

Yale Course 2018: 'Advanced Synchrotron and Electron Spectroscopy of Materials' **X-ray Powder Diffraction**

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Lecture Schedule

Yale Course 2018: Advanced Synchrotron and Electron Spectroscopy of Materials

APHY/ENAS/MENG 450/725, Becton Rm.227, Thursdays, 1:30 - 3:30 PM

Date	Lecturer	Topic	Confirmed
1/18/2018	Qun Shen	Introduction / X-ray Interaction with Matter	Y
1/25/2018	Robert Sweet	X-ray Diffraction & Crystallography	Y
2/1/2018	Timur Shaftan	Synchrotron Radiation	Y
2/8/2018	Eric Dooryhee	X-ray Powder Diffraction	Y
2/15/2018	Anibal Boscoboinik	X-ray Photoelectron Spectroscopy	Y
2/22/2018	Bruce Ravel	X-ray Spectroscopy	Y
3/1/2018	1st Exam		
3/8/2018	Yong Chu	X-ray Microscopy	Y
3/15/2018	Spring Break		
3/22/2018			
3/29/2018	Oleg Gang	Small-Angle & Wide-Angle X-ray Scattering	Y
4/5/2018	Ignace Jarrige	X-ray Emission Spectroscopy & RIXS	Y
4/12/2018	Wah-Keat Lee	X-ray Imaging	Y
4/19/2018	Eric Stach	Transmission Electron Microscopy & EELS	Y
4/26/2018	2nd Exam		
5/3/2018	NSLS-II/CFN tour ?		

My Bio

Education:

- University of Paris, PhD, 1987
- Res. Associate, Synchrotron Radiation Source, Daresbury, UK, 88-89

Professional Experience:

- Assoc. Scientist, Res. Center with Ions and Lasers, France, 90-95
- Scientist, European Synchrotron Radiation Facility, France, 96-00
- Scientist, Neel Institute, CNRS Grenoble, France, 01-09
- Scientist, beamline lead, Brookhaven National Lab., NSLS II, 09-

Background: Materials Science, Irradiation Physics, PD, SR

Current position: 'Diffraction & In Situ Scattering' Program Manager

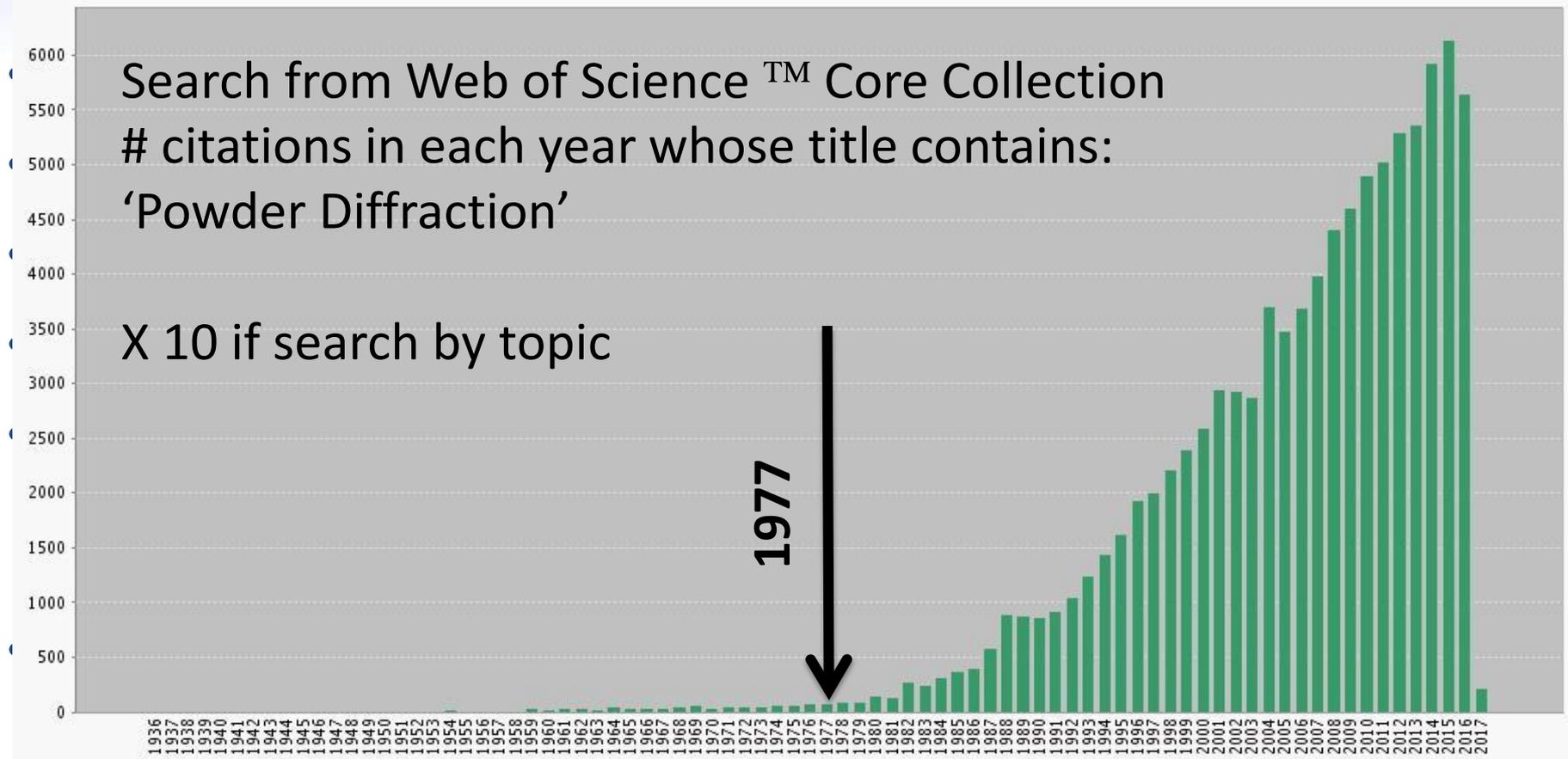
Outline

- 1. Applications for Powder Diffraction**
- 2. The basics of Powder Diffraction**
- 3. Experimental Methods for collecting PD data**
- 4. Crystal structure Determination**
- 5. Peak Profile Analysis for Microstructural Studies**
- 6. Advanced PD techniques**

Short History of Powder Diffraction

- **1895** Discovery of X-rays by Roentgen
- **1912** von Laue demonstrates that X-rays can be diffracted by crystals
- **1935** Le Galley constructs the first X-ray powder diffractometer
- **1947** Phillips introduces the first commercial powder diffractometer
- **1950-60's** Powder diffraction used primarily to study structural imperfections, phase identification, ... largely by metallurgists and mineralogists
- **1969** Hugo Rietveld develops a method for whole pattern analysis of neutron powder diffraction data
- **1977** Cox, Young, Thomas and others first apply Rietveld method to laboratory and synchrotron X-ray data

Short History of Powder Diffraction



- **1977** Cox, Young, Thomas and others first apply Rietveld method to laboratory and synchrotron X-ray data

Examples of Powders

*Multi-grained
i.e. each grain can
diffract x-rays,
neutrons or electrons*

*Polycrystalline
materials*

Examples of Powders



Wood fiber board



Glass ware



Metal parts



Ground coffee



Powdered milk



Cosmetic powder



Gunpowder

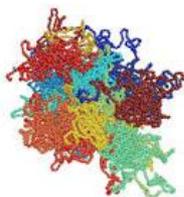


Powdered sugar



Flour

Polymers



Multi-grained
i.e. each grain can diffract x-rays, neutrons or electrons

Polycrystalline materials



Fine snow



Potassium bromate



Zymin



Brightening agent



Potassium bromate



Pharmaceuticals

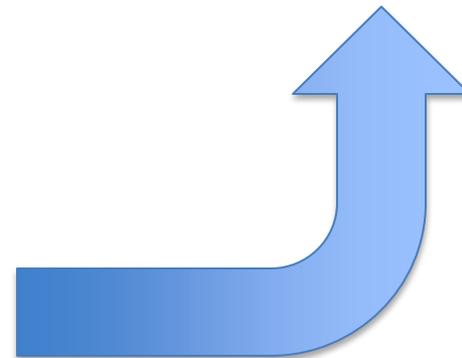


Ceramics

Examples of Powders

*Multi-grained
i.e. each grain can
diffract x-rays,
neutrons or electrons*
**Polycrystalline
materials**

**Highly disordered
Nanocrystalline
Amorphous**



Use of Powder Diffraction

Single Crystal Diffraction is preferred in most cases BUT:

- the product naturally occurs as a powder
- the functionalized (working) form of the product is a powder
- a single crystal of sufficient size or quality cannot be synthesized
- Lattice parameters and micro-structure
- Crystal may shatter during a phase transformation

Use of Powder Diffraction

1. Qualitative Analysis

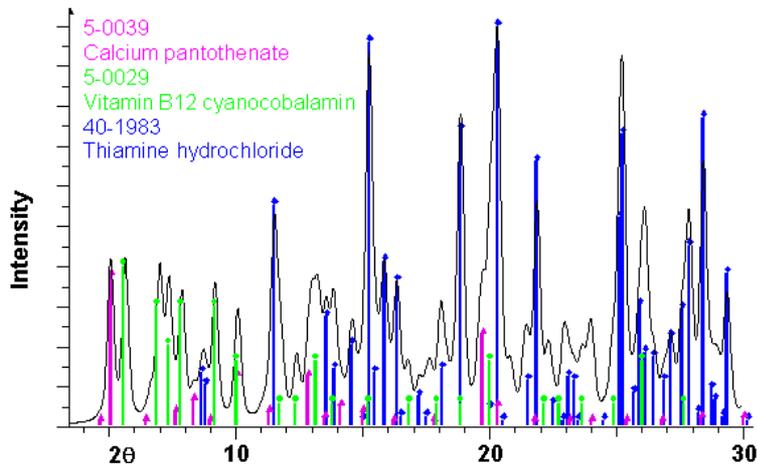
Phase ID

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Phase identification

e.g., **Mineralogy**: natural Granodiorite is a mix of Quartz, Feldspar, Albite, Biotite; lesser Clinocllore, Hornblende; traces of Zircon (see webmineral.com)

e.g., **Pharmaceutical** : Mannitol, Sucrose, DL-Valine, Starch, Nizatidine.



Use of Powder Diffraction

1. Qualitative Analysis

Phase ID

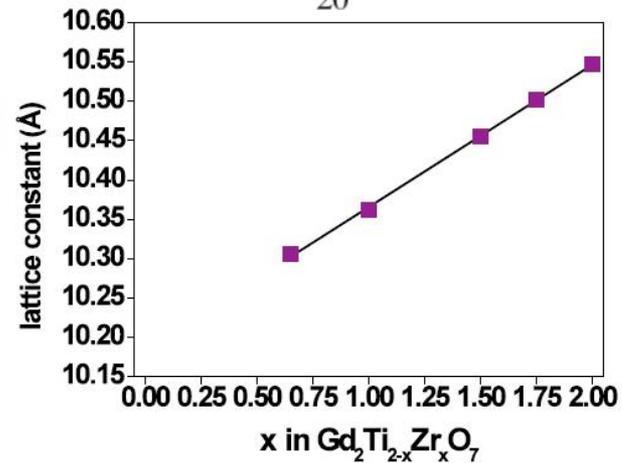
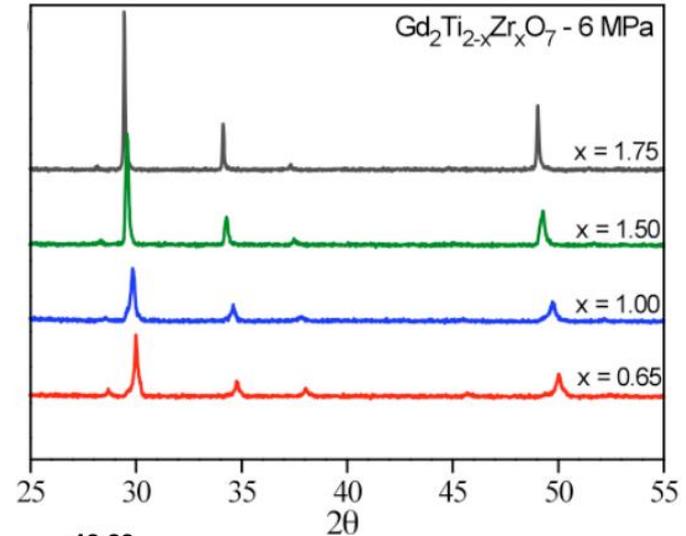
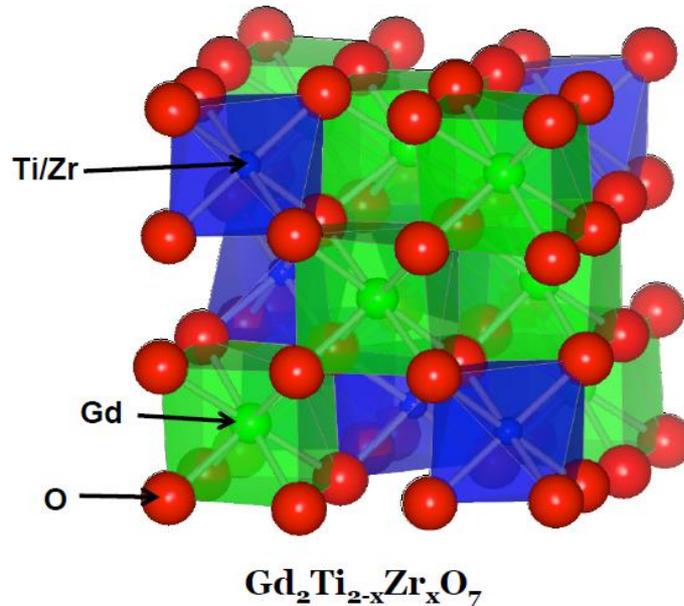
2. Quantitative Analysis

Lattice parameters (thermal expansivity, compressibility), indexing

Phase mass fraction (mixture)

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Effect of substitution

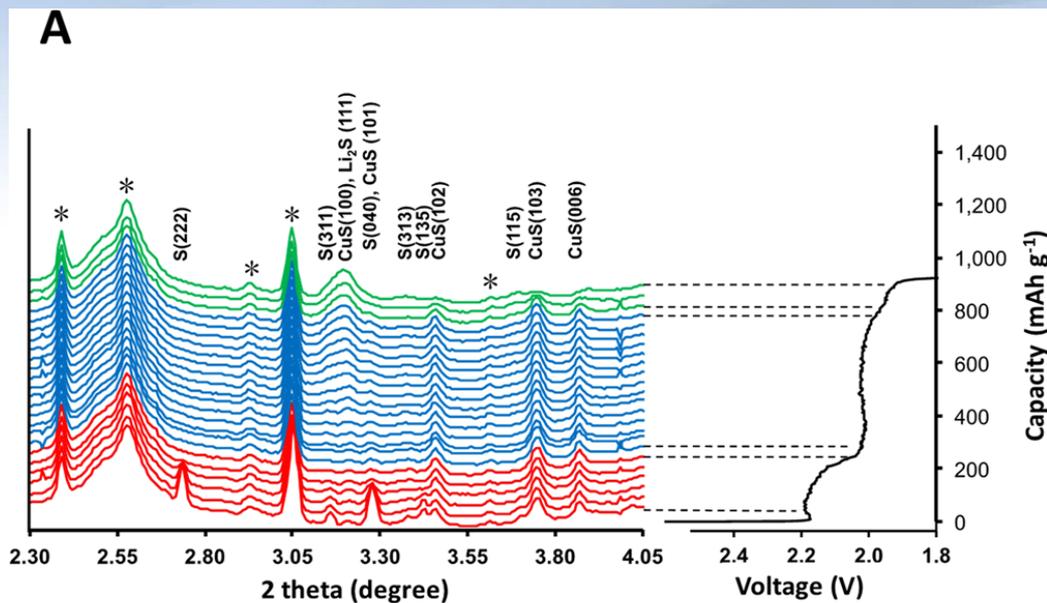


12

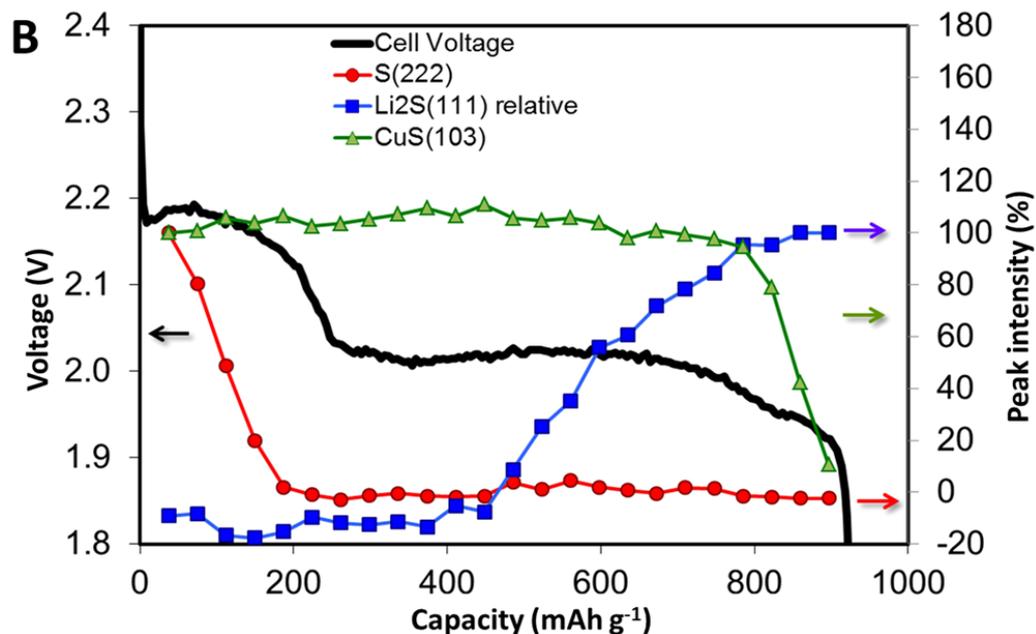
Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Quantitative analysis

- phase content: weight % of each phase inside a mixture
- $A + B \rightarrow A + C + D \rightarrow A + D$
identify and measure the kinetics: mass fractions $f(\text{time})$
- follow structural changes and phase transformations as a function of a process variable
- Structural behavior under complex environments



(A) *Operando* XPD patterns from the sulfur-CuS hybrid electrode aligned with its electrochemical capacity-voltage profile during lithiation; diffraction peaks from the cell are labelled with “*”



(B) Diffracted intensity of S (222), CuS (103), and Li₂S (111) are plotted as a function of depth of discharge in comparison with the cell discharge voltage profile (solid black lines)

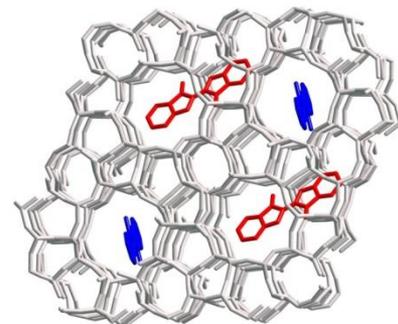
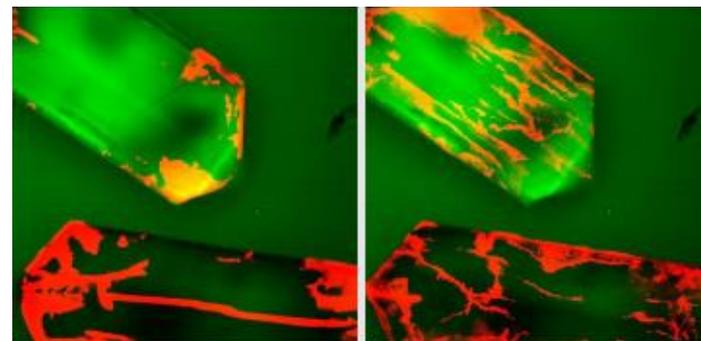
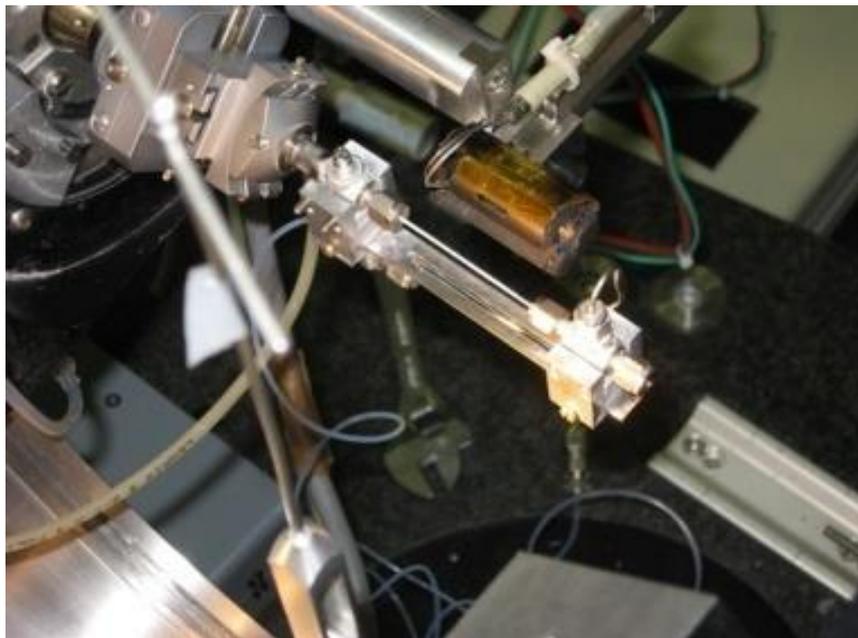
H. Gan et al. Scientific Reports (2017)

Stabilization of an organic dye in a mesoporous Al-Si mineral.

Dejoie, Martinetto, Dooryhee *et al.* Applied Spectroscopy 64 (2010) and Appl. Mater. Interfaces 2 (2010)

Design durable, non-toxic hybrid pigments (*e.g.* in replacement of fading paintings): stable color (irradiation, chemical).

In-situ synchrotron diffraction



Use of Powder Diffraction

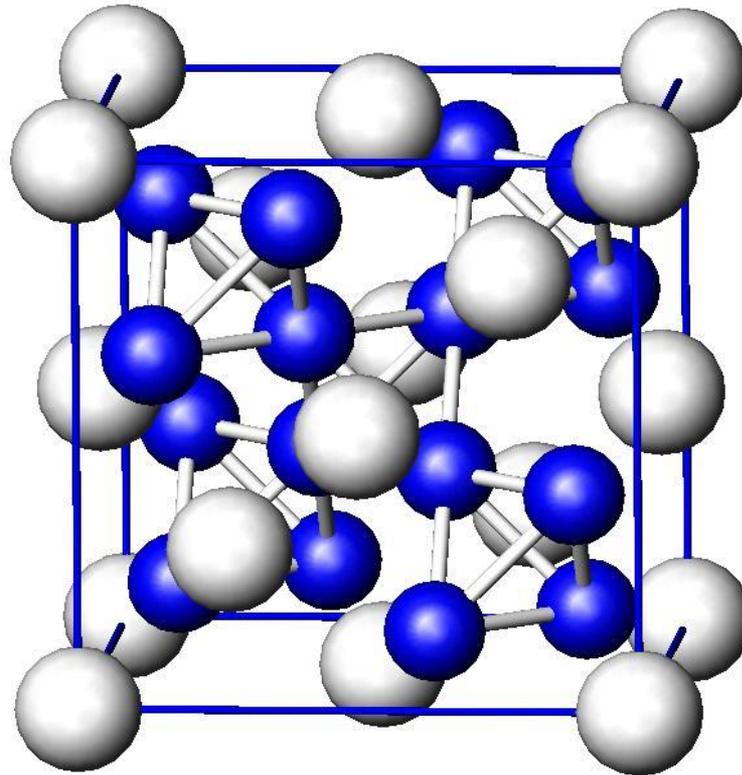
- 1. Qualitative Analysis** Phase ID
- 2. Quantitative Analysis** Lattice parameters (thermal expansivity, compressibility), indexing
Phase mass fraction (mixture)
- 3. Structure Determination** Reciprocal Space Methods (*ab initio*)
Real Space Methods

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Structure determination (crystallography: unit cell and lattice parameters, symmetry, atomic motif)

The entire, ordered and periodic stack of atoms can be generated once the above information is determined from the diffraction data.

The properties of the material are dependent on the crystal structure.



cubic
lattice constant a ,
space group is $Fd\bar{3}m$,
atomic motif is $MgCu_2$

Mg in grey, Cu in blue

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- 5. Microstructure and Texture** Crystallite Size Distribution
Preferred Orientation
Microstrain
Anti-phase domains, stacking faults,...

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Texture (orientation) of polycrystalline materials

Powder diffraction essentially assumes crystallites are randomly oriented
IF NOT, diffraction intensities are affected and this can be useful information:
texture, preferred orientation → found in many engineered materials, and
have a great influence on materials properties

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...



Large Superconducting cables
(American SuperConductor)

Texture (orientation) of polycrystalline materials

Powder diffraction essentially assumes crystallites are randomly oriented
IF NOT, diffraction intensities are affected and this can be useful information:
texture, preferred orientation → found in many engineered materials, and
have a great influence on materials properties

e.g., **HT superconductor thin films** deposited on metallic ribbons (critical
currents in $\text{YBa}_2\text{Cu}_3\text{O}_7$)

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...



Texture (orientation) of polycrystalline materials

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texture, preferred orientation → found in many engineered materials, and
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e.g., **geologic rocks** show texture due to their thermo-mechanic history of
formation processes

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...

Residual stress analysis

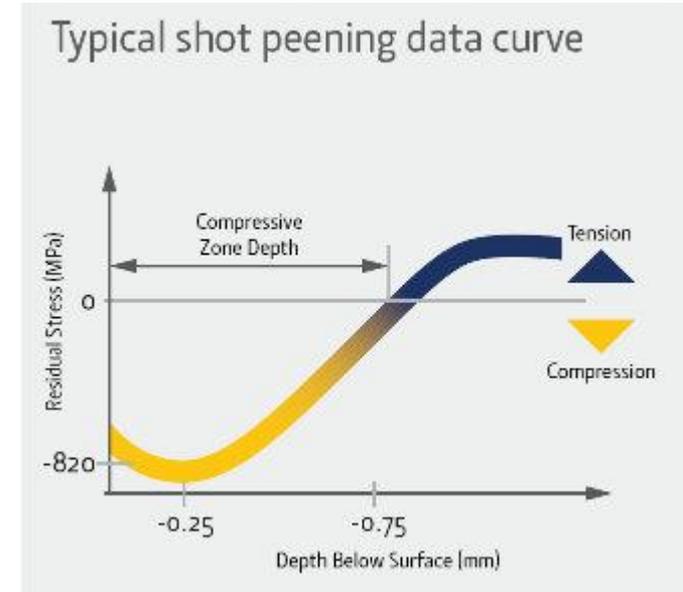
created during the manufacturing process, or it may accumulate in a structure over many years in operation.

stress can have a serious negative effect on a product's quality, durability and lifetime (QA, service)

Applications of Powder Diffraction: Physics, Chemistry, Geology, Industry,...



Peening is a working process to produce a compressive stress at the surface to overcome failure due to metal fatigue, stress corrosion cracking, fretting,...



Residual stress analysis

created during the manufacturing process, or it may accumulate in a structure over many years in operation.

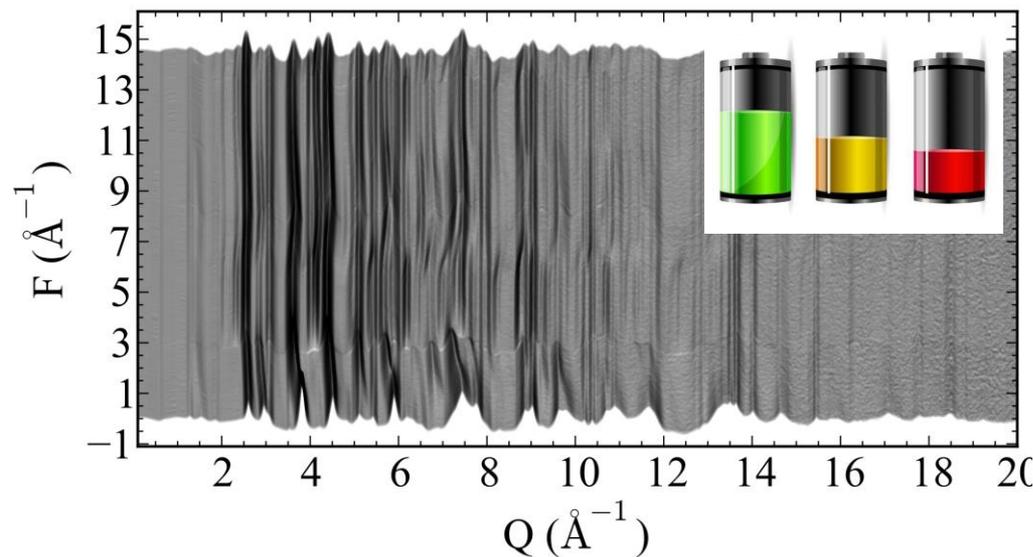
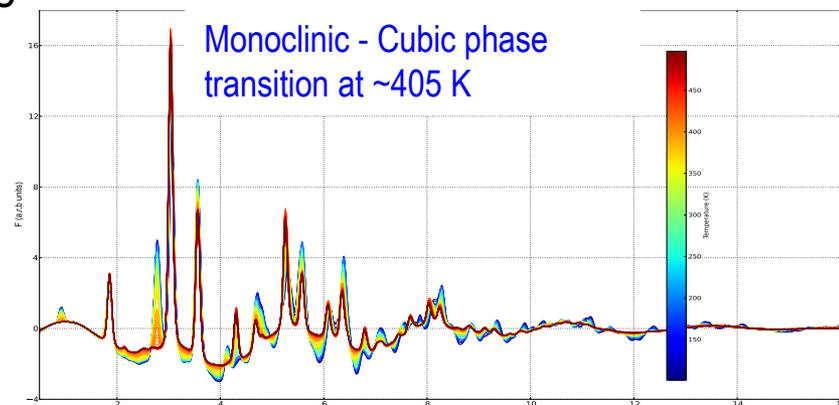
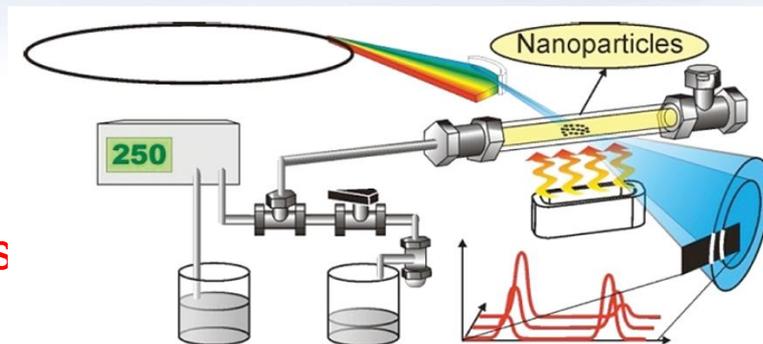
stress can have a serious negative effect on a product's quality, durability and lifetime (QA, service)

e.g., **laser peening** in aircraft engine blades

Applications of Powder Diffraction

A versatile and powerful technique for time-dependent phase transformations.

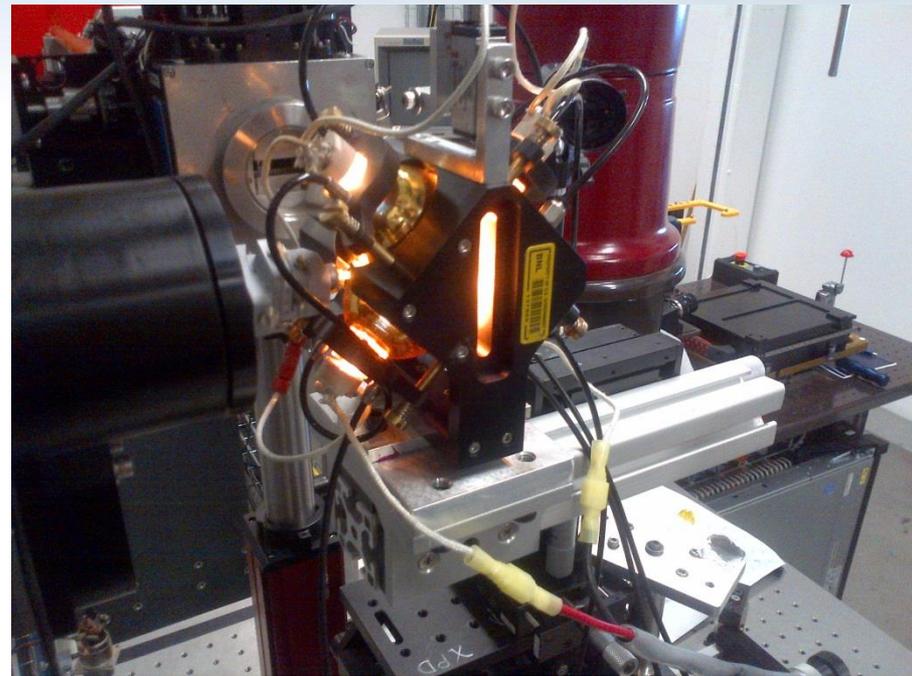
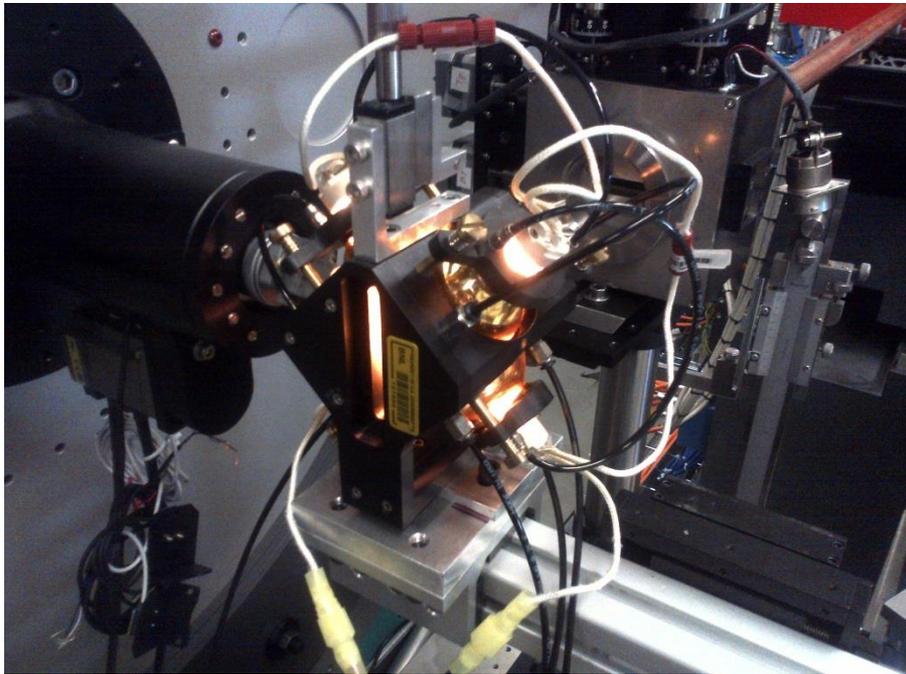
- Kinetics and mechanism of how structures form during reversible or irreversible **chemical reactions**
- How **materials work in devices** under real working conditions. e.g.: batteries, fuel cells.
- Functioning of **catalyst** during reactions.
- **Phase transitions.**



Develop oxidation-resistant Ceramics

In-situ, high temperature oxidation studies of HfB_2 composites using a quadrupole lamp furnace.

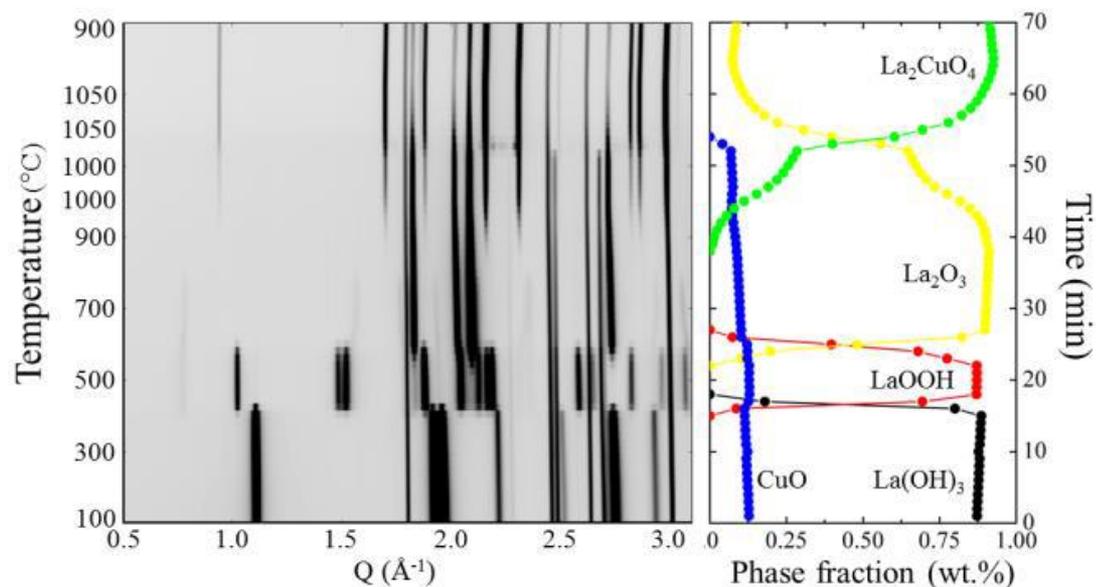
P. Sarin, Univ. of Oklahoma



Develop oxidation-resistant HfB_2 and HfB_2 -SiC composites as emerging **Ultra High Temperature Ceramics** for applications beyond 2,000°C (thermal protection for atmospheric re-entry vehicles)

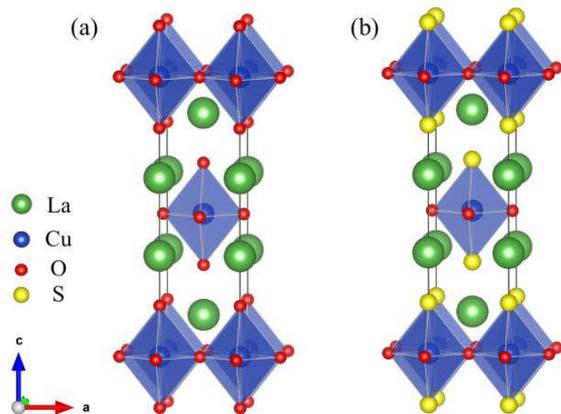
Solid-state syntheses → to accelerate materials discovery. Example: La-Cu-O-S system

- Theory to identify desired materials that are thermodynamically stable e.g. $\text{La}_2\text{CuO}_3\text{S}$ and $\text{La}_2\text{CuO}_2\text{S}_2$. However experimental realization of never reported.
- in situ XRD synthesis to pinpoint the feasible synthetic routes.



In situ powder XRD patterns showing how the diffraction peaks (left panel) and phase fractions (right panel) evolve as the temperature changes for synthesis from La_2O_3 + CuO .

Crystal structures of La_2CuO_4 (a) and the theoretical compound $\text{La}_2\text{CuO}_2\text{S}_2$ (b), which is derived by replacing the apical oxygen in La_2CuO_4 with sulfur.

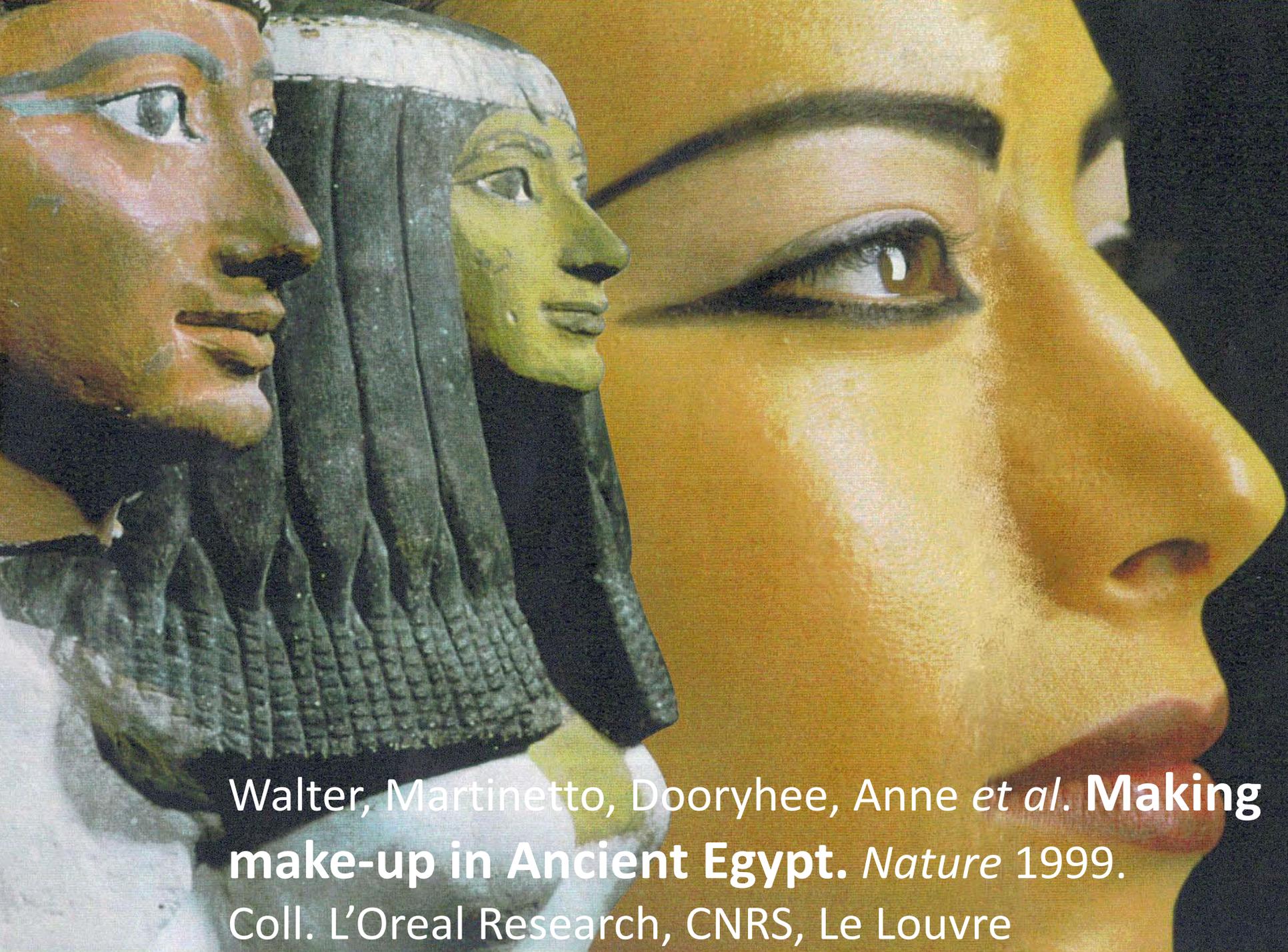


Hua He et al. Submitted to PNAS

Use of Powder Diffraction

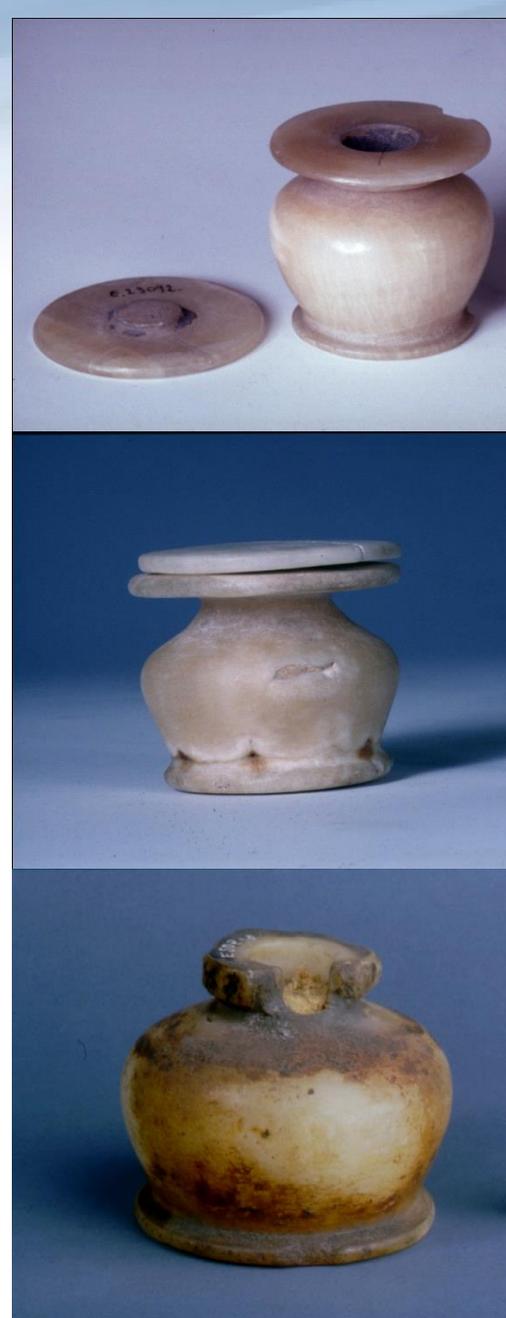
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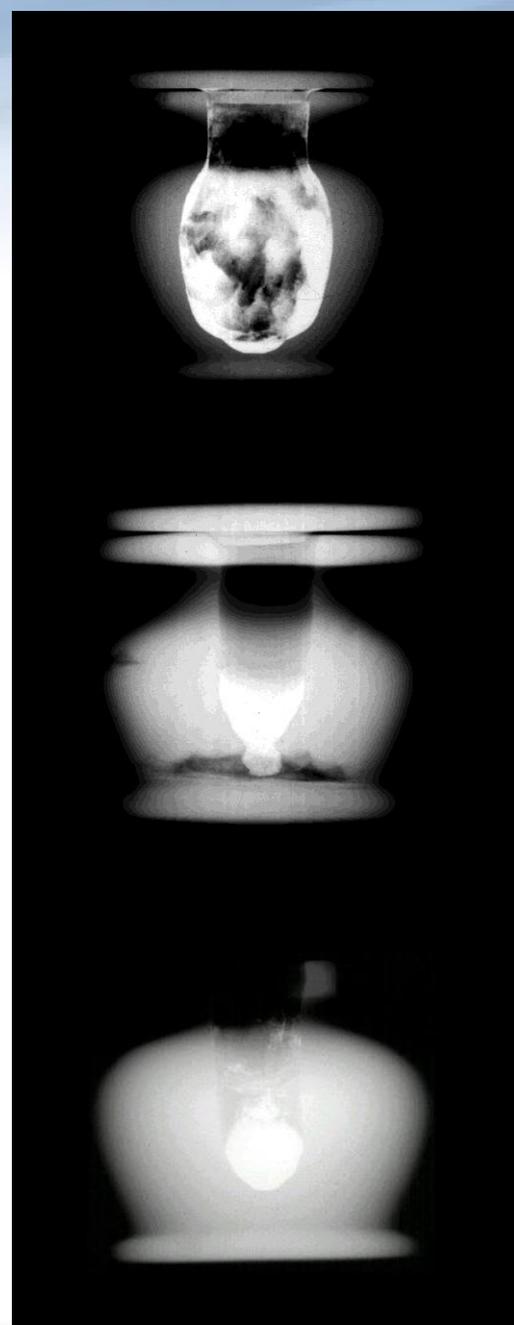
When Powder Diffraction meets Art



Walter, Martinetto, Dooryhee, Anne *et al.* **Making make-up in Ancient Egypt.** *Nature* 1999.
Coll. L'Oreal Research, CNRS, Le Louvre

Cosmetic,
social,
and ritual
functions
of the
make-up





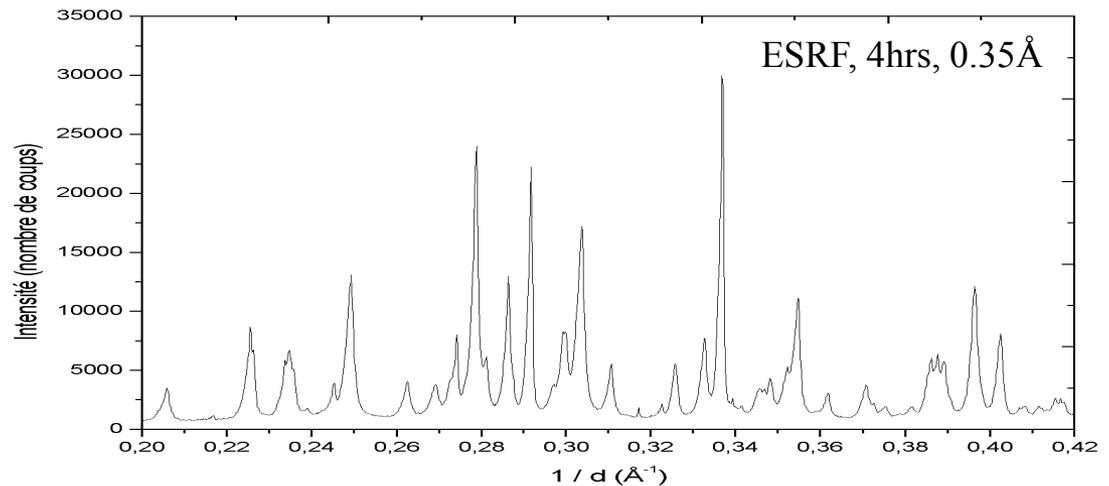
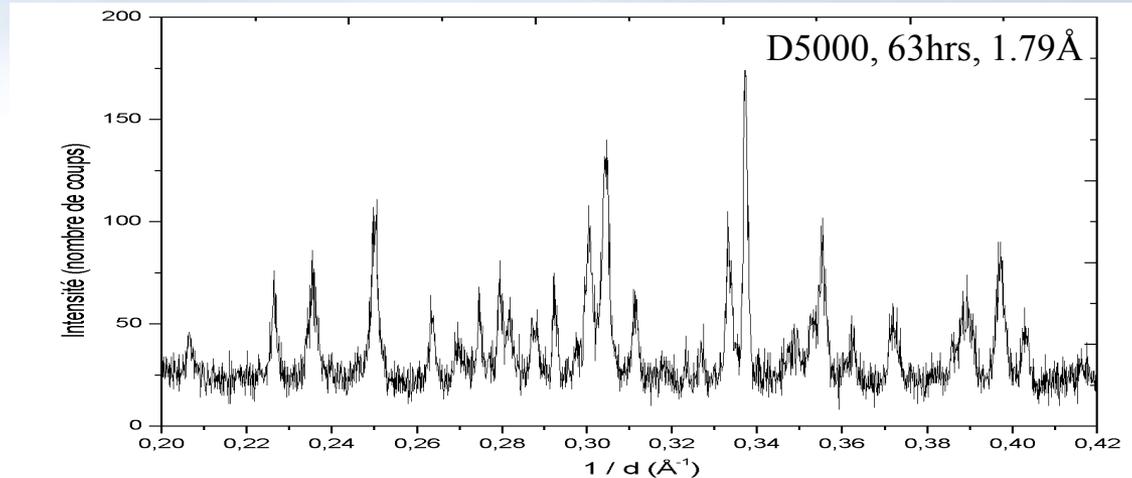
Egyptian Cosmetics

important role in everyday practices and during ritual and burial ceremonies

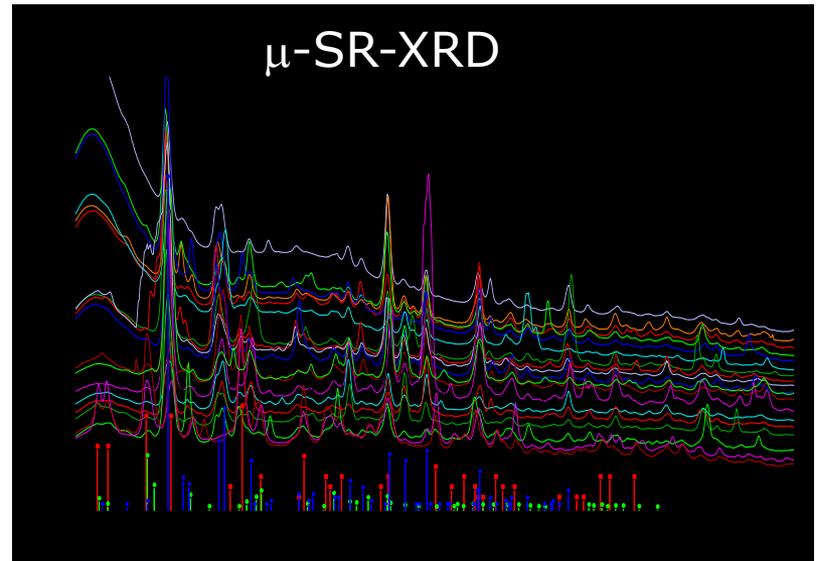
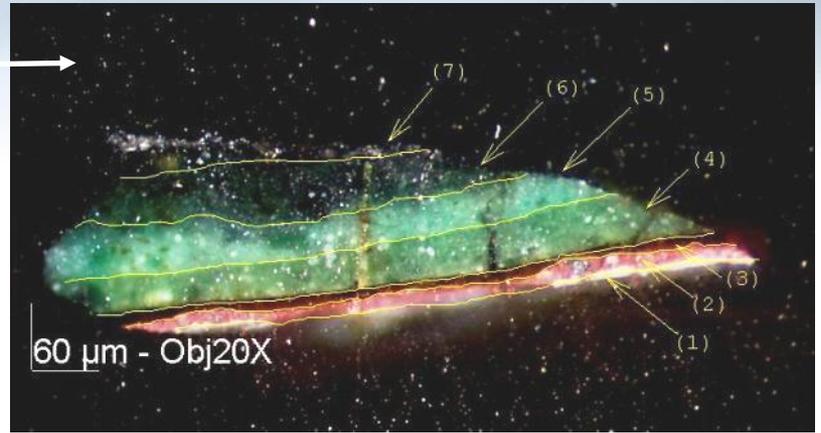
(from P. Martinetto)



Ph. Walter et al. Nature
397 (1999)



High resolution and flux (ESRF) and high energy (Pb-mixtures)

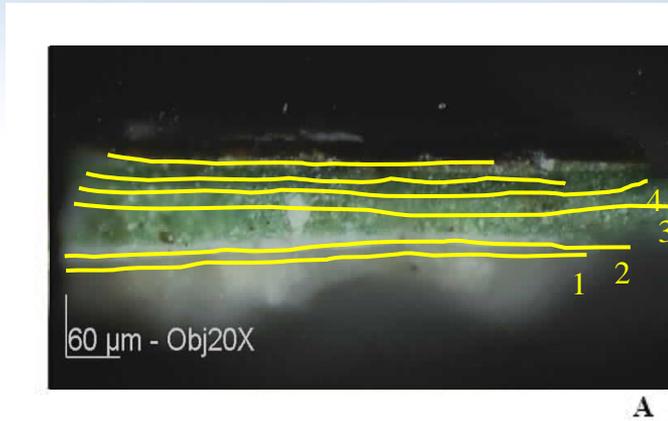
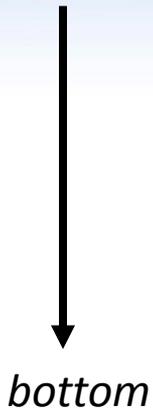


M. Grunewald – German Renaissance

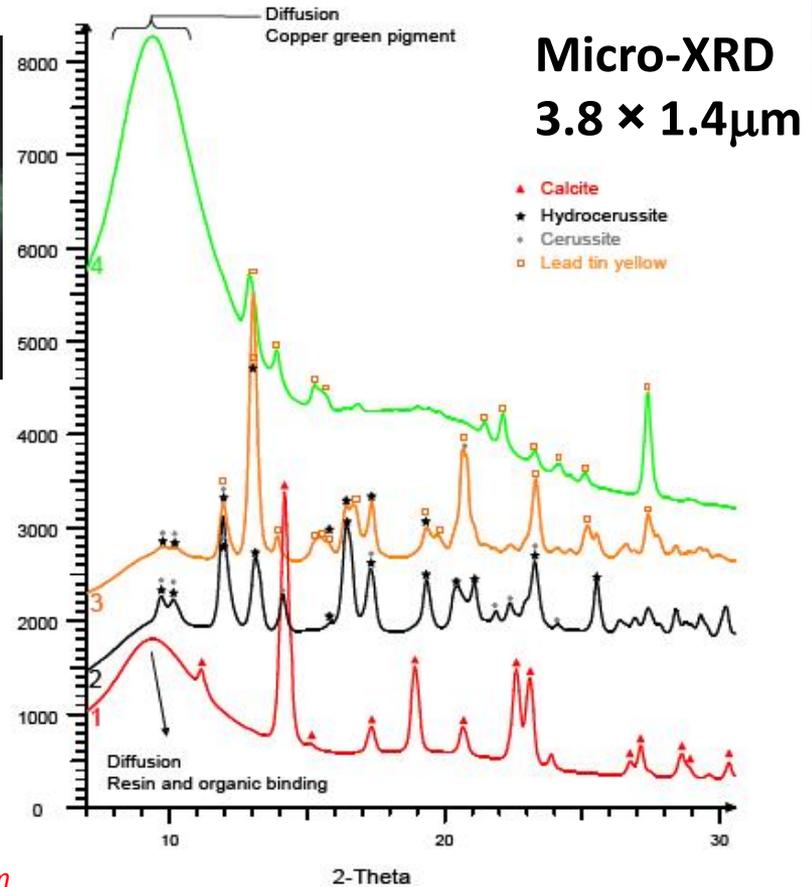
Welcomme E, Walter P, Bleuet P, Hodeau JL, Dooryhee E, Martinetto P, Menu M. Applied Physics A89 (2007)

Synchrotron Analysis of Oil Paintings

top



A

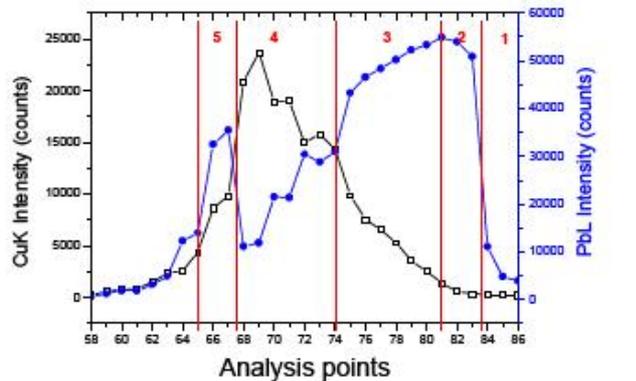


top



bottom

Micro-XRF
3.8x1.4 μm



B

top → bottom

Welcomme, Walter, Dooryhee *et al.* Appl. Phys. A83 (2006)

Layer 1: calcium carbonate preparation layer; Layer 2: priming layer made of lead carbonates (hydrocerussite, cerussite); Layers 3-5: green copper pigments with lead tin yellow $2PbO-SnO_2$ and lead carbonates; Layer 6: organic, varnish.

Outline

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- 2. The basics of Powder Diffraction**
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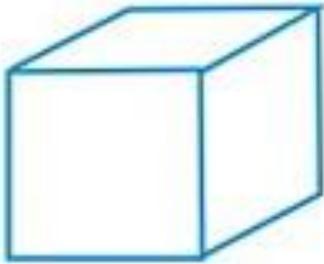
What is a **crystal** ?

- A **periodic arrangement** of objects (atoms, molecules) repeating in three dimensions
- **Crystal systems**

Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

What is a **crystal** ?

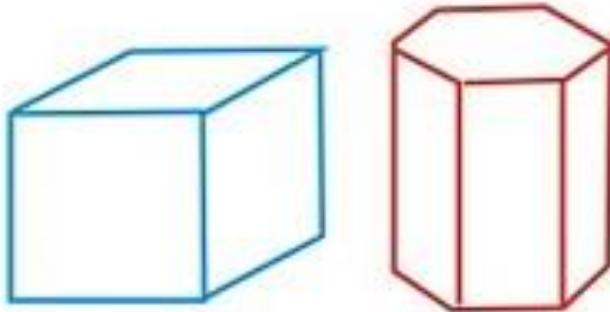
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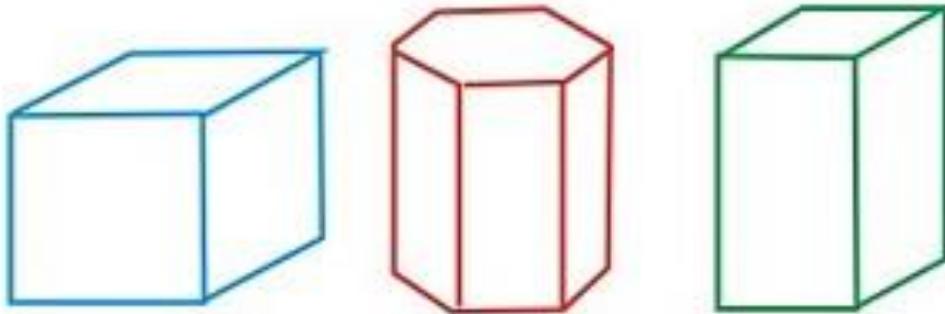
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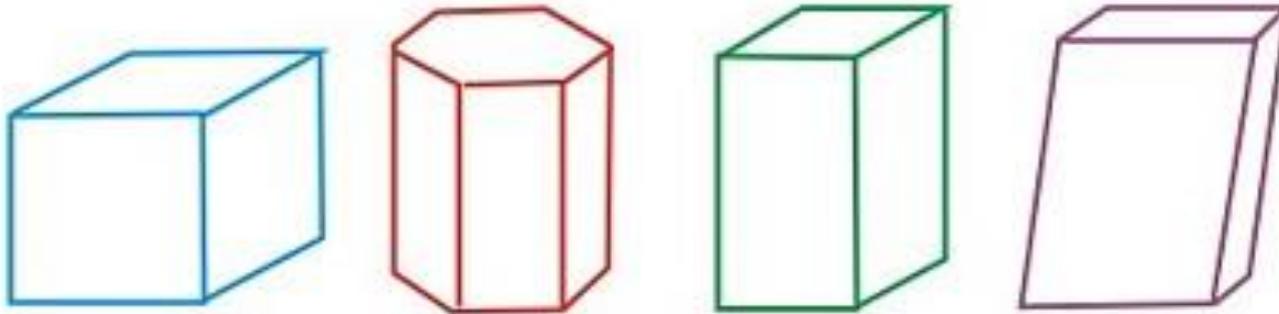
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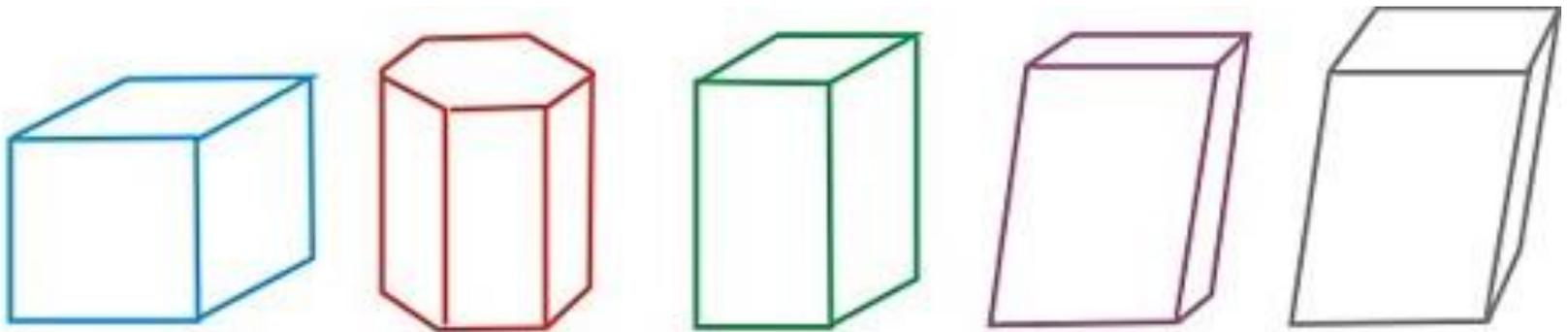
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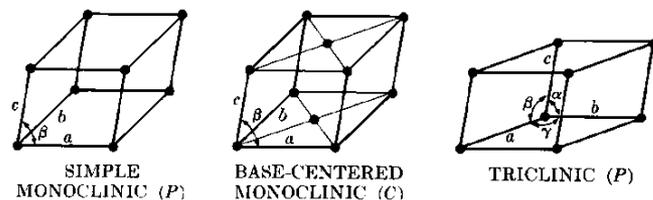
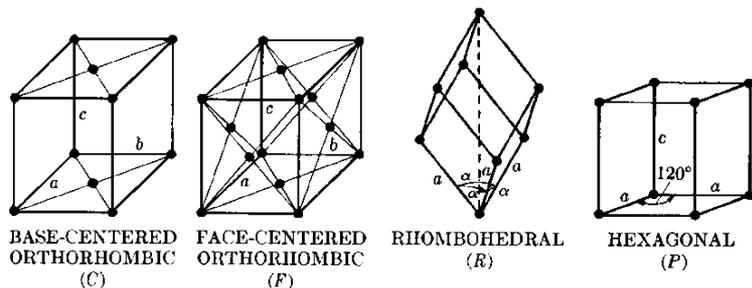
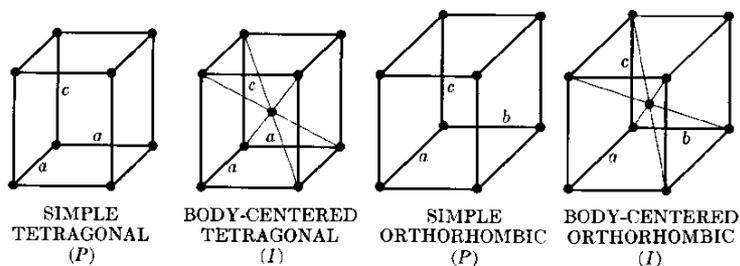
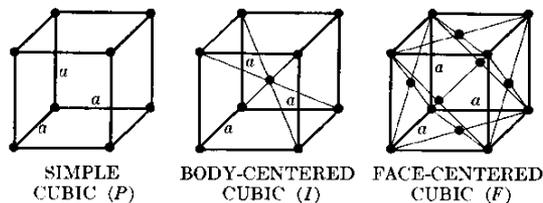
What is a **crystal** ?

- A **periodic arrangement** of objects (atoms, molecules) repeating in three dimensions
- **Crystal systems (7)**



Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

The 14 Bravais lattices



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

What is a **crystal** ?

Crystal System	Bravais Types	External Minimum Symmetry	Unit Cell Properties
Triclinic	P	None	a, b, c, α , β , γ ,
Monoclinic	P, C	One 2-fold axis, parallel b (b unique)	a, b, c, 90, β , 90
Orthorhombic	P, I, F, C	Three perpendicular 2-folds	a, b, c, 90, 90, 90
Tetragonal	P, I	One 4-fold axis, parallel c	a, a, c, 90, 90, 90
Trigonal	P*, R	One 3-fold axis	a, a, c, 90, 90, 120
Hexagonal	P*	One 6-fold axis	a, a, c, 90, 90, 120
Cubic	P, F, I	Four 3-folds along space diagonal	a, a, a, 90, 90, 90

Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

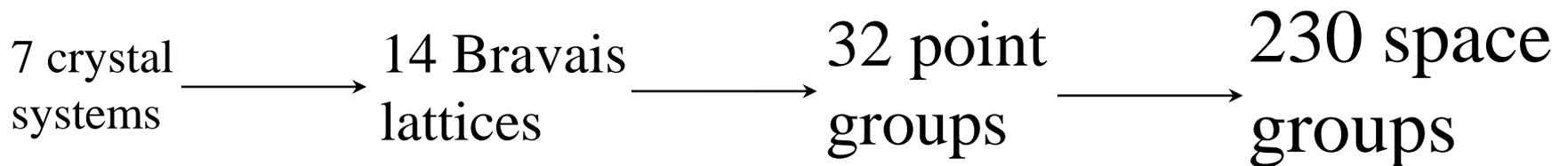
Combining symmetry operations

- ◆ An object can possess several symmetry elements
- ◆ Not all symmetry elements can be combined arbitrarily
 - for example, two perpendicular two-fold axes imply the existence of a third perpendicular two-fold
- ◆ Translational symmetry in 3D imposes limitations
 - only 2, 3, 4 and 6-fold rotation axes allow for space filling translational symmetry
- ◆ The allowed combinations of symmetry elements are called point groups
 - there are 32 point groups that give rise to periodicity in 3D

Courtesy of Cora Lind-Kovacs

Space groups

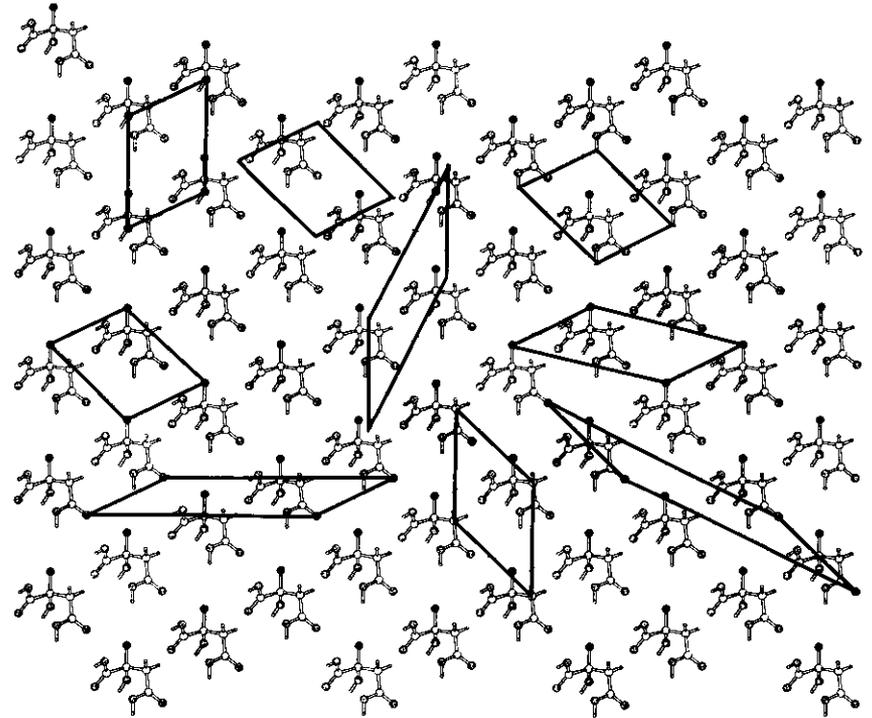
- ◆ Crystals are usually described by a *space group*
- ◆ Space groups are made up from
 - lattice symmetry (translational)
 - point symmetry (not translational)
 - glide and/or screw axes (some translational component)
- ◆ There are 230 space groups



Courtesy of Cora Lind-Kovacs

Lattice symmetry

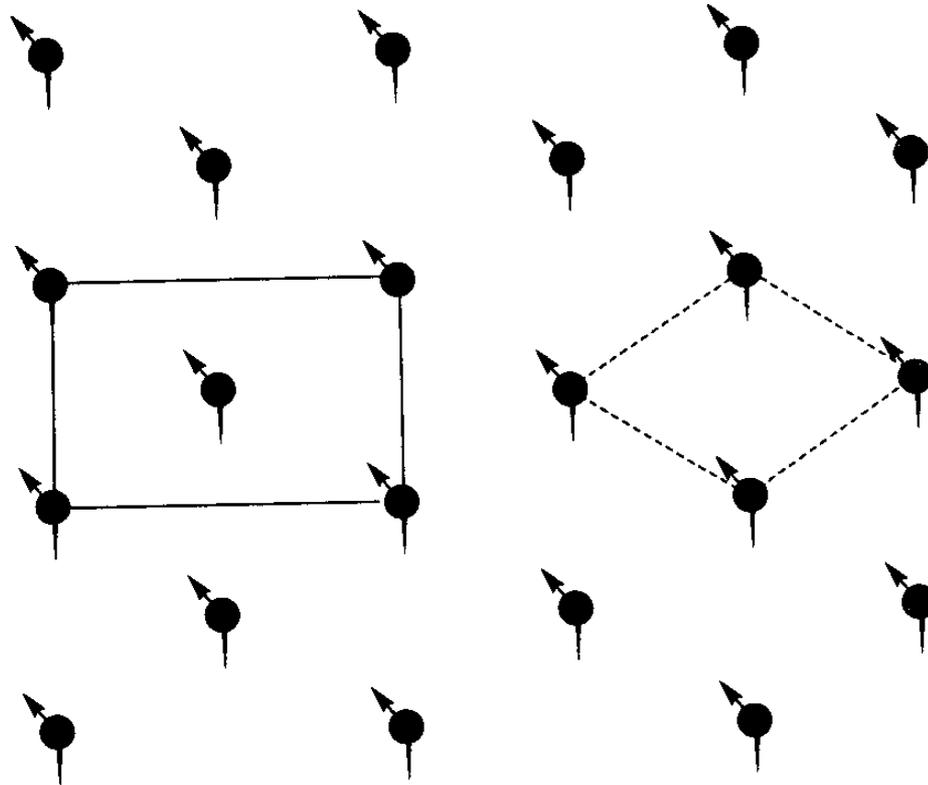
- ◆ Without rules, an infinite number of different unit cells could describe any given lattice
- ◆ Choose *smallest cell with highest symmetry*
 - Primitive or centered cells



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

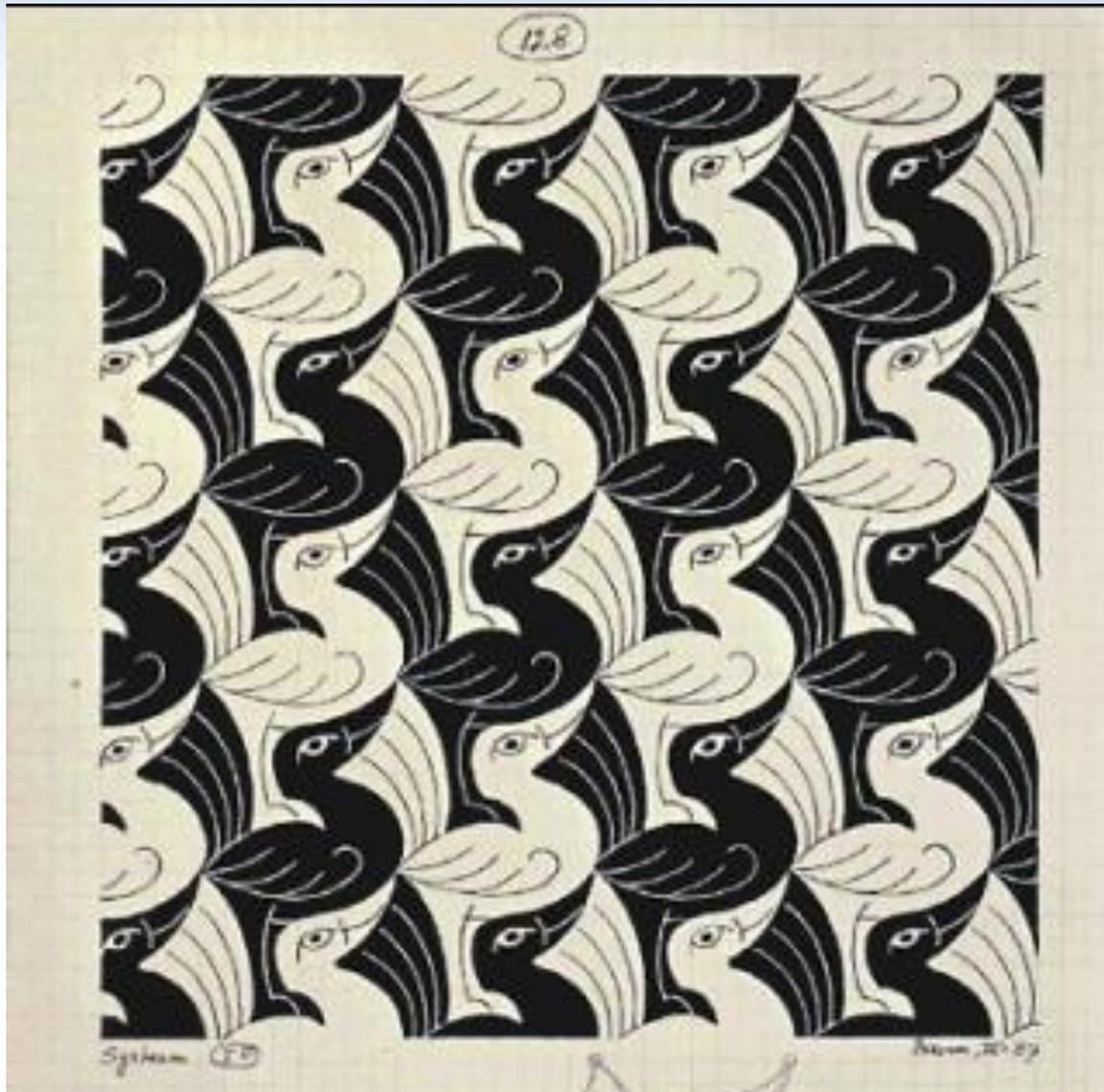
Courtesy of Cora Lind-Kovacs

An example of unit cell choice



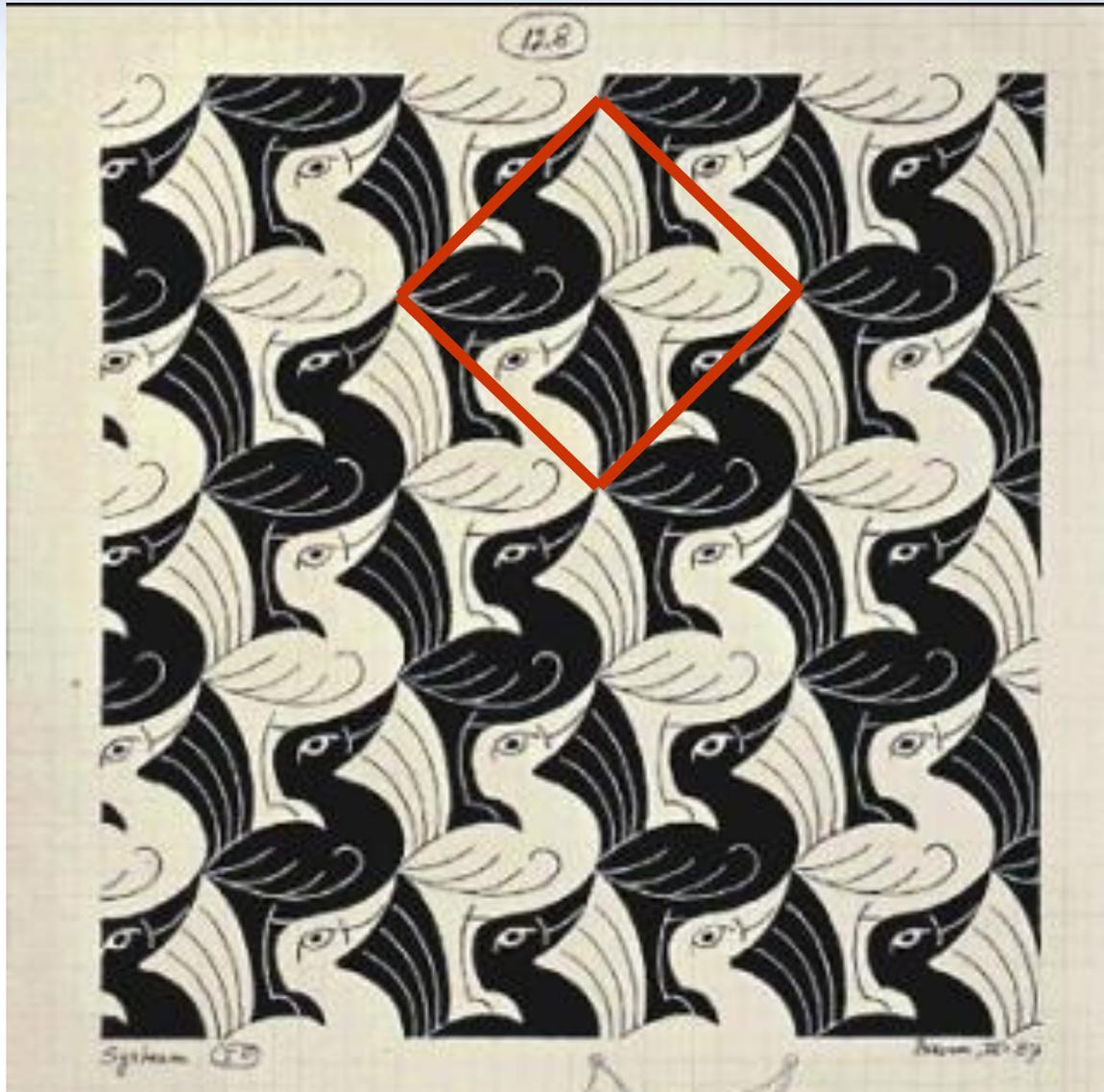
◆ According to our definitions, the centered cell would be preferred

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



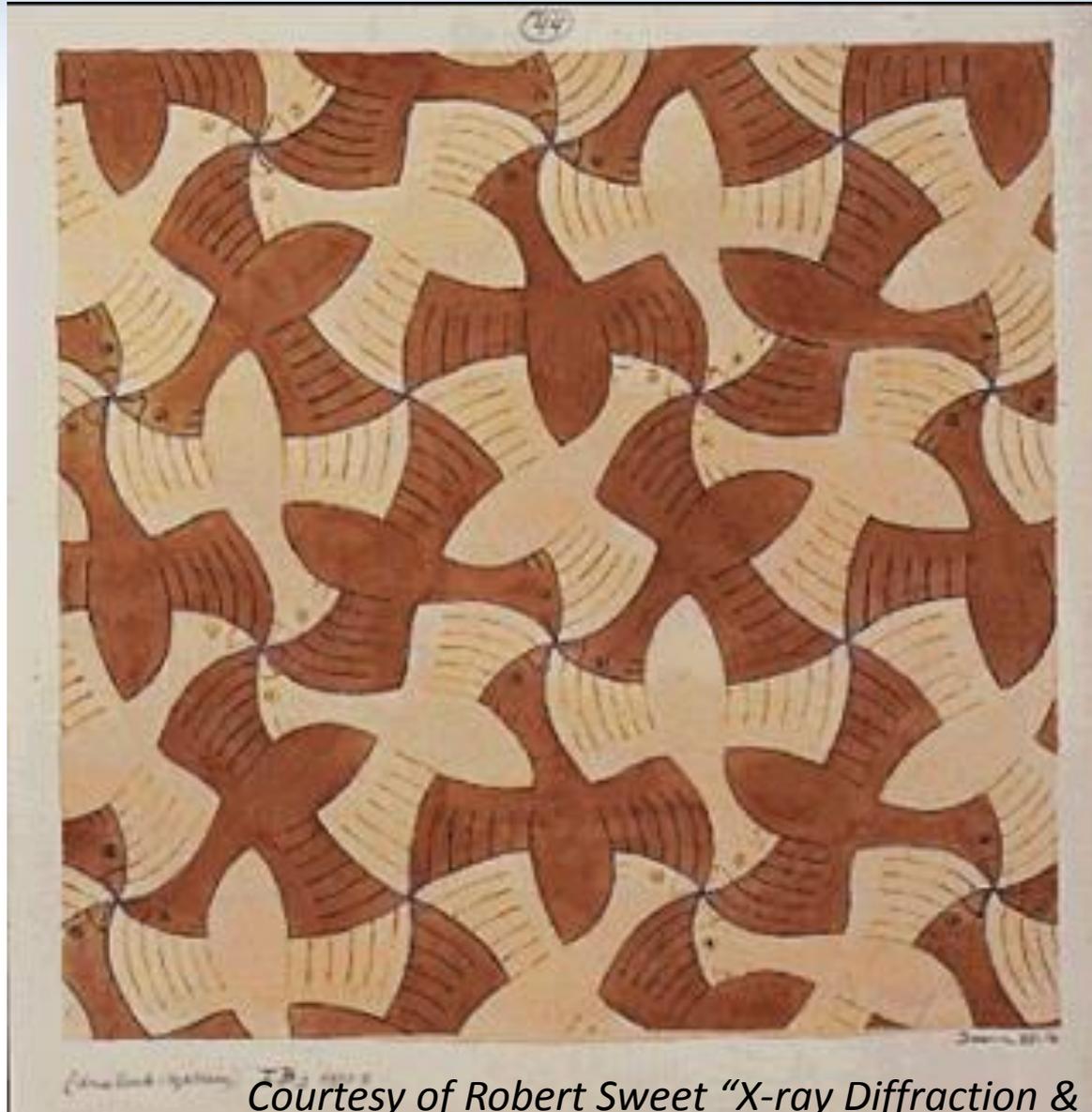
Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

Translation only.



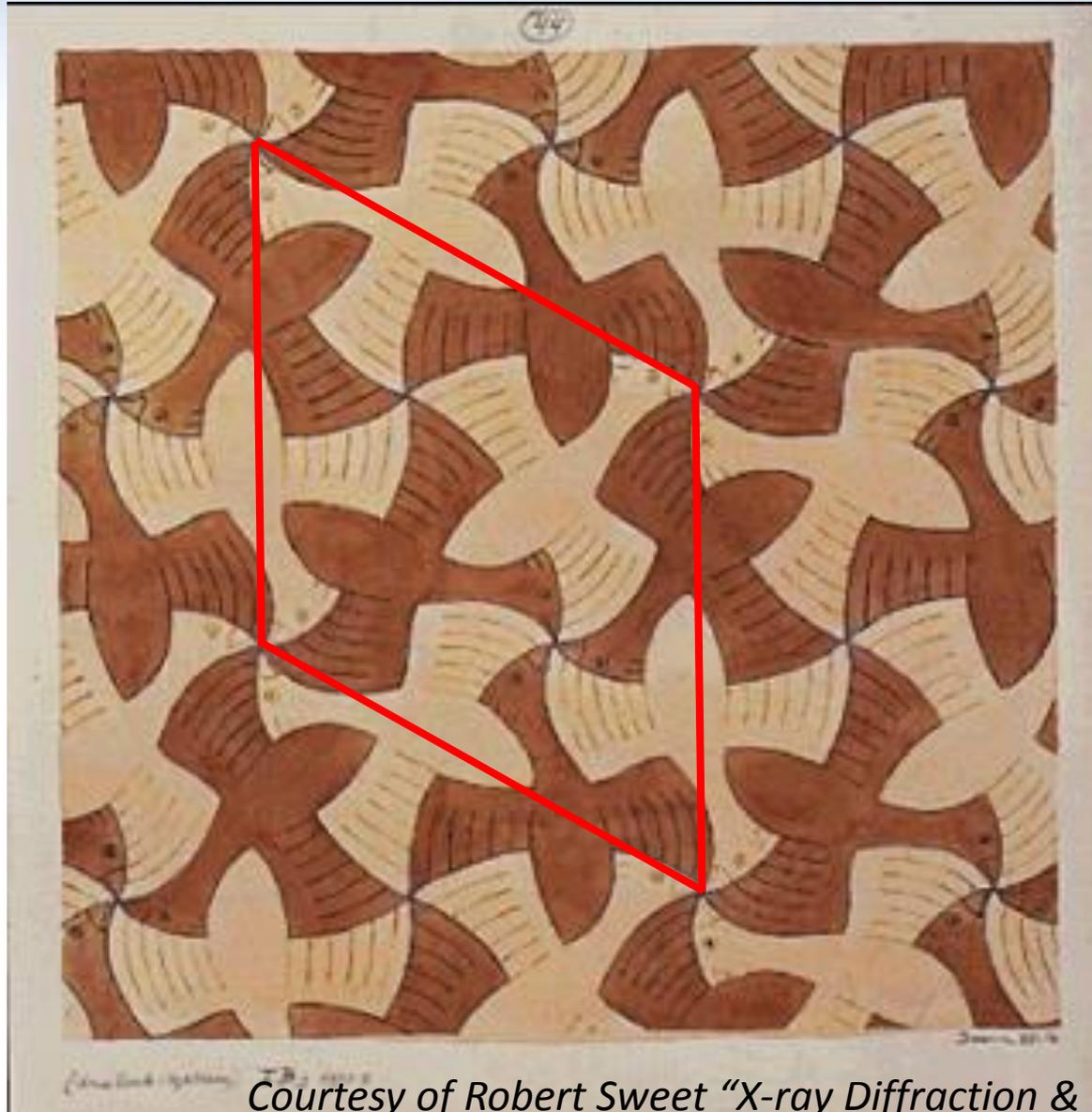
Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

What about here?



Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

Trigonal symmetry



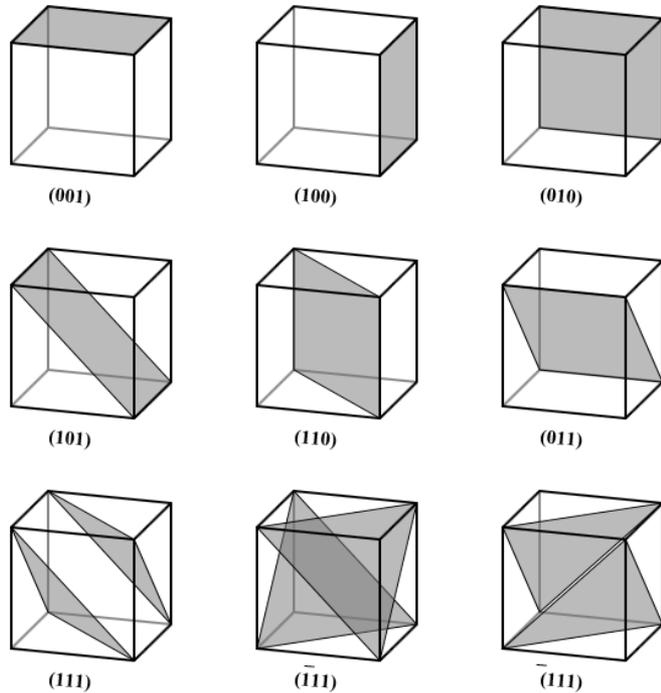
Notice the two other 3-fold axes in the cell.

Asymmetric unit is one white and one pink bird.

Order of the group is three.

Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

Crystal Structure

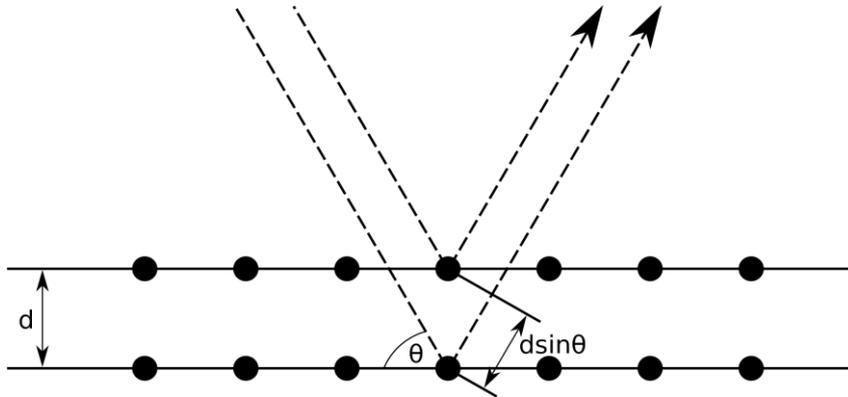


(hkl) indices	d-spacing
(111)	$a/\sqrt{3}$
(200)	$a/2$
(220)	$a/\sqrt{8}$
(311)	$a/\sqrt{11}$
<i>etc</i>	

where a is the length of the edge of the cube (lattice parameter)

The crystal structure of a material (the packing of atoms within the crystal) can be described by a discrete set of lattice planes (hkl). Lattice planes have specific inter-spacings $d(hkl)$.

Bragg Law



BRAGG LAW

$$2d(\sin\theta) = \lambda_0$$

where:

d = lattice interplanar spacing of the crystal

θ = x-ray incidence angle (Bragg angle)

λ = wavelength of the characteristic x-rays

- Bragg diffraction: constructive interference occurs when the path length difference is equal to an integer multiple of the wavelength of the incident radiation.
- **Translational symmetry and unit cell define the diffraction peak positions (in the 2θ space)**
- **Atomic distribution (atom locations) defines the diffracted peak intensities**

Remember the geometry -- if the lattice planes lie at an angle θ w/r to the incident x-ray beam, the **diffraction angle is 2θ (total beam deflection)**.

We can make an instrument to exploit that geometry.

BRAGG LAW

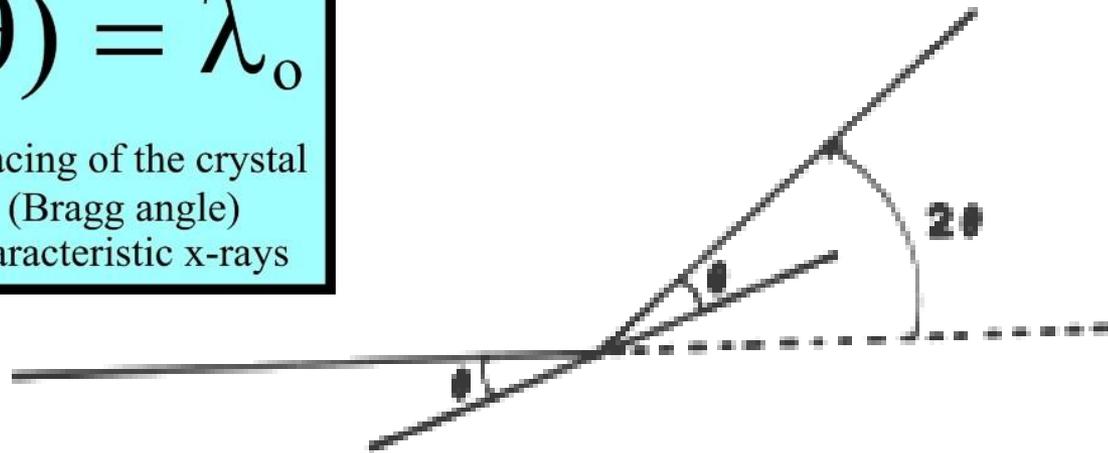
$$2d(\sin\theta) = \lambda_0$$

where:

d = lattice interplanar spacing of the crystal

θ = x-ray incidence angle (Bragg angle)

λ = wavelength of the characteristic x-rays



Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

And Remember the objective – We must view the crystal from every direction to recreate a three-dimensional image:

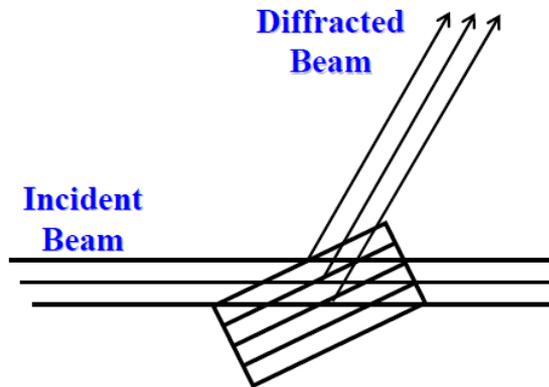
- We must obtain diffraction from **all** (or most) of the **lattice** planes (each set of h,k,l lattice planes produces a diffraction peak at the Bragg angle, extinction rules permitting)

i.e., sample **all** (or most) of the **reciprocal lattice**.

Courtesy of Robert Sweet “X-ray Diffraction & Crystallography”

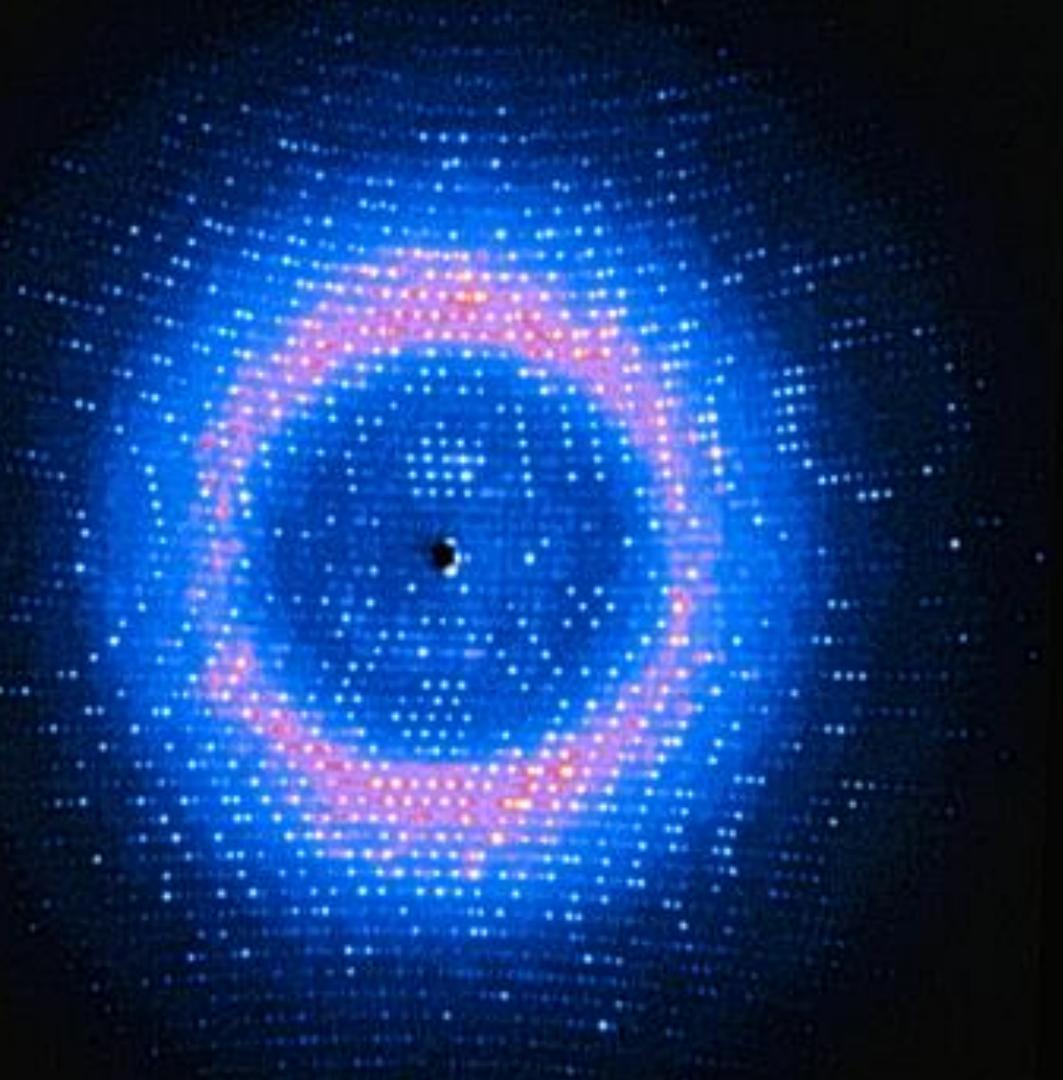
Single Crystal vs. Powder Diffraction

Single Crystal Diffraction



Precess the crystal in all orientations to get diffraction from all lattice planes

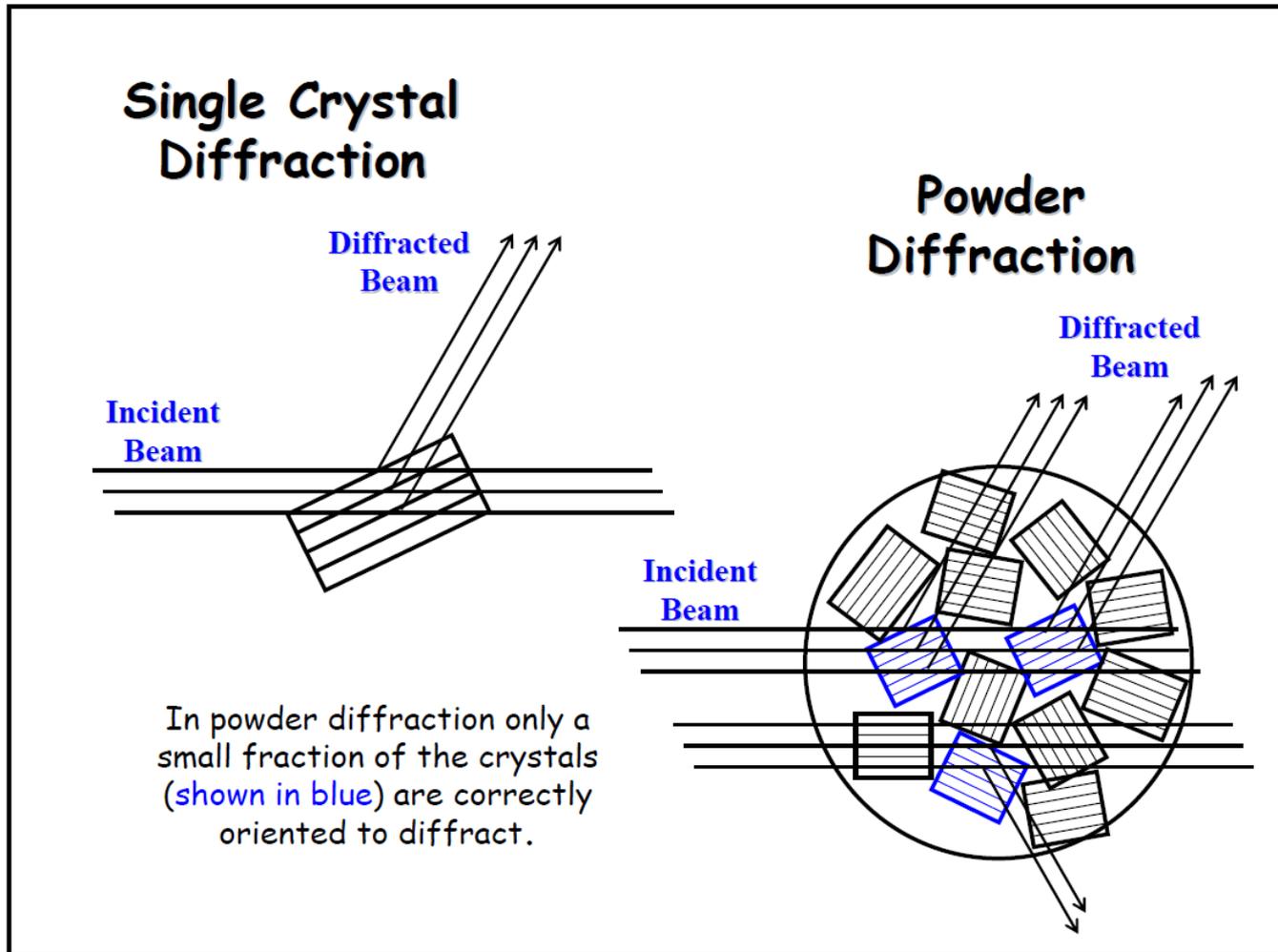
An antique rotation photograph of B-Phytoerythrin -- real x-ray film.



Courtesy of Robert Sweet "X-ray Diffraction & Crystallography"

Simple rotation geometry produces a complicated pattern that requires good software to interpret. Modern CCD-based detectors with four-circle diffractometers record such patterns and measure every spot intensity.

Single Crystal vs. Powder Diffraction

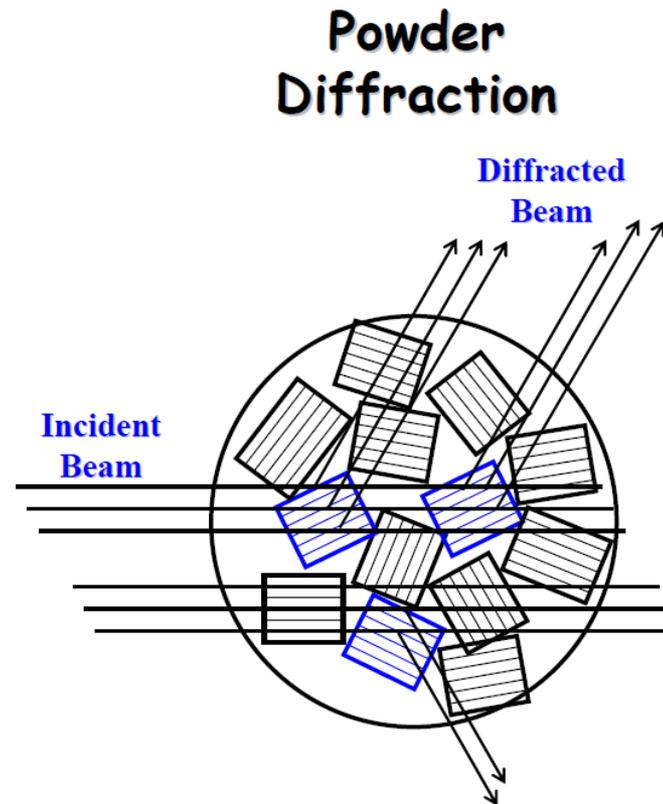


Single Crystal vs. Powder Diffraction

At any one diffraction angle, always the same family (or families) of lattice planes contribute(s).

When the sample is spun, different sub-sets of crystallites contribute but from the same family (or families) of lattice planes.

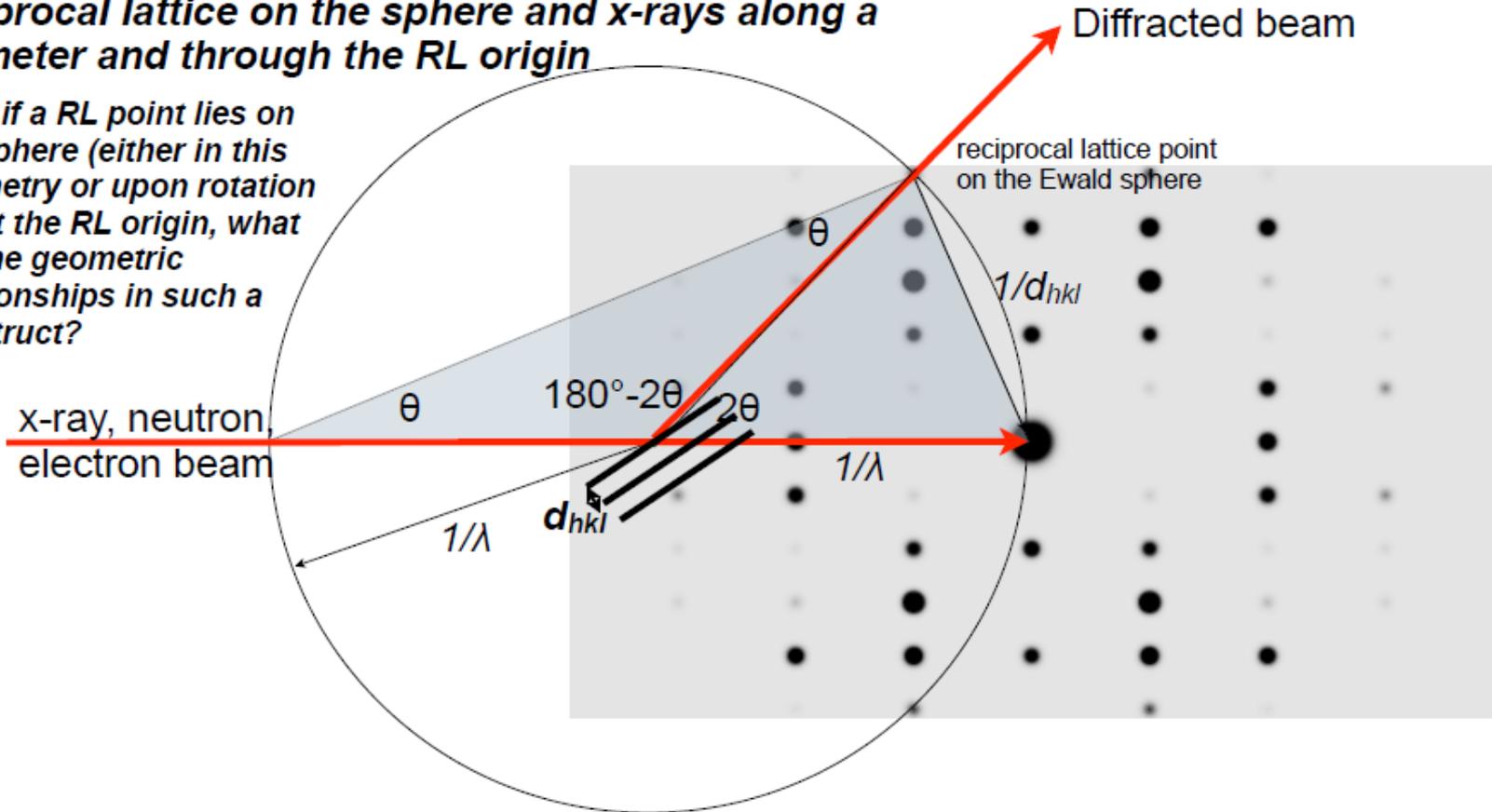
When the diffraction angle is changed, other lattice planes from other crystallites are probed.



The Ewald sphere construction

Construct a sphere of radius $1/\lambda$, with the origin of the reciprocal lattice on the sphere and x-rays along a diameter and through the RL origin

Now, if a RL point lies on the sphere (either in this geometry or upon rotation about the RL origin, what are the geometric relationships in such a construct?

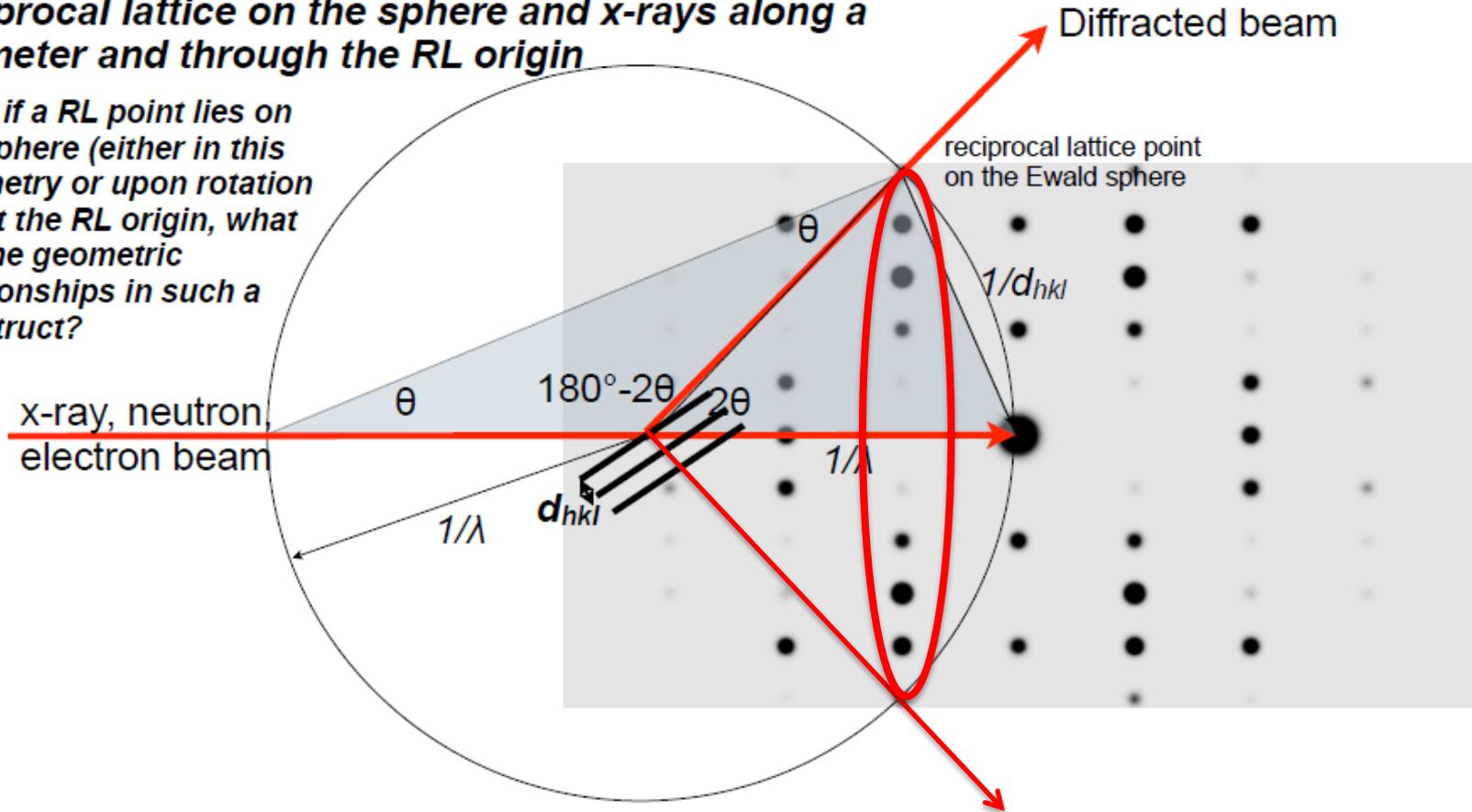


Courtesy of John Parise "Principles of diffraction"

The Ewald sphere construction

Construct a sphere of radius $1/\lambda$, with the origin of the reciprocal lattice on the sphere and x-rays along a diameter and through the RL origin

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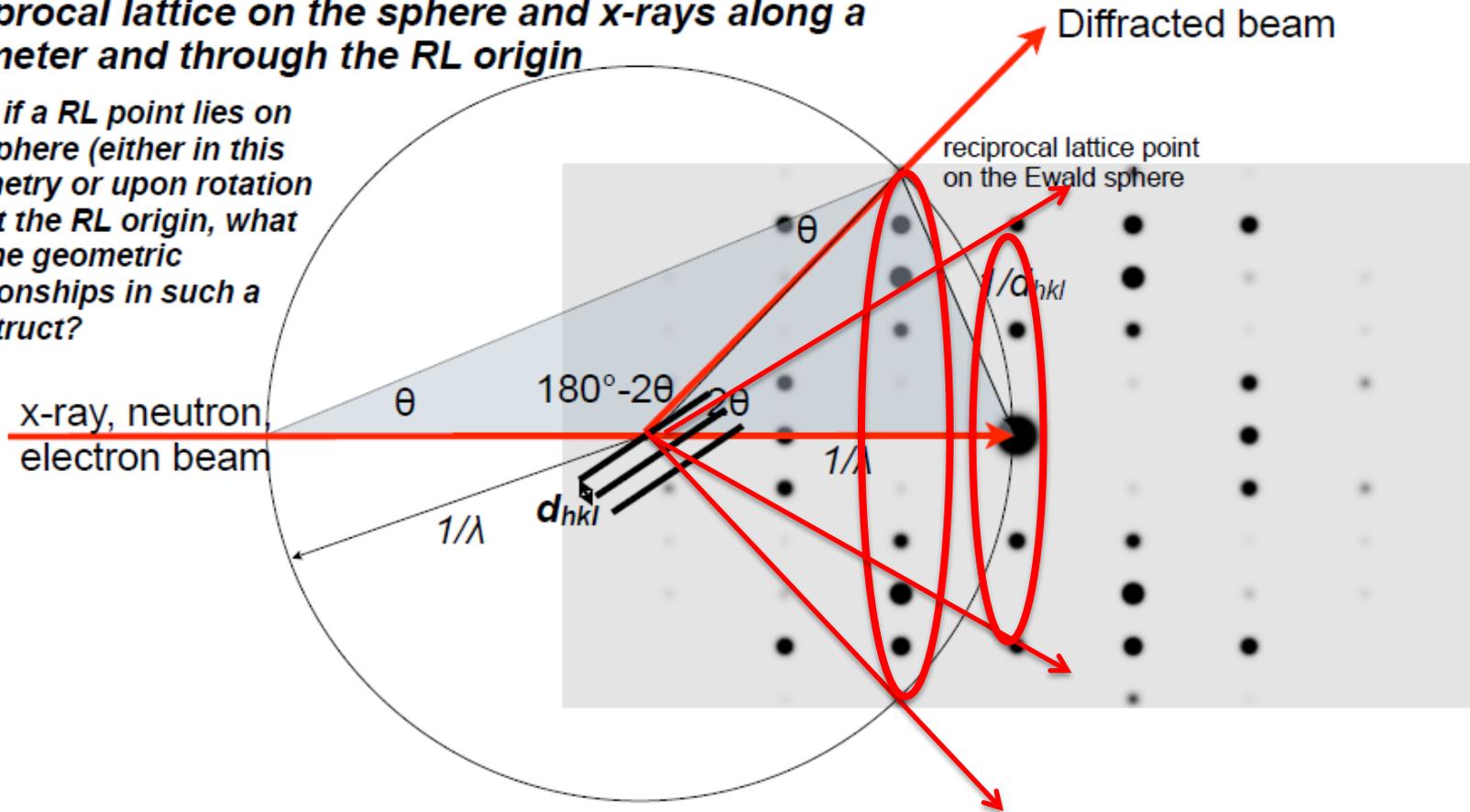


Courtesy of John Parise "Principles of diffraction"

The Ewald sphere construction

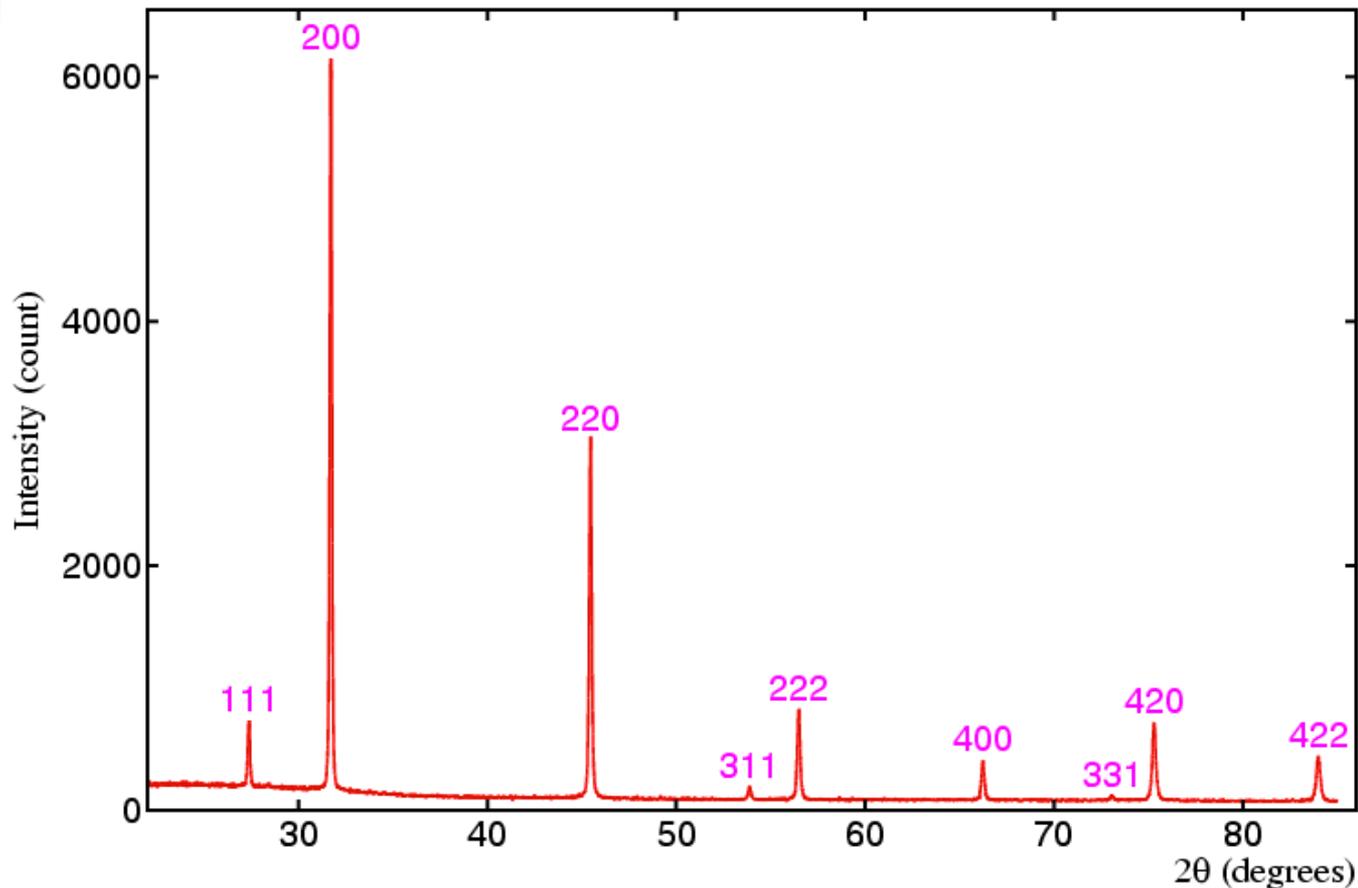
Construct a sphere of radius $1/\lambda$, with the origin of the reciprocal lattice on the sphere and x-rays along a diameter and through the RL origin

Now, if a RL point lies on the sphere (either in this geometry or upon rotation about the RL origin, what are the geometric relationships in such a construct?



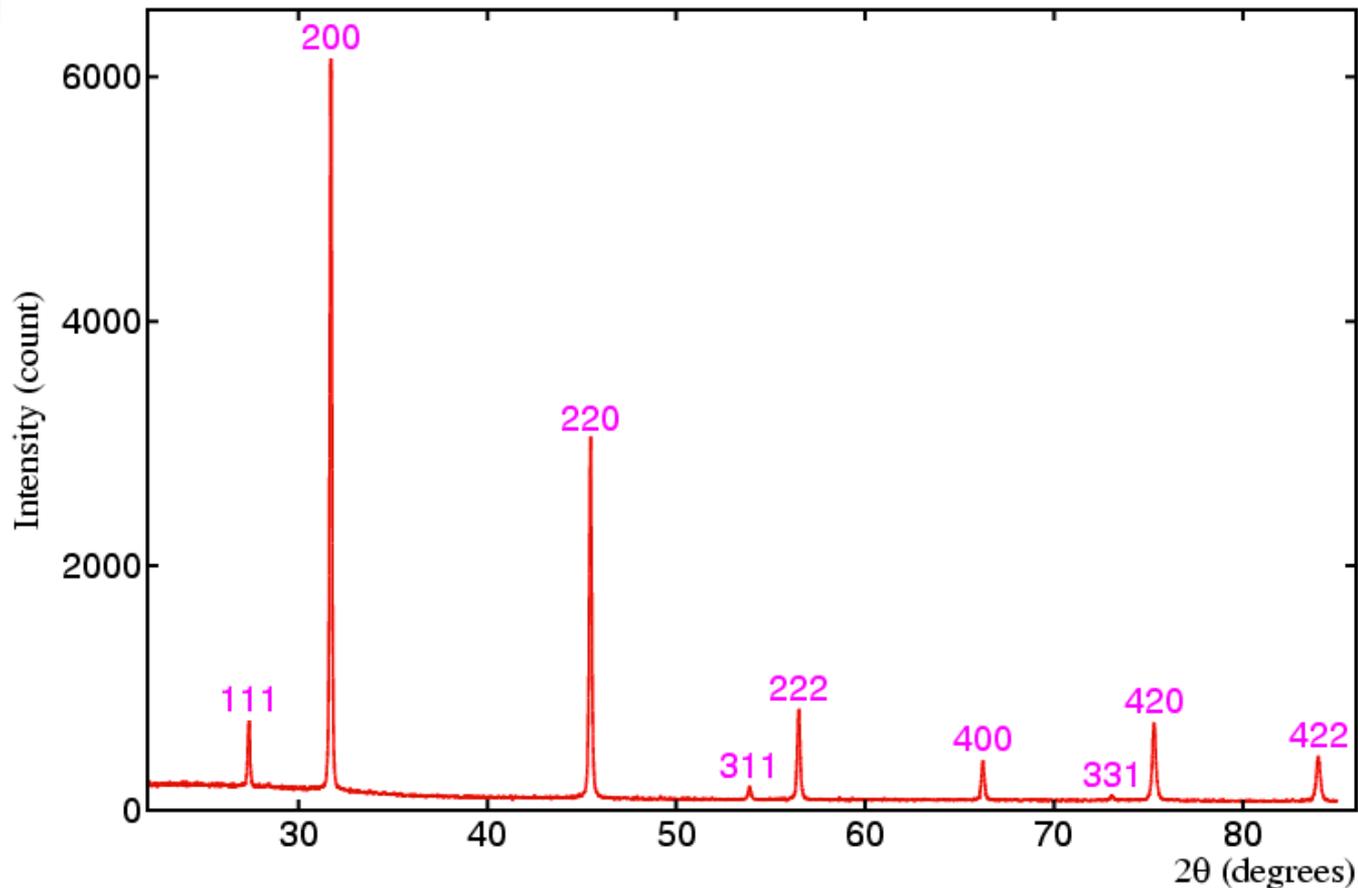
Courtesy of John Parise "Principles of diffraction"

Powder Diffraction Pattern



A series of intensity values (detector photon counts) vs. diffraction angle. Data are compressed into 1D (2θ scale): good and bad!!

Powder Diffraction Pattern



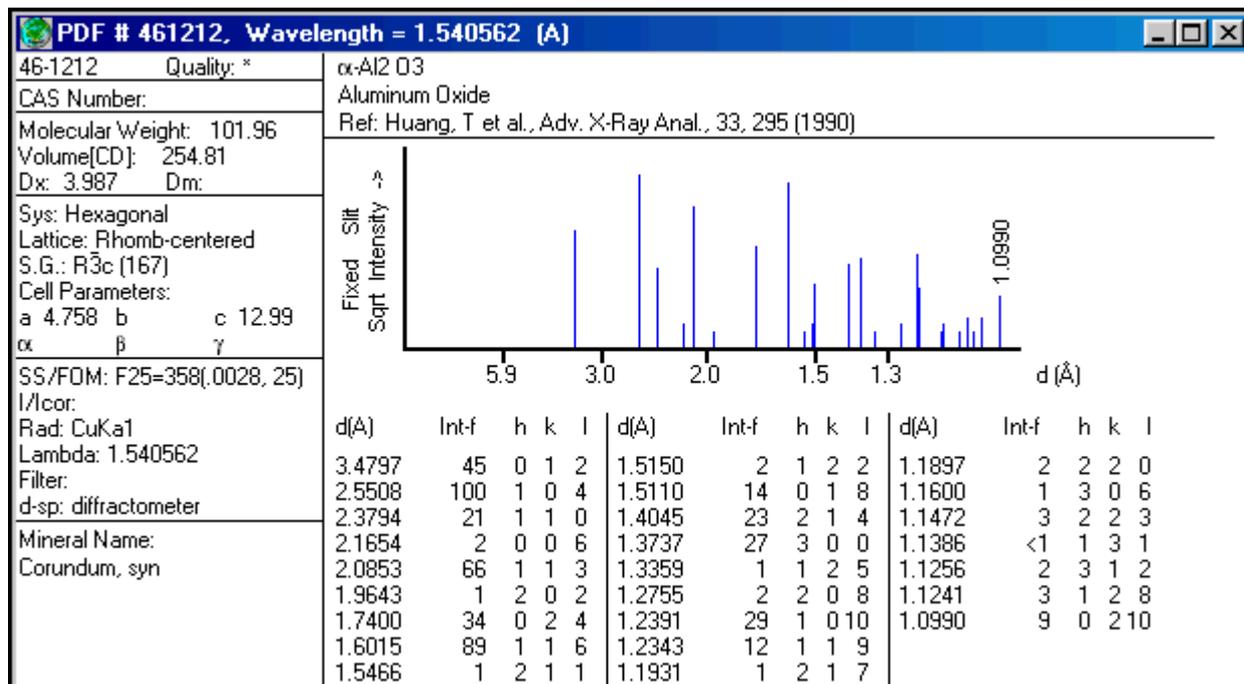
Diffraction pattern (not a spectrum): elastic process
no photon energy change

Qualitative Analysis

Any powder pattern can be used like a fingerprint to see if it matches the powder pattern of an already known compound.

The **International Centre for Diffraction Data (ICDD)** (PA) maintains a database of known powder diffraction patterns (www.icdd.com)

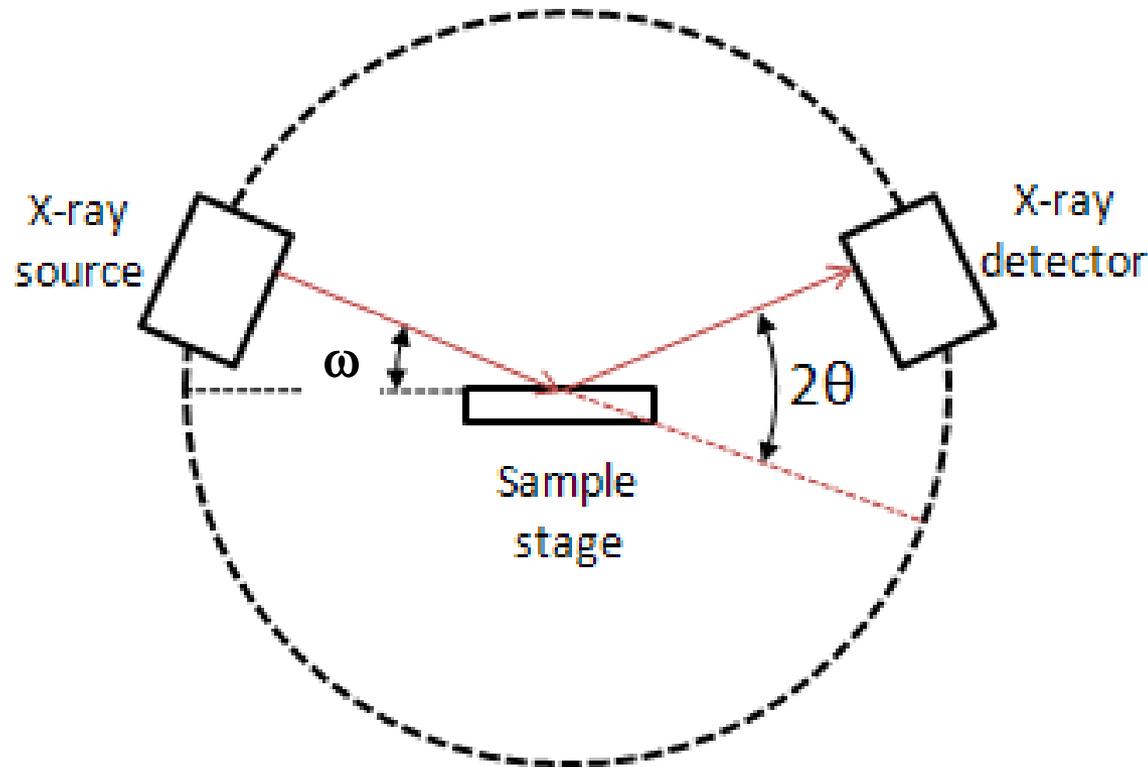
- 365,900 standardized entries
- 42,800 Minerals
- 502,000 Organics



Outline

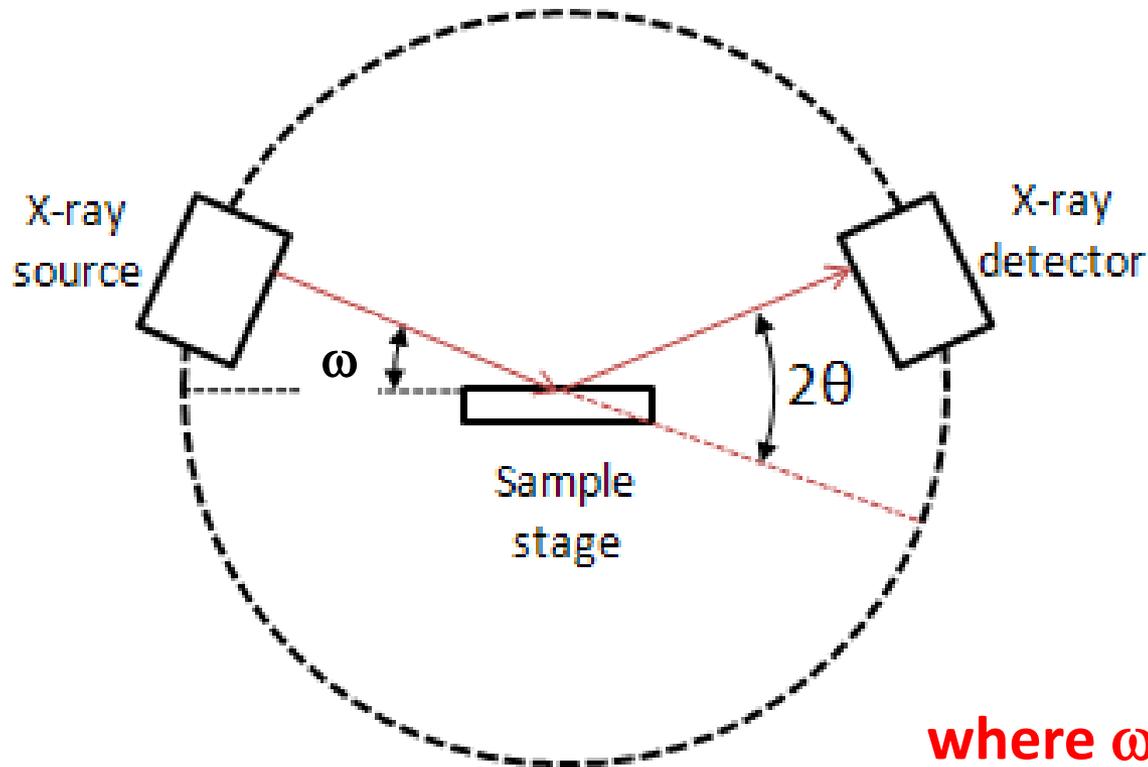
1. Applications for Powder Diffraction
2. The basics of Powder Diffraction
- 3. Experimental Methods for collecting PD data**
- 4. Crystal structure Determination**
- 5. Peak Profile Analysis for Microstructural Studies**
- 6. Advanced PD techniques**

Scheme of a One-Axis Diffractometer



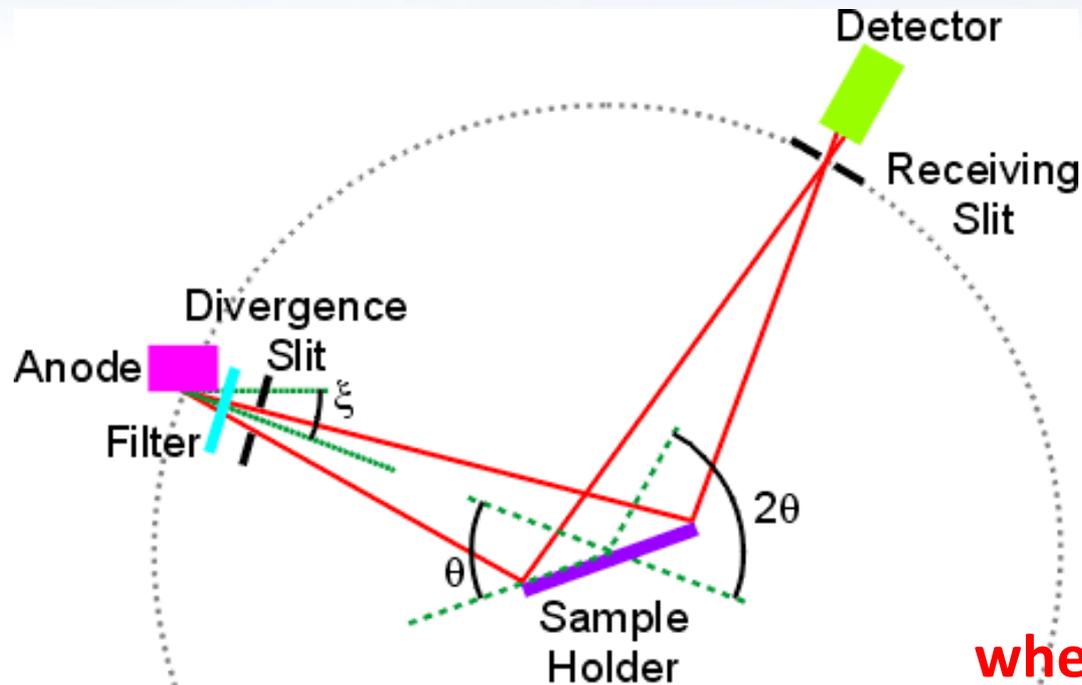
- X-ray source (tube, rotating anode)
- Powder sample (flat plate, capillary)
- X-ray detector (a variety of technologies: direct counting or conversion)
- Diffraction plane (contains the incident and the diffracted beam)

Scheme of a One-Axis Diffractometer



- X-ray source (tube, rotating anode)
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Scheme of a Two-Axis Diffractometer



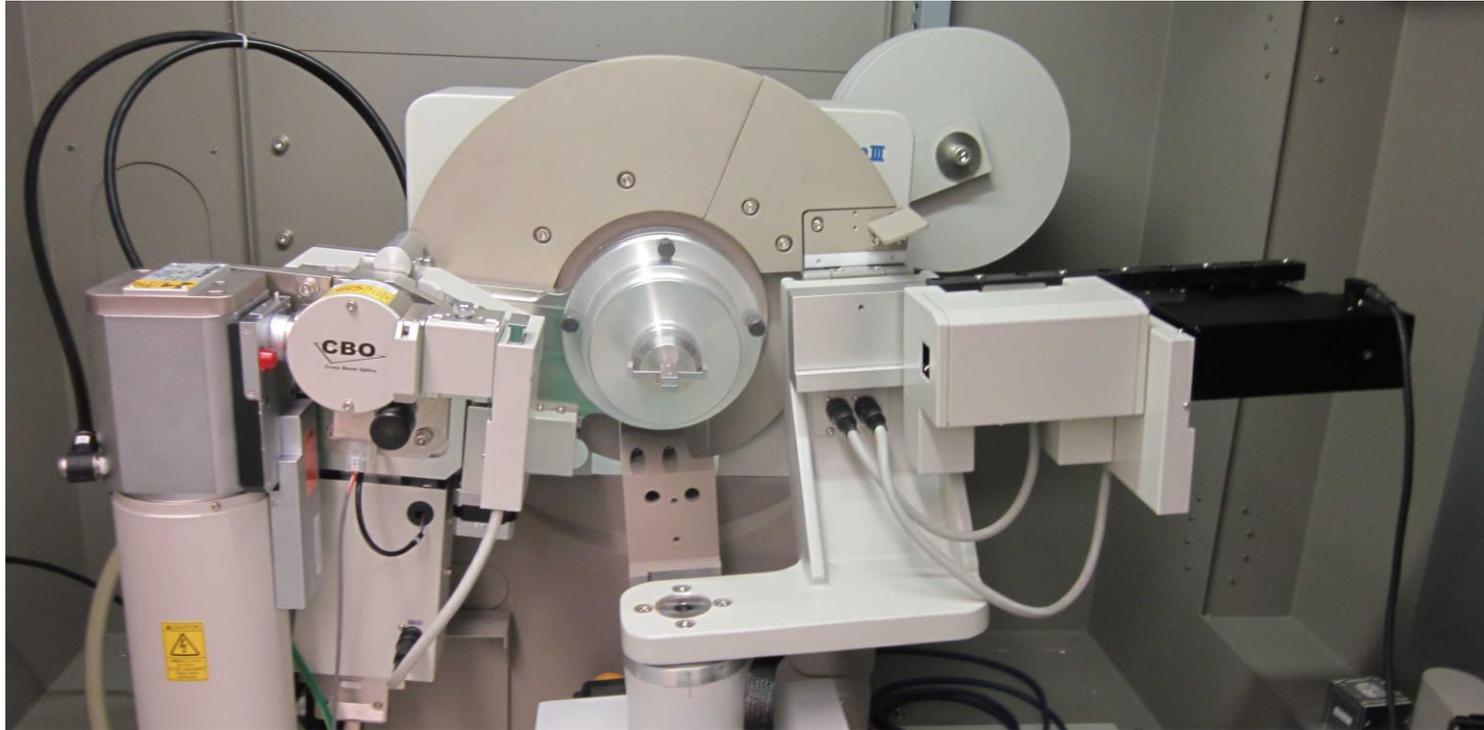
where $\omega = 2\theta/2$

Optics:

- Para-focusing optics (Bragg-Brentano: sample rotates by θ , detector rotates by $2\theta \rightarrow$ to keep detector on focusing (Rowland) circle
- Slits before the sample (divergence) and after (angle resolution $\Delta 2\theta$)

Scheme of a Three-Axis Diffractometer

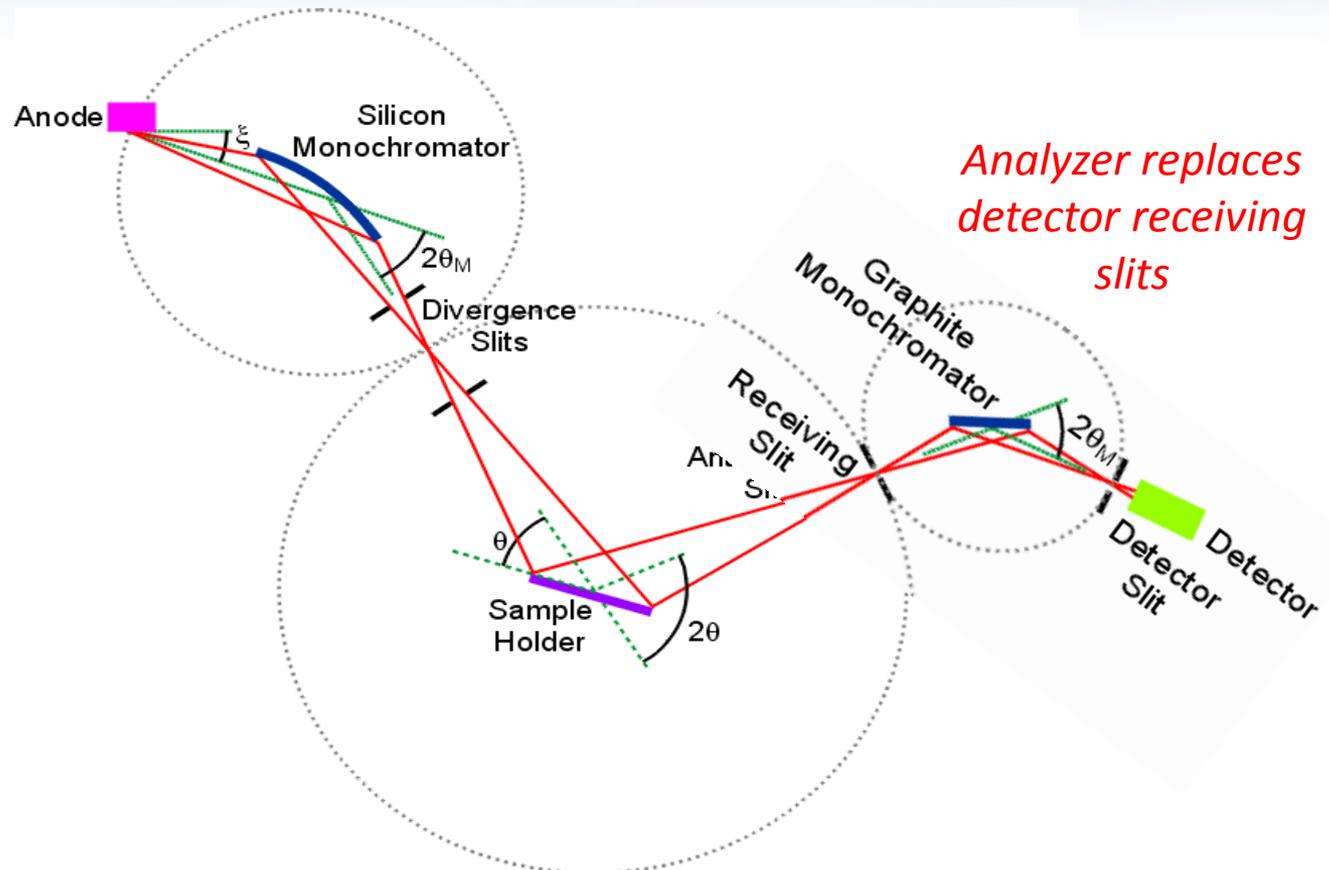
(do you see the additional 3rd circle?)



Spinning sample (in many cases):

- Average: as many crystallites (grains) in the beam as possible
- Randomize: eliminates preferred orientation
- Not an “ideal” powder (texture, poor sampling) → intensities are affected

Scheme of a Four-Axis Diffractometer



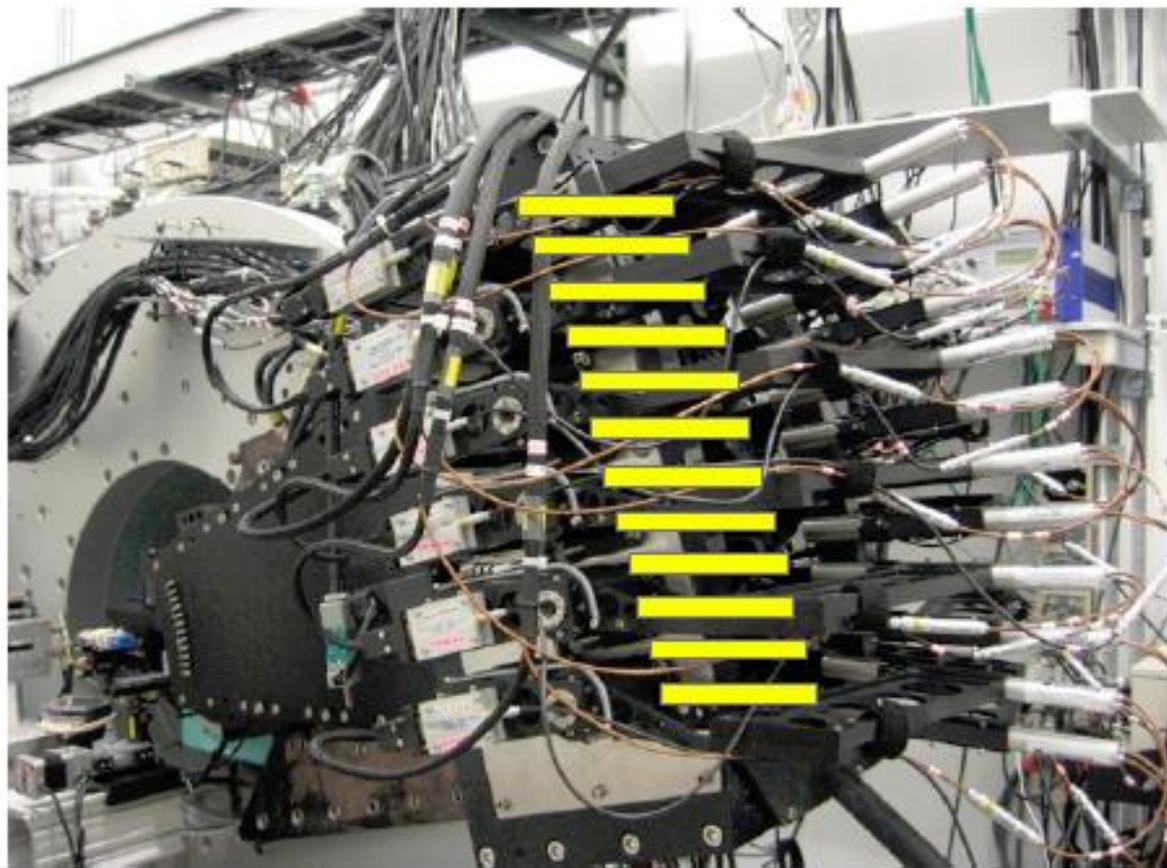
Can you see the 4 circles ?

Multi-Analyzer Crystal detector at Argone (IL)

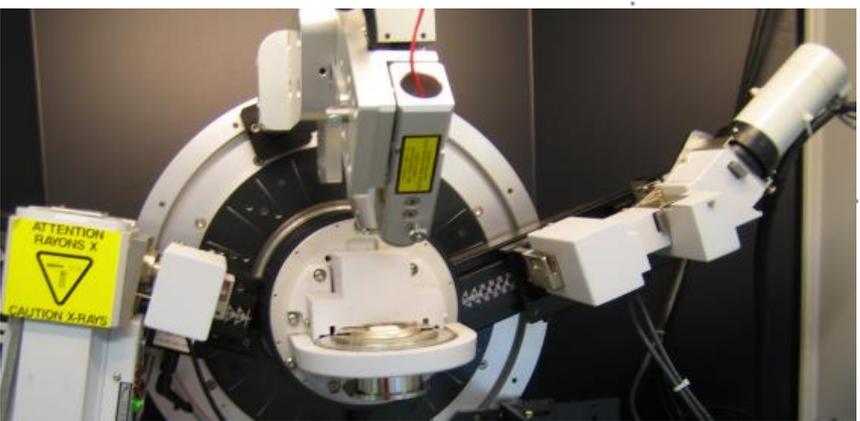
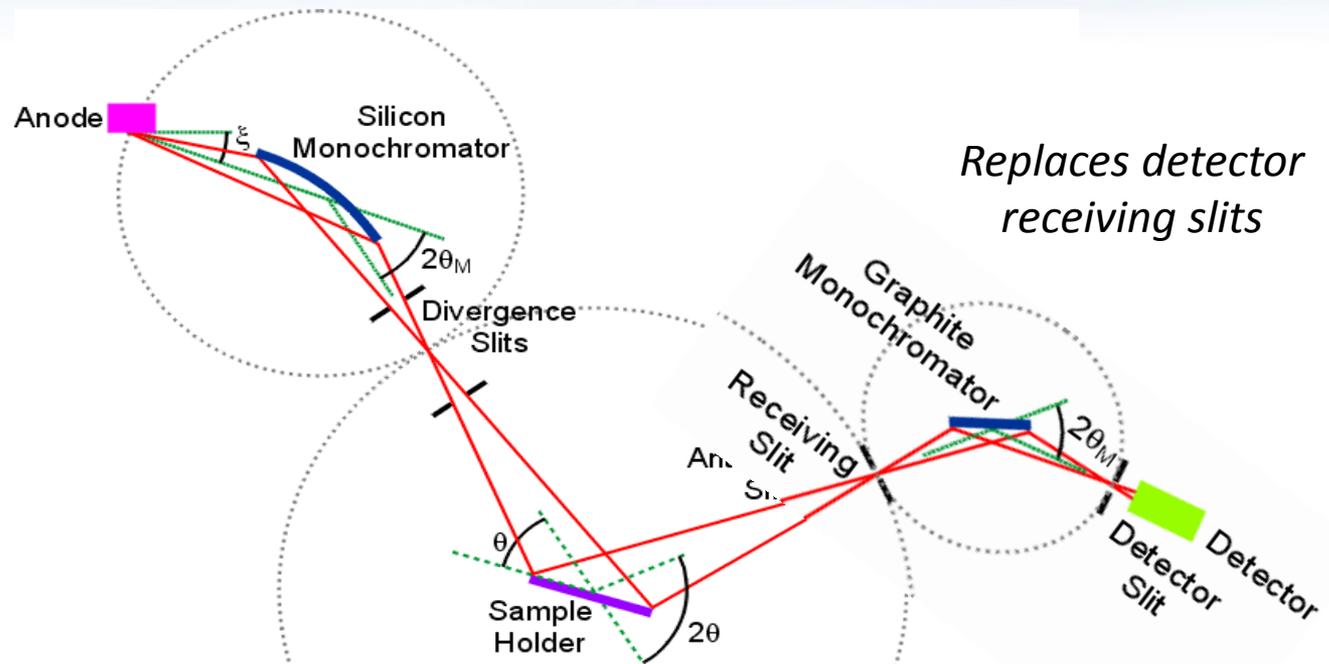
- 12 crystals/detectors
 - Individually adjusted
 - 2° separation
 - Total 2θ range = 22°
- Si(111) crystals
 - excellent discrimination
 - low background (& yield!)
 - LaCl_3 Scintillators

Multiple Observations

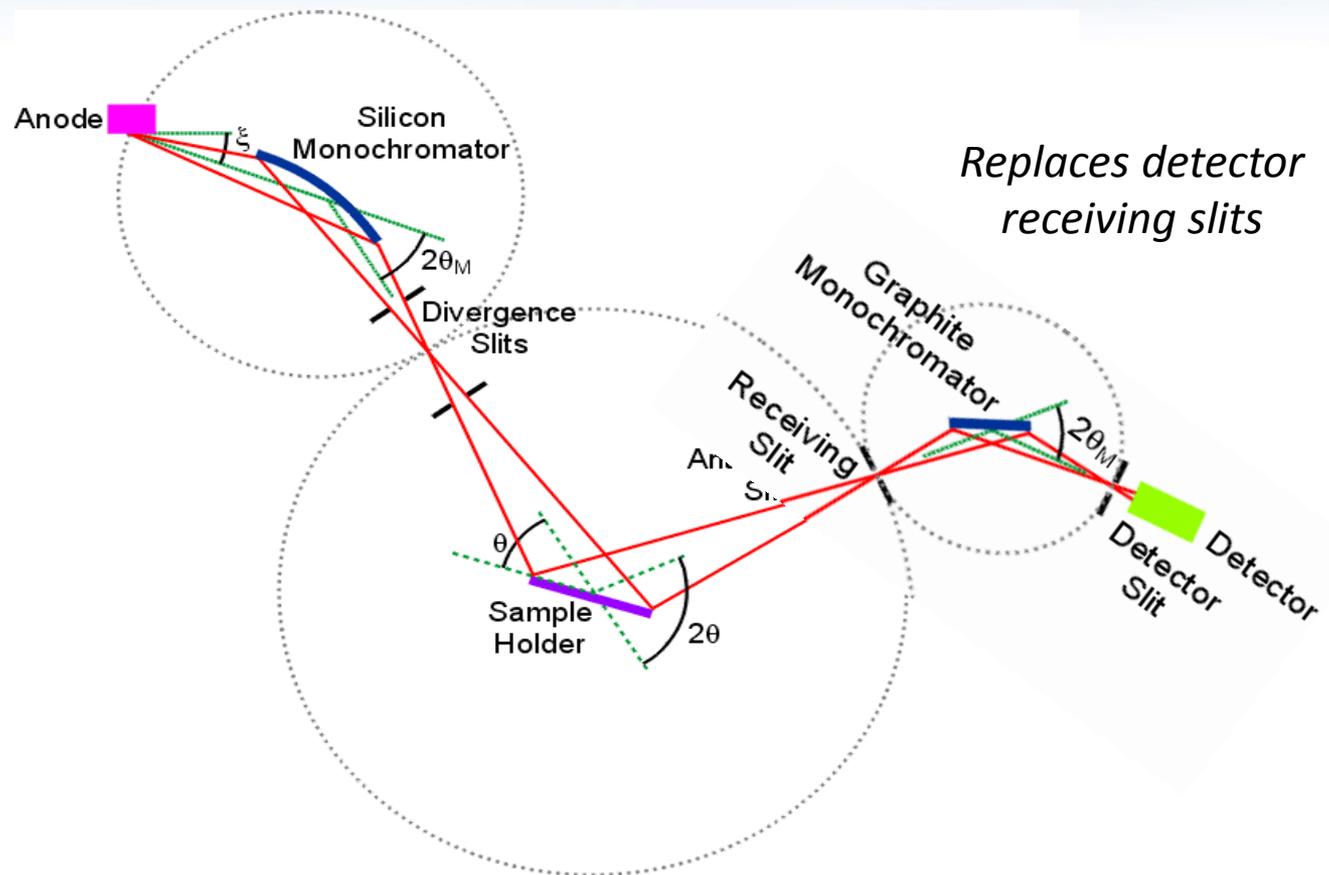
- Reduce Collection Time
- Improve Accuracy



Scheme of a Four-Axis Diffractometer



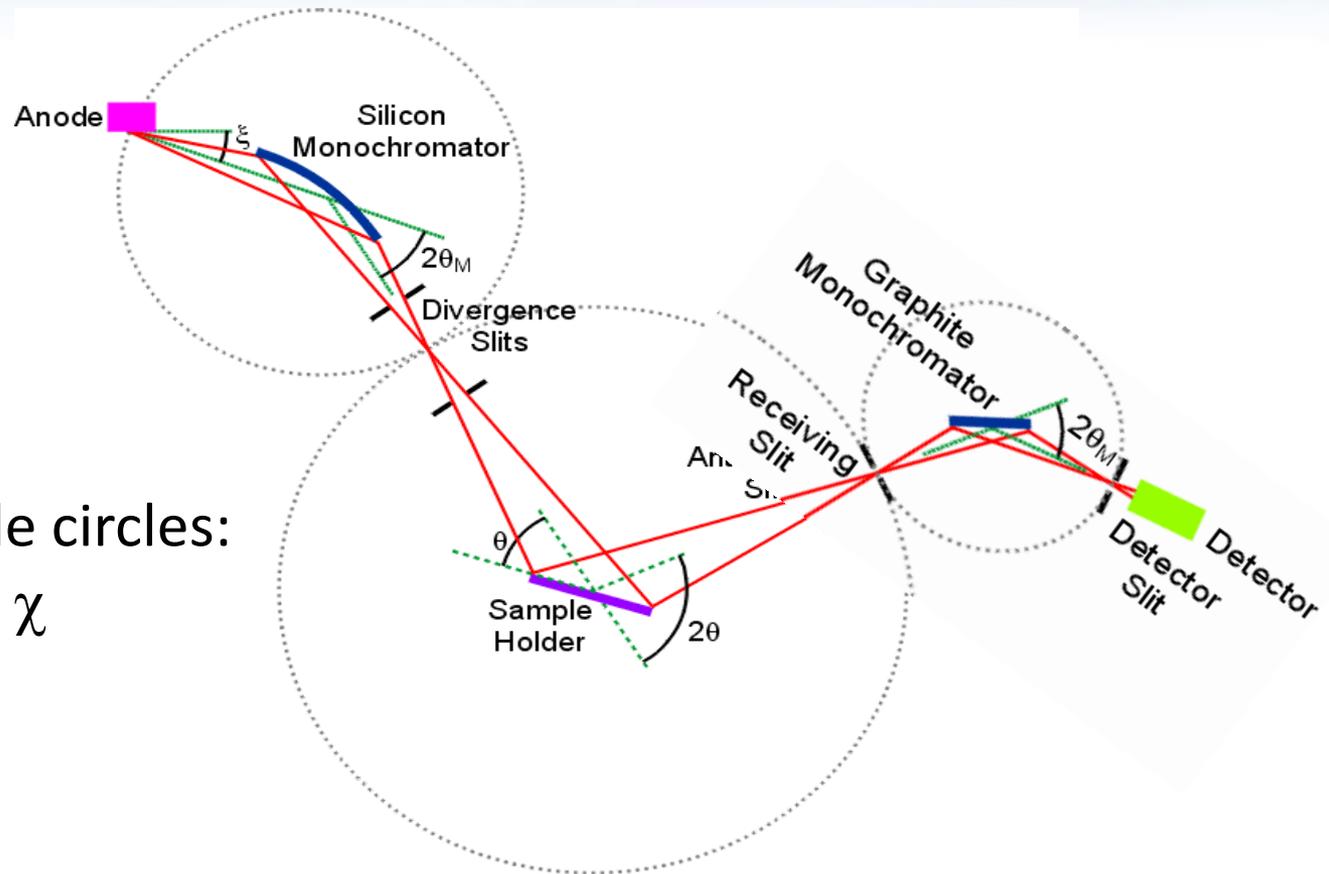
Scheme of a Four-Axis Diffractometer



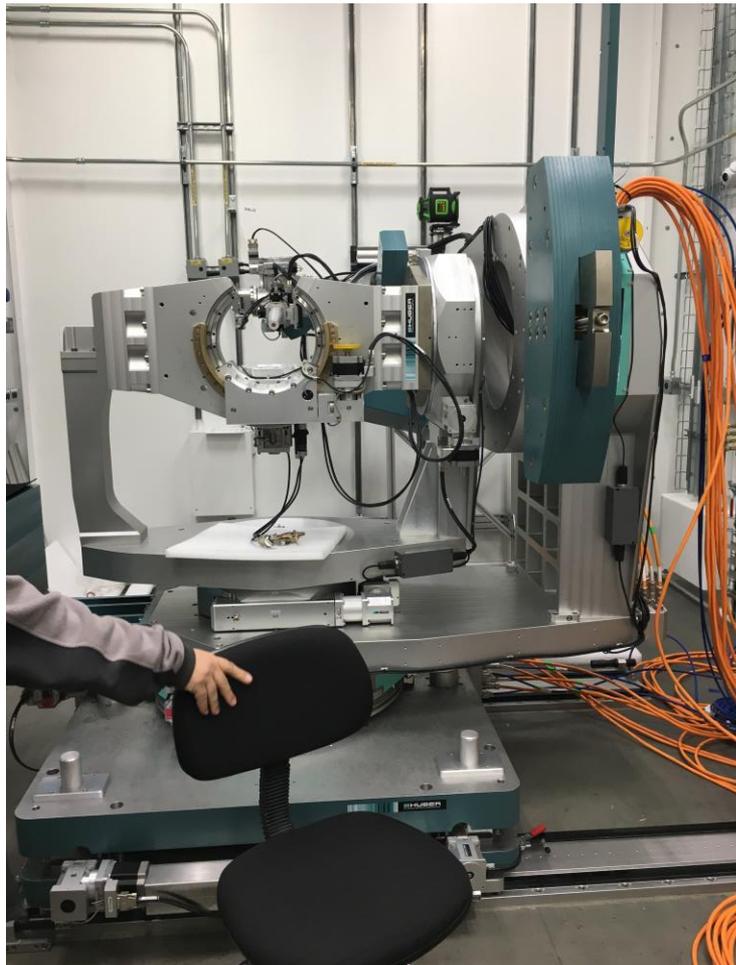
Can you find another 2 circles (not shown) ?

Scheme of a Six-Axis Diffractometer

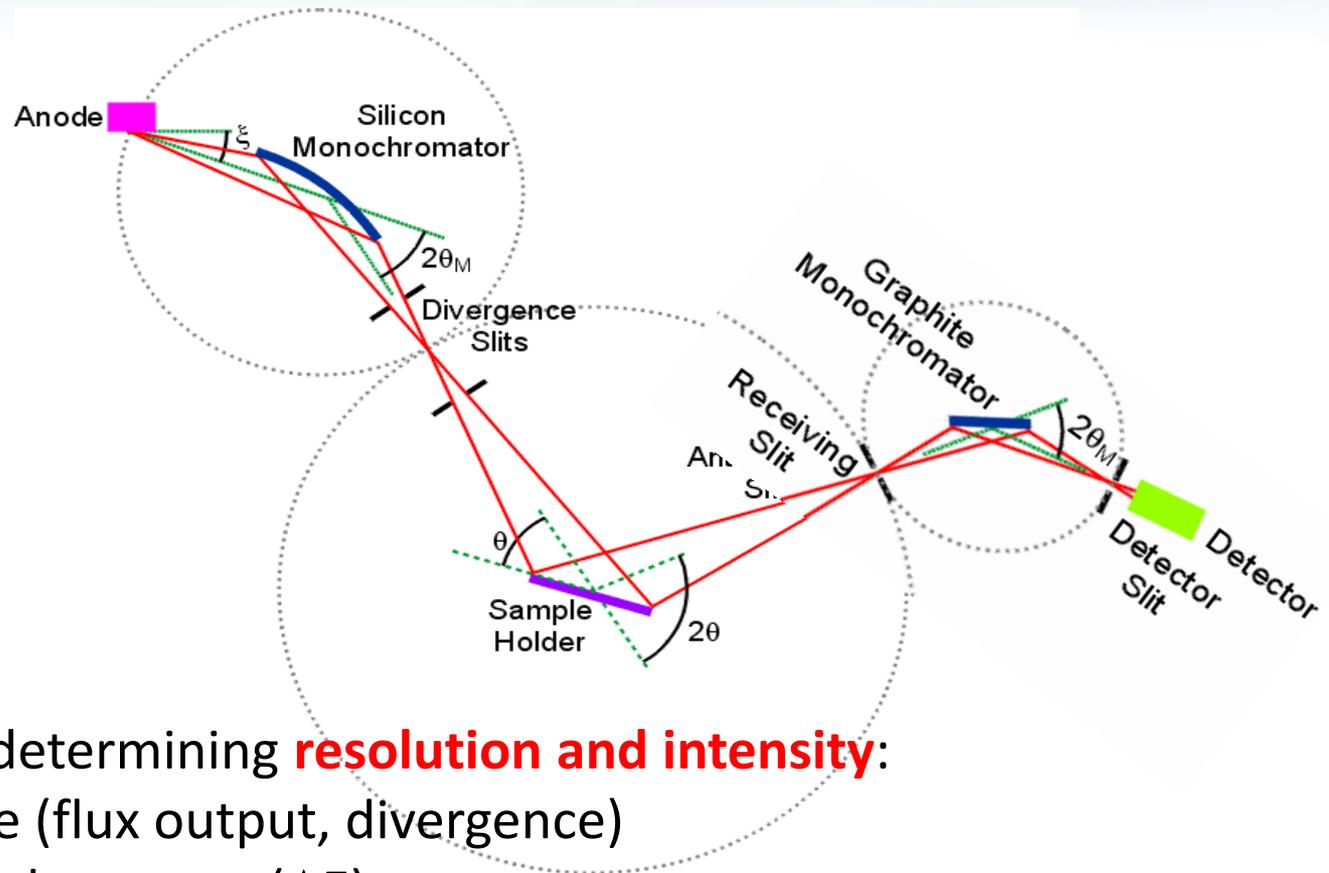
+ 2 sample circles:
 ϕ and χ



6-circle diffractometer at the NSLS-II (ISR)



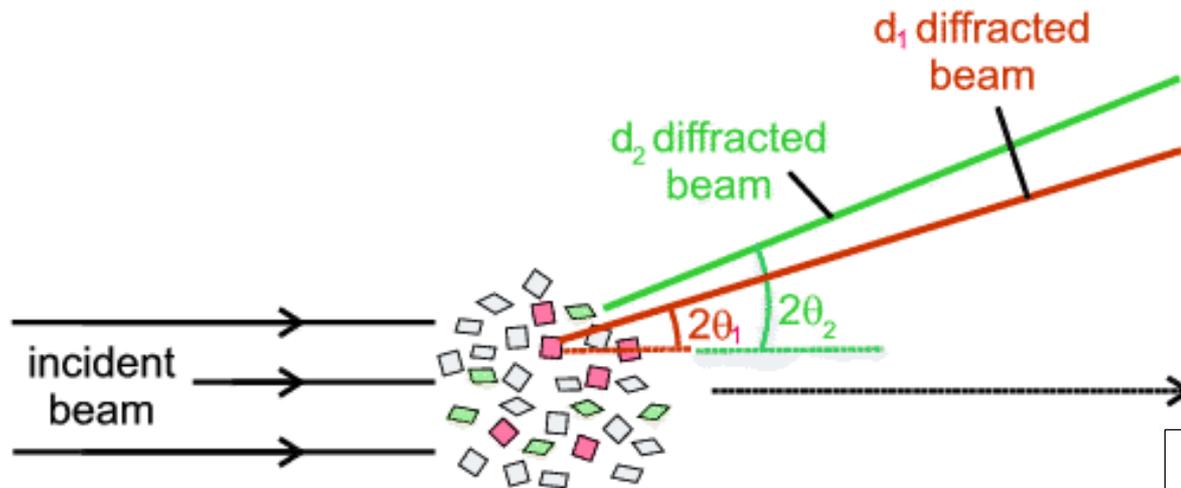
Scheme of a Six-Axis Diffractometer



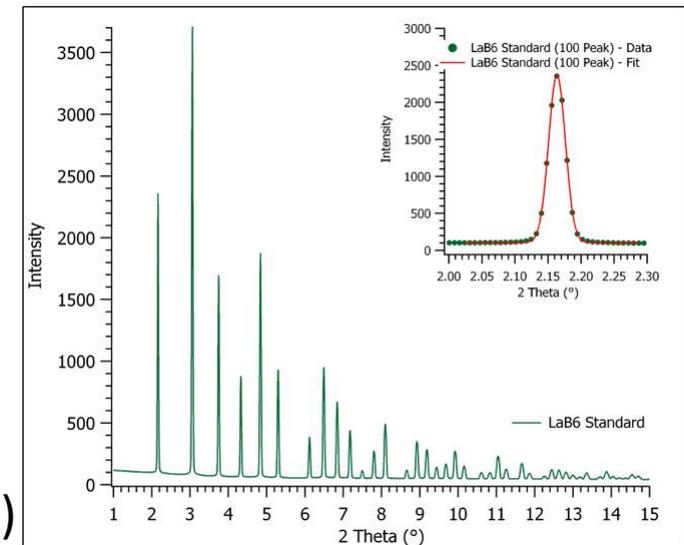
Parameters determining **resolution and intensity**:

- Source (flux output, divergence)
- Monochromator (ΔE)
- Mirror (focusing, parallel)
- Analyzer (discrimination, error immune)

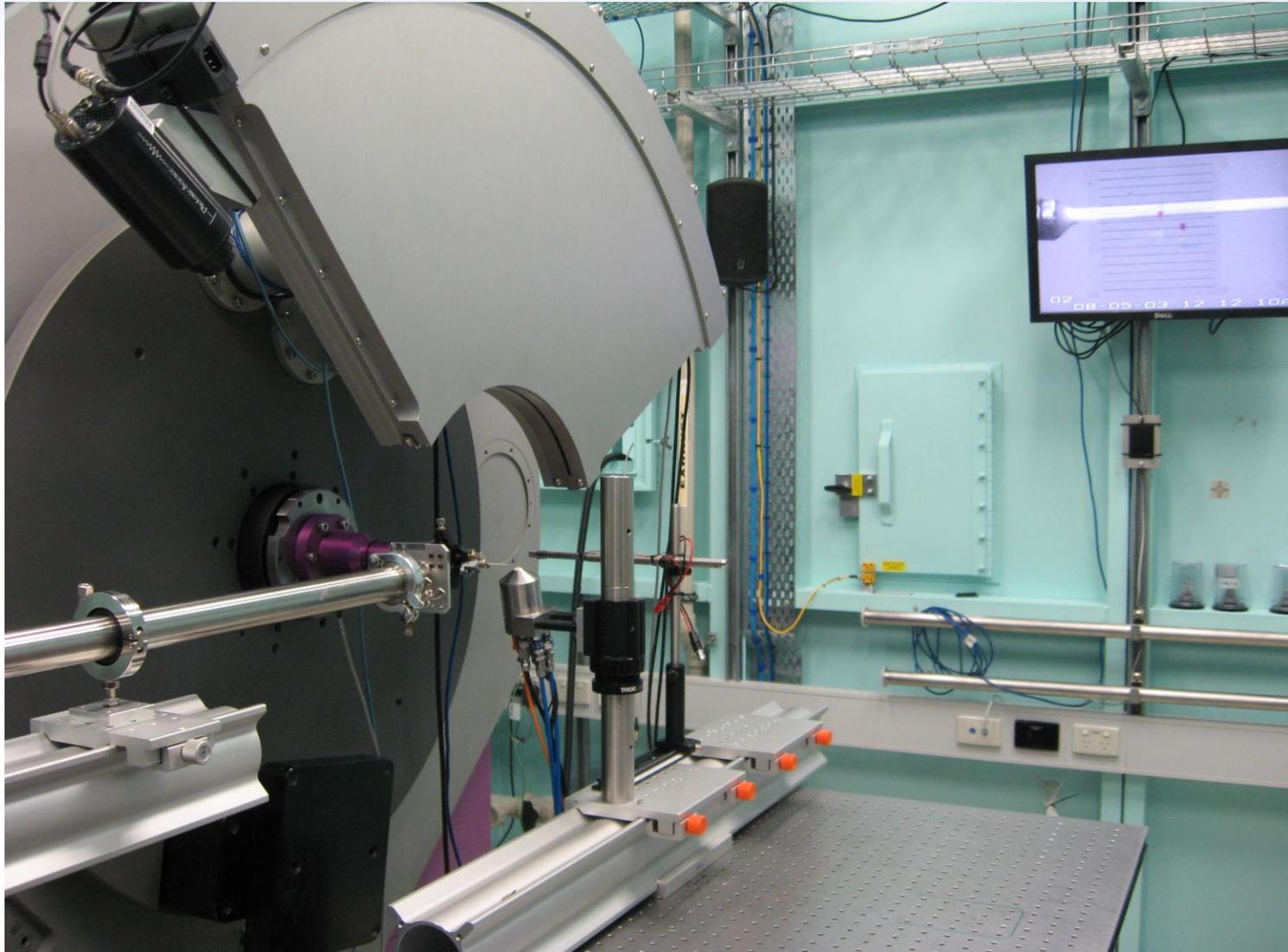
From 0-D Powder Diffraction...



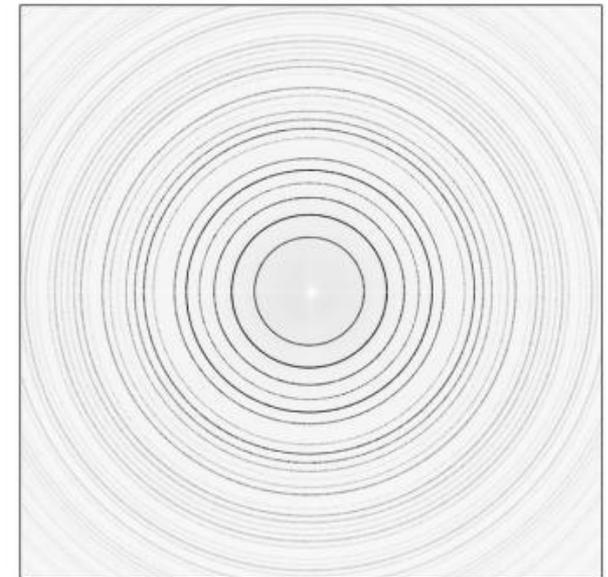
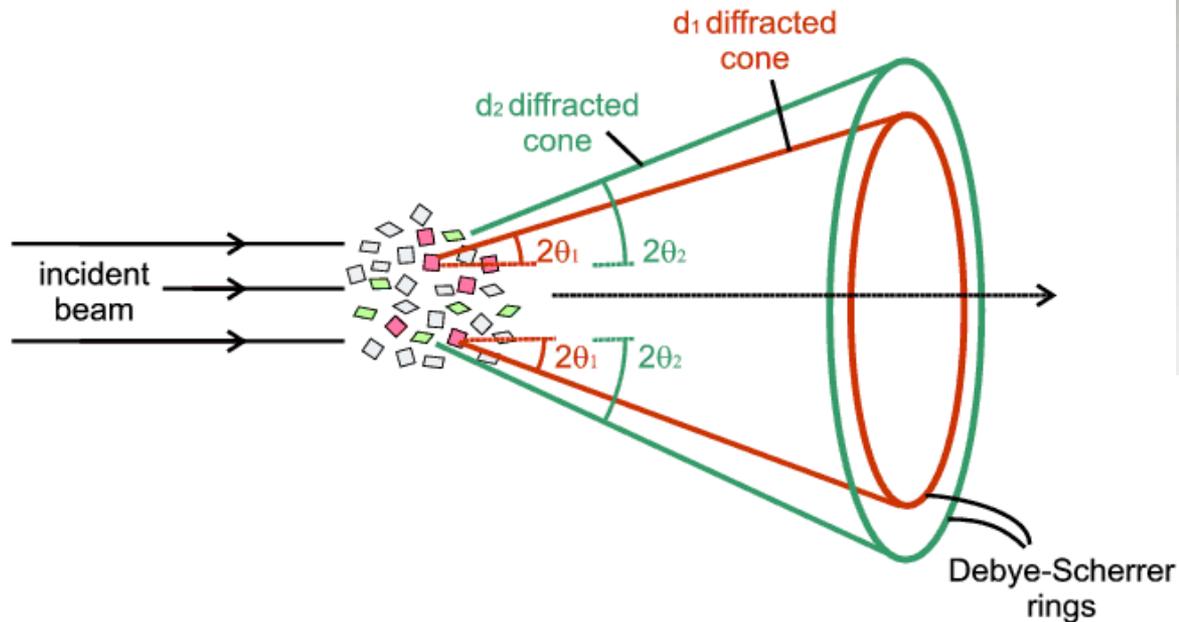
- 0-D (point) detector: scanning (slow and accurate)
- 1-D (line) detector: intercepts all 2θ angles (fast but lower res)



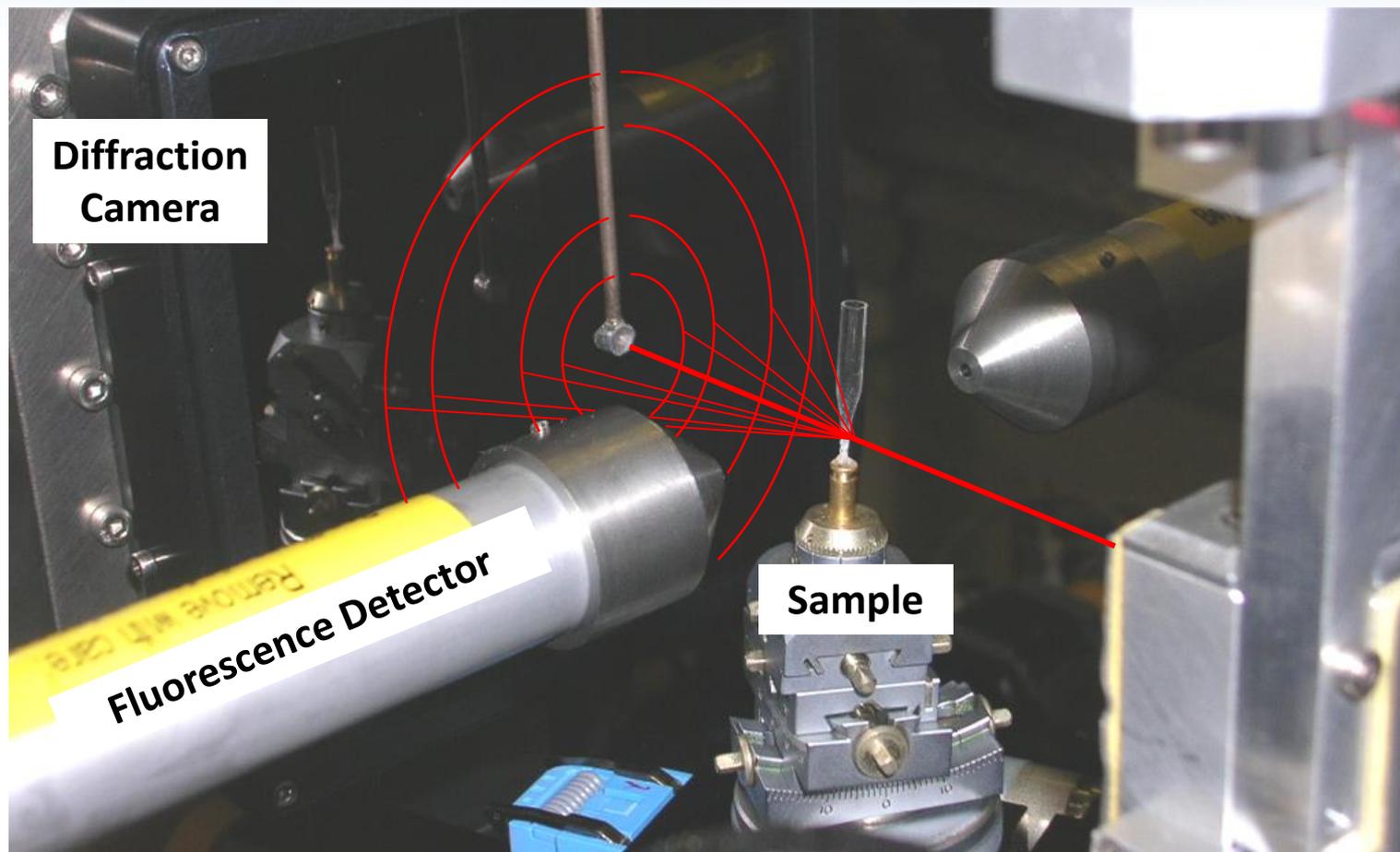
The 1-D detector at the Australian Synchrotron



...to 2-D Powder Diffraction



- 2-D (large-area) detector: **Debye-Scherrer rings** (large field of view in one single shot, background high, low resolution)
- **Non-random orientation generates non-uniform or spotty rings** (texture, pole figures)



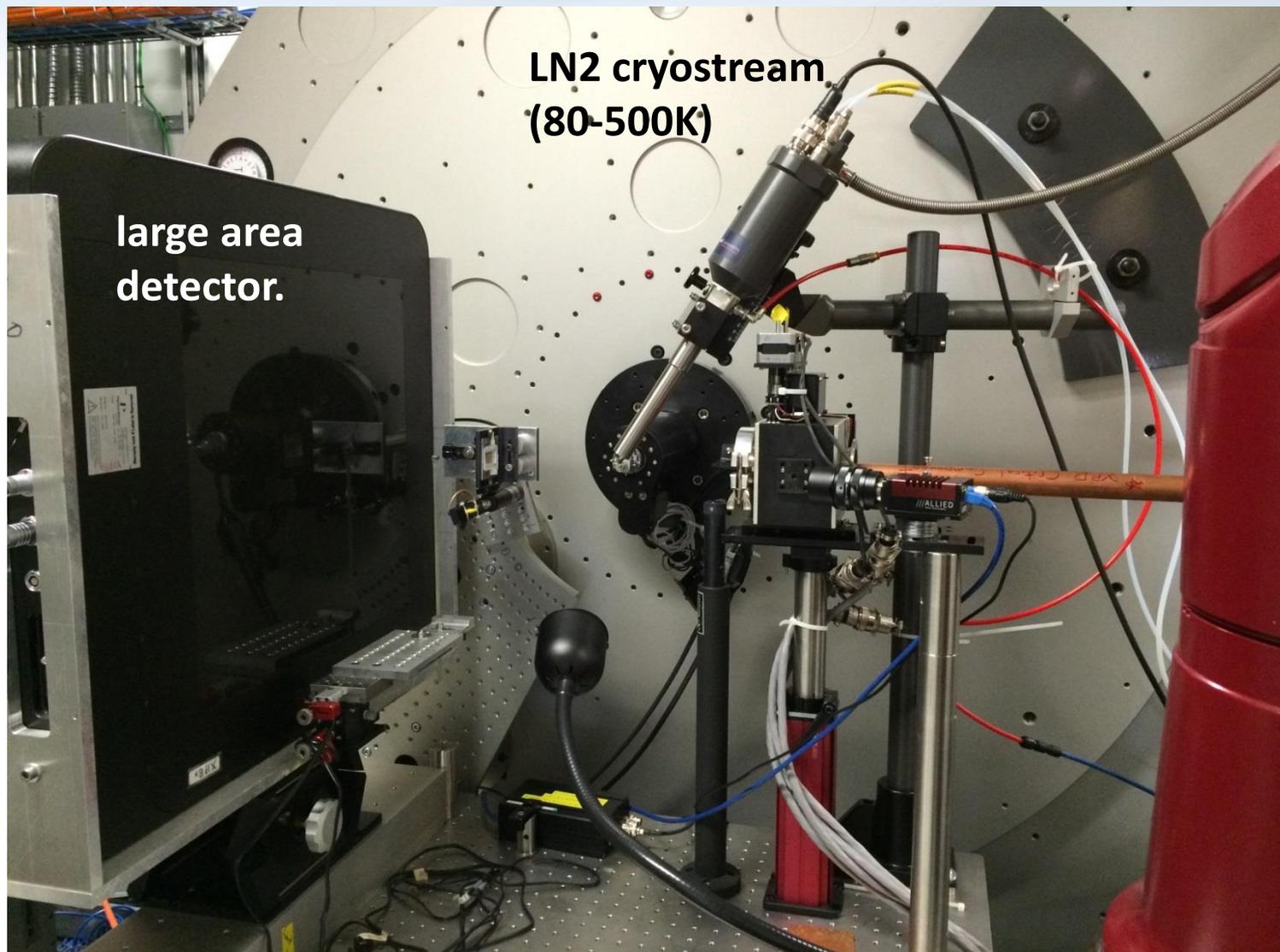
**Diffraction
Camera**

Fluorescence Detector

Sample

1cm

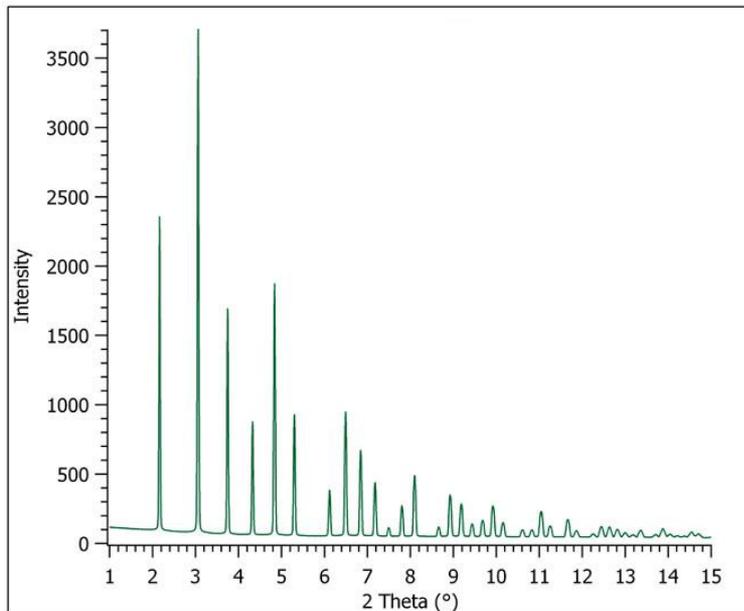
NSLS-II Experimental Set-Up



Comparison of 0D and 2D detection

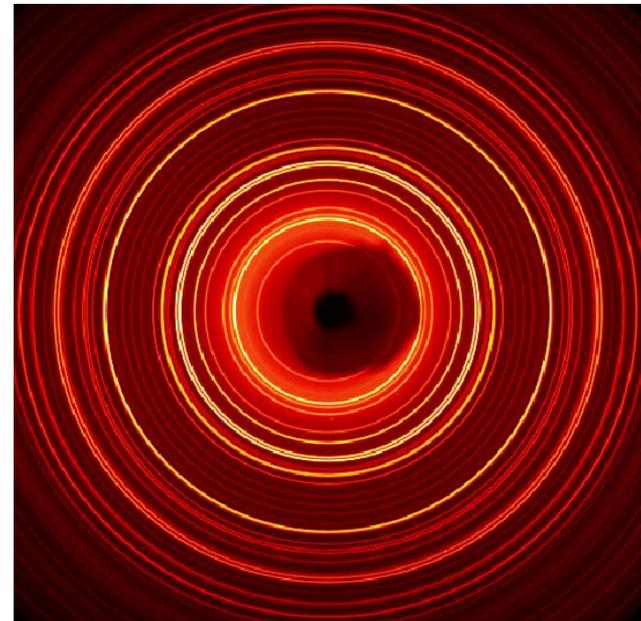
Point Detector

- Angle scanning (slow)
- Higher Resolution (peak shape, peak separation)
- Less Background



Area Detector

- Fixed
- Faster (single shot)
- Higher Statistics (full azimuth)



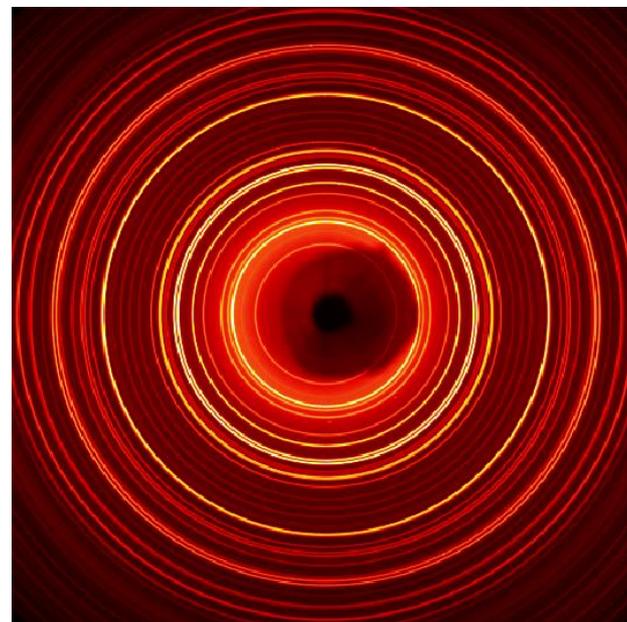
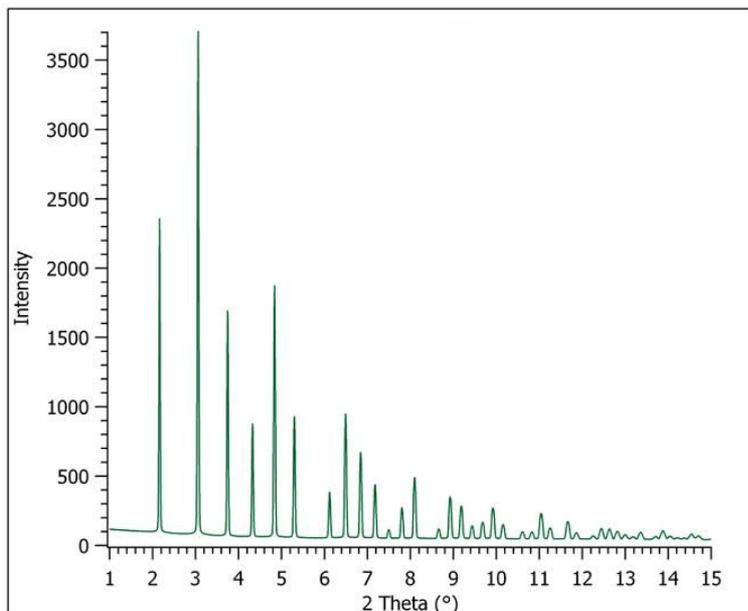
Courtesy of Saul Lapidus, Argonne

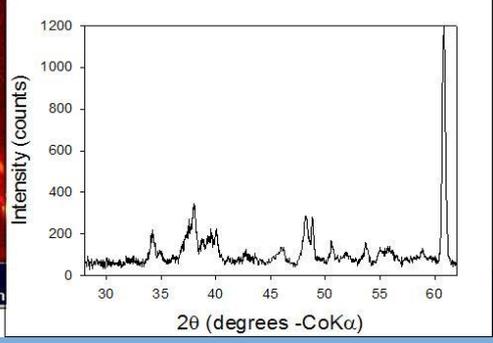
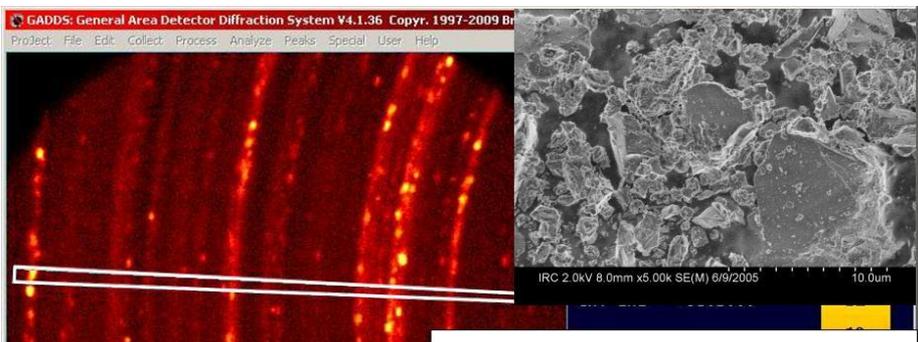
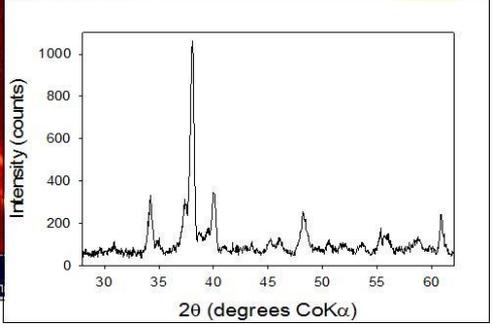
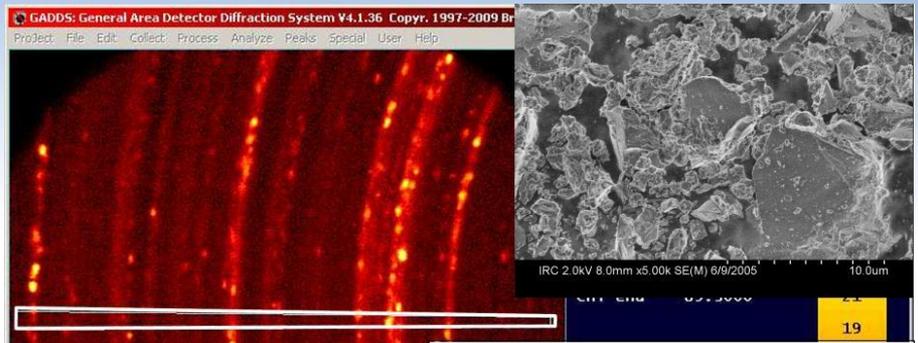
Comparison of 0D and 2D detection

1-D pattern

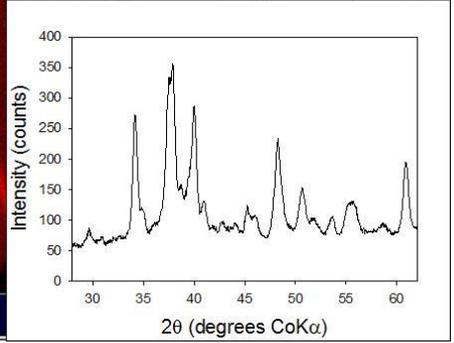
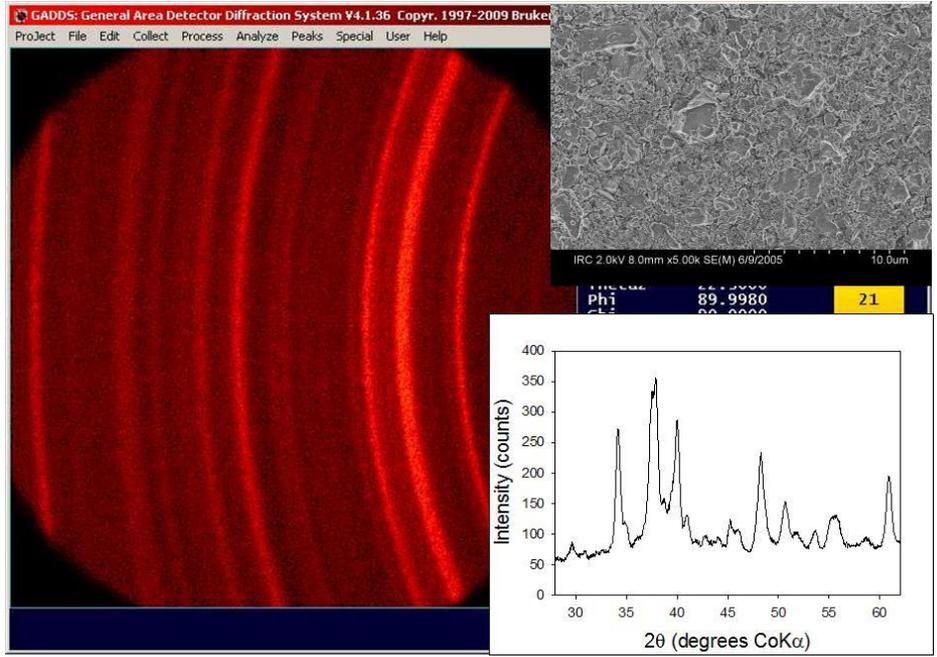
2-D pattern

Radial cross-section
or azimuthal integration





“Seeing” particle statistics



Courtesy of Pamela Whitfield, Oak Ridge (TN)

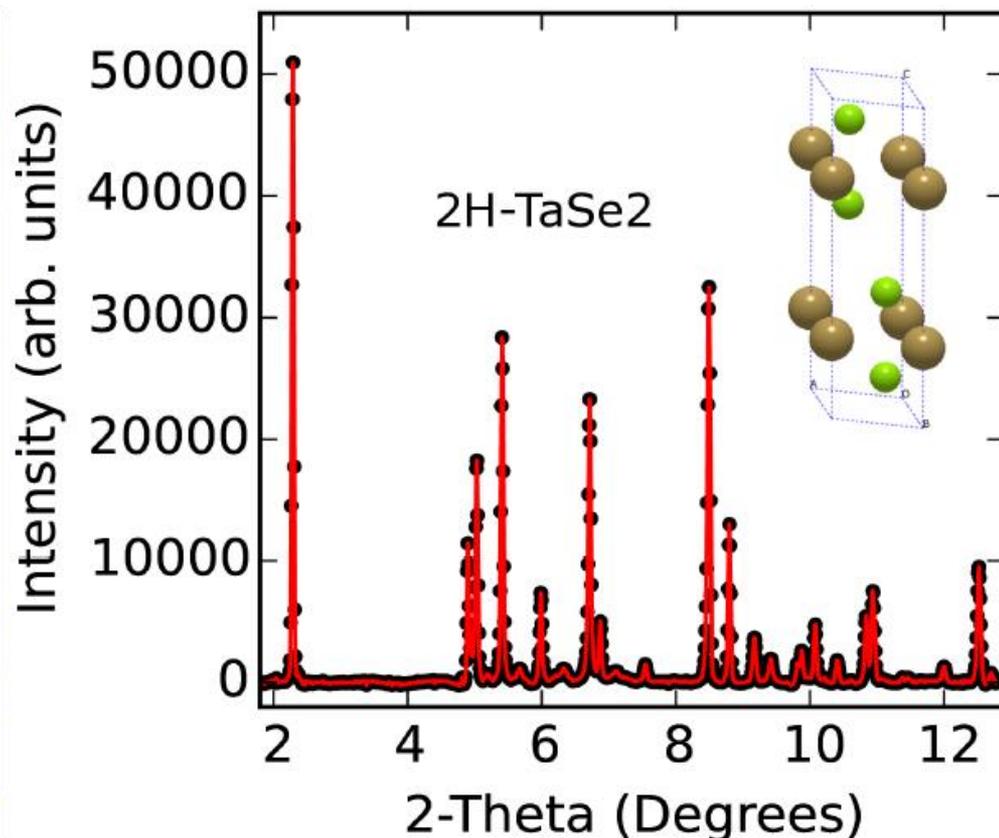
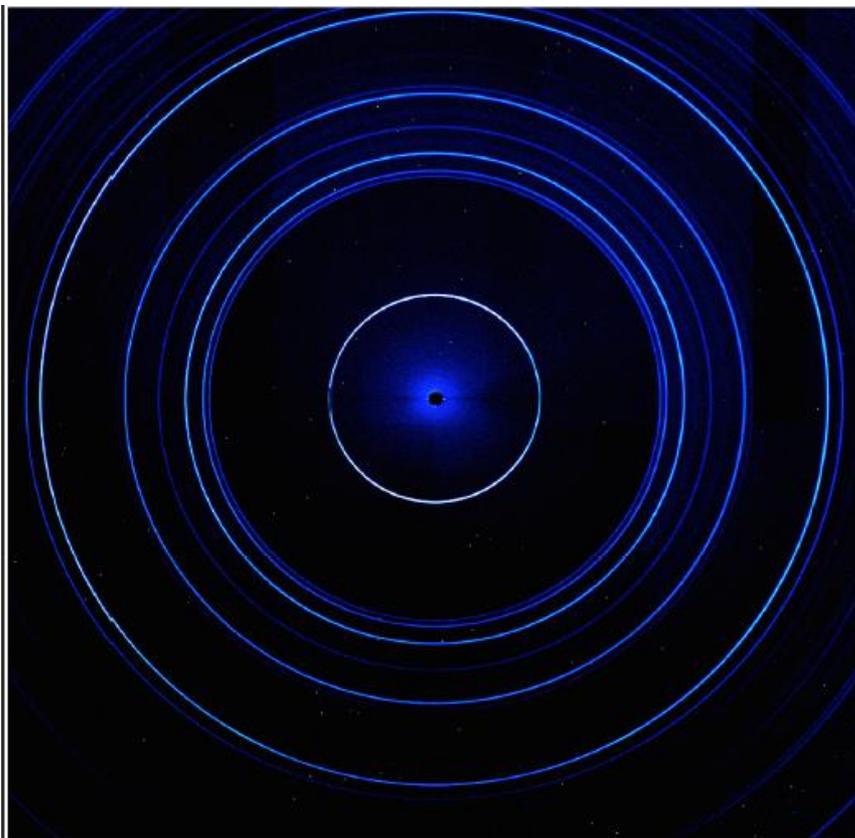
First Dataset at NSLS-II XPD beamline

Nov. 2014

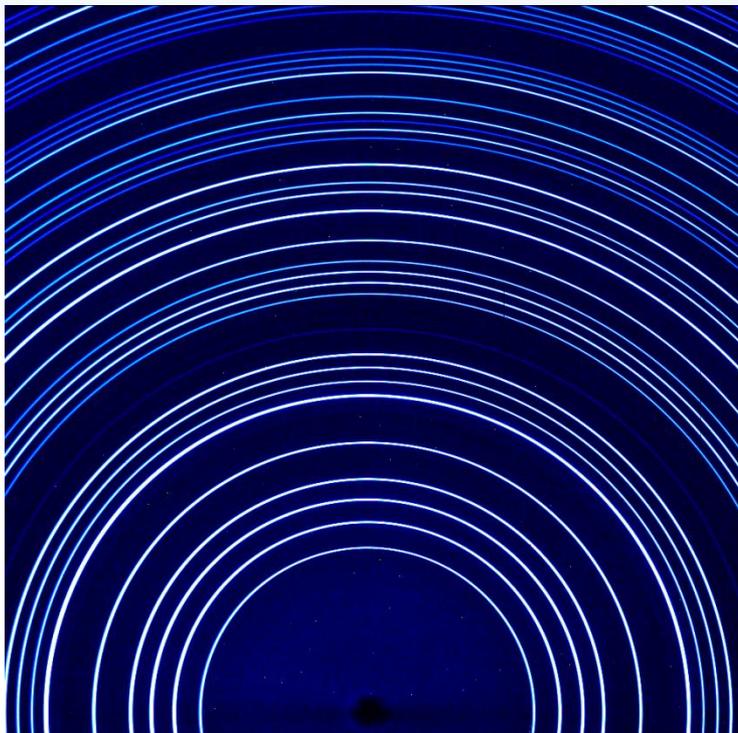
"Superconducting Order from Disorder in 2H-TaSe_{2-x}S_x"

npj Quantum Materials (2017)

Coll. BNL, APAM Columbia, Rutgers U (NJ), Institute of Solid State Phys. in China.

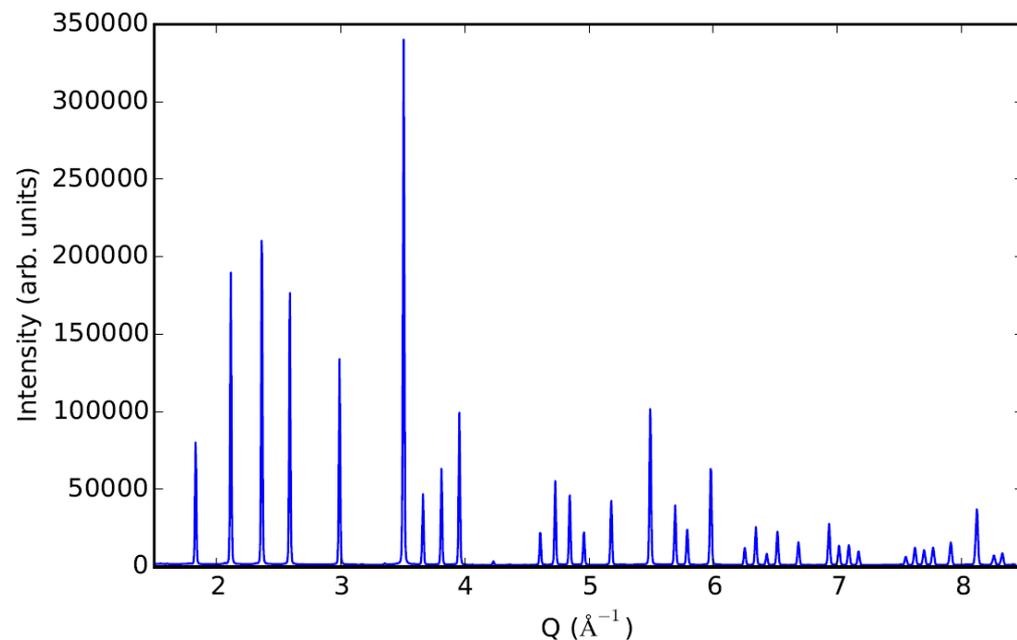


First Publication at NSLS-II



APL Materials 3 (2015) 041513.
Title: "Enhanced Thermoelectric Power and Electronic Correlations in RuSe₂"

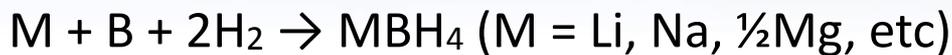
- 48 keV
- 60 sec acquisition per frame
- Collaboration: S. J. L. Billinge and C. Petrovic, Condensed Matter Physics and Materials Science Department, BNL and APAM Columbia.



Decision Making

	Angle-scanning	Large-area	Comments
Resolution	✓		Positron-resolved. Truly angle-resolved only if an analyzer is used. Crystallography
Acquisition rate		✓	In situ, operando
Statistics		✓	Azimuthal integration
Data quality	✓		Better discrimination of the signal, SNR Energy discrimination, parasitic scattering
Data processing	✓		Spatial distortion, intensity non-uniformity (flat field), parallax, dark image, ghost image, masking
Microstructure		✓	Point detector sees only a tiny portion of the scattering. Ring structure reveals preferred orientation, graininess, texture
Normalization		✓	More immune if beam decays or fluctuates, if sample degrades over time (<i>e.g.</i> , radiation damage)

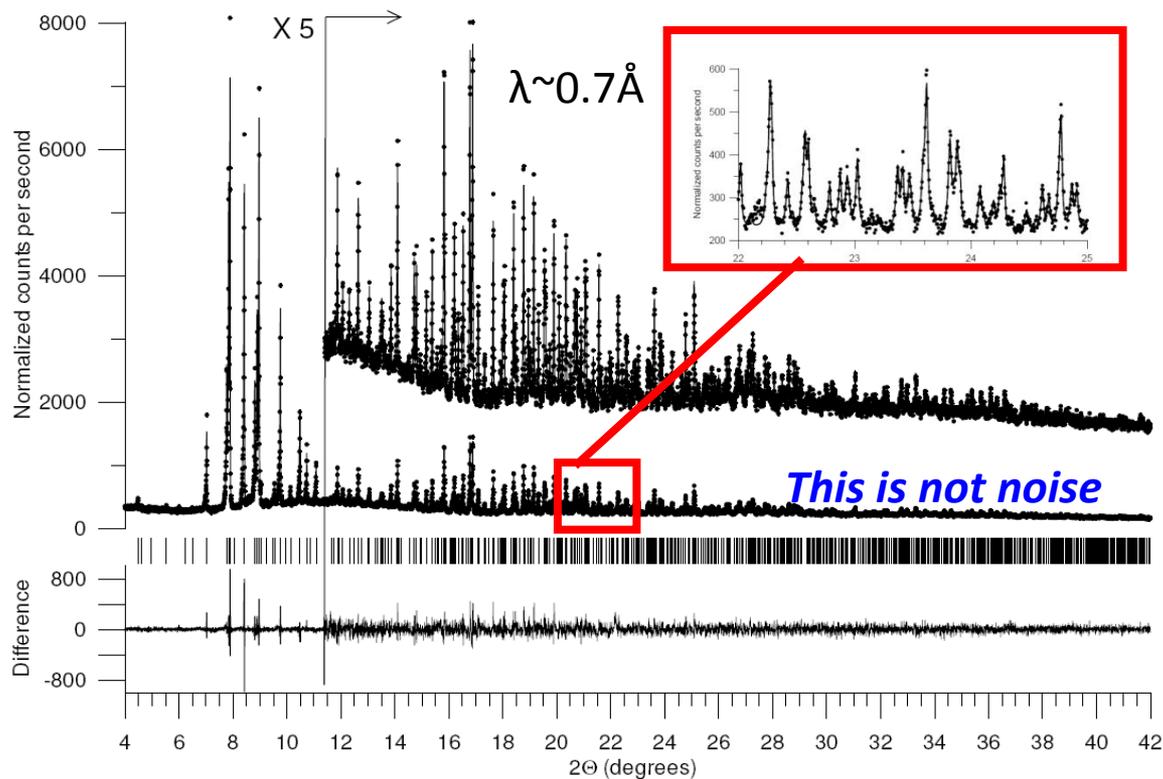
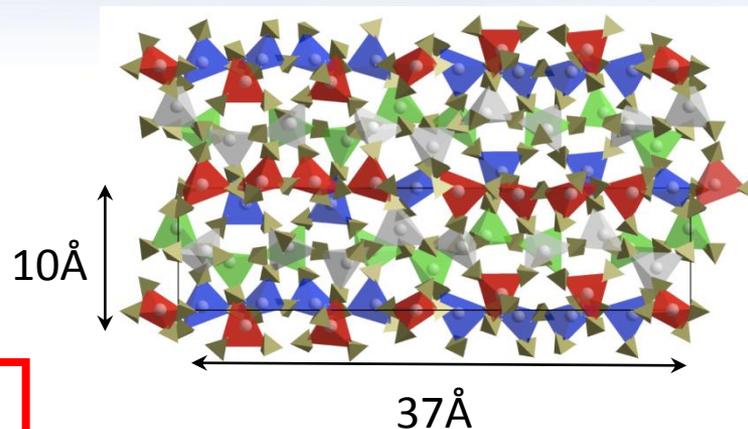
High-Resolution Powder Diffraction



550 - 700° C; 3 - 15 MPa (\approx 200 atm H₂)

Her et al (2007) Acta Crystallog., 63, 561

“Low Temperature” phase of Mg(BH₄)₂”



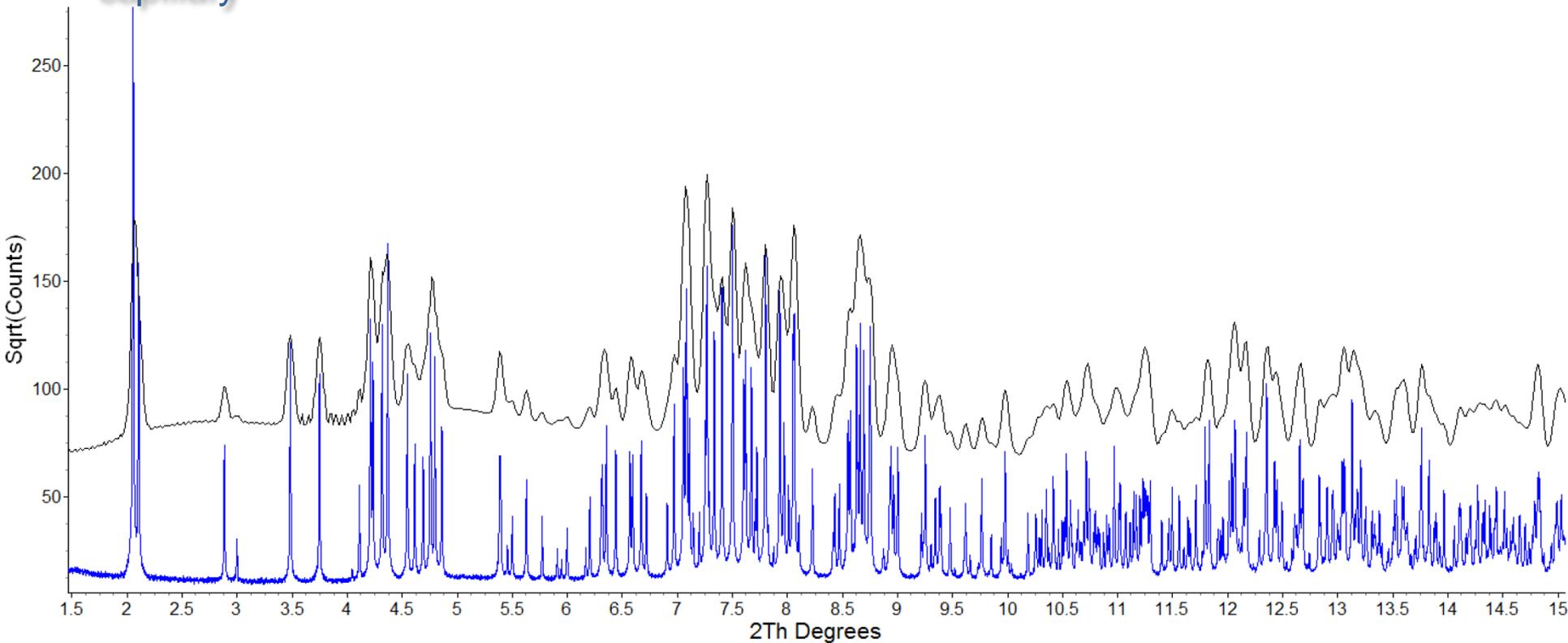
Today's challenge is to achieve this resolution and signal-to-noise performance from sample inside a reactor.

XPD (integrated 2D) vs. 11-BM data (hi-res)

S. Gill (Brookhaven), Greg. Morrison (University of South Carolina)

Gadolinium and Uranium based Salt Inclusion Materials (SIMs) for nuclear waste form applications

In situ ion-exchange of the SIM with NaCl/KCl solution at 90°C in a sealed capillary

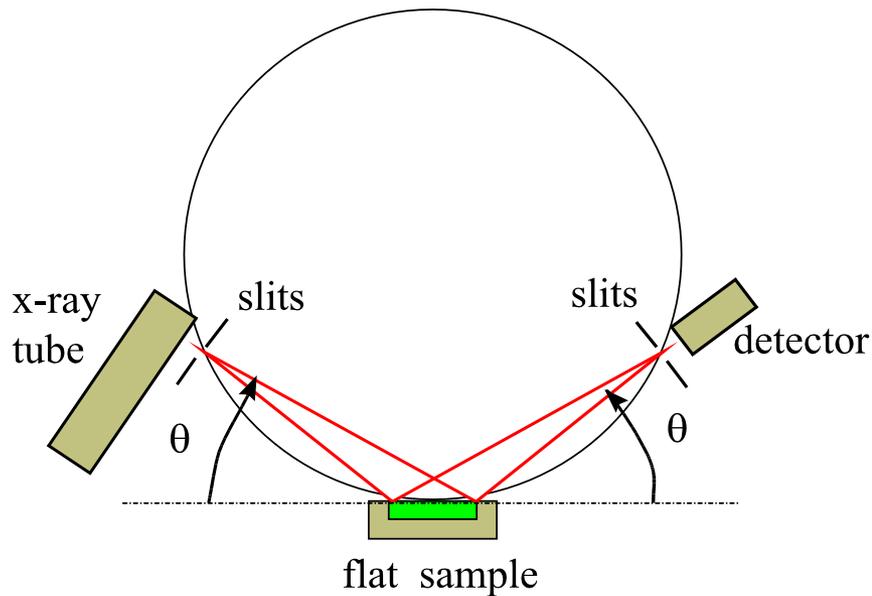


A bit of History

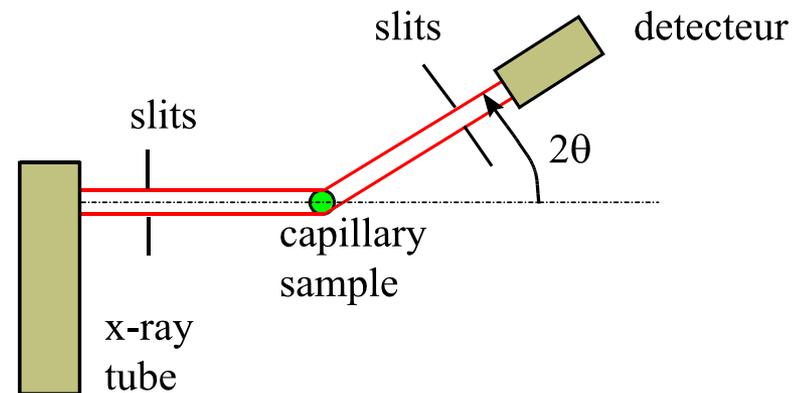
Detectors	Authors	Comments
2D X-ray films	Debye & Scherrer, 1916; Hull, 1917 Guinier, 1939	Debye–Scherrer camera Guinier camera: X-ray optics that focus the scattered beam onto the cylindrical film plane
0D	Brentano, 1925 Parrish, 1949 Cox , 1988 Fitch, 1996 Hodeau & Dooryhee, 1998	A goniometer sweeps a point detector through a range of 2θ angles A crystal analyzer is used with a parallel beam A bank of analyzers
1D PSD	Gabriel & Dupont, 1972 Gobel, 1982 Patterson, 2005	record intensity values as a function of the angular or linear distance along a flat or curved detection strip MYTHEN PSD detector from Dectris
Pixelated 2D	Cernik , 1992; Eikenberry, 2003; Lee, 2007	direct digital readout The intensity sensitivity is more linear than that of film Much larger dynamic range.

Reflection vs. Transmission

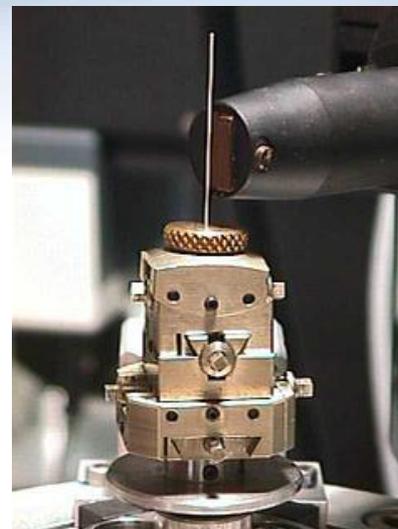
1. Reflection: low energy X-rays, thick or absorbing sample, surface
2. Transmission: high energy X-ray or transparent sample, bulk



Bragg-Brentano para-focusing geometry



Debye Scherrer geometry

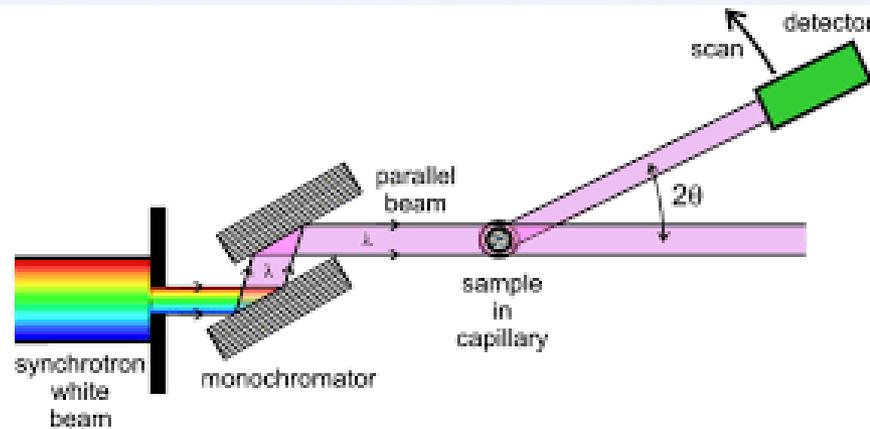


1. Capillary
2. Flat plate



Angle/Energy Dispersive Diffraction

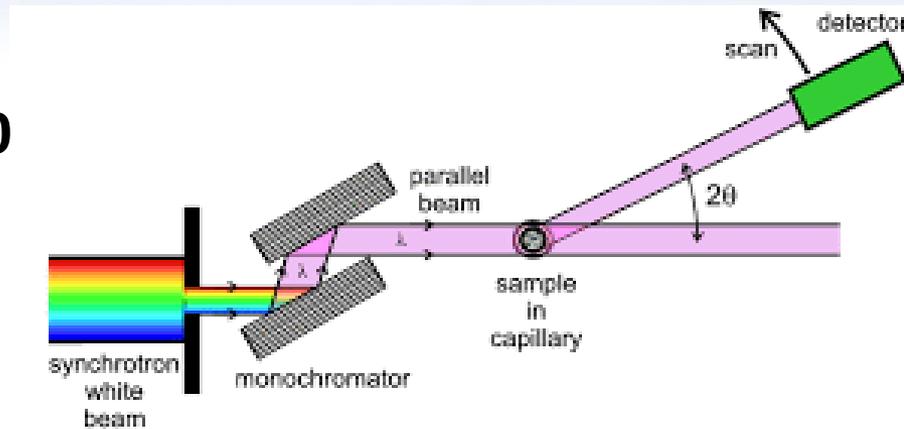
$$2 d \sin\theta = \lambda_0$$



1) Angle Dispersive I(2θ): monochromatic beam and detector is scanned to measure at all possible 2θ angles

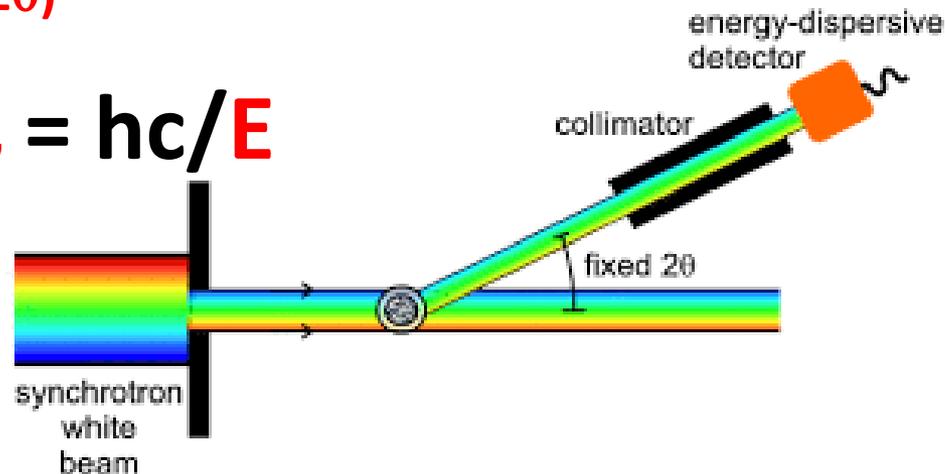
Angle/Energy Dispersive Diffraction

$$2 d \sin\theta = \lambda_0$$



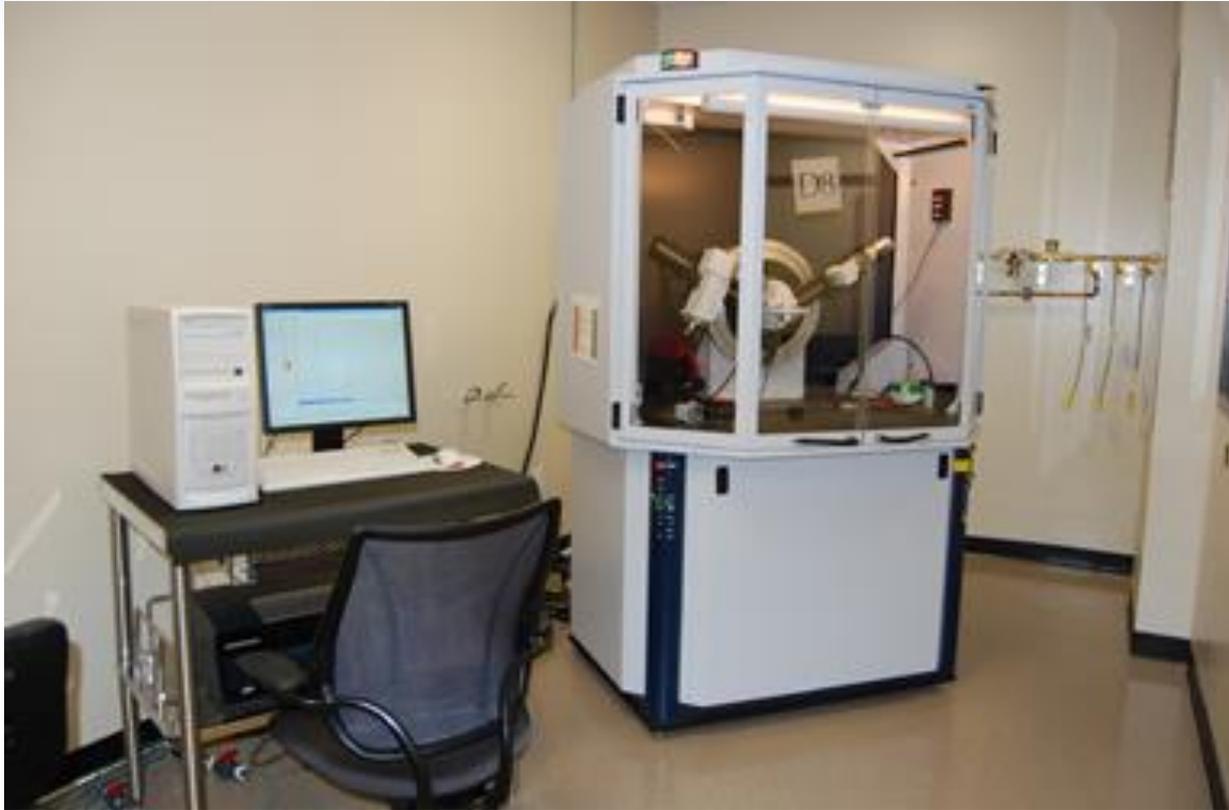
1) Angle Dispersive $I(2\theta)$

$$2 d \sin\theta_0 = \lambda = hc/E$$



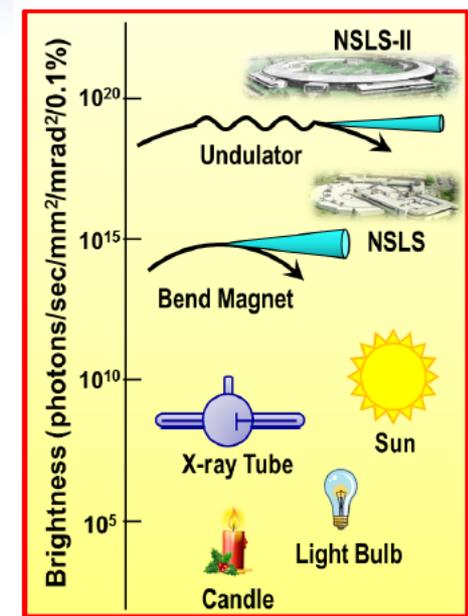
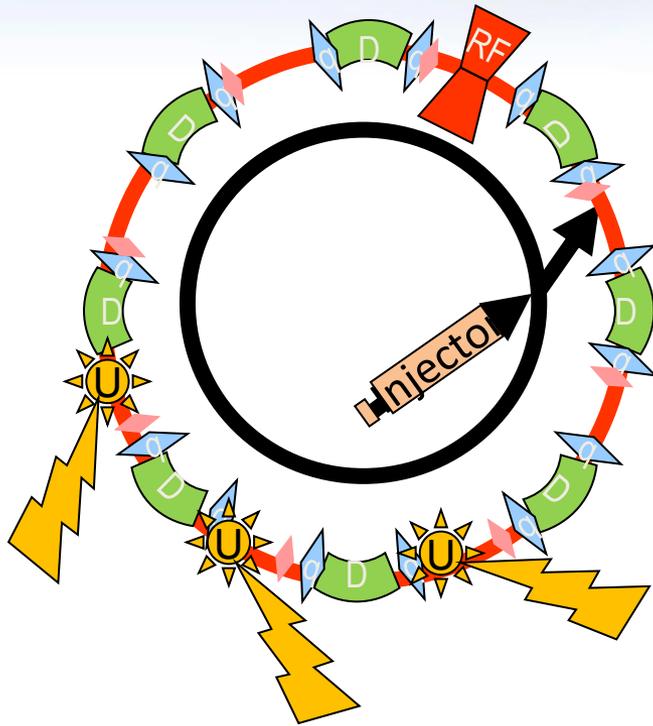
2) Energy Dispersive $I(E)$: polychromatic (white) beam (synchrotron) and detector (SSD with energy discrimination) is at a fixed angle.

From Laboratory Powder Diffraction...



- Flux (weak scatterer, diluted, time-dependent)
- Energy tunability (less absorption, anomalous, white beam)
- Optics (resolution, small beam size)

...to Synchrotron Powder Diffraction



- X-ray source: \$ cost and photon flux $\times 1,000,000$
- Capacity of NSLS-2: 60 beamlines

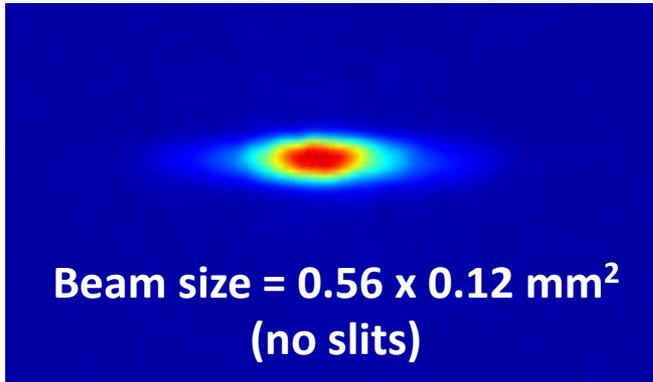
Source and Front End



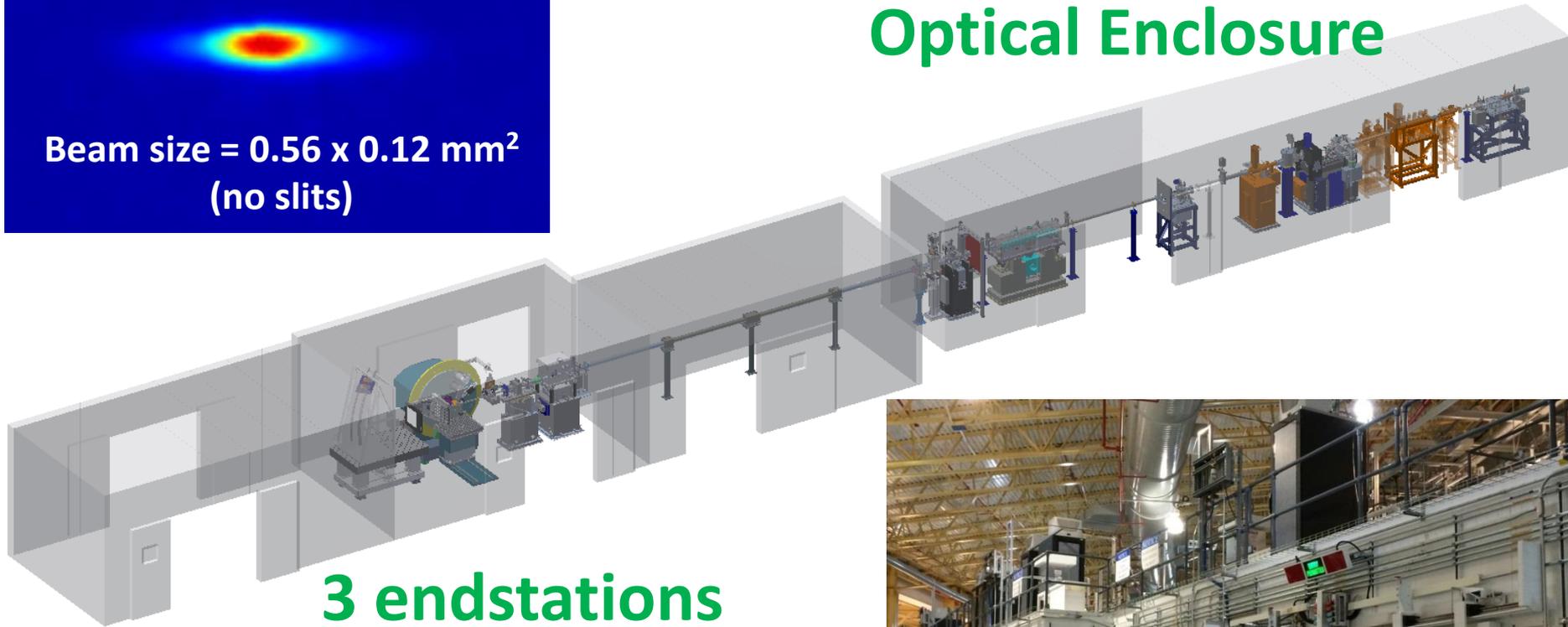
Damping Wiggler

Front End (shutter , masks, slits)

Powder Diffraction Beamline (XPD)



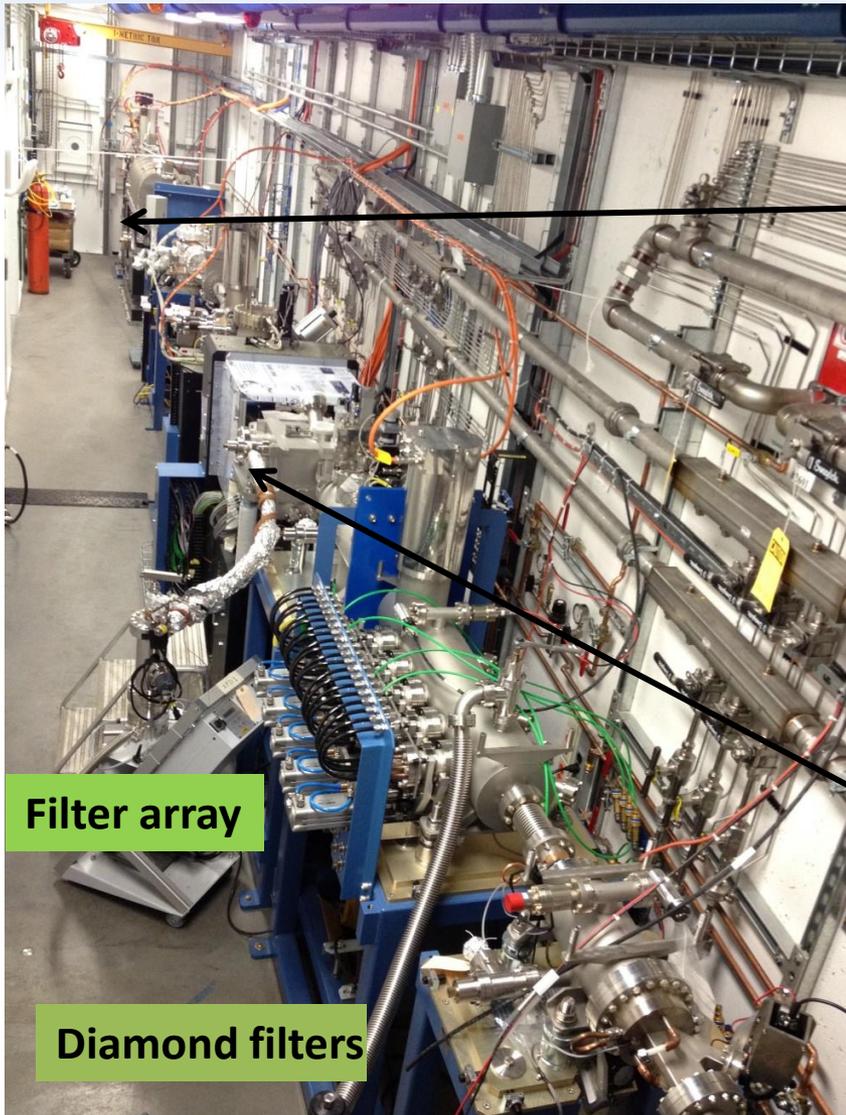
Optical Enclosure



5 years from design to start of operations

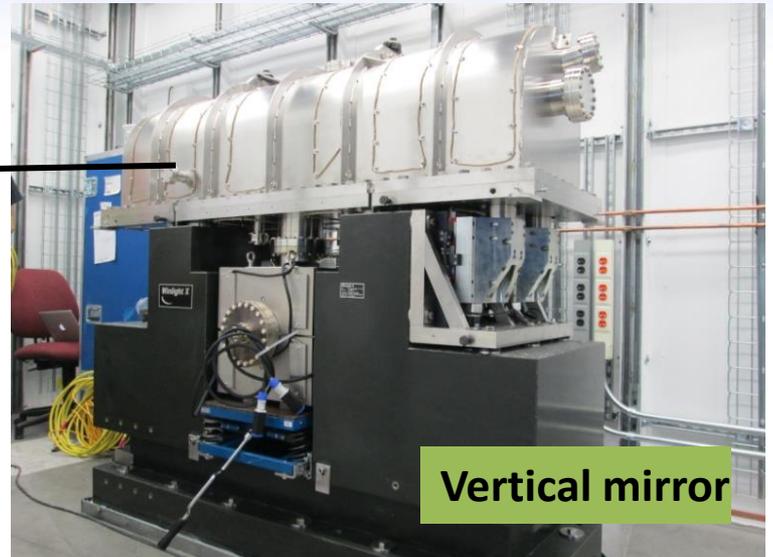


XPD OPTICAL ENCLOSURE

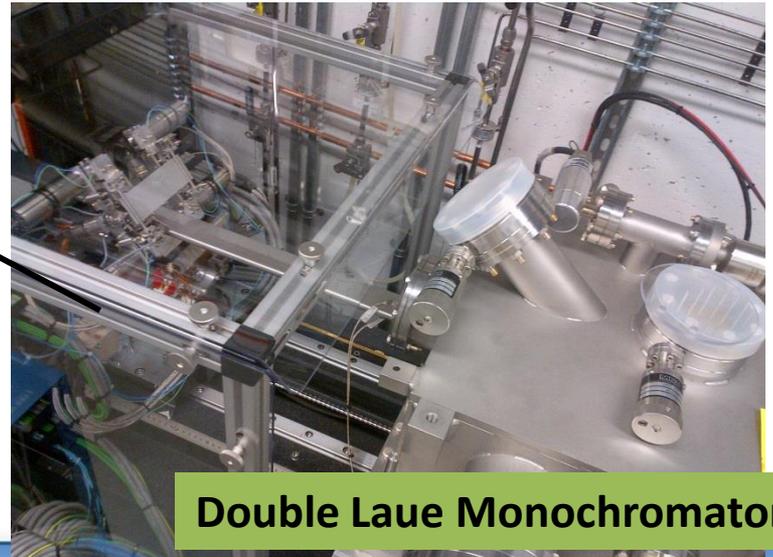


Filter array

Diamond filters



Vertical mirror



Double Laue Monochromator

VERTICALLY FOCUSING MIRROR



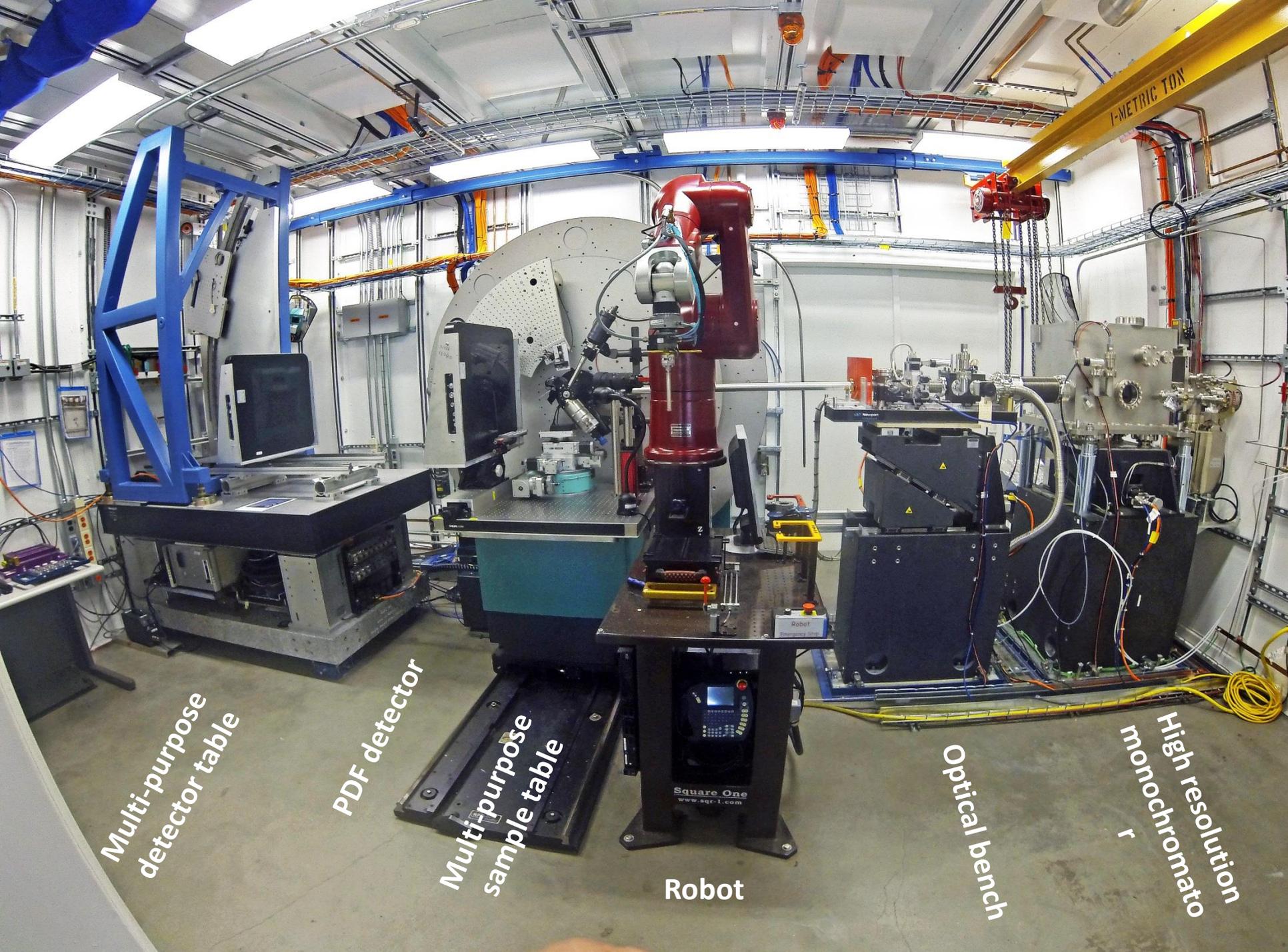
1.4 m long

Roughness
0.3nm rms

Bending 10 km
(anti-sag)

Slope Error:
0.36 microrad
(0.02 mdeg)

At 50 miles ?



Multi-purpose
detector table

PDF detector

Multi-purpose
sample table

Robot

Optical bench

High resolution
monochromato
r

ROBOTIC SAMPLE MANIPULATOR

1. Acquisition of a diffraction image takes seconds.

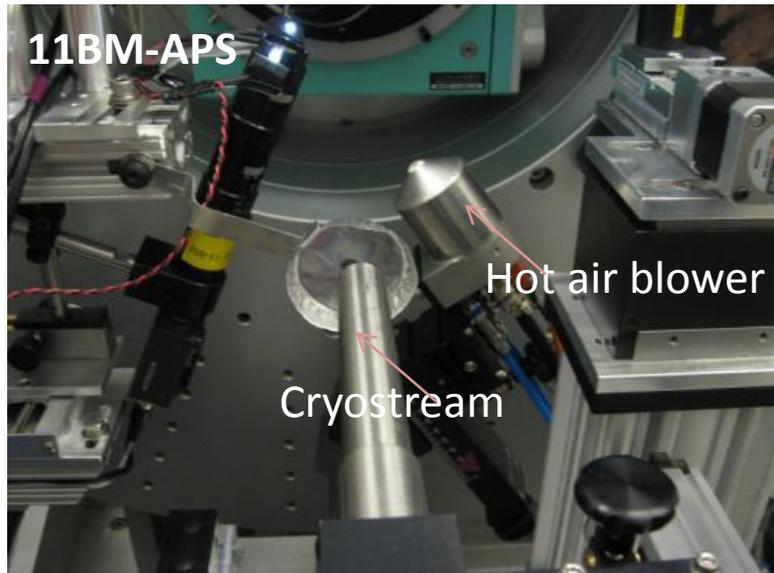
Breaking the interlock system, entering the hutch, changing the sample by hand, exiting the hutch, reactivating the interlock take ≥ 5 min.

2. Probability of operator's mistake during sample manipulation

3. Handling of radioactive samples (samples are kept in a shielded box at all times, personnel is never exposed during measurement)

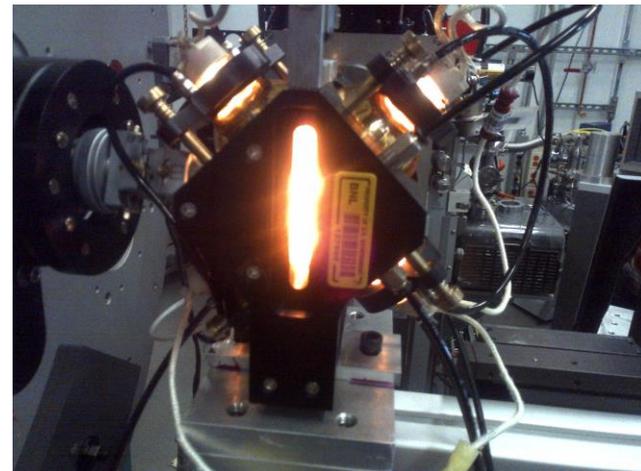
SAMPLE ENVIRONMENTS

LN2 cryostream, LHe cryostat, hot air blower, flat plate heater, quadrupole lamp furnace,...



Capabilities:

- Temperature Range: 80K, Ambient to 2,000°C
- Sample holder Type: capillary (compatible with robotic sample changer), or flat plate

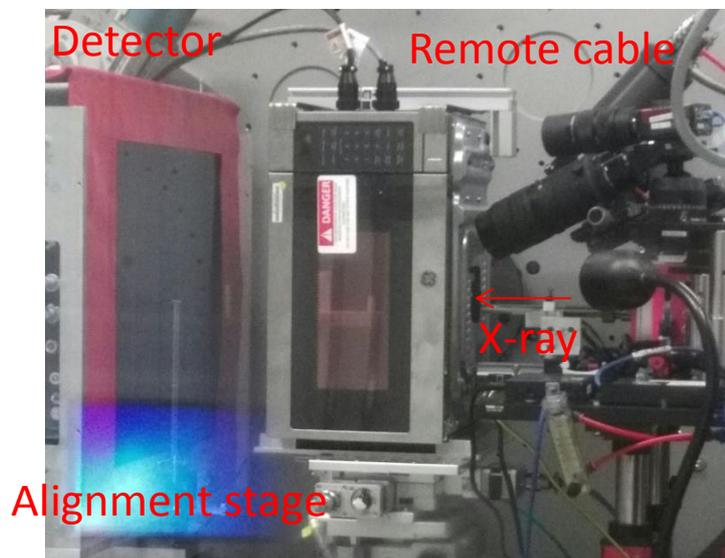


MICROWAVE REACTOR FOR IN SITU DIFFRACTION

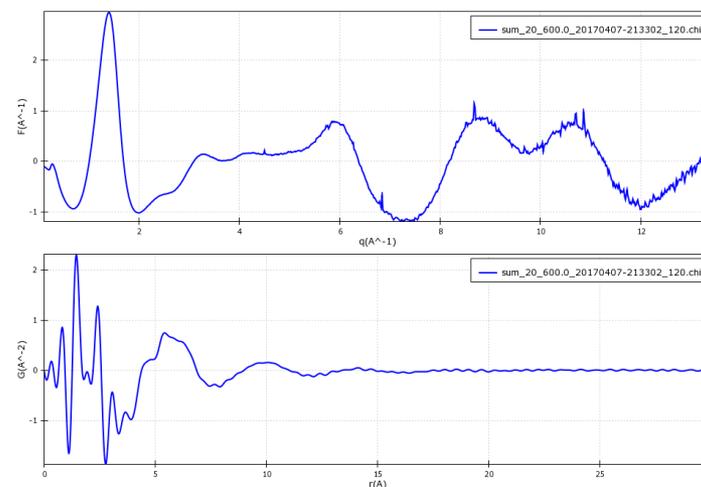
Science motivation: MW assisted materials synthesis is a fast growing field. The MW reactor enabled combined in situ PDF and XRD to study the crystallization process from the stage of nucleation.

Applications: Hydro-Solvothermal synthesis, solid state synthesis, thin film growing in solution

J. Bai



In Situ MW in operation with remote control



First in situ PDF measurement during MW assisted Titania (TiO_2) thin film growing @ 28-ID-2 (April 2017)

NEUTRONS vs. SYNCHROTRON

X-rays are scattered by

Neutrons are scattered by

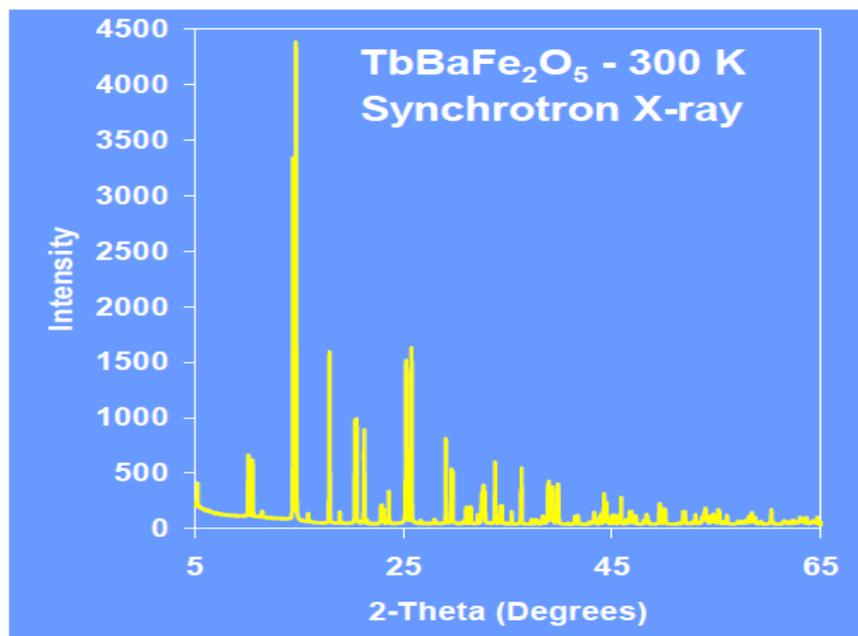
NEUTRONS vs. SYNCHROTRON

X-rays are scattered by the electrons in the atoms

Neutrons are scattered by the nuclei

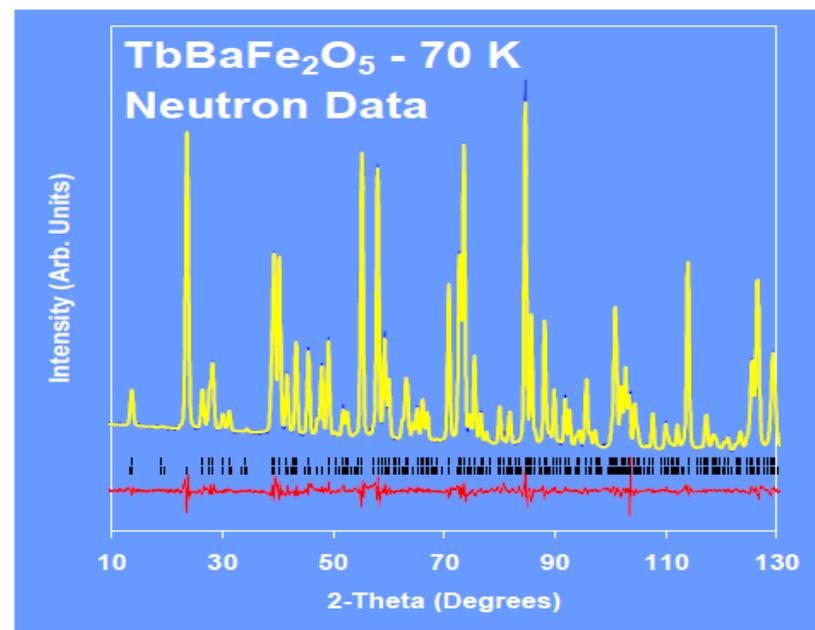
NEUTRONS vs. SYNCHROTRON

Synchrotron X-rays



The peak intensities drop off at high angles in an X-ray diffraction pattern because the form factor decreases

Neutrons



Neutrons are scattered from the nucleus and the form factor is not angle dependent. Intensities do not drop off at high angle.

NEUTRONS vs. SYNCHROTRON

Synchrotron X-rays

- Extremely **high resolution**
- Subtle distortions
- Indexing and Structure determination
- Anomalous scattering
- Texture effects

Neutrons

- scattering length contrast
- **More sensitive to light elements (H, N, O, F, C, etc.) and isotope distributions**
- Low absorption: **large samples**, easy sample environments
- **Magnetic** structures
- Moderate resolution

Outline

1. Applications for Powder Diffraction
2. The basics of Powder Diffraction
3. Experimental Methods for collecting PD data
- 4. Crystal structure Determination**
- 5. Peak Profile Analysis for Microstructural Studies**
- 6. Advanced PD techniques**

Crystal Structure: **lattice**

the unit cell

Crystal data

Formula

O₂ Ti

Crystal system

tetragonal

Space group

P 42/m n m (no. 136)

Unit cell dimensions

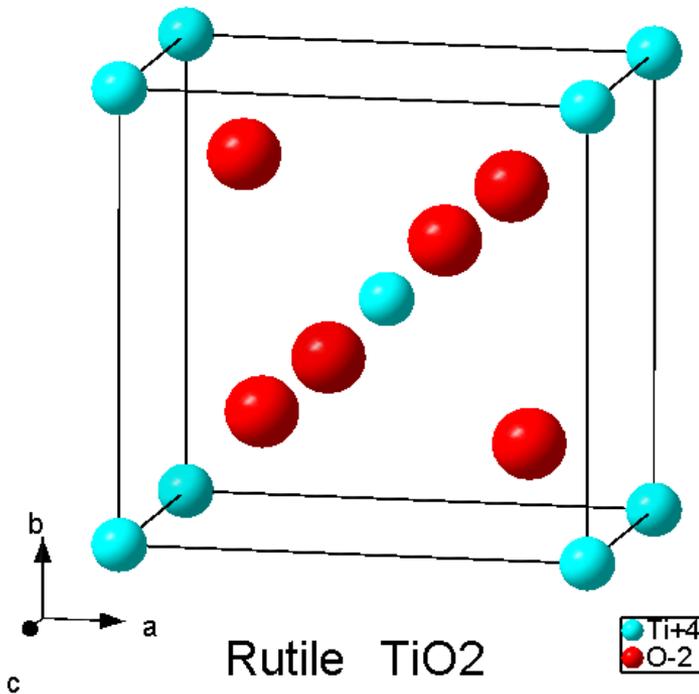
$a = 4.5937 \text{ \AA}$ $c = 2.9587 \text{ \AA}$

Cell volume

62.40 \AA^3

Z

2



Crystal Structure: + symmetry

the unit cell

the space group

Crystal data

Formula

O₂ Ti

Crystal system

tetragonal

Space group

***P 42/m n m* (no. 136)**

Unit cell dimensions

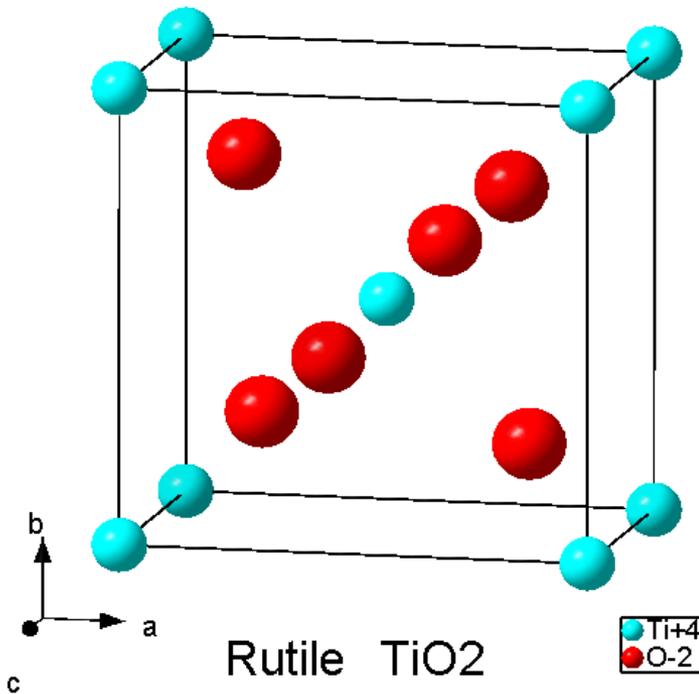
$a = 4.5937 \text{ \AA}$ $c = 2.9587 \text{ \AA}$

Cell volume

62.40 \AA^3

Z

2



Crystal Structure: + atomic motif

the unit cell

the space group

the asymmetric unit

Crystal data

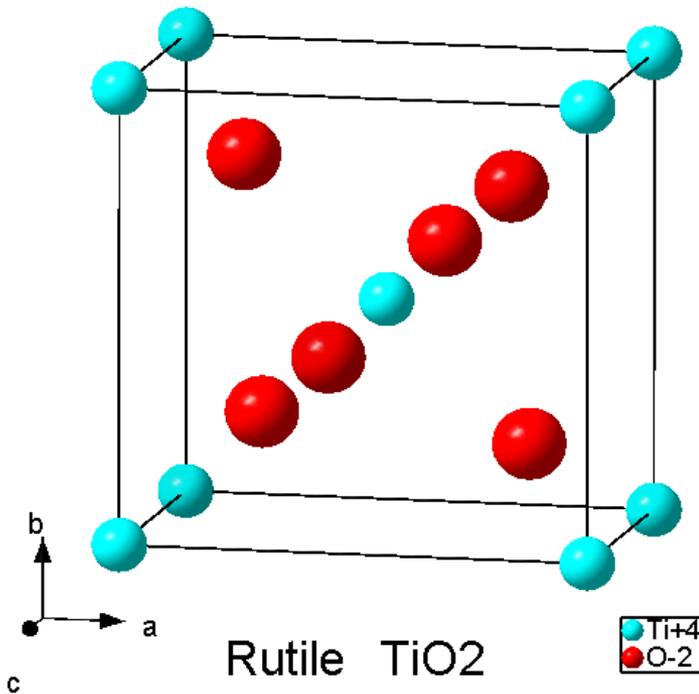
Formula	O ₂ Ti
Crystal system	tetragonal
Space group	<i>P</i> 42/m n m (no. 136)
Unit cell dimensions	$a = 4.5937 \text{ \AA}$ $c = 2.9587 \text{ \AA}$
Cell volume	62.40 \AA^3
Z	2

Atomic coordinates

Atom	Ox.	Wyck.	x	y	z
Ti	+4	2 <i>a</i>	0	0	0
O	-2	4 <i>f</i>	0.30469	0.30469	0

Bond lengths, bond angles (\AA , $^\circ$)

Ti—O	1.949(0)	Ti—Ti	2.959
O—O	2.959	Ti—Ti	3.569



Crystal Scattering Amplitude

A crystal = convolution of an **atomic motif** $M(\vec{r})$ and a **lattice** $R(\vec{r})$

$$C(\vec{r}) = R(\vec{r}) * M(\vec{r})$$

The scattered amplitude corresponds to the Fourier transform of the scattering object

$$C(\vec{h}) = TF\{C(\vec{r})\} = TF\{R(\vec{r}) * M(\vec{r})\} = R(\vec{h}) \times F(\vec{h})$$

The Fourier space is called **reciprocal space**

$R(\vec{h})$ = FT of the (direct) lattice: **Reciprocal Lattice**

$F(\vec{h})$ = FT of the atomic motif: **Structure factor**

Knowing the reciprocal lattice and the structure factors is equivalent to knowing the structure in real space

Real and reciprocal space

CRYSTAL SPACE

DIFFRACTION SPACE

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

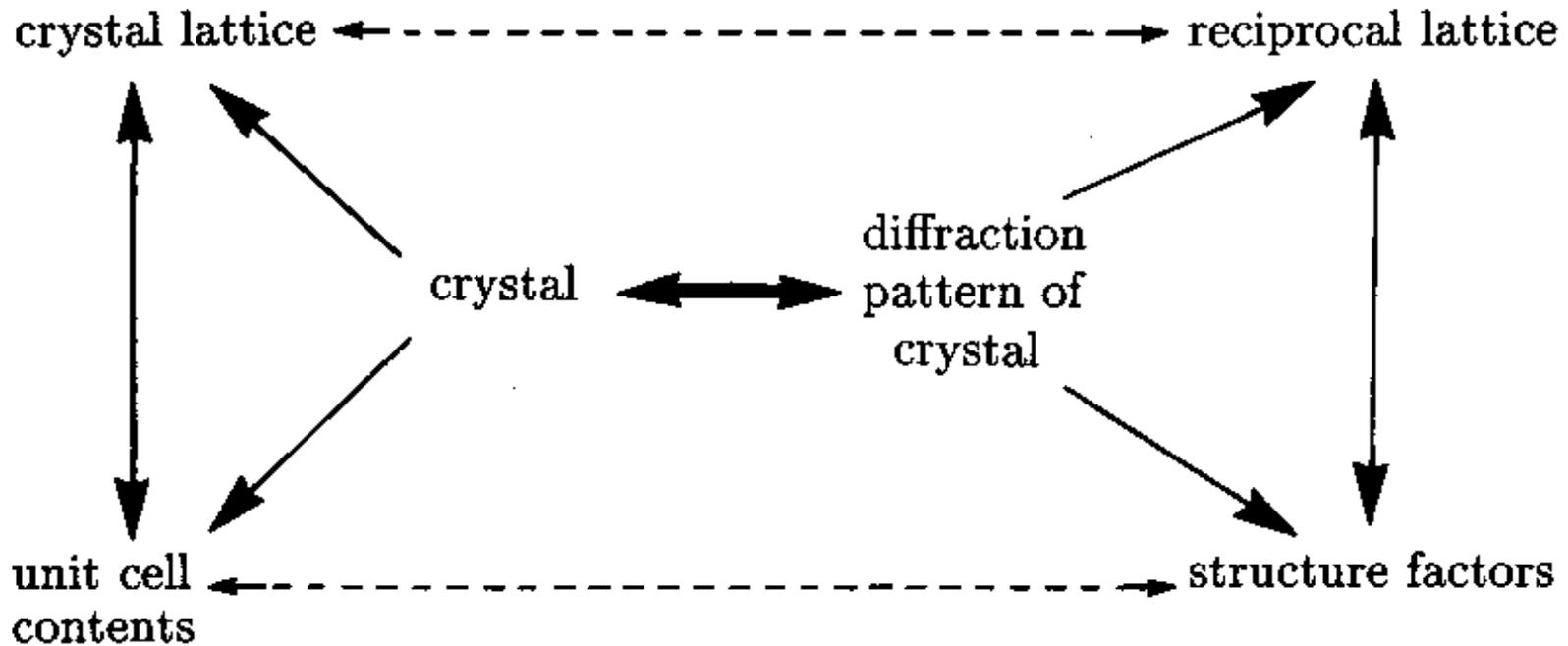


FIGURE 3.11. Relationships between a crystal (in crystal space) and its reciprocal lattice (in diffraction space).

Reciprocal lattice constants

- ◆ The mathematical definition of the reciprocal lattice constants is

- $$\mathbf{a}^* = \frac{\mathbf{b} \times \mathbf{c}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

where $\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c}) = V$

- $$\mathbf{b}^* = \frac{\mathbf{c} \times \mathbf{a}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

$V =$ volume of the unit cell

- $$\mathbf{c}^* = \frac{\mathbf{a} \times \mathbf{b}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

It follows that $|\mathbf{d}| = \frac{1}{|\mathbf{d}^*|}$

- ◆ Alternative formulation:

- $$\mathbf{g}_i \cdot \mathbf{h}_j = \delta_{ij}$$

Courtesy of Cora Lind-Kovacs

Reciprocal lattice constants

- ◆ The mathematical definition of the reciprocal lattice constants is

- $$\mathbf{a}^* = \frac{\mathbf{b} \times \mathbf{c}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

- $$\mathbf{b}^* = \frac{\mathbf{c} \times \mathbf{a}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

- $$\mathbf{c}^* = \frac{\mathbf{a} \times \mathbf{b}}{\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c})}$$

where $\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c}) = V$

V = volume of the unit cell

It follows that $|\mathbf{d}| = \frac{1}{|\mathbf{d}^*|}$

- ◆ Alternative formulation:

- $$\mathbf{g}_i \cdot \mathbf{h}_j = \delta_{ij}$$

What is the symmetry of the reciprocal lattice ?

Courtesy of Cora Lind-Kovacs

Crystal Structure Factor

F_{hkl} : structure factor for reflection (hkl) = TF{ $M(\vec{r})$ }, atomic motif

F_{hkl} : *Fourier transform of the scattering (electron) density of the crystal*

$$F_{hkl} = \sum_{j \in \text{cell}} f_j T_j \exp(2i\pi(hx_j + ky_j + lz_j))$$

f_j : **scattering factor** for atom j with coordinates (x_j, y_j, z_j) . FT of the scattering (electron) density of each individual atom j in ONE unit cell.

For x-rays, f_j is the atomic form factor,

For neutrons, $f_j = b_j$ is the Fermi length.

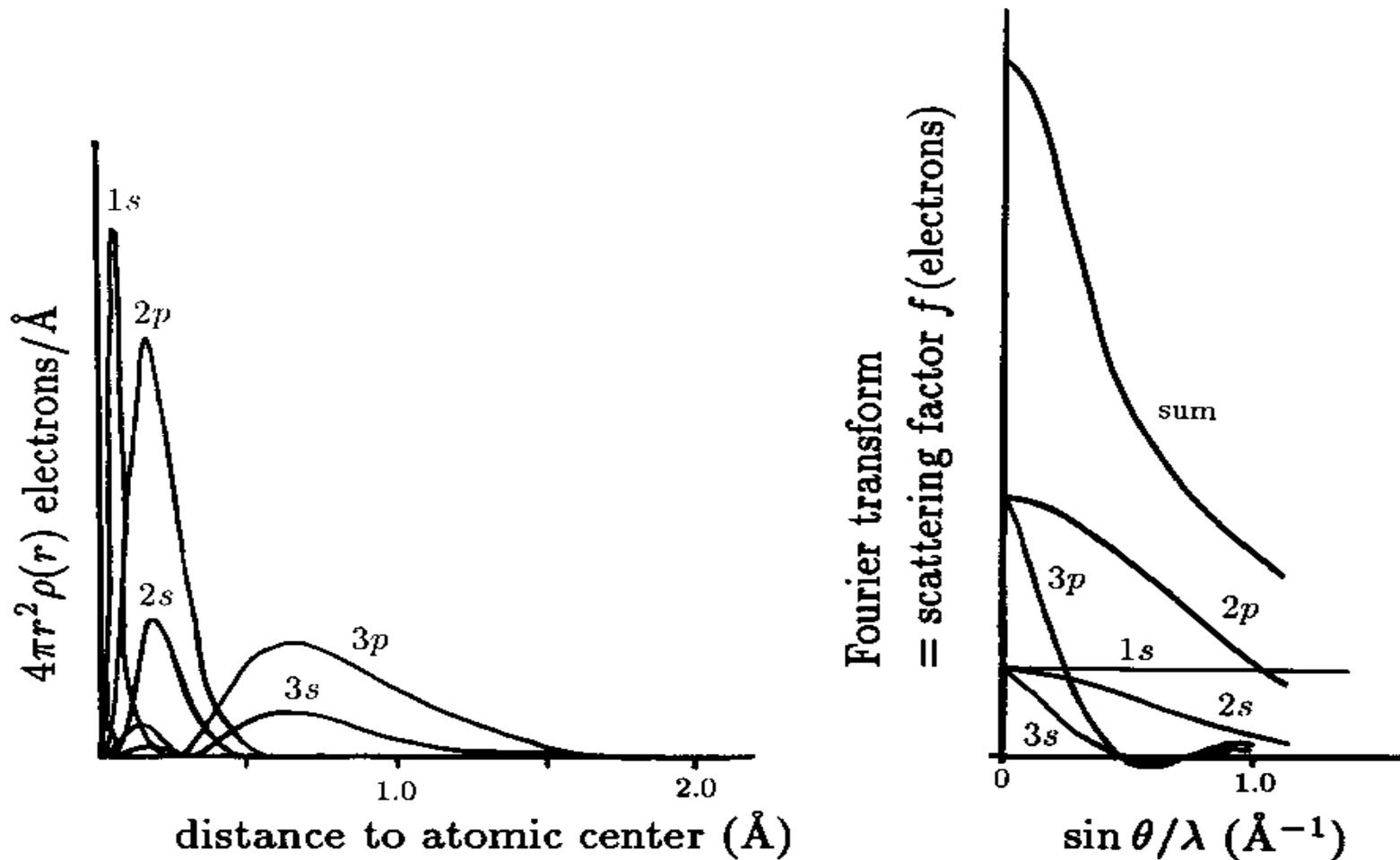
We can evaluate (compute) this complex number f_j as a trigonometric function:

$$\mathbf{f} = A_0 e^{i\varphi} = A_0 \{ \cos\varphi + i \sin\varphi \}$$

Notice that λ and the unit cell parameters are **NOT** part of this.

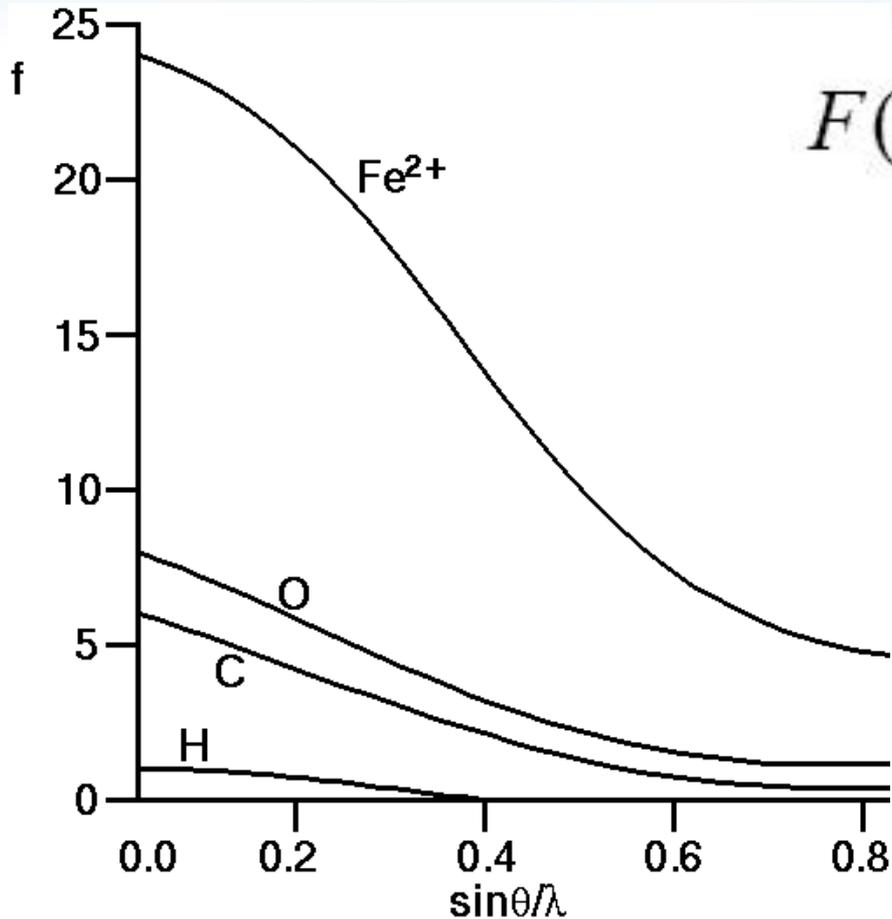
F depends only on the hkl of the planes, nature of the atoms j and their fractional positions.

The FT relationship between f and ρ_{atomic}



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

Atomic Form Factors



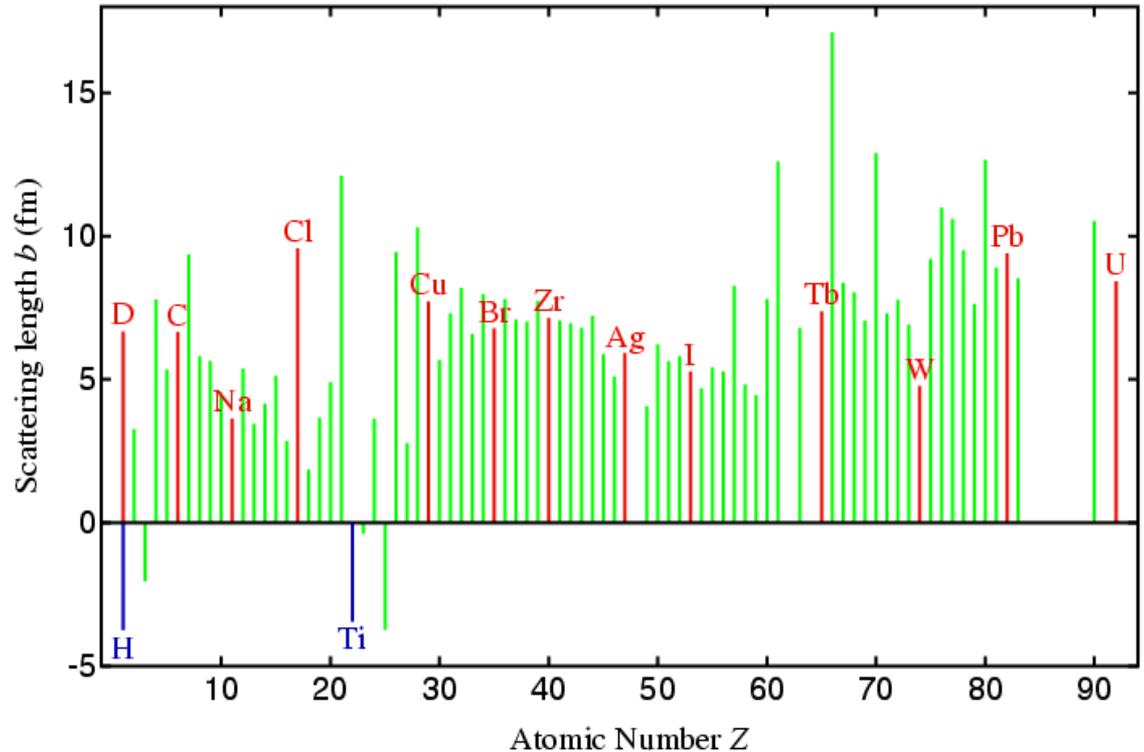
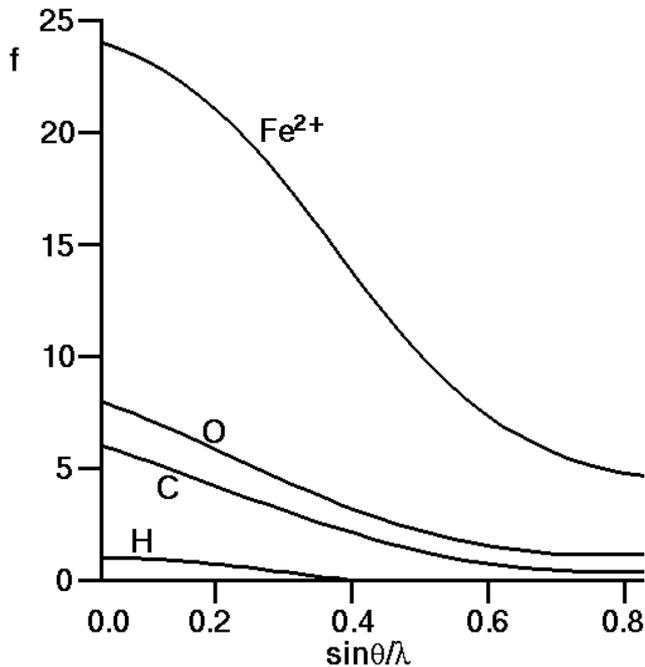
Atomic form factor = scattering amplitude

$$F(hkl) = \sum_j f_j e^{2\pi i(hx_j + ky_j + lz_j)}$$

Structure factor is the Fourier transform of the electron density. It takes into account the individual scattering f of all j atoms contained in the crystal unit cell.

Notice that data are plotted as a function of $\sin\theta/\lambda$. This is the scattering vector in reciprocal units \AA^{-1} .

X-ray vs. neutron Scattering Length



The neutron scattering length does not vary with $\sin\theta/\lambda$. It varies with Z and isotope number.

Crystal Structure Factor

F_{hkl} : structure factor for reflection (hkl) = TF{ $M(\vec{r})$ }, atomic motif

F_{hkl} : *Fourier transform of the scattering density of the crystal*

$$F_{hkl} = \sum_{j \in \text{cell}} f_j T_j \exp(2i\pi(hx_j + ky_j + lz_j))$$

f_j : **scattering factor** for atom j with coordinates (x_j, y_j, z_j) . FT of its scattering density of each individual atom j .

For x-rays, f_j is the **atomic form factor**,

For neutrons, $f_j = b_j$ is the **Fermi length**.

The **amplitude** of scattering depends on the electron density (atoms).

The **phase** depends on the fractional distance of the atoms to the lattice plane.

$T_j =$ **Debye-Waller** factor related to atomic displacements (thermal, positional,...).

The atomic “Temperature” factor

Can we describe the **vibration** of an individual atom?

If the vibration is harmonic, the probability distribution will be Gaussian with a form $\exp(-a \cdot x^2)$. The diffraction pattern of a Gaussian is another Gaussian, where the argument is related to $1/d$ -spacing

We usually use this in the Structure Factor expression:

$$\mathbf{T} = \exp(-B_j \mathbf{Q}^2) = \exp(-B_j \sin^2\theta / \lambda^2)$$

where $B = 8\pi^2 \langle u^2 \rangle$ where \mathbf{u} is the r.m.s. displacement of a scattering center, and $\langle \dots \rangle$ denotes time averaging.

We call B the “**Temperature factor**” because it rises with the temperature of the crystal. You see that it’s **more properly a measure of mean square displacement**, whatever the temperature. → Anisotropic Displacement Factor

Courtesy of Robert Sweet “X-ray Diffraction & Crystallography”

Crystal Diffracted Intensity

The diffracted intensity is the quantity accessible in a diffraction experiment (proportional to the number of diffracted photons/particles reaching the detector)

In the **kinematic approximation** (neglecting double diffraction), one has :

$$I_{hkl} = S \cdot C_{hkl} \cdot |F_{hkl}|^2$$

S : scale factor (amount of sample, count time, incident flux,...)

C_{hkl} : experimental corrections

instrument (*Lorentz, polarization, diffraction geometry...*)

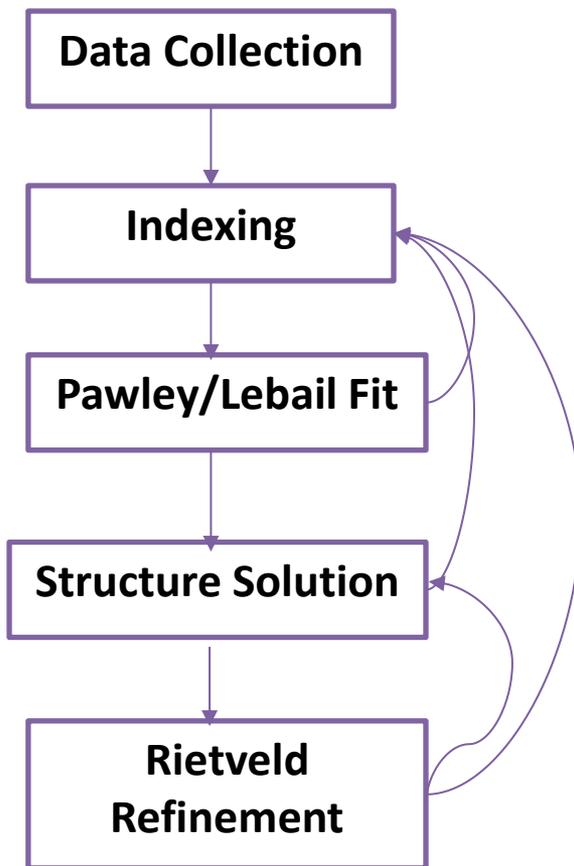
sample (*multiplicity, absorption, preferred orientation, extinction...*)

$$F_{hkl} = \sum_{j \in \text{cell}} f_j T_j \exp(2i\pi(hx_j + ky_j + lz_j))$$

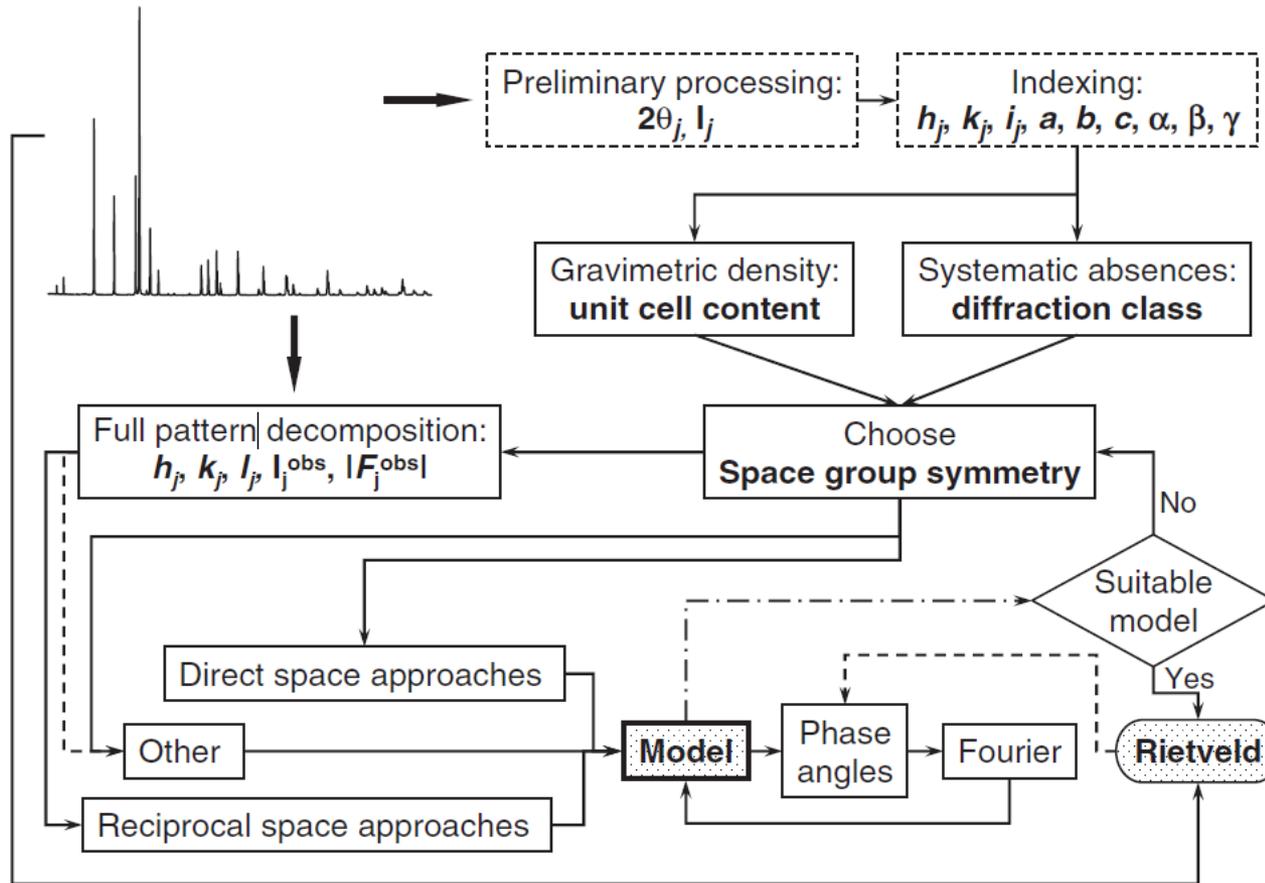
F_{hkl} : complex number => **phase of F_{hkl} not measured**. Information is incomplete

To solve the structure, one must retrieve the phases of F_{hkl} , one way or another...

Flowchart illustrating crystal structure determination from powder diffraction.



Flowchart illustrating crystal structure determination from powder diffraction.



Source: V.K. Pecharsky, P.Y. Zavalij, *Fundamentals of Powder Diffraction and Structural Characterization of Materials*, © Springer Science 2009

Ab Initio Structure Solving

- 1) Needs a high quality sample already well characterized (composition, density, ...)
- 2) Carry out one or more *well adapted* diffraction experiments
Optimize the data quality (low background, low peak overlapping,...)
(x-rays and/or neutrons and/or electrons, choice of resolution, wavelength, *etc...*)

Ab Initio Structure Solving

- 1) Needs a high quality sample already well characterized (composition, density, ...)
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Optimize the data quality (low background, low peak overlapping,...)
(x-rays and/or neutrons and/or electrons, choice of resolution, wavelength, *etc...*)
- 3) Find unit cell and index Bragg reflections (Dicvol, Treor, Ito, *etc...*). See www.ccp14.ac.uk/solution/indexing/. See 'Peter W. Stephens (SBU): *Indexing Powder Diffraction Data*' tutorial on youtube.

also see '*Discussion of the indexing algorithms within TOPAS*'. A. Coelho & A. Kern (2005) CPD Newsletter No. 32, 43-45

→ finding the unit cell symmetry and dimensions that best describe the position of the diffraction peaks on the powder pattern

Ab Initio Structure Solving

- 1) Needs a high quality sample already well characterized (composition, density, ...)
- 2) Carry out one or more *well adapted* diffraction experiments
Optimize the data quality (low background, low peak overlapping,...)
(x-rays and/or neutrons and/or electrons, choice of resolution, wavelength, *etc...*)
- 3) Find unit cell and index Bragg reflections (Dicvol, Treor, Ito, *etc...*). See www.ccp14.ac.uk/solution/indexing/. See ‘Peter W. Stephens: *Indexing Powder Diffraction Data*’ tutorial on youtube.
- 4) Obtain intensities of Bragg reflections and determine the space group
(LeBail technique + CheckGroup)

Systematic absences (extinctions)

TABLE 4.1. Limiting Conditions for Unit-Cell Type

Unit-cell type	Limiting conditions	Translations associated with the unit-cell type
<i>P</i>	None	None
<i>A</i>	<i>hkl</i> : $k + l = 2n$	$b/2 + c/2$
<i>B</i>	<i>hkl</i> : $h + l = 2n$	$a/2 + c/2$
<i>C</i>	<i>hkl</i> : $h + k = 2n$	$a/2 + b/2$
<i>I</i>	<i>hkl</i> : $h + k + l = 2n$	$a/2 + b/2 + c/2$
<i>F</i>	<i>hkl</i> : $h + k = 2n$	$a/2 + b/2$
	<i>hkl</i> : $k + l = 2n$	$b/2 + c/2$
	<i>hkl</i> : $(h + l = 2n)^a$	$a/2 + c/2$
	<i>hkl</i> : $-h + k + l = 3n$ (obv)	$a/3 + 2b/3 + 2c/3$
		$2a/3 + b/3 + c/3$
<i>R</i> _{hex} ^b	or	
	<i>hkl</i> : $h - k + l = 3n$ (rev)	$a/3 + 2b/3 + c/3$
		$2a/3 + b/3 + 2c/3$

^a This condition is not independent of the other two, as may be shown easily.

^b See page 70 and Table 2.3.

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

TABLE 4.2. Limiting Conditions for Screw Axes

Screw axis	Orientation	Limiting condition	Translation component
2 ₁	<i>a</i>	<i>h</i> 00: $h = 2n$	$a/2$
2 ₁	<i>b</i>	0 <i>k</i> 0: $k = 2n$	$b/2$
2 ₁	<i>c</i>	00 <i>l</i> : $l = 2n$	$c/2$
3 ₁ or 3 ₂	<i>c</i>	000 <i>l</i> : $l = 3n$	$c/3, 2c/3$
4 ₁ or 4 ₃	<i>c</i>	00 <i>l</i> : $l = 4n$	$c/4, 3c/4$
4 ₂	<i>c</i>	00 <i>l</i> : $l = 2n$	$2c/4(c/2)$
6 ₁ or 6 ₅	<i>c</i>	000 <i>l</i> : $l = 6n$	$c/6, 5c/6$
6 ₂ or 6 ₄	<i>c</i>	000 <i>l</i> : $l = 3n$	$2c/6, 4c/6(c/3, 2c/3)$
6 ₃	<i>c</i>	000 <i>l</i> : $l = 2n$	$3c/6(c/2)$

TABLE 4.3. Limiting Conditions for Glide Planes

Glide plane	Orientation	Limiting condition	Translation component
<i>a</i>	⊥ <i>b</i>	<i>h</i> 0 <i>l</i> : $h = 2n$	$a/2$
<i>a</i>	⊥ <i>c</i>	<i>h</i> <i>k</i> 0: $h = 2n$	$a/2$
<i>b</i>	⊥ <i>a</i>	0 <i>k</i> <i>l</i> : $k = 2n$	$b/2$
<i>b</i>	⊥ <i>c</i>	<i>h</i> <i>k</i> 0: $k = 2n$	$b/2$
<i>c</i>	⊥ <i>a</i>	0 <i>k</i> <i>l</i> : $l = 2n$	$c/2$
<i>c</i>	⊥ <i>b</i>	<i>h</i> 0 <i>l</i> : $l = 2n$	$c/2$
<i>n</i>	⊥ <i>a</i>	0 <i>k</i> <i>l</i> : $k + l = 2n$	$b/2 + c/2$
<i>n</i>	⊥ <i>b</i>	<i>h</i> 0 <i>l</i> : $h + l = 2n$	$a/2 + c/2$
<i>n</i>	⊥ <i>c</i>	<i>h</i> <i>k</i> 0: $h + k = 2n$	$a/2 + b/2$

Ab Initio Structure Solving

- 1) Needs a high quality sample already well characterized (composition, density, ...)
- 2) Carry out one or more *well adapted* diffraction experiments
Optimize the data quality (low background, low peak overlapping,...)
(x-rays and/or neutrons and/or electrons, choice of resolution, wavelength, *etc...*)
- 3) Find unit cell and index Bragg reflections (Dicvol, Treor, Ito, *etc...*). See www.ccp14.ac.uk/solution/indexing/. See 'Peter W. Stephens: *Indexing Powder Diffraction Data*' tutorial on youtube.
- 4) Obtain intensities of Bragg reflections and determine the space group
(LeBail technique + CheckGroup)
- 5) Find approximate structural model *ab initio* using Direct Methods (reciprocal space) or Heavy Atom (Patterson) Methods, or methods in direct space.

also use Difference Fourier analysis to locate missing atoms

Ab Initio Structure Solving by Direct Methods

- 1) It requires preliminary determination of the diffraction intensity of the individual reflections (**as accurately as possible**), which is done using profile fitting or full pattern decomposition methods (G. S. Pawley: J. Appl. Cryst. (1981), or A. Le Bail: Mat. Res. Bull. (1988)).
- 2) Any substantial overlapping of the reflections lowers the quality and makes the true solution less distinguishable.
- 3) The same programs that are used to solve single crystal structures are used to solve structure from the powder diffraction data, *e.g.*, SHELXS or EXPO

Ab Initio Structure Solving in Direct Space

Plausible structure models are randomly generated by moving atoms or groups of atoms in crystal coordinates within the pre-determined unit cell. The calculated diagram is compared to the observed one.

- **Trial-and-error approach**, that can be computer-intensive and can be guided by prior chemical or crystallographic knowledge and structural database mining (*e.g.*, search for isostructural or analogous compounds)
- In case of overlapping reflections, the direct space search, *e.g.*, global optimization method, becomes much more advantageous.
- Prior to simulations, eliminate experimental biases, *e.g.*, huge preferred orientation or absorption effects, and other factors that decrease intensity or increase broadening of the peaks.
- Avoid a priori impossible configurations (too small distances, unrealistic angles, *etc.*) *e.g.*, **apply “common” chemical sense, check the calculated density** ($Z \times M/V$)
- Avoid to get blocked in a local minimum

Ab Initio Structure Solving in Direct Space

Plausible structure models are randomly generated by moving atoms or groups of atoms in crystal coordinates within the pre-determined unit cell. The calculated diagram is compared to the observed one.

One possible method: **simulated annealing** (FOX, PowderSolve)

For each configuration, one calculates a *cost function*, based on a pseudo-potential energy to minimize, and on the agreement between experiment and model :

$$C = \alpha E_{pot} + (1-\alpha) R$$

E_{pot} may depend on the bond type (ionic, covalent, intermetallic...), Lennard-Jones, Coulomb, ...

$$R = 100 \sum_{2\theta} |I_{exp}(2\theta) - I_{calc}(2\theta)| / \sum_{2\theta} I_{exp}(2\theta)$$

One goes from config. C_i to C_{i+1} following a **Monte-Carlo** type algorithm:

If $C_{i+1} < C_i$, move accepted

If $C_{i+1} > C_i$, move accepted with probability : $exp[-(C_{i+1}-C_i)/K]$

$K \approx$ temperature, decreases slowly during search.

And more...

Complexity of the materials, and hence the data, increases...

Performance of the measuring methods increases...

Effectiveness of numerical data processing techniques increases...

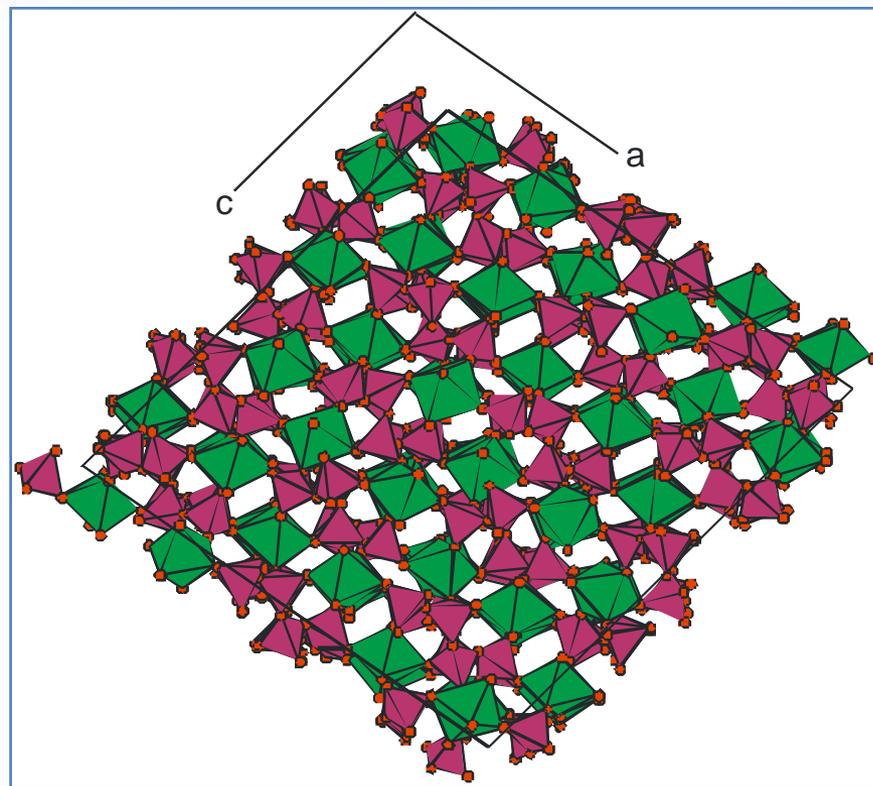
- Maximum Entropy (1996)
- Maximum Likelihood (1998)
- Genetic Algorithm (1999)
- Simulated Annealing (2000)
- Parallel Tempering (2002)
- Charge Flipping (2008) (truly ab initio; no information about atom types, composition, charge, *etc.* is required; starting symmetry can be kept as low as P1)

Structure Determination of $\text{Mo}_2\text{P}_4\text{O}_{15}$

One of the largest structures solved with TOPAS (simulated annealing)

Single crystal data
(Bruker AXS SMART 6000)

- SG: Pn (7)
- $a = 24.1134(6) \text{ \AA}$
- $b = 19.5324(5) \text{ \AA}$
- $c = 25.0854(6) \text{ \AA}$
- $\beta = 100.015(1)^\circ$
- $V = 4450.9 \text{ \AA}^3$
- 441 atoms in asymmetric unit



Lister et al., Chem. Commun., 2004, 2540

Ab Initio Structure Solving

- 1) Needs a high quality sample already well characterized (composition, density, ...)
- 2) Carry out one or more *well adapted* diffraction experiments
Optimize the data quality (low background, low peak overlapping,...)
(x-rays and/or neutrons and/or electrons, choice of resolution, wavelength, *etc...*)
- 3) Find unit cell and index Bragg reflections (Dicvol, Treor, Ito, *etc...*). See www.ccp14.ac.uk/solution/indexing/. See 'Peter W. Stephens: *Indexing Powder Diffraction Data*' tutorial on youtube.
- 4) Obtain intensities of Bragg reflections and determine the space group
(LeBail technique + CheckGroup)
- 5) Find approximate structural model *ab initio* using Direct Methods (reciprocal space) or Heavy Atom (Patterson) Methods, or methods in direct space.
- 6) Rietveld refinement and Fourier recycling to obtain a complete and accurate structure using Fullprof or GSAS

Pattern Refinement

- ✓ cell dimensions define the position of the peaks
- ✓ atomic parameters (the only parameters optimized in the single crystal refinement) determine the diffraction intensities
- ✓ microstructural parameters drive the shape of the diffraction peaks (*e.g.*, crystallite size, defects)
- ✓ Instrumental parameters affect the diffracted intensity (*e.g.*, preferred orientation and absorption) and the peak shape

Pattern Refinement

The method consists of refining a crystal (and/or magnetic) structure by minimizing the weighted squared difference between the observed y_i and the calculated y_{ci} intensities at all diffraction angles $1 \leq i \leq n$.

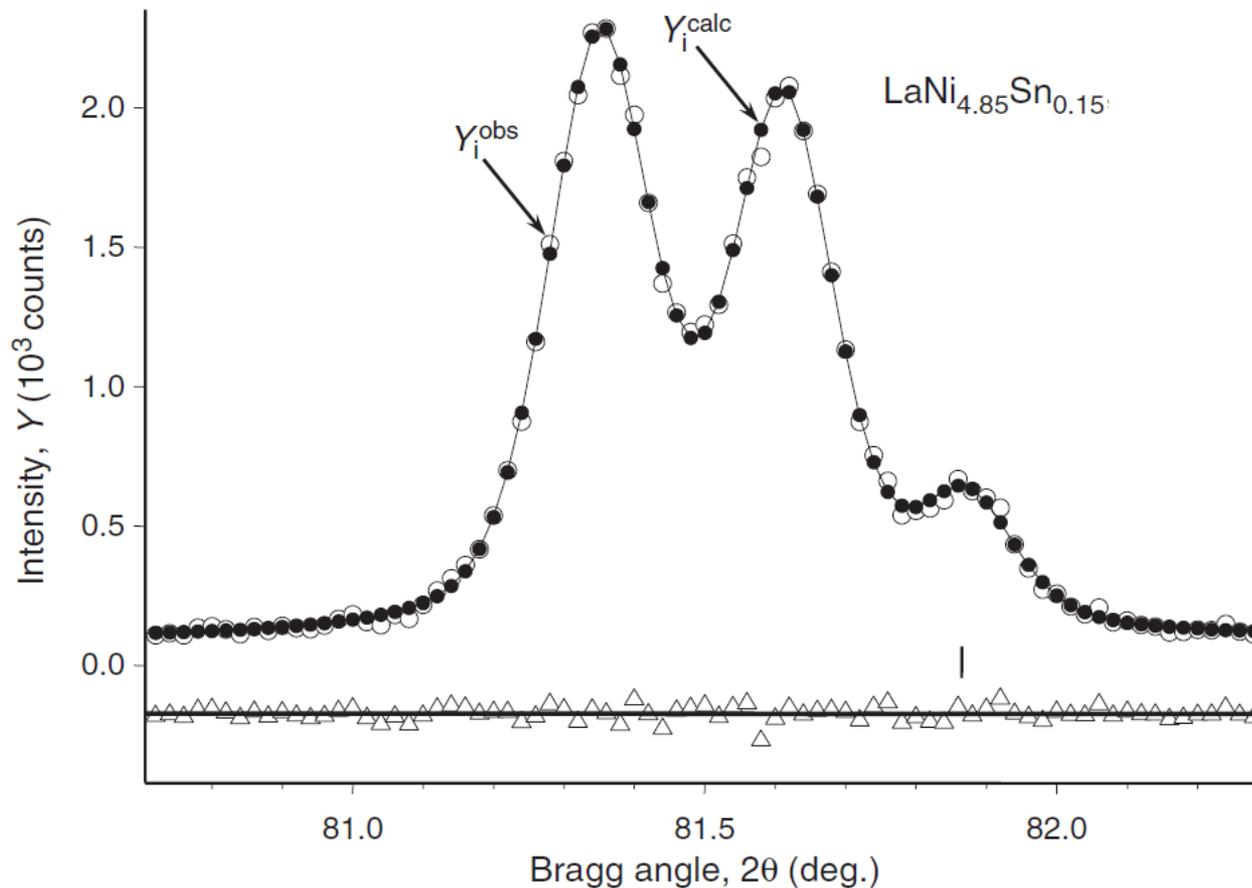
$$\chi^2 = \sum_{i=1}^n w_i \{y_i - y_{ci}(\beta)\}^2$$

$$w_i = \frac{1}{\sigma_i^2}$$

σ_i^2 : is the variance of the "observation" y_i

The model to calculate a powder diffraction pattern is:

$$y_{ci} = \sum I_h \Omega(T_i - T_h) + b_i \quad \begin{array}{l} T = 2\theta \\ \text{diffraction angle} \end{array}$$



Pattern Refinement

The model to calculate a powder diffraction pattern is:

$$y_{ci} = \sum_{\mathbf{h}} I_{\mathbf{h}} \Omega(T_i - T_{\mathbf{h}}) + b_i \quad \begin{array}{l} T = 2\theta \\ \text{diffraction angle} \end{array}$$

y_{ci} is the calculated diffraction intensity at the diffraction angle T_i

$I(h)$ is the diffracted intensity from a particular set of planes (h).

$I(h)$ is driven by the atomic motif *i.e.*, by the structure factor $F(h)$.

Several peaks $h - I(h)$, angle T_h - might contribute to the measured intensity at any particular T_i (diffraction) angle.

$I(h)$ is distributed over a range of T_i (diffraction) angles around T_h .

Pattern Refinement

The model to calculate a powder diffraction pattern is:

$$y_{ci} = \sum_{\mathbf{h}} I_{\mathbf{h}} \Omega(T_i - T_{\mathbf{h}}) + b_i \quad \begin{array}{l} T = 2\theta \\ \text{diffraction angle} \end{array}$$

$$\int_{-\infty}^{+\infty} \Omega(x) dx = 1$$

Profile function characterized by its full width at half maximum (FWHM= H) and *shape* parameters (η , m , ...)

$$\Omega(x) = g(x) \otimes f(x) = \text{instrumental} \otimes \text{intrinsic profile}$$

Pattern Refinement

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$$\Omega(x) = g(x) \otimes f(x) = \text{instrumental} \otimes \text{intrinsic profile}$$

- **Profile refinement**: pattern matching is constrained by peak positions (symmetry, unit cell) and peak profile (instrument, microstructure). Each $I(h)$ is not constrained by the atomic motif but is a free parameter.

Rietveld Refinement

The model to calculate a powder diffraction pattern is:

$$y_{ci} = \sum_{\mathbf{h}} I_{\mathbf{h}} \Omega(T_i - T_{\mathbf{h}}) + b_i \quad \begin{array}{l} T = 2\theta \\ \text{diffraction angle} \end{array}$$

$$\int_{-\infty}^{+\infty} \Omega(x) dx = 1$$

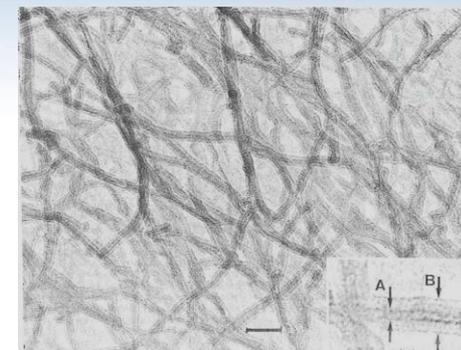
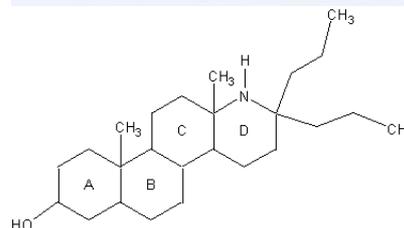
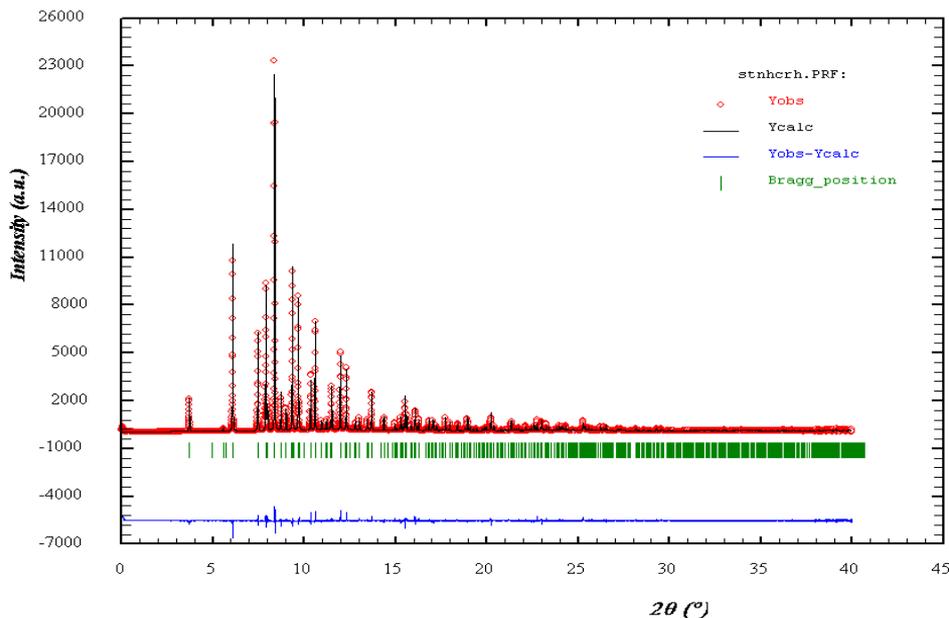
Profile function characterized by its full width at half maximum (FWHM= H) and *shape* parameters (η , m , ...)

$$\Omega(x) = g(x) \otimes f(x) = \text{instrumental} \otimes \text{intrinsic profile}$$

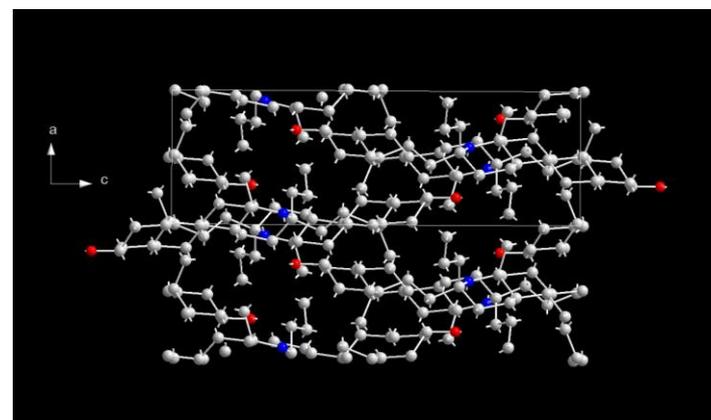
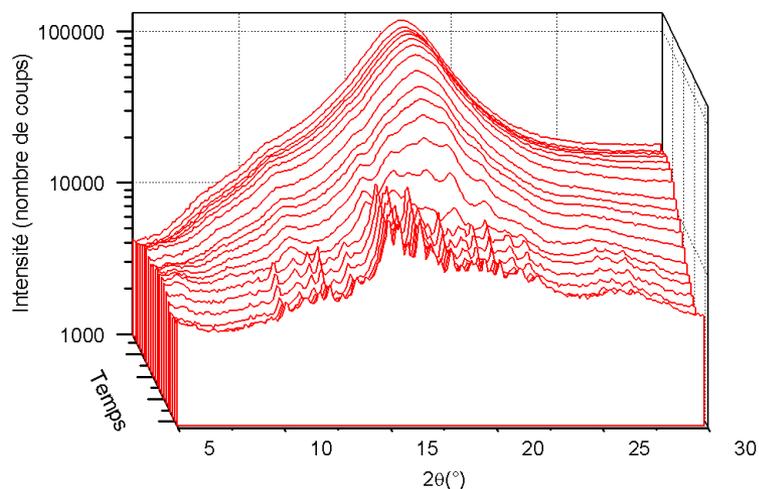
- **Rietveld refinement**: diffraction peak intensities are driven by the atomic model (atom coordinates within the unit cell). Each $I(h)$ is constrained by the atomic motif *i.e.*, $F(h)$

see for example "The Rietveld Method", R.A. Young (IUCr Monograph no.5) - 1994

Rietveld Structure refinement



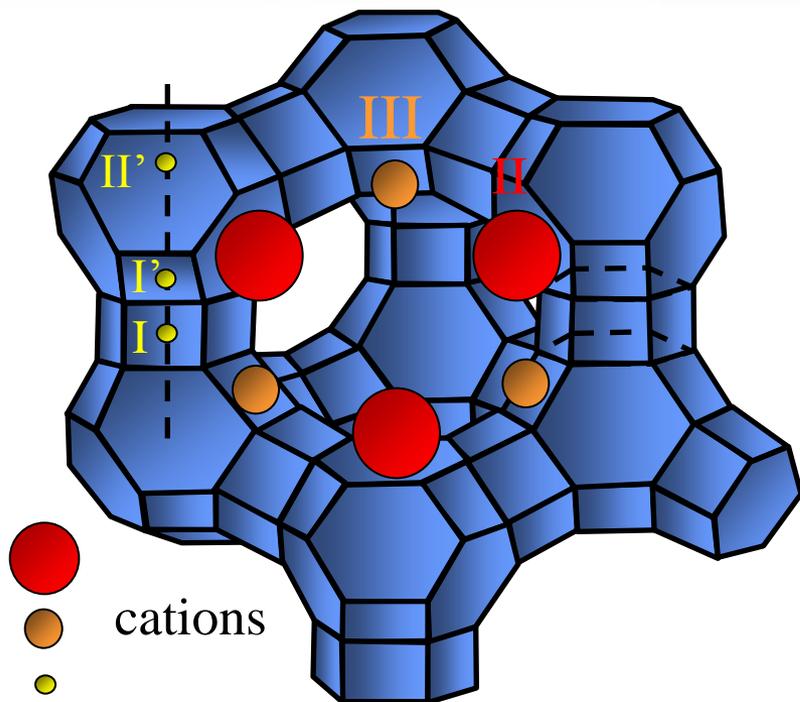
$a = 6,5358 \text{ \AA}$
 $b = 17,6040 \text{ \AA}$
 $c = 19,7070 \text{ \AA}$
 $P 2_1 2_1 2_1$
 $\chi^2 = 1,73$



Hydrogen linked network

Rietveld Structure refinement

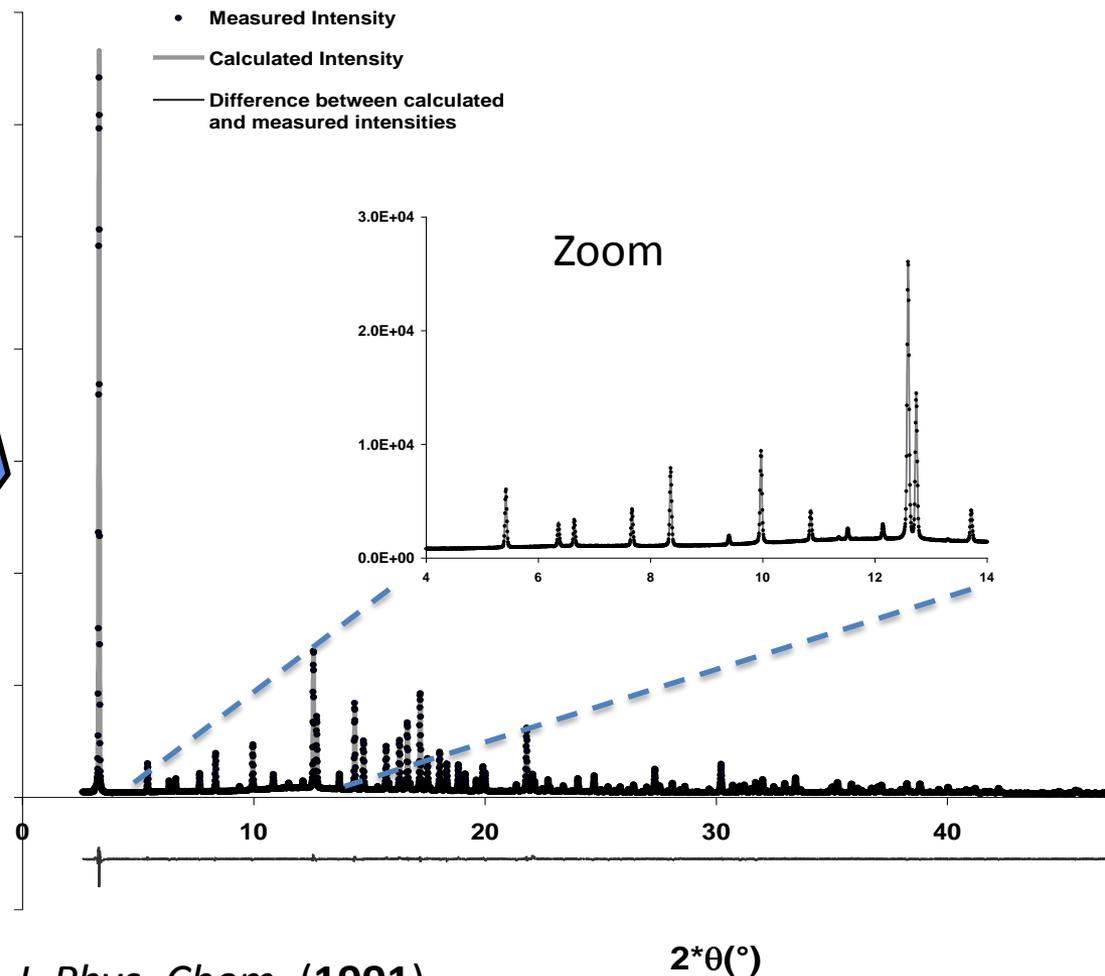
Cation sorption in a zeolite (filter, membrane, trap)



Formula: $\text{Si}_{110}\text{Al}_{84}\text{O}_{384}, n\text{H}_2\text{O}$

Space Group: $Fd\bar{3}m$

Unit cell parameter: 25 \AA



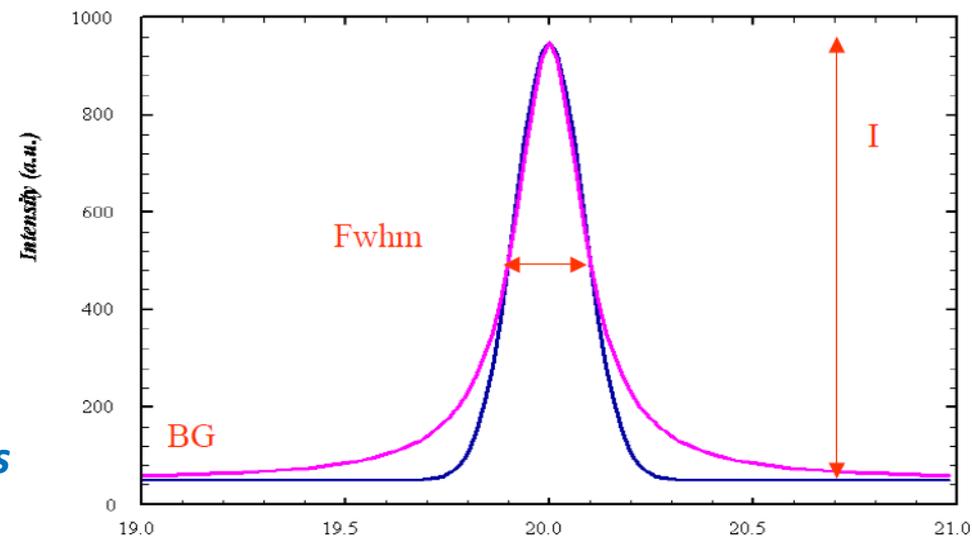
J. Phys. Chem. (1991)

$2\theta(^{\circ})$

Rietveld Refinement

- 1) Peak position fitting: **unit cell parameters** (thermal expansivity, macroscopic strain)
- 2) Intensity fitting of individual reflections to extract the structure factors: they contain the phase fractions (**quantitative analysis**) and the structural parameters (**crystal structure**: atomic positions)
- 3) Isotropic or anisotropic temperature or displacement factors
- 4) **Peak shape fitting** of individual from the perfect crystal
 - crystallite size
 - shape (needle, platelet,...)
 - microstrain

Peak broadens when crystallite size decreases



Rietveld Refinement

L.B. McCusker, et al, "*Rietveld Refinement Guidelines*", J. Appl. Cryst. (1999), 32, 36-50

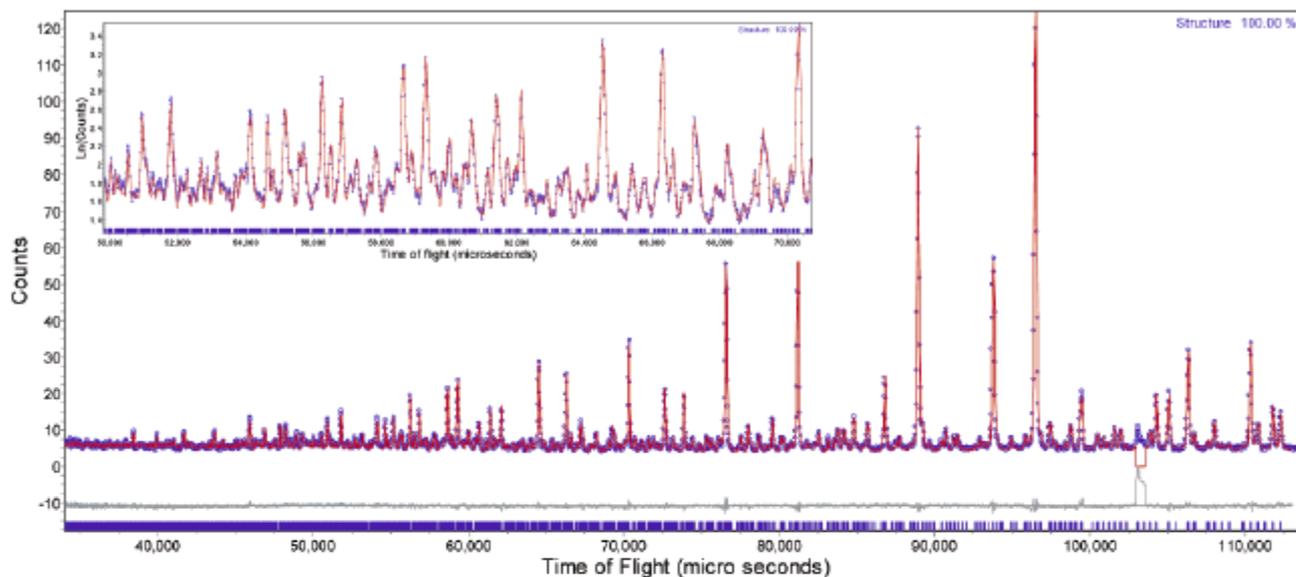
see Peter W. Stephens (SBU): '*Rietveld Refinement*':
<https://www.youtube.com/watch?v=YF5EHwMGJFc>

Technique played major role in every 'hot' material in solid-state chemistry in past 30 years, *e.g.*, high T_c , C60, giant magnetoresistance, batteries, fuel cells, *etc*

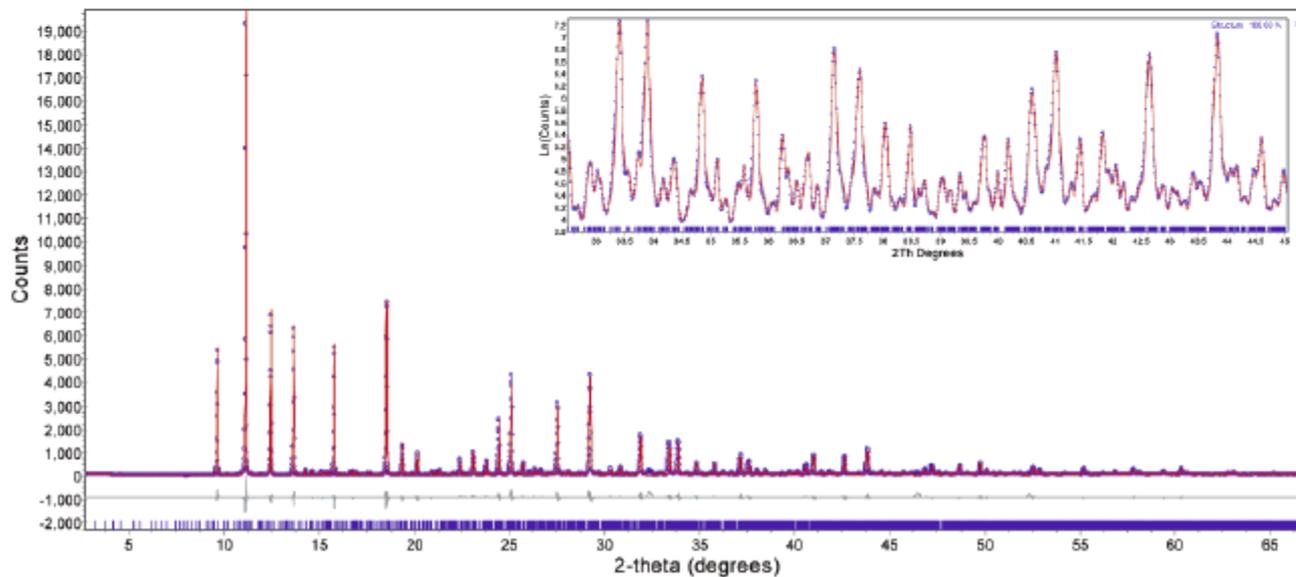
Programs: TOPAS, GSAS, FullPROF, ...

ZrP₂O₇ – a ‘peak-fest’

Stinton et al.



TOF neutron data
(HRPD, ISIS)
21195 reflections



Synchrotron data
(X7A, NSLS, 0.8Å)
22015 reflections

Courtesy of Pamela
Whitfield, Oak Ridge (TN)

Outline

1. Applications for Powder Diffraction
2. The basics of Powder Diffraction
3. Experimental Methods for collecting PD data
4. Crystal structure Determination
- 5. Peak Profile Analysis for Microstructural Studies**
- 6. Advanced PD techniques**

Peak Profile Fitting

- 3 contributions into each individual peak profile:
 1. Instrument
 2. Crystallite size
 3. Lattice strain

E. J. Mittemeijer, P. Scardi (eds.)

Diffraction Analysis of the Microstructure of Materials, Springer Verlag, Heidelberg 2004.

Peak Profile Fitting

- 3 contributions into each individual peak profile:
 1. Instrument
 2. Crystallite size
 3. Lattice strain

- XRD Crystallite Size using the Scherrer Formula
<http://www.d.umn.edu/~bhar0022/dpcalculator>

$$D = \frac{K\lambda}{\beta \cos \theta}$$

β is the FWHM of the Bragg peak after correction for instrumental broadening

The Scherrer formula applies if strain broadening is negligible.

Peak Profile Fitting

- Modeling the Gaussian and Lorentzian components of the Bragg profile function: provides microstructural parameters

$$H_G^2 = (U + (1 - \xi)^2 D_{ST}^2) \tan^2 \theta + V \tan \theta + W + \frac{I_G}{\cos^2 \theta}$$

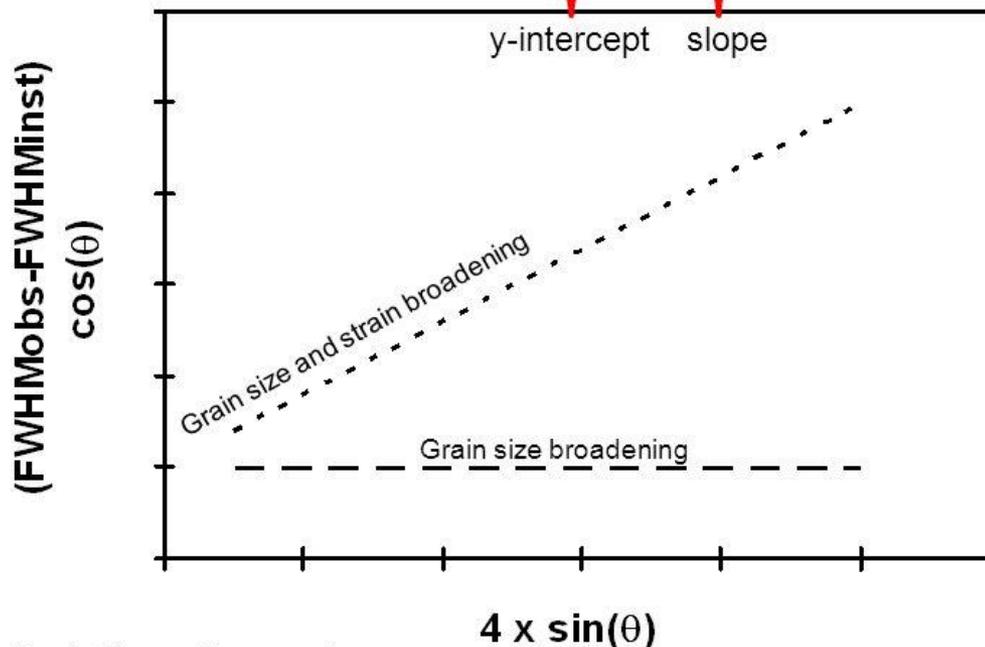
$$H_L = (X + \xi D_{ST}) \tan \theta + \frac{[Y + F(S_Z)]}{\cos \theta}$$

Microstructure Analysis from *Diffraction* Edited by R. L. Snyder, H. J. Bunge, and J. Fiala, International Union of Crystallography, 1999

Size and Strain in Reciprocal Space

Size and strain may be separated in favorable cases by measuring several peaks (*i.e.*, over a portion of the reciprocal space)

$$FWHM \times \cos(\theta) = \underbrace{\frac{K \times \lambda}{\text{Size}}}_{\text{y-intercept}} + \underbrace{\text{Strain} \times 4 \times \sin(\theta)}_{\text{slope}}$$



grain (actual particle of matter) \neq crystallite (coherent domain)

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6. **Advanced PD techniques**

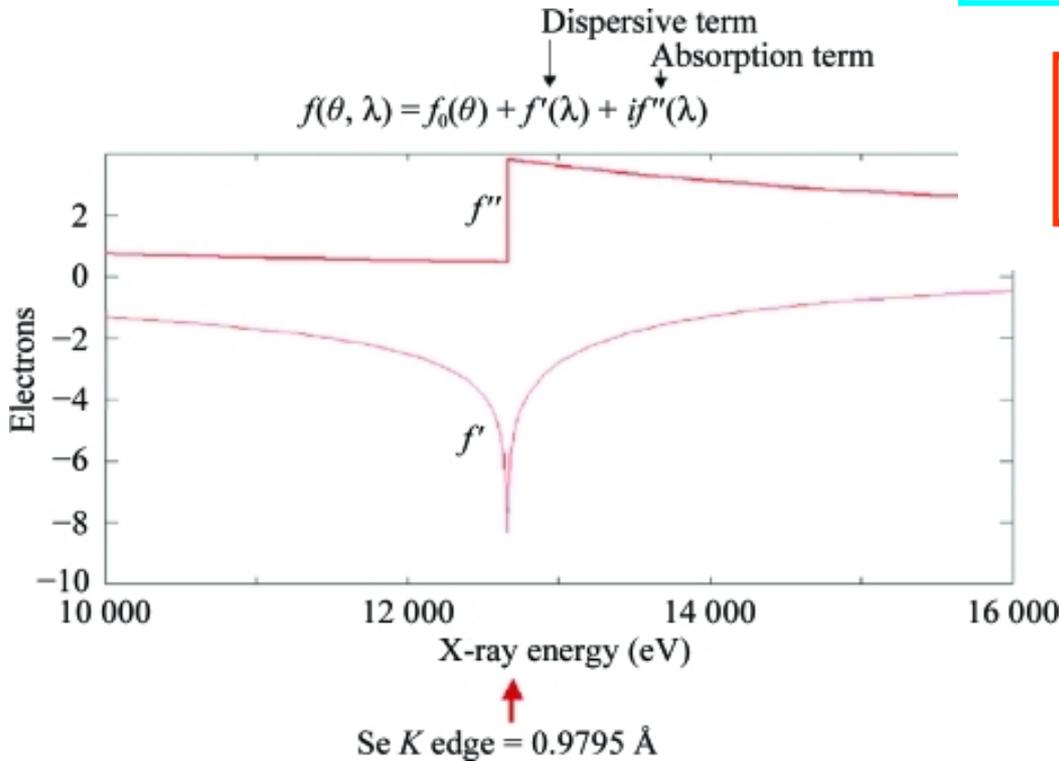
Anomalous Powder Diffraction

$$F_{hkl} = \sum_{i=1}^n f^i \cdot e^{2\pi i(hx_i + ky_i + lz_i)}$$

Electronic property of atom

two terms

Structural property (position)



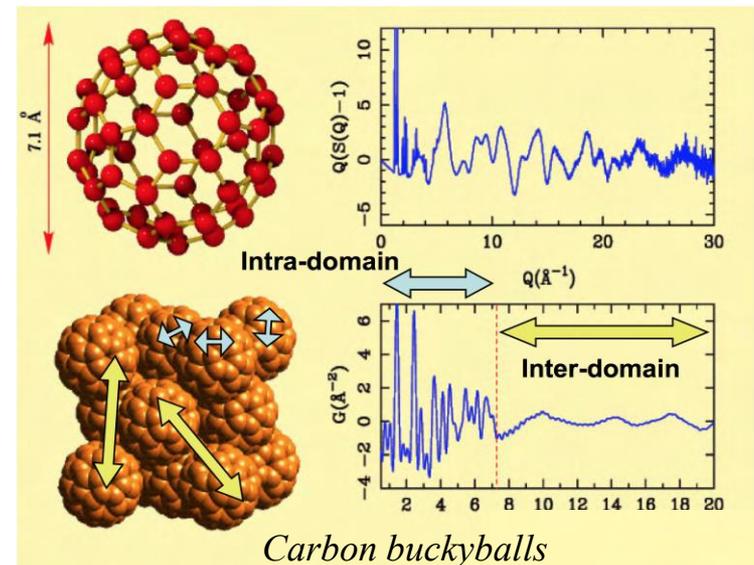
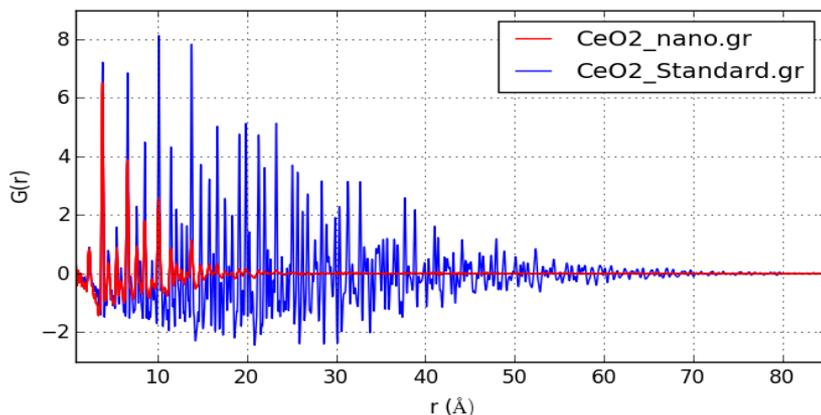
See Q. Shen's lecture

Total Scattering

- “Conventional” powder diffraction data analysis is insufficient or fails in particular for non crystalline materials (disorder, defects) and nanomaterials (short structural coherence length, broad Bragg peaks)
- PD patterns of complex materials often shows **diffuse scattering** smeared in the reciprocal under the Bragg peaks.

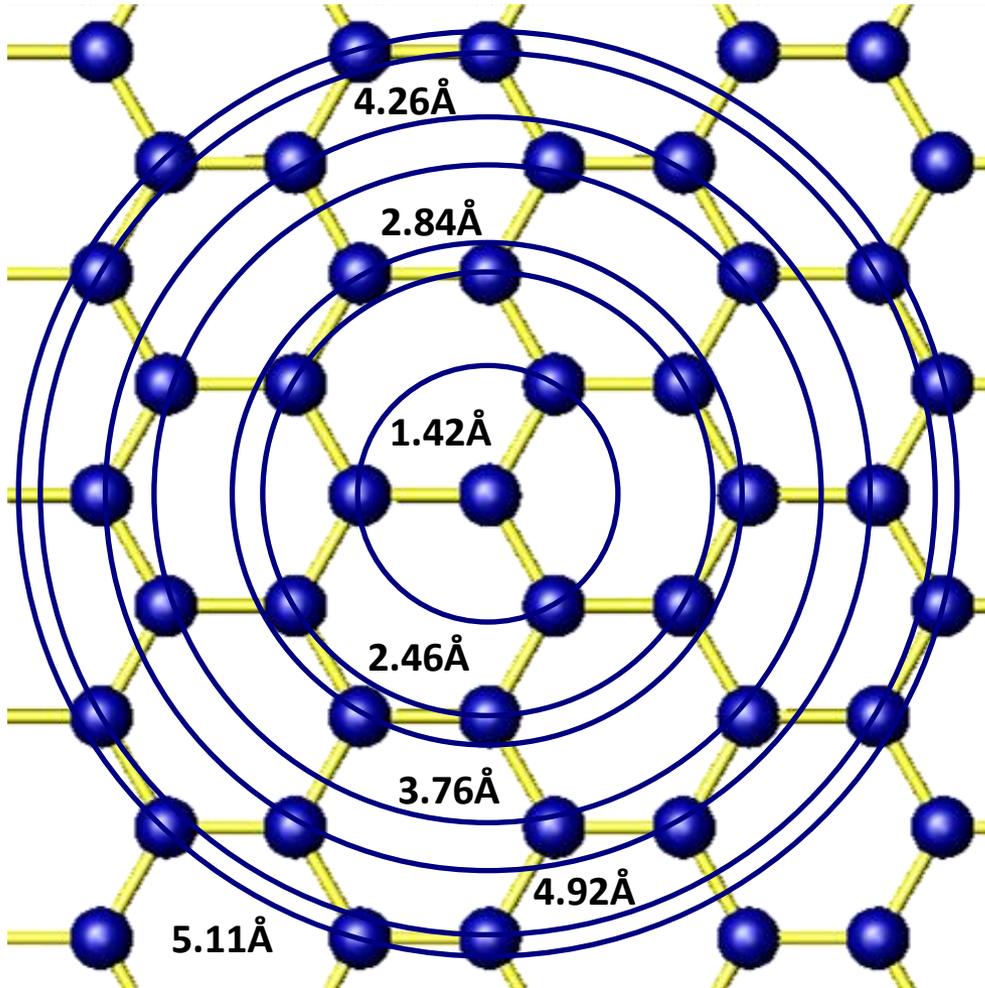
Total Scattering

- “Conventional” powder diffraction data analysis is insufficient or fails in particular for non crystalline materials (disorder, defects) and nanomaterials (short structural coherence length, broad Bragg peaks)
- PD patterns of complex materials often shows diffuse scattering smeared in the reciprocal under the Bragg peaks.
- FT of the total powder diffraction data (Bragg + diffuse) gives the PDF.



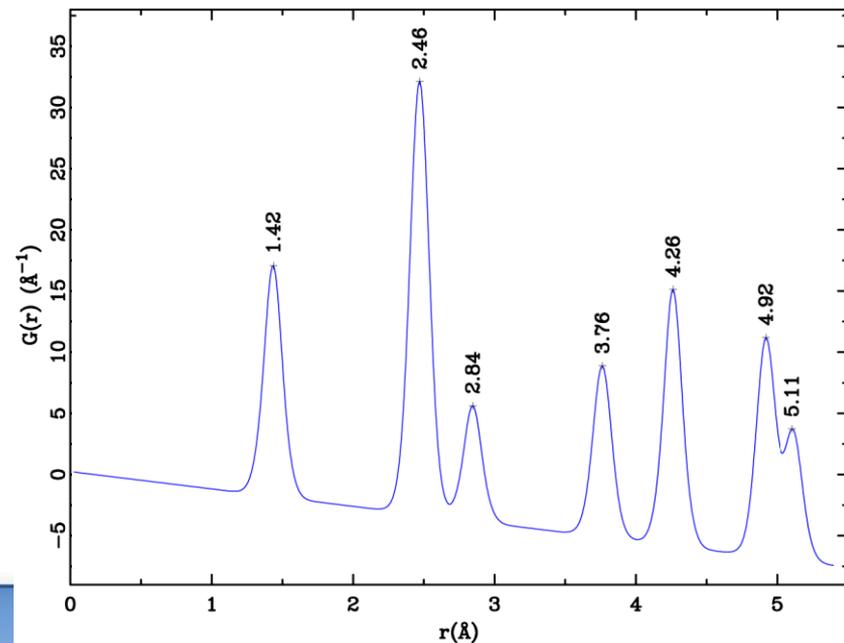
Schematic representation of the PDF

$$Q = 4\pi \sin(\theta)/\lambda$$

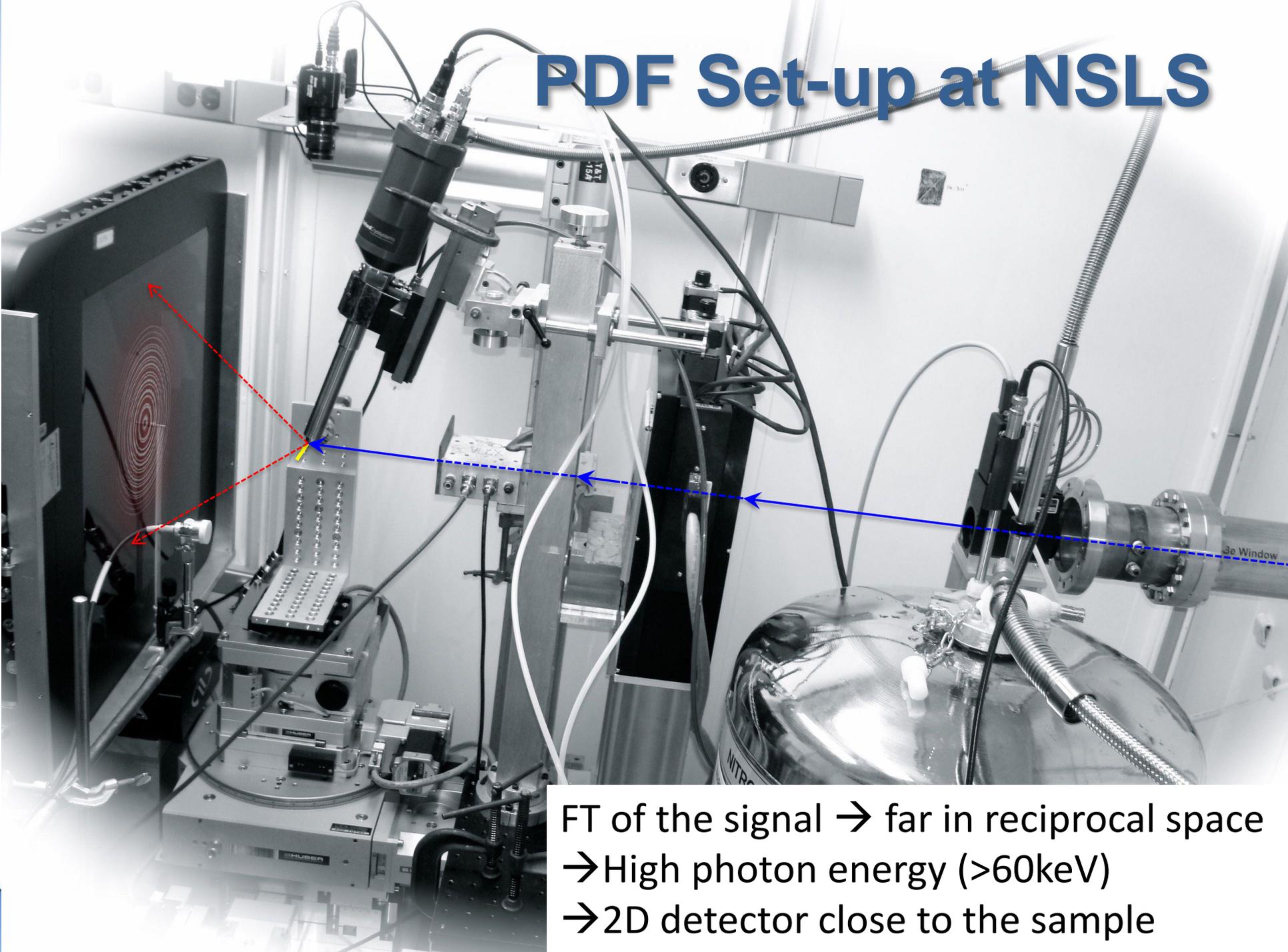


Pair distribution function (PDF) gives the probability of finding a neighbor atom at a distance “r” from a given atom.

$$G(r) = (2/\pi) \int_{Q_{\min}}^{Q_{\max}} Q[S(Q) - 1] \sin(Qr) dQ$$



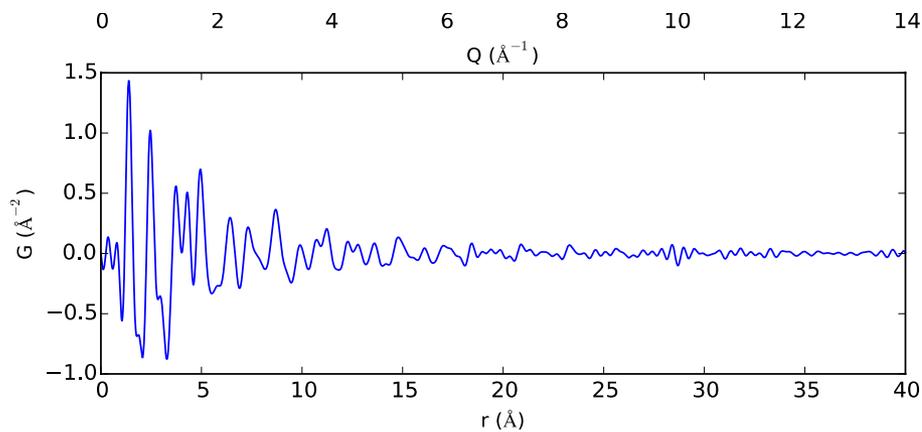
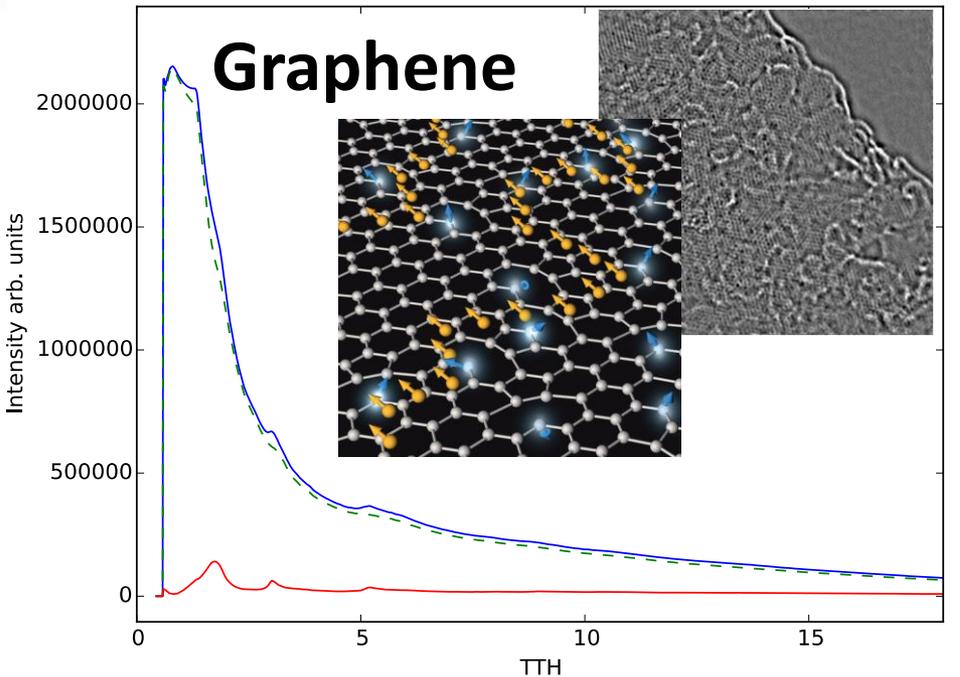
PDF Set-up at NSLS



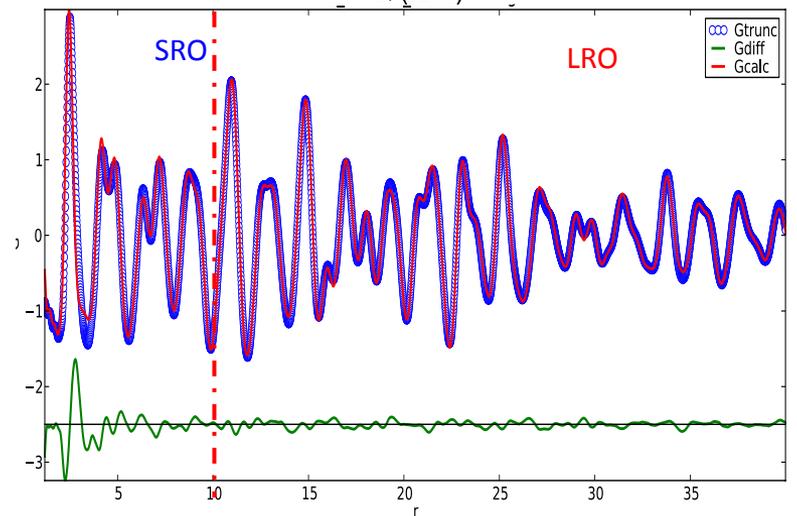
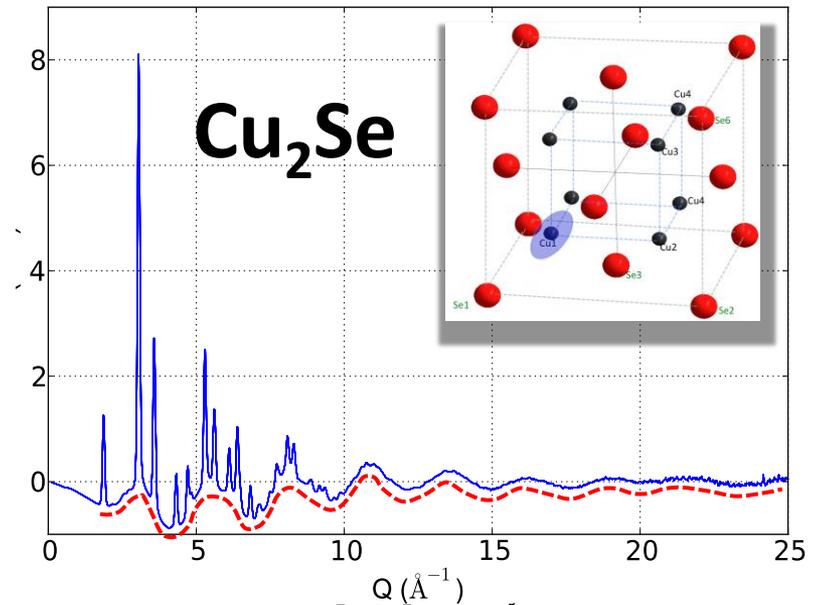
FT of the signal \rightarrow far in reciprocal space
 \rightarrow High photon energy ($>60\text{keV}$)
 \rightarrow 2D detector close to the sample

PDF Analysis Examples

Graphene



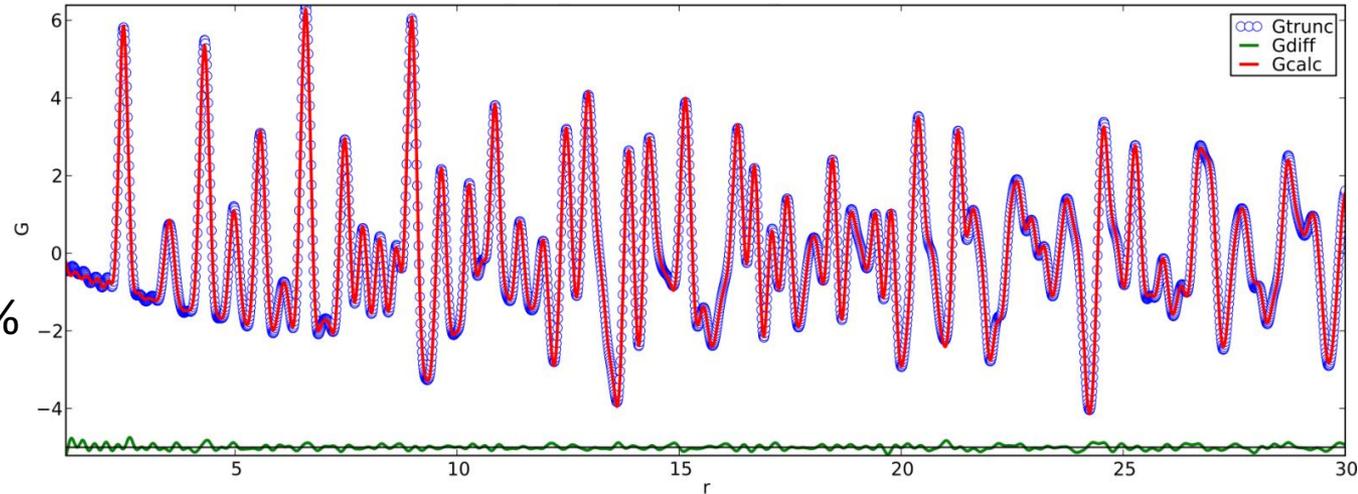
Cu_2Se



PDF Benchmarking NSLS-II (Columbia, BNL)

65.7 keV (300 μ m)²
1 min @ 25mA

Measured (blue) and calculated (red) Pair Distribution Functions. Green curve shows the difference.

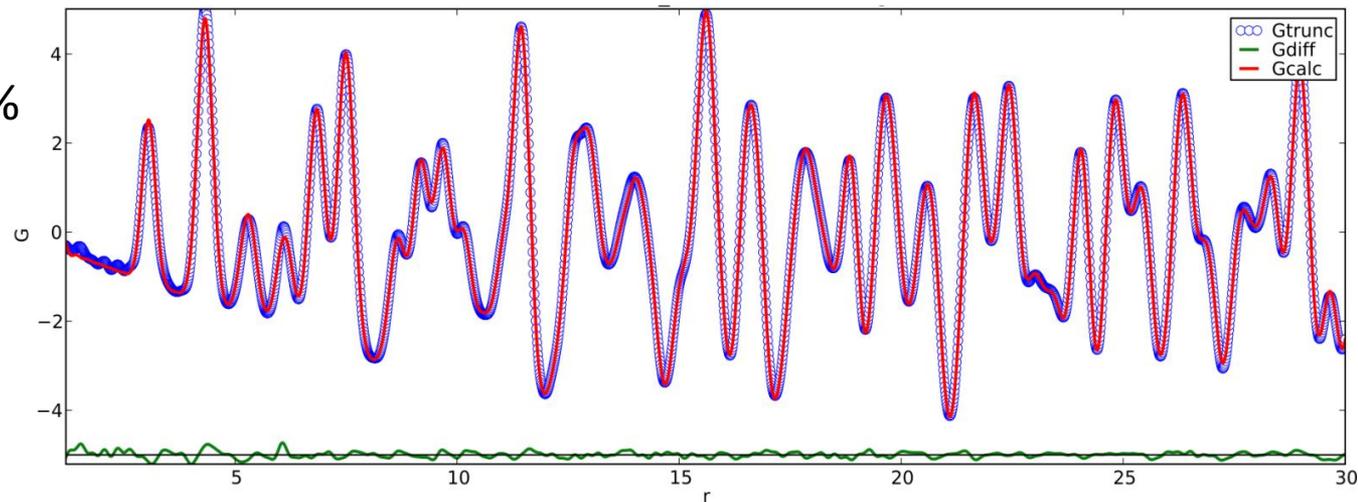


Ni Standard

$Q_{\text{max}} = 25 \text{ \AA}^{-1}$, $R_w = 3.6\%$

PbSe

$Q_{\text{max}} = 25 \text{ \AA}^{-1}$, $R_w = 3.8\%$



Total Scattering

- “Conventional” **Powder Diffraction** is applied to **crystalline materials** (**long range order**) with limited disorder. PD only looks at the Bragg peaks, background is treated as a baseline that is subtracted.
- **Pair Distribution Function** extends the PD technique over to **non-crystalline materials**, materials that contain a high degree of disorder and defects, and nanomaterials (**short range order**). PDF treats the signal that is contained both in the Bragg peaks and in the background.

‘Underneath the Bragg Peaks: Structural Analysis of Complex Materials’ (Pergamon Materials Series) 2nd Edition, by. Takeshi Egami and Simon J.L. Billinge.

Total Crystallography

Applies to samples that contain too many crystallites for “conventional” single crystal diffraction and too few for powder diffraction (randomization is insufficient).

This is a recurrent issue as probing beams get smaller and smaller. Software is being developed (see FABLE project) to manage this case.

TIME TO CONCLUDE...

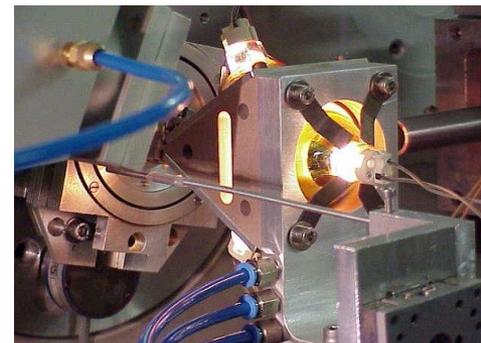
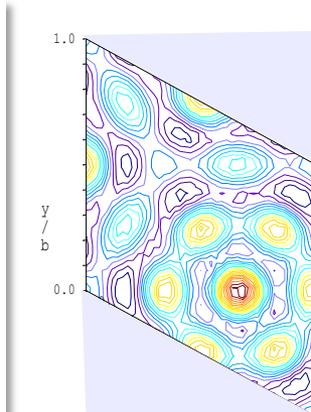
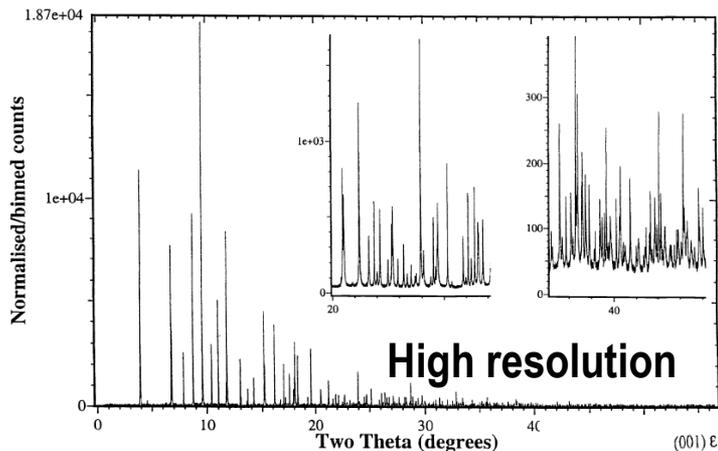
My hope is that you will leave with a sense of why or how powder XRD might be applicable to your research.

Where is the information in powder data?

1. **Peak positions** *(easy to measure, interpret)*
 - *Lattice parameters determination*
 - *Thermal expansivity or compressibility*
 - *Macro-stress*
 2. **Peak widths and shapes** *(easy to measure / hard to interpret)*
 - *Microstructure (crystallite size, stacking faults, etc)*
 3. **Peak intensities** *(challenging to measure, interpret)*
 - *Identification of fully occupied sites*
 - *Quantification of vacancies / mixed sites*
 - *Accurate DPs / ADPs*
 - *Structure solution*
 4. **Between the Bragg peaks (PDF)** *(even more challenging)*
 - *Short range order*
- All problems are affected by quality of data
 - Ask yourself how many parameters contribute to intensity of peaks
 - A light atom may contribute 1% of total peak intensity (small vs. noise)
 - Data quality \sim number of independent peaks, and esds in their intensities

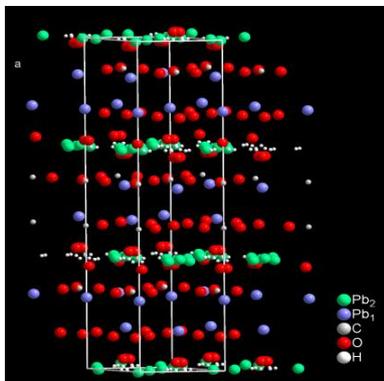
Summary of the Scientific Scope

Powder Diffraction addresses a wide range of topics from Fundamental (*incl.* metallurgy, physics, chemistry, biology *etc.*) to Applied Research → controlling complexity (real devices, heterogeneous)

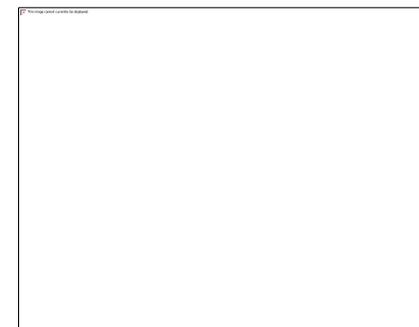
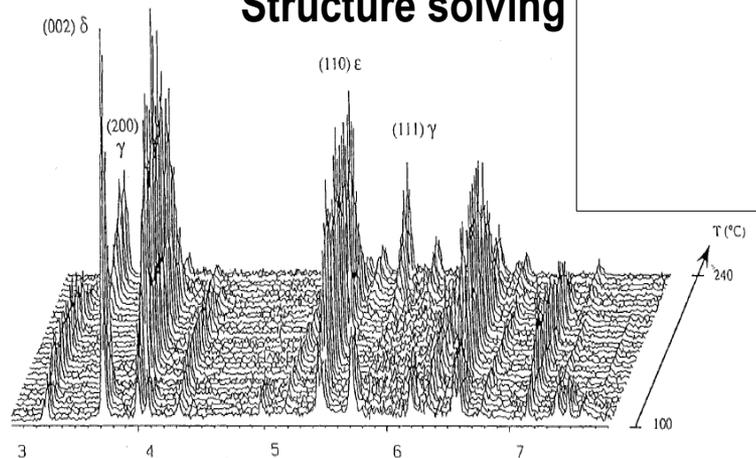


Extreme environments (T, P, gas, B, E)

Crystallography



Structure solving



In situ
Operando

Kinetics / time-resolved
High throughput

Current Trends in Powder Diffraction

Complex materials:

- Photovoltaics with improved efficiency – *Nanoparticles in the light collecting layer*
- High energy density batteries – *Electrodes, Electrolytes*
- Fuel cells – *Electrodes, Electrolytes*
- Catalysts – *Higher-order oxides, graphitic carbon, nanoparticles*
- Hydrogen storage – *MOF's*
- Carbon sequestration – *Functionalized mesoporous materials*
- Strongly correlated materials – *Materials doped with impurities*

Complexity: *Defects, Surfaces, Interfaces, Nano and multi-scale structures, heterogeneities, nanoscale disorder and surface relaxation etc.*

Properties in relation with LRO (average) and SRO (deviations)

Current Trends in Powder Diffraction

- **Real materials** \neq lab prepared (synthetic)
- **Real conditions** (*in situ*, operando) \neq room T, P
- **Real time** \neq static observation

Powder Diffraction Resources

- Commission on Powder Diffraction: <http://www.iucr.org/iucr-top/comm/cpd/>
- <http://11bm.xray.aps.anl.gov/resources.html> Instructional Resources, Software and Tools, Textbooks & Journal Articles, PD Beamlines.
- http://www.aps.anl.gov/Xray_Science_Division/Powder_Diffraction_Crystallography/ This page references a series of demos and lectures that introduce Rietveld analysis techniques, incl. Flash movie presentations.
- Modern Methods in Rietveld Refinement and Structural Analysis https://sites.google.com/a/stonybrook.edu/mmrrsa_2016/home
- <http://www.ccp14.ac.uk/mirror/mirror.htm> Collaborative Computational Project Number 14 Tutorials and Program Repository.
- <http://www.lightsources.org/> news and science highlights from each facility, as well as photos and videos, education and outreach resources, a calendar of conferences and events
- <https://www.bnl.gov/ps/beamlines/beamline.php?r=28-ID-2> XPD at NSLS-II

Powder Diffraction Software Packages

Structure Refinement Software (Rietveld, Le Bail etc)

- GSAS-EXPGUI
- FullProf
- JANA
- TOPAS Professional
- TOPAS Academic
- MAUD
- BGMN

Structure Solution from Powder Diffraction Software

- FOX/ObjCryst++
- EXPO2009
- DIFFRACplus (TOPAS)
- Superflip (JANA)

Crystal structure drawing & visualization software

- DRAWxtl
- ATOMS
- VESTA
- CrystalMaker
- Diamond
- Mercury

And many more...

International Tables for Crystallography

- Home page = <http://it.iucr.org/>
- A. Space-group symmetry
Symmetry relations between space groups
- B. Reciprocal space
- C. Mathematical, physical and chemical tables
- D. Physical properties of crystals
- E. Subperiodic groups
- F. Crystallography of biological macromolecules
- G. Definition and exchange of crystallographic data
- H. Powder Diffraction (new: 50 chapters, 800 pages)
 1. Introduction to the principles of powder diffraction.
 2. Instrumentation for laboratory X-ray studies, synchrotron, neutron and electron diffraction, 2D diffraction, and special environments (temperature, pressure, magnetic fields, reaction cells).
 3. Different methodologies used in powder diffraction.
 4. Structure determination and validation.
 5. Defects, texture and microstructure: stress and strain, grain size and thin films.
 6. Review of available software.
 7. Applications to: macromolecules, zeolites, mining, ceramics, cement, forensic science, archaeology and pharmaceuticals.
- I. X-ray Absorption Spectroscopy

Acknowledgments

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- Pamela Whitfield (Oak Ridge National Laboratory, TN)
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Thank you for your attention