

Lesson 3 Sample Preparation & Problems

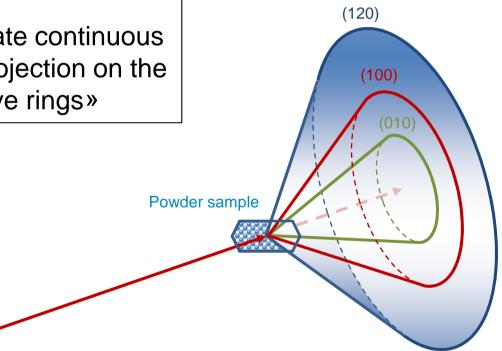
Nicola Döbelin RMS Foundation, Bettlach, Switzerland



February 11 – 14, 2013, Riga, Latvia

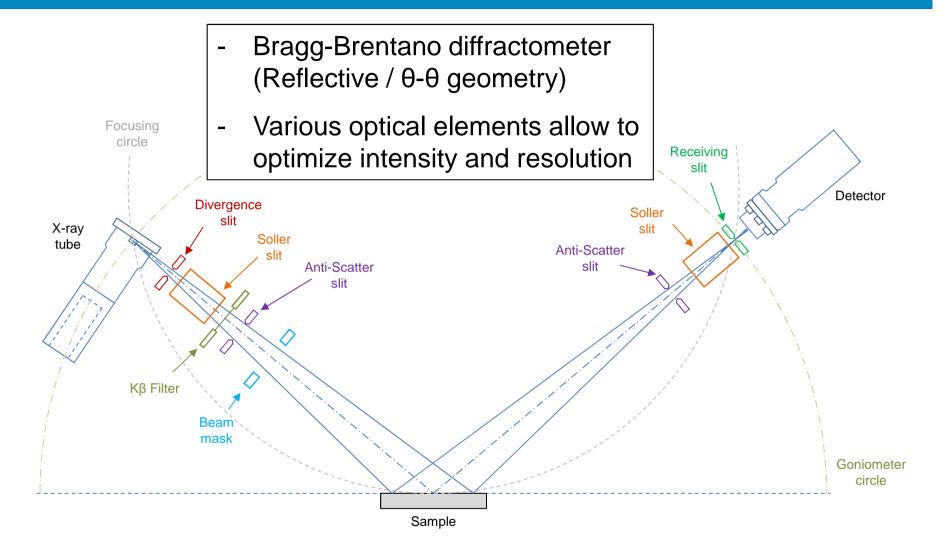
Repetition: Powder XRD Samples

- A fine powder contains particles in all possible orientations
- The orientation distribution is random, no orientation is preferred
- Powder samples generate continuous diffraction cones, the projection on the detector is called «Debye rings»





Repetition: Bragg-Brentano Diffractometer



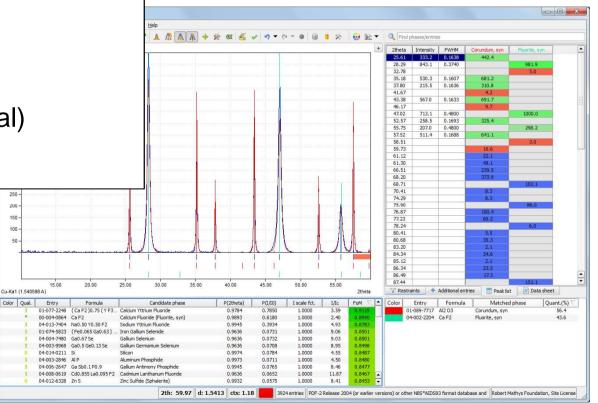


Repetition: Phase Identification

200 150

100

- Crystalline phases are identified by comparing peak positions with a database (Search/Match)
- Qualitative (sometimes semi-quantitative) results
- Databases: -
 - PDF-2/4 (commercial)
 - COD (open-access)





Sample Preparation

Sample preparation is **ABSOLUTELY CRUCIAL** for a good diffraction pattern!

Some problems encountered during Rietveld refinement are inherent to the sample.

Some others are related to sample preparation errors.

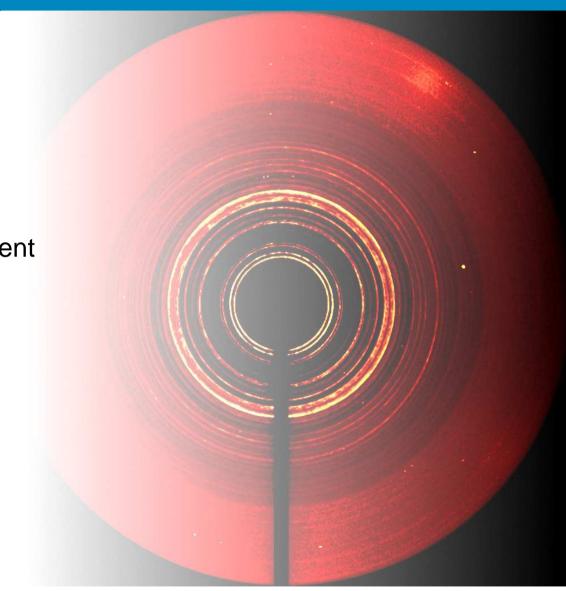




Problems

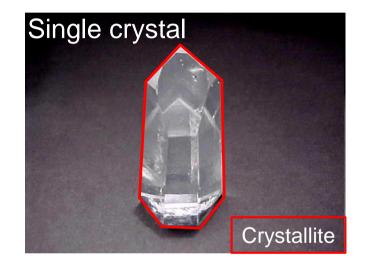
- Graininess
- Micro-absorption
- Texture
- Crystallite size
- Sample height displacement
- Surface roughness
- Sample transparency





Particles & Crystallites

Crystallite = Domain of coherent crystal structure





Particle size = Crystallite size

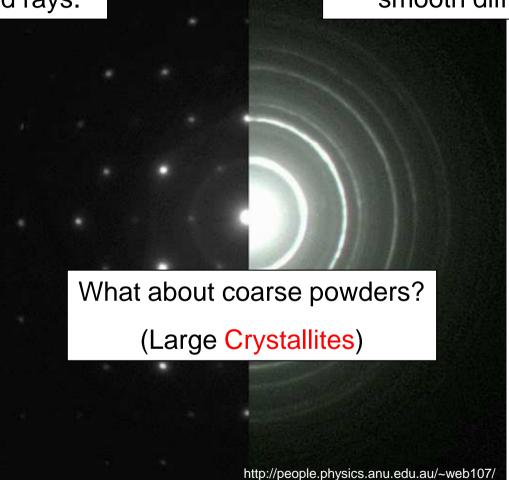
Particle size > Crystallite size



Graininess

Single crystals generate spotty diffracted rays.

Fine powders crystals generate smooth diffraction rings.

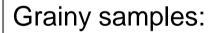




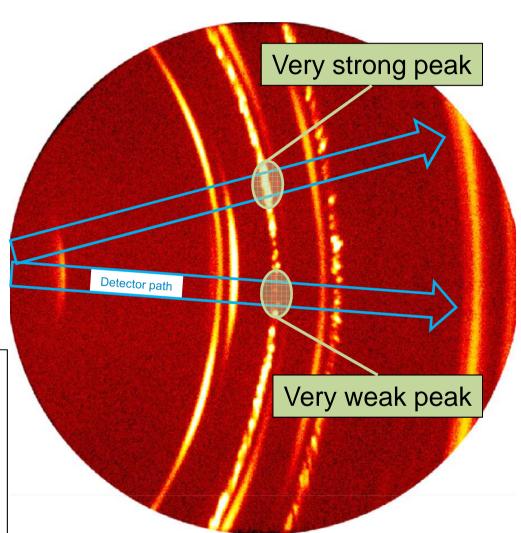
Graininess

Spotty diffraction rings

The same sample, at the same 20 position, but different intensities!



- non-reproducible intensities
- «phantom» peaks
- «missing» peaks



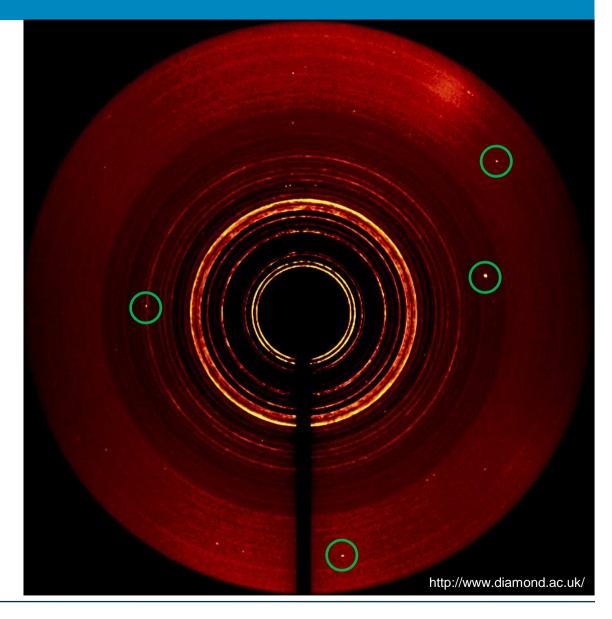
Bruker AXS



Graininess: Rocks in Dust

«Rocks in Dust»: A few large crystals in a fine matrix

Usually invisible, but if scanned: Strong peaks out of nowhere!

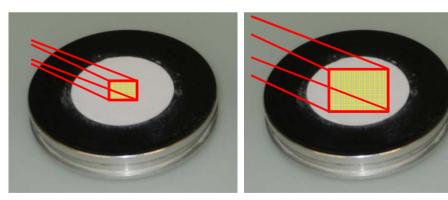




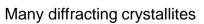
Graininess

Reducing graininess:

- Grinding / milling
- Adjust divergence slit and beam mask for largest possible irradiated area (= more particles contribute to diffraction pattern)
- Use spinning sample stage (= better randomisation)



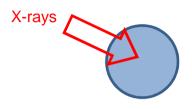
Few diffracting crystallites



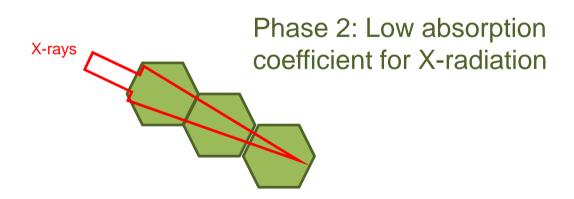


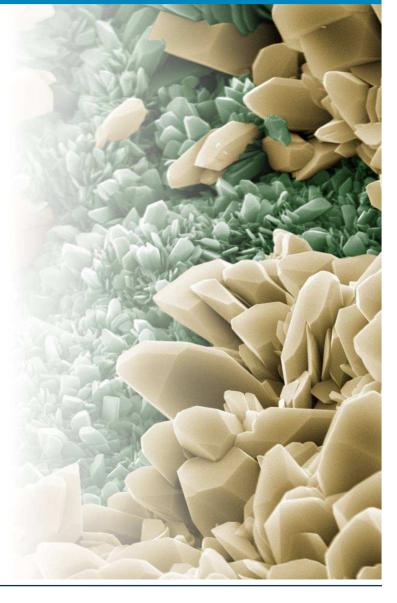


Micro-absorption



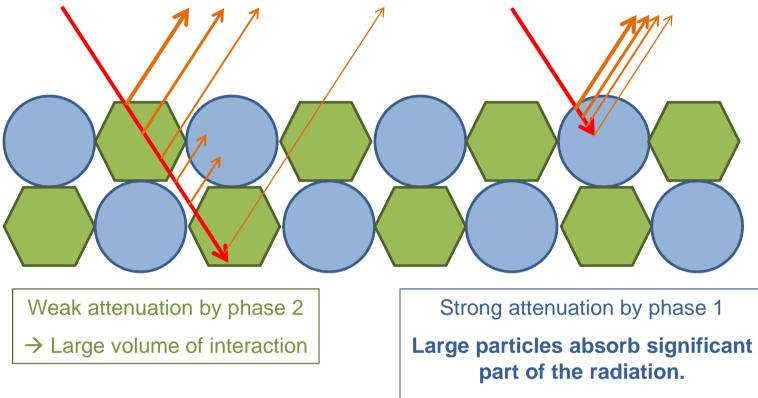
Phase 1: High absorption coefficient for X-radiation







Micro-absorption

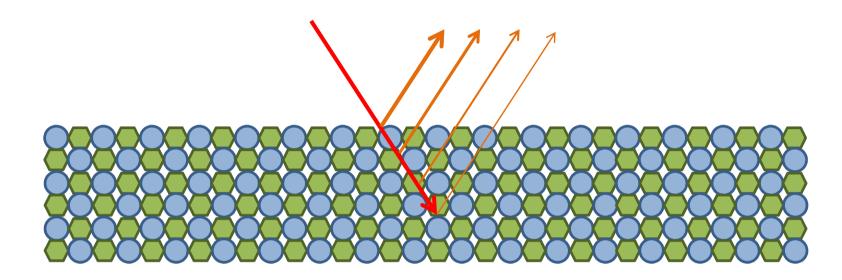


 \rightarrow Small volume of interaction

Phase quantification biased for phase 2!



Micro-absorption



Small particles absorb insignificant part of the radiation.

- → Similar volumes of interaction with phases 1 & 2
 - \rightarrow Correct phase quantification



Micro-absorption occurs in samples with...

- ... large particles (not crystallites!)
- ... phases with large differences in absorption coefficients

Reducing micro-absorption:

- Grinding / milling to reduce particle size

Mathematical correction:

- Difficult, but possible to a certain degree (more about «Brindley correction» in the lesson on «Rietveld refinement»)



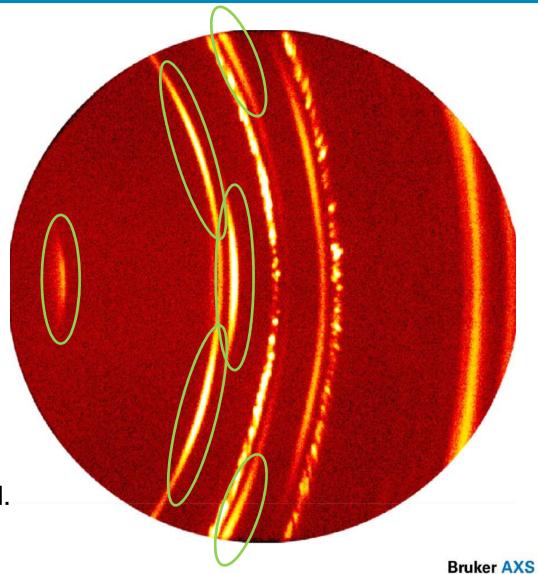


Texture, Preferred Orientation

Smooth, but non-continuous diffraction rings

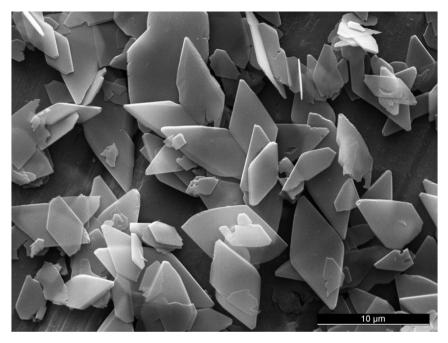
Reason:

- Orientation of crystallites is not random.
- Some orientations are over-represented, others are under-represented.

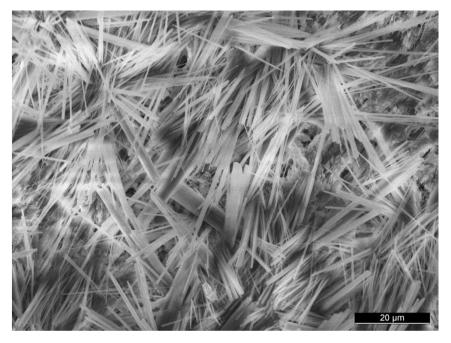




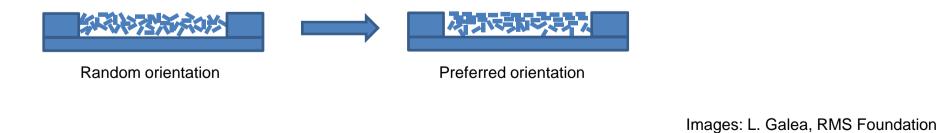
Texture, Preferred Orientation



Platelets



Needles, Fibers, Whiskers



RMS

Texture, Preferred Orientation

Try to avoid orientation at the surface of the sample:

- Press powder without «rubbing» the surface
- Use back-loading sample holder
- Disorder surface with textured stamp
- Various creative solutions can be found on the internet (involving Vaseline, hair spray, ...)

PO can be corrected mathematically, but phase quantification will be biased. (more on this in the lesson on «Rietveld refinement»)





More sources in ordered arrangement

More distinct interference pattern

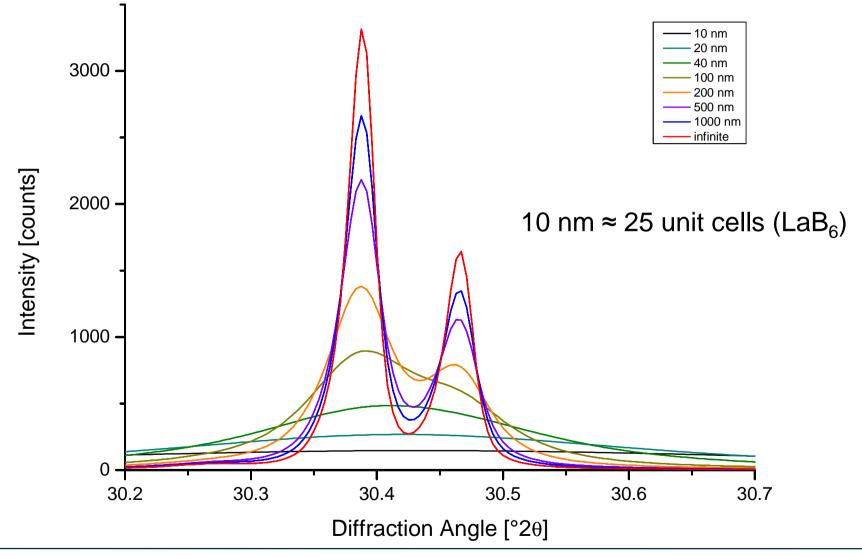
Larger domains of coherent crystal structure = More distinct diffraction pattern

Image: http://www.forbes.com/

••••• Testing • Research • Consulting

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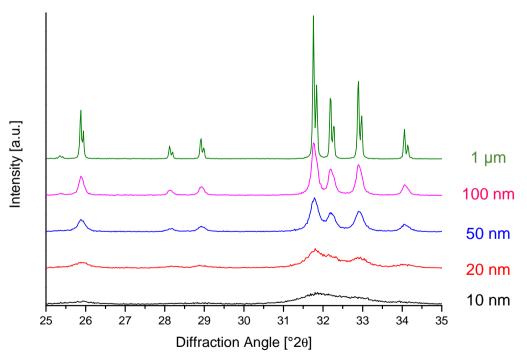


Small crystallites generate broad peaks

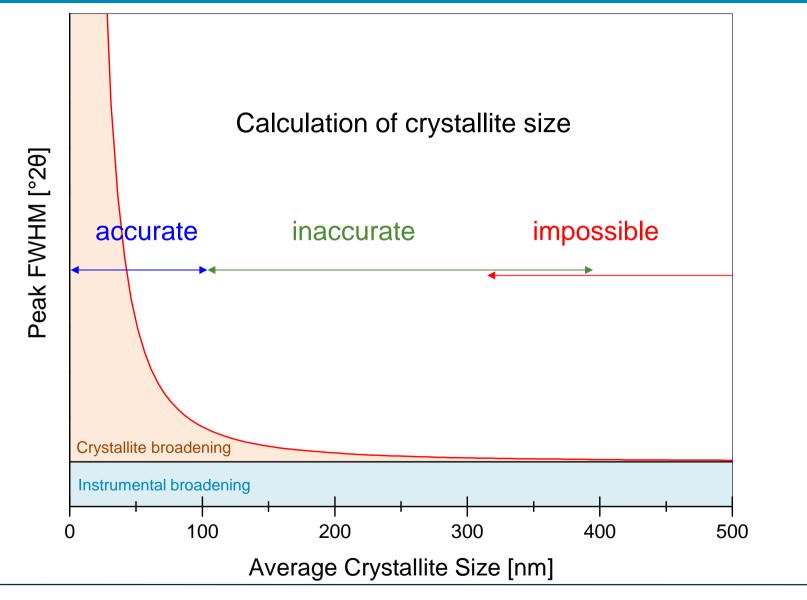
Crystallite size can be calculated from peak broadening

Severe peak broadening:

- Excessive peak overlap
- Unusable diffraction pattern

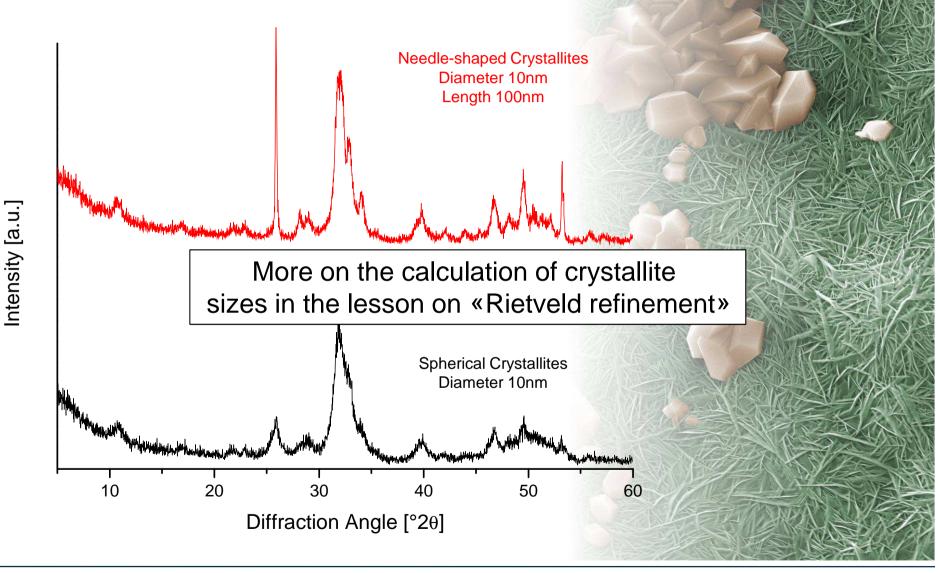








Anisotropic Crystallites



RMS

Summary: Ideal Particle Size

- Ideal particle and crystallite size: 1–5 µm
- Larger particles: Micro-absorption
- Larger crystallites: Grainy sample
- Caution: High-energy milling (e.g. planetary mill) generates:
 - Lattice defects and strain in the crystal structure
 - Peak broadening due to reduction of crystallite size
 - Amorphous fraction (invisible to XRD)
- Manual milling in agate mortar is usually recommended







RMS

Automatic mill for XRD

«Industry standard» for automatic XRD sample milling: McCrone Micronizing Mill

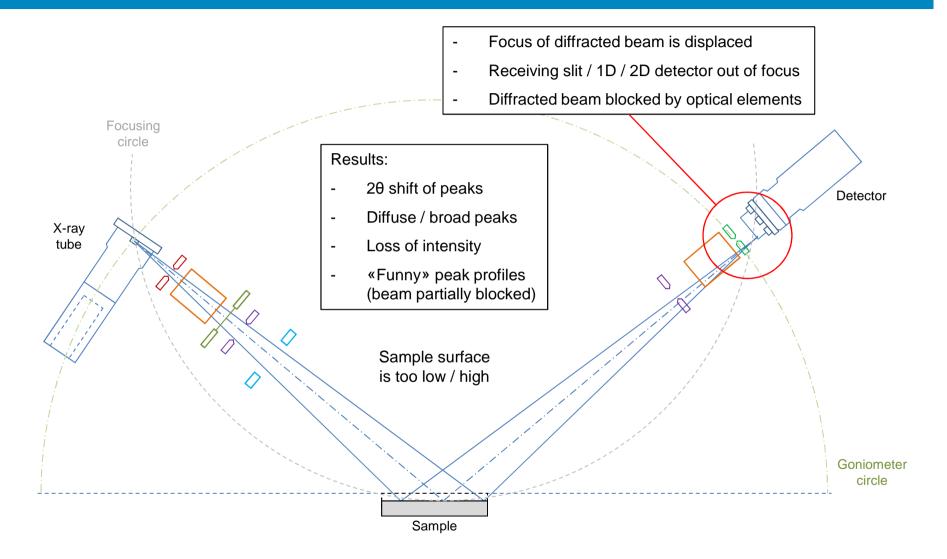


http://www.powderbulksolids.com



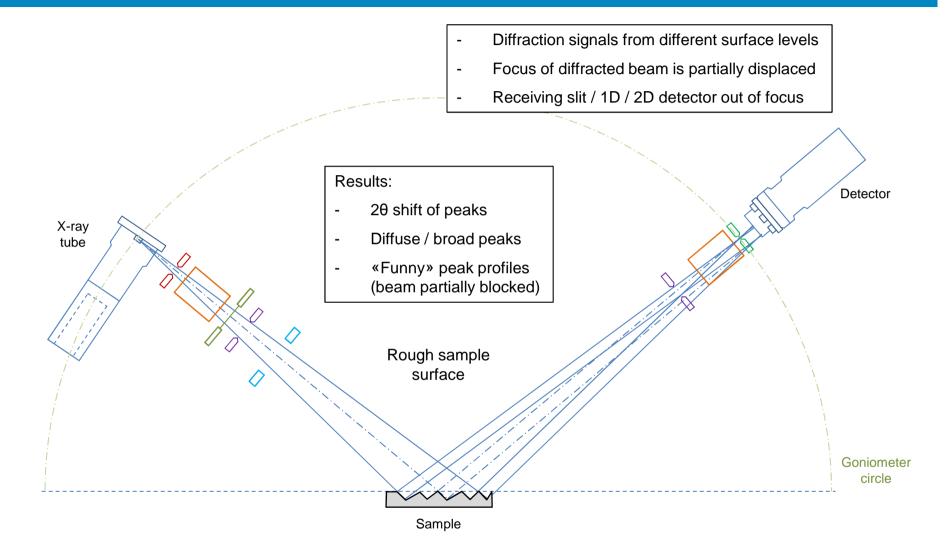


Sample Height Displacement



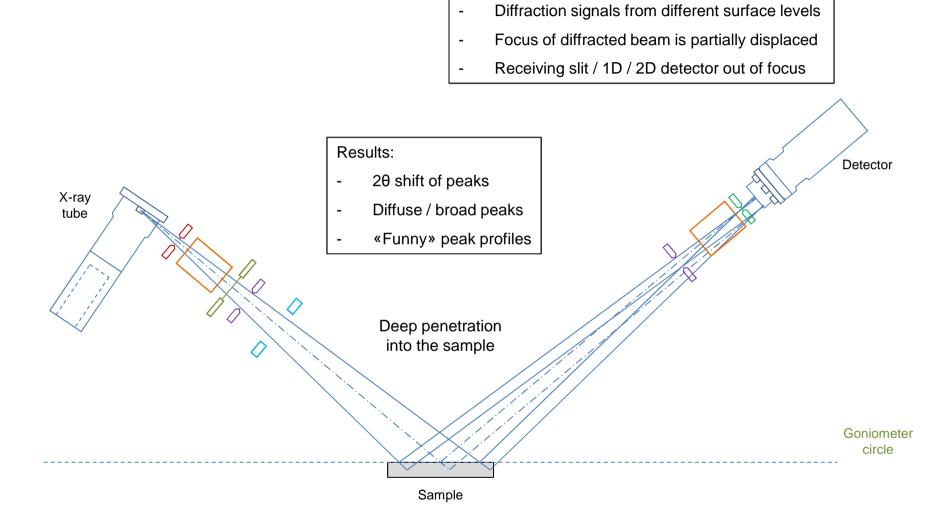


Rough Sample Surface





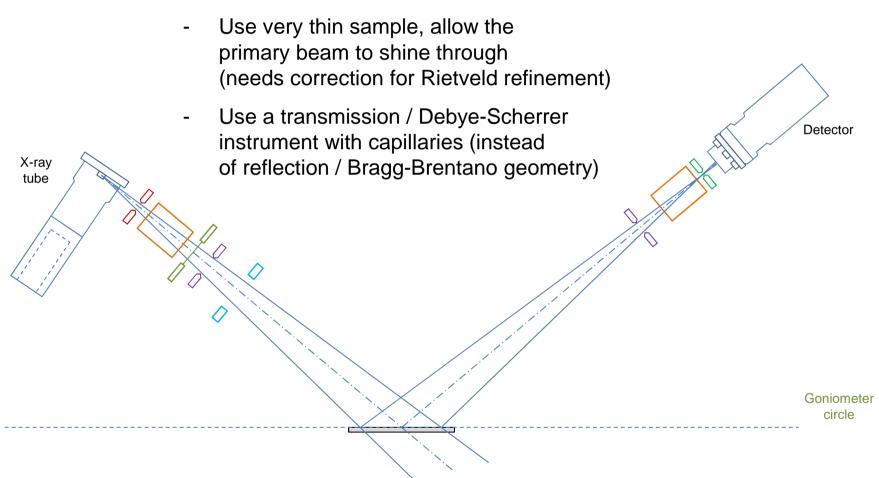
Sample Transparency





Sample Transparency

Possible solutions:



Summary: The Perfect Sample

The perfect sample for Bragg-Brentano diffractometers:

- Crystallites and particles of 1-5 µm size
- Perfectly random orientation
- Perfectly flat surface
- Surface precisely centered in the goniometer
- Short penetration depth





Software Download

For tomorrow:

Download software and examples from:

http://www.doebelin.org/nic/xrd

Login name: Riga2013

Password: Rietveld

Software installation sessions:

- Today, 12:00-12:30
- Tomorrow, 08:45-09:15

