

ECM-30, Basel, CH  
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# Serial Crystallography using Electron Diffraction

Stef Smeets

Dept. of Materials and Environmental Chemistry

Stockholm University, Sweden



SWISS NATIONAL SCIENCE FOUNDATION



Stockholm  
University

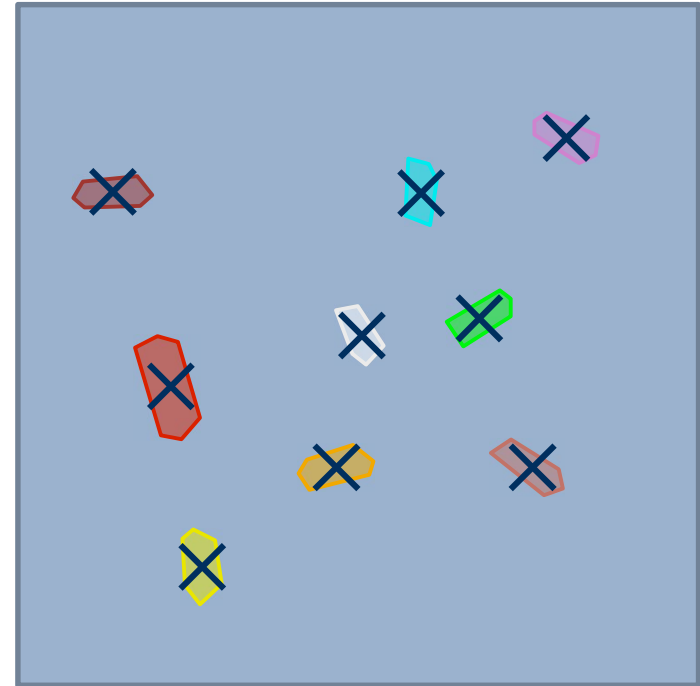
# Serial crystallography

- Randomly oriented crystals
- 1 crystal = 1 diffraction pattern
- Collect and combine data on large number of crystals
- Single-crystal-like data set



# Why use a TEM for serial crystallography?

- Electron beam is very intense
- Crystals can be located directly
- All images contain information
- There is one in many labs



# Advantages of serial electron diffraction

- Beam damage is avoided
- No (re)alignment required
- No rotation needed
- Fully automatic data collection

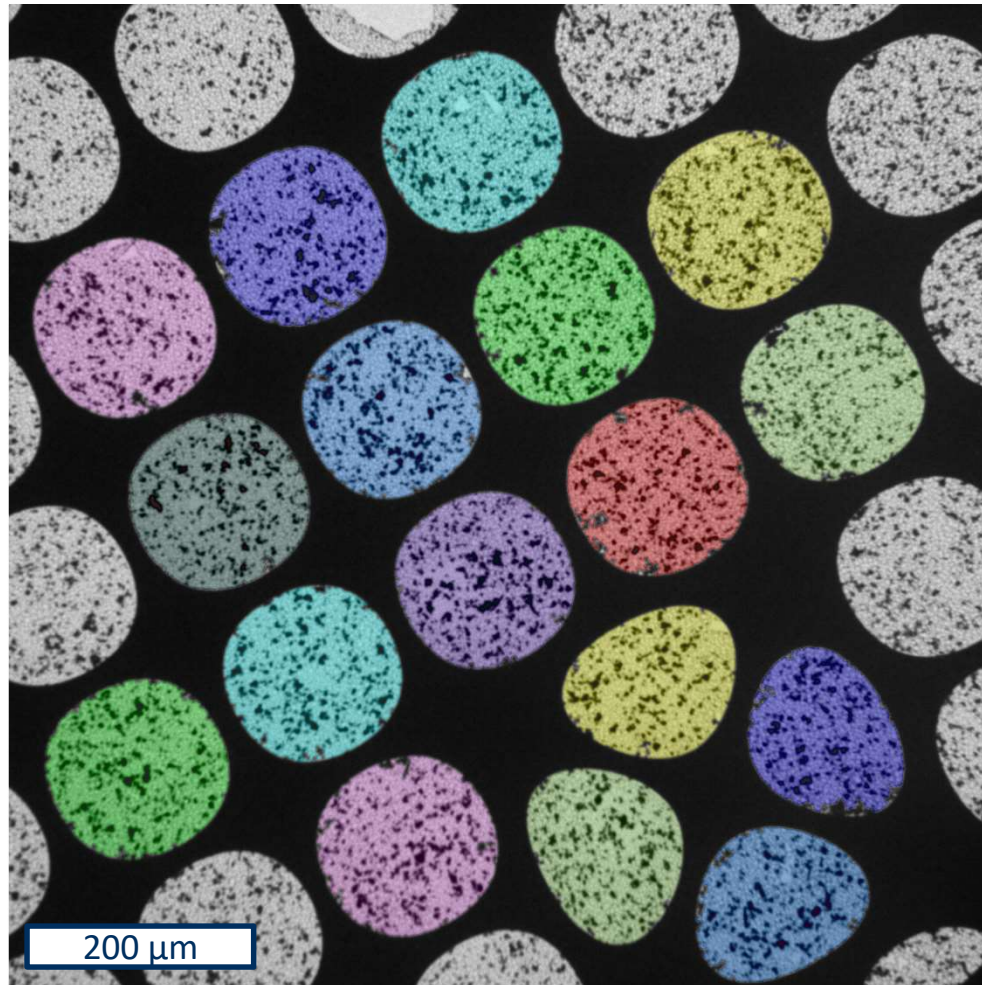
**Develop serial crystallography using electron diffraction  
for beam-sensitive materials**

- Organic/inorganic materials, pharmaceuticals
- Small unit cells ( $<25000\text{\AA}$ )

# Challenges

- Data collection
  - How to collect ED data on 500-5000 crystals?
  - How to locate and shoot crystals automatically?
- Indexing
  - Can single frames be indexed reliably?
- Data processing
  - How to merge and scale data collected on a large number of crystals?
- Structure solution
  - Are the data good enough for structure determination?

# Data collection: mapping the grid

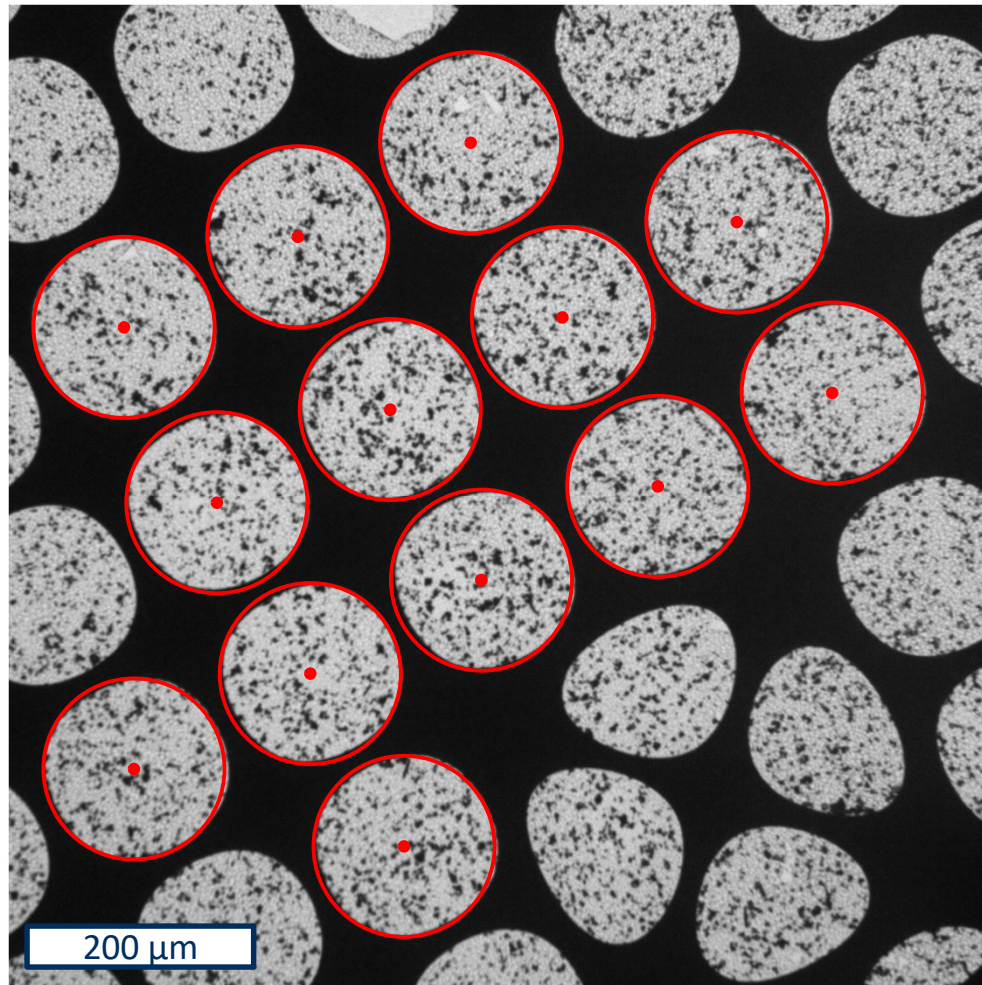


100x magnification  
150µm holes

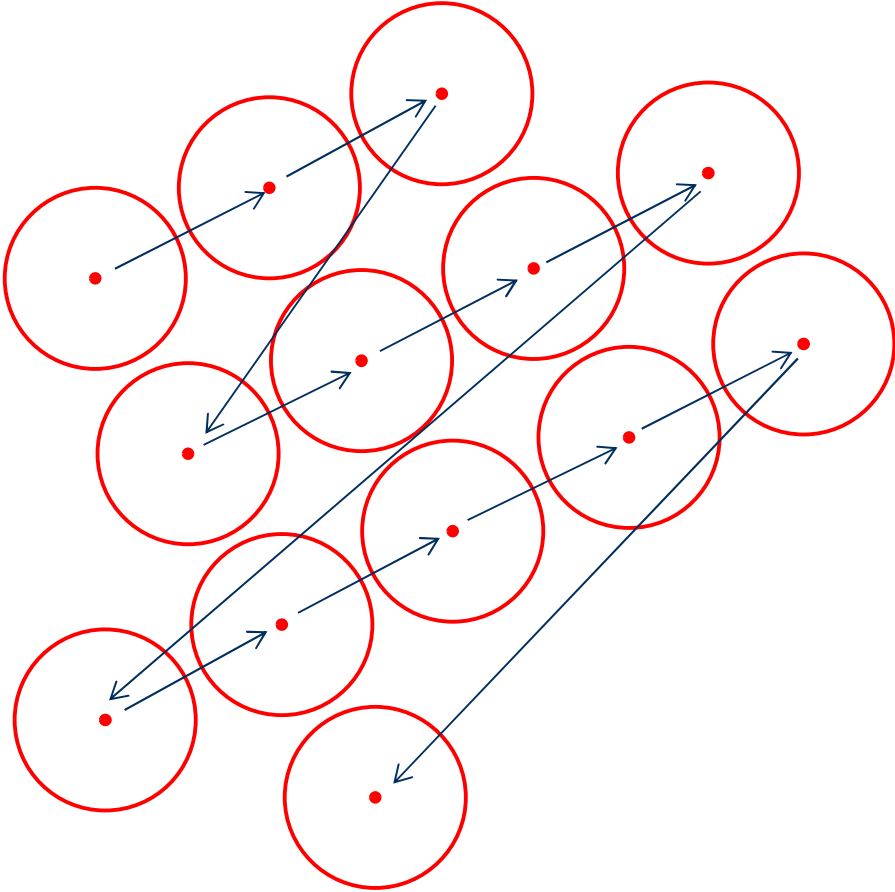
JEOL JEM2100-LaB<sub>6</sub>  
200 kV  
Orius CCD

Image processing:  
scikit-image

## Data collection: map the grid

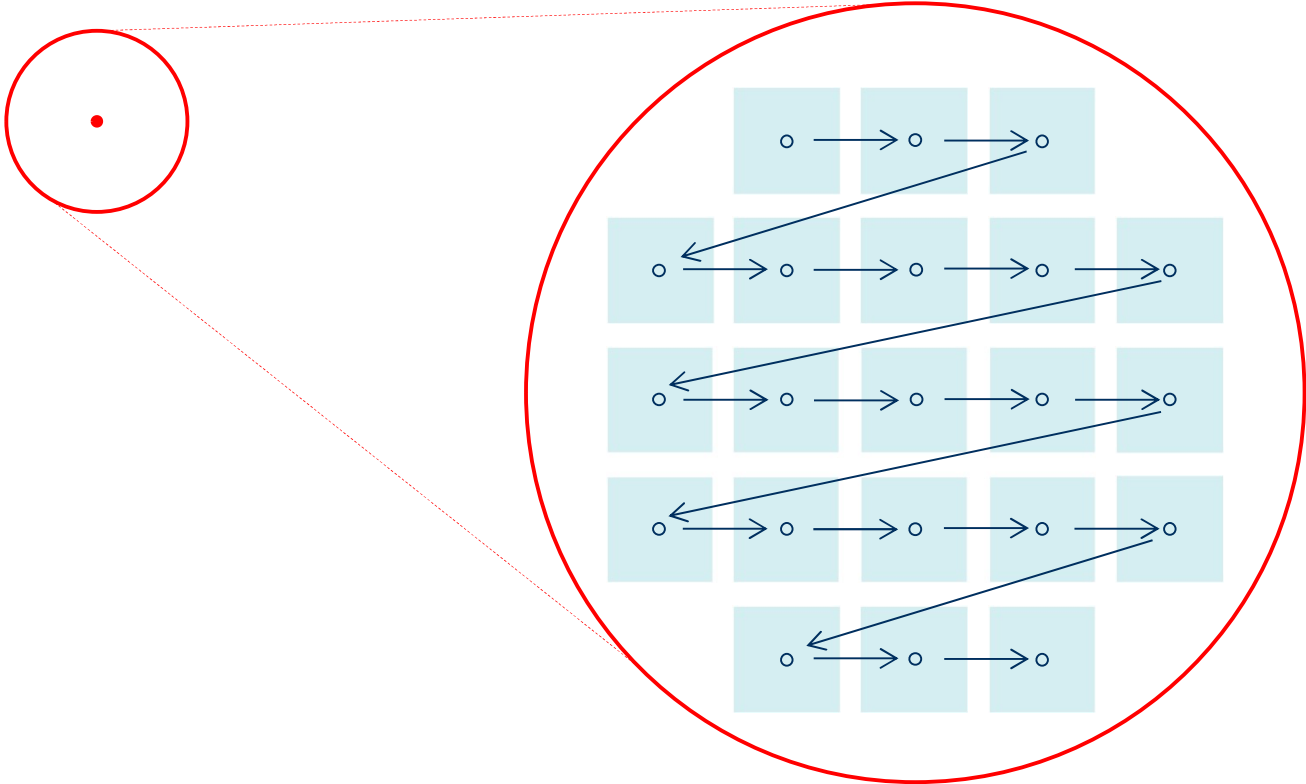


# Data collection: map the grid



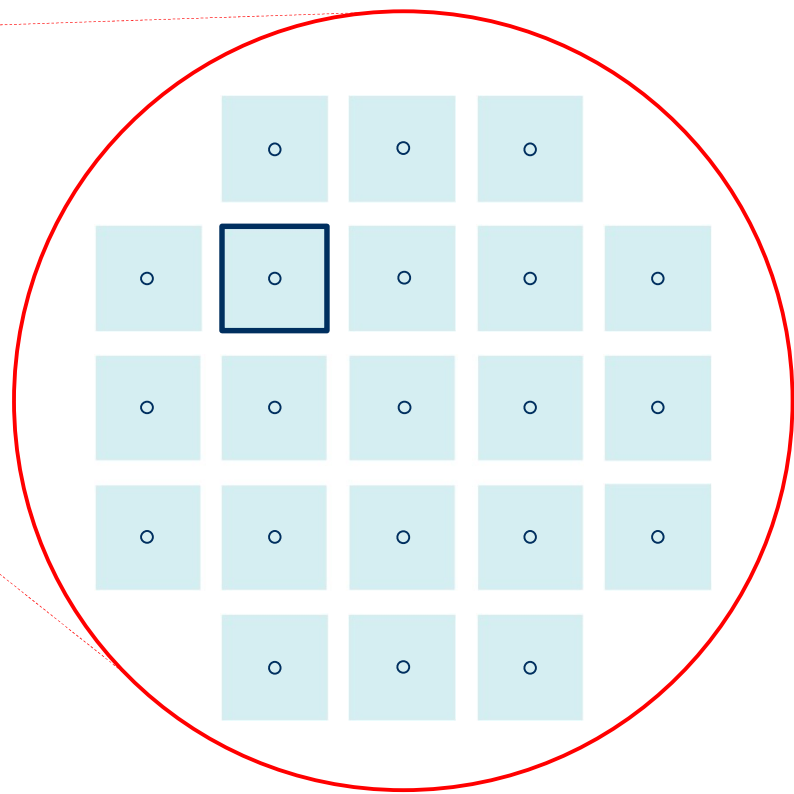
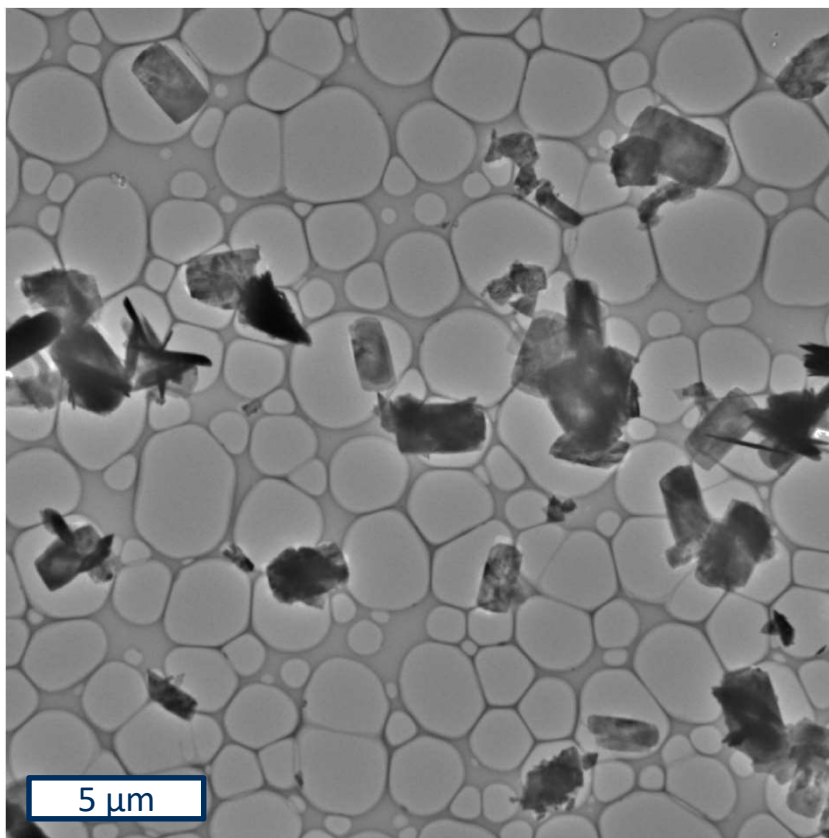


# Data collection: map hole



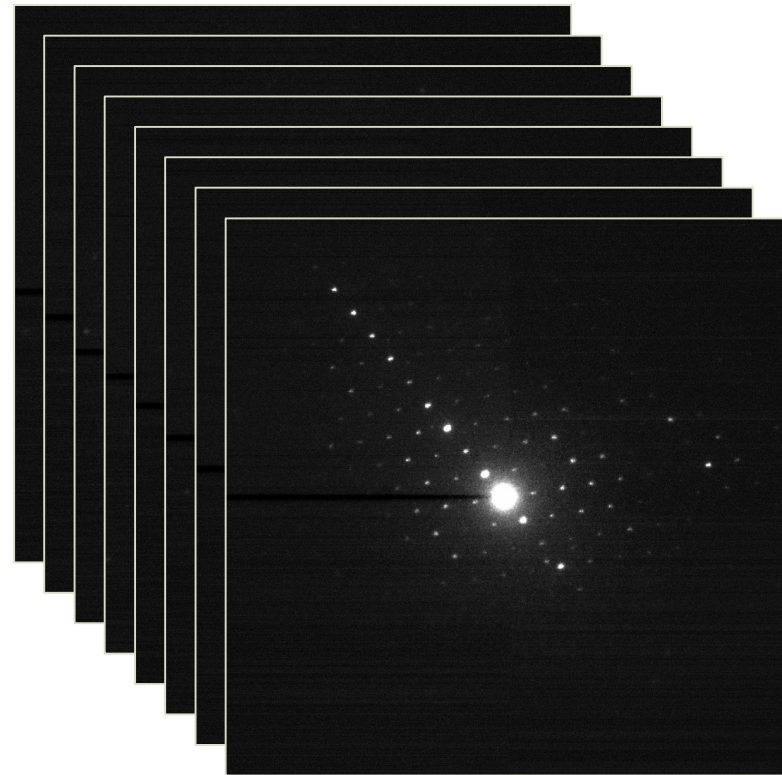
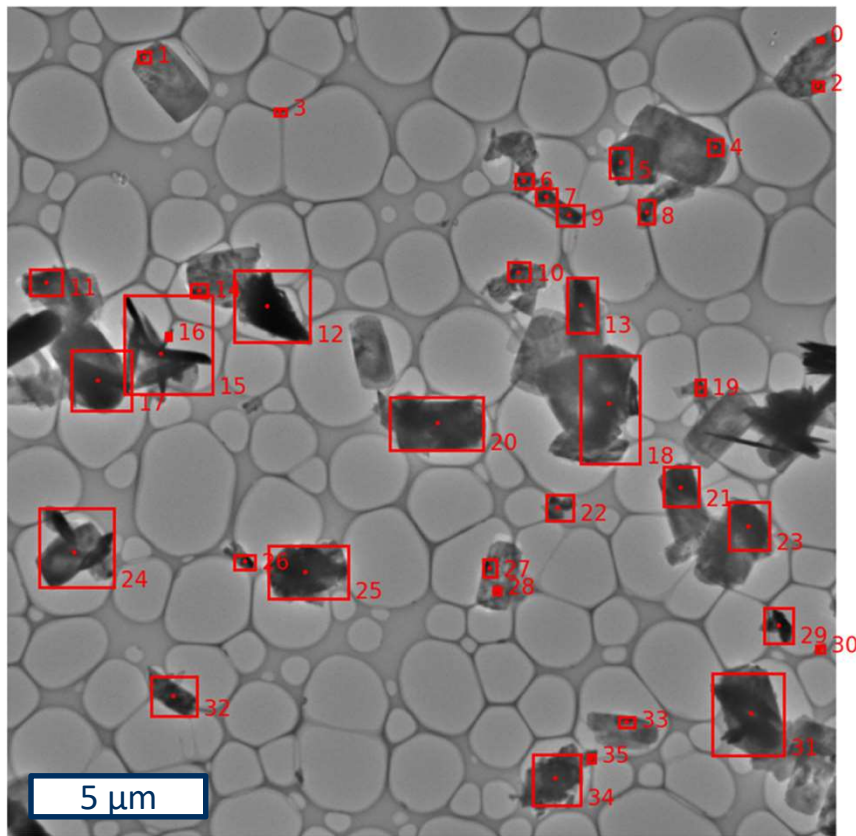
# Data collection: find crystals

2500x magnification



# Data collection: diffraction

- Beamshift -> Focused nanobeam to collect ED data
- Potentially: 13 holes x 21 positions x 20 crystals = 5500 data sets

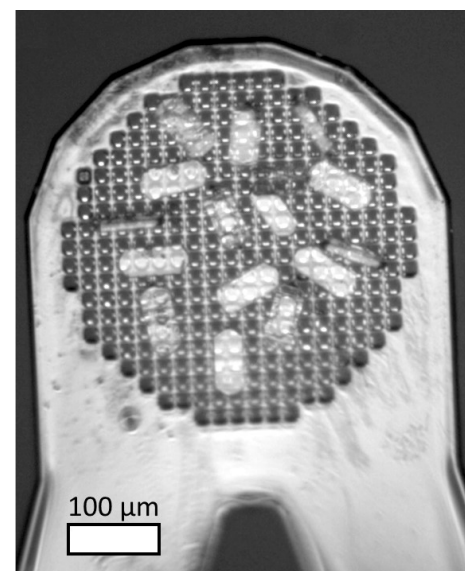


# Indexing

## *Challenges*

- Get accurate crystal orientations from a single frame
- Small unit cells (limited number of reflections)
- Crystals can overlap
- Find criteria to accept/reject orientations
  
- Assume unit cell is known

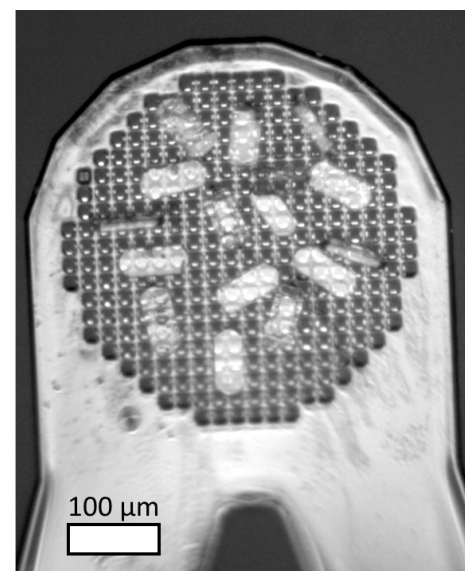
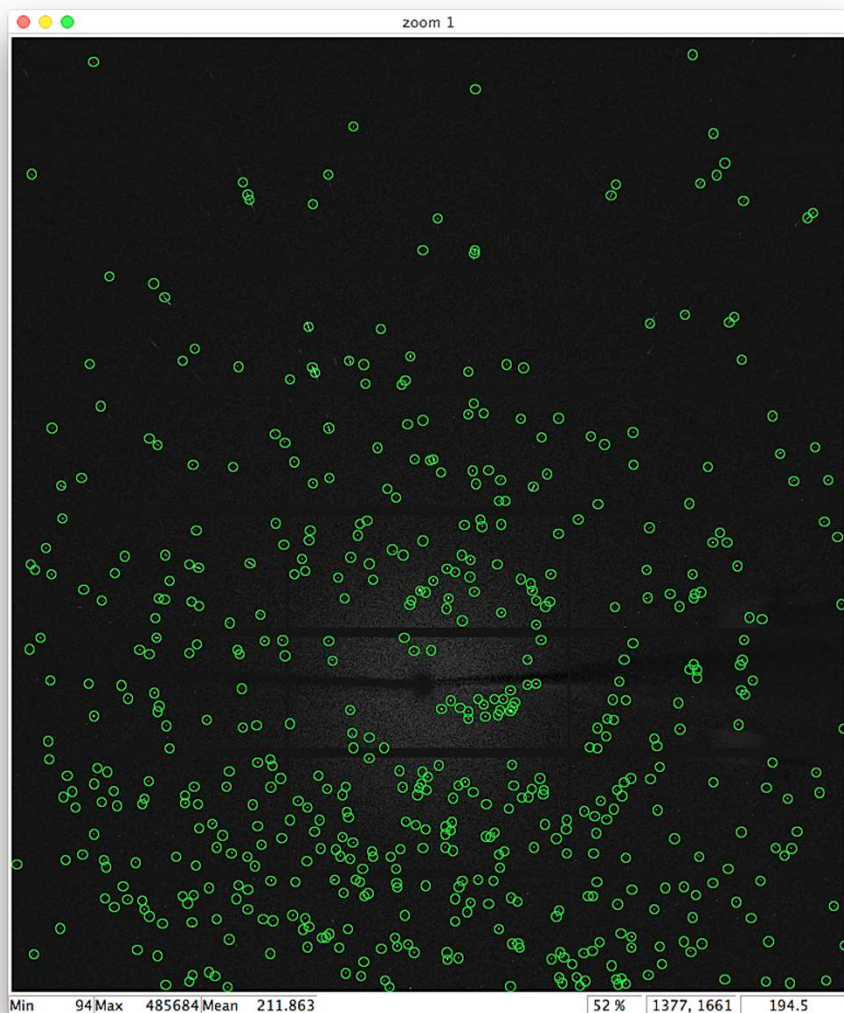
# Indexing



15 crystals of ZSM-5

SNBL (ESRF),  $\varphi$ -scan,  $1^\circ$  / step  
4.6% bandwidth

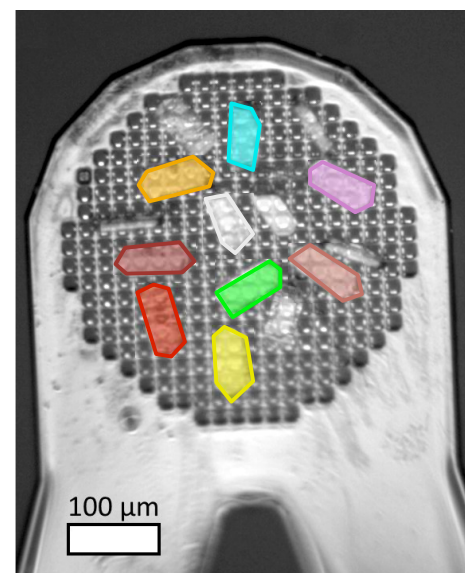
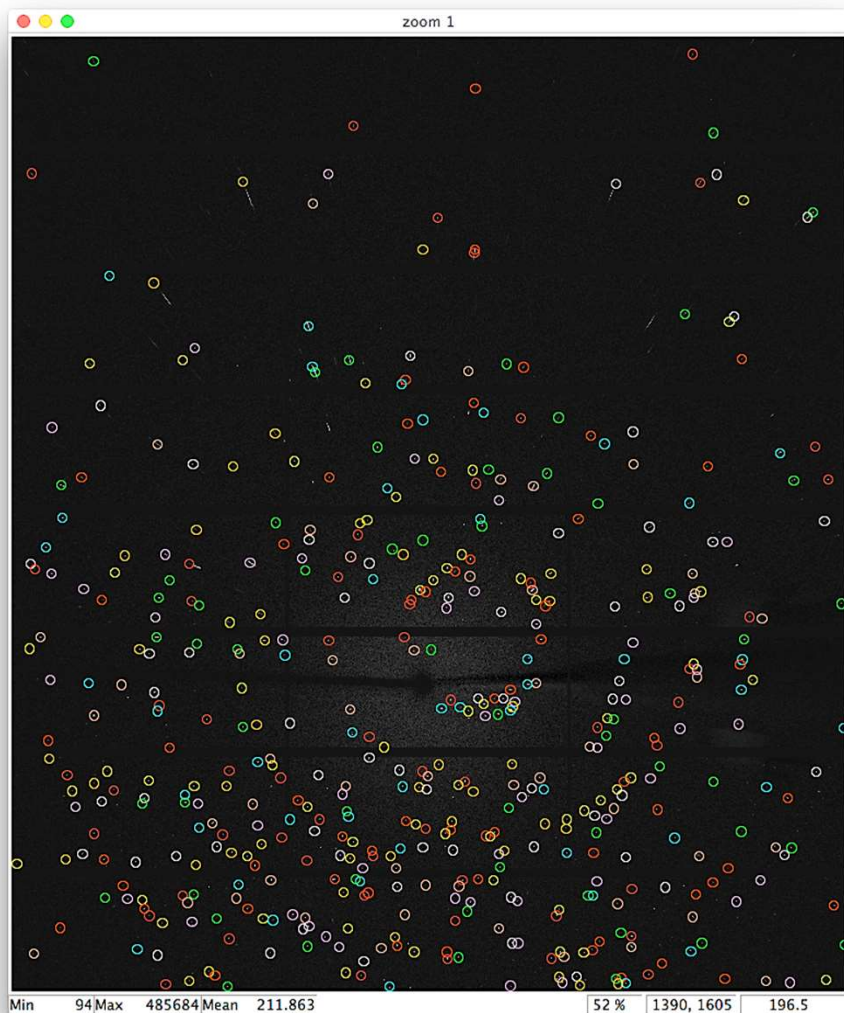
# Indexing: peak finding



15 crystals of ZSM-5

548 reflections

# Indexing: brute force algorithm



15 crystals of ZSM-5

548 reflections (9 crystals)

# Indexing: single frames

Tests using single-crystal rotation electron diffraction (RED) data

Tilt range:  $\sim 120^\circ$

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sample	space group	volume ( $\text{\AA}^3$ )	# frames	indexed
Garnet	<i>Ia-3d</i>	1737	1204	96%
ZSM-5	<i>Pnma</i>	5327	1471	90%
Ni-Se-O-Cl system <sup>1</sup>				

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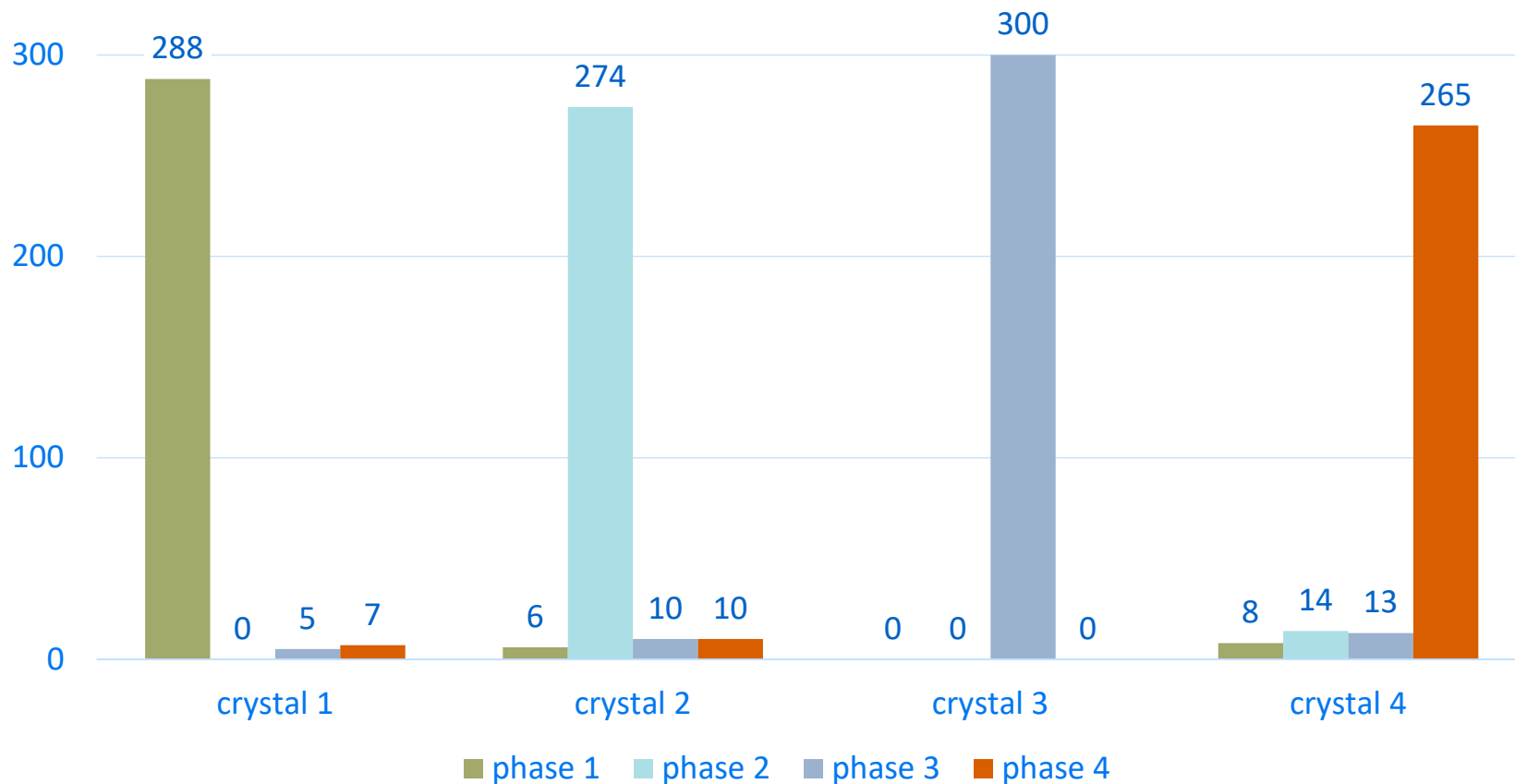
<sup>1</sup>Y. Yun *et al.* (2014) *J. Appl. Cryst.* **47**, 2048



# Indexing: phase matching

Tests using RED data on multiphase Ni-Se-O-Cl system

Tilt range:  $\pm 15^\circ$  (300 frames)



# Data processing: merging & scaling

How to merge *hkl* listings of 500+ data frames?

## *Challenges*

- Variations in diffracting volume (scaling)
- Dynamical scattering
- Missassignment of reflection indices
- Partial intensities
  
- *Structure solution of oxides from zone axes precession diffraction data* (Klein, Z. Kristallogr., 2012)
- *Refining structures against reflection rank: an alternative metric for electron crystallography* (Eggeman & Midgley, Acta Cryst. A, 2012)

# Data processing: ranking

- *SerialRank: Spectral Ranking using Seriation* (Fogel et al., ArXiv, 2014)

# Data processing: ranking

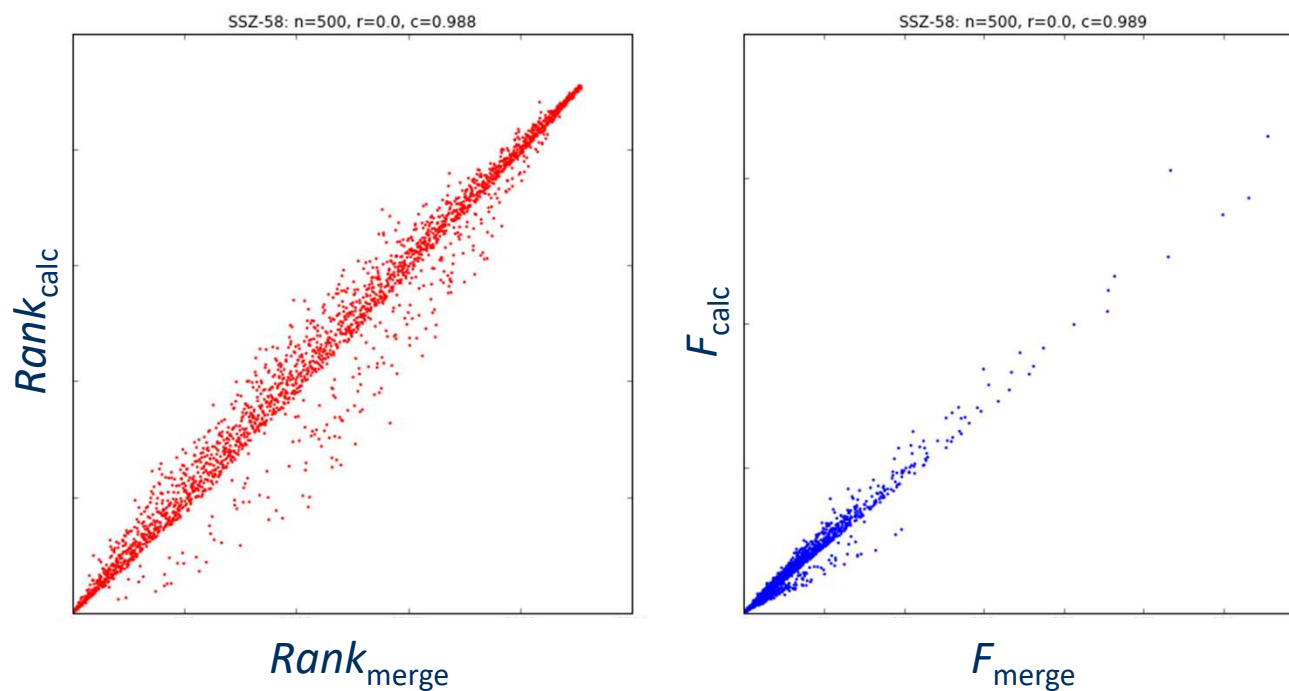
- *SerialRank: Spectral Ranking using Seriation* (Fogel et al., ArXiv, 2014)

	h k l	Frame 1 Rank	Frame 2 Rank	Frame 3 Rank	Frame 4 Rank	Frame 5 Rank	Combined Rank
Strong ↓ weak	3 1 1	1	4	1	4	1	2
	3 1 3	3	7	3	1	3	3
	0 0 2	2	1	2	3	2	1
	1 1 7	4	2	4	2	5	4
	1 3 1	5	6	9	10	4	6
	2 2 2	6	3	5	9	6	5
	1 1 3	9	10	6	7	8	8
	1 1 1	7	5	10	5	7	7
	1 1 5	10	8	7	6	10	9
	2 0 2	8	9	8	8	9	10

- Find rank -> Assign intensities based on reflection rank

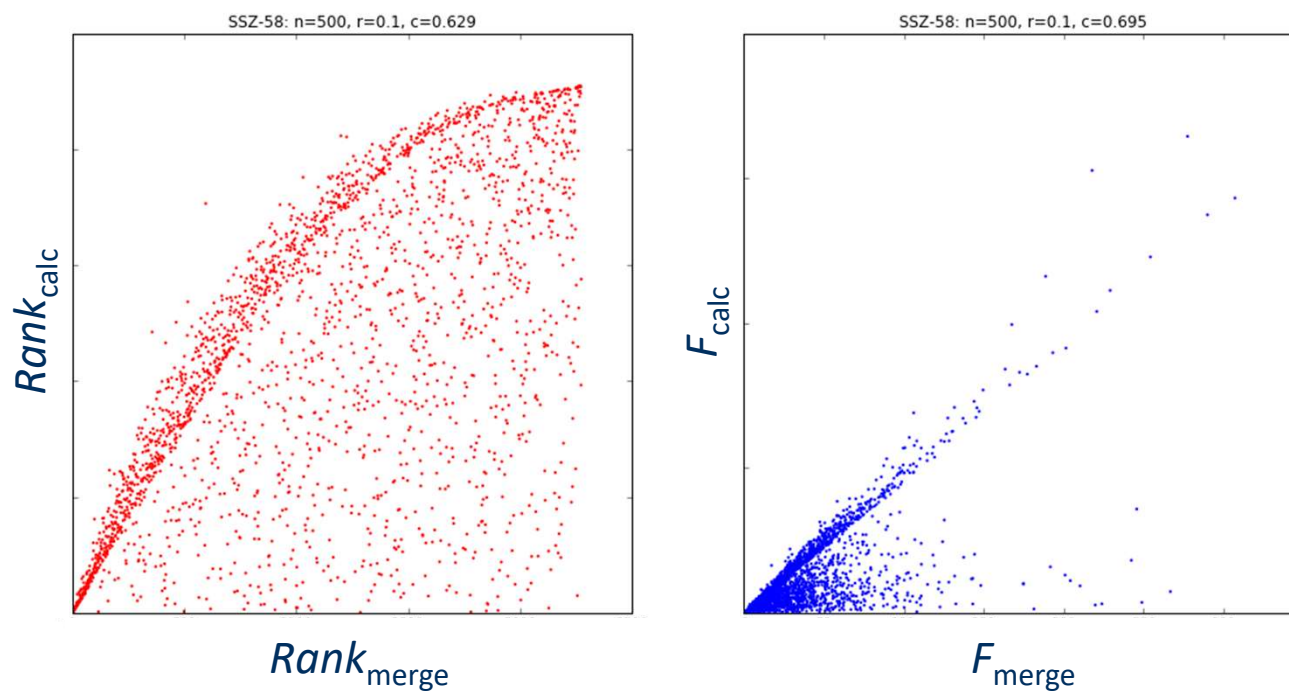
# Data processing: conventional merge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale



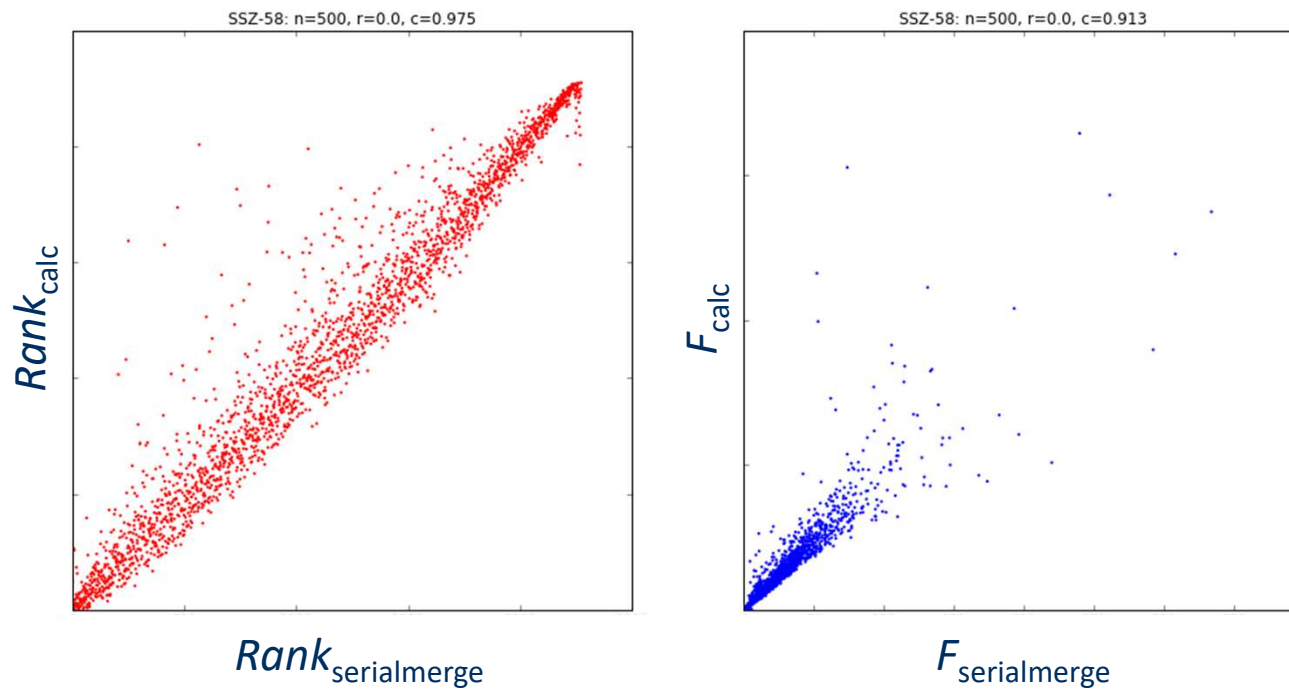
# Data processing: conventional merge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 10% of indices



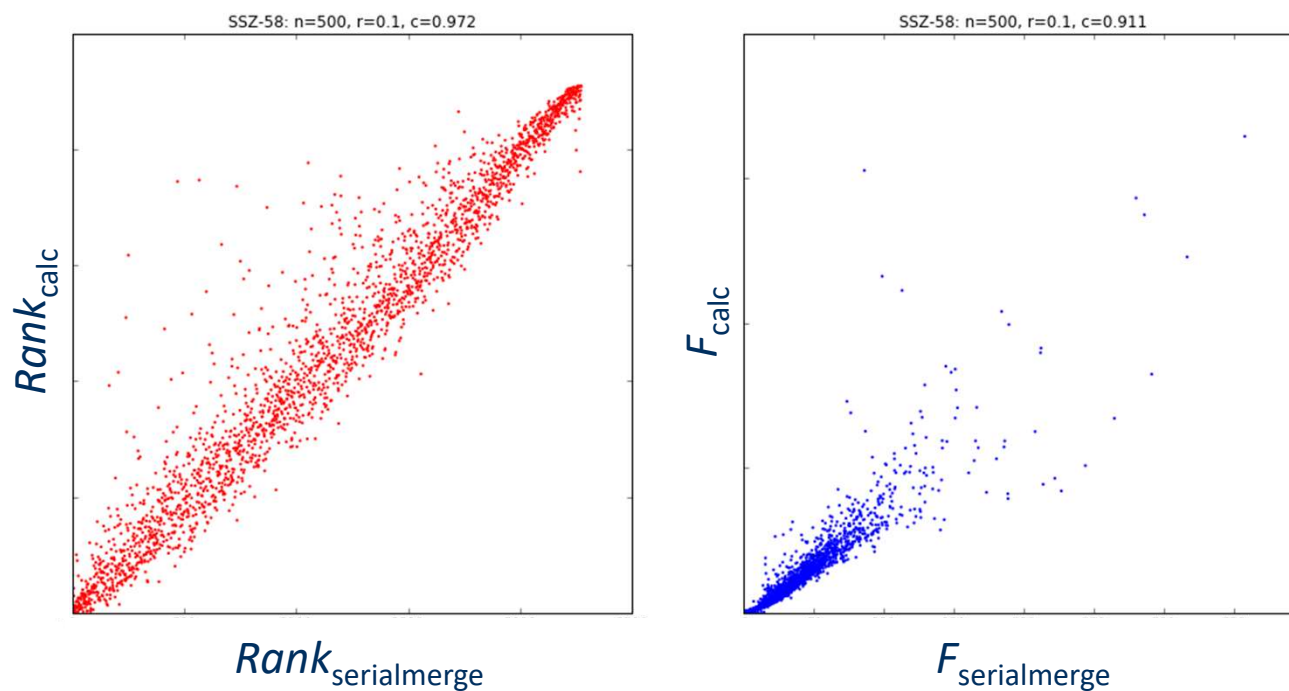
# Data processing: SerialMerge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale



# Data processing: SerialMerge

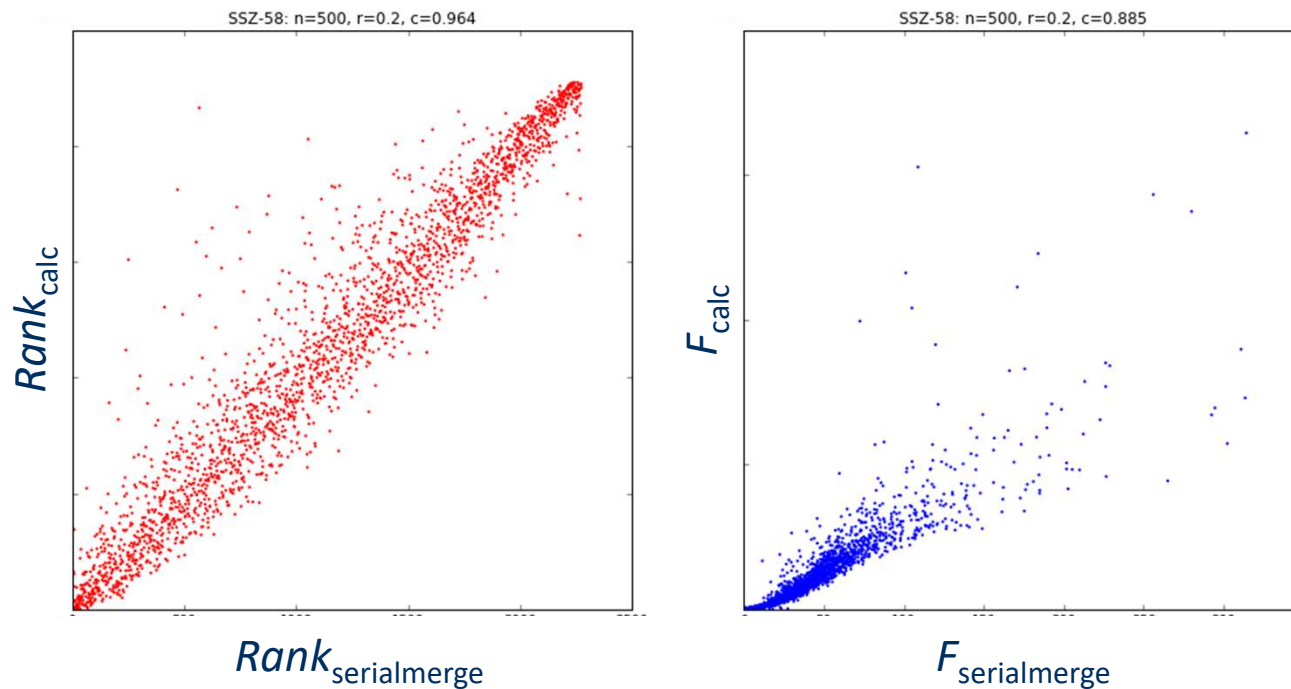
- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 10% of indices





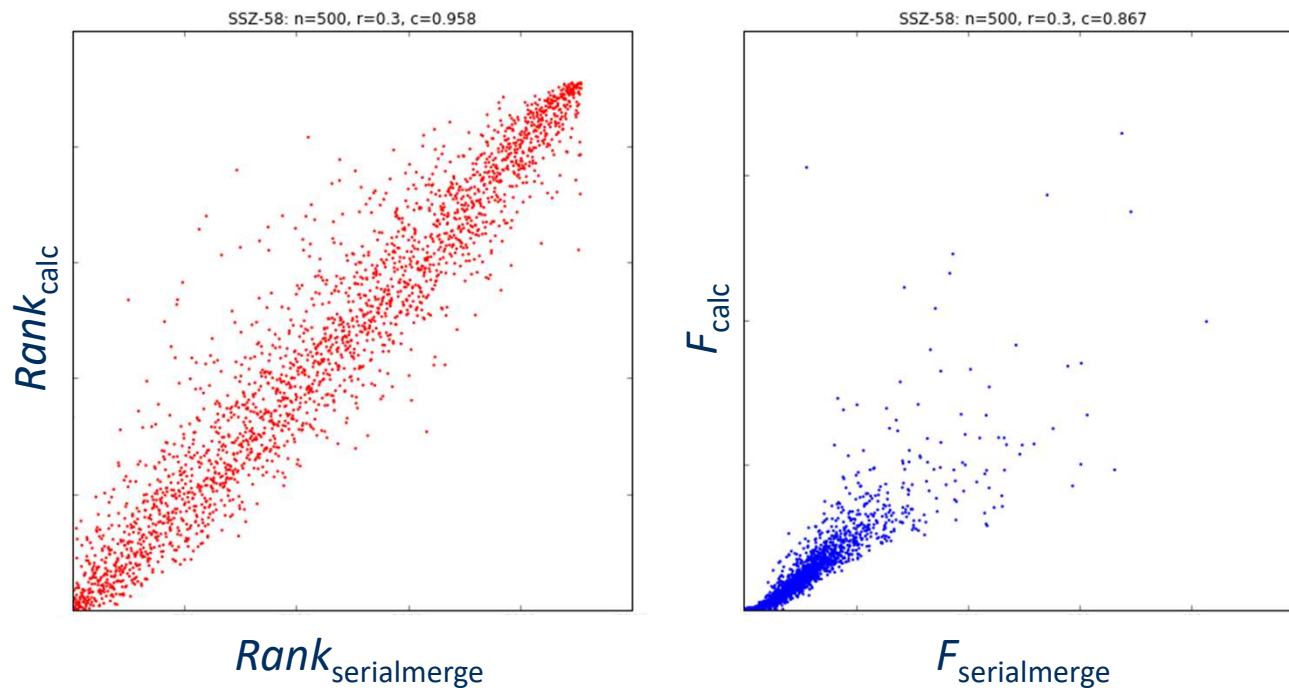
# Data processing: SerialMerge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 20% of indices



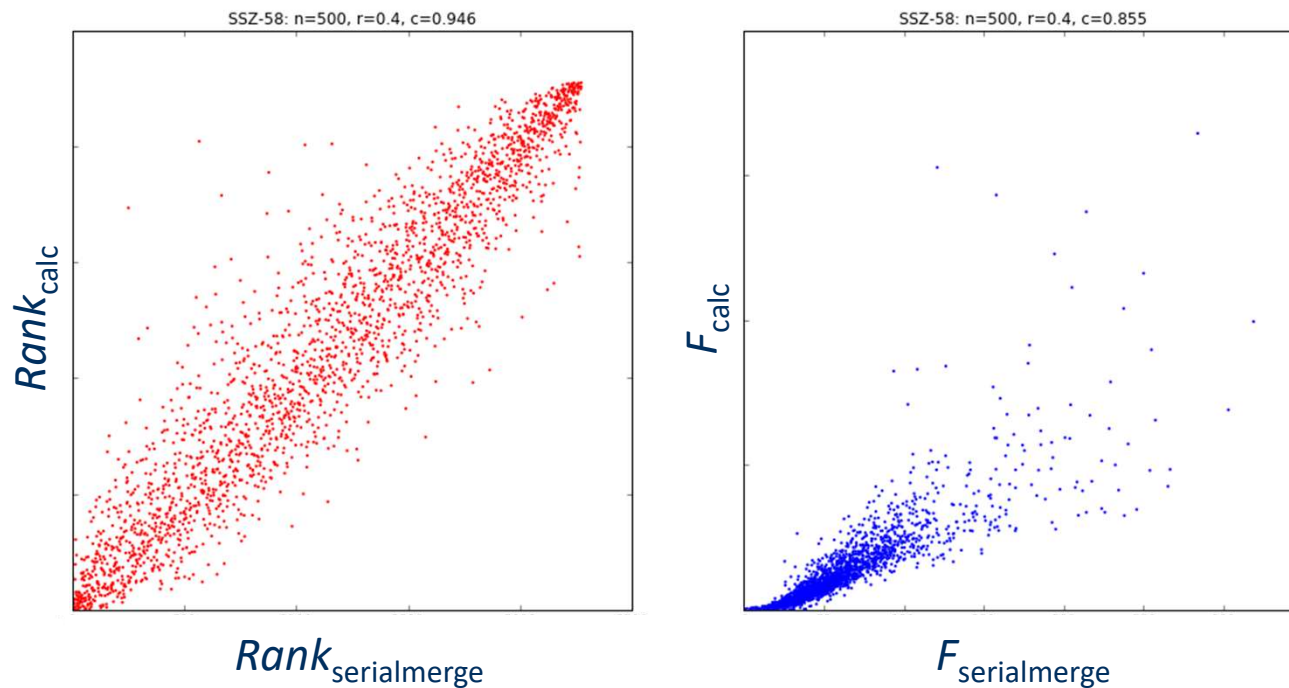
# Data processing: SerialMerge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 30% of indices



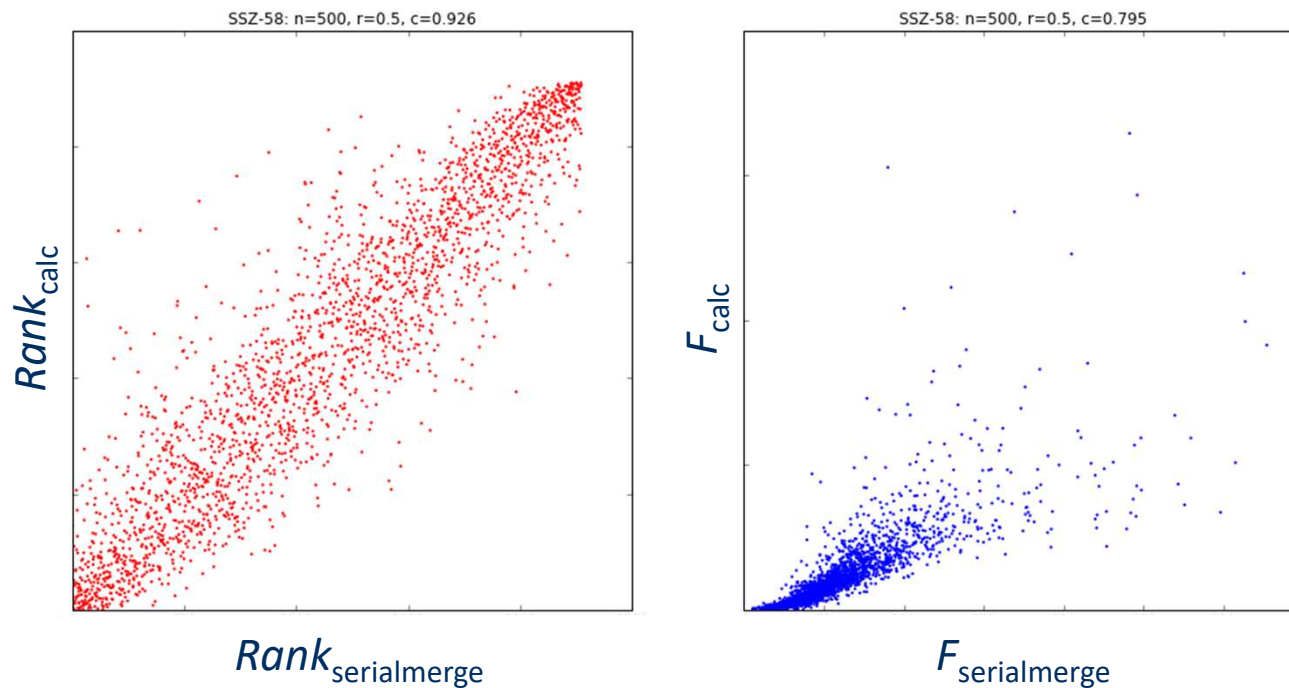
# Data processing: SerialMerge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 40% of indices



# Data processing: SerialMerge

- 500 simulated frames on zeolite SSZ-58
- Partial intensities,  $d_{\min} = 1\text{\AA}$ , 2250 uniq. refs.
- Random scale, shuffle 50% of indices



# Structure solution: simulated data

- Simulate 500 frames of 6 zeolites
- Random scale, shuffle up to 80% of indices
- Program FOCUS

	# atoms	Spgr	Vol. (Å <sup>3</sup> )	10%	20%	30%	40%	50%	60%	80%
<b>SSZ-55</b>	11	<i>C2221</i>	1396	yes	yes	yes	yes	yes	yes	yes
<b>SSZ-59</b>	24	<i>P21nm</i>	891	yes	yes	yes	yes	yes	no	no
<b>SSZ-53</b>	25	<i>C2/c</i>	3562	yes	yes	yes	yes	yes	no	no
<b>SSZ-45</b>	29	<i>Fmmm</i>	10697	yes	yes	yes	no	no	no	no
<b>SSZ-58</b>	38	<i>Pmma</i>	4051	yes	yes	yes	no	no	no	no
<b>SSZ-56</b>	45	<i>P2<sub>1</sub>/m</i>	3277	yes	yes	yes	no	no	no	no

# Summary & Conclusions

- Data collection
  - Work in progress
- Indexing
  - Single frames can be indexed reliably
- Data processing
  - SerialMerge algorithm avoids scaling and is insensitive to intensity inaccuracies
- Structure solution
  - Merged data quality good enough for structure determination
- Software
  - Automated data collection: *Instamatic*
  - Data processing/indexing: *SnapKit*
- Applications
  - Structure determination of beam sensitive materials
  - Phase analysis and identification