

IUCr-2021

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# In the age of electrons, do we still need powder diffraction?

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netherlands  
**eScience center**



Lukáš Palatinus

ECM-31 (Oviedo, 23-08-2018)

Crystallography in the 21<sup>st</sup> Century, the age of electrons?

2021: Dedicated equipment



**Science**  
 2018 BREAKTHROUGH OF THE YEAR  
 Development cell by cell

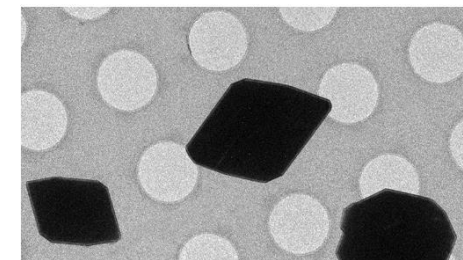
**RUNNERS-UP**  
 Messengers from a far-off galaxy  
 Molecular structures made simple  
 Ice age impact  
 #MeToo makes a difference  
 An archaic human 'hybrid'  
 Forensic genealogy comes of age  
 Gene-silencing drug approved  
 Molecular windows into primeval worlds  
 How cells marshal their contents

**BREAKDOWNS**  
 Climate-fueled disasters rise  
 An ethically fraught gene-editing claim  
 Brazilian science gutted

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**Molecular structures made simple**



Structures can now be gleaned from micrometer-size crystals (black), seen here on an electron microscope slide. (GONEN LAB)

Two research teams simultaneously published papers in October revealing a new way to determine the molecular structures of small organic compounds in just minutes, rather than the days, weeks, or months required by traditional methods.

For decades, the gold standard for molecular mapping has been a technique known as x-ray crystallography, which involves firing a beam of x-rays at a crystal containing millions of copies of a molecule lined up in a common orientation. Researchers then track the way x-rays bounce off

2018: Runner-up breakthrough of the year

2019: The revolution



Outlook  
 Cite This: *ACS Cent. Sci.* 2019, 5, 1315–1329  
<http://pubs.acs.org/journal/acscii>

**3D Electron Diffraction: The Nanocrystallography Revolution**

Mauro Gemmi,<sup>\*,†,‡</sup> Enrico Mugnaioli,<sup>†</sup> Tatiana E. Gorelik,<sup>‡</sup> Ute Kolb,<sup>§,||</sup> Lukas Palatinus,<sup>⊥</sup> Philippe Boullay,<sup>#,¶</sup> Sven Hovmöller,<sup>∇</sup> and Jan Pieter Abrahams<sup>○,◆,¶</sup>

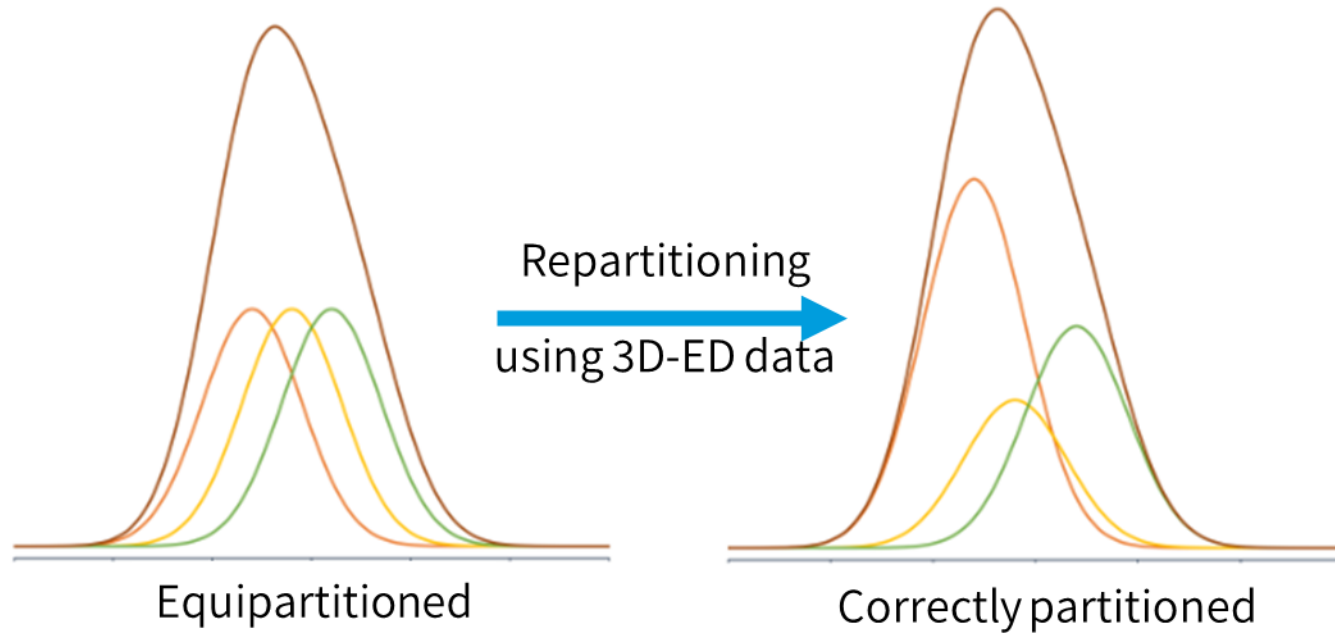
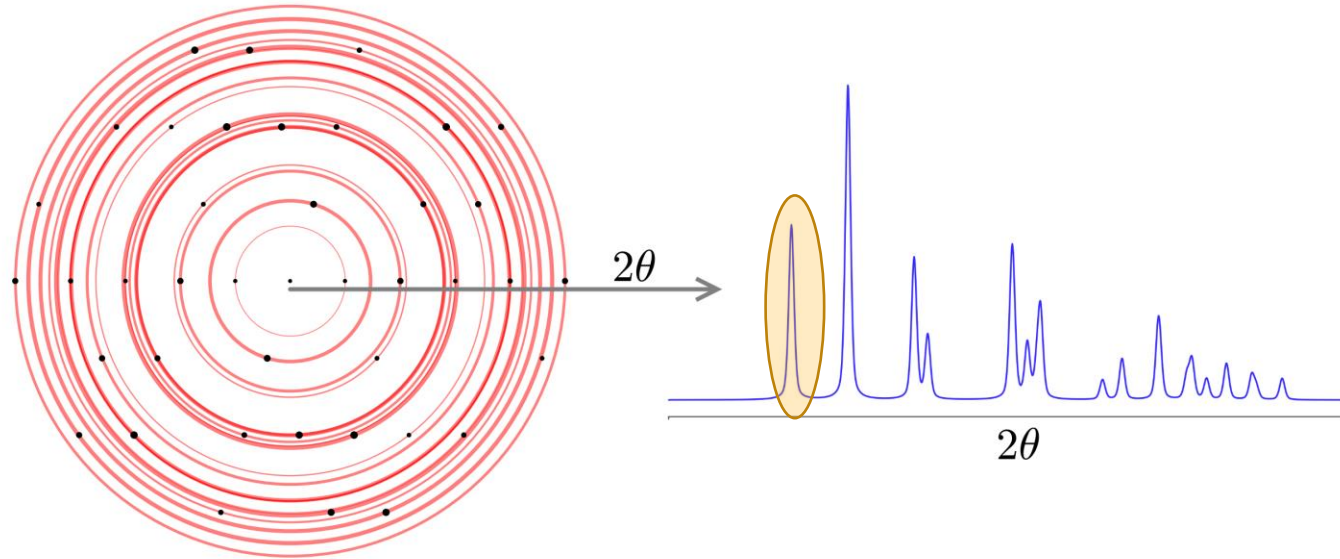
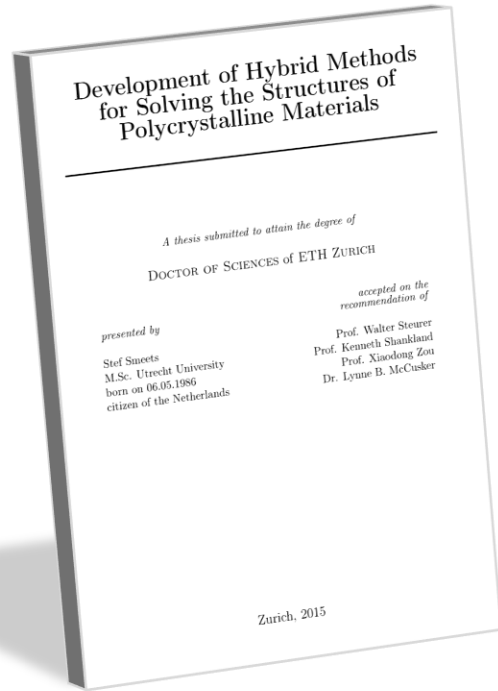


**A** ABSTRACTS  
*Acta Cryst.* (2019). **A75**, a261

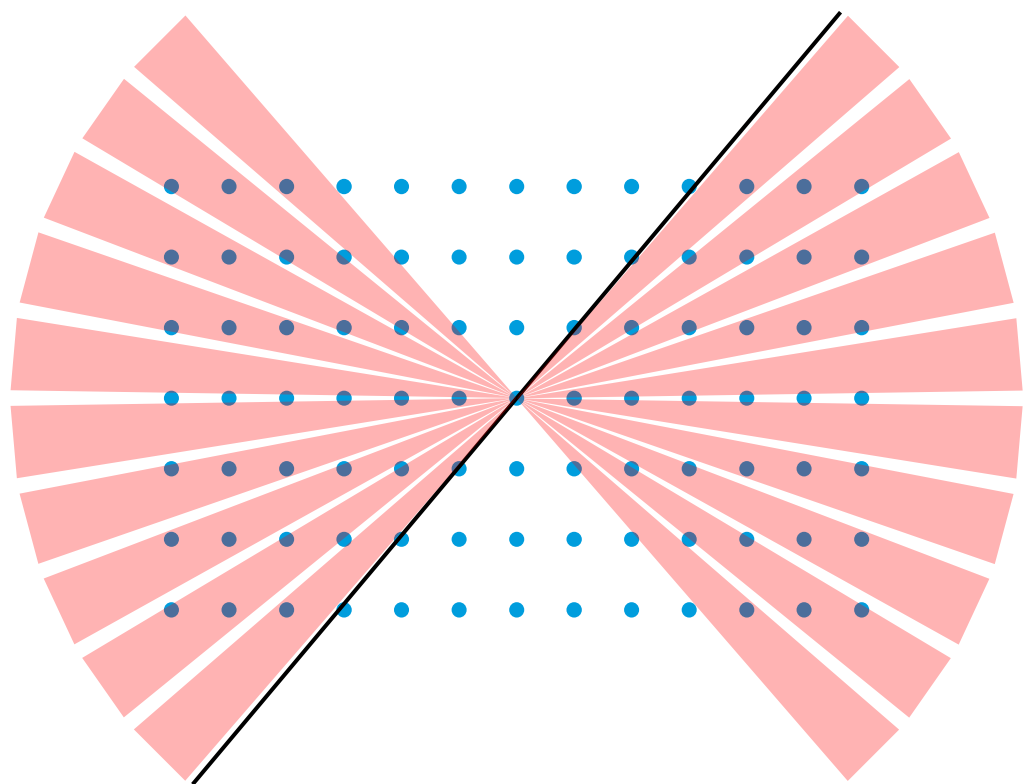
**The death of powder – micro-electron diffraction with EIGER**

C. Schulze-Briese, A. Förster, P. Hofer, S. De Carlo, L. Piazza, J. T. C. Wennmacher and T. Grüne

Also 2019:



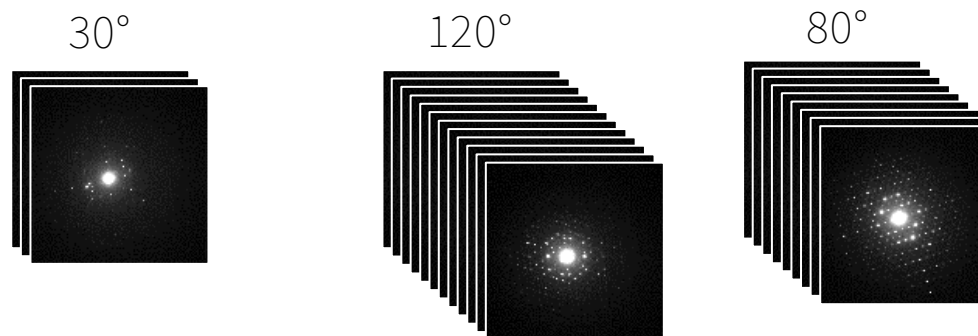
# Typical workflow



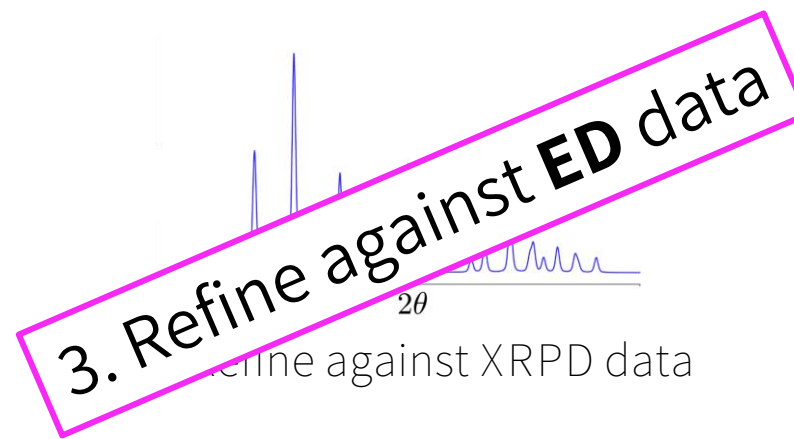
3D-ED / microED  
Data collection



1. Collect ED data on a few crystals



2. Solve with SHELXS/SHELXT/Superflip/SIR



# In the age of electrons, do we still need powder diffraction?

- [SSZ-27](#) – What if our crystal selection is biased?
- [SerialED](#) – Can we do quantitative phase analysis using ED?
- [SerialRED](#) – Can we do phase identification using ED?



# Zeolite SSZ-27

What if our crystal selection is biased?

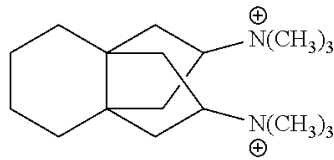


# Zeolite SSZ-27

## Synthesis

S.I. Zones, D. Xie, and R.J. Saxton, US patent 9,586,829 B2 (2017)  
*Molecular sieve SSZ-27*

Aluminosilicate, Si:Al = ~14

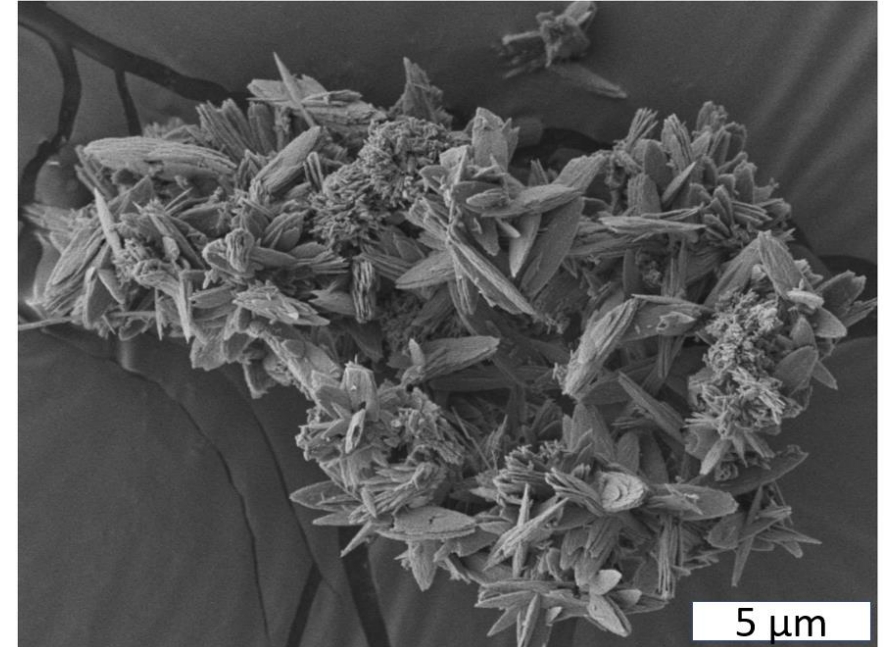


hexamethyl [4.3.3.0] propellane-8, 11-diammonium cation

Organic template

## Applications

- Adsorbent for gas separation
- Catalyst MTO/amine synthesis
- DeNO<sub>x</sub>
- Hydrocarbon trap (C<sub>3</sub>)



Microcrystalline powder



# Electron diffraction data collection

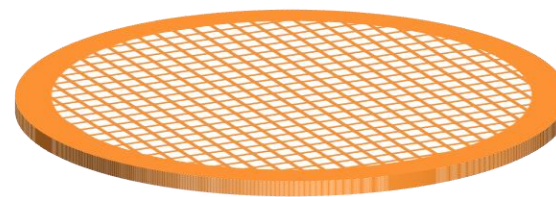
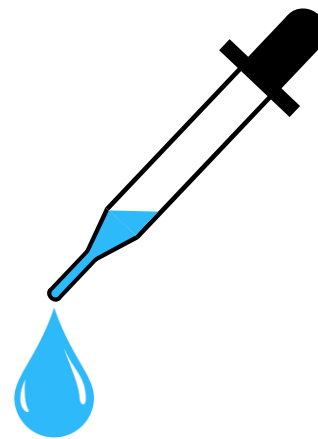
Pick a single crystal for data collection

150  $\mu\text{m}$

## Sample prep



Crystalline powder  
(Disperse in EtOH)



3 mm Cu grid



# Structure analysis

*3D electron diffraction*

Collected by Lukas Palatinus (2012)

$C2/m$

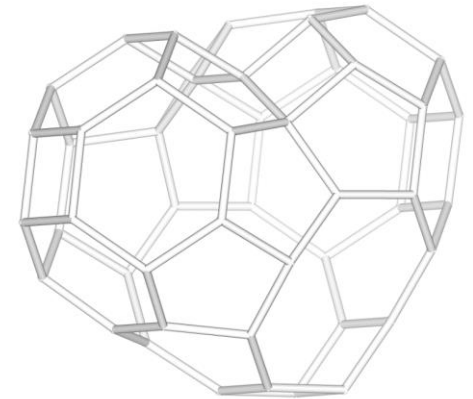
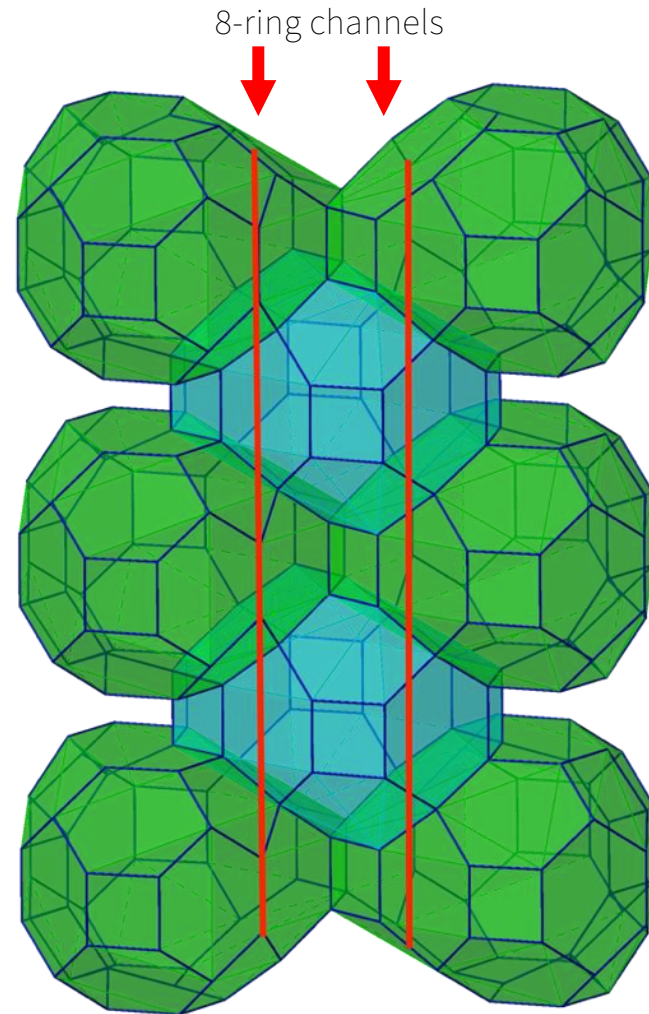
$a = 23.95 \text{ \AA}$

$b = 13.79 \text{ \AA}$

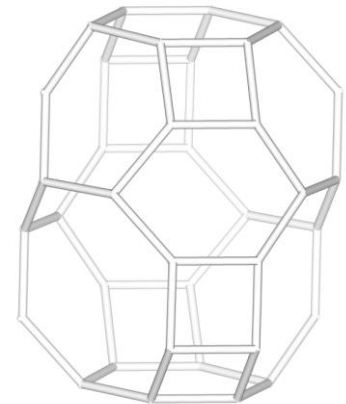
$c = 24.96 \text{ \AA}$

$\beta = 115.7^\circ$

Solved with Superflip and FOCUS



Two types of cavities



# Structure did not match powder data

*3D electron diffraction*

Collected by Lukas Palatinus (2012)

$C2/m$

$a = 23.95 \text{ \AA}$

$b = 13.79 \text{ \AA}$

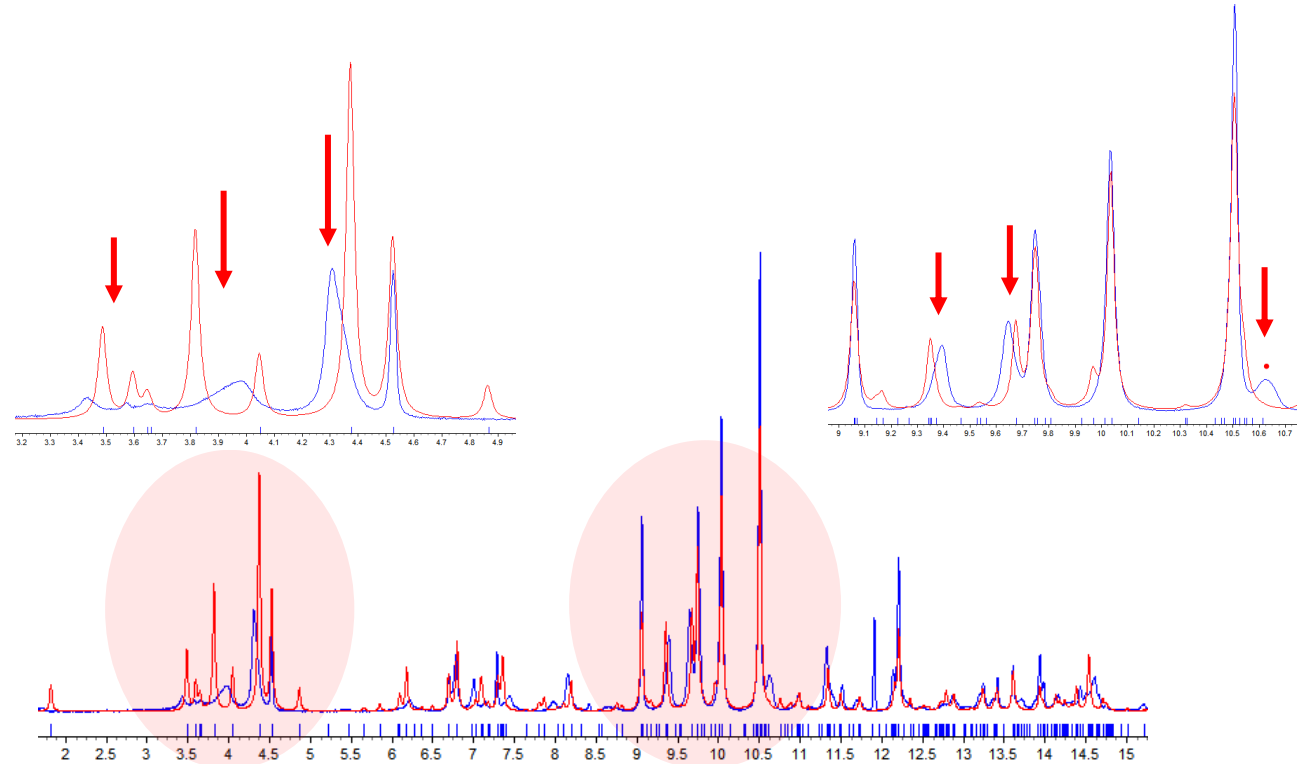
$c = 24.96 \text{ \AA}$

$\beta = 115.7^\circ$

Solved with Superflip and FOCUS

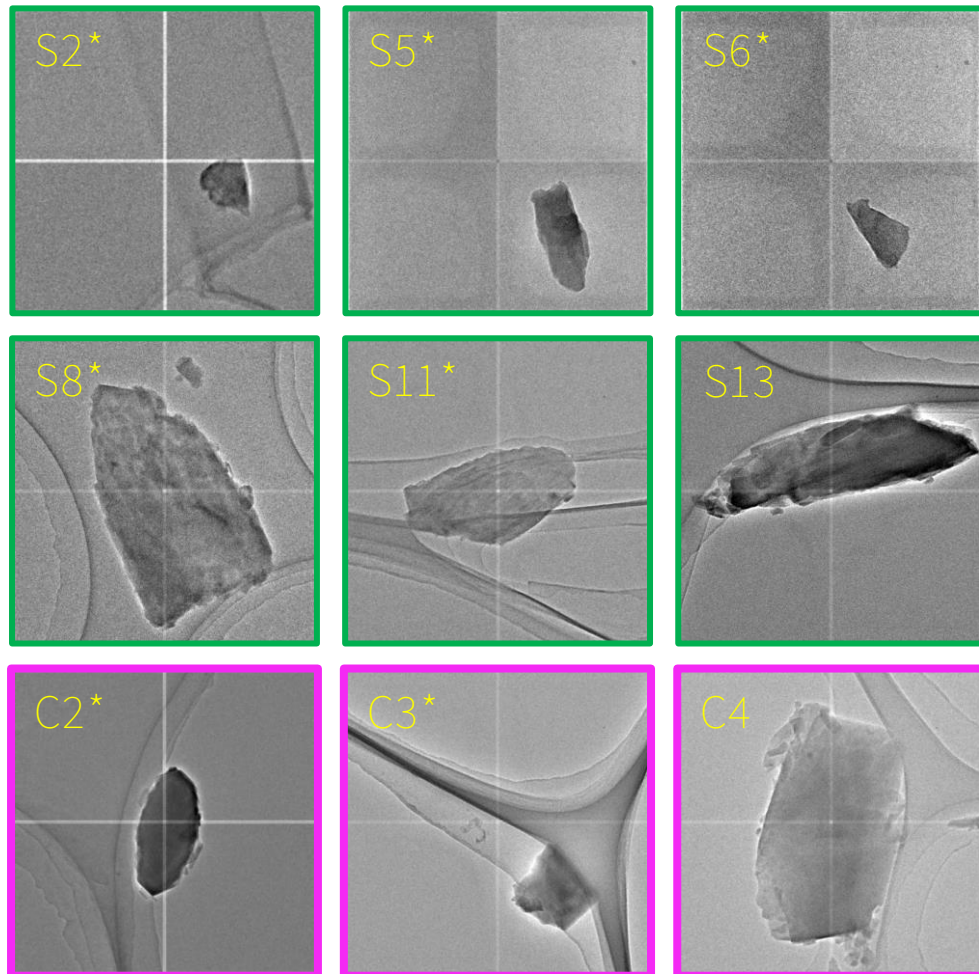
*Synchrotron X-ray powder diffraction*

Profile refinement failed



# Collect ED data on a few more crystals

Total: 18 crystals



400 nm

Phase 1 (14 crystals)

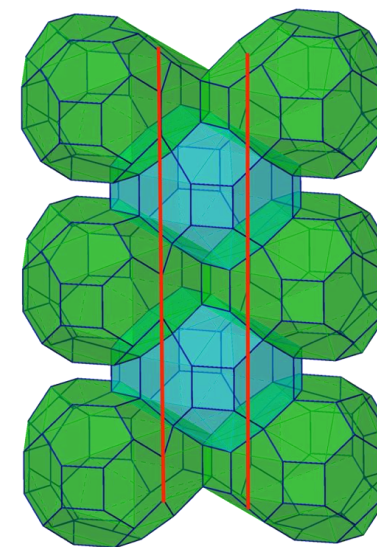
$C2/m$

$$a = 24.12 \text{ \AA}$$

$$b = 13.81 \text{ \AA}$$

$$c = 25.07 \text{ \AA}$$

$$\beta = 115.19^\circ$$



New phase (SSZ-27)

Phase 2 (4 crystals)

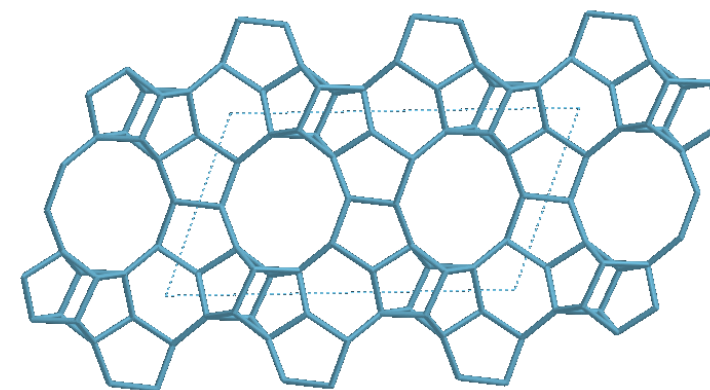
$C2/m$

$$a = 23.40 \text{ \AA}$$

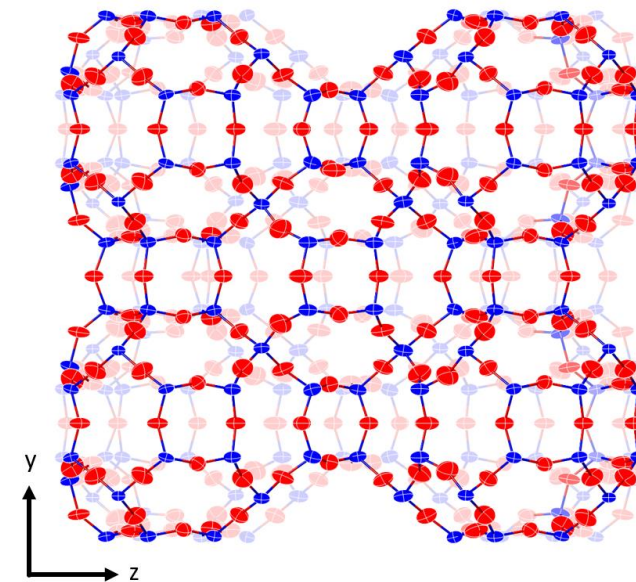
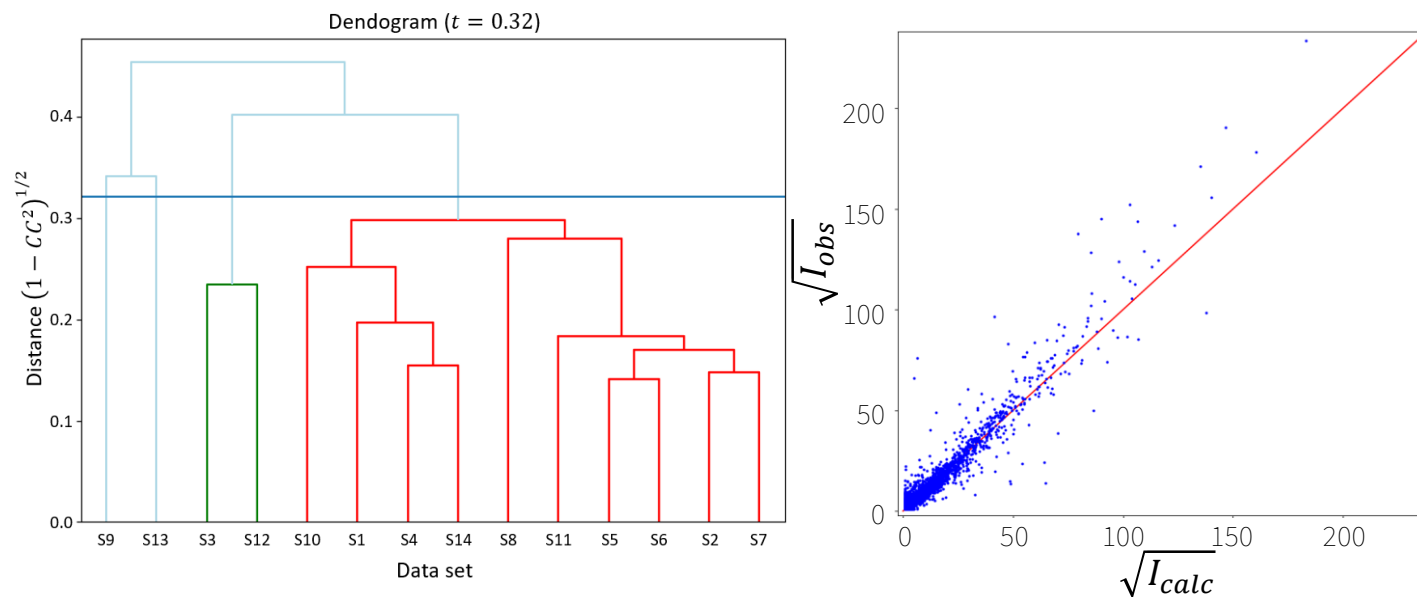
$$b = 13.77 \text{ \AA}$$

$$c = 12.73 \text{ \AA}$$

$$\beta = 110.89^\circ$$

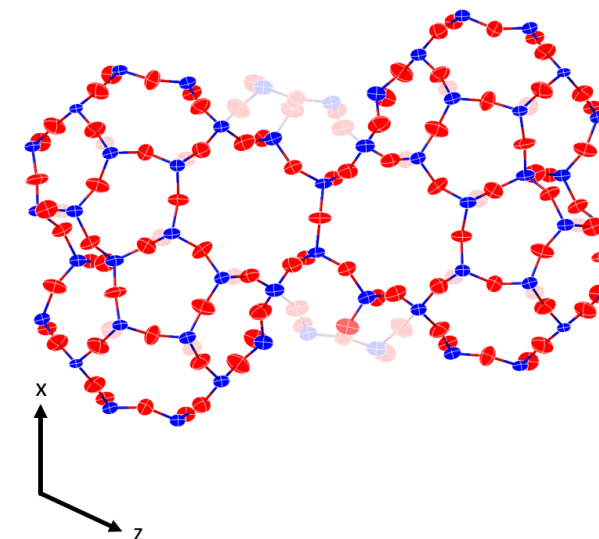


Known phase (CON)

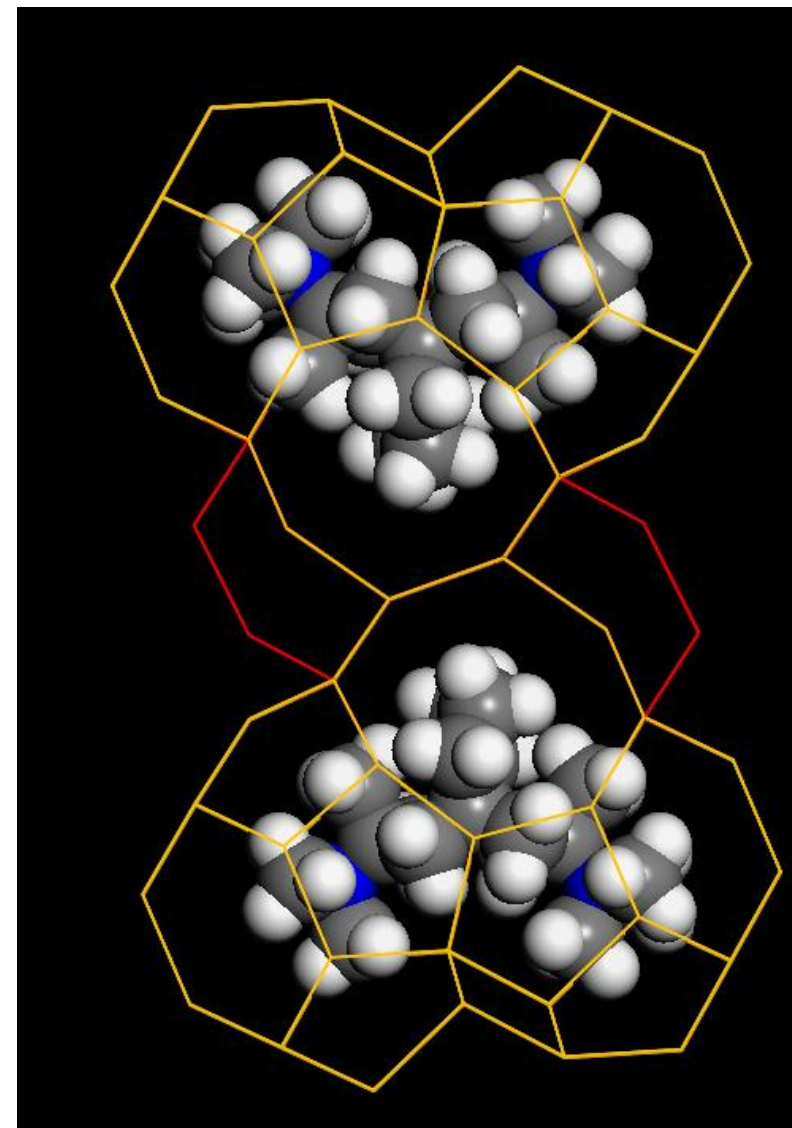
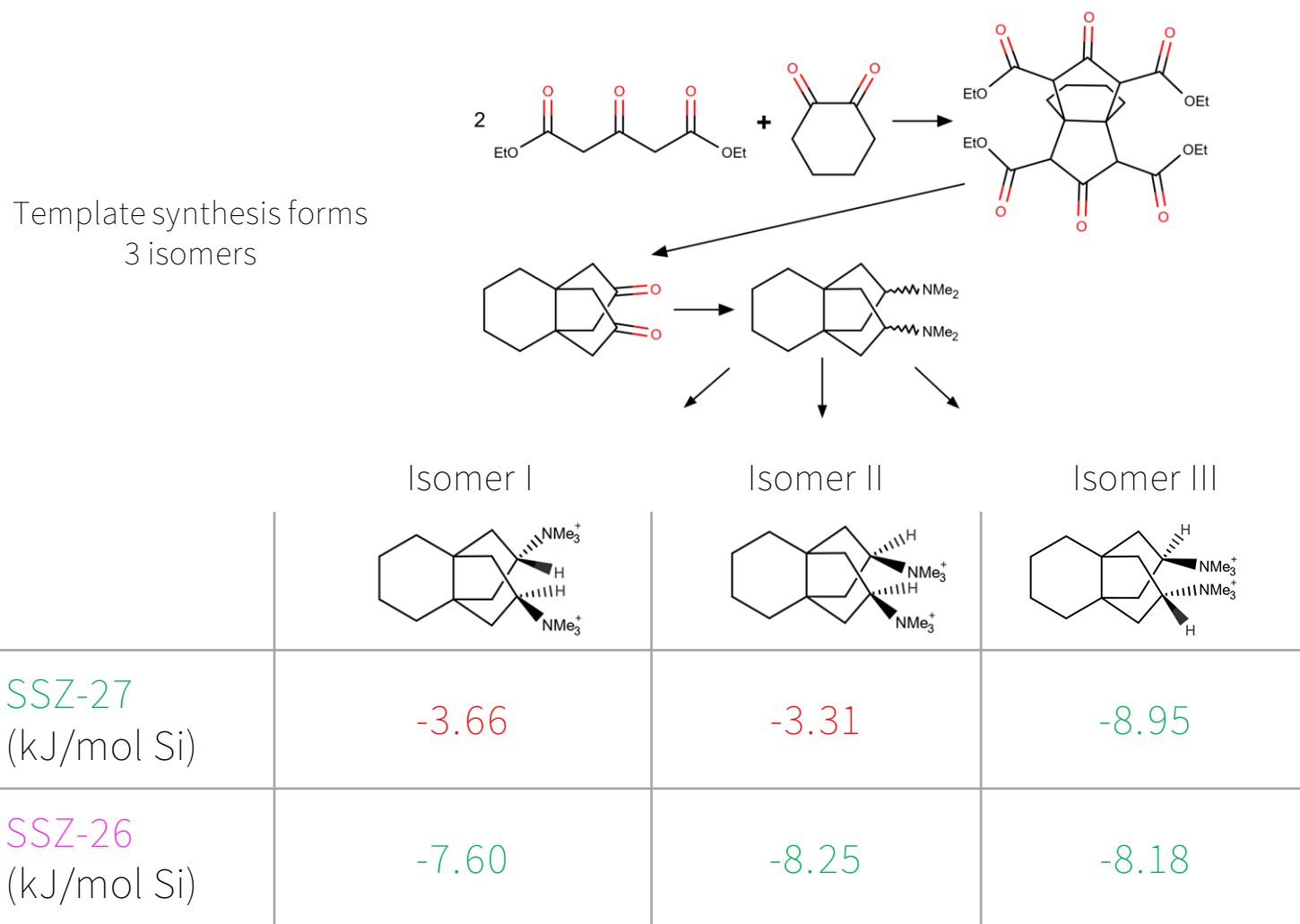


Combine 10 crystals for higher redundancy/completeness using hierarchical cluster analysis

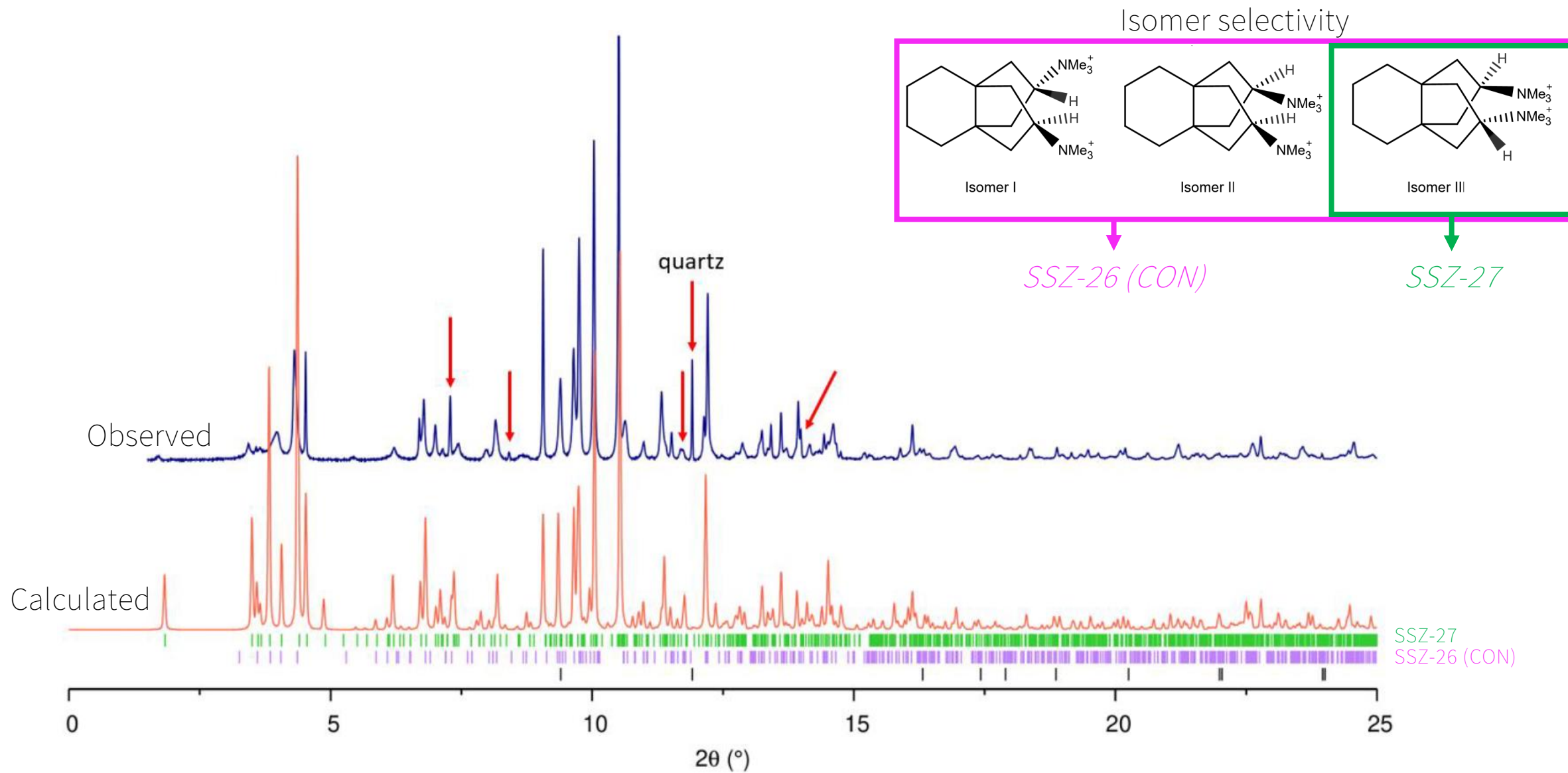
# Refs. (total)	130755	Completeness	98.8
# Refs. (unique)	7913	$CC_{1/2}$	98.8
# Refs. (obs.)	4129	$R_{int}$	0.287
Resolution (Å)	0.77	$R1 [I > 2\sigma(I)]$	0.178



# Molecular modelling confirms isomer specificity



# Profile refinement against XRPD data



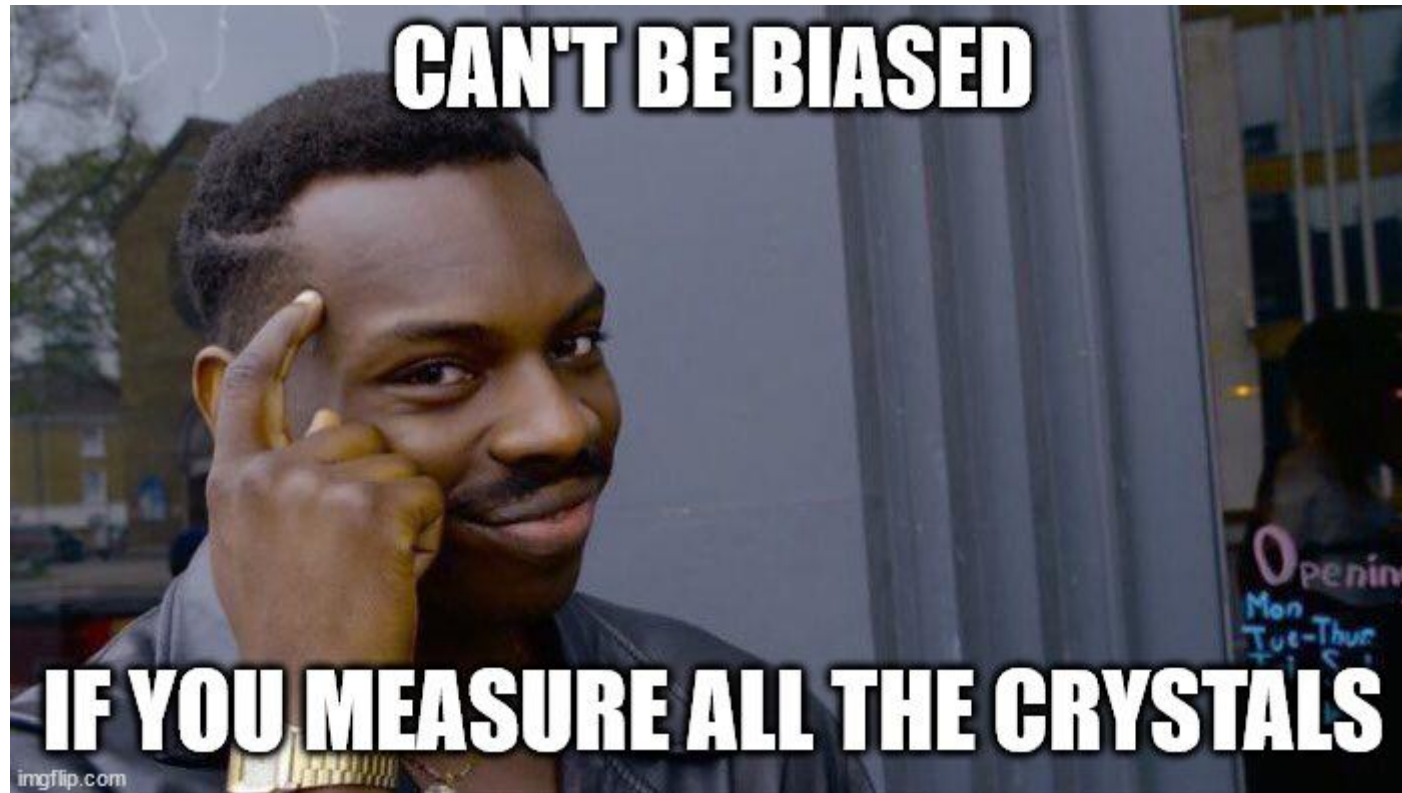
# Summary SSZ-27

- 3D-ED data: [crystal](#) structure
- XPRD data: [bulk](#) structure
  
- ED data [can be biased](#) by crystal selection
- XRPD showed our model was not complete
- Combined data revealed new purification pathway



# Serial electron diffraction

Can we do quantitative phase analysis using electron diffraction?





## Serial electron diffraction

1. Global map
2. Medium mag map (roi)
3. Image segmentation
4. Get particle coordinates
5. Acquire data for each crystal

150  $\mu\text{m}$



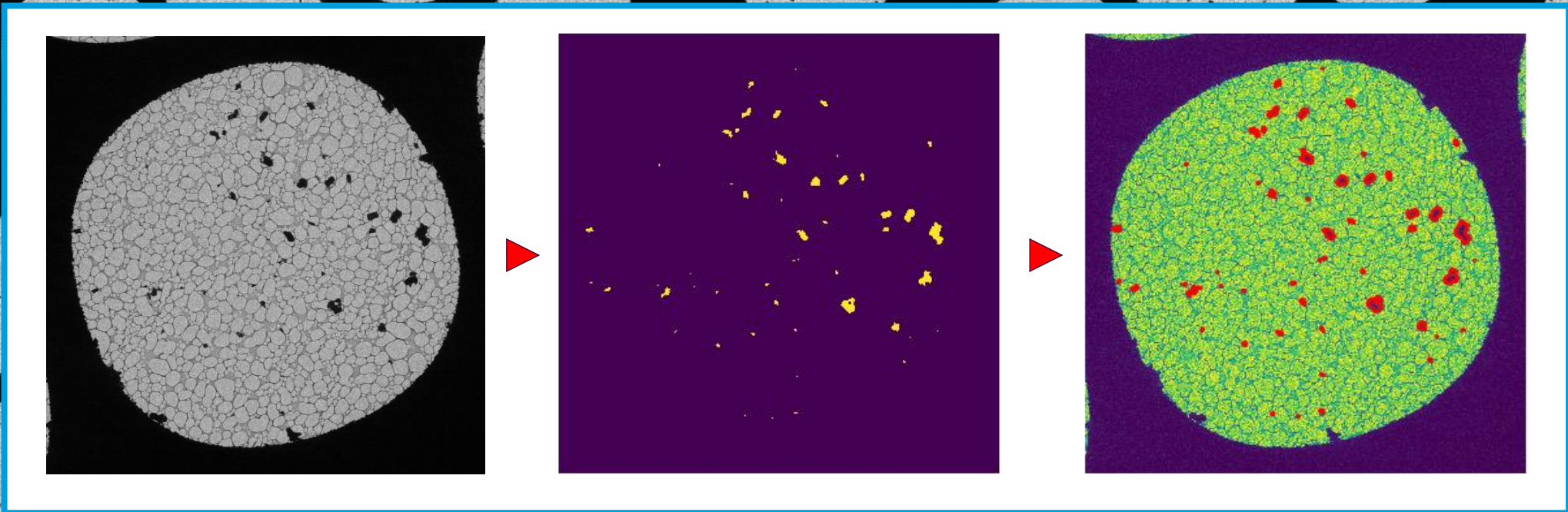
**CRYSTALS**



**CRYSTALS EVERYWHERE**

# Serial electron diffraction

1. Global map
2. Medium mag map (roi)
3. Image segmentation
4. Get particle coordinates
5. Acquire data for each crystal



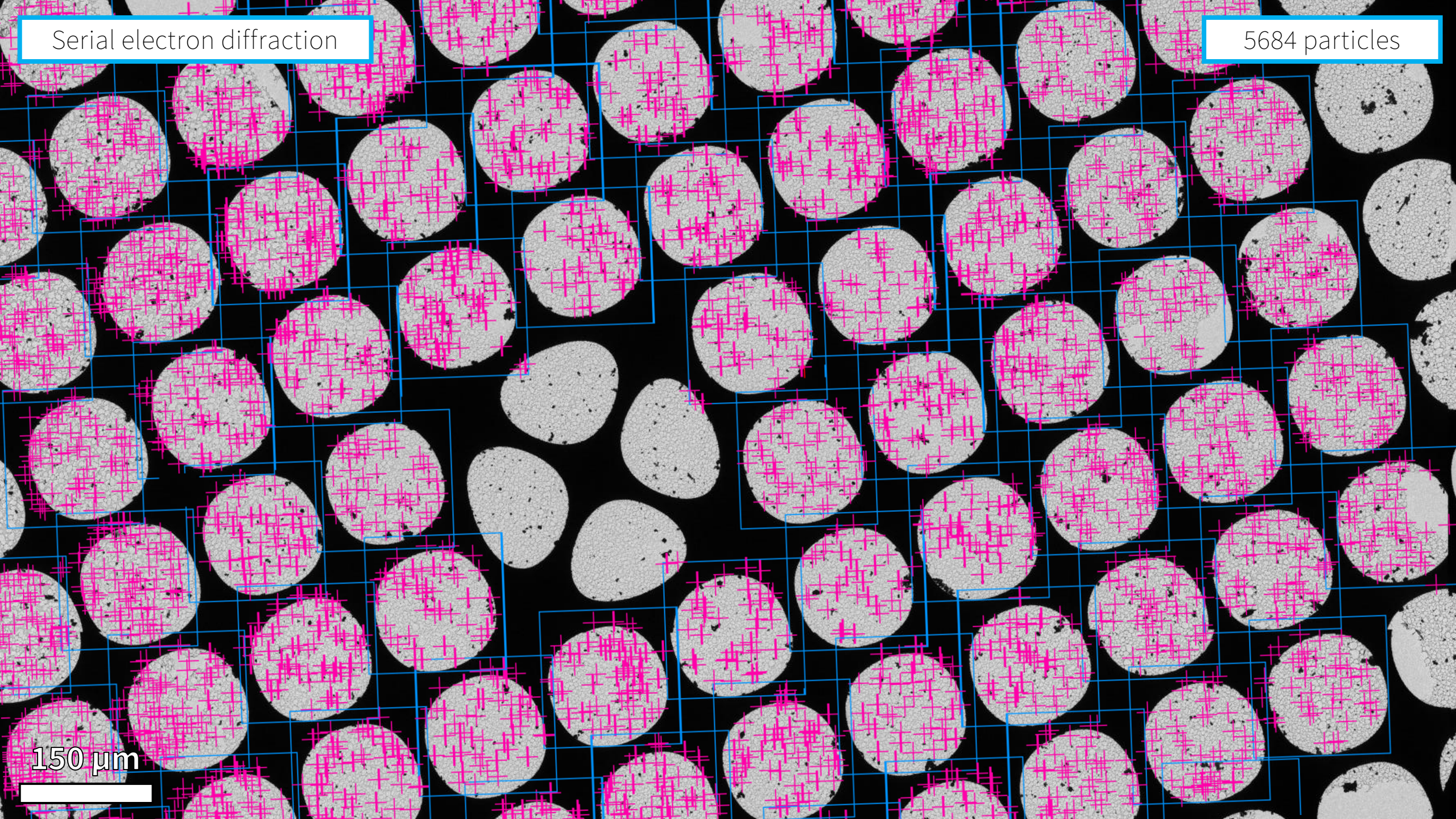
150  $\mu\text{m}$



Serial electron diffraction

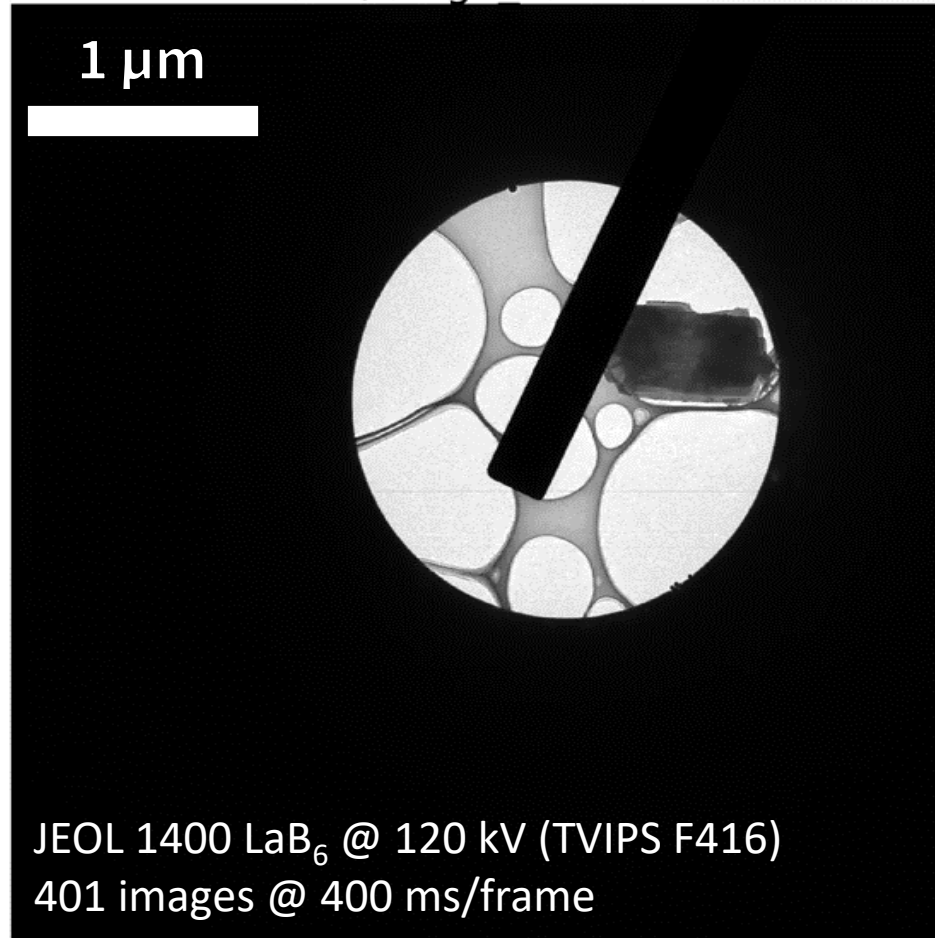
5684 particles

150  $\mu\text{m}$

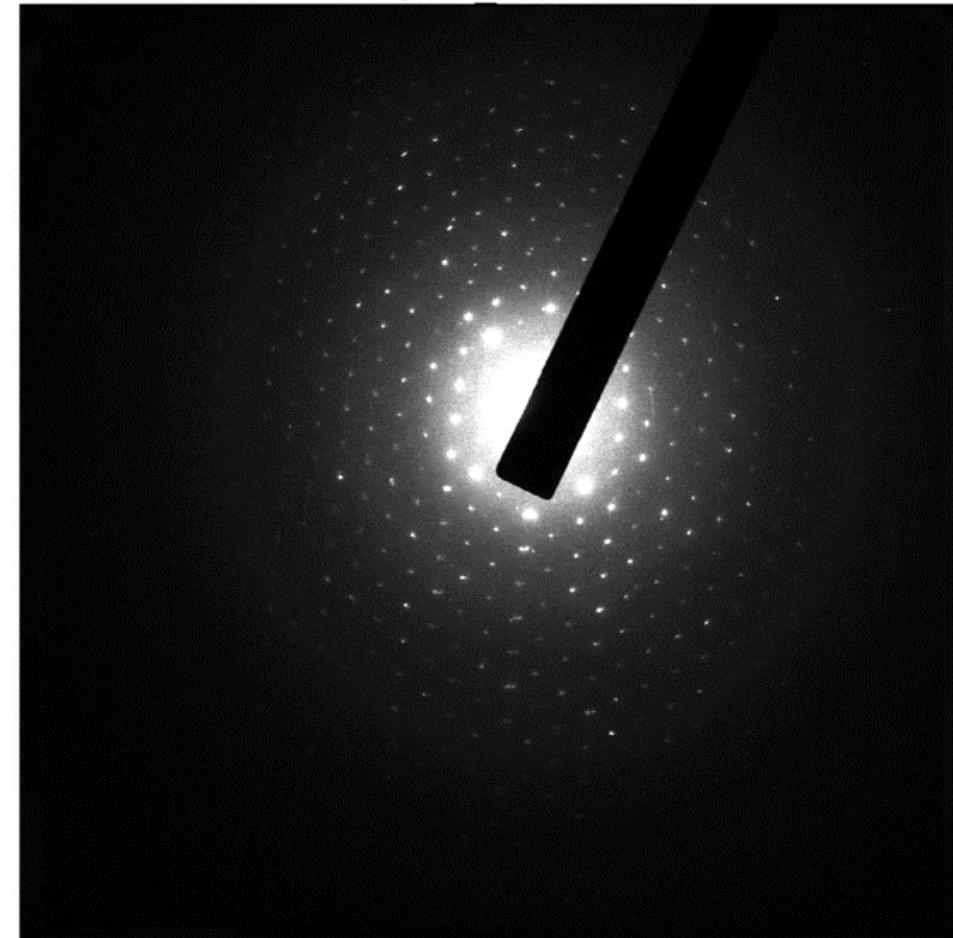


# SerialED data collection (Zeolite)

diff\image 232.tiff

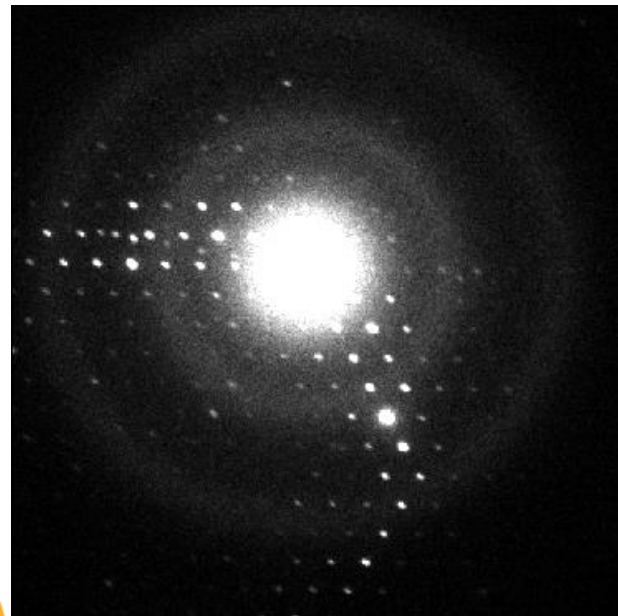


diff\diff 232.tiff

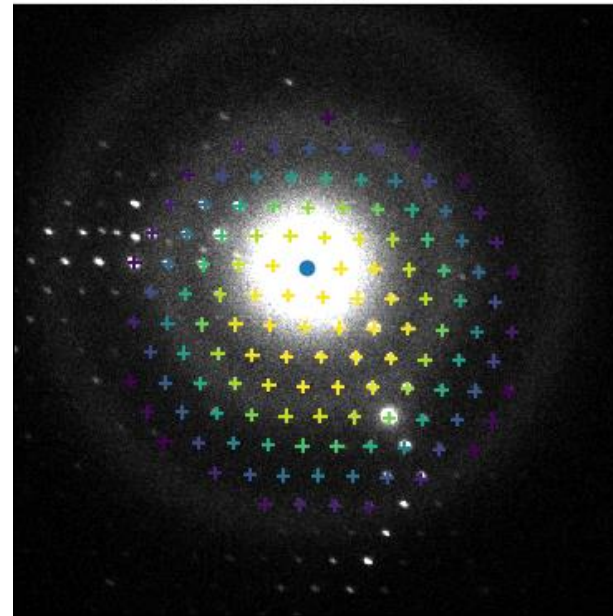


# Indexing: orientation finding

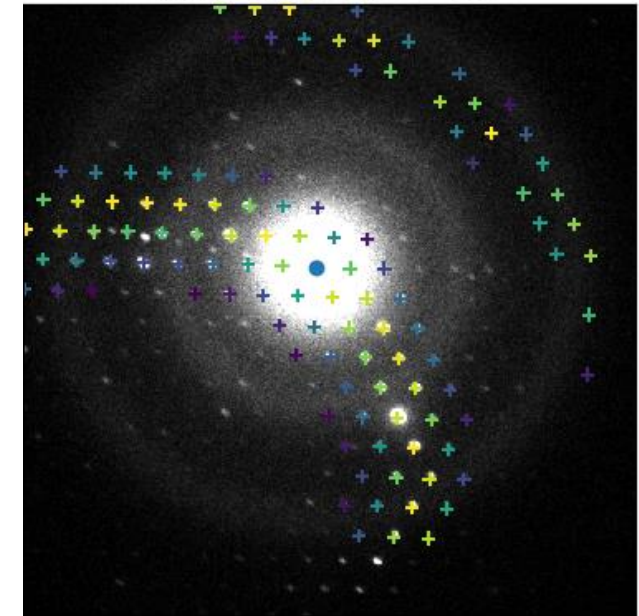
- Forward projection model using known lattice parameters
- Generate pattern library of all possible orientations (~1.5M in **P1**)
- Match best orientation and index data



al: 0.96, be: 0.78, ga: 0.81  
score = 7363.9, scale = 251.6  
proj = 700, phase = LTA

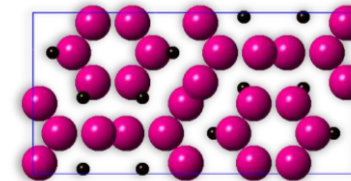
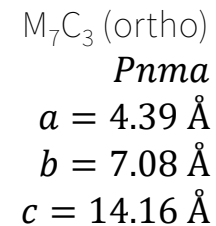
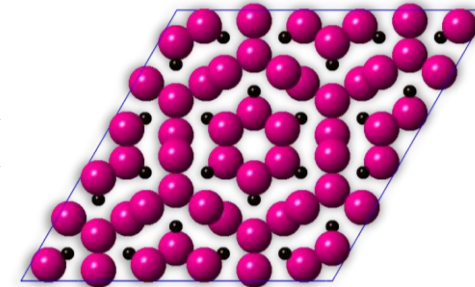
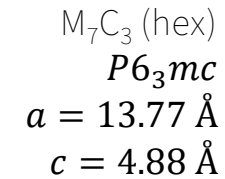
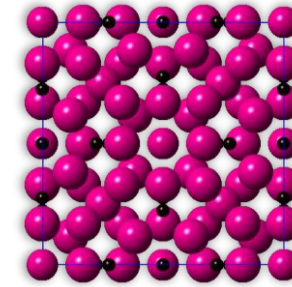
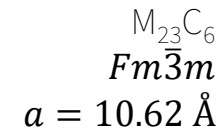


al: 0.98, be: 0.80, ga: 2.90  
score = 11996.2, scale = 251.1  
proj = 732, phase = LTA

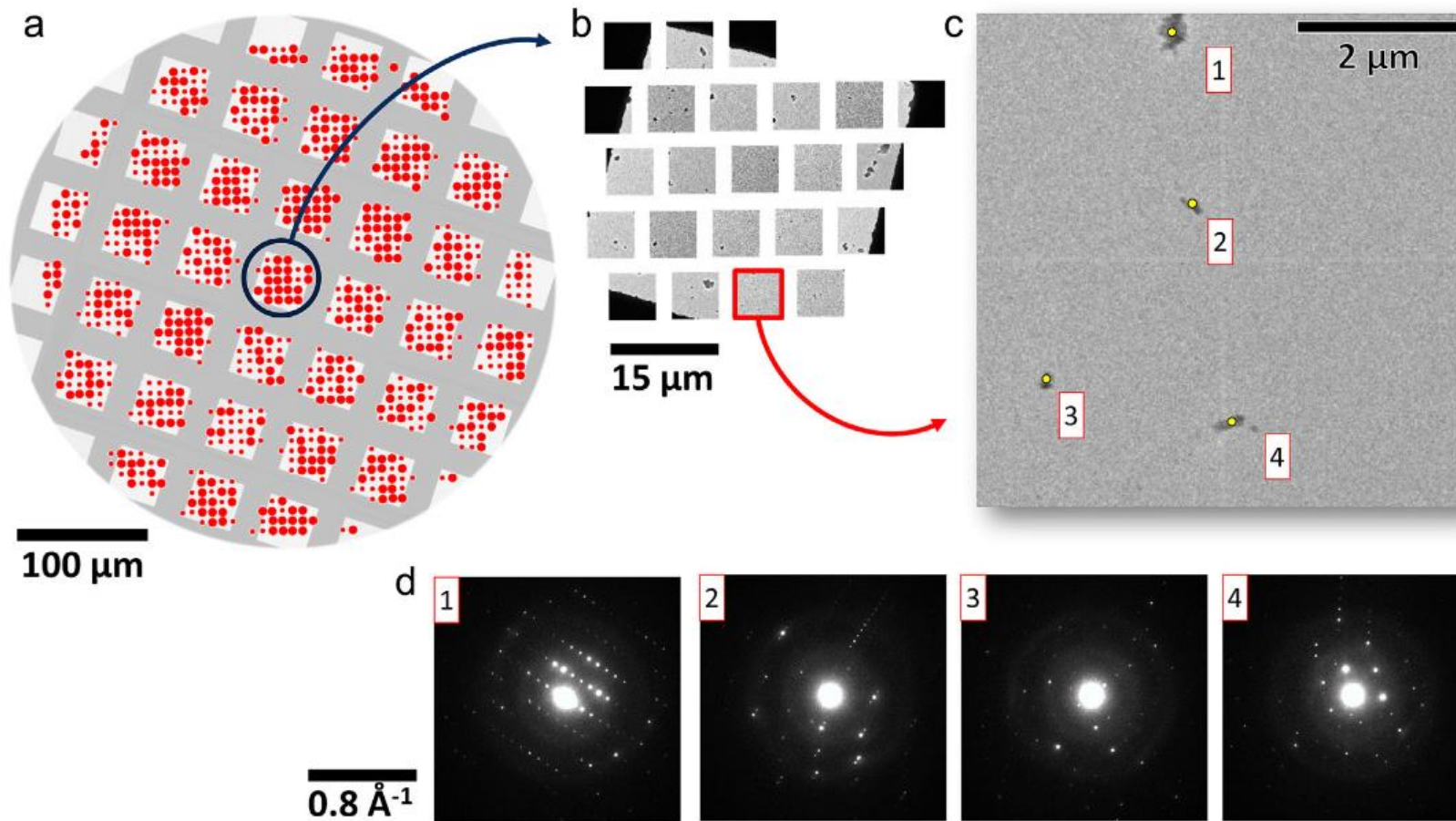


# Study on Cr Carbides

- Carbide:  $M_xC_y$  (M=Cr, Fe, Mo, ...)
- Additive in steel manufacturing ( $\leq 5$  wt%)
- Influences strength, ductility, corrosion resistance, etc
- Sample:
  - Phase isolates from martensitic stainless steel



# SerialED data collection



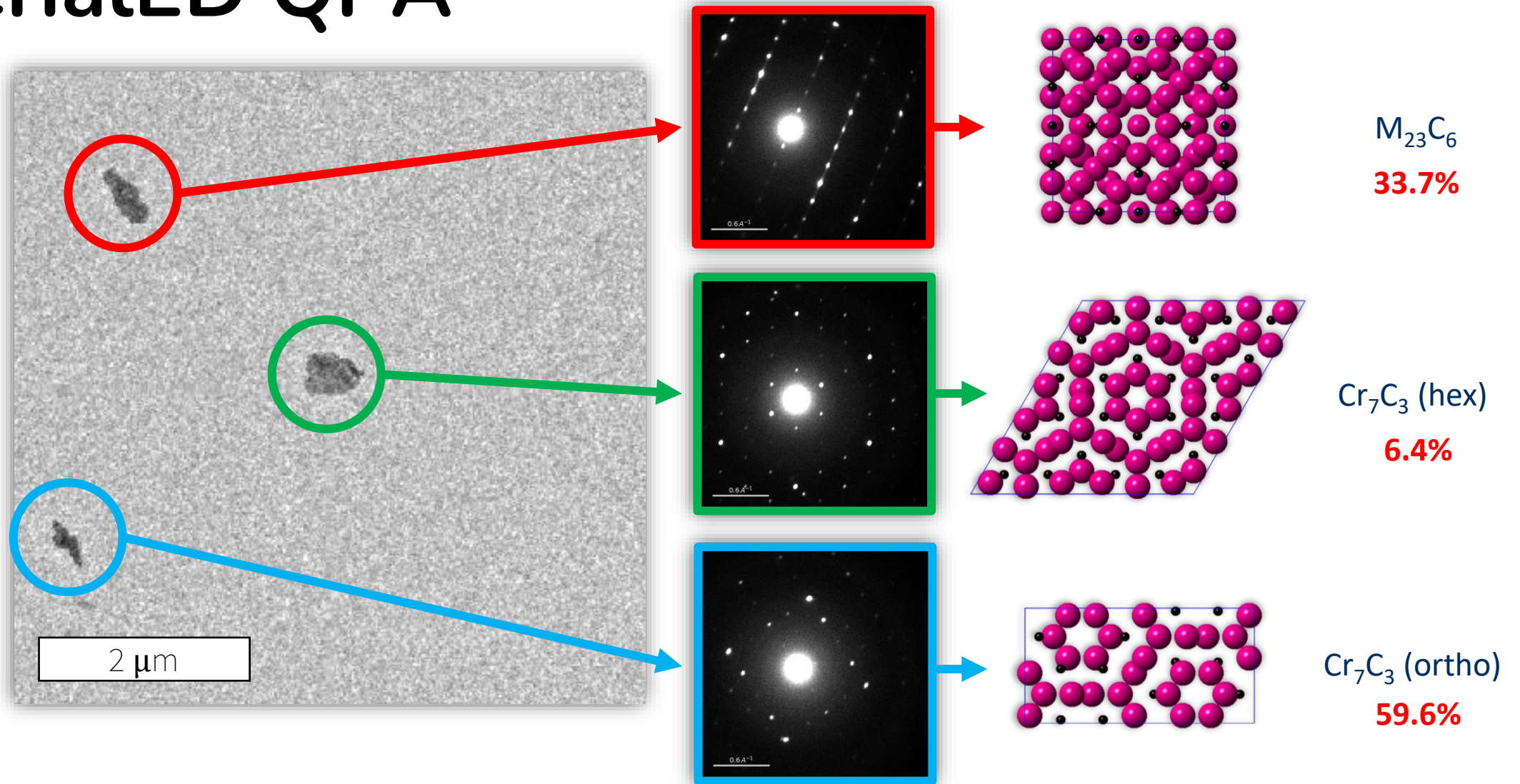
4 sessions  
3939 particles  
~4 hours total  
1000 crystals/hr

Collected with Instamatic  
JEOL 2100 with ASI Timepix

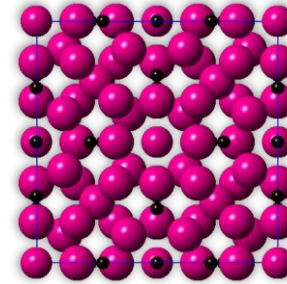
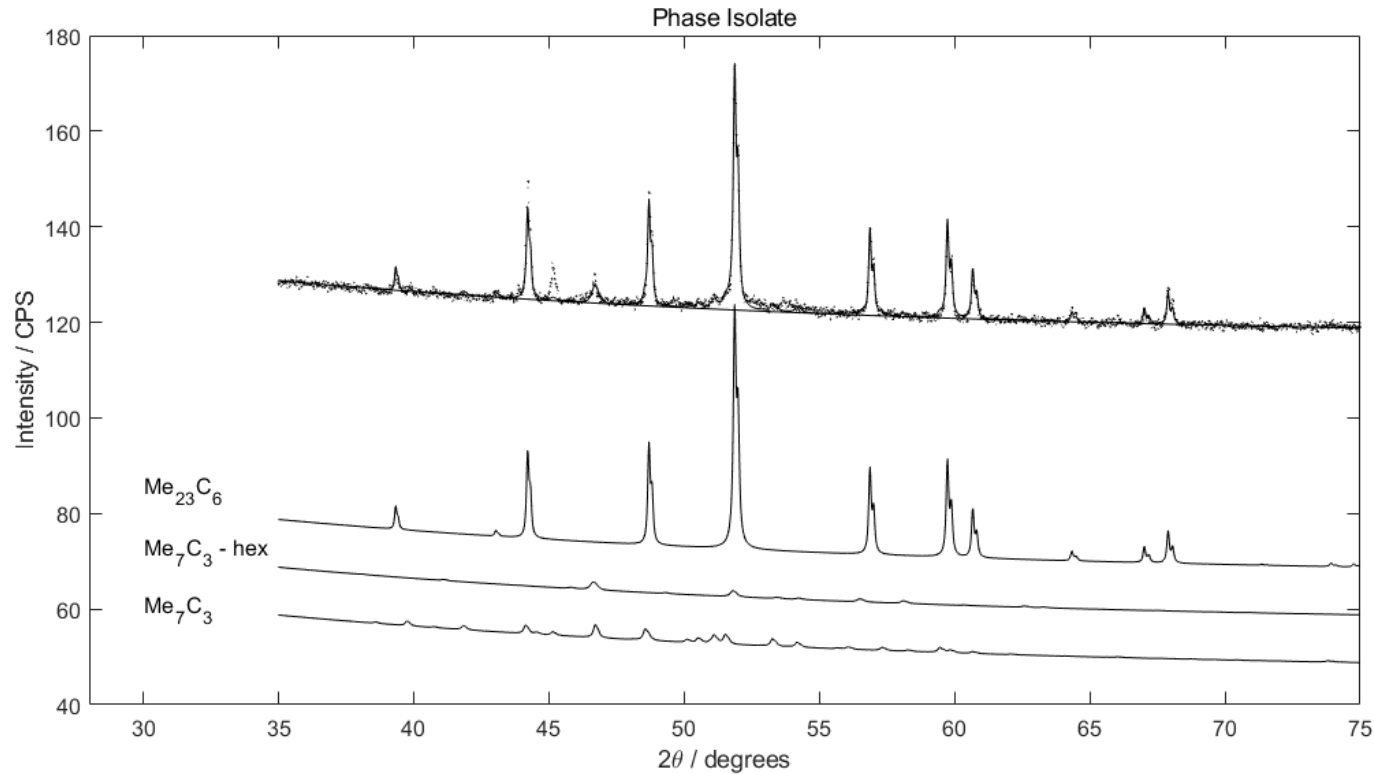




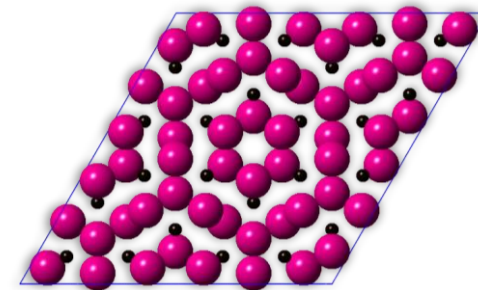
# SerialED QPA



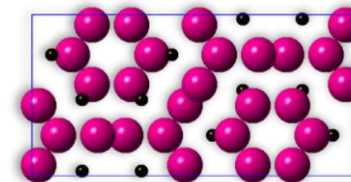
# Powder QPA



$\text{Me}_{23}\text{C}_6$   
XRPD: 84.49 %  
SerialED: 33.7 %



$\text{Me}_7\text{C}_3$  (hex)  
XRPD: 3.09 %  
SerialED: 6.4 %



$\text{Me}_7\text{C}_3$  (ortho)  
XRPD: 12.42 %  
SerialED: 59.6 %



# Summary SerialED

- SerialED data: [crystal counts](#)
- XPRD data: [bulk composition](#)
  
- Indexing still ED frames is an unsolved problem(?)
- Difficult to beat simplicity of XRPD



# Serial **rotation** electron diffraction

Can we do phase identification using electron diffraction?

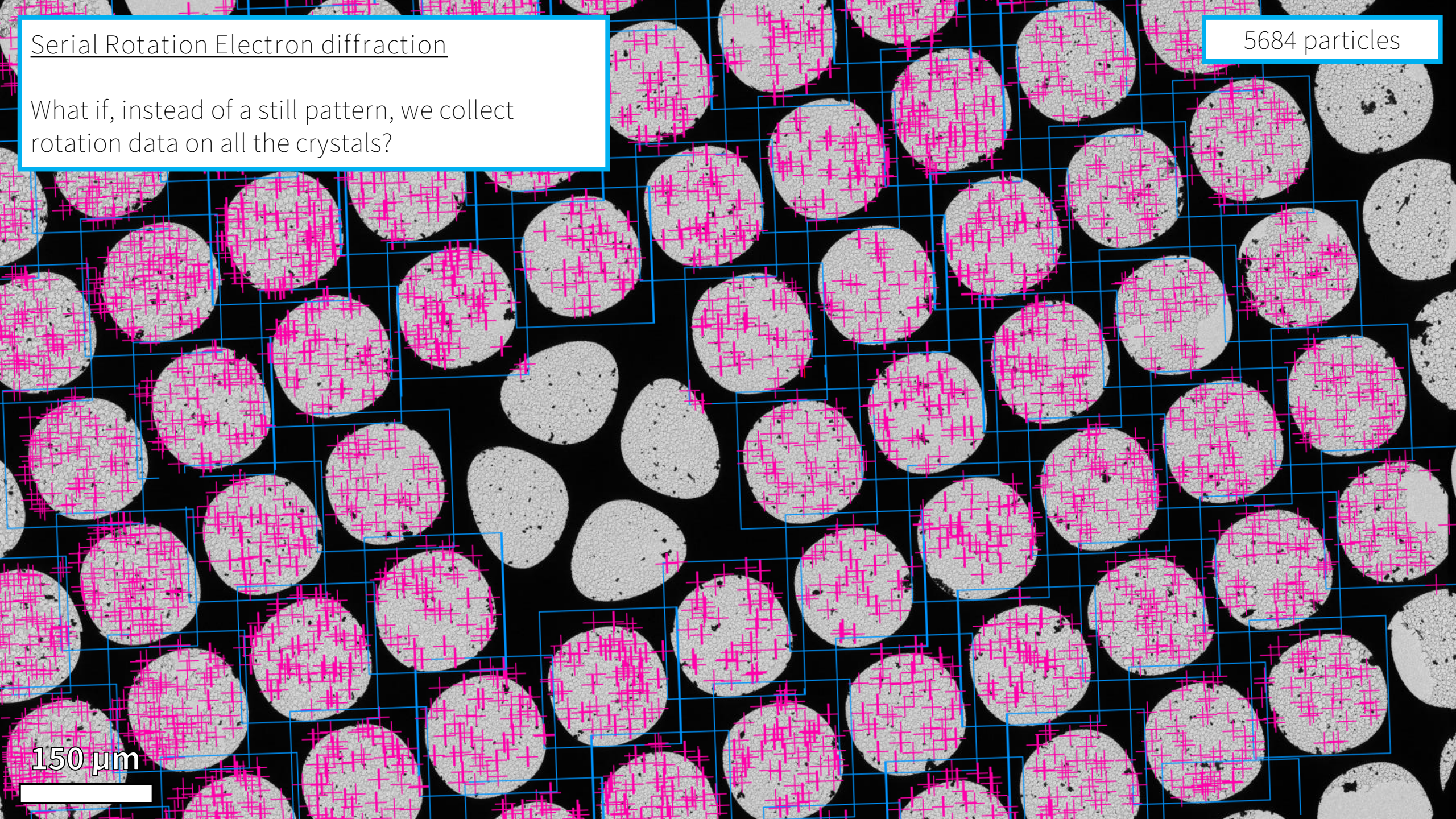


# Serial Rotation Electron diffraction

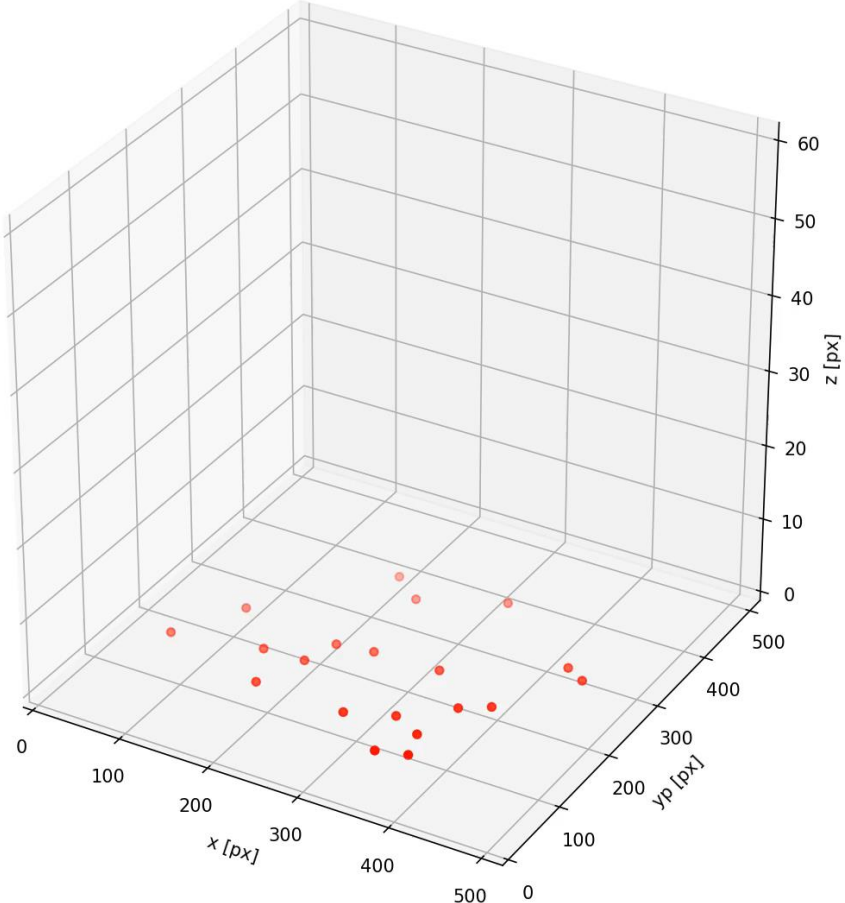
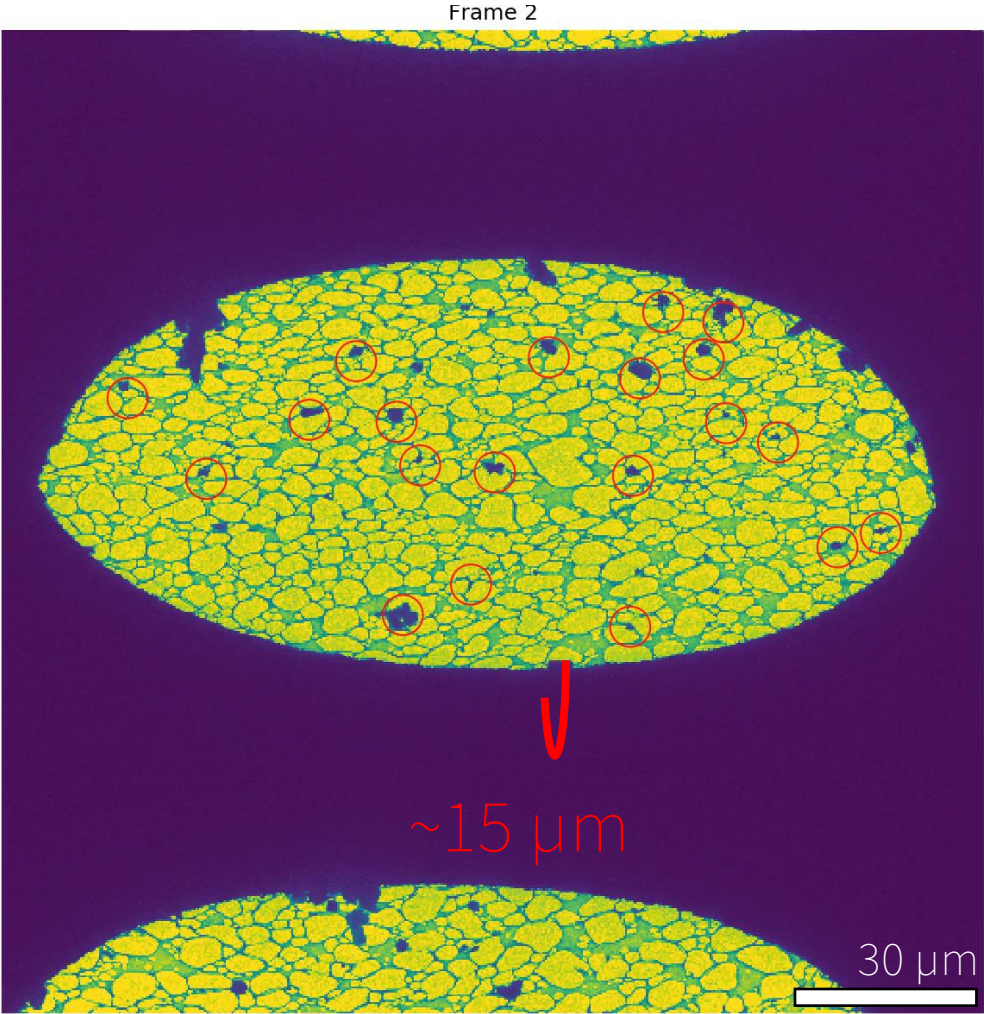
5684 particles

What if, instead of a still pattern, we collect rotation data on all the crystals?

150  $\mu\text{m}$



# The crystal tracking problem

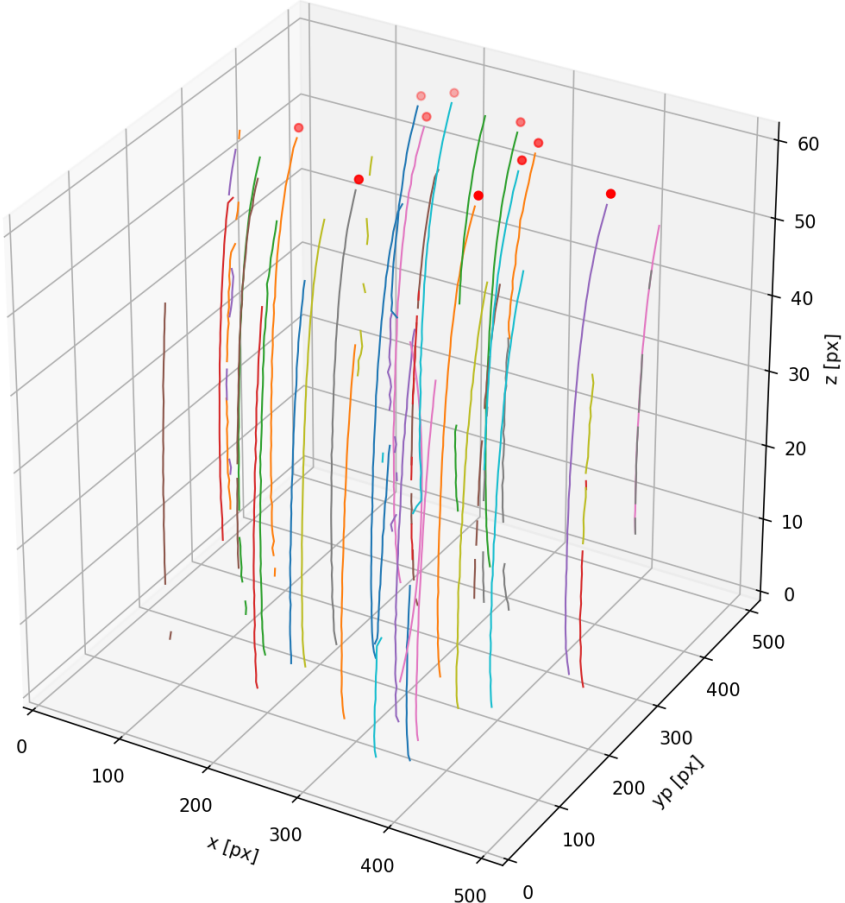
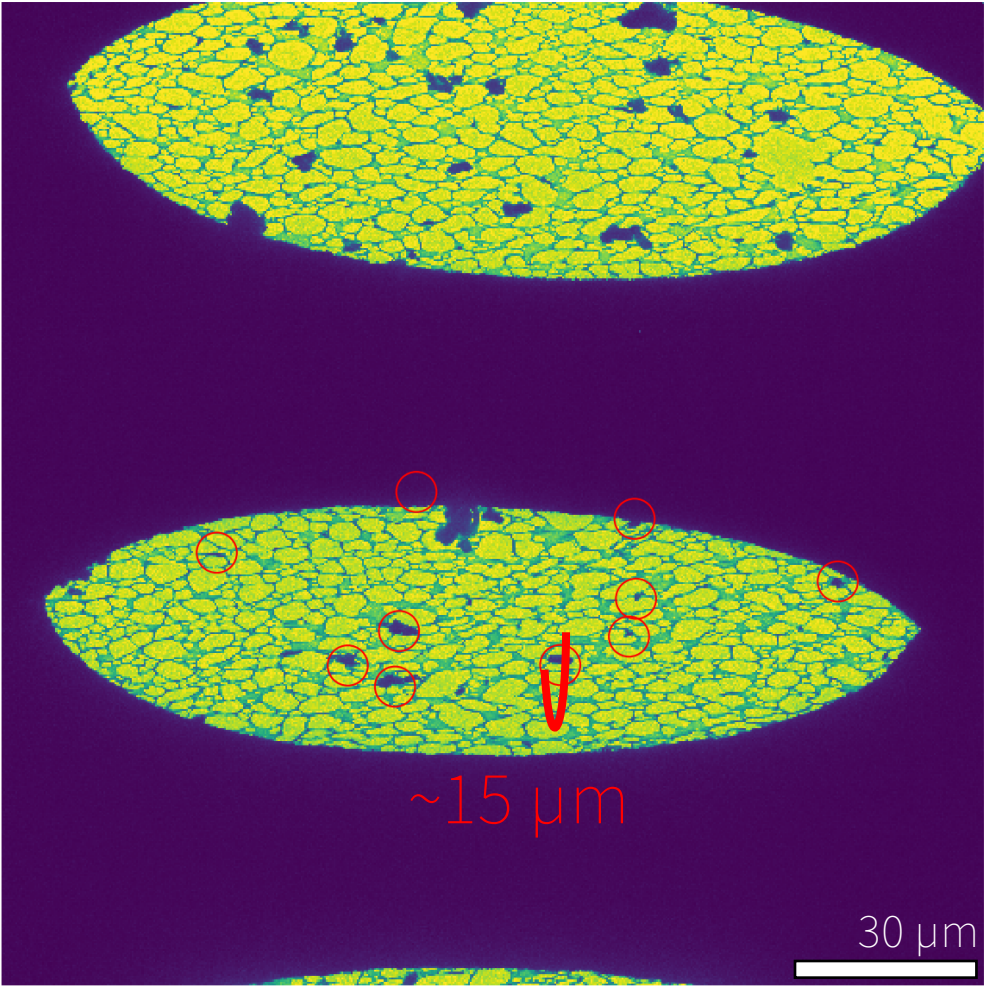


Crystal movement



# The crystal tracking problem

Frame 60

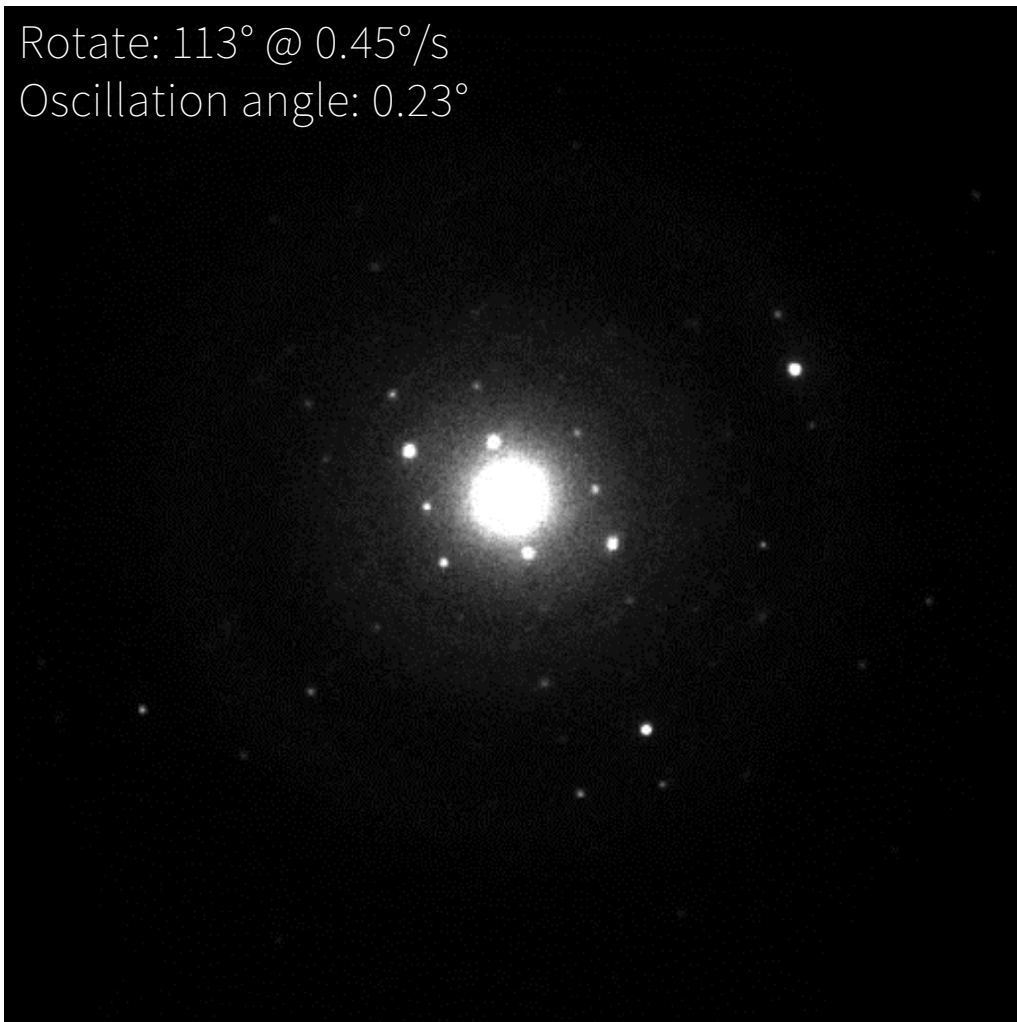


Crystal movement



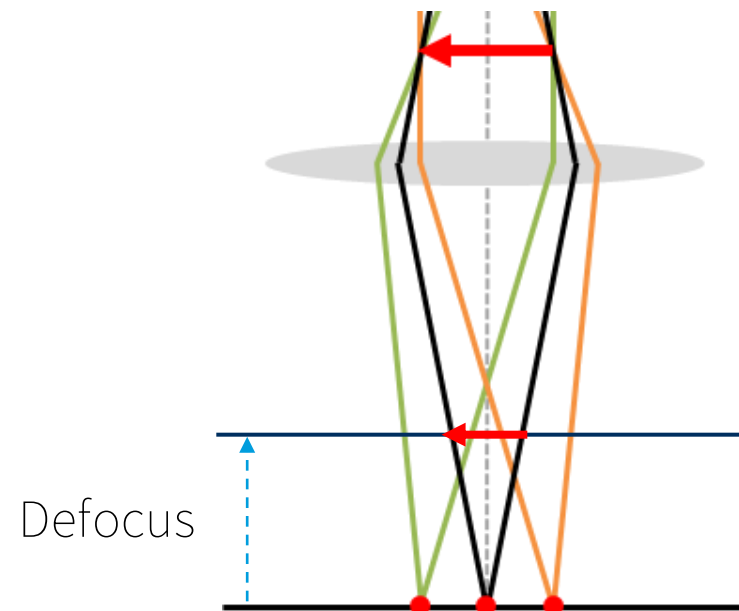
# Using defocus for tracking

Rotate:  $113^\circ$  @  $0.45^\circ/\text{s}$   
Oscillation angle:  $0.23^\circ$



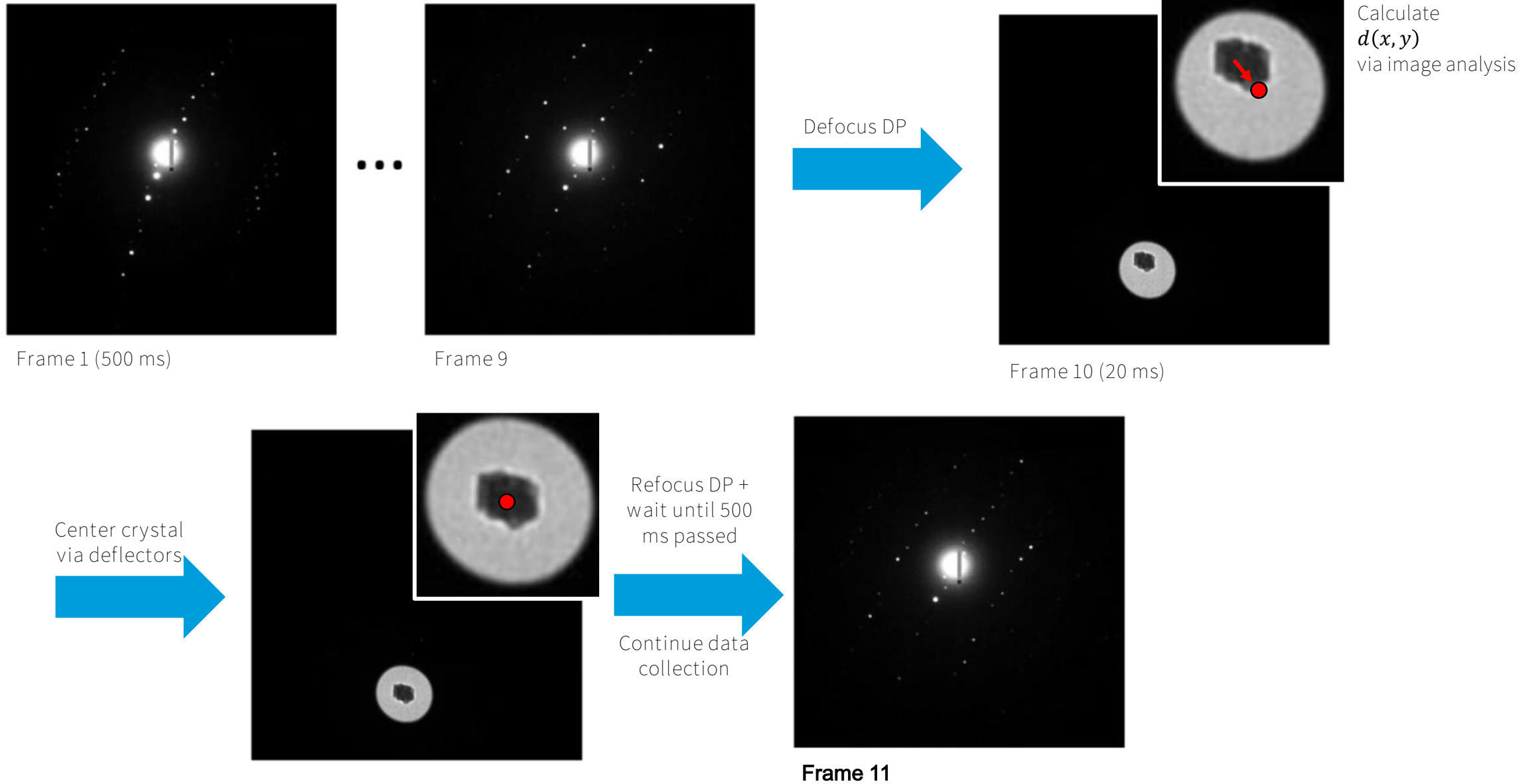
Tracking

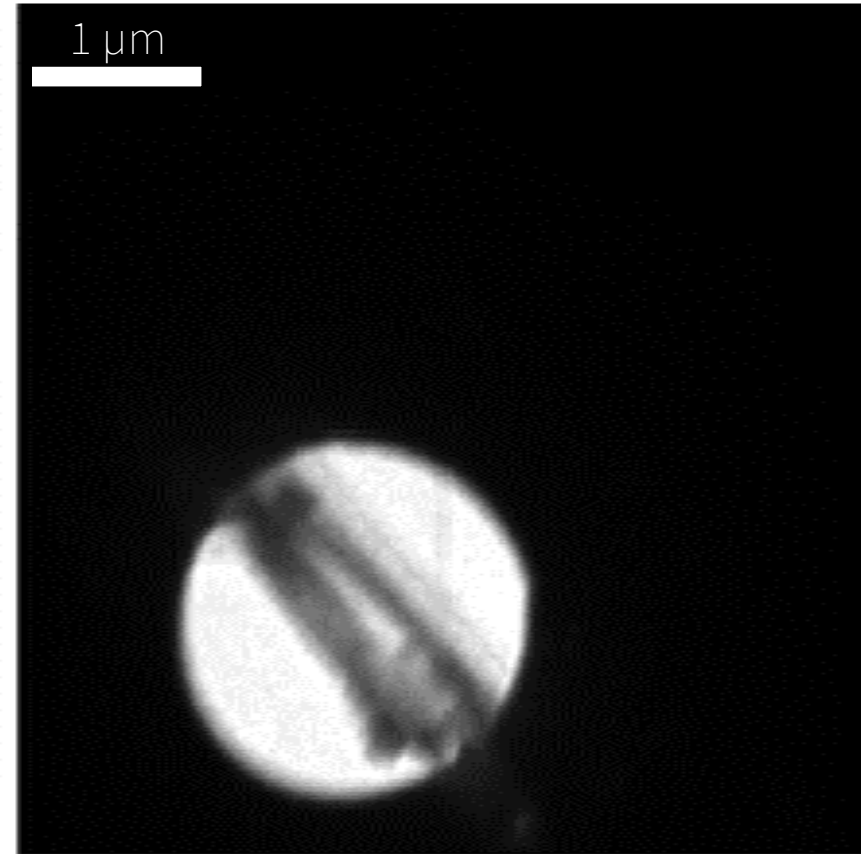
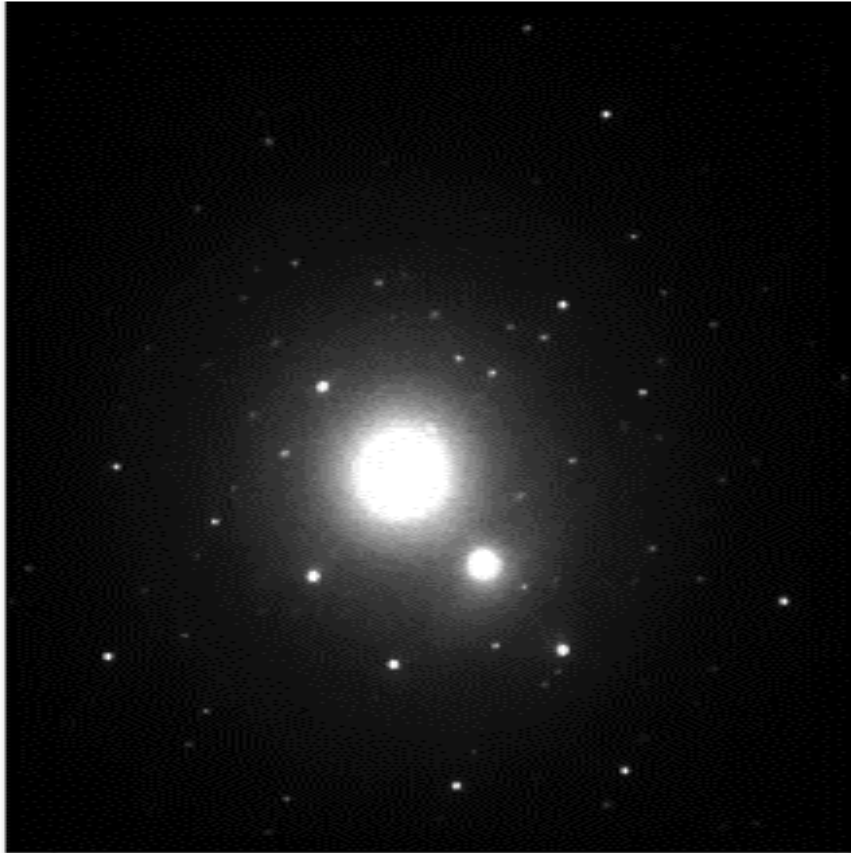
- Defocus every 10<sup>th</sup> image (IL1)
- Manually control stage position





# Automated crystal tracking strategy



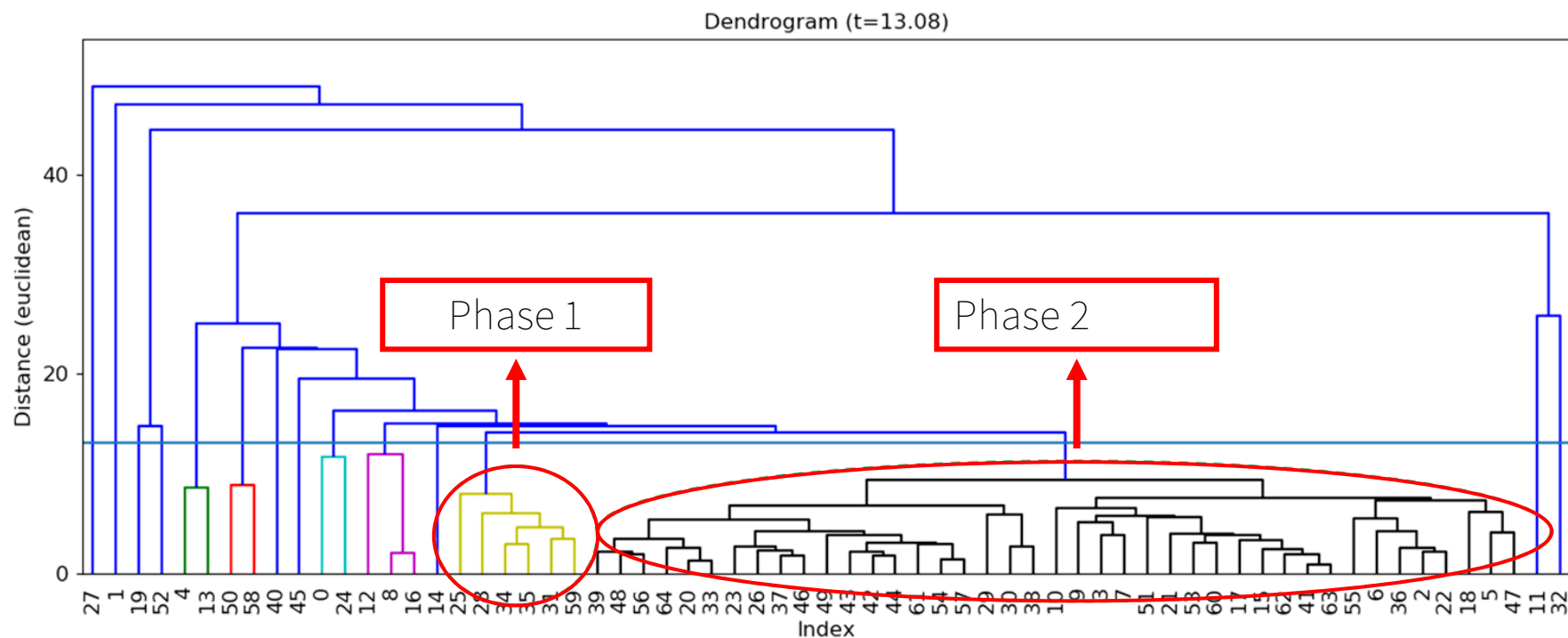


JEOL 2100-LaB<sub>6</sub> @ 200 kV (Timepix)  
Rotation: -44.0 to 47.4° @ 0.76°/s (91.4°)  
Exposure: 0.5 s, oscillation angle: 0.39°

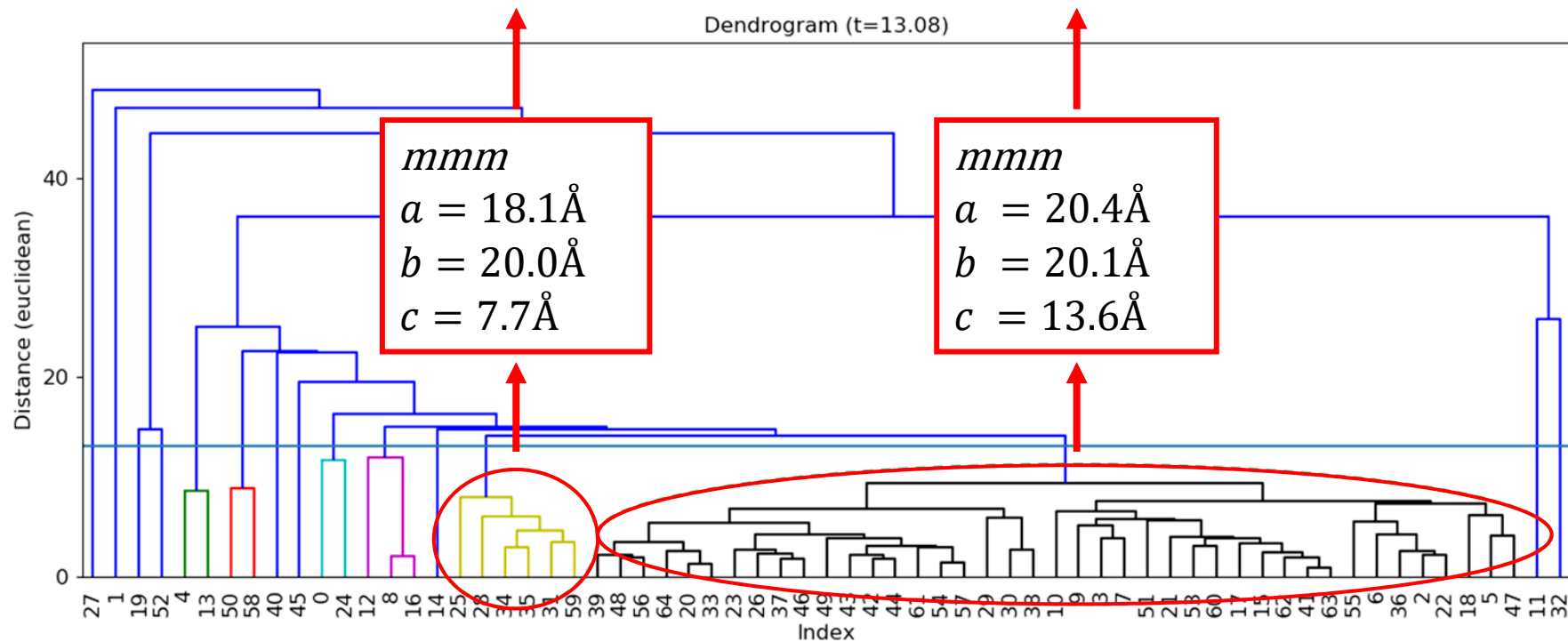
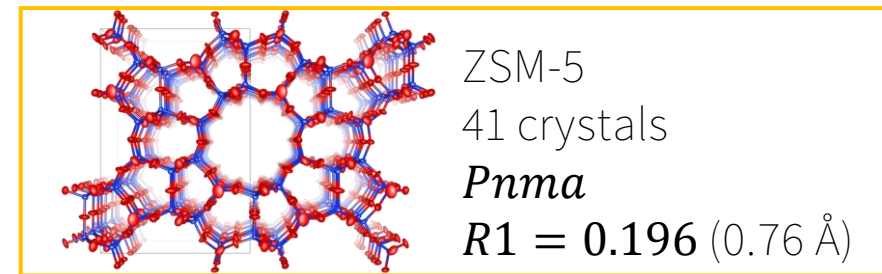
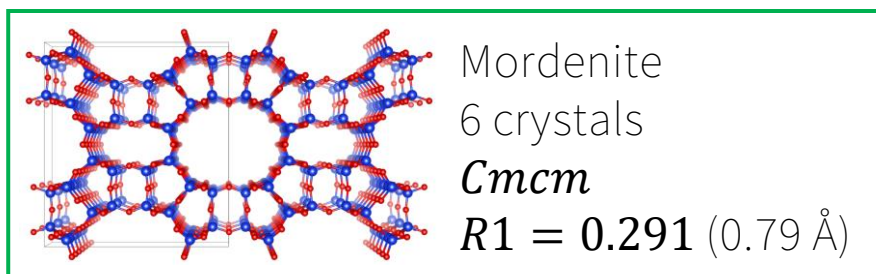


# Lattice-based clustering

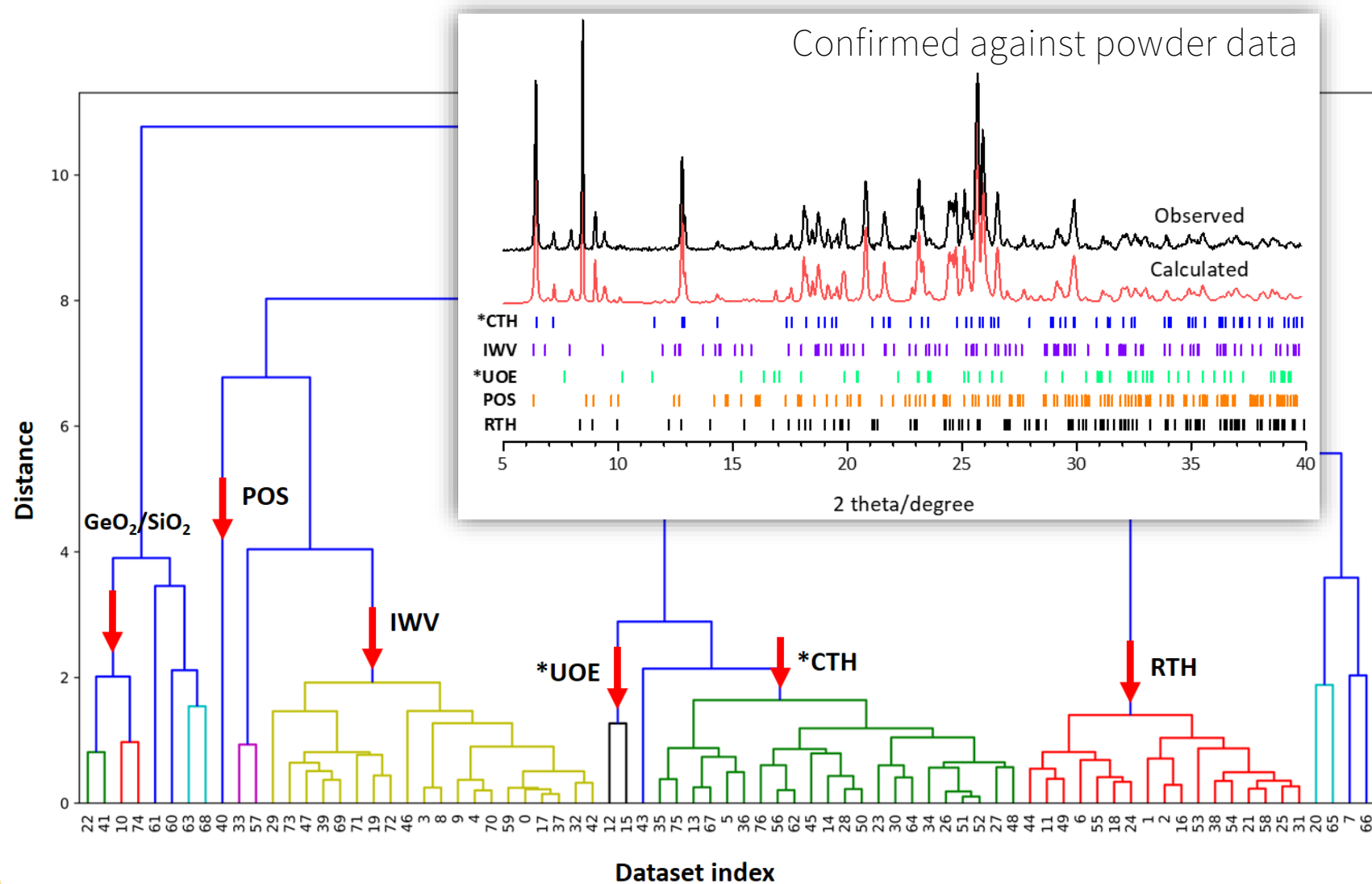
- Polycrystalline mixture of 2 phases
- SerialRED data from 89 crystals indexed using XDS



# Lattice-based clustering



# Phase identification on a real-world sample



Sample  
Single batch  
1 organic template

Data collection  
JEOL 2100  
Timepix camera

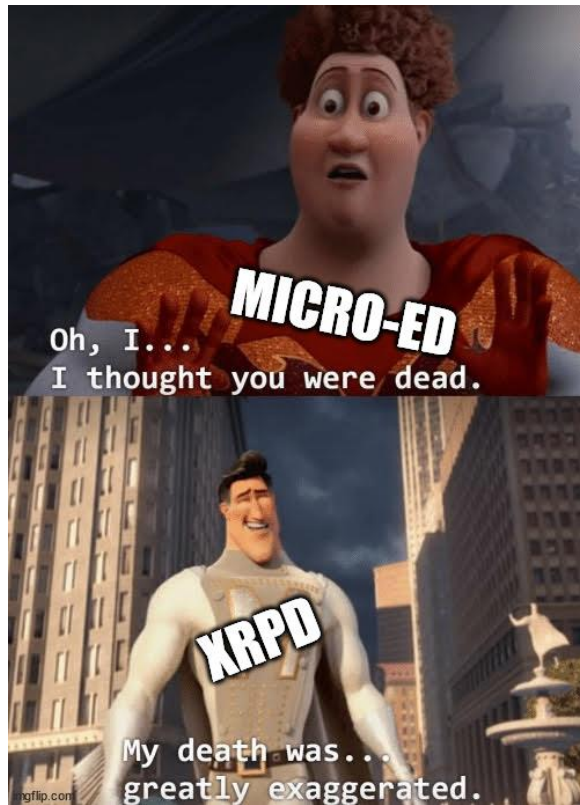
7 hours measurement  
321 crystals  
122 >20° rotation  
74 Indexed (DIALS)



**In the age of electrons, do we still need powder diffraction?**



# In the age of electrons, do we still need powder diffraction? – **Absolutely!**



- The data are mostly **complementary**
- XRPD methods are well-established for bulk material characterization
- Structure determination from XRPD data only has always been difficult
- ED data are great for single-crystal structure determination
- Potential for applying ED to phase identification problems
- Choice of method depends on the question



# Let's stay in touch



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e