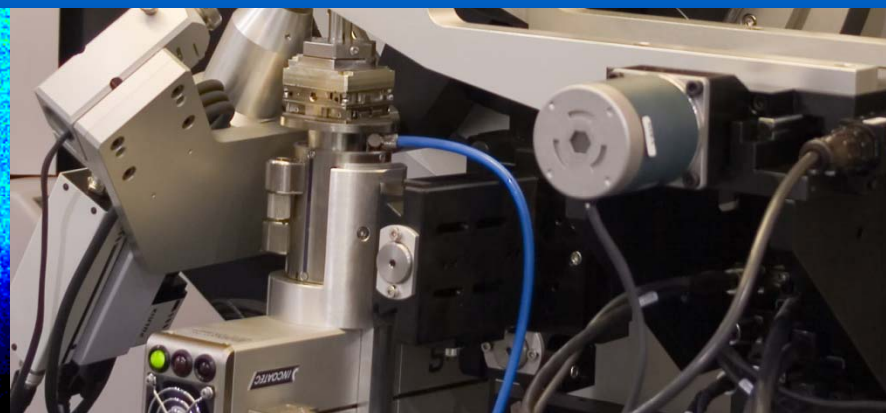
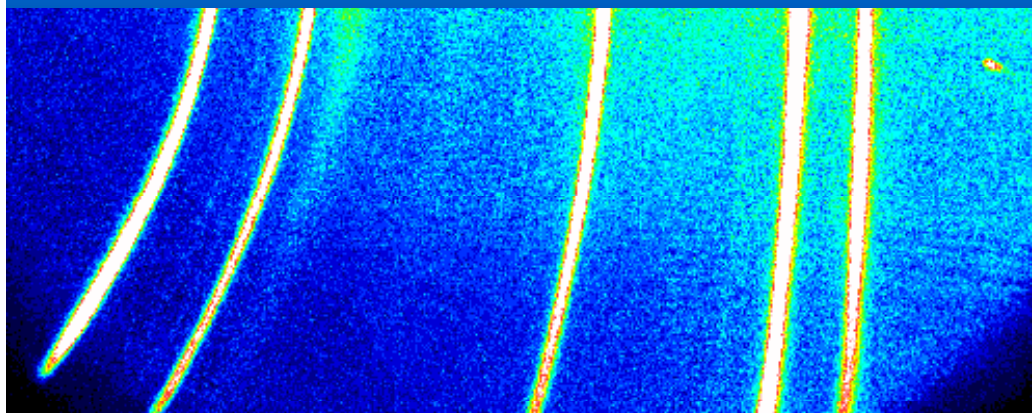


Geometry, Polarization and Absorption Corrections for Two-dimensional X-ray Diffraction

Bob He Bruker AXS, Inc.



This document was presented at PPXRD - Pharmaceutical Powder X-ray Diffraction Symposium

Sponsored by The International Centre for Diffraction Data

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ICDD Website - www.icdd.com

XRD²: What do we know about two-dimensional XRD?



Most recognized features of XRD² for pharmaceutical:

- High speed: 10^2 higher than XRD with a point detector
- Reliable info: Integration in γ (ring) direction
- Micro scale sample: Point beam and 2D pattern

Most concerns with XRD²:

- Resolution and geometry defocusing
- Relative intensity is different from Bragg-Brentano
- Higher instrument cost (but much higher productivity)

This presentation interprets the differences and suggests the best use of XRD² systems

X-ray Applications for typical pharmaceutical samples



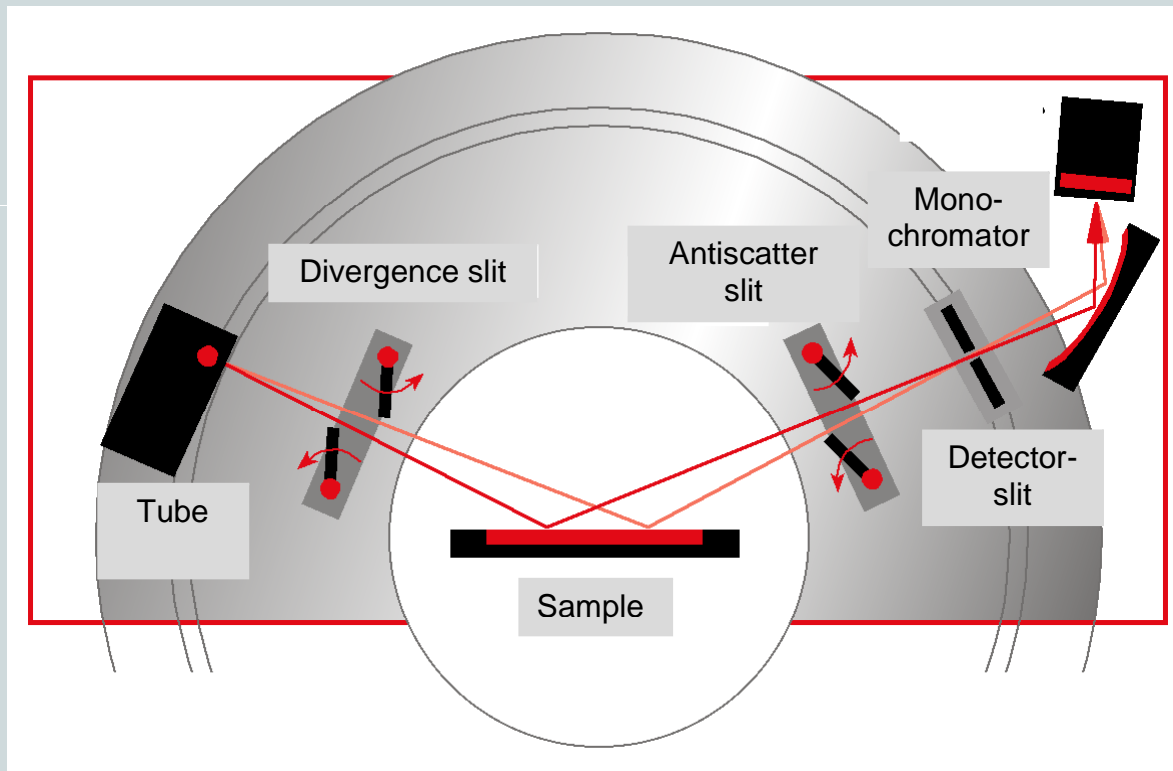
XRD & XRD ²	Single Crystal	Several Grains	Powder	Finished Product	Solutions
Qualitative Phase ID	✓Φ⊕	✓Φ⊕	✓Φ	✓Φ	✓Φ
Quantitative Rietveld analysis			✓		
Quantitative analysis with standards		✓	✓	✓	
X-ray movie, Non-Ambient	✓Φ	✓Φ	✓Φ	✓Φ	✓Φ
Structure solution, Indexing	✓Φ		✓		
Microdiffraction/ Mapping		✓Φ⊕	✓Φ⊕	✓Φ⊕	
Shape analysis			✓Φ	✓Φ	✓Φ
HTS	✓Φ⊕	✓Φ⊕	✓Φ⊕		
Grain-Size det.		✓Φ	✓Φ		
%Crystallinity		✓Φ⊕	✓Φ⊕	✓Φ⊕	✓Φ⊕

✓ - can be performed by either XRD or XRD²

Φ - better with XRD²

⊕ - accept performance and accurate results only with XRD²

Bragg-Brentano-Geometry



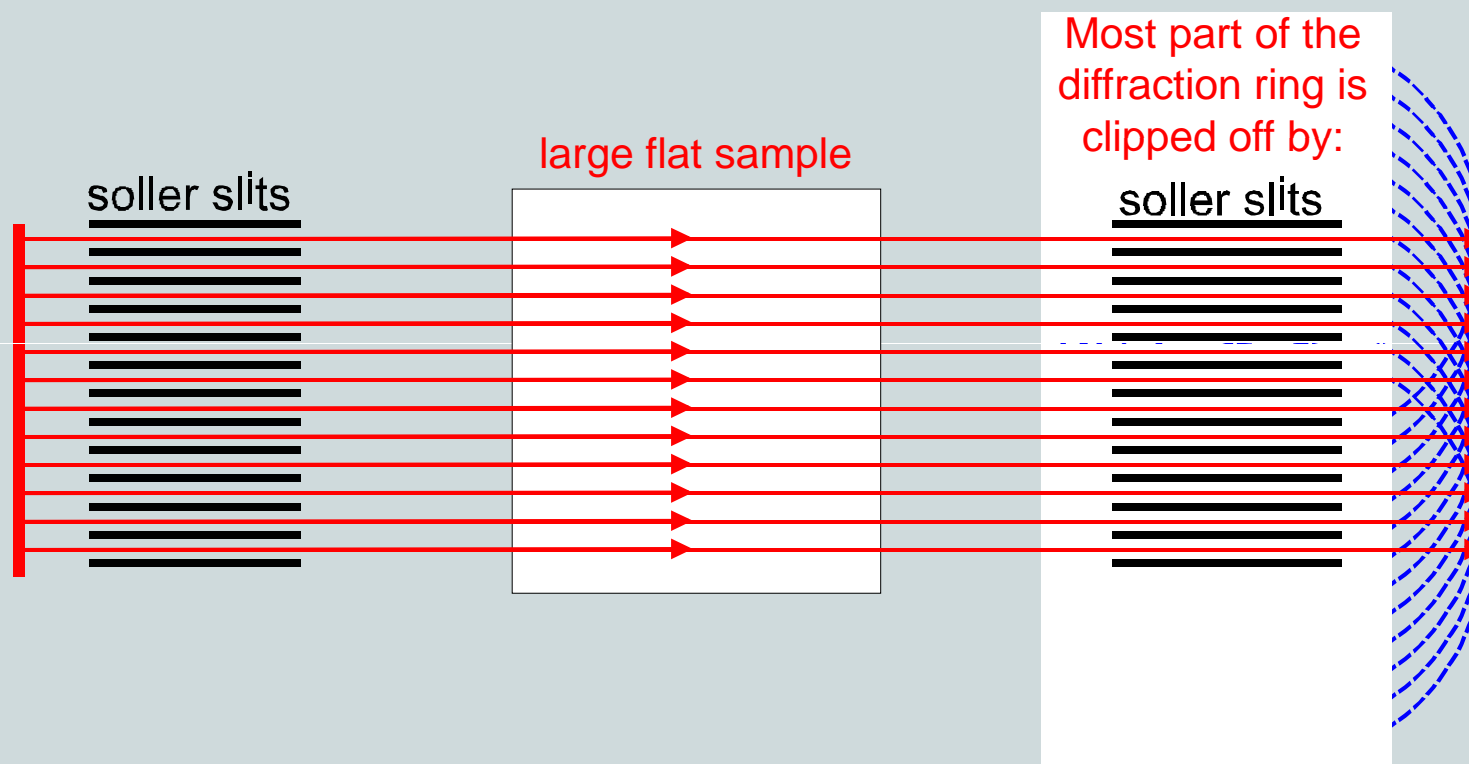
Advantages

- Resolution
- Less expensive

Disadvantages

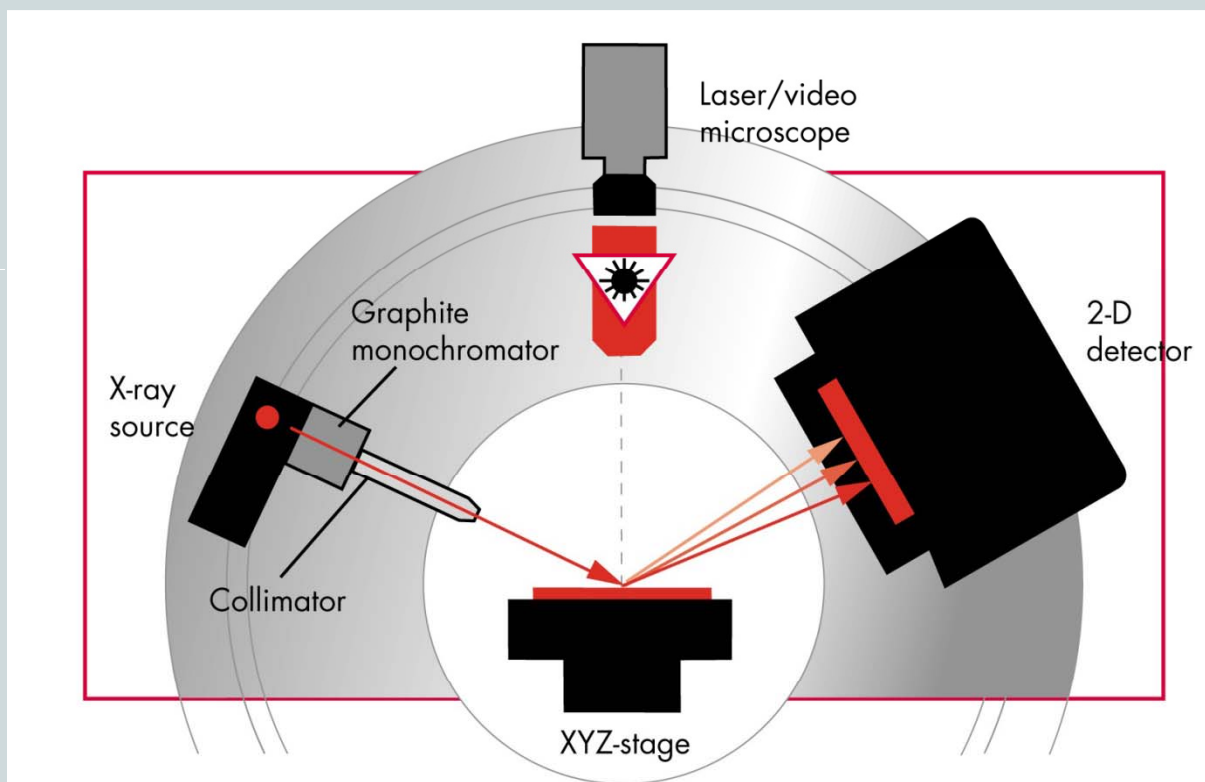
- Requires flat sample surface
- Requires bulky powder sample
- Slow

Soller slits are used to control axial divergence



- In Bragg-Brentano geometry, the line focus beam can be considered as a superposition of point beams.
- All in parallel with the diffractometer plane and the same geometry condition separated by soller slit foils.

XRD²: Two-dimensional X-ray Diffraction



Advantages

- High speed
- Micro scale sample
- Virtual oscillation (large grain size & preferred orientation)

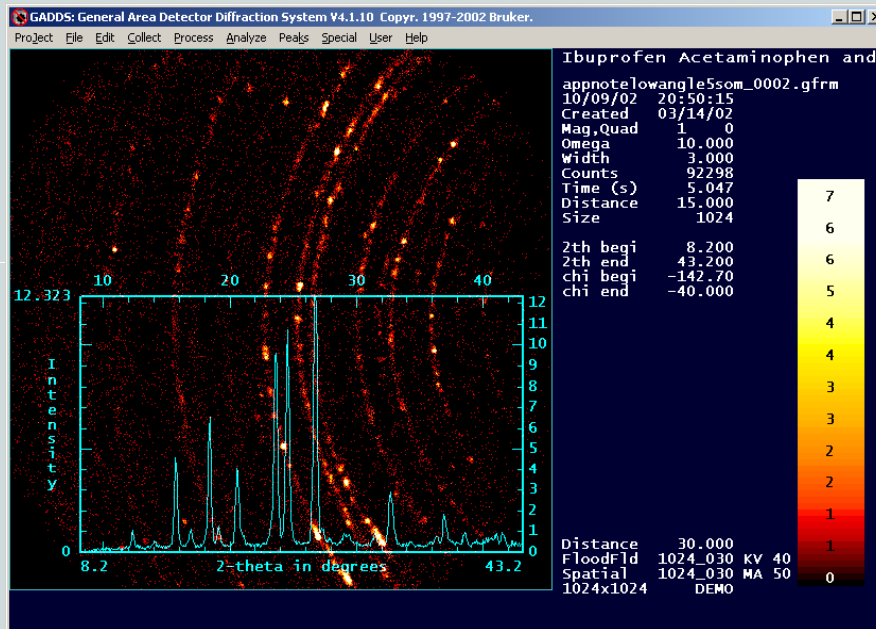
Disadvantages

- Resolution is limited by the detector PSF
- Defocusing at low angle in reflection
- Expensive

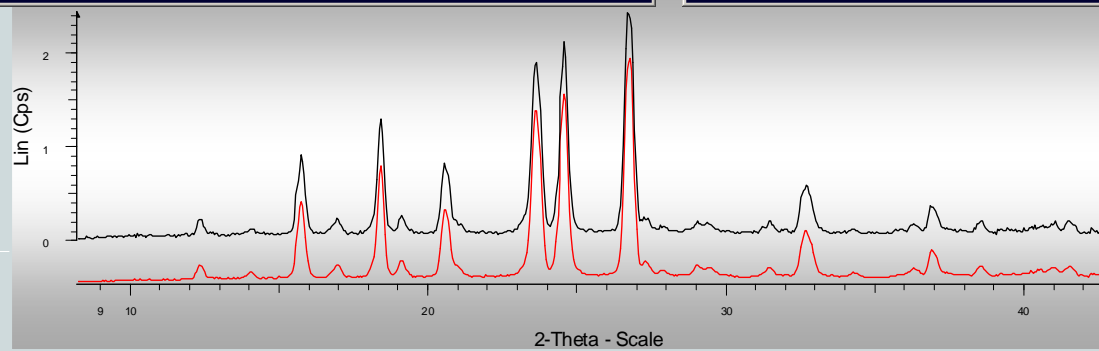
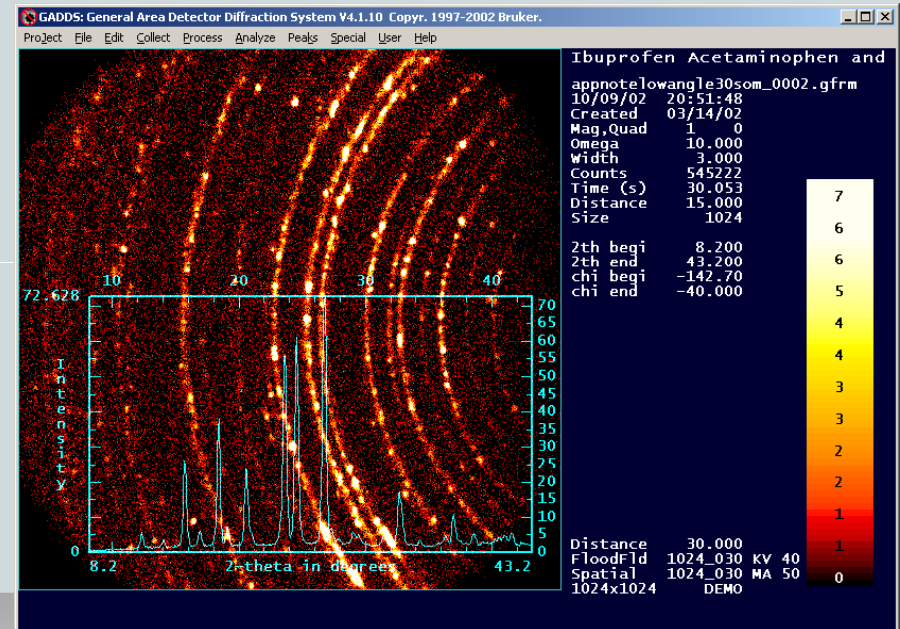
XRD²: Data Collection:

Acetaminophen powder

5 second data collection



30 second data collection

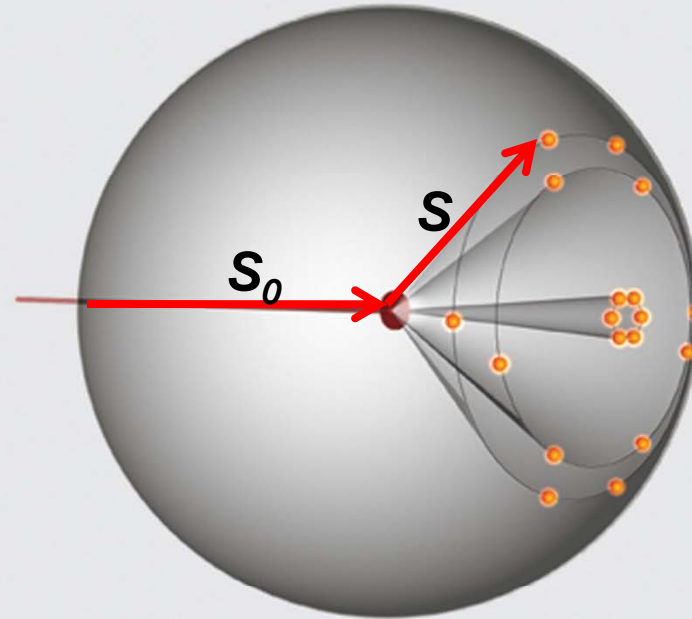
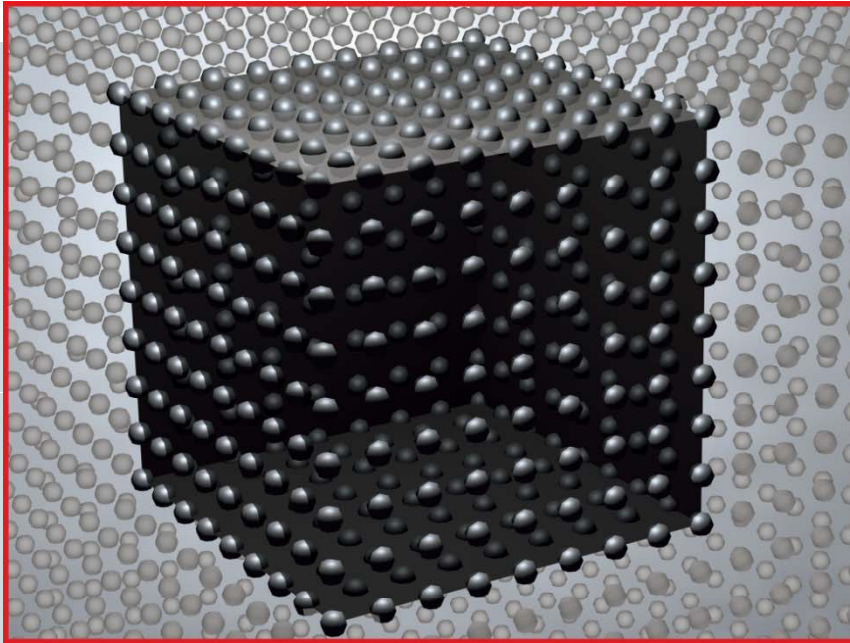




XRD²: Diffraction vector approach

Applications	Vector approaches
Phase Identification:	Polarization and absorption correction
Texture Analysis:	Orientation mapping angles; Data collection strategy (scheme)
Stress Measurement:	Fundamental equation derived by second order tensor transformation; Data collection strategy (scheme)
Crystal Size Analysis:	Equations for the effective volume calculation at both reflection and transmission modes.

XRD² & Single Crystals

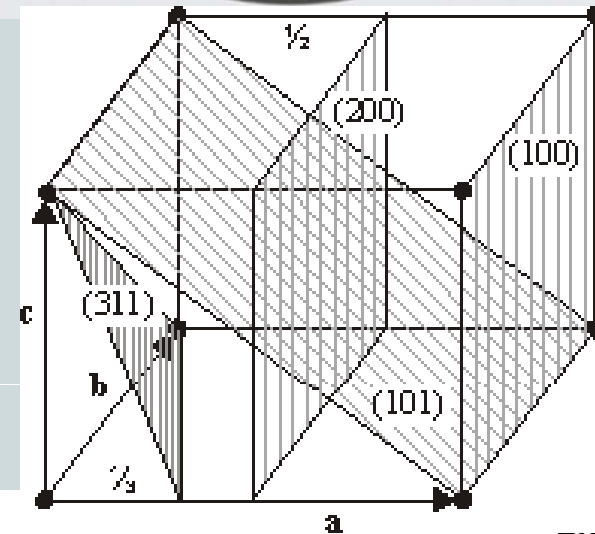


Laue equation

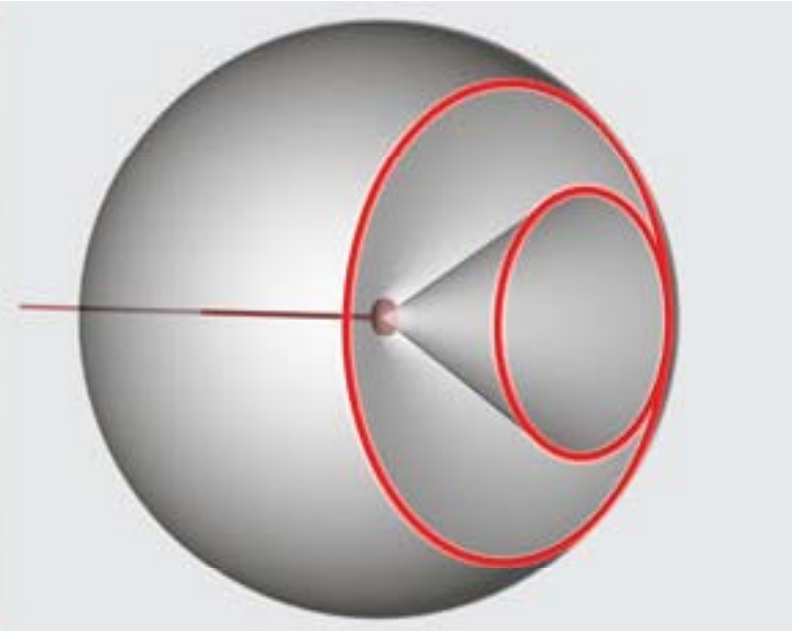
$$\mathbf{a} \cdot (\mathbf{s} - \mathbf{s}_0) = h\lambda$$

$$\mathbf{b} \cdot (\mathbf{s} - \mathbf{s}_0) = k\lambda$$

$$\mathbf{c} \cdot (\mathbf{s} - \mathbf{s}_0) = l\lambda$$

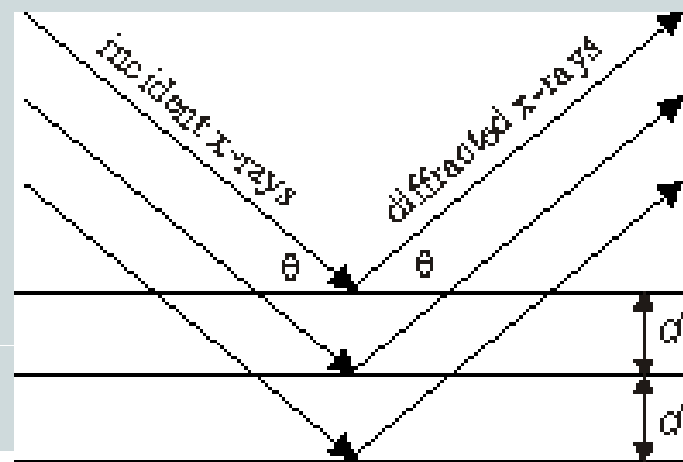


XRD² & Powders

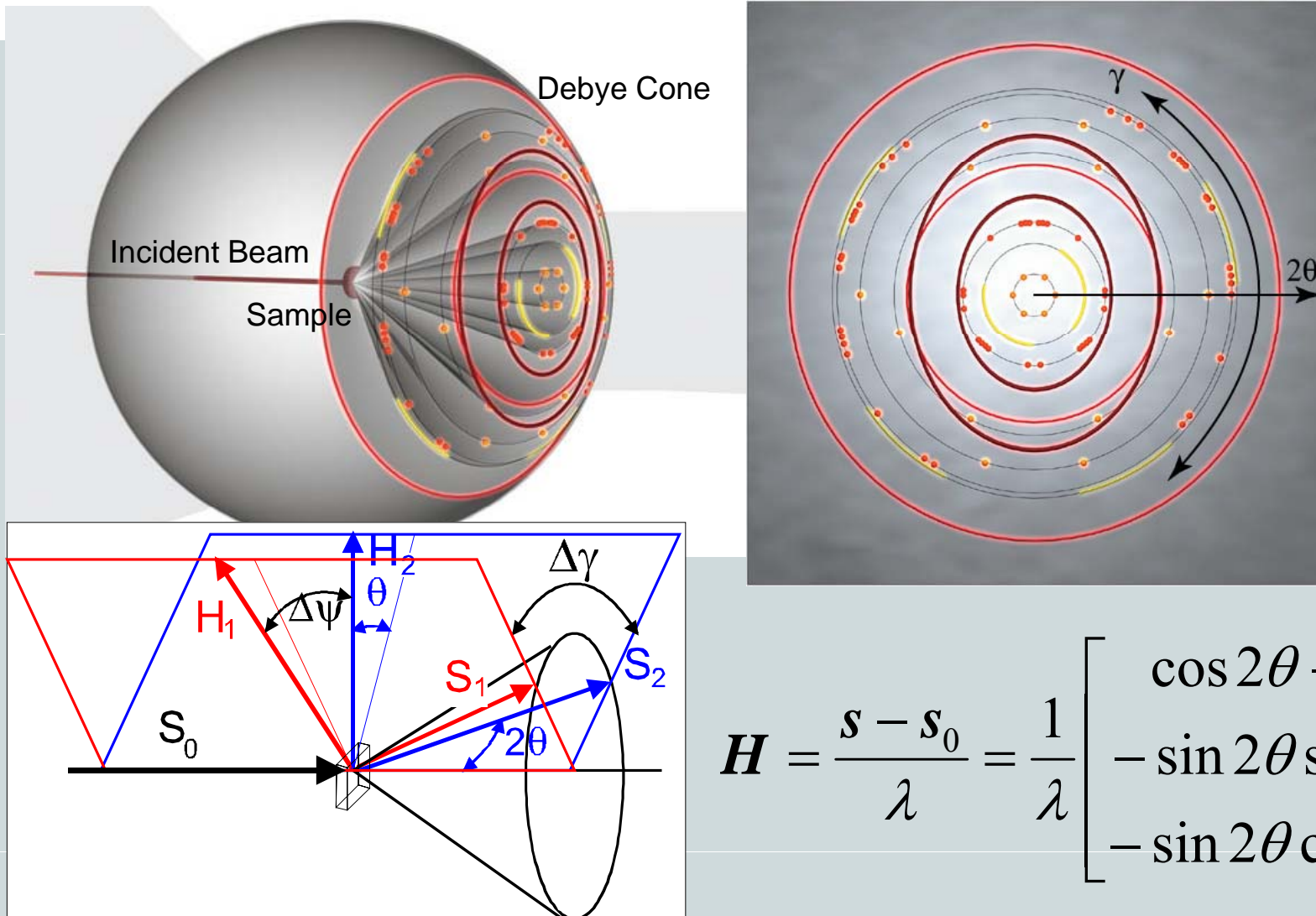


Bragg law

$$n\lambda = 2d \sin \theta$$

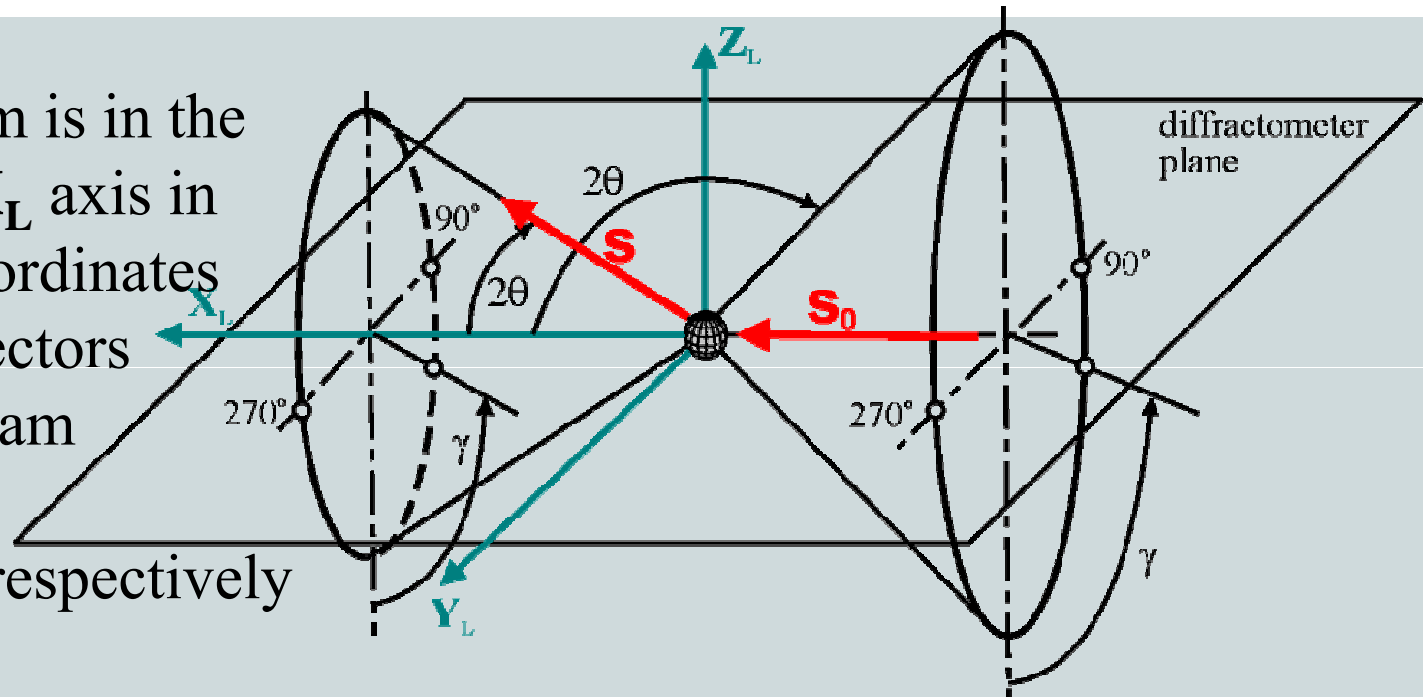


XRD²: Diffraction pattern with both γ and 2θ information



XRD²: Diffraction Space & Laboratory coordinators

The incident beam is in the direction of the \mathbf{X}_L axis in the laboratory coordinates so that the unit vectors of the incident beam and diffracted beams are given respectively by:



$$\mathbf{S}_0 = \begin{bmatrix} S_{0x} \\ S_{0y} \\ S_{0z} \end{bmatrix} = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} \quad \mathbf{S} = \begin{bmatrix} S_x \\ S_y \\ S_z \end{bmatrix} = \begin{bmatrix} \cos 2\theta \\ -\sin 2\theta \sin \gamma \\ -\sin 2\theta \cos \gamma \end{bmatrix}$$

XRD²: Diffraction Vector & Unit Diffraction Vector

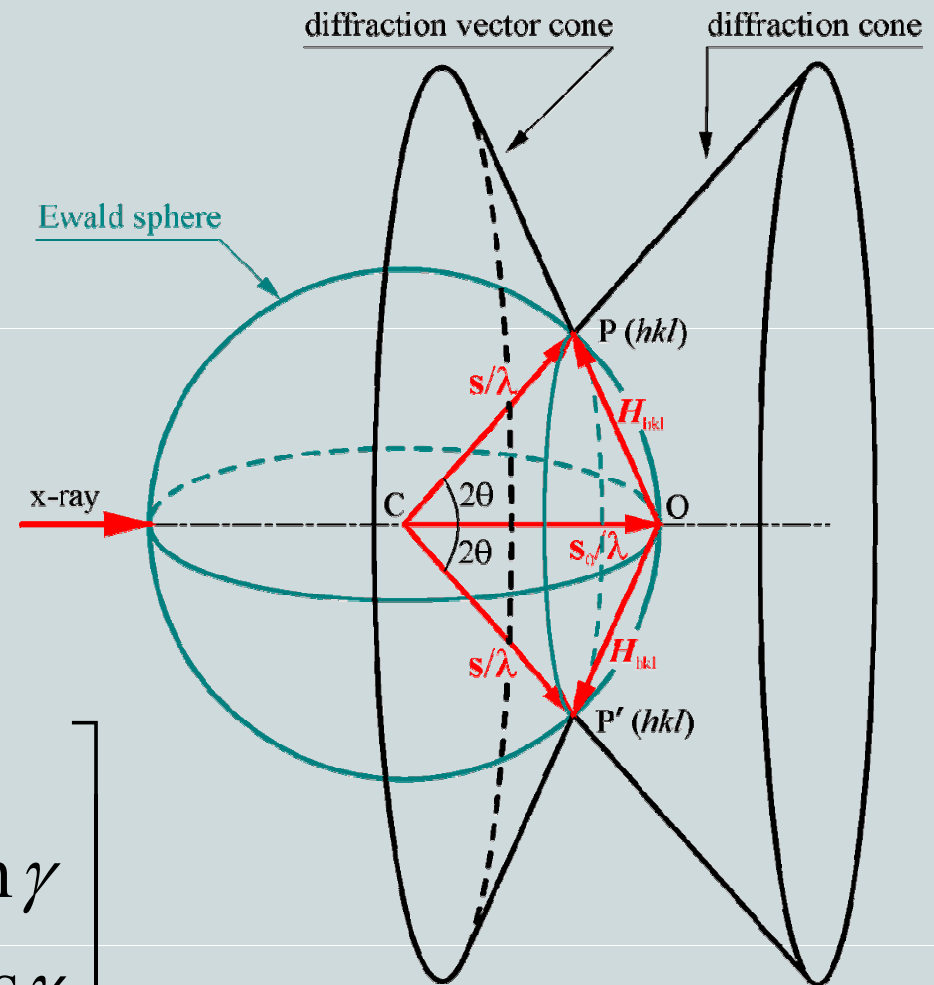


The diffraction vector is given in laboratory coordinates by

$$\mathbf{H} = \frac{\mathbf{s} - \mathbf{s}_0}{\lambda} = \frac{1}{\lambda} \begin{bmatrix} s_x - s_{0x} \\ s_y - s_{0y} \\ s_z - s_{0z} \end{bmatrix} = \frac{1}{\lambda} \begin{bmatrix} \cos 2\theta - 1 \\ -\sin 2\theta \sin \gamma \\ -\sin 2\theta \cos \gamma \end{bmatrix}$$

The direction of each diffraction vector can be represented by its unit vector given by:

$$\mathbf{h}_L = \frac{\mathbf{H}}{|\mathbf{H}|} = \begin{bmatrix} h_x \\ h_y \\ h_z \end{bmatrix} = \begin{bmatrix} -\sin \theta \\ -\cos \theta \sin \gamma \\ -\cos \theta \cos \gamma \end{bmatrix}$$

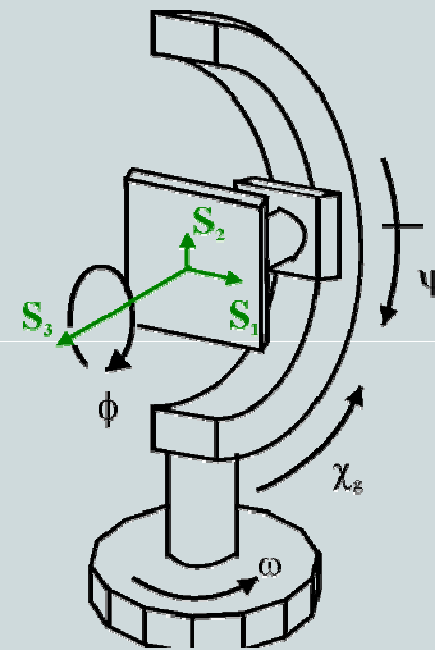
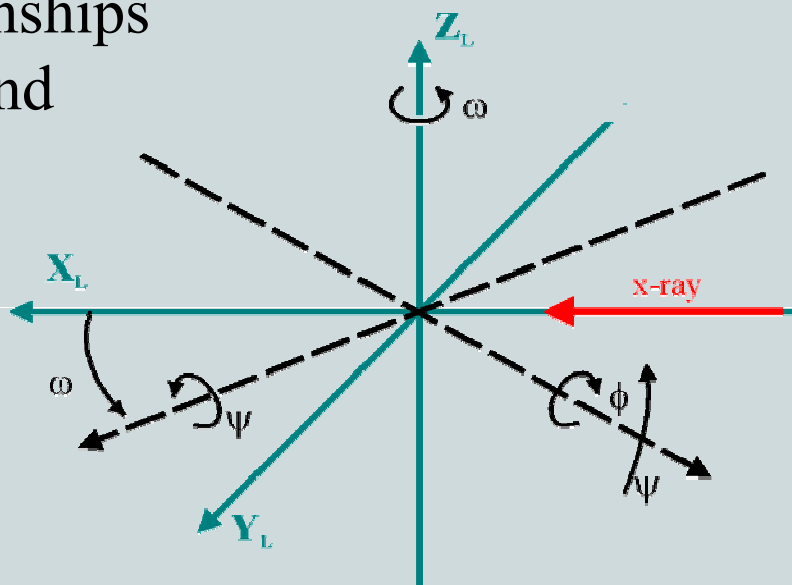


XRD²: Sample Space & Eulerian Geometry



The angular relationships between $X_L Y_L Z_L$ and $S_1 S_2 S_3$ are:

	X_L	Y_L	Z_L
S_1	a_{11}	a_{12}	a_{13}
S_2	a_{21}	a_{22}	a_{23}
S_3	a_{31}	a_{32}	a_{33}



The transformation matrix from the diffraction space to the sample space is:

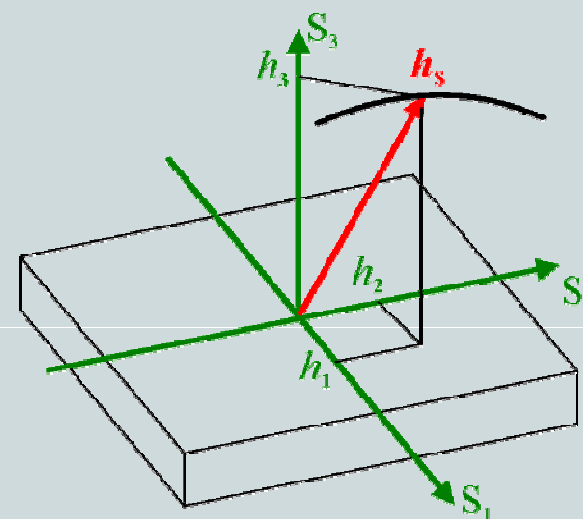
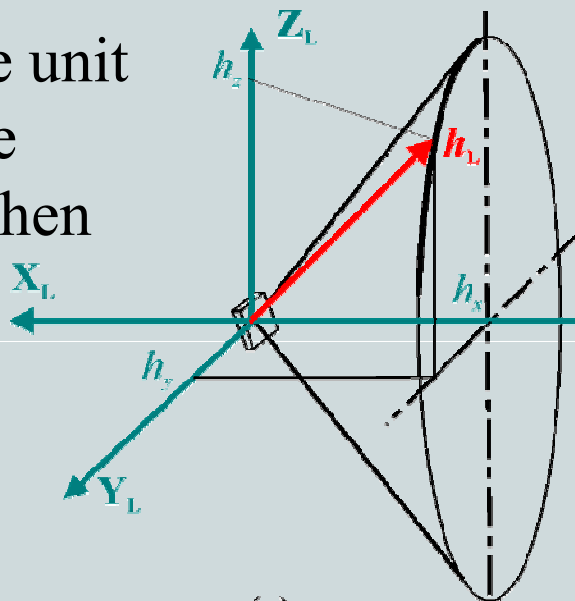
$$\begin{bmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{33} \end{bmatrix} = \begin{bmatrix} -\sin \omega \sin \psi \sin \phi & \cos \omega \sin \psi \sin \phi & -\cos \psi \sin \phi \\ -\cos \omega \cos \phi & -\sin \omega \cos \phi & \cos \psi \cos \phi \\ \sin \omega \sin \psi \cos \phi & -\cos \omega \sin \psi \cos \phi & \sin \psi \\ -\cos \omega \sin \phi & -\sin \omega \sin \phi & \sin \psi \\ -\sin \omega \cos \psi & \cos \omega \cos \psi & \sin \psi \end{bmatrix}$$

XRD²: Sample Space & Unit Diffraction Vector



The components of the unit vector \mathbf{h}_S in the sample coordinates $\mathbf{S}_1\mathbf{S}_2\mathbf{S}_3$ is then given by

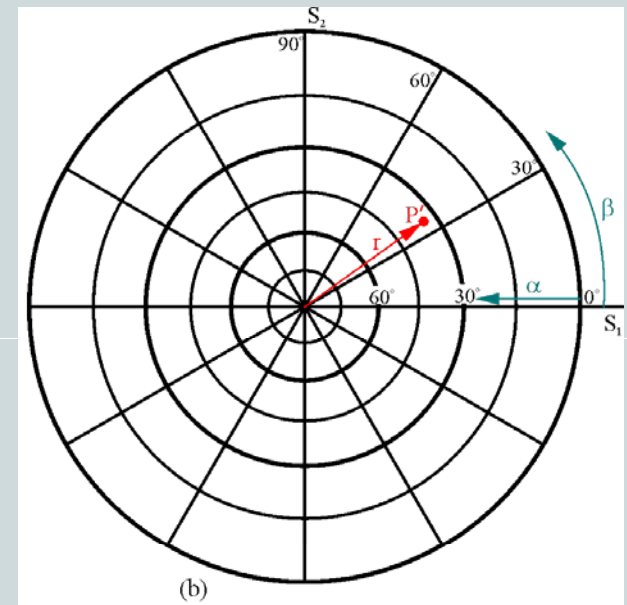
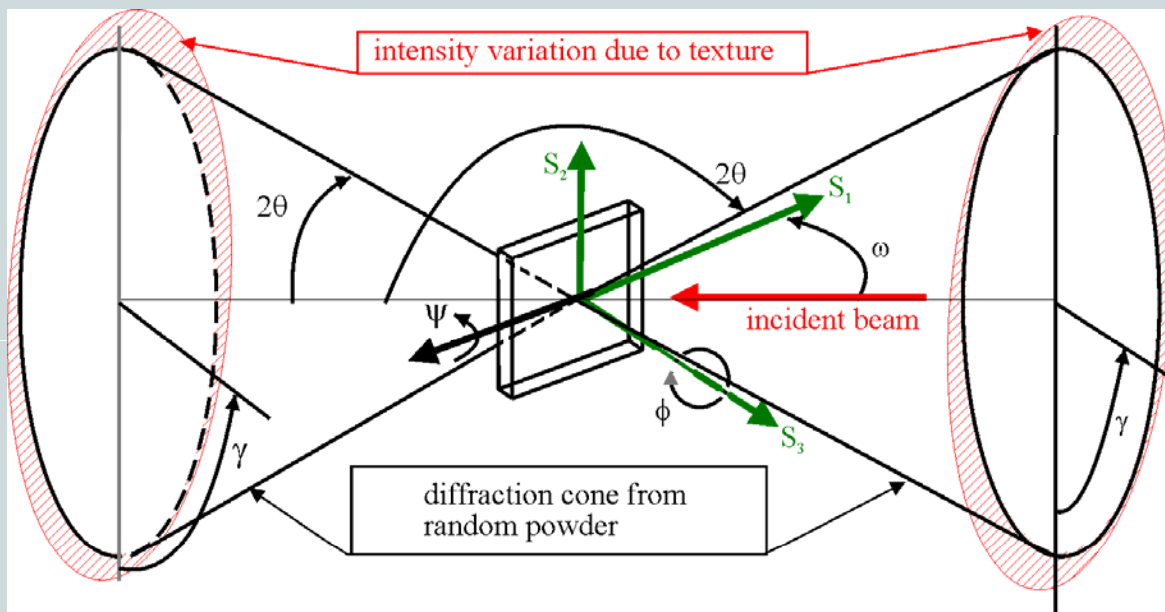
$$\begin{bmatrix} h_1 \\ h_2 \\ h_3 \end{bmatrix} = \begin{bmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{33} \end{bmatrix} \begin{bmatrix} h_x \\ h_y \\ h_z \end{bmatrix}$$



Or
in expanded form:

$$\begin{aligned} h_1 &= \sin\theta(\sin\phi\sin\psi\sin\omega + \cos\phi\cos\omega) + \cos\theta\cos\gamma\sin\phi\cos\psi \\ &\quad - \cos\theta\sin\gamma(\sin\phi\sin\psi\cos\omega - \cos\phi\sin\omega) \\ h_2 &= -\sin\theta(\cos\phi\sin\psi\sin\omega - \sin\phi\cos\omega) - \cos\theta\cos\gamma\cos\phi\cos\psi \\ &\quad + \cos\theta\sin\gamma(\cos\phi\sin\psi\cos\omega + \sin\phi\sin\omega) \\ h_3 &= \sin\theta\cos\psi\sin\omega - \cos\theta\sin\gamma\cos\psi\cos\omega - \cos\theta\cos\gamma\sin\psi \end{aligned}$$

XRD²: Fundamental Equation for Texture Analysis



The pole figure angles (α, β) can be calculated from the unit vector components by the pole mapping equations:

$$\alpha = \sin^{-1} |h_3| = \cos^{-1} \sqrt{h_1^2 + h_2^2}$$

$$\beta = \pm \cos^{-1} \frac{h_1}{\sqrt{h_1^2 + h_2^2}} \quad \begin{cases} \beta \geq 0^\circ & \text{if } h_2 \geq 0 \\ \beta < 0^\circ & \text{if } h_2 < 0 \end{cases}$$

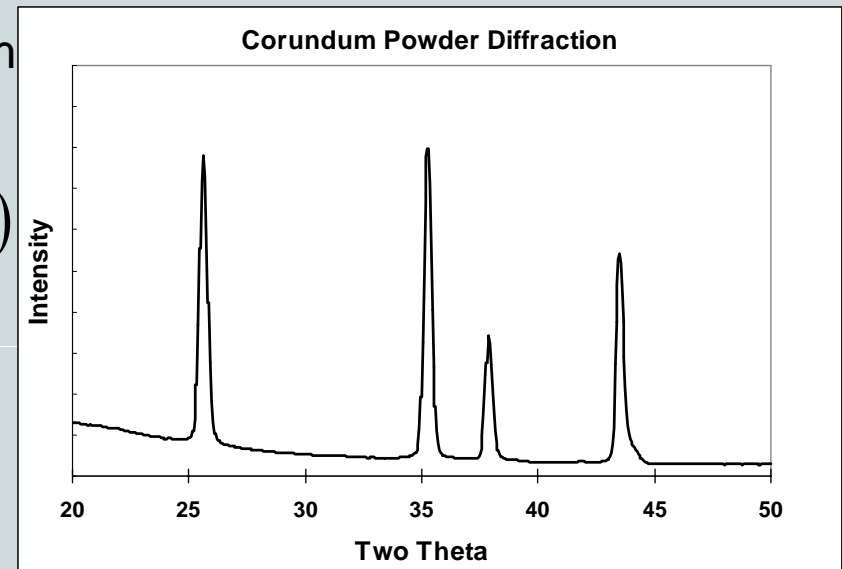
XRD²: Relative Intensity of Powder Pattern

The integrated intensity diffracted from random polycrystalline materials is given by:

$$I_{hkl} = k_I \frac{p_{hkl}}{v^2} (LPA) \lambda^3 F_{hkl}^2 \exp(-2M_t - 2M_s)$$

where:

- ⊕ k_I - instrument constant;
- p_{hkl} - the multiplicity of the planes;
- v - the volume of the unit cell;
- ⊕ (LPA) - the Lorentz-polarization and absorption factors;
- F_{hkl}^2 - the structure factor of the crystal plane (hkl) and
- $\exp(-2M_t - 2M_s)$ - the attenuation factor due to lattice thermal vibrations and weak static displacements.



-
- ⊕ Denotes the factors which are different between Bragg-Brentano geometry and XRD² geometry. k_I is determined by the source, optics and detector (PPXRD-7) (LPA) will be given in this presentation.

XRD²: Polarization Correction

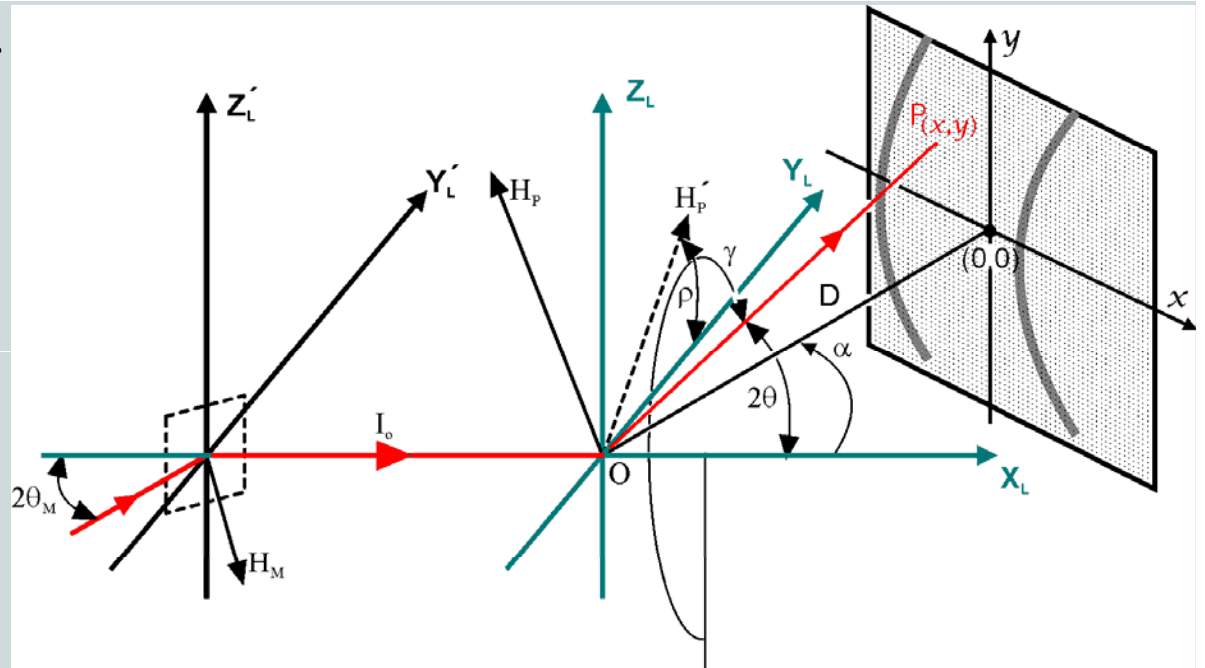
The polarization factor for Bragg-Brentano geometry with incident beam monochromator is:

$$P_I = \frac{1 + \cos^2 2\theta_M \cos^2 2\theta}{1 + \cos^2 2\theta_M}$$

where $2\theta_M$ is the Bragg angle of the monochromator.

The general polarization factor for the diffracted beam to point P is:

$$P_G = \frac{(\cos^2 2\theta \cos^2 \rho + \sin^2 \rho) \cos^2 2\theta_M + \cos^2 2\theta \sin^2 \rho + \cos^2 \rho}{1 + \cos^2 2\theta_M}$$



Geometric relationship between the monochromator and detector in laboratory coordinates.

XRD²: Polarization Correction

The unit vector of the diffraction vector \mathbf{H}_p and its projection on Y_L - Z_L plane, \mathbf{H}'_p , in the laboratory system are given respectively as:

$$\mathbf{h}_L = \begin{bmatrix} h_x \\ h_y \\ h_z \end{bmatrix} = \begin{bmatrix} -\sin\theta \\ -\cos\theta\sin\gamma \\ -\cos\theta\cos\gamma \end{bmatrix} \quad \mathbf{h}'_L = \begin{bmatrix} 0 \\ h'_y \\ h'_z \end{bmatrix} = \begin{bmatrix} 0 \\ -\sin\gamma \\ -\cos\gamma \end{bmatrix}$$

The unit vector of Y_L is $\mathbf{y}_L = [0, 1, 0]$, then:

$$\cos\rho = \cos(\mathbf{h}'_L, \mathbf{y}_L) = \mathbf{h}'_L \cdot \mathbf{y}_L = -\sin\gamma$$

Therefore, $\cos^2\rho = \sin^2\gamma$ and $\sin^2\rho = \cos^2\gamma$

The polarization factor for XRD² can then be given as a function of both θ and γ :

$$P(\theta, \gamma) = \frac{(1 + \cos^2 2\theta_M \cos^2 2\theta) \sin^2 \gamma + (\cos^2 2\theta_M + \cos^2 2\theta) \cos^2 \gamma}{1 + \cos^2 2\theta_M}$$

XRD²: Sample Absorption Correction



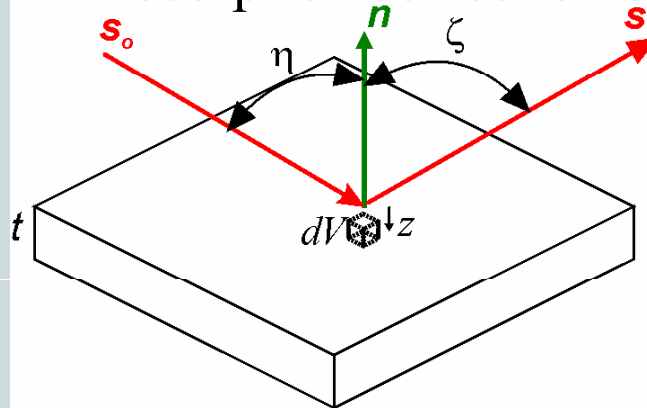
The absorption can be measured by the transmission coefficient:

$$A = \frac{1}{V} \int_V e^{-\mu\tau} dV$$

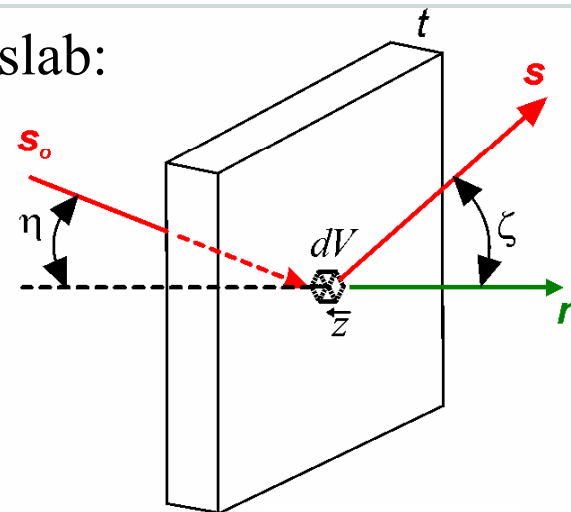
where τ is the total beam path and A is the average over all the element dV . For Bragg-Brentano geometry, we have:

$$A_{BB} = 1/(2\mu)$$

Absorption correction of flat slab:



(a) reflection

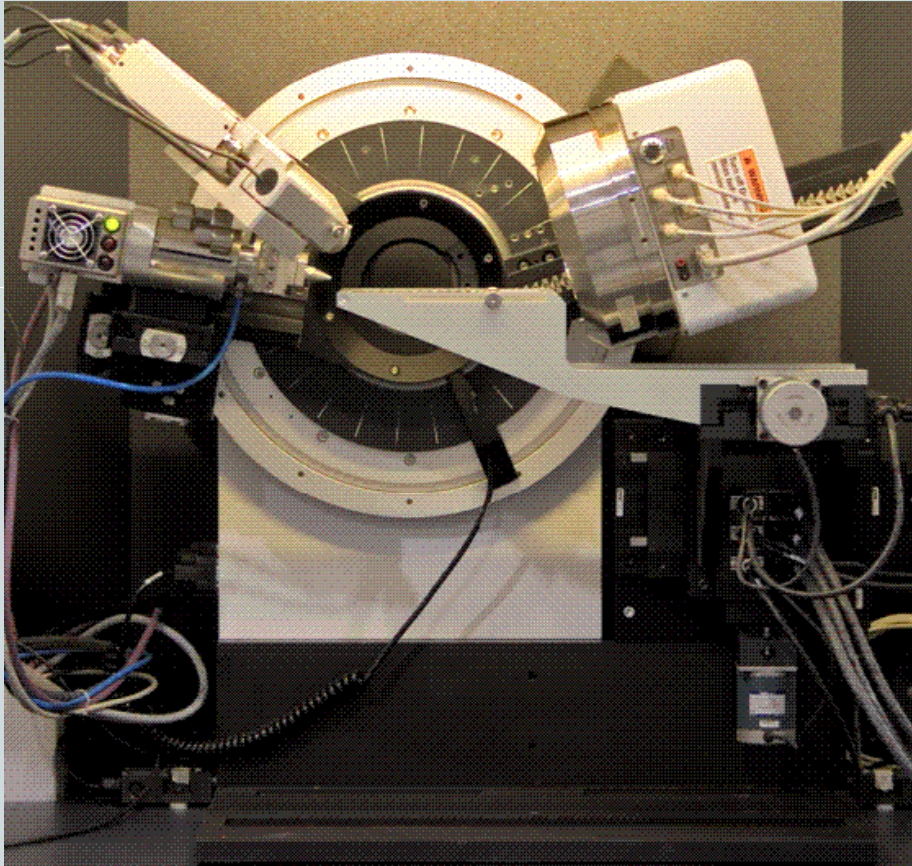


(b) transmission.

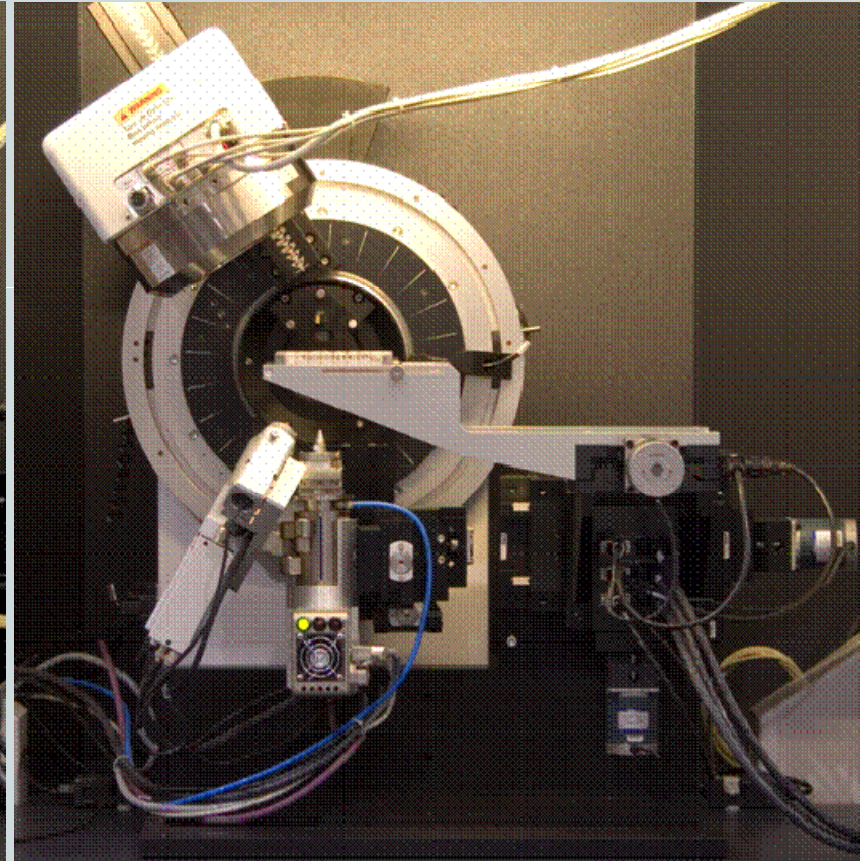
To make the relative intensity comparable to Bragg-Brentano geometry, we introduce a normalized transmission coefficient T :

$$T = A / A_{BB} = 2\mu A$$

D8 DISCOVER with GADDS HTS ($I\mu S$): Reflection & Transmission (Pharmaceutical Delight)



■ Reflection mode



■ Transmission mode

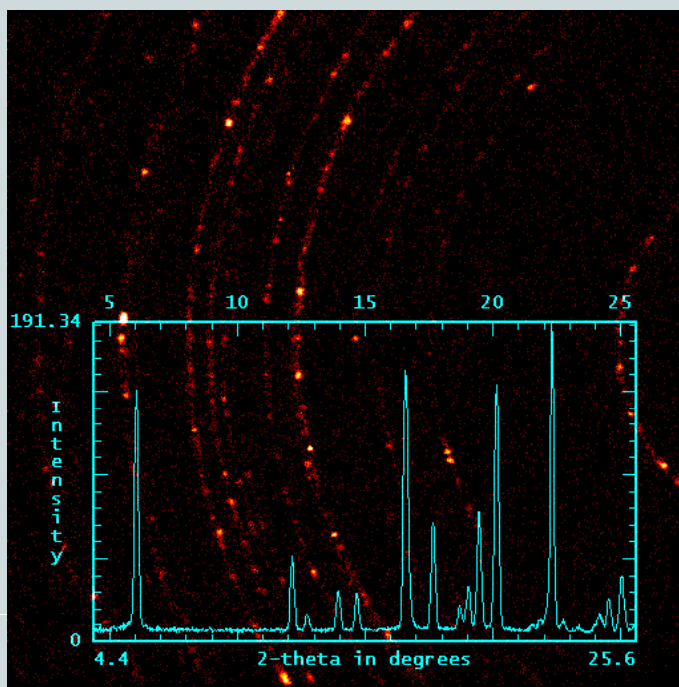
Comparison: Ibuprofen I μ S & VÅNTEC-2000 vs. Clasical set-up



Sealed Tube

- 0.3 mm collimator
- Sample-Detector distance 29 cm

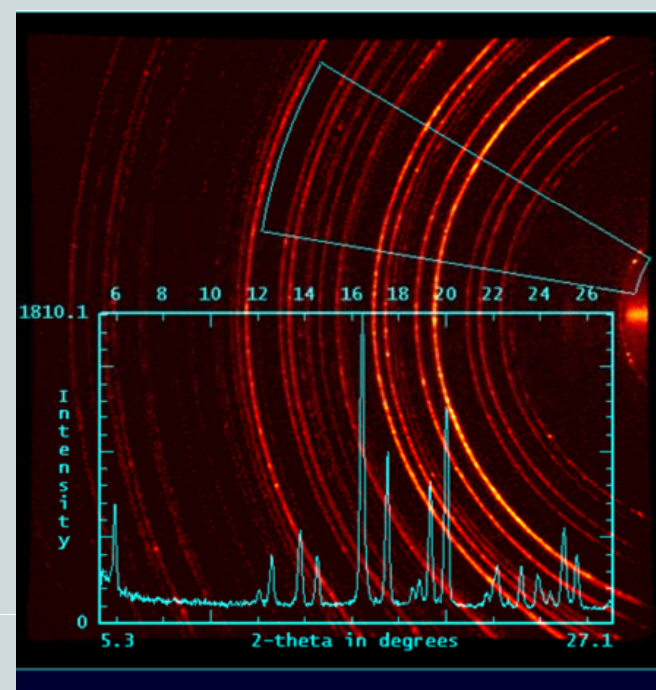
120 sec collection time



I μ S – XRD² – focus

- 2mmX2mm on sample, and 200um spot focused on detector
- small slice for integration to obtain better resolution

15 sec collection time



XRD²: Sample Absorption Correction

For reflection mode diffraction with a thick plate:

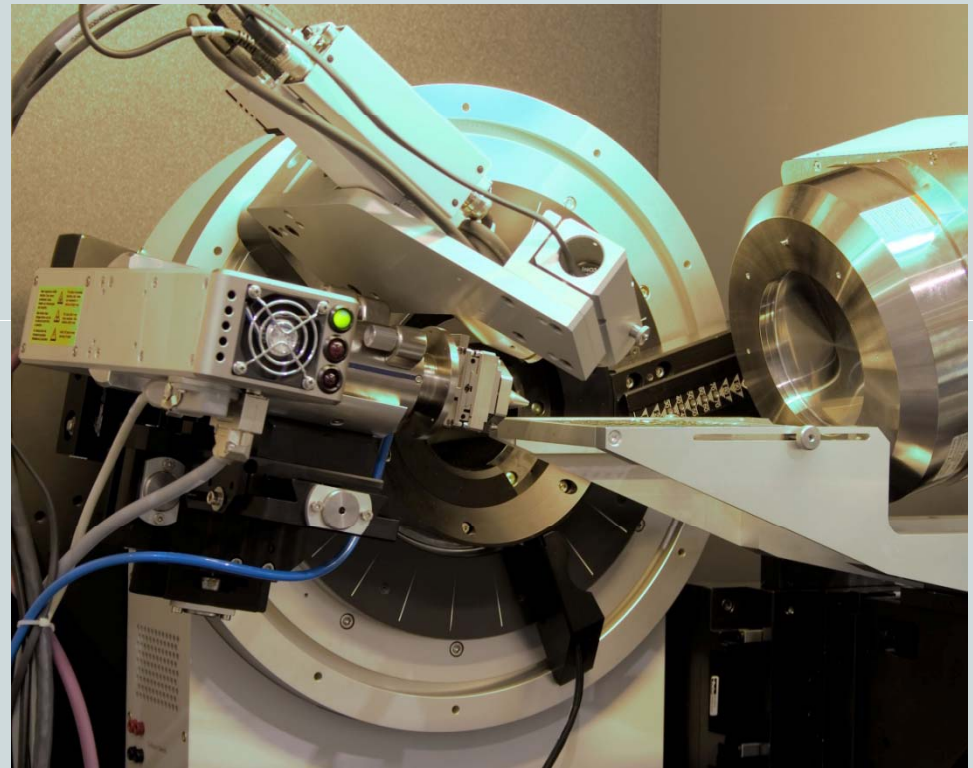
$$s_o = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} \quad s = \begin{bmatrix} \cos 2\theta \\ -\sin 2\theta \sin \gamma \\ -\sin 2\theta \cos \gamma \end{bmatrix}$$

and

$$n = \begin{bmatrix} -\sin \omega \cos \psi \\ \cos \omega \cos \psi \\ \sin \psi \end{bmatrix}$$

The normalized transmission coefficient:

$$T = \frac{2 \cos \eta}{(\cos \eta + \cos \zeta)} \quad \text{with} \quad \begin{aligned} \cos \eta &= -s_o \cdot n = \sin \omega \cos \psi \\ \cos \zeta &= s \cdot n = -\cos 2\theta \sin \omega \cos \psi \\ &\quad -\sin 2\theta \sin \gamma \cos \omega \cos \psi - \sin 2\theta \cos \gamma \sin \psi \end{aligned}$$



XRD²: Sample Absorption Correction

For transmission mode:

$$s_o = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix} \quad s = \begin{bmatrix} \cos 2\theta \\ -\sin 2\theta \sin \gamma \\ -\sin 2\theta \cos \gamma \end{bmatrix}$$

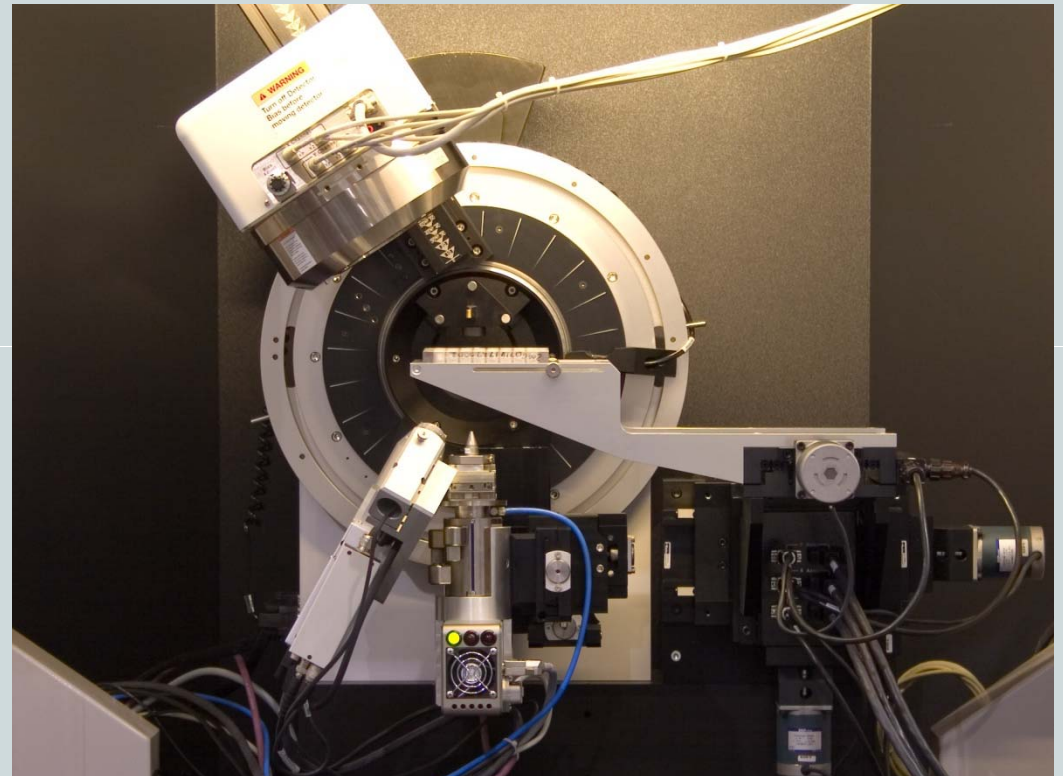
and

$$n = \begin{bmatrix} \sin \omega \sin \psi \sin \phi + \cos \omega \cos \phi \\ -\cos \omega \sin \psi \sin \phi + \sin \omega \cos \phi \\ \cos \psi \sin \phi \end{bmatrix}$$

The normalized transmission coefficient :

$$T = \frac{2 \sec \eta [\exp(-\mu t \sec \eta) - \exp(-\mu t \sec \zeta)]}{\sec \zeta - \sec \eta}$$

$$\begin{aligned} \cos \eta &= s_o \cdot n = \sin \omega \sin \psi \sin \phi + \cos \omega \cos \phi \\ \cos \zeta &= s \cdot n = (\sin \omega \sin \psi \sin \phi + \cos \omega \cos \phi) \cos 2\theta \\ &\quad + (\cos \omega \sin \psi \sin \phi - \sin \omega \cos \phi) \sin 2\theta \sin \gamma \\ &\quad - \cos \psi \sin \phi \sin 2\theta \cos \gamma \end{aligned}$$



XRD²: Texture Effect and Correction



The integrated intensity with texture is:

$$I_{hkl} = k_I \frac{P_{hkl}}{v^2} (LPA) \lambda^3 F_{hkl}^2 g_{hkl}(\alpha, \beta) \exp(-2M_t - 2M_s)$$

where $g()$ is the normalized pole density function.

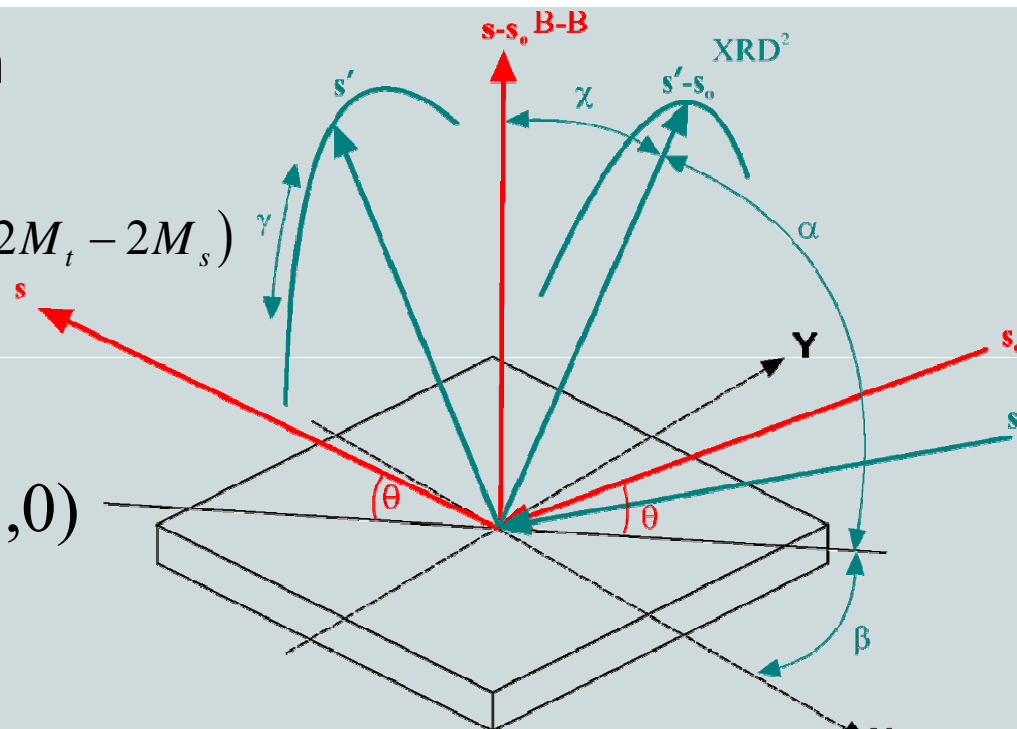
For the BB geometry, $g_{hkl}(\frac{\pi}{2}, 0)$

The texture effect for XRD²:

$$I_{hkl}^c = \frac{I_{hkl}^m}{\langle g_{hkl}(\Delta\gamma) \rangle}$$

Correct to the B-B equivalent with a texture effect:

$$I_{hkl}^{BB} = \frac{g_{hkl}(\frac{\pi}{2}, 0) I_{hkl}^m}{\langle g_{hkl}(\Delta\gamma) \rangle}$$



For fiber texture: $\int_{\gamma_1}^{\gamma_2} g_{hkl}[\chi(\gamma)] \gamma d\gamma$

$$\langle g_{hkl}(\Delta\gamma) \rangle = \frac{\int_{\gamma_1}^{\gamma_2} g_{hkl}[\chi(\gamma)] \gamma d\gamma}{\gamma_2 - \gamma_1}$$

and $\chi = \cos^{-1} |h_3|$

XRD²: Summary and Suggestions

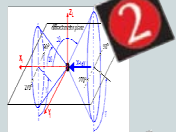


- Two-dimensional X-ray diffraction has many advantages over the conventional diffraction (speed, completeness and accuracy, micro-sample volume).
- The discrepancy between XRD² and Bragg-Brentano in geometry, polarization, absorption and preferred orientation can be interpreted and corrected.
- Fortunately, most of the discrepancies can be ignored without affecting the specific application.
- Or the corrections are already built in the software, so users do have to work on the correction details.

XRD²: Theory, System and Applications



X-ray Diffraction (XRD)



Geometry Conventions



System and Configuration



Phase Identification



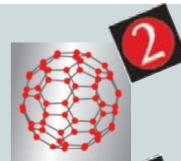
Quantitative Analysis



Texture



Stress



Small Angle X-ray Scattering



Micro Diffraction



Mapping



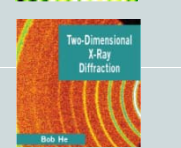
Thin Films



High-Throughput Screening



Forensics and Archaeology



Reference

XRD²: Fundamentals, theory and applications

THE FUNDAMENTALS, THEORY, AND WIDE-RANGING APPLICATIONS OF TWO-DIMENSIONAL X-RAY DIFFRACTION

Two-Dimensional X-Ray Diffraction is proving itself as an ideal non-destructive, analytical method for measuring the atomic arrangement of materials and extracting an array of information beyond the limitations of conventional X-ray diffraction. Researchers in materials science, chemistry, physics, pharmaceuticals, and related fields will find this introductory reference invaluable in understanding and applying two-dimensional X-ray diffraction for examining a broad range of samples.

Two-Dimensional X-ray Diffraction shows how two-dimensional X-ray diffraction can be a useful tool for the examination of metals, polymers, ceramics, semiconductors, thin films, coatings, paints, biomaterials and composites for material science researches, molecular structure determination and polymorphism study for drug discovery and processing, and samples with micro volume or micro-area for forensic analysis, and archaeology analysis, to name just a few of the method's applications.

The text covers:

- The fundamentals of X-ray diffraction and its extension to two-dimensional X-ray diffraction
- The geometry conventions and diffraction vector approach for diffraction data interpretation, data correction, and process algorithms for various applications
- Instrumentation technologies, including the critical components, such as X-ray source and optics, two-dimensional detectors, goniometer, and sample stages
- The configurations of the two-dimensional X-ray diffraction systems for various applications, such as phase identification, texture, stress, microstructure analysis, crystallinity, thin film analysis, and combinatorial screening
- Experimental examples in materials research, pharmaceuticals, materials processing, and quality control

Written by one of the pioneers in the field, *Two-Dimensional X-Ray Diffraction* brings readers up to speed on a fast-rising, state-of-the-art method for materials characterization.

BOB BAOPING HE is the Director of R&D and Engineering at Bruker AXS (formerly Siemens AXS). Mr. He holds a PhD -in materials science from Virginia Tech and holds twelve U.S. patents.

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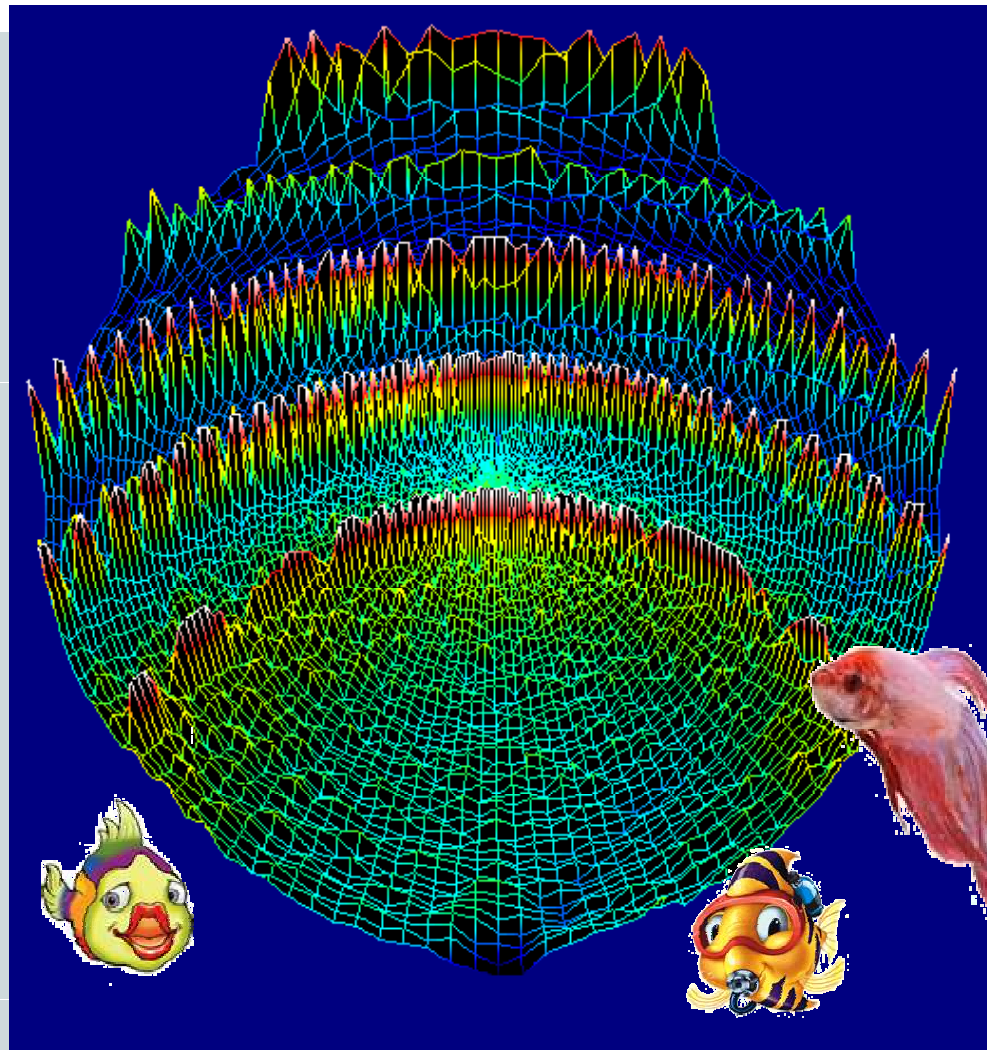
He
Two-Dimensional X-Ray Diffraction

Two-Dimensional
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