

Electron crystallography

3D electron diffraction for crystal structure determination

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Outline

- What is 3D ED
- From diffraction patterns to crystal structure
- Challenges in 3D ED:
 - Experimental techniques
 - Crystal tracking
 - Distortions in diffraction patterns
 - Dealing with dynamical diffraction effects
- Some examples and highlights

Electron crystallography

General definition: a scientific field which retrieves crystallographic information by using electrons as a radiation probe

In a stricter sense: crystal structure determination predominantly by means of electron diffraction

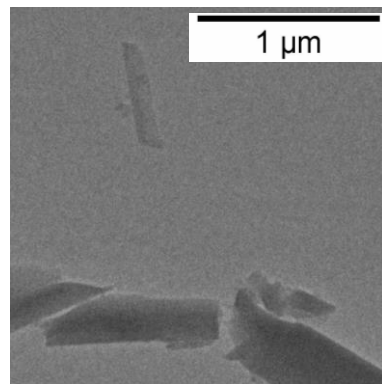
Information obtainable (in principle) from crystallographic investigation:

- crystal structure
- chemical composition
- polymorphism
- molecular connectivity
- molecular structure including absolute configuration
- nature of molecular species (salt/co-crystal)
- bonding ...

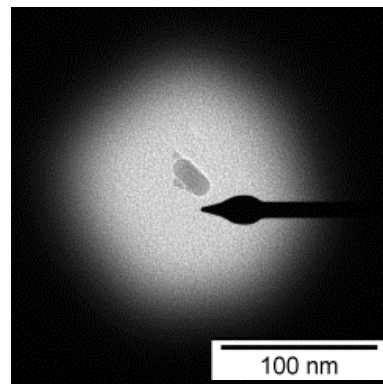
Electrons interact strongly with atoms

--> possibility to analyze small crystals.

For a crystallographer,
this is small



...and this is really
really small.



Diffraction and crystal structure

Basic relationship: diffracted intensity is proportional to the square of the amplitude of the diffracted wave:

$$I \propto A^2$$

The diffracted wave from a (small) crystal is proportional to the Fourier transform of the scattering power density in the unit cell of the crystal. This quantity is called the *structure factor*:

$$F_{\mathbf{h}} = \int_{V_{UC}} \rho(\mathbf{r}) \exp(2\pi i \mathbf{h} \cdot \mathbf{r}) dV$$

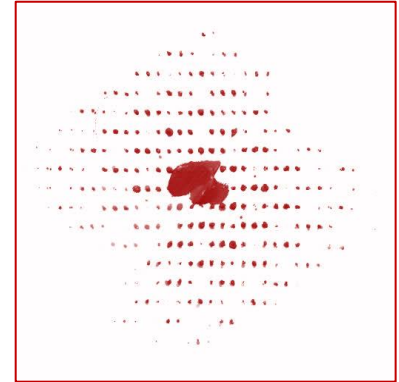
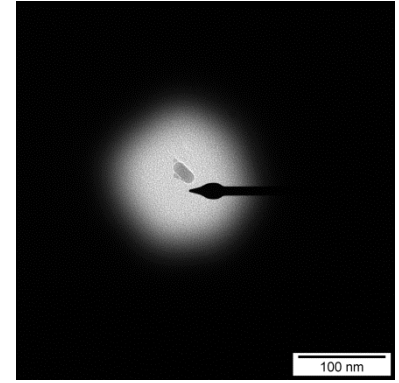
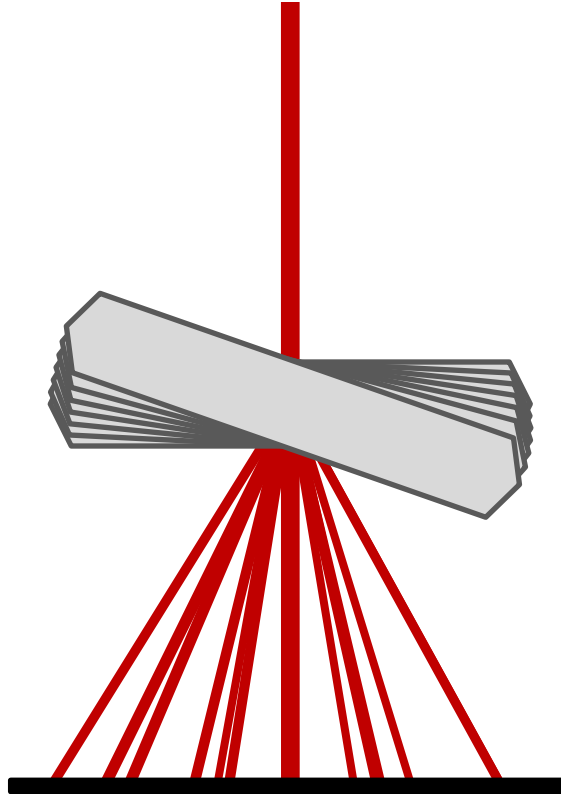
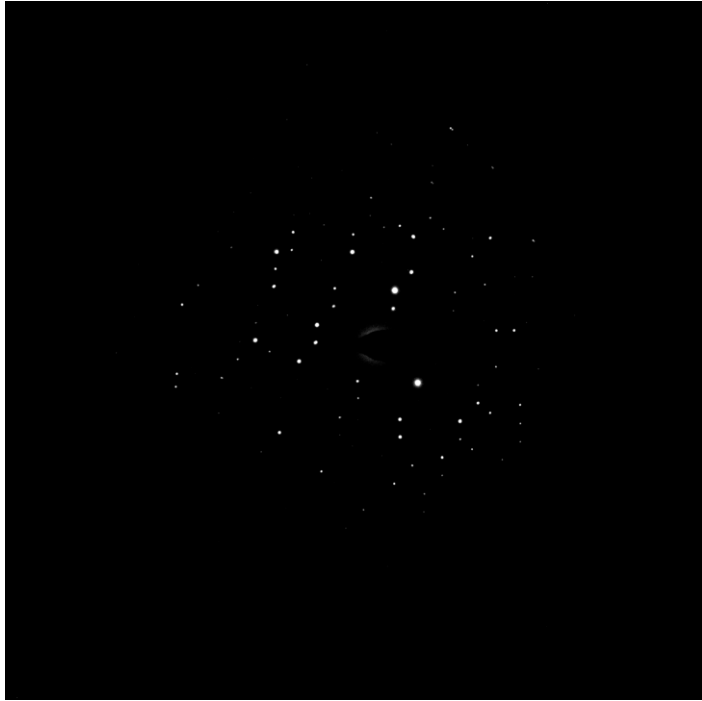
$$\mathbf{h} = h\mathbf{a}^* + k\mathbf{b}^* + l\mathbf{c}^*$$

Hence:
$$I_{\mathbf{h}} \propto |F_{\mathbf{h}}|^2$$

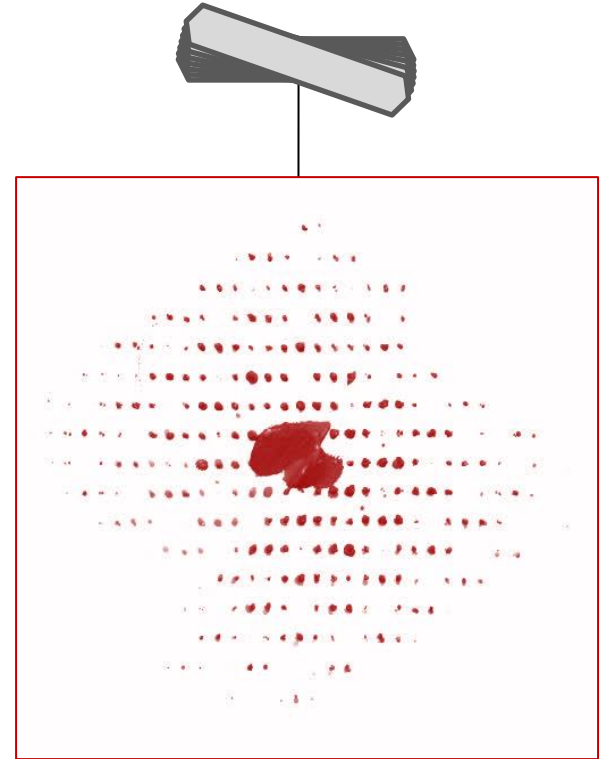
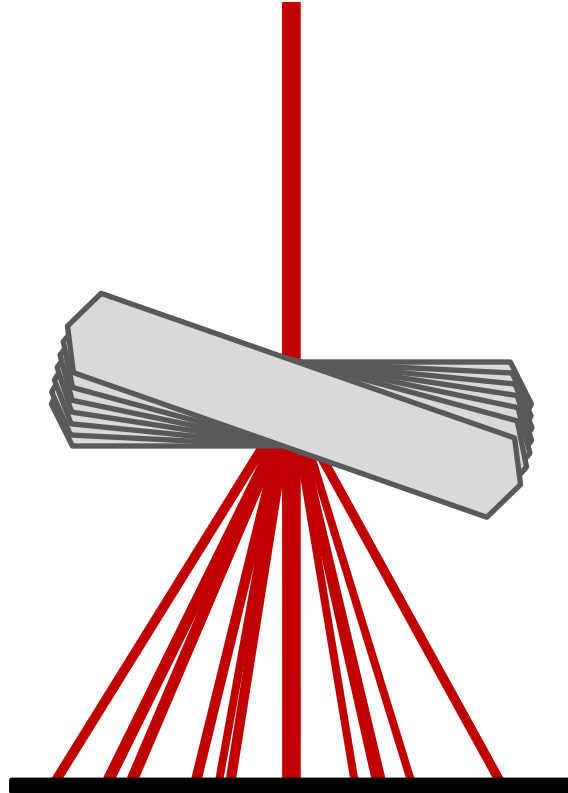
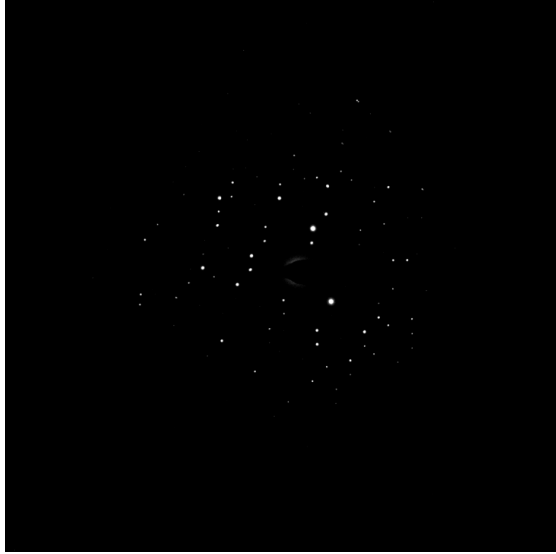
Basic assumption for the validity of the relationship: The crystal is very small. It is so small, that each quantum of radiation gets scattered at most once.

The theory based on this assumption is called the *kinematical diffraction theory*

3D electron diffraction (3D ED)



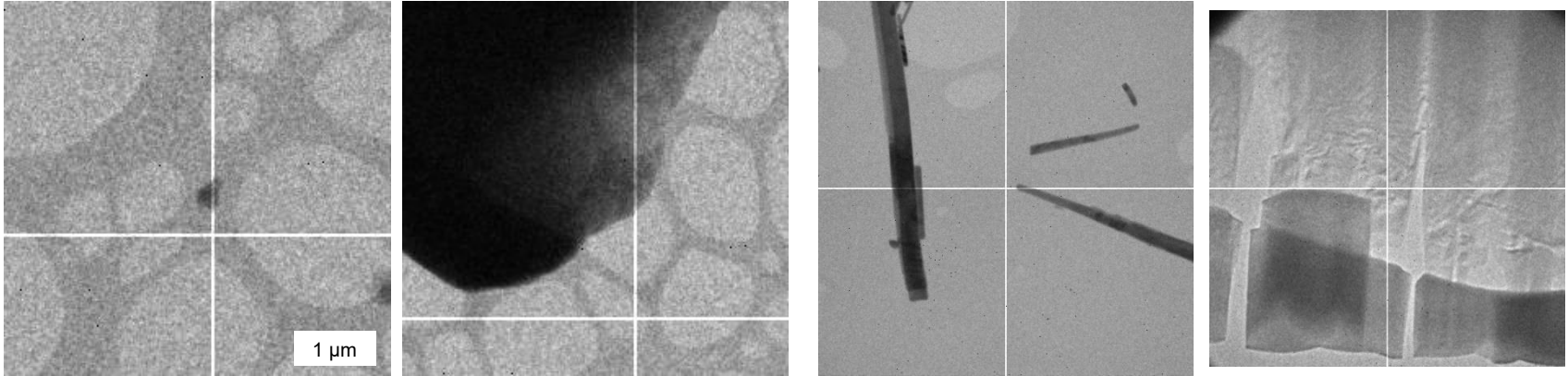
3D electron diffraction (3D ED)



From sample to crystal structure

Step 0: prepare the sample

You need small, but especially THIN crystals. Ideally $<100\text{nm}$



From sample to crystal structure

Step 1: collect data

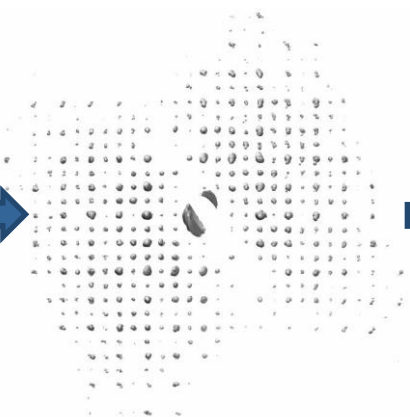
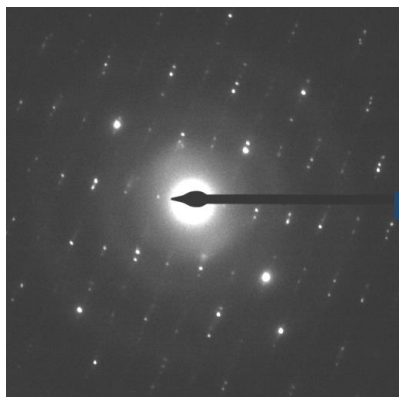
- option **a**: you have an adapted microscope with suitable data collection software. Well, then use it!
- option **b**: you have a dedicated electron diffractometer. Even better!
- option **c**: No special microscope, no special software. You can still collect the data manually.
Tedious, but works fine for beam-stable materials!

What microscope do you need:

- large tilt – at least $\pm 40^\circ$
- precession capability useful, but not necessary
- radiation hard detector, ideally with fast readout (not 100% necessary)
- no need for special features, no high resolution, no aberration correction, monochromator or filter...

From sample to crystal structure

Step 2: process the data



1	0	6	398.55	1.61
2	0	6	31.94	1.70
0	1	6	111.23	1.72
-2	2	6	17.69	0.86
4	-1	7	0.21	0.30
5	-1	7	0.08	0.72
1	0	7	21.21	0.52
2	0	7	74.14	1.59
3	0	7	16.37	2.11

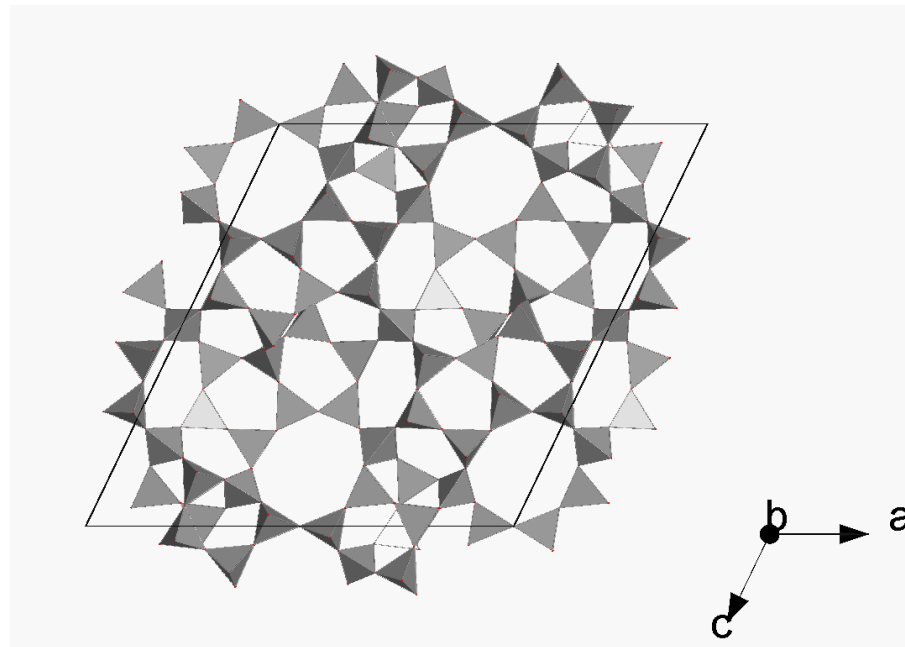
Several programs available for this purpose:

PETS2, XDS,
CrysAlis PRO
Dials, APEX4,
eADT

From sample to crystal structure

Step 3: solve the structure

1	0	6	398.55	1.61
2	0	6	31.94	1.70
0	1	6	111.23	1.72
-2	2	6	17.69	0.86
4	-1	7	0.21	0.30
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1	0	7	21.21	0.52
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3	0	7	16.37	2.11



Available software: SHELXS/T/D
Superflip
SIR
XLENS

From sample to crystal structure

Step 4: refine the structure

Structure refinement = optimization of structure model against experimental data.
Performed by least-squares minimization

$$S = \sum_{\mathbf{h}} w_{\mathbf{h}} \left(I_{obs}^{\mathbf{h}} - I_{calc}^{\mathbf{h}}(s, \{x, y, z, U\}) \right)^2$$

$$\Delta p = (J^T[w]J)^{-1} J^T[w] \Delta I$$

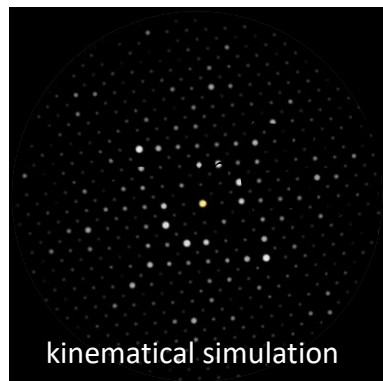
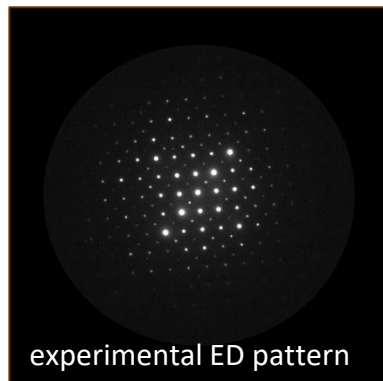
$$J_{ij} = \frac{\partial I_i(p_1 \dots p_n)}{\partial p_j}$$

The key problem: Calculation of $I_{calc}^{\mathbf{h}}$!

Option 1: Ignore multiple scattering =>
use kinematical diffraction theory =>
kinematical refinement

Option 2: Include multiple scattering =>
use dynamical diffraction theory =>
dynamical refinement

History of structure analysis by ED = history of fight with multiple scattering



Kinematical approximation:

$$I_{\mathbf{h}} \propto |F_{\mathbf{h}}|^2$$

Dynamical theory:

1) Find all reflections that contribute to diffraction

2) Build structure matrix \mathbf{A} :

$$a_{ii} = 2KS_{\mathbf{g}_i}, i = 1, N_{\text{beams}}$$

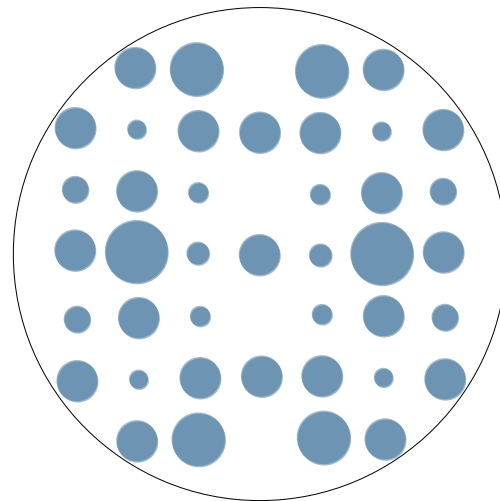
$$a_{ij} = U_{\mathbf{g}_i - \mathbf{g}_j}, i, j = 1, N_{\text{beams}}; i \neq j$$

3) Calculate scattering matrix \mathbf{S} :

$$\mathbf{S} = \exp\left(\frac{2\pi i t \mathbf{A}}{2K_n}\right)$$

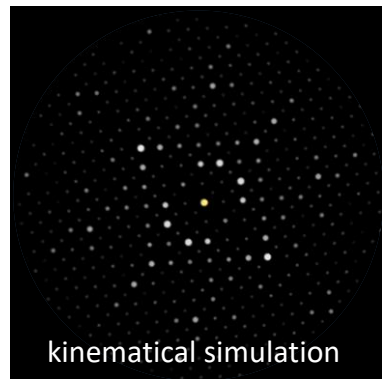
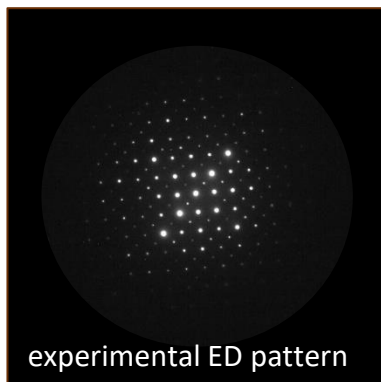
4) Calculate intensities from the first column of \mathbf{S} :

$$I_{\mathbf{h}_i} = |s_{i1}|^2$$



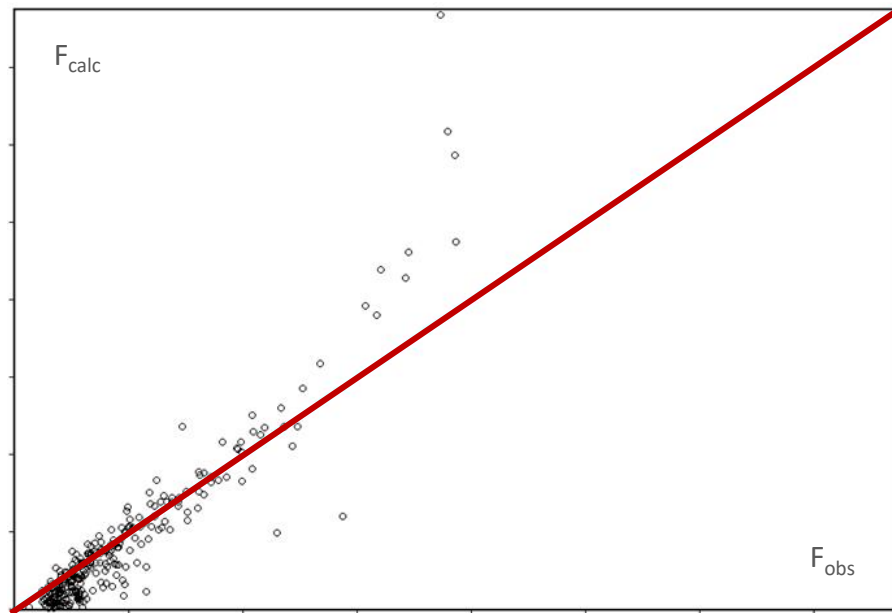
Dealing with the dynamical effects in the refinement

- 1) Ignore
- 2) Average out
- 3) Correct approximately
- 4) Take into account



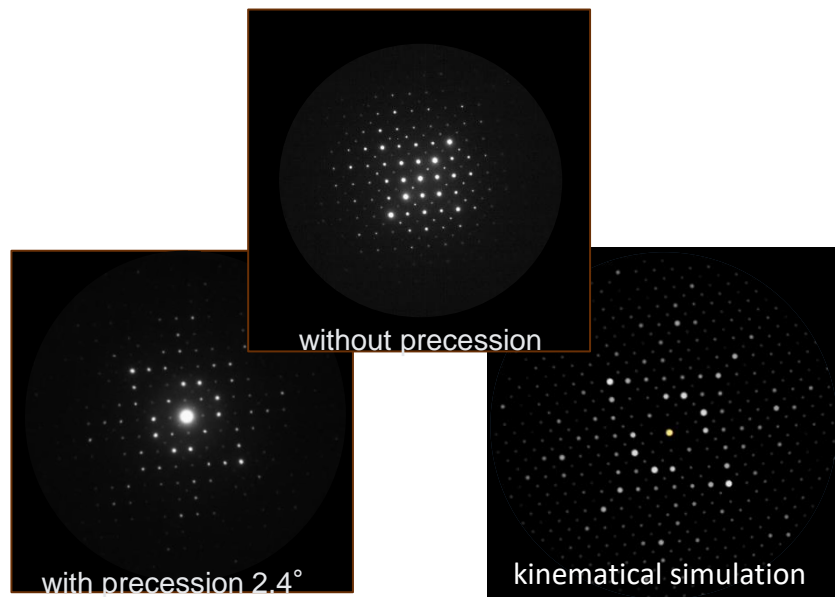
The easiest approach.

Leads to the highest figures of merit.
Potentially inaccurate structure models,
low sensitivity to details.



Dealing with the dynamical effects in the refinement

- 1) Ignore
- 2) Average out
- 3) Correct approximately
- 4) Take into account



Averaging helps a lot.

Dynamical effects critically depend on the crystal orientation and can vary sharply.

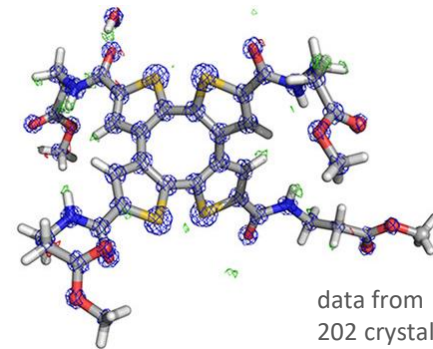
However, averaging does not remove the effects completely, no matter how much data you average.

Averaging options:

During data collection: continuous rotation or precession

Post data collection: averaging several datasets.

Averaging a large number of crystals is particularly useful

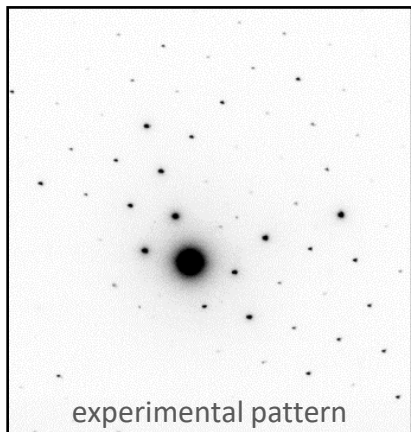


data from
202 crystals

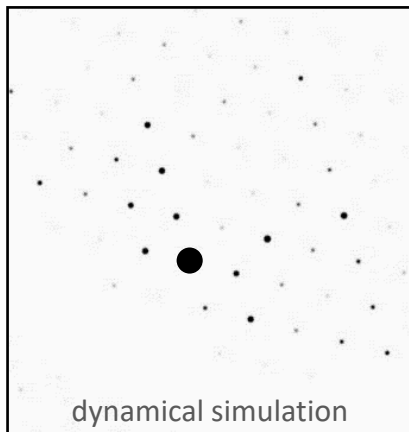
Dealing with the dynamical effects in the refinement

- 1) Ignore
- 2) Average out
- 3) Correct approximately
- 4) Take into account

This approach is called **dynamical refinement** for short and gives the best results in most cases.



experimental pattern



dynamical simulation

Downside: long(ish) computing times

- 1) Refinement uses frame-based intensities
- 2) Filtering of reflections based on their proximity to Ewald sphere
- 3) I_{calc} is a function of thickness and crystal orientation
- 4) Integration still necessary to limit sensitivity to crystal imperfections

The main challenges

fundamental

methods

technical

Size of
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Optical
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Beam sensitive materials

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Beam sensitive materials

Electrons are less damaging than x-rays per single elastic event.

However, the probed volume is much smaller in electron diffraction.

Therefore, **radiation damage is a much bigger issue for electron crystallography!**

What is “beam sensitive”? Indicative limiting doses for the loss of crystallinity

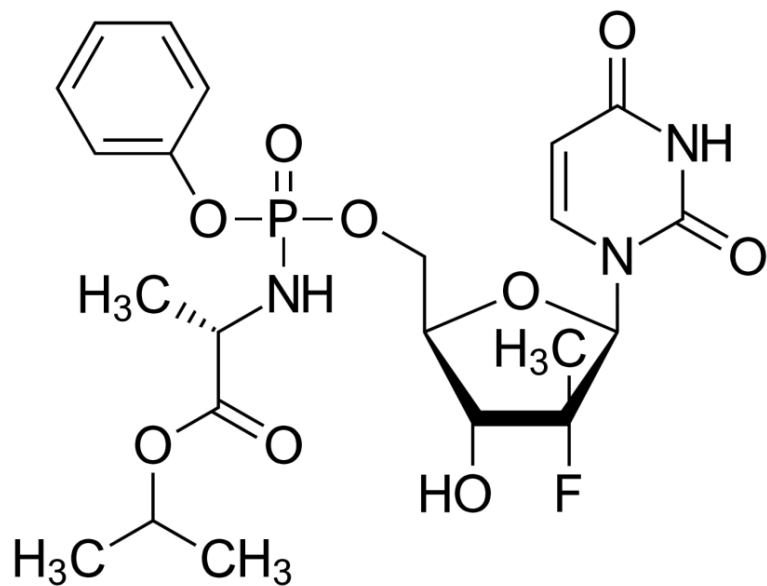
zeolites	$>100 \text{ e}/\text{\AA}^2$
MOFs	$5\text{-}15 \text{ e}/\text{\AA}^2$
protein crystals	$1\text{-}10 \text{ e}/\text{\AA}^2$
crystals of small organic molecules with hydrogen bonds	$0.5\text{-}10 \text{ e}/\text{\AA}^2$
crystals of small organic aliphatic molecules with van der Waals bonds only	$0.01\text{-}0.5 \text{ e}/\text{\AA}^2$

SOLUTION:

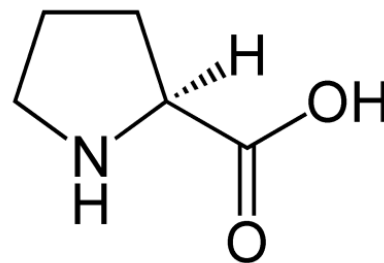
- 1) Use fast data collection with the modern sensitive direct detection cameras
- 2) Collect data on different parts of a large(r) crystal or use serial electron crystallography

Sofosbuvir L-prolin

a cocrystal of L-proline and an anti-hepatitis drug (both chiral)



Sofosbuvir – antiviral



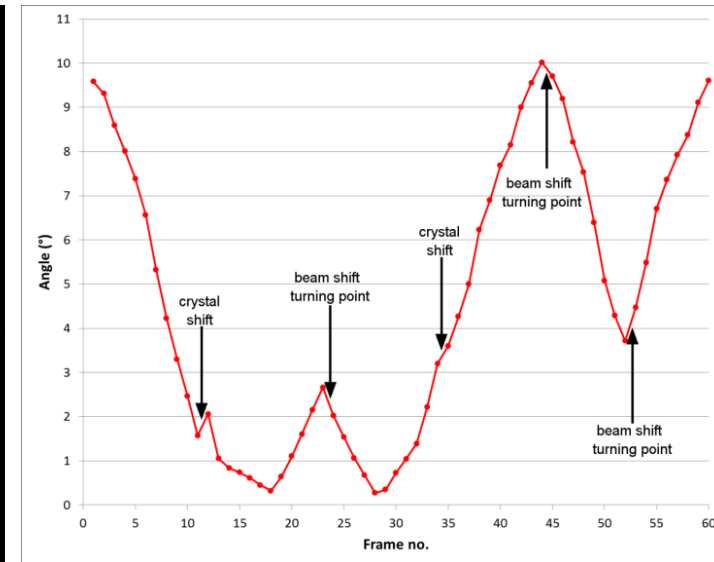
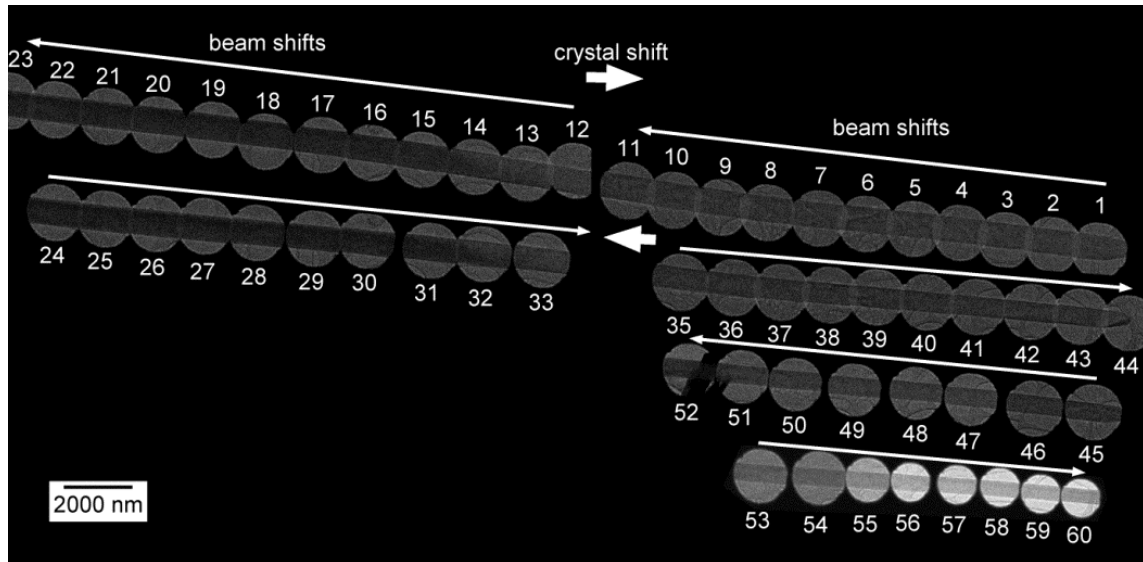
L-proline – amino acid

Sofosbuvir L-prolin

A cocrystal of L-prolin and an anti-hepatitis drug.

Extremely beam-sensitive, most crystals deteriorate after $<0.08 \text{ e}/\text{\AA}^2$.

Crystals form long ribbons.

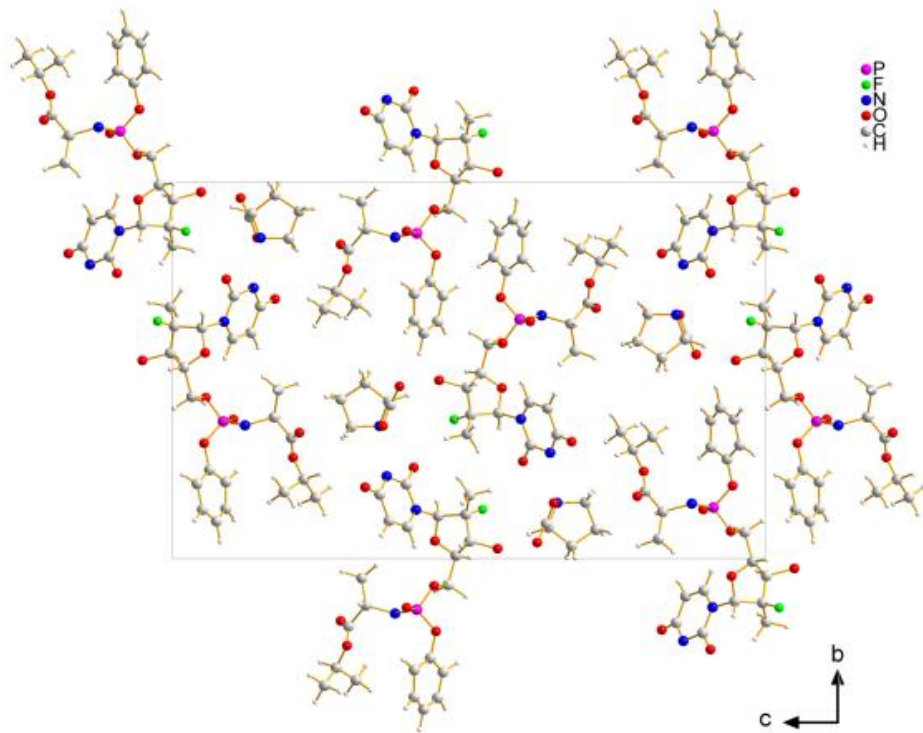


Sofosbuvir L-prolin

A cocrystal of L-prolin and an anti-hepatitis drug.
Extremely beam-sensitive, most crystals deteriorate after $<0.08 \text{ e}/\text{\AA}^2$.
Crystals form long ribbons.



Sofosbuvir L-prolin



Formula: $C_{27}H_{38}N_4O_{11}FP$
44 independent non-H atoms

Space group $P2_12_12_1$

$a=5.35\text{\AA}$, $b = 19.60\text{\AA}$, $c = 29.87\text{\AA}$

Unit cell volume: 3127 \AA^3

Kinematical refinement	$R_{\text{obs}} = 19.7\%$
Dynamical refinement	$R_{\text{obs}} = 9.7\%$

Data collection techniques

fundamental

methods

technical

Size of
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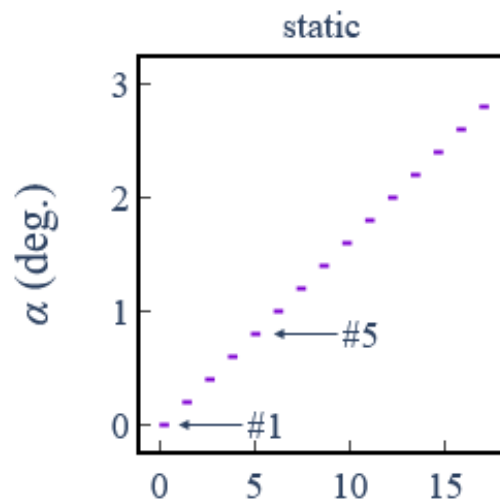
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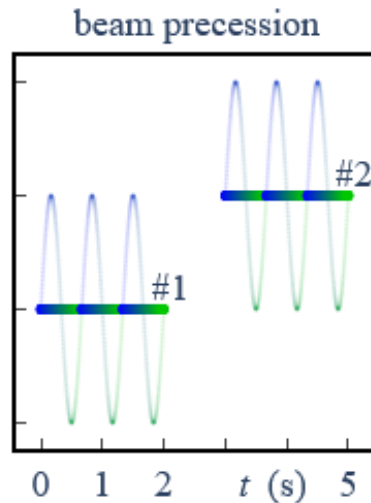
Optical
stability
and
accuracy

3D ED variants



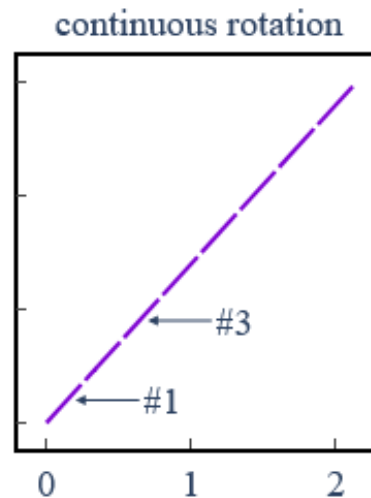
static-beam 3D ED

ADT/EDT
(Automated/Electron
Diffraction Tomography)
MicroED (version from 2013)



precession-assisted 3D ED

PEDT
(Precession El. Diffr. Tomography)
precession ADT



continuous-rotation 3D ED

cRED (continuous RED)
IEDT (integrated EDT)
MicroED (current version)

Crystal tracking

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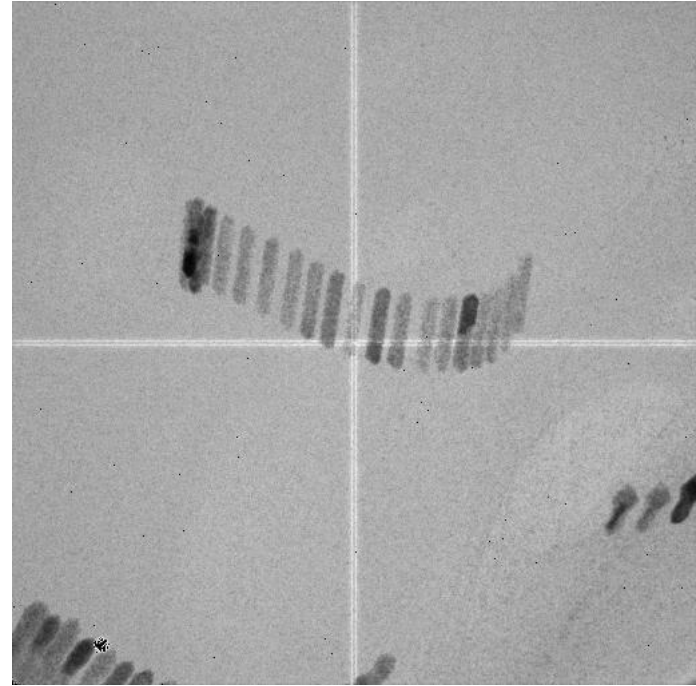
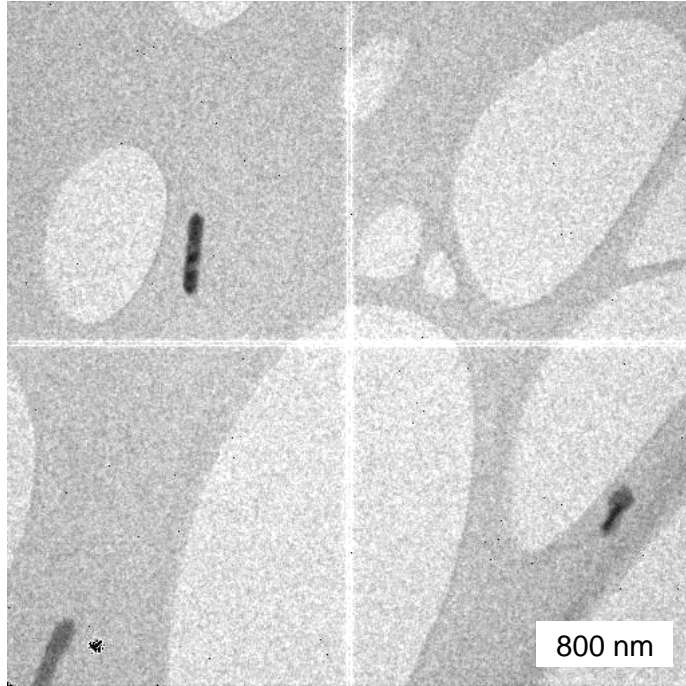
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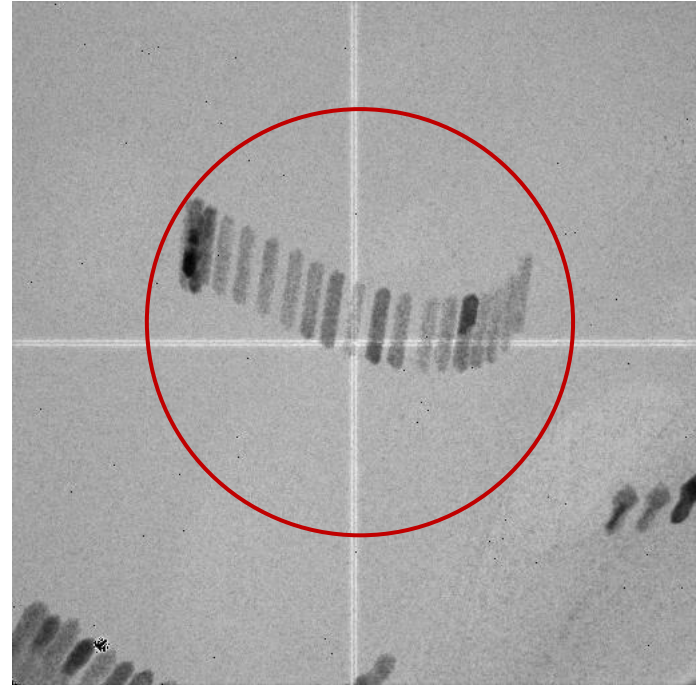
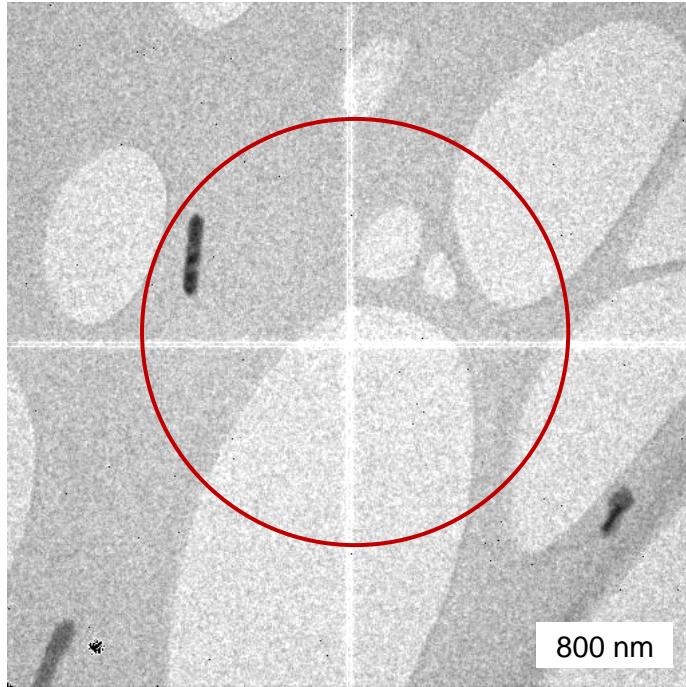
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Crystal tracking

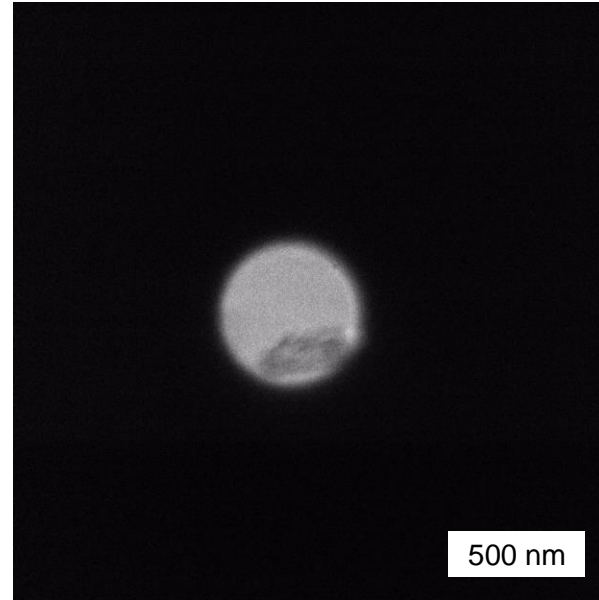
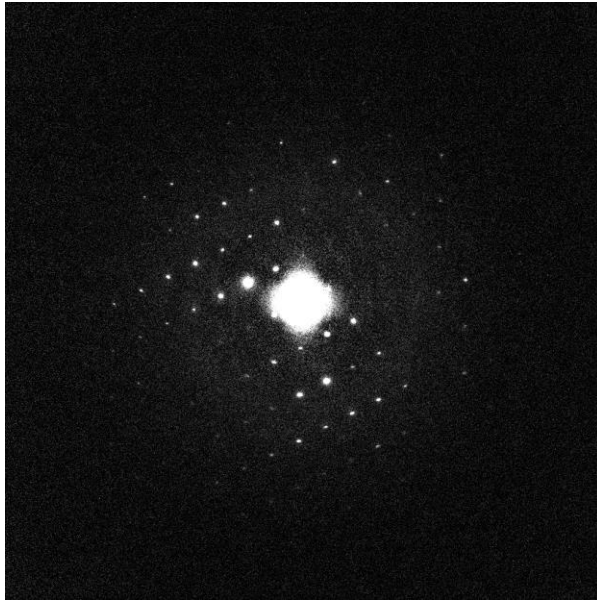


Crystal tracking



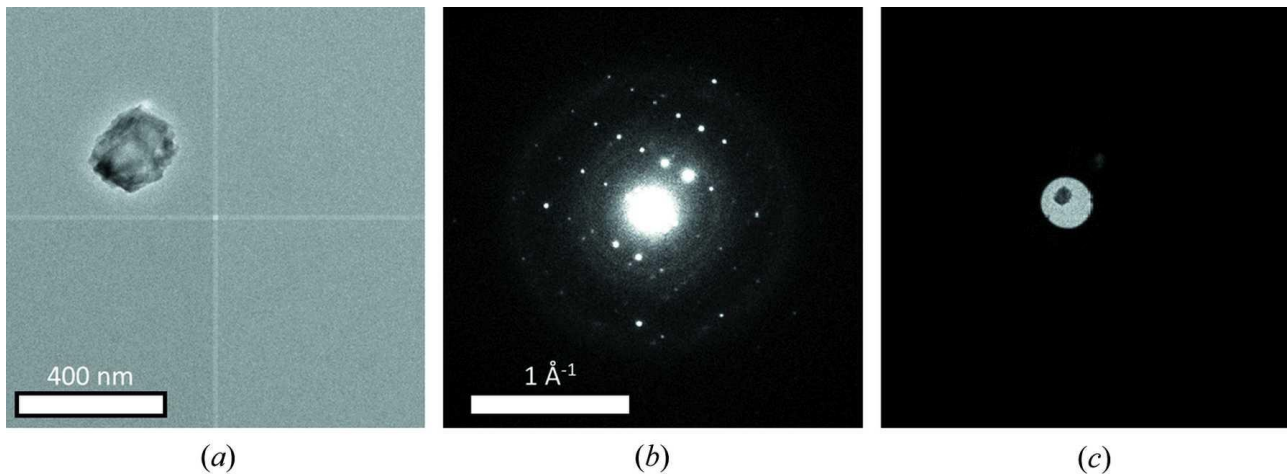
Option 1: use a large beam. Downside: high background, illumination of more than one crystal, varying illumination of large crystals

Crystal tracking



Option 2: switch from diffraction to imaging to track the crystal. Downside: lens hysteresis, incompatible with continuous rotation

Crystal tracking



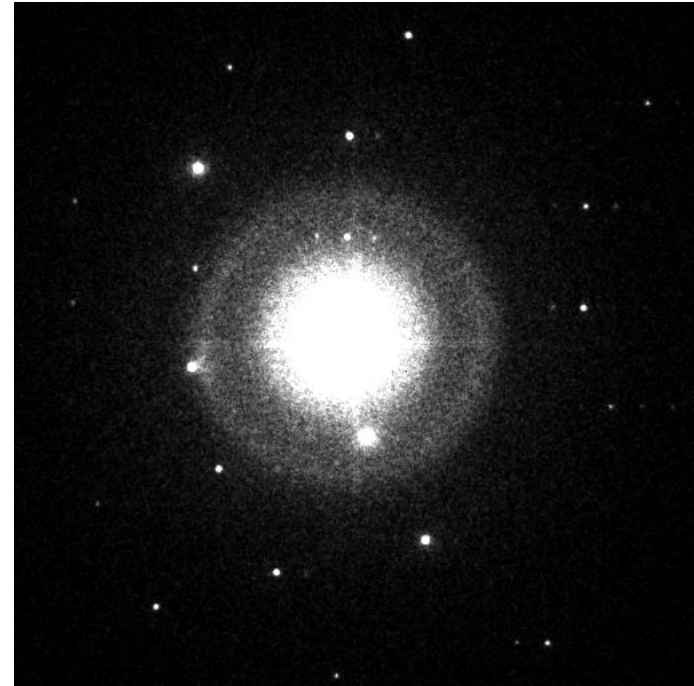
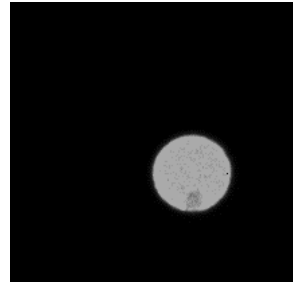
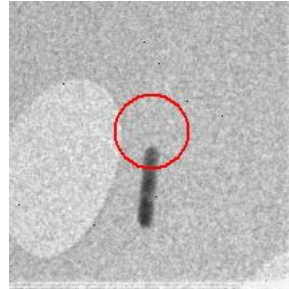
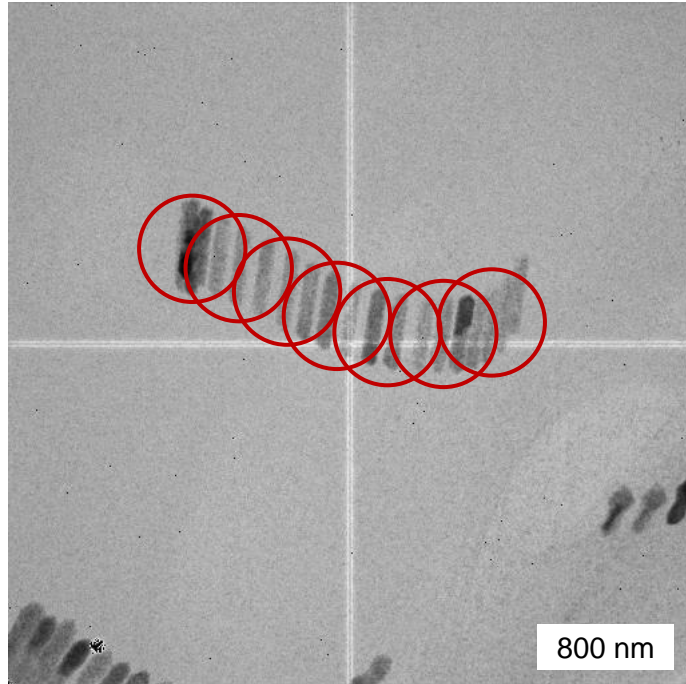
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APPLIED
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• Cichocka *et al.*

• Volume 51 | Part 6 | December 2018 | Pages 1652–1661 | 10.1107/S1600576718015145

Option 3: defocus the diffraction to see the crystal in the central beam. Downside: lens hysteresis, gaps in the data

Crystal tracking



Option 4: pre-record the path of the crystal, then track it using the pre-recorded path. Downside: additional time (not much), additional illumination of the crystal (not much, may be avoided), potentially not perfectly reproducible path

Optical distortions and lattice parameters

fundamental

methods

technical

Size of
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crystal

Size of
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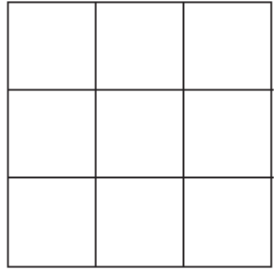
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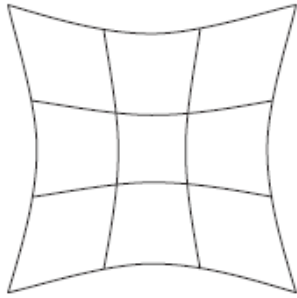
Optical
stability
and
accuracy

Optical distortions and lattice parameters

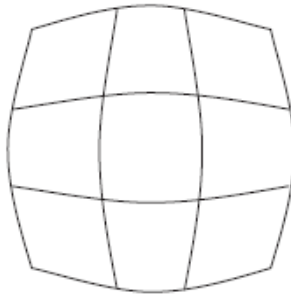
undistorted image



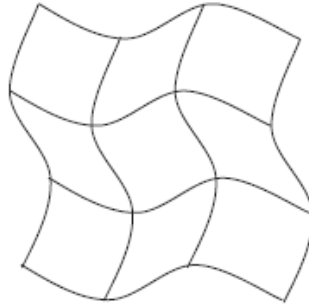
pincushion distortion



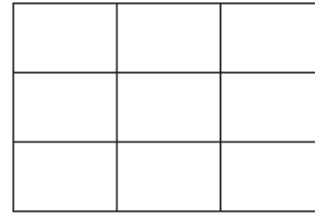
barrel distortion



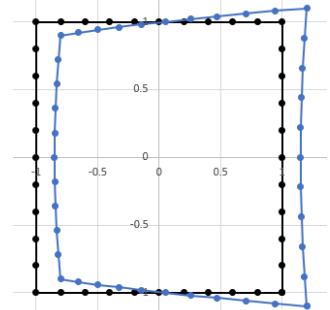
spiral distortion



elliptical distortion



parabolic distortion



Optical distortions and lattice parameters

Example:

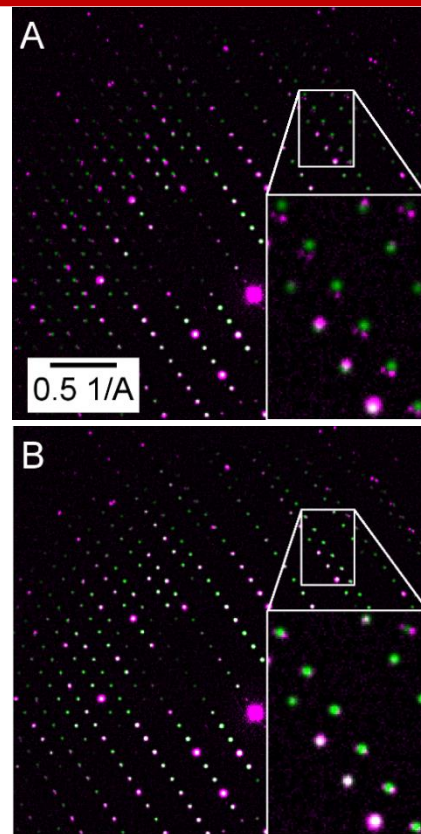
- Lutetium aluminium garnet, Bravais class cF, $a=11.9084 \text{ \AA}$
- Main reason for distorted unit cell: elliptical distortion

Without distortion correction:

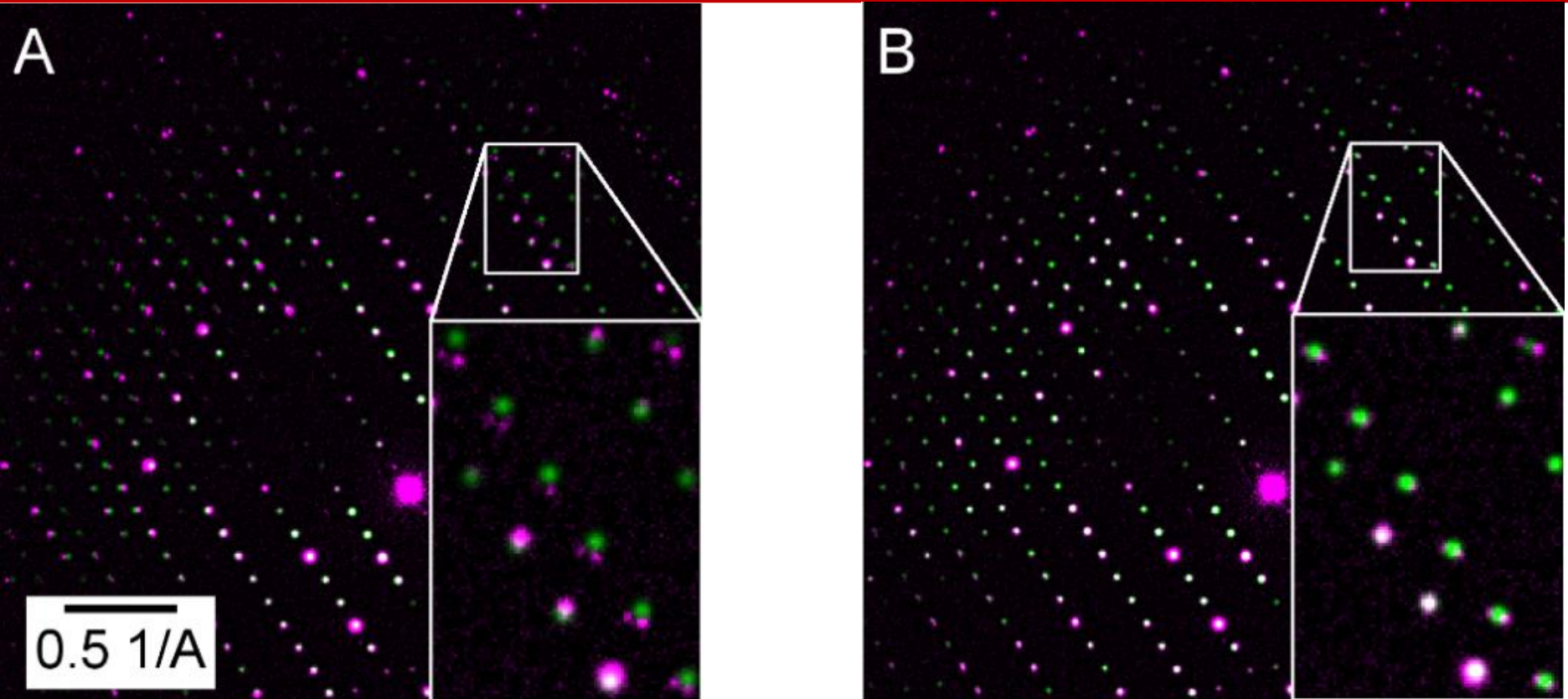
	a (Å)	b (Å)	c (Å)	alpha (°)	beta (°)	gamma (°)
DS1	11.848(1)	11.858(3)	11.907(1)	90.21(1)	90.33(1)	89.97(1)
DS2	11.842(1)	11.881(2)	11.888(2)	89.77(2)	89.75(1)	89.75(1)
DS3	11.854(1)	11.863(3)	11.890(1)	89.80(1)	90.40(1)	90.11(1)

With distortion correction:

	a (Å)	b (Å)	c (Å)	alpha (°)	beta (°)	gamma (°)
DS1	11.909(1)	11.910(2)	11.908(1)	90.00(1)	90.00(1)	90.02(1)
DS2	11.908(1)	11.914(2)	11.908(2)	90.03(2)	90.00(1)	89.98(1)
DS3	11.907(1)	11.912(2)	11.909(1)	90.001(3)	90.00(1)	90.00(1)



Optical distortions and lattice parameters



Calculation of intensities

fundamental

methods

technical

Size of
the
crystal

Size of
the
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Symm-
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analysis

Beam
sensit-
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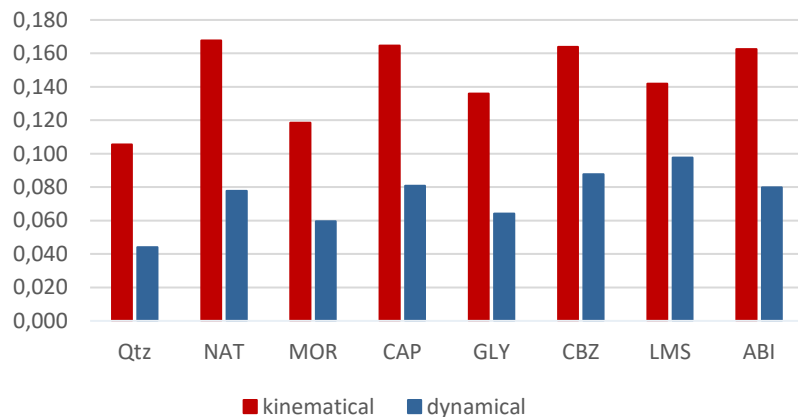
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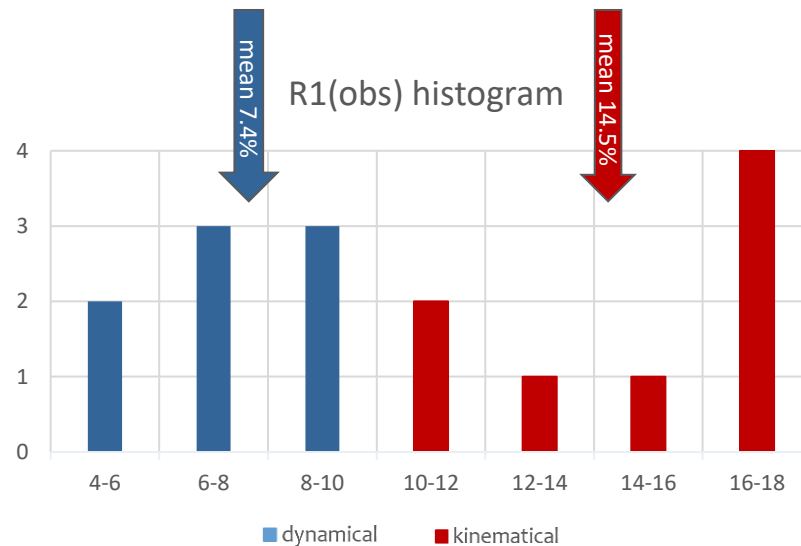
The R-factor gap in electron crystallography

8 structures carefully refined kinematically and dynamically from the same data

R1(obs) kinematical vs dynamical



R1(obs) histogram



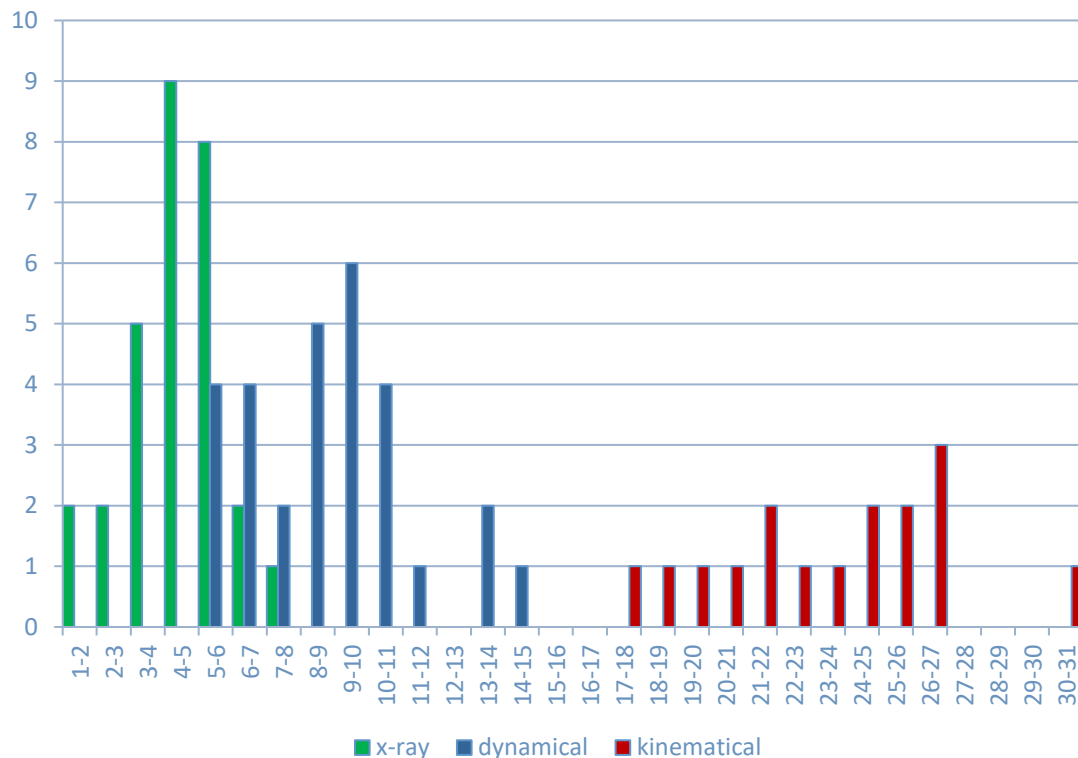
Dynamical: refined against unmerged data, but R-factors calculated on merged data for comparison with the kinematical factors.

The R-factor gap in electron crystallography

The fit to 3DED data is not as good as for x-ray data even for dynamical refinement.

Reason: diffraction comes from imperfect, often irregular crystals, but theory assumes perfect crystals

Solution: modify the refinement to account for crystal imperfections



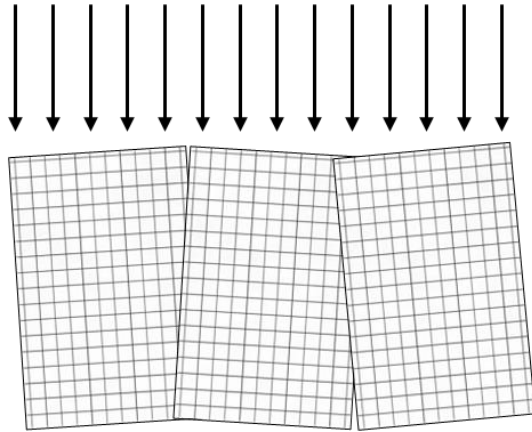
Mosaicity in the dynamical case

Aspects to consider:

- The intensity depends on the crystal orientation
- Electrons passing from one mosaic block to the next „know“ their history
- Column approximation may come to the rescue

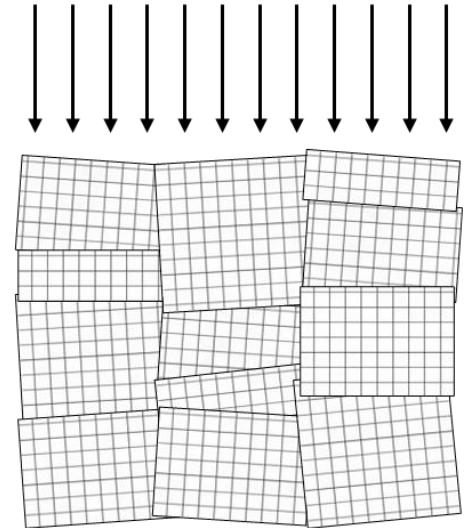
Case 1: large blocks

Incoherent superposition of diffraction from individual blocks



Case 2: small blocks

Coherent superposition of diffraction from individual blocks in each column + incoherent superposition of individual columns

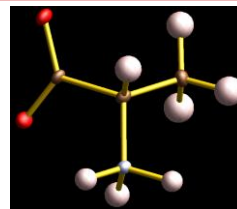


Mosaicity in the dynamical case

L-alanine: simple organic molecule

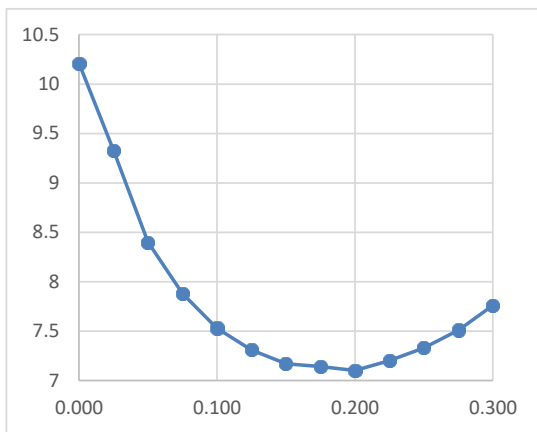
orthorhombic, $P2_12_12_1$

relatively high mosaicity of 0.29° (from data reduction)



Test 1: Incoherent isotropic mosaicity

0.000	10.21
0.025	9.33
0.050	8.4
0.075	7.88
0.100	7.53
0.125	7.31
0.150	7.17
0.175	7.14
0.200	7.1
0.225	7.2
0.250	7.33
0.275	7.51
0.300	7.76



Test 2: Combined coherent and incoherent mosaicity

		μ_i										
		0.000	0.025	0.050	0.075	0.100	0.125	0.150	0.175	0.200	0.225	0.250
μ_o	0.00	10.18	9.33	8.40	7.95	7.58	7.36	7.25	7.18	7.27	7.32	7.51
	0.05	10.02	9.15	8.33	7.90	7.54	7.32	7.22	7.16	7.24	7.29	7.49
	0.10	9.62	8.84	8.15	7.78	7.49	7.28	7.18	7.13	7.21	7.26	7.46
	0.15	9.07	8.48	7.96	7.66	7.41	7.22	7.14	7.09	7.16	7.23	7.42
	0.20	8.57	8.16	7.78	7.54	7.32	7.16	7.10	7.08	7.14	7.23	7.42
	0.25	8.24	7.94	7.63	7.42	7.24	7.11	7.08	7.07	7.12	7.24	7.44
	0.30	7.94	7.71	7.47	7.32	7.17	7.09	7.09	7.10	7.15	7.30	7.52
	0.35	7.69	7.54	7.38	7.29	7.19	7.16	7.16	7.18	7.25	7.43	7.67
	0.40	7.56	7.47	7.37	7.31	7.28	7.31	7.31	7.34	7.43	7.64	7.85

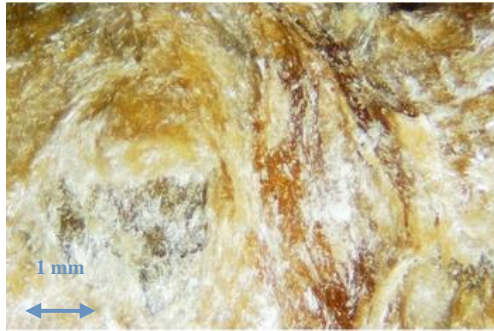
$\mu_{comb} = \mu_i^2 + \mu_o^2 = 0.31^\circ$, very close to the experimental value

Example – eveslogite



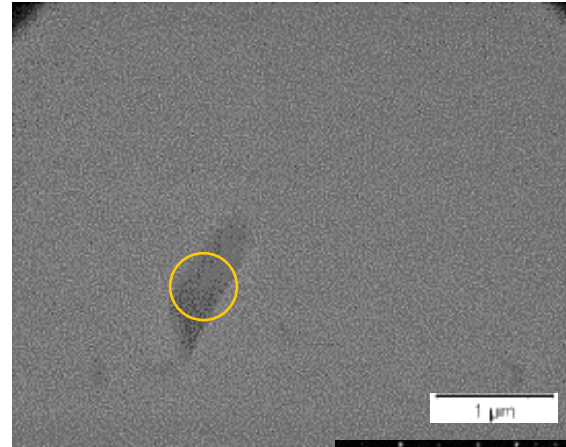
Eveslogite, etc.

Fersman Gorge, Eveslogchorr Mt, Khibiny Massif, Murmansk Oblast, Russia



Eveslogite

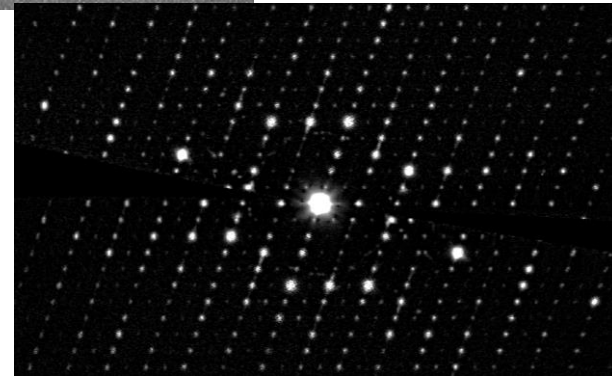
Eveslogchorr Mt, Khibiny Massif, Murmansk Oblast, Russia



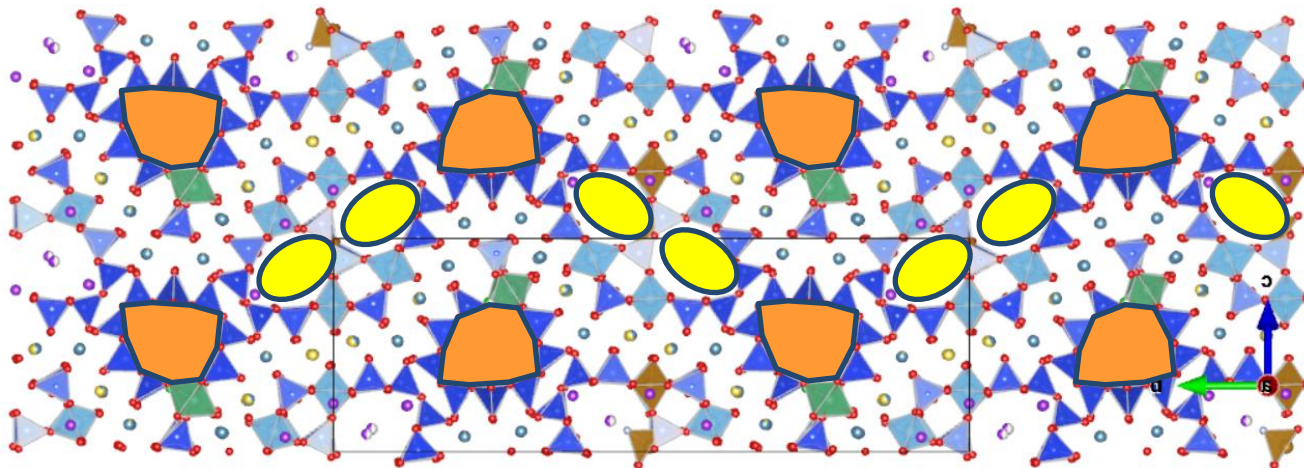
Chemical Properties of Eveslogite

Formula: $(\text{Na}, \text{K}, \text{Ca}, \text{Sr}, \text{Ba})_{48} [(\text{Ti}, \text{Nb}, \text{Mn}, \text{Fe}^{2+})_{12} \text{Si}_{48} \text{O}_{144} (\text{OH})_{12}] (\text{F}, \text{OH}, \text{Cl})_{14}$

IMA Formula: $(\text{Ca}, \text{K}, \text{Na}, \text{Sr}, \text{Ba})_{48} (\text{Ti}, \text{Nb}, \text{Fe}, \text{Mn})_{12} (\text{OH})_{12} \text{Si}_{48} \text{O}_{144} (\text{OH}, \text{F}, \text{Cl})_{14}$



Example – eveslogite



space group $P2_1$
 a 14.1898 Å
 b 44.7704 Å
 c 15.9111 Å
 β 109.4677°
 V_{uc} 9530.171 Å³

~ 360 atoms in the asymmetric unit

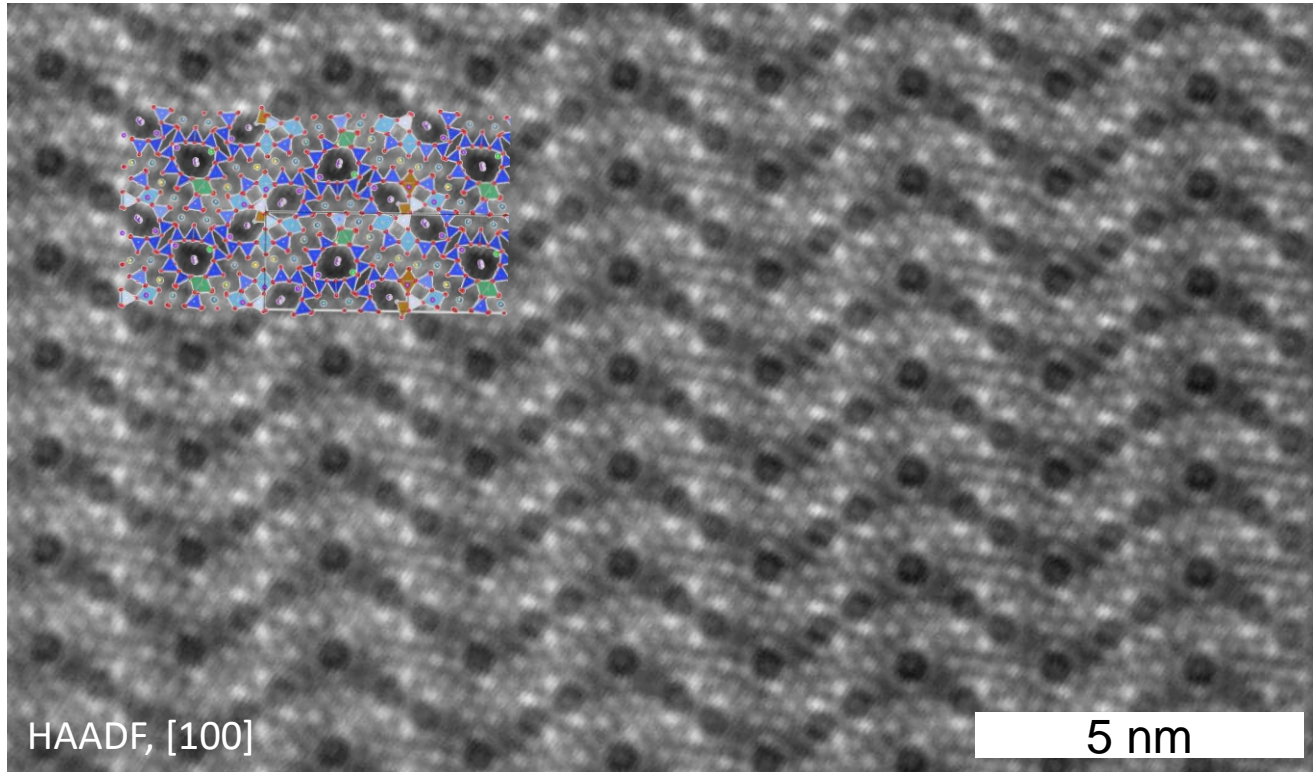
115 842 reflections $R_{obs} = 19.82$

$R_{int} = 17.24$

$R_{all} = 24.70$

Na	
K	
Ca	
Nb	
Ti	
Si	

Example – eveslogite

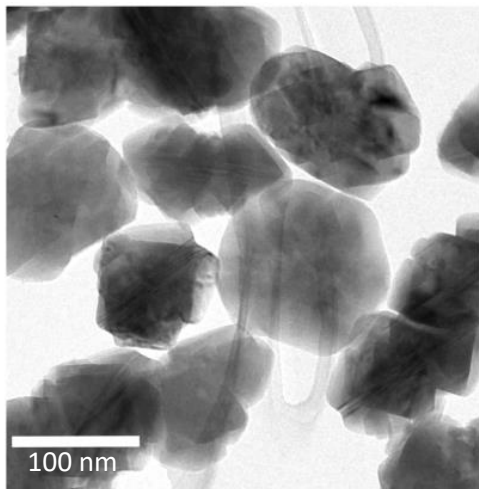


Example – CO₂-loaded zeolite

CO₂-loaded CHABAZITE

Synthesis of an industrially important zeolite in nanocrystalline form without OSDA.

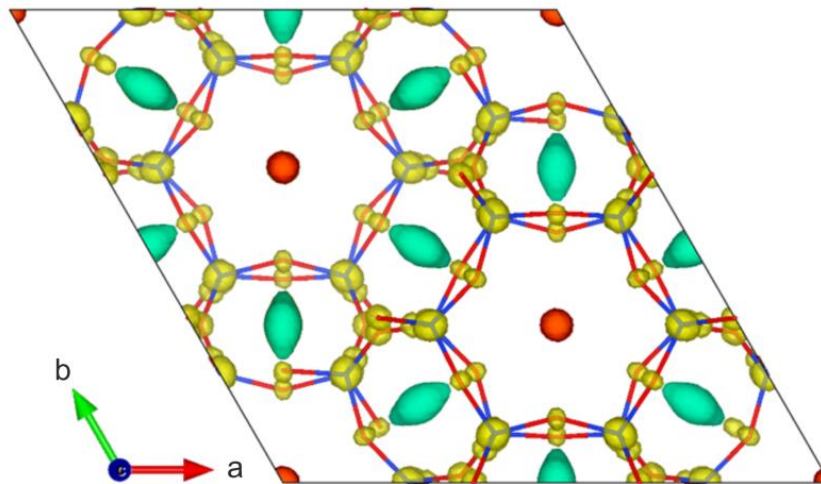
Extra-framework cations: Na⁺, K⁺, Cs⁺.



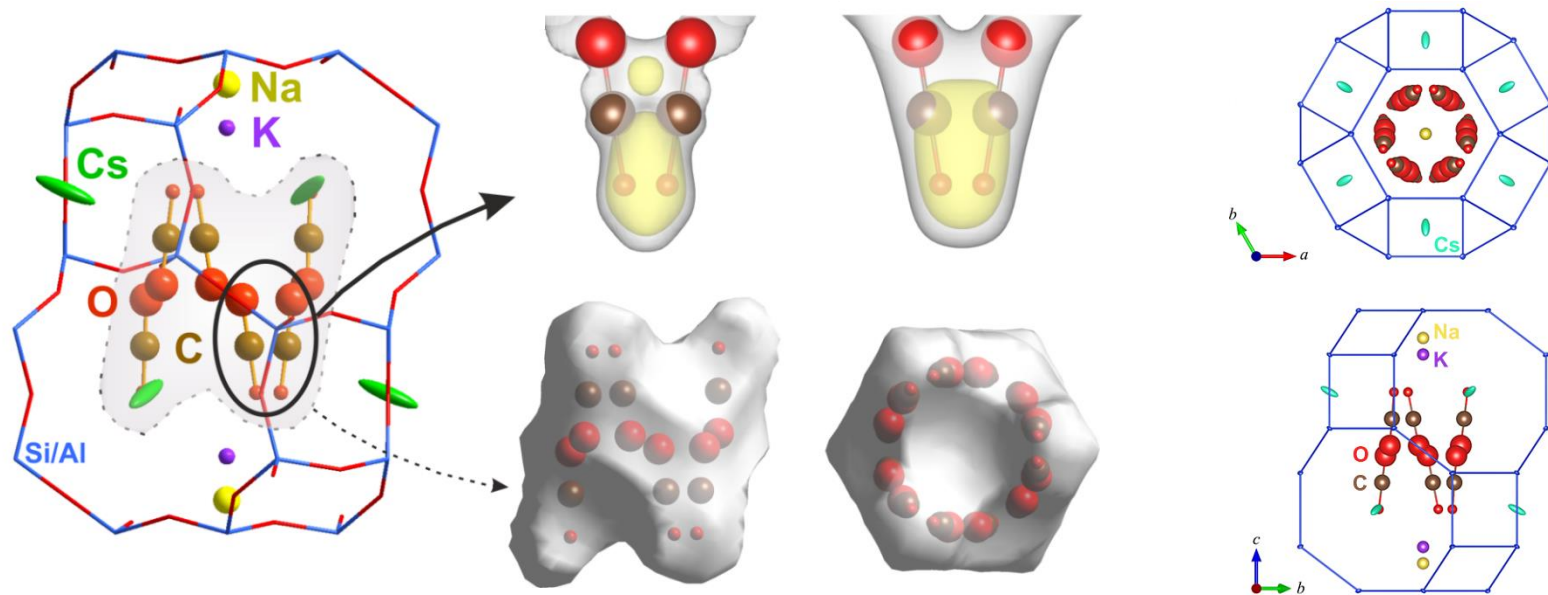
Chabazite has a very good CO₂ adsorption and selectivity towards CH₄.

Crystallographic question:

Can we locate the CO₂ molecules in the chabazite structure?



Example – CO₂-loaded zeolite



Theoretical maximum adsorption capacity: 9 CO₂ molecules per unit cell

Experimental adsorption capacity: 8 CO₂ molecules per unit cell

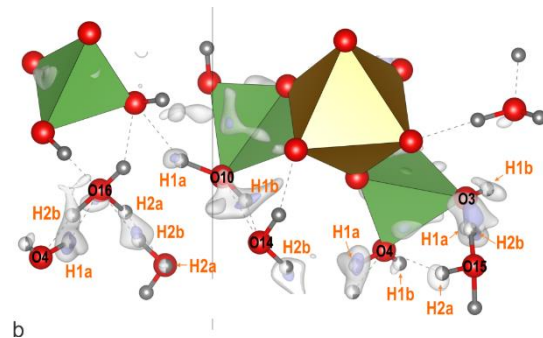
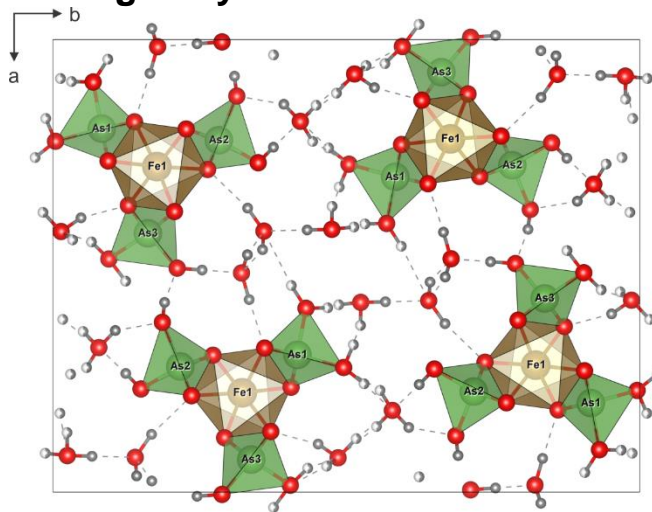
Cationic composition is crucial for the capacity and selectivity of CO₂ adsorption

Example – disordered hydrogen atoms



Hydrogen disorder in kaatialaite $\text{Fe}[\text{AsO}_2(\text{OH})_2] \cdot 5\text{H}_2\text{O}$

The structure of synthetic kaatialaite known (Boudjada & Guitel, 1981) but the hydrogen sites remained **undetected** from X-ray single-crystal data.



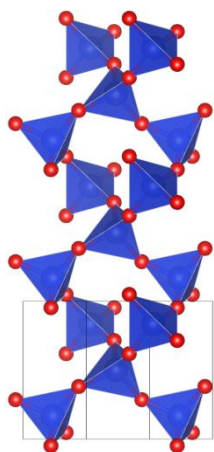
Map after dynamical refinement of the structure including the non-disordered hydrogen

22 independent hydrogen positions, out of them 12 disordered

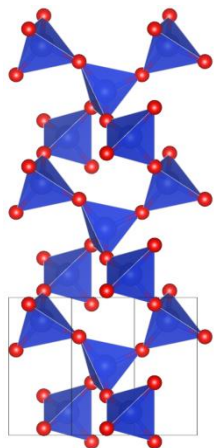
Absolute structure and absolute configuration

Absolute structure is a specification of the orientation of a non-centrosymmetric crystal structure under the operation of inversion (Online dictionary of Crystallography)

A non-centrosymmetric crystal may or may not be composed of chiral species.

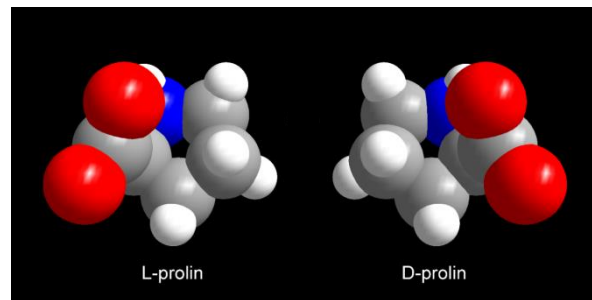


left quartz ($P3_121$)



right quartz ($P3_221$)

Absolute configuration is a specification of the spatial arrangement of atoms in a molecule containing *chiral centers*. Such molecules are not superimposable onto their mirror images. Different absolute configuration may mean (and often means) different biological function of the molecule.



Absolute configuration of molecules is most reliably and most often determined by determining the **absolute structure** of crystals containing the molecule.

Absolute structure and absolute configuration

Breaking Friedel's law

Friedel's law: In kinematical approximation, opposite structure factors have equal amplitudes: $|F_{-h}| = |F_h|$
Consequence: It is impossible to determine absolute structure from kinematical diffracted intensities

X-rays:

Resonant scattering shifts the phase of scattered photons from atoms \rightarrow Friedel's law does not hold exactly.

Strength of resonant scattering depends on the degree of non-centrosymmetry, on the wavelength and atomic number.

Light atoms have very low resonant scattering \rightarrow difficulties in determination of absolute structure of organic species.

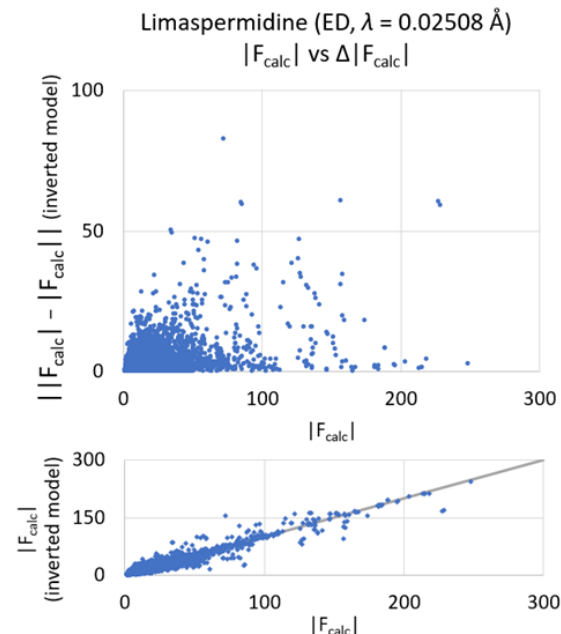
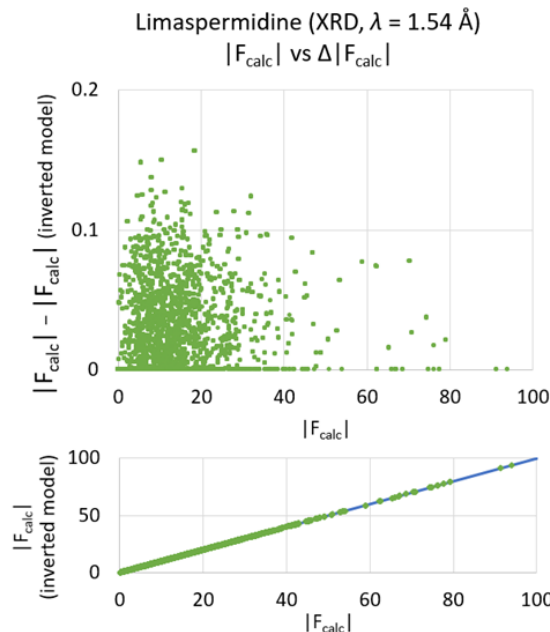
Electrons:

Electron diffraction is dynamical = coherent interference of more than one diffracted beam \rightarrow Friedel's law does not hold.
In three-beam approximation:

$$I_h - I_{-h} \propto F_h F_g F_{h-g} \sin\varphi$$

where φ is the sum of structure factor phases

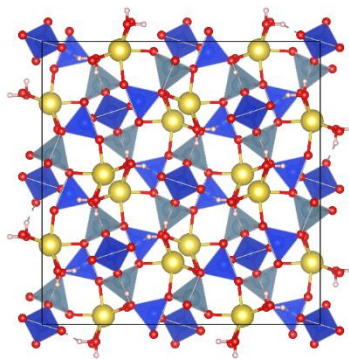
Strength of the breaking of Friedel's law depends only on the degree of non-centrosymmetry (deviation of $\sin\varphi$ from 0) not on the atomic number. Absolute structure is equally easily determined for light and heavy atoms.



Absolute structure and absolute configuration

Absolute structure determination

Natrolite
 $\text{Na}_2[\text{Al}_2\text{Si}_3\text{O}_{10}] \cdot 2\text{H}_2\text{O}$
Fdd2
 $V = 2252 \text{ \AA}^3$
 $T = 293 \text{ K}$



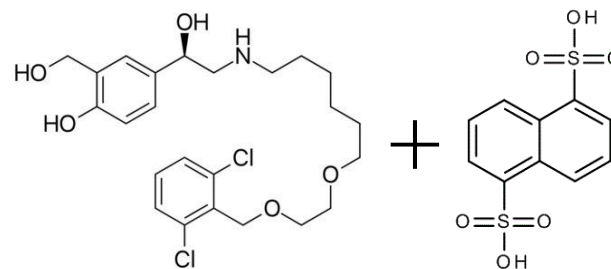
Dynamical refinement:

Correct enantiomorph: $R_{\text{obs}} = 5.95\%$

Wrong enantiomorph: $R_{\text{obs}} = 15.81\%$

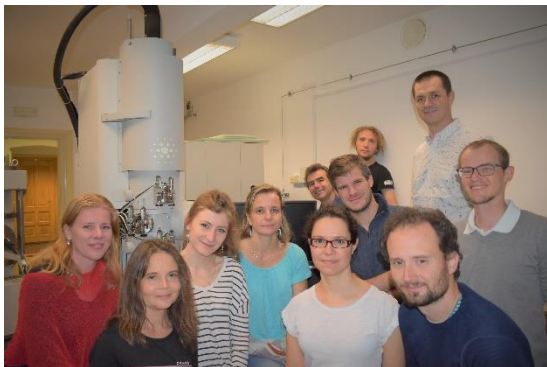
z-score (confidence level): **36.13σ**

vilanterol + 1,5-naphthalenedisulfonic acid



Datas et	R(obs) (%) (R-form)	R(obs) (%) (S-form)	Confidence sigma level (z-score)	Probability of <i>R</i> -form (%)
1	10.93	12.48	5.809	100.000
2	11.21	12.71	8.970	100.000
3	11.16	12.17	7.727	100.000
4	10.82	12.06	8.471	100.000
5	11.91	13.04	6.931	100.000
6	12.18	12.57	5.011	100.000
Comb.			17.336	100.000

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Summary

- ✓ Structure determination by 3D electron diffraction methods has become an almost routine method applicable to all classes of materials
- ✓ Many applications can be done on almost any commercial TEM
- ✓ Optimized TEM setup and software customizations are needed for the best results
- ✓ Dedicated instruments are available on the market, making the method much more easily accessible
- ✓ Many challenges remain. There is a potential for further significant improvement of the method
- ✓ Take-home-message: 3D ED can solve many hard problems in material science. Do not lose time attacking these problems with „familiar“ methods, spend it rather on learning 3D ED (or contact someone to help you)...