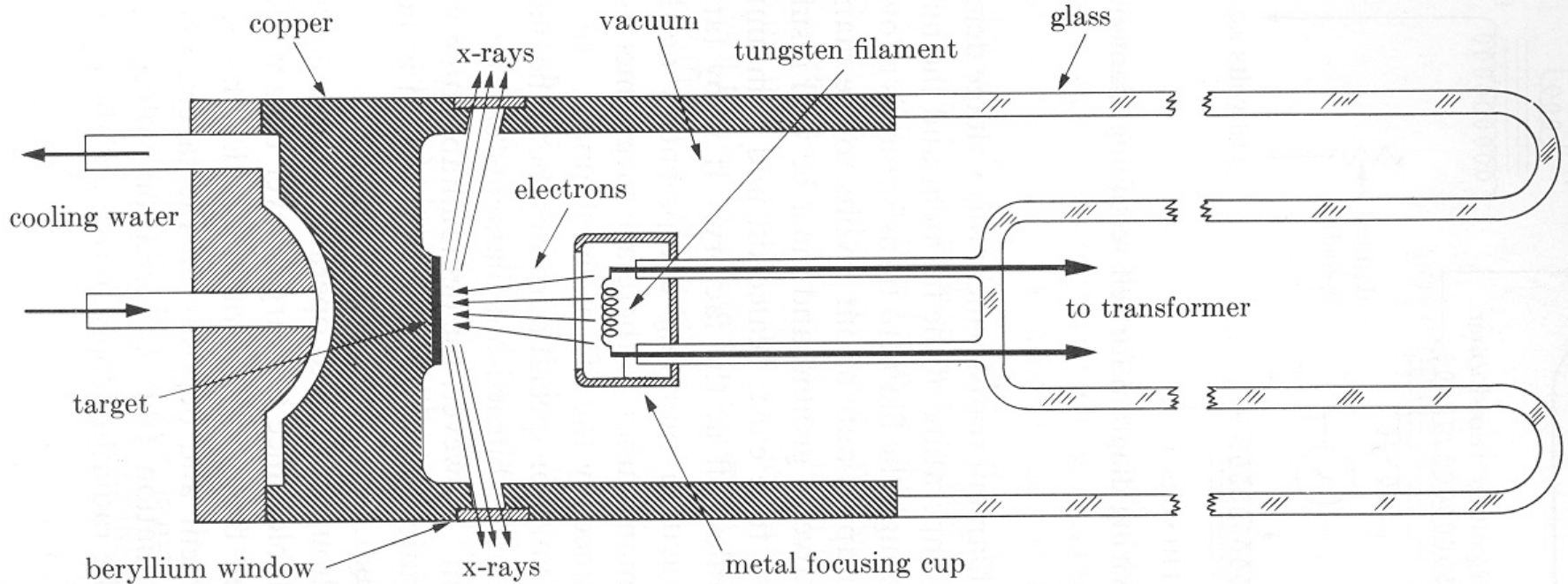
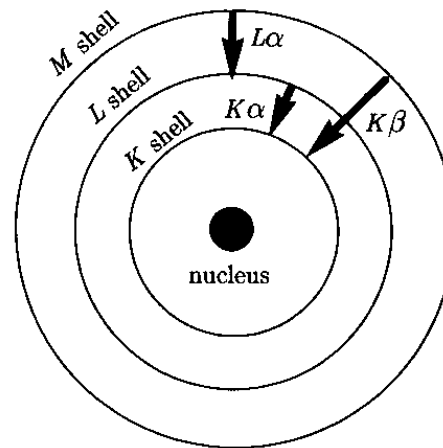
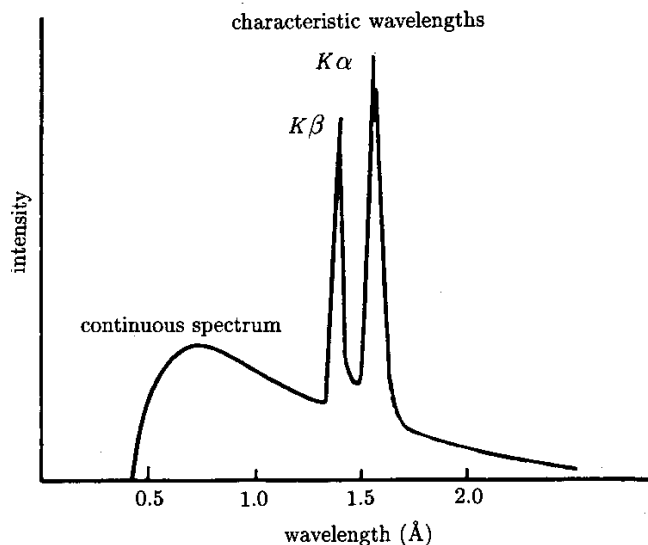


Fig. 1-15 Cross section of sealed-off filament x-ray tube (schematic).

"Elements of X-ray Diffraction", Cullity and Stock,  
Prentice Hall College Div., 3rd edition, 2001.

# Radiation from X-ray tubes

- X-rays are produced by hitting a metal target with an electron beam
  - white radiation from deceleration of electrons in target, called *Bremsstrahlung*
  - characteristic radiation from electronic transitions after an inner shell electron is knocked out



“Elements of X-ray Diffraction”,  
Cullity and Stock, Prentice Hall  
College Div., 3rd edition, 2001.



# Monochromatic X-rays

- **The majority of lab X-ray experiments requires monochromatic (single wavelength) radiation**
  - Exception: Laue patterns
- **X-ray tubes produce white and characteristic radiation**
  - White radiation has low intensity, but having more than one characteristic line would lead to superimposed patterns!
- **We need to monochromate the beam to get easily analyzable data**
  - Always done on incident beam for single crystal experiments
  - Incident beam, diffracted beam, and detector options are possible for powder setups
- **Generally, the  $K\alpha$  line is used (most intense)**





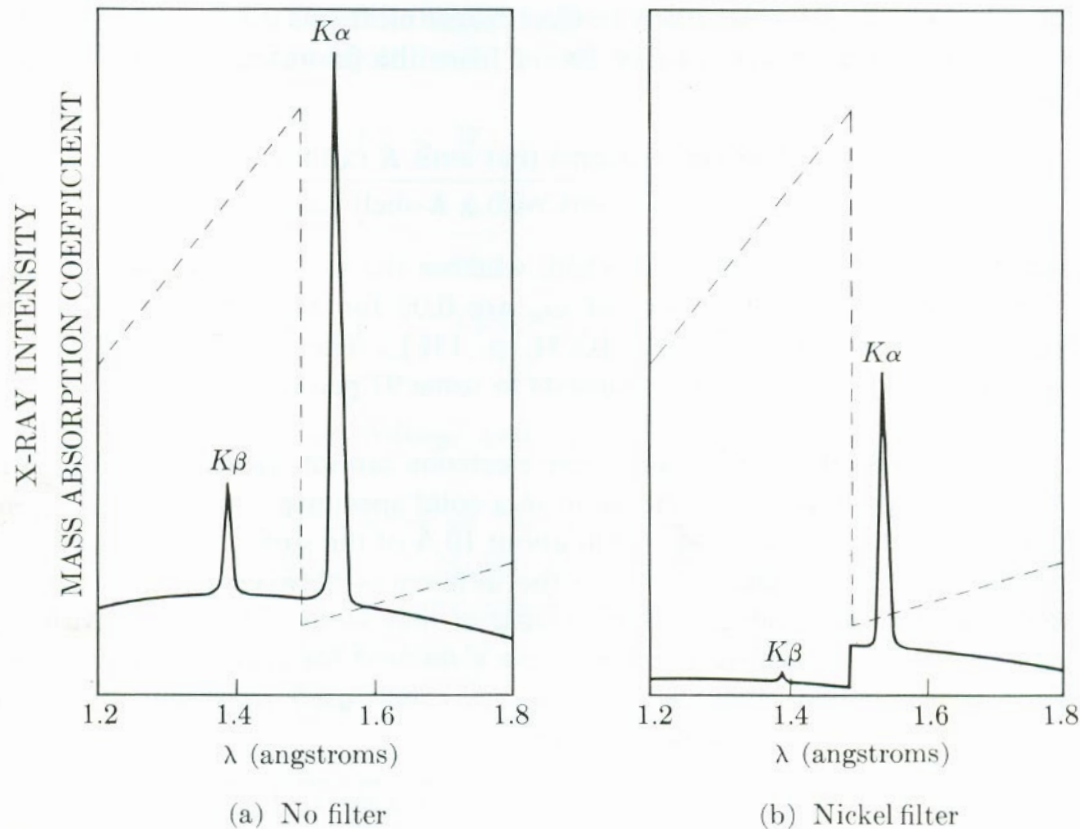
# Making Radiation Monochromatic

- **Two common options:**
  - X-ray filters
  - Monochromators
- ***X-ray filters absorb X-rays of unwanted wavelength***
  - Nickel filter for copper tubes, zirconium or niobium for molybdenum tubes
- ***Monochromators diffract the desired wavelength***
  - Graphite 200 or others
  - Usually 2-bounce or 4-bounce setups
- **For powder experiments: Energy-sensitive detectors**
  - No monochromation necessary, only X-rays of the desired wavelength are counted
  - Very low background, very good intensity

# X-ray Filters

- X-ray filters absorb unwanted radiation
- Filters must be chosen to match the target
  - Ni for Cu
  - Zr or Nb for Mo
- Choose a filter with an absorption edge between the desired and all undesirable wavelengths
  - Atomic number just below atomic number of the target element
- A filter will not *eliminate*, but significantly reduce the intensity of unwanted wavelengths
  - The thicker the filter, the more of the unwanted wavelengths is absorbed
  - But some  $K\alpha$  intensity is also absorbed → tradeoff

# X-ray Filters

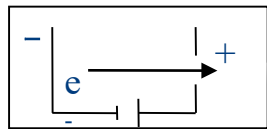


**Fig. 1-13** Comparison of the spectra of copper radiation (a) before and (b) after passage through a nickel filter (schematic). The dashed line is the mass absorption coefficient of nickel.

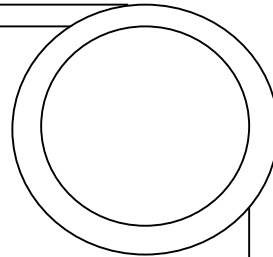
Cullity; "Elements of X-ray Diffraction"

# Synchrotron radiation

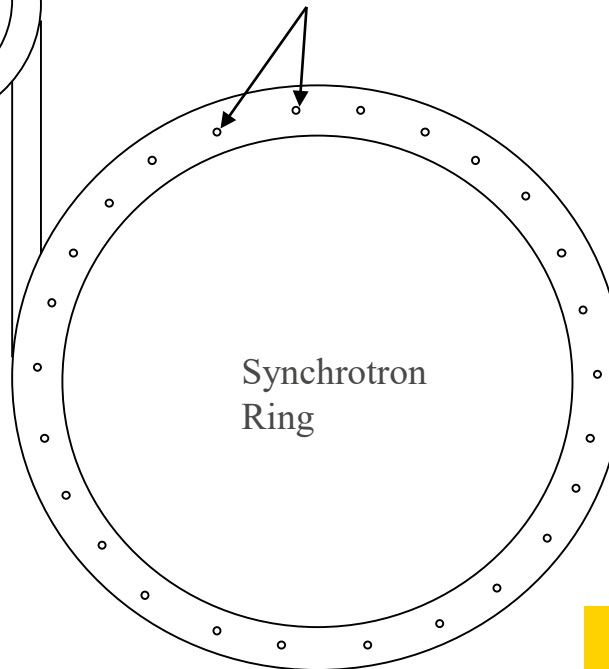
Linear accelerator



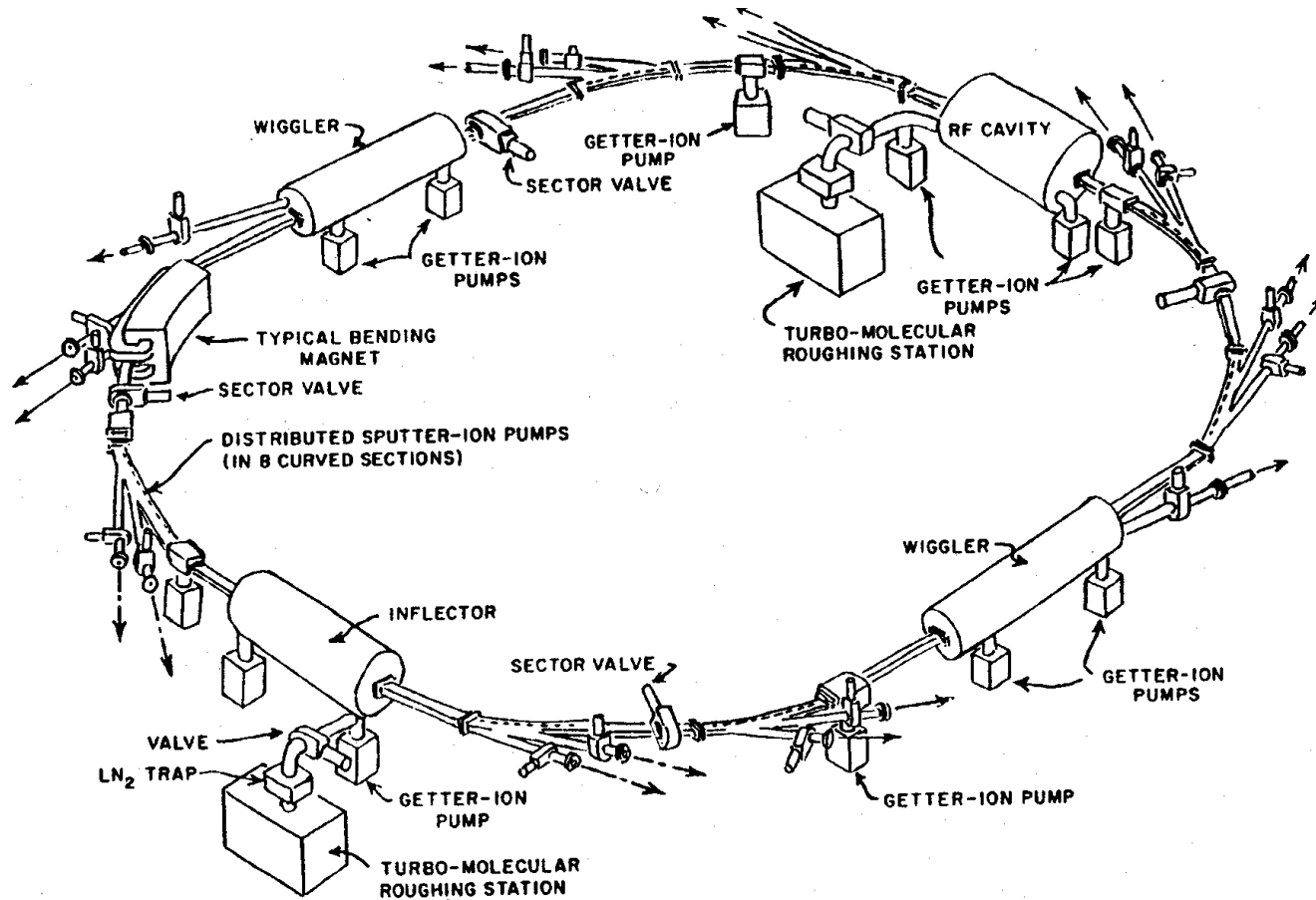
Booster  
Ring



electron bunches



# Producing Synchrotron Radiation



Winick, Doniach; "Synchrotron Radiation Research"

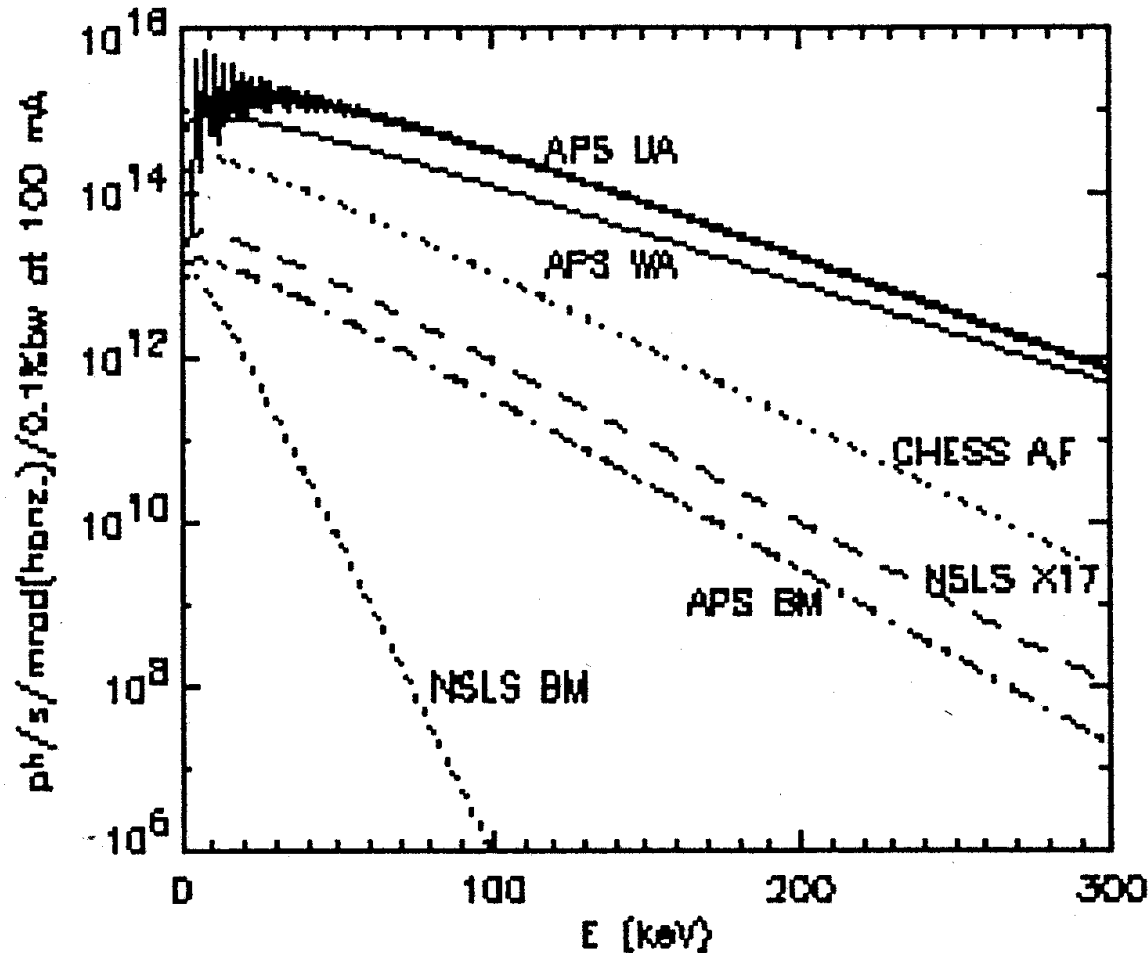


# The Advanced Photon Source



[http://www.aps.anl.gov/About/APS\\_Overview/](http://www.aps.anl.gov/About/APS_Overview/)

# Energy spectrum of synchrotron radiation



- High intensity
- Plane polarized
- Intrinsically collimated
- White radiation
- Has time structure



# Neutron sources

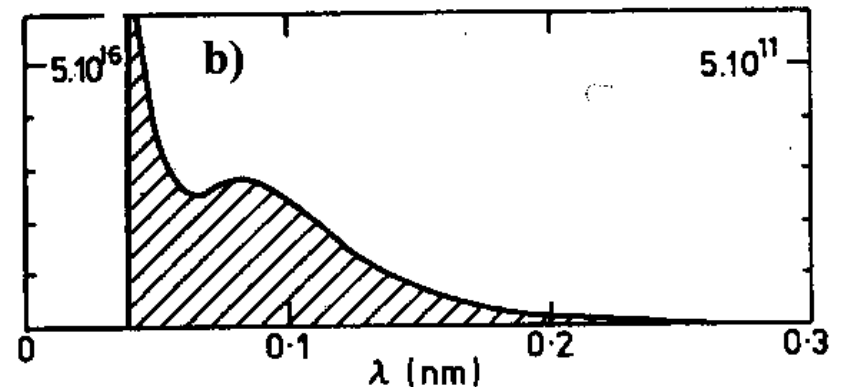
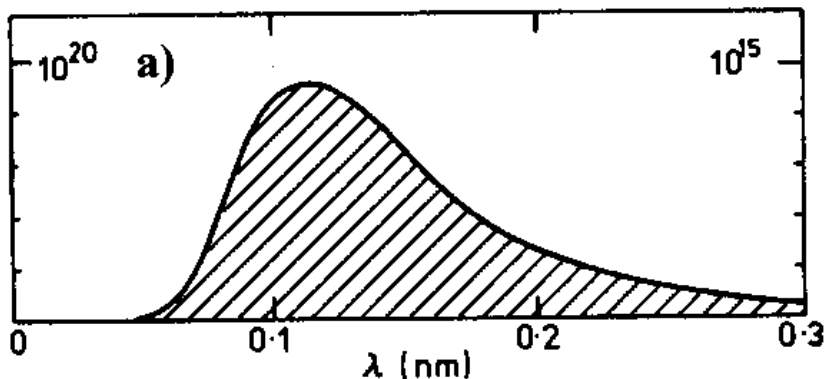
## Reactor sources

- Neutrons are produced by nuclear chain reaction
- Neutrons must be slowed down by moderator for use in diffraction
- Neutron wavelength distribution is thermal equilibrium distribution from moderator
- Monochromator needed => uses only small portions of the produced neutrons

## Spallation sources

- Neutrons are produced by bombarding a metal target with protons
- Different wavelength distribution from reactor
- High peak flux, low average flux
- Due to time structure, all neutrons can be used

Windsor, C. G. *Pulsed Neutron Scattering*; Taylor & Francis Ltd.: London, 1981.





# The Spallation Neutron Source at ORNL

(Image source: <http://neutrons.ornl.gov>)



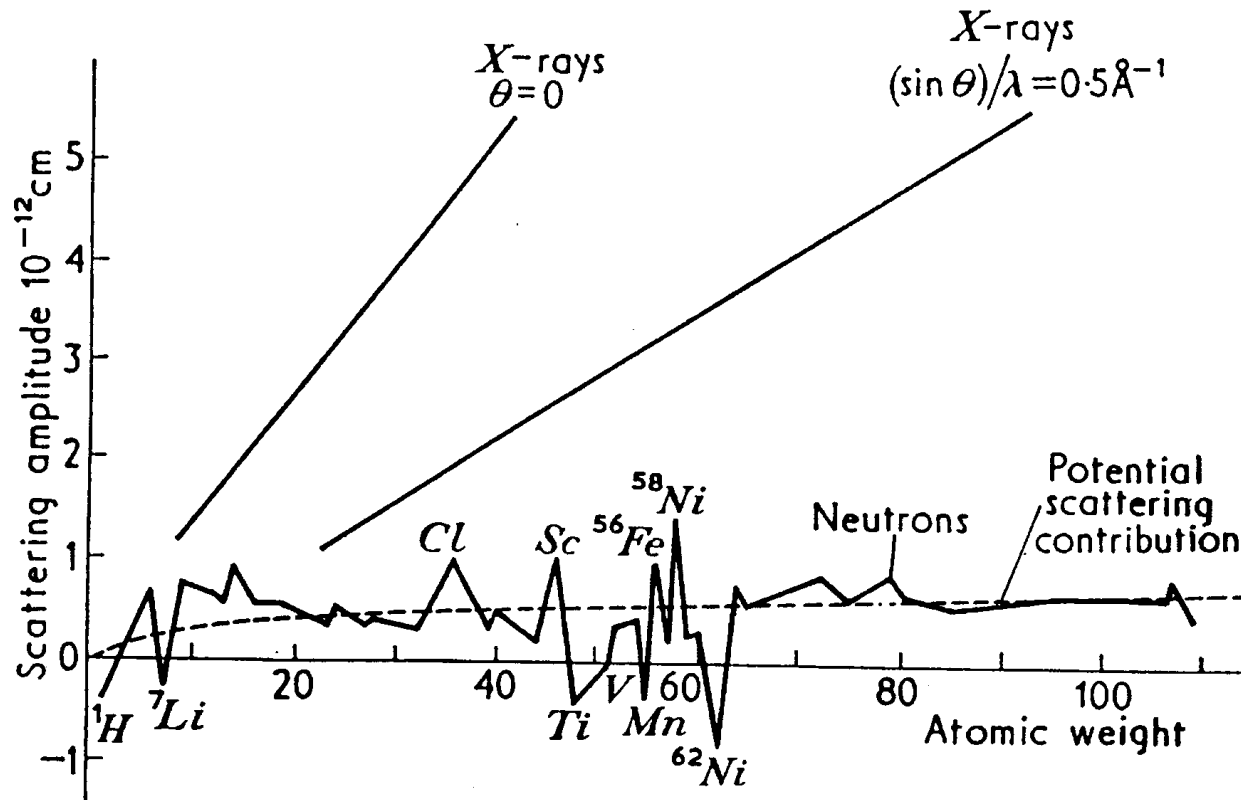
# Why neutrons?

- **Neutrons are scattered by nuclei, not by electrons**
  - form factor does not show rapid fall off at high angles
  - thermal motion of nuclei is much smaller than thermal motion of the electron cloud
- **Neutron form factors are not dependent on Z**
  - more sensitive to light elements than X-rays
  - isotopes or neighboring elements can often be distinguished
- **Neutrons interact with magnetic moments**
  - magnetic structure of materials can be determined by neutron diffraction
- **Most elements do not strongly absorb neutrons**

\*Form factor/scattering lengths: Measure of how sensitive an element is to radiation, e.g., how strongly it scatters it (e.g., elements with more electrons scatter X-rays more)



# Neutron scattering lengths



# Specimen preparation - powder

- **A good powder sample should consist of fairly small particles**
  - should definitely be below 20 microns
  - ideal: 1-5 microns
  - usually achieved by grinding and passing through sieves
  - caution: do not overgrind your sample!
- **Sample morphology can be important**
  - e.g., rod like particles tend to result in preferred orientation
- **Most commonly used configurations: Flat plate samples or capillaries**
  - capillaries allow experiments with air sensitive samples or in a special environment (liquid, gas)
  - capillary size depends on absorption properties of the sample

# Specimen selection – single crystal

- **For lab experiments, crystals of ~50 to 200 micron size are generally needed**
  - exact size depends on scattering power
  - at a synchrotron, much smaller crystals might be sufficient
- **Quality is important**
  - only true single crystals will give a single diffraction pattern
  - use microscope and polarizers to make sure crystals are not split or twinned
- **If possible, crystals with approximately equal dimensions are chosen**
  - straightforward absorption correction

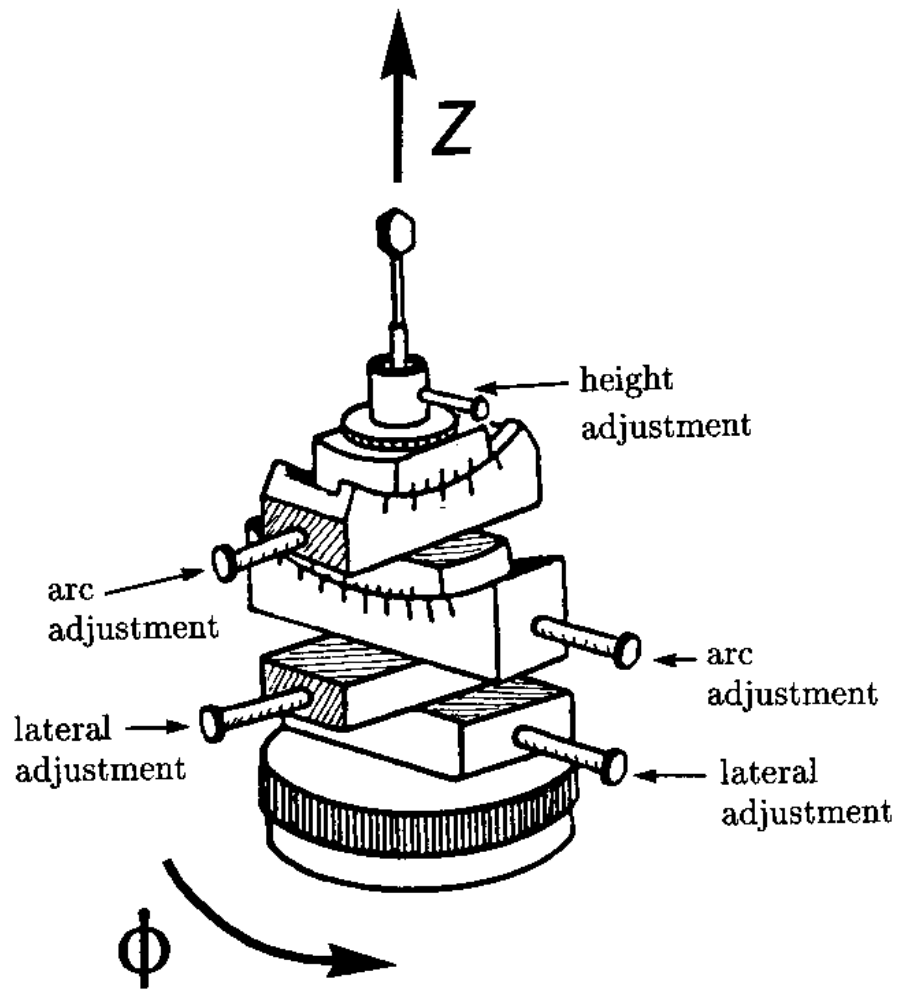


# Specimen preparation – single crystal

- Crystals can be attached to a glass fiber, a loop (common: MiTeGen loops), or sealed inside a capillary tube
  - capillary protects air sensitive samples and could include mother liquor
- Traditionally, crystals were glued to glass fibers with epoxy, but since the advent of low temperature methods, crystals are often mounted using grease or viscous oils
  - the epoxy and the crystal will separate during cooling if their thermal expansion is considerably different
  - grease or oil will become rigid at low temperatures and immobilize the crystal
  - oil can often be used for slightly air sensitive crystals
- The glass fiber or capillary are mounted on a goniometer head



# Goniometer heads



“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.



# Detector options

- **Photographic film is darkened by X-rays**
  - oldest method
  - fairly sensitive, but difficult to get integrated intensities
  - small dynamic range
- **Image plates**
  - developed in the 1980's
  - very sensitive, wide dynamic range (six orders of magnitude)
  - can be read out by computer
  - reusable
  - photosensitive phosphor powder with organic binder
  - latent image of diffracted X-rays is stored and can be read out by scanning a laser across the image plate
    - photostimulated luminescence, detectable by a photomultiplier





# Film

- Can be used for powder and single crystal methods

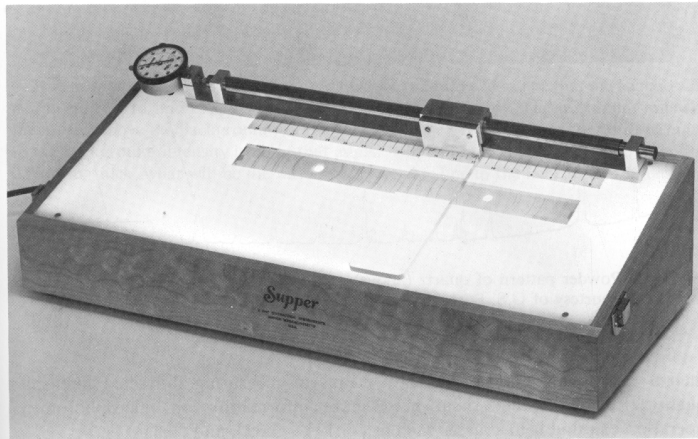


Fig. 6-18 Film-measuring device. (Courtesy of Charles Supper Company.)

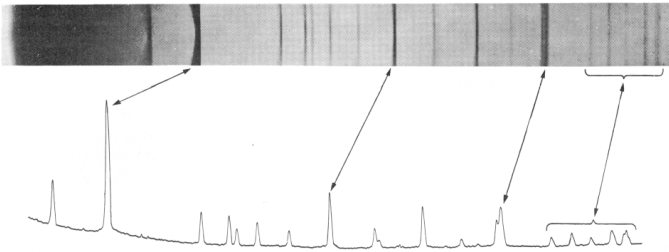


Fig. 6-19 Powder pattern of quartz (above) and corresponding microphotometer trace (below). (Courtesy of U.S. Bureau of Mines.) [6.7]

Cullity, "Elements of X-ray Diffraction"

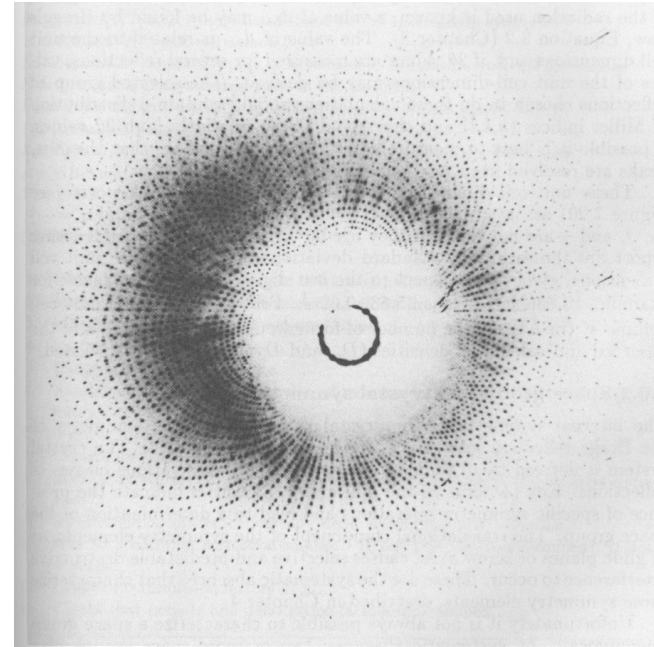
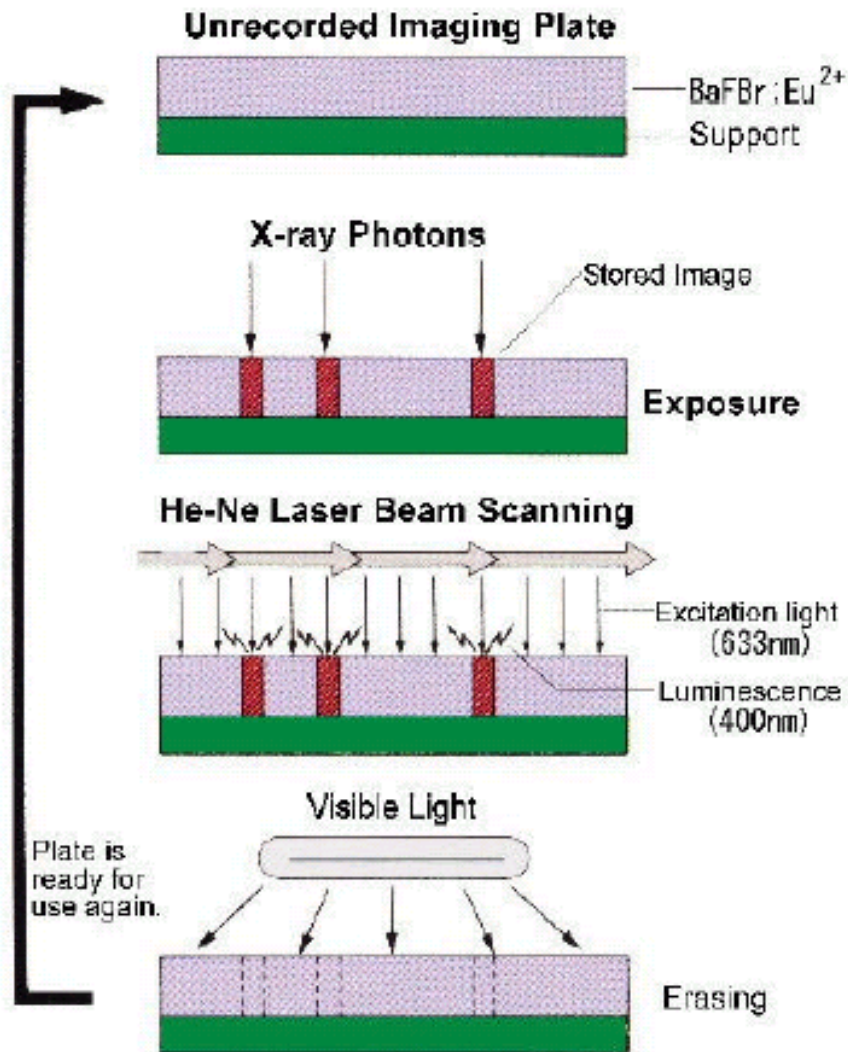


FIGURE 7.19. Laue photograph of lysozyme, 100 msec photograph, with synchrotron radiation 0.2 – 2.0 Å wavelength, taken at CHESS (Cornell High Energy Synchrotron Source). (Courtesy Edwin Westbrook).

"Crystal Structure Analysis for Chemists and Biologists",  
Glusker, Lewis and Rossi, VCH, 1994.



# Image plates



<http://xray0.princeton.edu/~phil/Facility/Guides/XrayDataCollection.html>



# More detector options

## ▪ Point detectors

- works well for powder samples
  - only 1D information collected anyway
- rather slow for single crystal experiments
- proportional, scintillation or solid-state counters

## ▪ Area detectors

- collection of multiple spots at once
- standard setup of most modern (small molecule) single crystal diffractometers
  - CCD detector = charge coupled device detectors
  - CMOS detector = Complementary Metal Oxide Semiconductor; also referred to as “active pixel detector”

# Powder setups (1)

- **Oldest method: Debye-Scherrer camera**
  - capillary sample surrounded by cylindrical film
  - simple, cheap setup

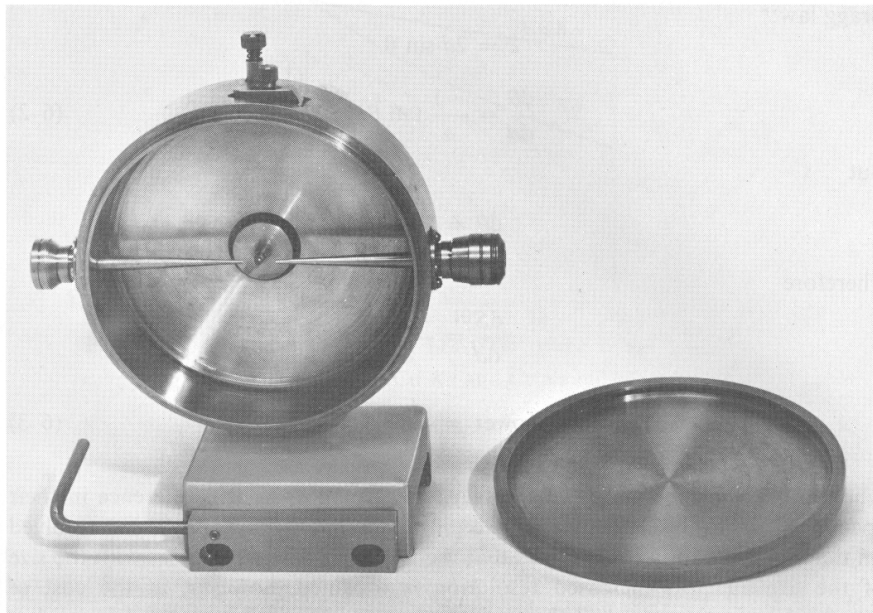


Fig. 6-1 Debye-Scherrer camera, with cover plate removed. (Courtesy of Philips Electronic Instruments, Inc.)

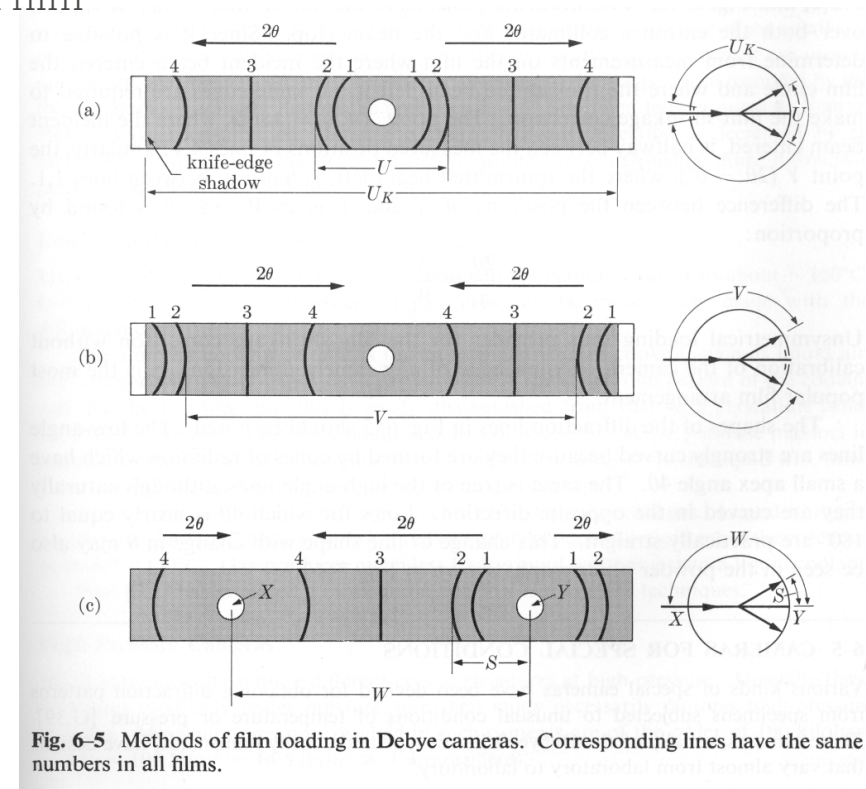


Fig. 6-5 Methods of film loading in Debye cameras. Corresponding lines have the same numbers in all films.

Cullity, "Elements of X-ray Diffraction"

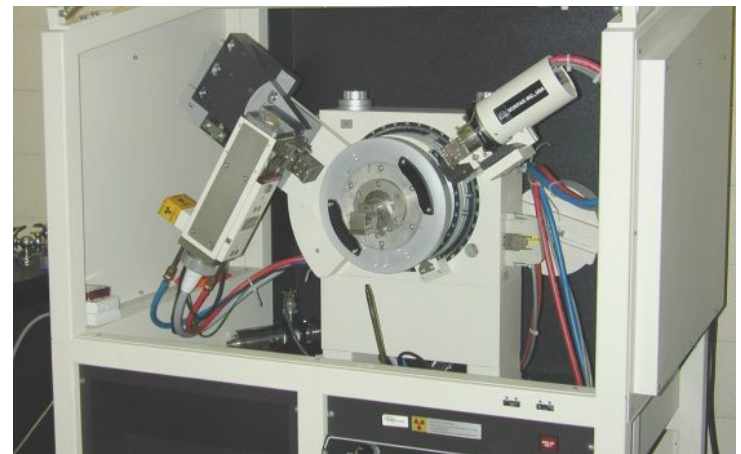




# Powder setups (2)

- Powder diffractometers
  - theta-theta or theta-2theta
  - point or area detectors

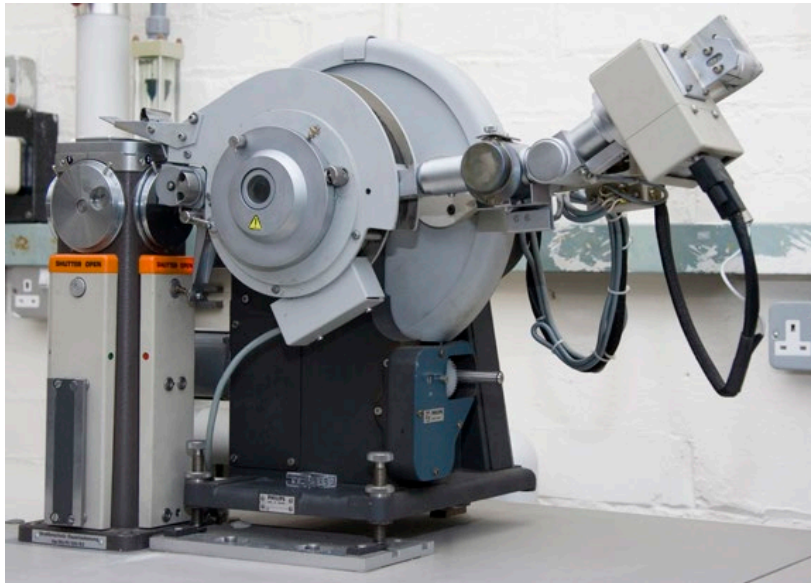
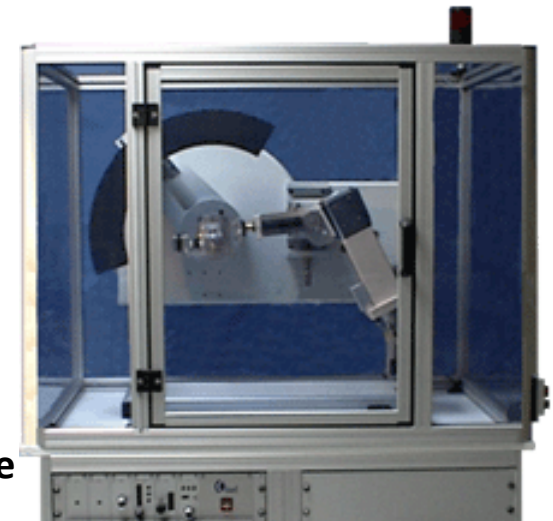
<http://www.msm.cam.ac.uk/xray/images/pdiff3.jpg>



**Scintag theta-theta diffractometer with Peltier cooled solid-state detector**

[www.inel.com](http://www.inel.com)

**Inel diffractometer with 120° PSD (position sensitive detector)**



**Philips PW1050 – first commercial diffractometer (1947)**

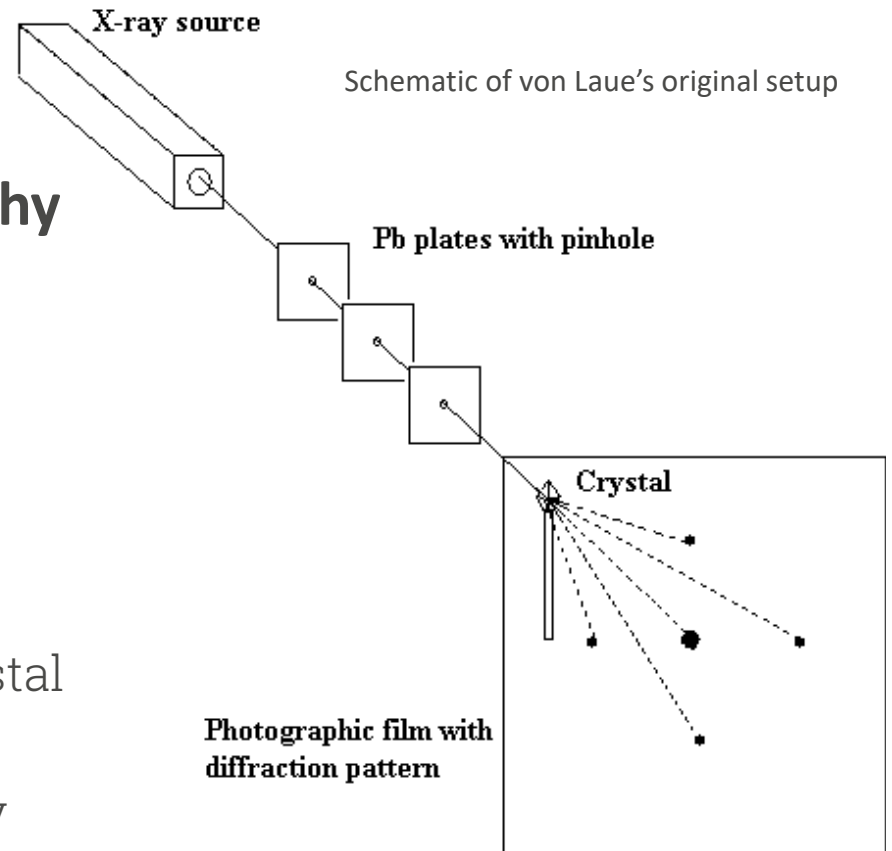
# Single crystal setups: Historical methods

- **Oldest method: Laue photography**

- still in use nowadays for determining orientation and indexing faces of crystals

- **Weissenberg, oscillation and precession methods**

- used for a long time to check crystal quality before tying up diffractometer time (unnecessary with modern, fast instruments)



# Oscillation and Weissenberg techniques

- Use monochromatic radiation
- Crystal is rotated around the spindle axis by a few degree
- Spot patterns are recorded on film
  - if a crystal axis is aligned with the rotation axis, the diffraction spots display a series of straight lines that are called *layer lines*
- In the Weissenberg method, a metal screen allows selection of only one layer line
  - oscillation is coupled to motion of film, which leads to better spot separation
- Neither method gives a direct view of the reciprocal lattice!
  - lattice spacings can be determined indirectly

# Oscillation and Weissenberg techniques (2)

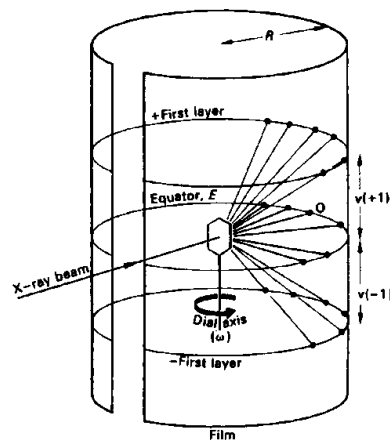


FIGURE 3.26. Basic geometry of the oscillation method, showing how diffraction spots are recorded on a cylindrical film placed around the crystal.

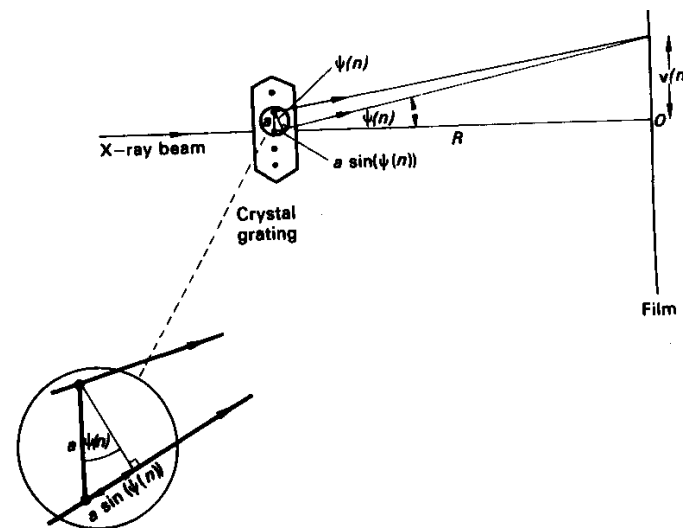


FIGURE 3.28. Diffraction-grating analogy explaining the layer-line spacings on oscillation photographs. Monochromatic x rays are incident normal to  $a$  (oscillation axis) in the crystal. The size of any spot at height such as  $v(n)$  depends upon the experimental conditions.

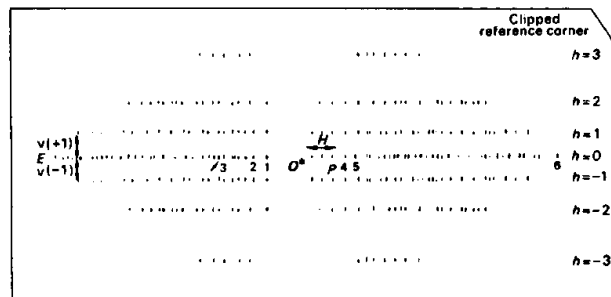


FIGURE 3.27 Sketch of a  $15^\circ$  oscillation photograph of an orthorhombic crystal mounted on the  $z$  axis ( $a = 6.167 \text{ \AA}$ ); the camera radius  $R$  is 30.0 mm and  $\lambda$  ( $\text{Cu K}\alpha$ ) = 1.2542  $\text{\AA}$ . The film is flattened out and the right-hand corner, looking toward the x-ray source, is clipped in order to provide a reference mark.  $P$  represents any equatorial reflection at a distance  $OP$  ( $= H$  mm) from the center  $O$ . Reflections numbered 1–6 on the zero level are indexed by the method given on page 156*f*. The linear scale of the diagram is (1/1.78).

$$a \cdot \sin \psi(n) = n\lambda \quad \tan \psi(n) = v(n) / R$$

$$a = \frac{n\lambda}{\sin \left\{ \tan^{-1} \left[ v(n) / R \right] \right\}}$$

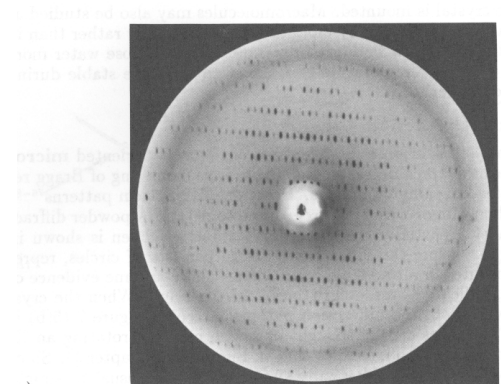
“Structure Determination by X-ray Crystallography”, Ladd and Palmer, Plenum, 1994.



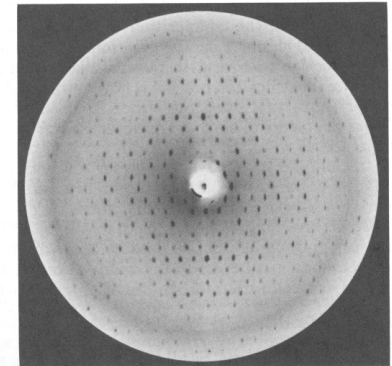


# Precession method

- Uses monochromatic radiation
- Crystal axis precesses about the X-ray beam
- Film is always kept perpendicular to the crystal axis
  - very complex mechanical setup!
- Spot patterns are recorded on film
  - gives true picture of reciprocal lattice
  - useful for lattice and space group determination
  - no longer used since advent of fast diffractometers



a)



(b)

FIGURE 7.14. Two precession photographs of a protein crystal (lac repressor). (a)  $0k\ell$  photograph. (b)  $hk0$  photograph. The  $hk0$  photograph was also shown in Figure 3.15 (Chapter 3).

“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.



# Laue method

- Uses white radiation
- Stationary crystal
  - Fulfillment of Bragg condition is achieved by varying the wavelength
- Spot patterns are recorded on film
  - still in use for determining crystal orientations
  - useful for determining Laue groups from pattern symmetry

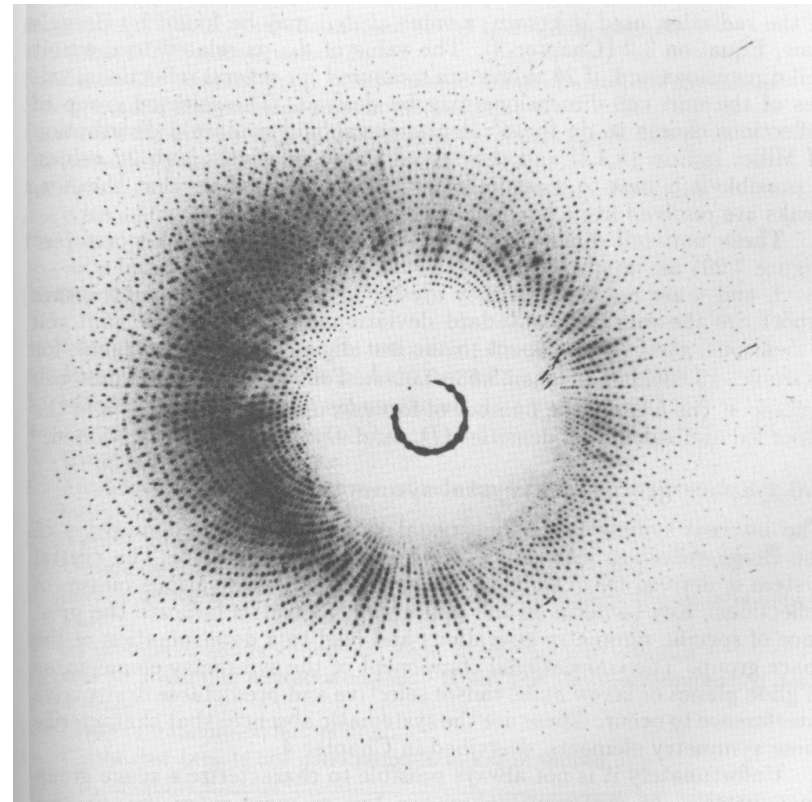
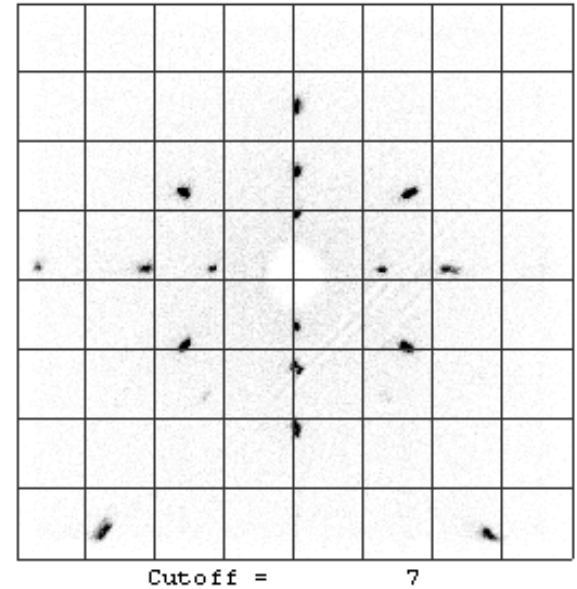
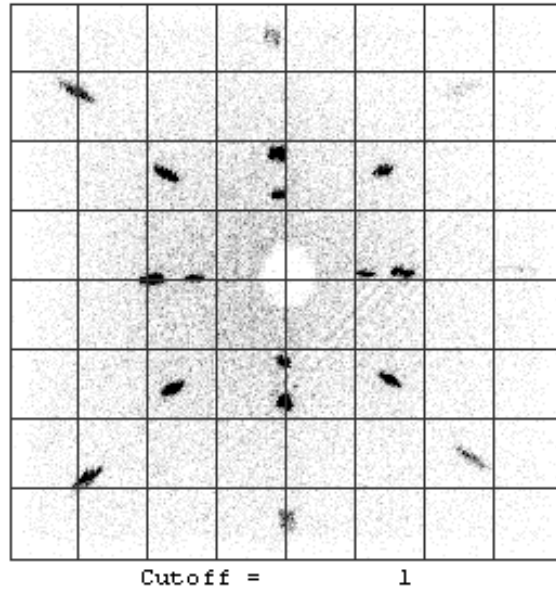
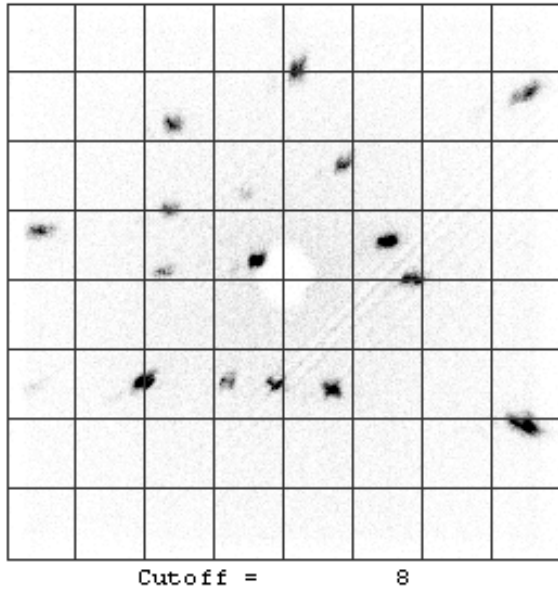


FIGURE 7.19. Laue photograph of lysozyme, 100 msec photograph, with synchrotron radiation 0.2 – 2.0 Å wavelength, taken at CHESS (Cornell High Energy Synchrotron Source). (Courtesy Edwin Westbrook).

“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.



# Laue photographs



PbF<sub>2</sub> Laue photographs: a) unoriented, b) 100, c) 110.

Courtesy of Jonathan P. Wrubel

# Laue groups

- The symmetry in Laue patterns allows to determine the *Laue group* of a crystal
  - can be very useful for narrowing down possible space groups!

For PbF<sub>2</sub>: Reflection conditions are

- hkl: h+k, h+l, k+l
- Okl: k, l
- hhl: h+l
- 00l: l

CUBIC, Laue classes  $m\bar{3}$  and  $m\bar{3}m$

Reflection conditions (Indices are permutable, apart from space group No. 205) <sup>†</sup>				Extinction symbol	Laue class				
					$m\bar{3} (2/m \bar{3})$		$m\bar{3}m (4/m \bar{3} 2/m)$		
					Point group				
hkl	Ok <sup>†</sup>	hhl	00l		23	$m\bar{3}$	432	$\bar{4}3m$	$m\bar{3}m$
			$l$	$P---$ $\{P2_1---$ $P4_2---$ $P4_1---$	$P23 (195)$ $P2_13 (198)$	$Pm\bar{3} (200)$	$P432 (207)$	$P\bar{4}3m (215)$	$Pm\bar{3}m (221)$
	$k^†$ $k+l$ $k+l$	$l$	$l$	$P--n$ $Pa--$ $Pn--$ $Pn--$		$Pa\bar{3} (205)$ $Pn\bar{3} (201)$	$\{P4_232 (208)$ $\{P4_332 (213)$ $\{P4_332 (212)\}$	$P\bar{4}3n (218)$	$Pm\bar{3}n (223)$ $Pn\bar{3}m (224)$ $Pn\bar{3}n (222)$
$h+k+l$	$k+l$	$l$	$l$	$l---$	$[I23 (197)]$ $[I2_13 (199)]$	$Im\bar{3} (204)$	$I432 (211)$	$I\bar{4}3m (217)$	$Im\bar{3}m (229)$
$h+k+l$	$k+l$	$l$	$l=4n$	$I4_1---$			$I4_132 (214)$	$I\bar{4}3d (220)$	
$h+k+l$	$k+l$	$2h+l=4n, l$	$l=4n$	$l---d$					
$h+k+l$	$k, l$	$l$	$l$	$la--$		$la\bar{3} (206)$			
$h+k+l$	$k, l$	$2h+l=4n, l$	$l=4n$	$la-d$					
$h+k, h+l, k+l$	$k, l$	$h+l$	$l$	$F---$	$F23 (196)$	$Fm\bar{3} (202)$	$F432 (209)$	$F\bar{4}3m (216)$	$Ia\bar{3}d (230)$ $Fm\bar{3}m (225)$
$h+k, h+l, k+l$	$k, l$	$h+l$	$l=4n$	$F4_1---$			$F4_132 (210)$		
$h+k, h+l, k+l$	$k, l$	$h, l$	$l$	$F- -c$				$F\bar{4}3c (219)$	$Fm\bar{3}c (226)$ $Fd\bar{3}m (227)$ $Fd\bar{3}c (228)$
$h+k, h+l, k+l$	$k+l=4n, k, l$	$h+l$	$l=4n$	$Fd--$		$Fd\bar{3} (203)$			
$h+k, h+l, k+l$	$k+l=4n, k, l$	$h, l$	$l=4n$	$Fd-c$					

<sup>†</sup> For No. 205 only cyclic permutations permitted. Conditions are Okl: k = 2n; h0l: l = 2n; hkl: h = 2n.

"International Tables for Crystallography, Vol. A", Kluwer, 1993.



# Single crystal diffractometers

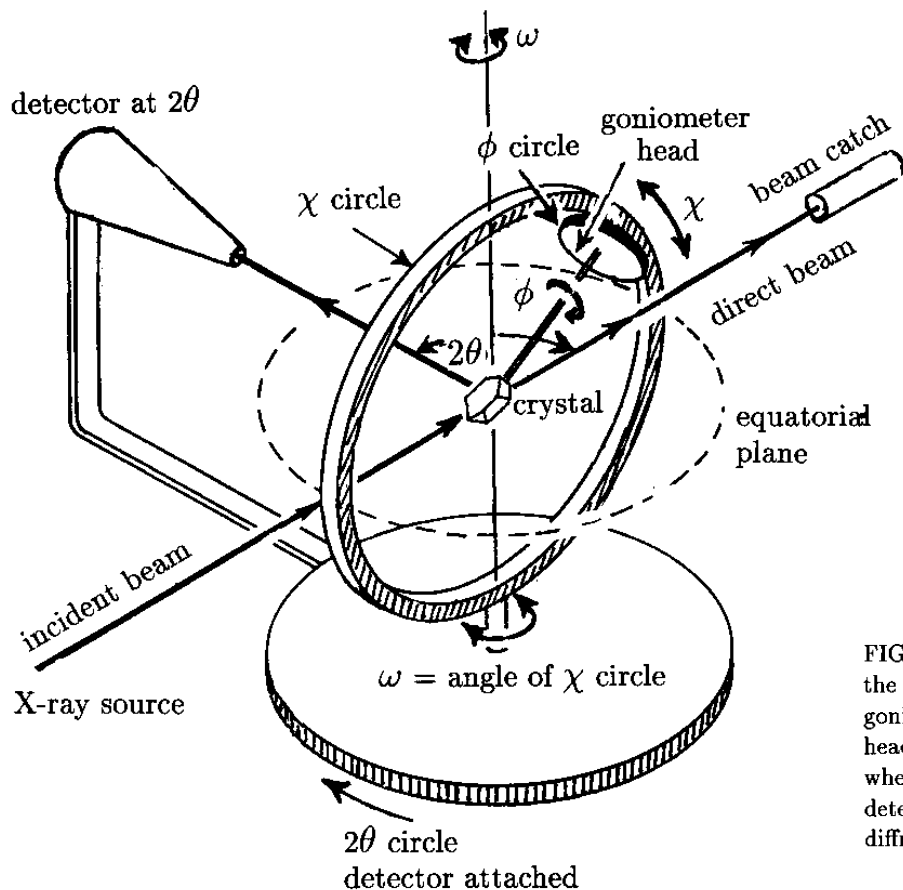


FIGURE 7.16. Diagram of an automatic diffractometer. The crystal is mounted on the goniometer head, which is attached, in turn, to the  $\chi$  circle. The spindle axis of the goniometer head is  $\phi$ . The angle  $\chi$  is the angle between the  $\phi$  axis of the goniometer head and the base of the diffractometer. The  $\chi$  circle can be rotated about the  $\omega$  axis, where  $\omega$  is the angle between the diffraction vector and the plane of the  $\chi$  circle. The detector is moved on the  $2\theta$  circle, where  $2\theta$  is the angle between the incident and diffracted beams.

“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.

# Single crystal diffractometry

- **Uses monochromatic radiation**
  - Mo tube for small molecule crystallography; Cu is useful for absolute structure
  - Nowadays: Mainly Cu for macromolecules
- **Normal approach: Collect a rotation matrix**
  - a series of 60-90 frames that the program will try to index
  - if you cannot index the matrix, the crystal is probably not worth collecting on!
- **Check spot quality**
  - should be single, spherical spots
  - streaky spots or shoulders usually point to low crystal quality, defects, twinning...
- **If everything looks good: Collect a full data set!**
  - normal exposure times: 10 to 60s
  - number of frames depends on crystal symmetry



# Judging spot quality

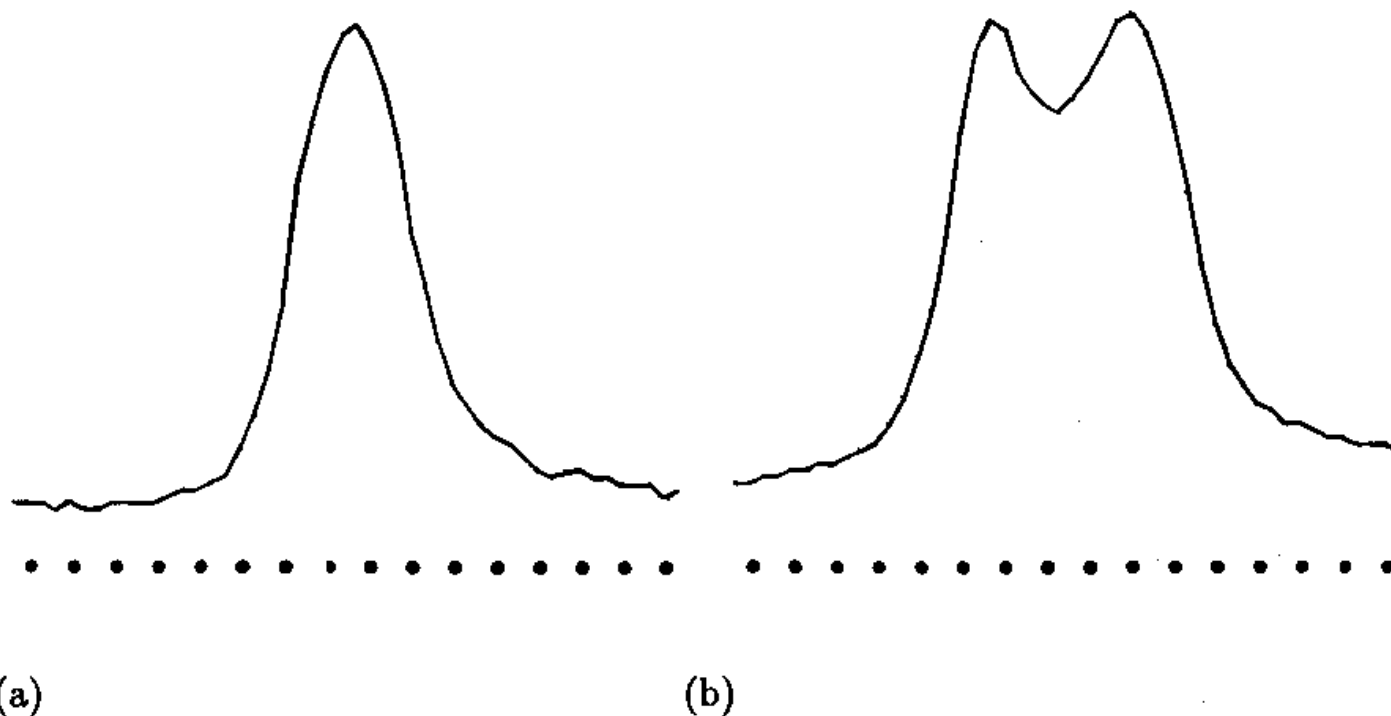


FIGURE 7.18. (a) A good and (b) a bad peak profile.

“Crystal Structure Analysis for Chemists and Biologists”, Glusker, Lewis and Rossi, VCH, 1994.