

# Chemical Crystallography and Structural Chemistry

(VO 270287)

Lecture 11

25<sup>th</sup> June 2020

Dr. Tim Grüne

Centre for X-ray Structure Analysis

Faculty of Chemistry

University of Vienna

tim.gruene@univie.ac.at

## Previous Lecture

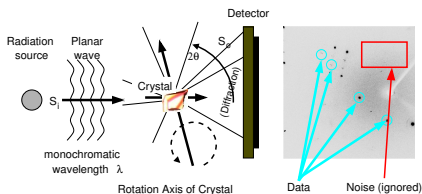
1. Absolute structure / Chirality
  - anomalous signal: breakdown of Friedel's law
  - Flack-Parameter, Parsons' Q-value
  - heavy atom method
2. Charge density refinement
  - limits of the independent atom model
  - multipole expansion
  - chemical features of electron density

## Today's Lecture

1. Structure Determination with Crystallography
2. Electron and X-rays
3. Applications for Electron Crystallography
4. Practical Aspects
5. Radiation Damage
6. Dynamic Scattering

# Structure Determination by Single Crystal Diffraction

## Data Collection for Single Crystal Structure Analysis



- Rotation (>1966): **Contiguous** recording of reciprocal space
- hybrid pixel detectors (>2002): shutterless, *i.e.* continuous data collection
- Spot position: wave type independent
- Intensity: wave type dependent
- Radiation source (wave): X-rays, electrons, or neutrons

## 3D Electron Crystallography

- > late 1990s, as opposed to 2D electron crystallography
- confusingly many terms (ADT , RED, EDT, PEDT, MicroED, ...)
- Historical “dispute”, Ute Kolb, Mainz University,  $\approx$  2007 (ADT), Xiaodong Zou & Sven Hovmöller, Stockholm University  $\approx$  2011 (RED)
- technical term: “3D Electron Diffraction”, Enrico Mugnaioli (PSI 2017; IUCrJ (2019), 6, 178–188)
- “3D”:
  1. Collection of 3D reciprocal space
  2. 3D crystals:  $\geq$  10–15 unit cell in each direction; typically 200–1000nm per dimension

## Spot Position

- Spots positions according to Laue Conditions and orientation of Unit Cell:

$$(\vec{S}_o - \vec{S}_i) \cdot \vec{a} = h$$

**and**  $(\vec{S}_o - \vec{S}_i) \cdot \vec{b} = k$

**and**  $(\vec{S}_o - \vec{S}_i) \cdot \vec{c} = l$

- Monochromatic wave:  $\vec{S} = (\vec{S}_o - \vec{S}_i)$  depends on wavelength  $\lambda$  and experimental geometry
- Spot **position**  $\Leftrightarrow$  Crystal lattice, independent from radiation type
- Resolution  $d_{hkl}$  of a spot from position on detector *via* Bragg's law,  $\lambda = 2d_{hkl} \sin(\theta)$

## Spot Intensity

- Spots **intensity** depends on physics of interaction

**X-rays** interact with electrons, crystallographic map corresponds to electron density (number of electron per Volum,  $e^-/A^3$ ).

**Electrons** interact with electrostatic potential from electrons + nucleus  
( $\varphi(\vec{r})$ )

**Neutrons** interact with nucleus *via* weak interaction, and magnetic moment.

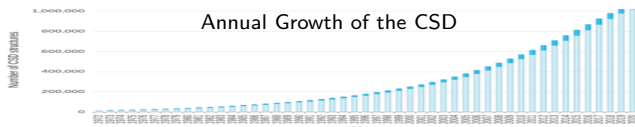
- Spot intensity  $\Leftrightarrow$  Unit cell content: where are the atoms, what type of atoms are they



## Dominance of X-ray Crystallography

- most advanced (pipelines from data collection to structure refinement)
- typical wavelength:  $\lambda = 0.8\text{--}1.9\text{\AA}$
- standard structure determination

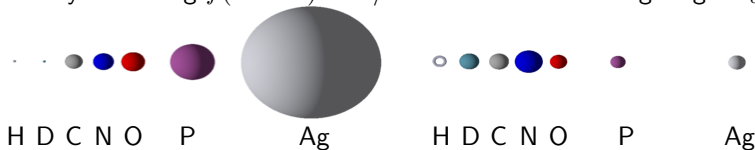
	PDB (MX)	CSD (SX)
X-ray	>140,000	>1,000,000
neutron	161	1,500
electron	111	10'ish



## Neutron crystallography

X-ray scattering  $f(2\theta = 0) = Z/\text{Å}$

Neutron scattering length  $b_c$



1. visualisation of hydrogen atoms
2. adjacent elements (e.g.  $K^+$  vs.  $Cl^-$ ,  $Zn^{2+}$  vs.  $Cu^+$ )
3. (virtually) no radiation damage
4. requires large crystals ( $\geq 1\text{mm}^3$ )

## 3D Electron Crystallography (3D ED)

- Electrons interact with electrostatic potential
- Electrons interact **much,much** stronger with matter than X-rays

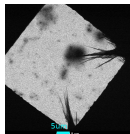
⇒ Much smaller crystals

⇒ problematic: dynamic diffraction,  $|F| \neq \sqrt{I}$

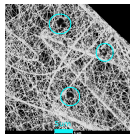
- Electron optics enable some special applications and tiny beam (5 nm diameter)

# 3D Electron Crystallography

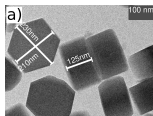
## 3D ED: small crystals



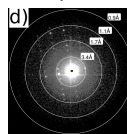
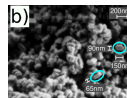
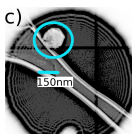
organic compound



sucrose



Silicalite-1 / ZSM-5 (Teng Li)



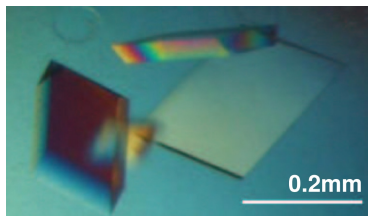
## Small Crystals

Main advantage for electron crystallography:

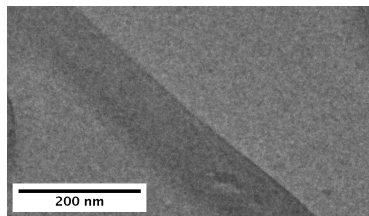
Diffraction from very small crystal ( $< 1\mu m$ )

Some instruments provide 5–10 nm beam diameter

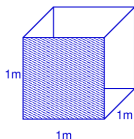
## How small is “nano”?



typical protein crystal size for X-rays



typical protein crystal size for electrons,  $100 \times 140 \times 1,700 \text{ nm}^3$



volumes compare like  $1\text{m}^3$  or 6 bath tubs of water vs.  $10\mu\text{l}$

## Effects of Crystal Volume on Diffraction Data

Reducing crystal volume reduces the resolution by (at least) two effects:

1.  $I(hkl) \propto V_{\text{crystal}}$ : 1/10 volume = 1/10 intensity
2. Henderson / Garman limit: maximum dose per volume before resolution is halved: 1/10 volume = 1/10 dose before radiation damage destroys crystal

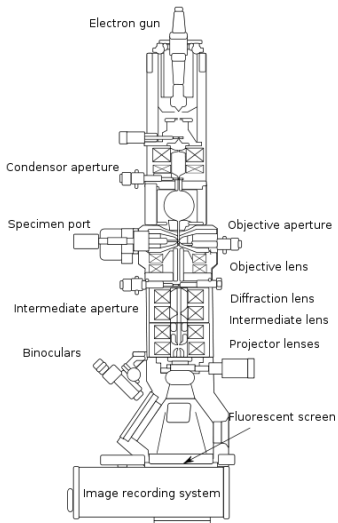
From (1): In order to record the same quality diffraction pattern from a 10 times smaller crystal requires 10 times more intense beam.

From (1)+(2): This makes the crystal die 100 times faster



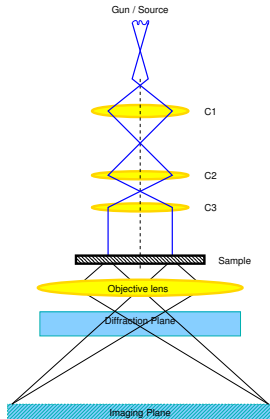
# Instrumentation for Electron Diffraction

## Electron Microscopes



(Wikipedia)

## The Lens System

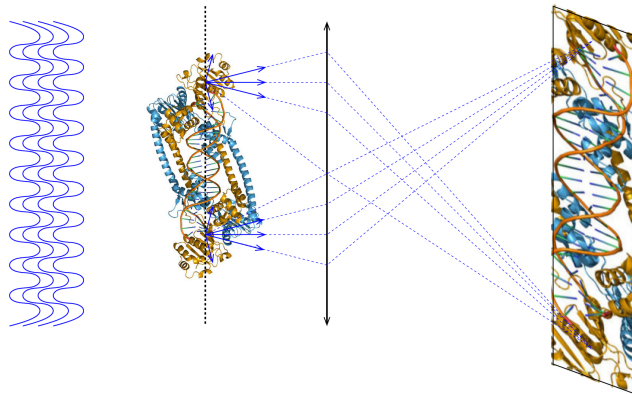


- Lenses C1–C3 shape beam
- Crystallography: Parallel beam
- Objective lens: sets effective detector distance to backfocal plane = diffraction mode
- C3 not present in all microscopes

**Lenses cause distortions.**

see e.g. Zuo & Spence, “Advanced Transmission Electron Microscopy”, Springer  
Carter & Williams, “Transmission Electron Microscopy”, Springer

## Electron Microscope: Imaging Mode



Plane Wave

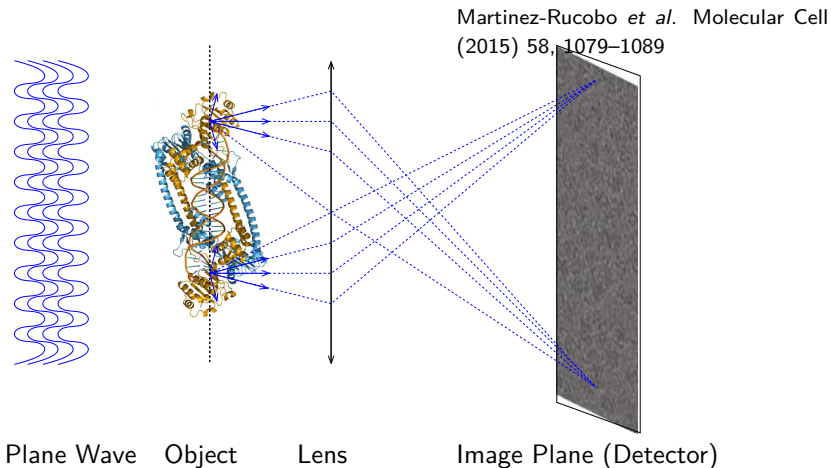
Object

Lens

Image Plane (Detector)

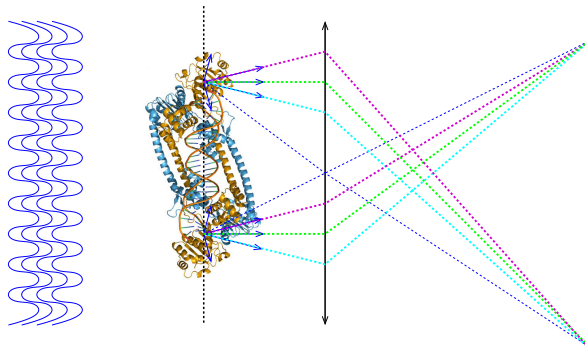
Rays of **equal origin** focus on detector

## Electron Microscope: Imaging Mode



Detector noise and radiation sensitivity  
require low contrast images

## Electron Microscope: Diffraction Mode

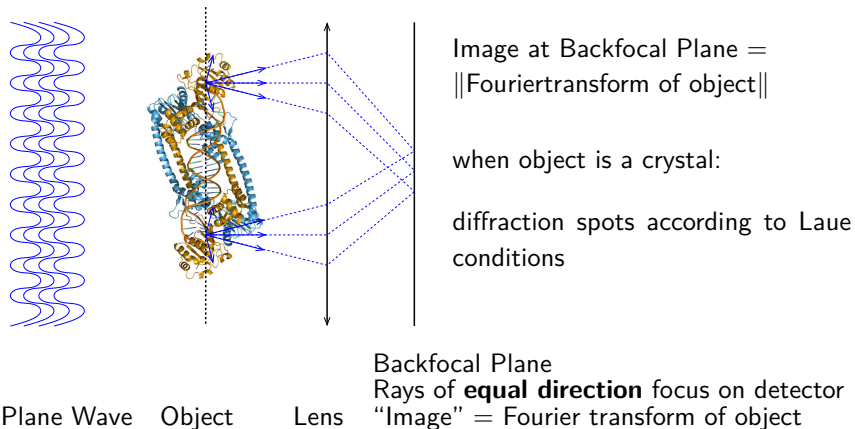


Plane Wave

Object

Lens

## Electron Microscope: Diffraction Mode



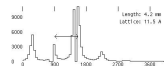
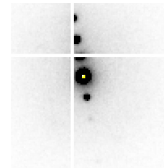
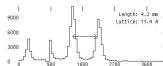
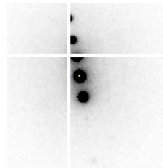
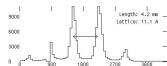
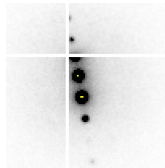
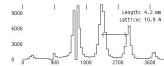
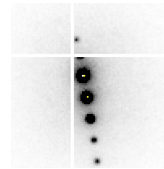
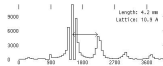
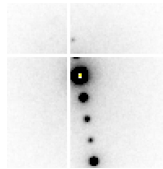
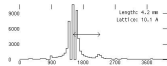
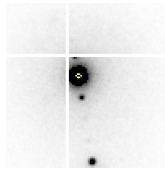
# Dynamic Scattering



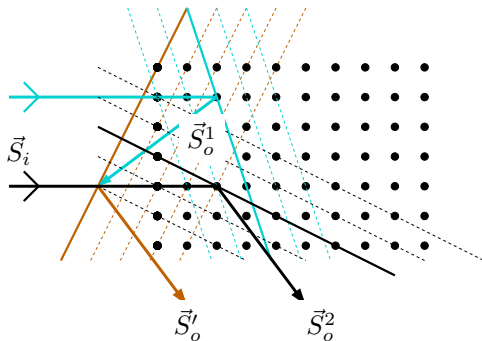
## Dynamic Scattering

- Kinematic Theory of Diffraction: Every photon / electron / neutron scatters once in the crystal
- $|F_{\text{ideal}}(hkl)| \propto \sqrt{I_{\text{exp}}(hkl)}$
- Dynamic Scattering: Multiple Scattering events occur
- Electron Diffraction: Multiple Scattering occurs even with nanocrystals

# Dynamic Scattering

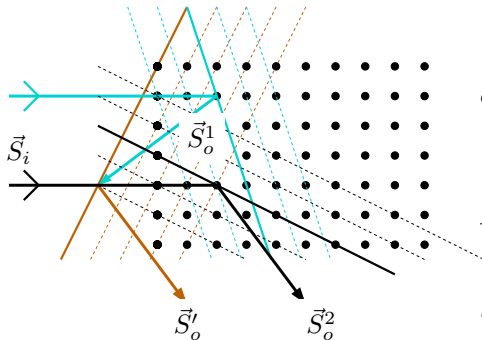


## Multiple (Dual) Scattering



- First reflection  $\vec{S}_o^1$  acts as source of second reflection  $\vec{S}_o^2$ .
- Second reflection  $\vec{S}_o^2$  overlaps with another reflection  $\vec{S}_o^2'$ .

## Multiple (Dual) Scattering



Laue Conditions (accordingly  $\vec{b}$  and  $\vec{c}$ ):

$$(\vec{S}_o^1 - \vec{S}_i) \cdot \vec{a} = h_1$$

$$(\vec{S}_o' - \vec{S}_o^1) \cdot \vec{a} = h'$$

$$\frac{(\vec{S}_o' - \vec{S}_i) \cdot \vec{a} = h_1 + h' = h_2}{}$$

$\vec{S}_o' - \vec{S}_i$  fulfills the Laue conditions, hence the secondary reflection  $\vec{S}_o' - \vec{S}_o^1$  overlaps with the "ordinary" reflection  $\vec{S}_o' - \vec{S}_i$ .

## Experimental Treatment of Dynamic Scattering for Organic Crystals

- Exact calculation very complicated
- Not feasible for complex molecules (more than a few atoms)
- Ignorance of dynamic scattering, i.e. assumption of kinematic scattering (as in X-rays) provides reliable structures
- Data statistics and model statistics poor, despite reliable structures

## Data Collection for Single Crystal Structure Analysis

- Rotation (>1966): **Contiguous** recording of reciprocal space
- hybrid pixel detectors (>2002): shutterless, *i.e.* continuous data collection
- Radiation source: X-ray, electron, or neutron crystallography
- Spot position: wave type independent
- Intensity: wave type dependent

## 3D Electron Crystallography

- > late 1990s, as opposed to 2D electron crystallography
- confusingly many terms (ADT , RED, EDT, PEDT, MicroED, ...)
- Historical “dispute”, Ute Kolb, Mainz University,  $\approx$  2007 (ADT), Xiaodong Zou & Sven Hovmöller, Stockholm University  $\approx$  2011 (RED)
- technical term: “3D Electron Diffraction”, Enrico Mugnaioli (PSI 2017; IUCrJ (2019), 6, 178–188)
- “3D”:
  1. Collection of 3D reciprocal space
  2. 3D crystals:  $\geq$  10–15 unit cell in each direction; typically 200–1000nm per dimension

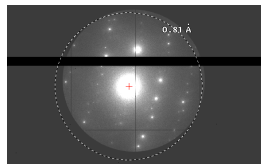
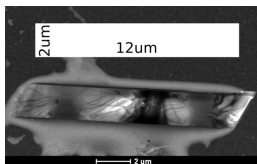
# Example Structures



# Single Crystal Structure from a Pharmacy Powder

Gruene *et al.*, *Angew. Chemie. Int. Ed.* (2018), 57, 16313–16317

## Grippostad<sup>®</sup>, STADA



### active compounds

paracetamol  
 ascorbic acid  
 caffeine  
 chlorphenamine maleate

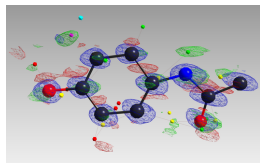
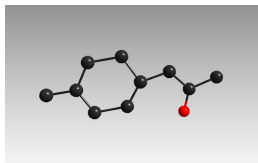
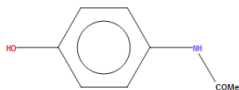
### non-active compounds

gelatine  
 glycerol tristearate  
 lactose monohydrate  
 quinoline yellow (E104)  
 erythrosine (E127)  
 titanium dioxide (E171)

- powder from capsule deposited on sample grid
- Crystal dimensions  $2\mu\text{m} \times 12\mu\text{m} \times \approx 300\text{nm}$
- $d_{\text{min}} < 0.8\text{\AA}$

## Single Crystal Structure from a Pharmacy Powder

1. Data from single crystal: Completeness < 40%
2. Cell parameters:  $a = 6.9$ ,  $b = 9.4$ ,  $c = 11.6$ ,  $\alpha = 90.6$ ,  $\beta = 98.4$ ,  $\gamma = 89.8$   
CSD search  $a = 7.1$ ,  $b = 9.3$ ,  $c = 11.7$ ,  $\alpha = 90.0$ ,  $\beta = 97.7$ ,  $\gamma = 90.0$   
CCDC HXACAN04,  $P2_1/n$ , Paracetamol,
3. SHELXT solves structure
4. Difference map reveals hydrogen atoms: data sensitivity

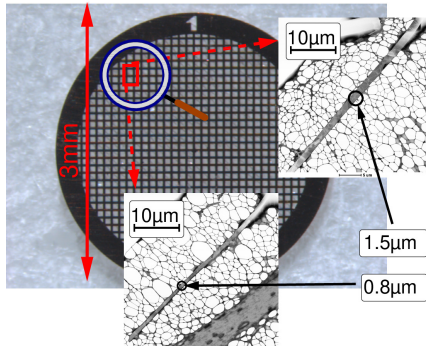


**Future: Complete crystallographic analysis from powder blends**

# Drug Design: Structure of a New Methylene Blue Derivative MBBF<sub>4</sub>

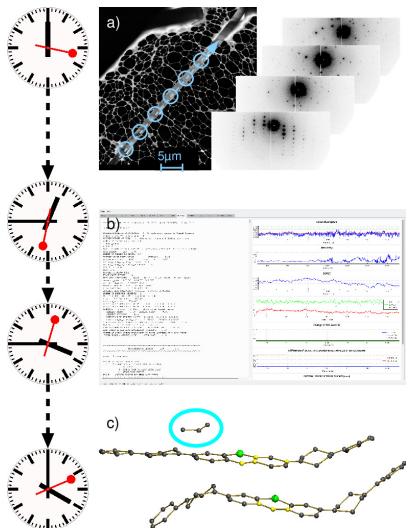
Collaboration Dr. J. Holstein & Prof. G. Clever, TU Dortmund  
Gruene *et al.*, *Angew. Chemie. Int. Ed.* (2018), 57, 16313–16317

## MBBF<sub>4</sub>-nanoCrystal (Holstein/Clever, TU Dortmund)



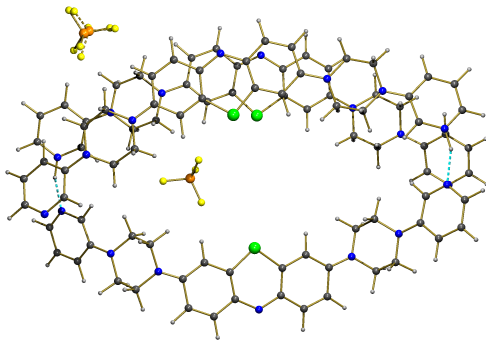
Tip of thin MBBF<sub>4</sub> needle on a TEM sample grid

# MBBF<sub>4</sub> — EIGER and a TEM make a Synchrotron



- 60 – 120° @ 3°/s = 40 s / data set
- 45 min for 16 data sets on both grids
- manual processing  $\approx$  4h to structure solution

## Structure of MBBF<sub>4</sub> (Refinement J. Holstein, TU Dortmund)



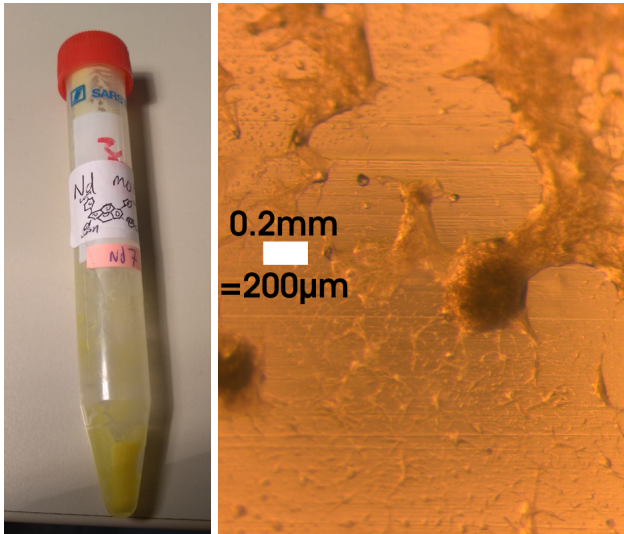
- $R1 = 22.7\%(2941F_o > 4\sigma_F)$
- $R1 = 27.2\%(4832F_o)$
- $Goof = 1.5$

# Nd-MOF

Prof. Jia Min Chin & Prof. Michael Reithofer, University of Vienna

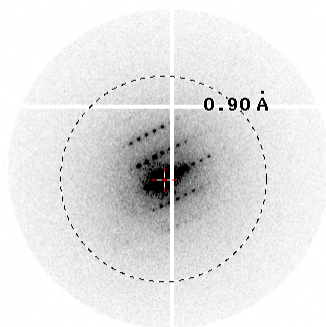
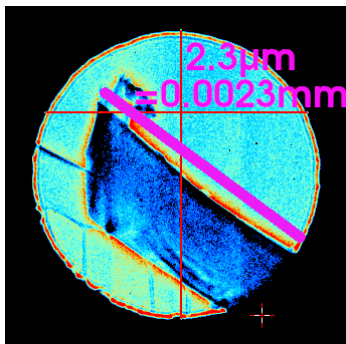


## Jewels in the mud



photographs courtesy Dipl.-WIng. A. Roller

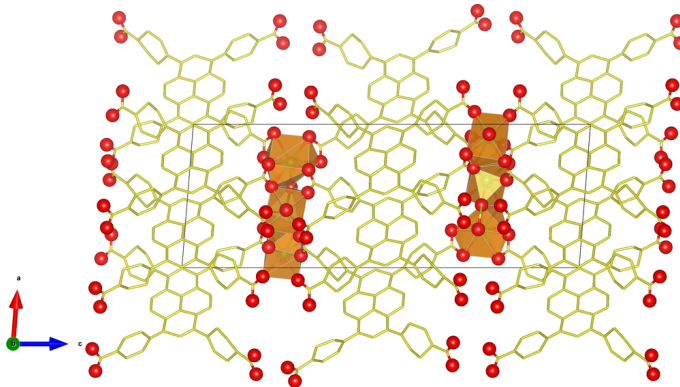
## Powerful electron diffraction



Sample preparation: A. Roller & N. Gajic

At DESY, the strongest X-ray source in the world, this crystal would probably not show any diffraction.

## Nd-MOF structure from 5 crystals

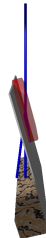
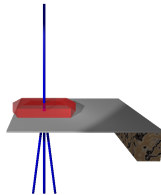
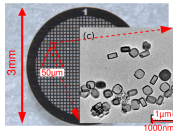


Courtesy Jia Min Chin & Michael Reithofer, unpublished data  
Room temperature measurement, under vacuum

# Preferred Crystal orientation & the Missing Wedge Problem

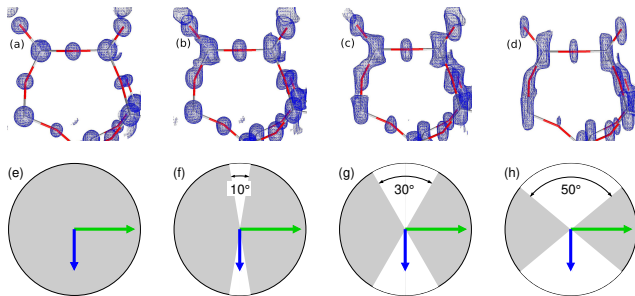
Wennmacher *et al.*, Nat. Comm. (2019), 10, 3316; (Patent EP 18 202 868)

## Missing Wedge in Electron Diffraction



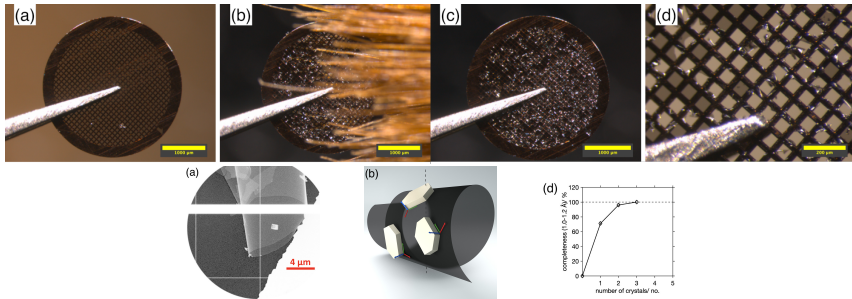
- Crystals very often have a **flat shape**: always the same orientation
- Sample support stabilised by Cu-grid
- Copper grid too thick: intransparent for electrons
- Limited rotation range
- Crystals typically have low symmetry space group

## Effect of Systematically Missing Data



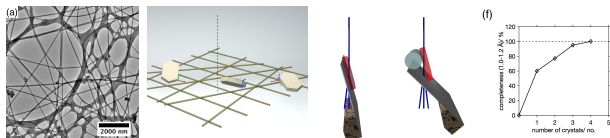
- as little as  $10^\circ$  degree of missing data lead to shearing of the experimental map
- Shearing of experimental map results in unreliable coordinates for structure

## Solution 1 — Coiled carbon grids



- Brush Stroke causes carbon layer to coil
- Visual selection of orientation from carbon curvature
- Complete data from 5-ish crystals

## Solution 2 — Nylon Fibres



- Nylon fibres ( $\approx 100nm$  diameter) disturb preferred orientation
- Orientation less obvious from visual inspection — possibly more screening required
- Complete data from 5-ish crystals
- Nylon grids adaptable to sample size and shape



## Acknowledgements

- My colleagues at the X-ray Centre: A. Roller & N. Gajic
- My collaborators from the nanoArgovia project A3EDPI (*cf.* references)
- H. Stahlberg (C-CINA), N. Blanc (ScopeM), C. Schulze-Briese (DECTRIS Ltd.)
- B. Luethi, L. Wagner, L. Piazza, D. Mayani (all DECTRIS)
- B. Schmitt (PSI), E. Frojdh (PSI), G. Tinti (PSI), S. Vetter (all PSI detector group)
- R. Mezzenga (ETHZ), U. Shimanovich (Weizmann Inst.), I. Adrianssens Martiel (PSI), I. Schlichting (MPI Heidelberg), K. Diederichs (Uni Konstanz), J. Lübben (Bruker AXS), M. Clabbers (Uni Stockholm)
- J. Chin, M. Reithofer (both Uni Vienna)



DECTRIS QUADRO: to be in-  
stalled at Vienna University

**DECTRIS**  
detecting the future

nanoArgovia A3EDPI

SNF Project 169258