
Methods of High-Pressure Single-Crystal X-ray Diffraction

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- **Technical challenges**
 - **Overview of methods**
 - **Overview of the workshop**

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Single crystal and powder

- **Single-crystal data is 3 dimensional not 1 dimensional, intrinsically superior to powder diffraction:**
 - No peak overlaps
 - No resolution problems
 - No preferred orientation problems
 - No “powder average” problems
 - Signal to noise is higher

- **Single crystal diffraction allows:**
 - The unambiguous determination of minute structural changes
 - The determination of small structural distortions (phase transitions)
 - Acentric crystals
 - Measurement of diffuse scattering and incommensurate structures
 - Reliable displacement parameters

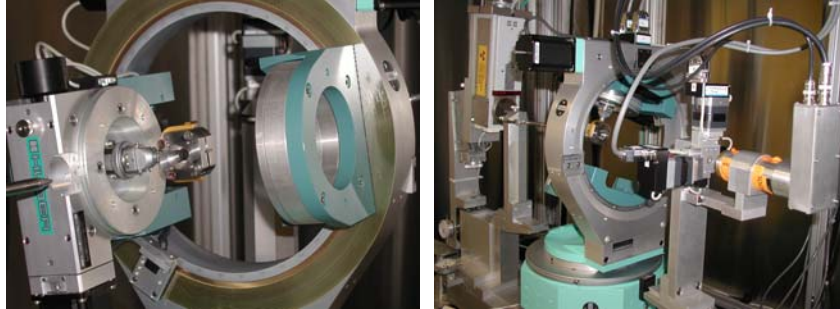
- **“A bad single crystal is better than a good powder” (McMahon)**

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Cell parameters at high pressure

Precise diffraction angle data to measure small changes in cell parameters at high pressures:



In-house single-crystal X-ray diffraction.

Eulerian cradle diffractometers (Huber, P4, Stadi4), Single software Cell parameters to a precision of 1 part in 30,000

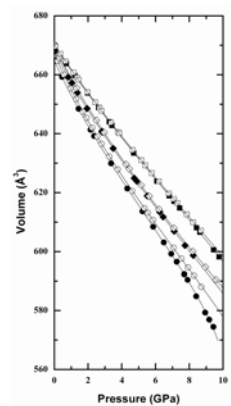
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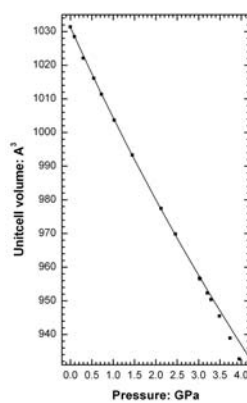
Equations of State and Phase Transitions

Precise cell parameter data allows studies of small changes in EoS and of phase transitions:

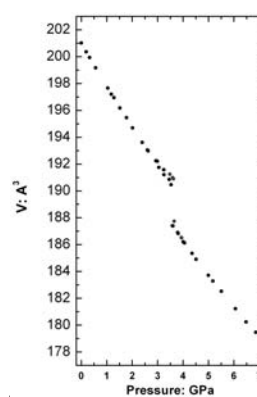
*Order/disorder in plagioclase
Sochalski-Kolbus et al. (2011)*



*Softening in CuCO₃
Spencer et al. (2011)*



Isosymmetric phase transition in kalsilite. Gatta et al. (2011)

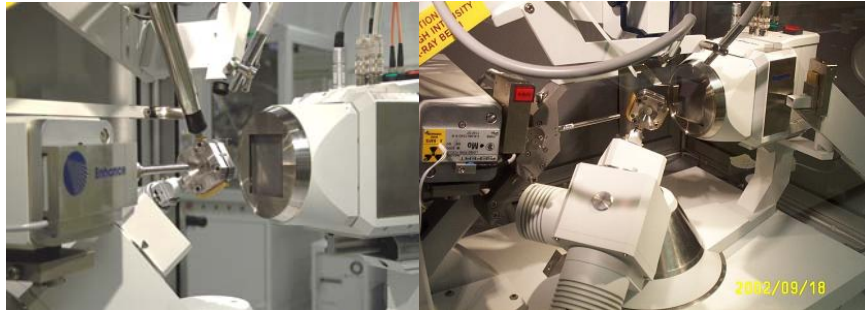


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Determining structures at high pressure

Precise intensity data to measure small changes in structures at high pressures:



Single-crystal X-ray diffraction.

Point and CCD detectors.

Beamlines and in-house

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Determining structures at high pressure

Very precise data to measure very small changes in structures at high pressures.

Single crystal diffraction is only worth the extra effort if it produces data significantly superior to powder diffraction.

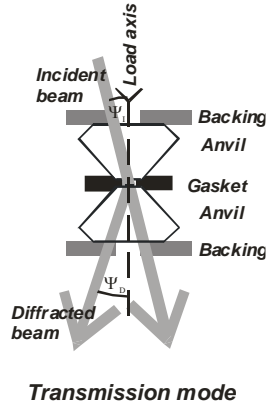
Challenges:

- **Limited access**
- **Low signal (but higher than powder)**
- **Background scattering**
- **Absorption by cell components**
- **Maintaining hydrostatic conditions**

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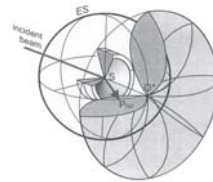
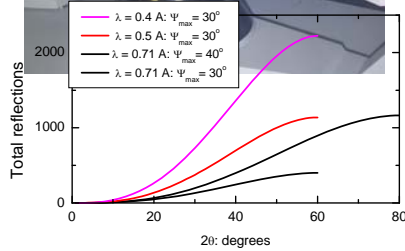
Diamond-anvil geometry



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Limited access



Miletich, RIM volume

Fewer reflections means:

- more sensitivity to individual data
- higher chance of systematic error
- higher influence of random error
- less resolution

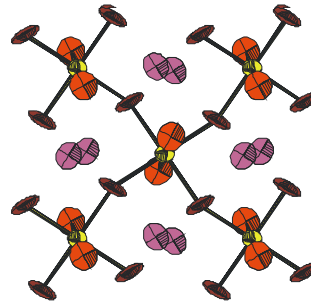
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Limited access

Lab: partially overcome by simulations to:

- predict the effects of restricted access
- choose best orientation

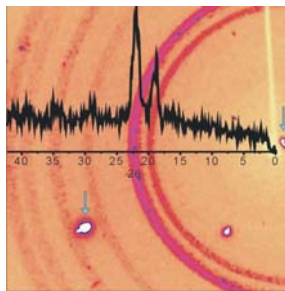


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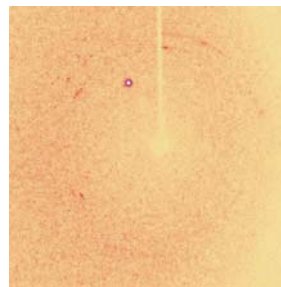
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Reducing background

Everything you see is background!



Conventional DAC with Be seats, steel gasket



DAC with steel seats and rhenium gasket

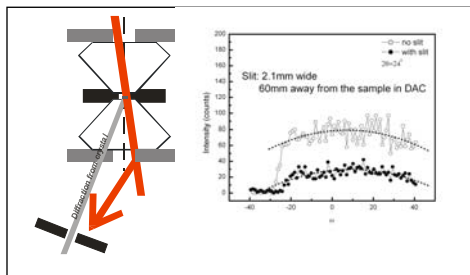
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Reducing background

Use a point detector:

- Additional collimation



- Optimised scan speed

- Step scans

- Profile fitting

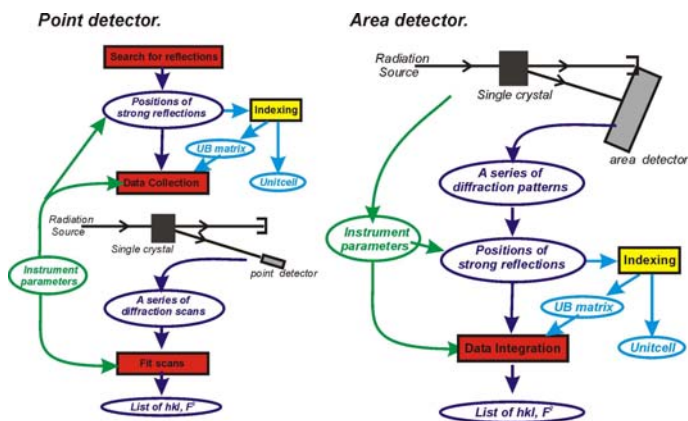
- Recovery of weak data



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Intensity data collection



Area detector is “shoot first and ask questions later”

Point detector needs more work at the diffractometer

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Intensity data integration with area detectors

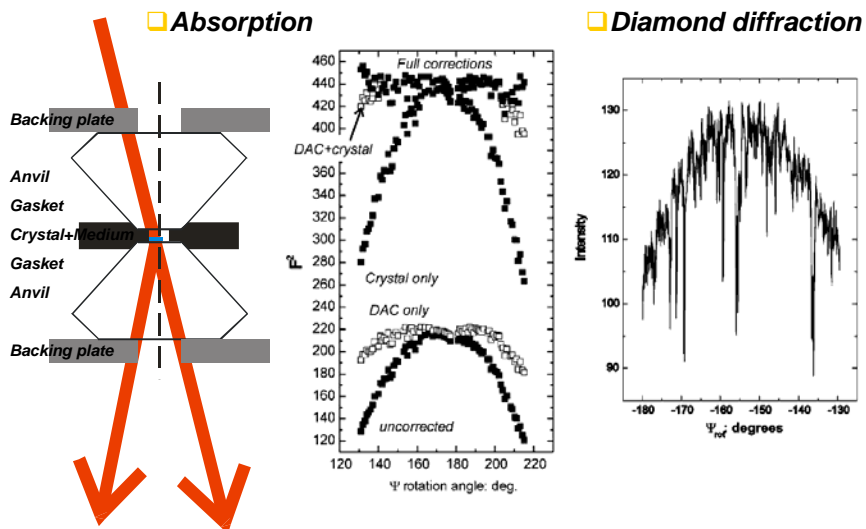
- Indexing
 - Identification of sample spots for indexing
 - Indexing algorithms to handle 2-d data
 - Masking of blank areas of detector
- Integration
 - Background subtraction: global + local
- Commercial and non-commercial software solutions available
- These processes can be automated



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Post-integration data corrections

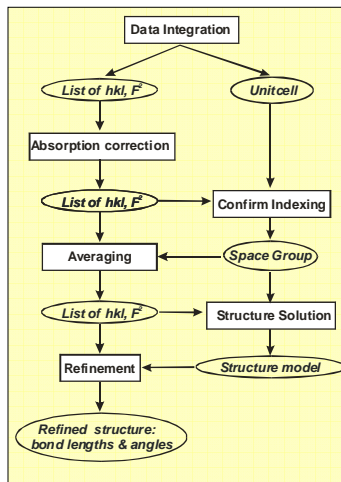


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Karen & Ross: tomorrow **VTX**

Post-integration steps

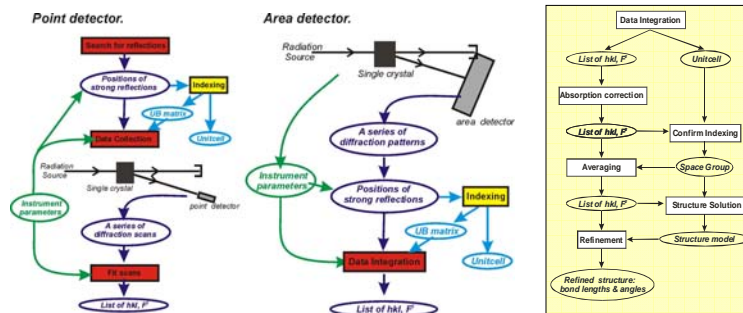
- Data reduction
 - Critical for HP data
 - Absorption corrections
 - Averaging
- Structure solution
 - Limited data coverage
 - Often not ab-initio
- Refinement
 - Robust-resistant weighting
 - Restraints
 - Careful evaluation of outliers
- Structure validation



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Karen, Ross & Tonci: tomorrow VTX

Workshop Overview



- Today
 - Experiment design
 - Data Collection
 - Data integration

- Tomorrow
 - Equations of State
 - Data reduction
 - Refinement
 - Structure validation
 - Problem solving

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