

# **XRPD:**

## **Challenging organic disordered structures**

**Prof. Dr. Martin U. Schmidt**  
**Goethe-Universität Frankfurt**  
**Inst. f. Anorgan. und Analyt. Chemie**  
**[m.schmidt@chemie.uni-frankfurt.de](mailto:m.schmidt@chemie.uni-frankfurt.de)**

*Mainz, 7-12-2022*

# Content

## 1. Iso-butyl-lithium

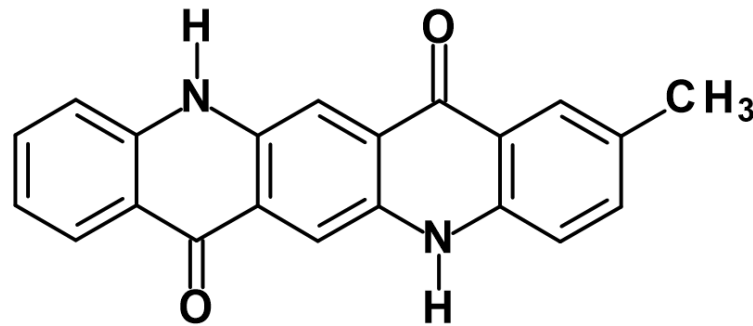
Ordered phase and rotator phase

## 2. NaOEt

Disordered structure with unclear space group

## 3. Monomethyl-quinacridone

Head-to-tail disorder



# <sup>i</sup>BuLi iso-Butyl-lithium

**Crystal structure determined by  
X-ray powder diffraction**

Alexander Bodach, Lothar Fink, Martin U. Schmidt\*:

*Crystal structures of ordered and plastic-crystalline phases of iso-butyllithium by X-ray powder diffraction,*

*Chem. Commun.*, **2018**, 54, 10734-10737.

# Preparation and measurement

**<sup>i</sup>BuLi: highly sensitive to air and moisture  
All procedures under N<sub>2</sub> or Ar**

**<sup>i</sup>BuLi obtained as solution in hexane**

**2x recrystallised**

**Evaporated**

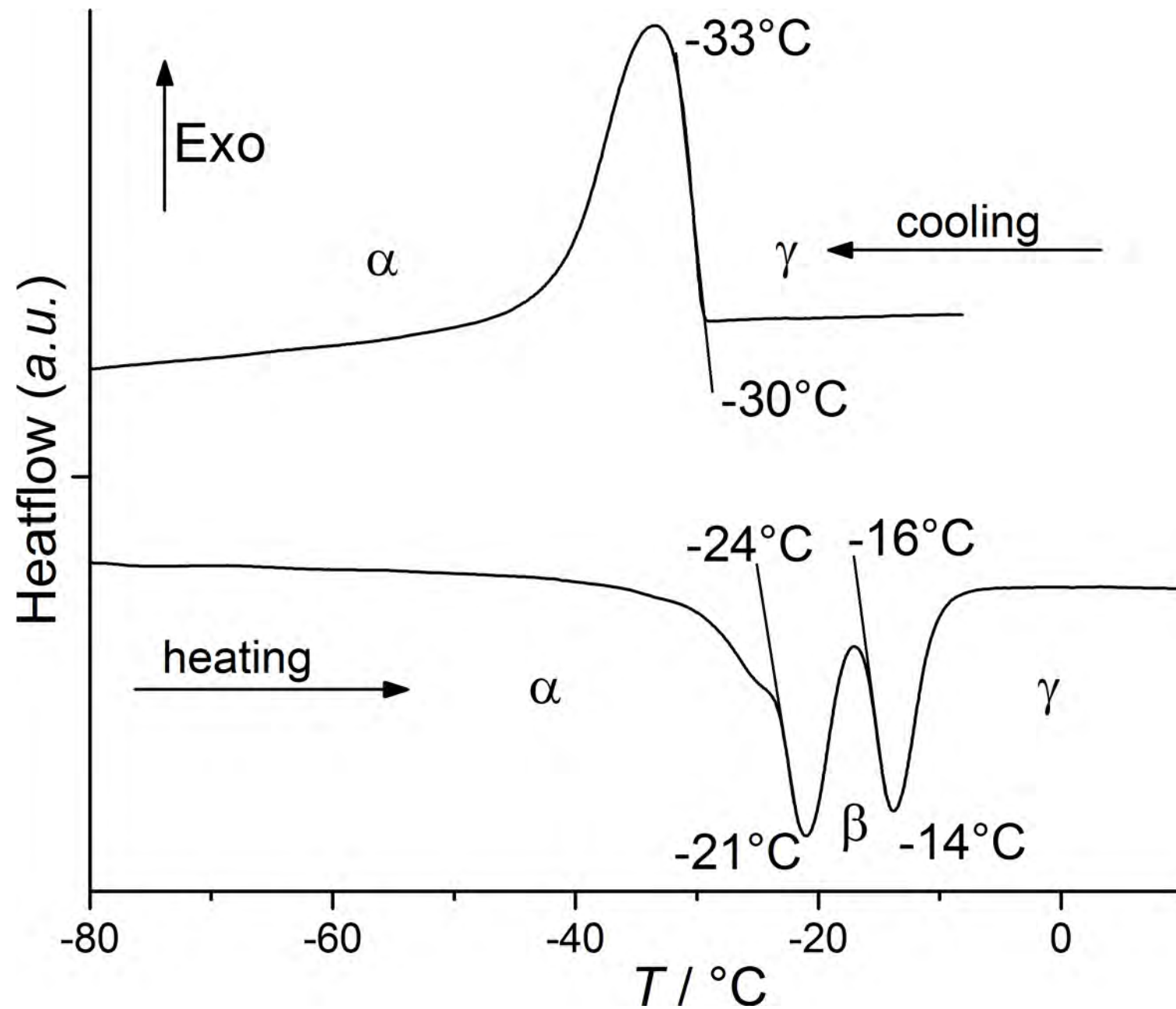
**Filled in capillary and sealed**

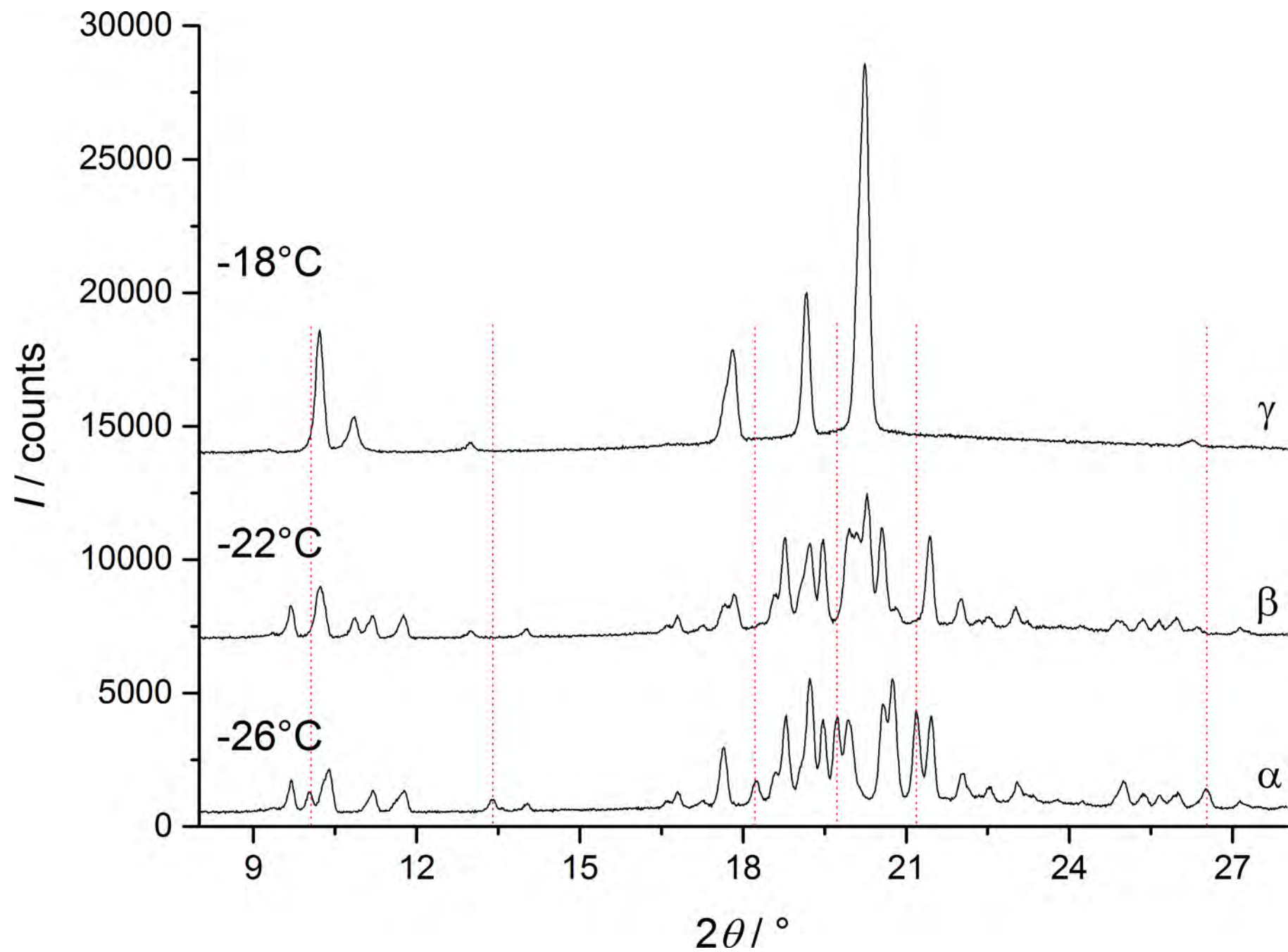
**Measured in sealed capillary**

**STOE Stadi-P diffractometer, Cu-K $\alpha_1$  radiation**

**Temperature control by Oxford Cryostream**

# DSC





# Structure determination of the low-temperature phase ( $\alpha$ )

Indexing: triclinic

Space group  $P\ 1$  or  $P\ \bar{1}$ ?

From unit cell volume:  $Z = 12$   
 $\Rightarrow$  3 tetramers or 2 hexamers?

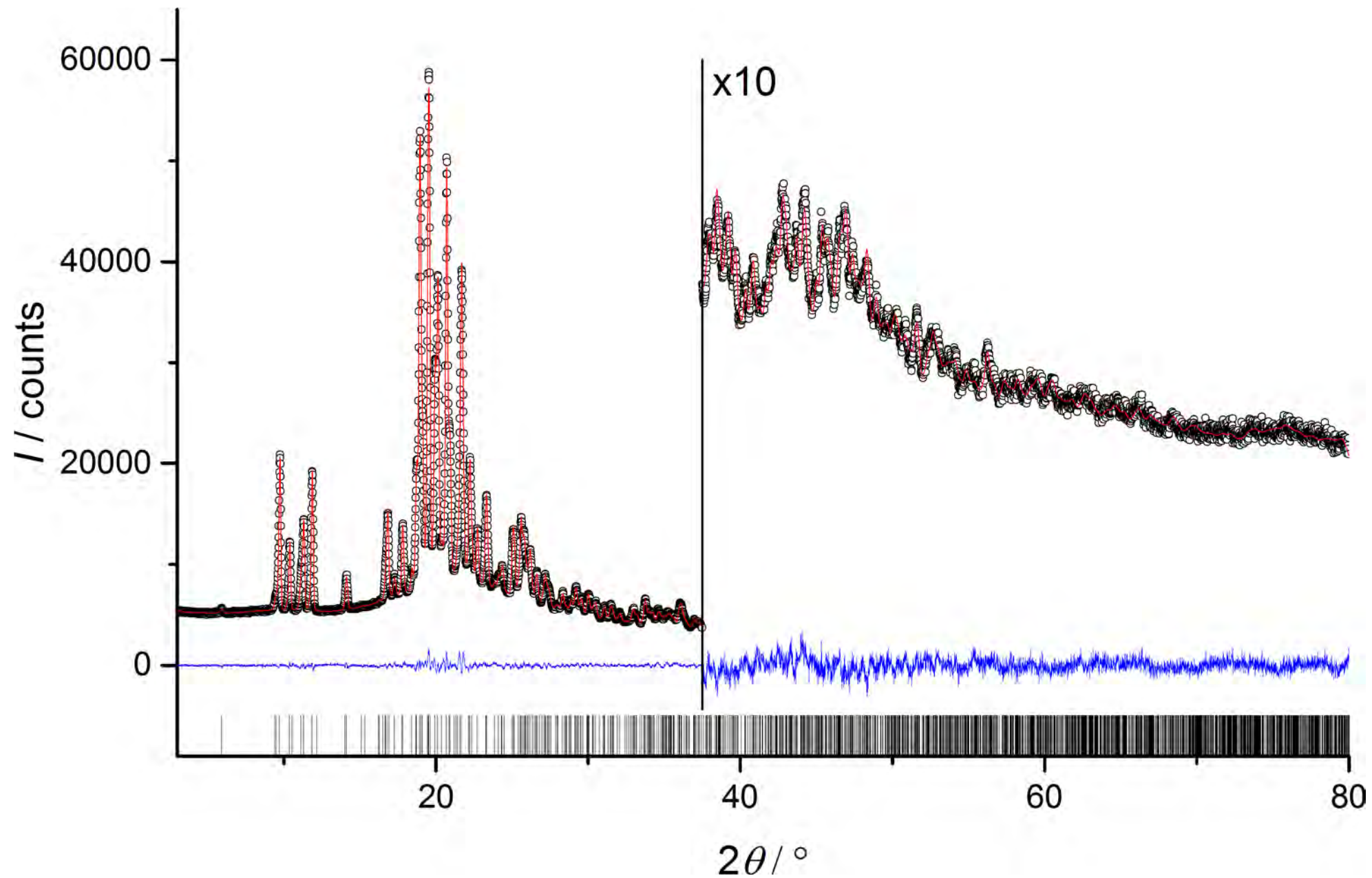
Structure solution by direct-space methods with DASH

- with tetramers in  $P\ 1$  and  $P\ \bar{1}$
- with hexamers in  $P\ 1$  and  $P\ \bar{1}$

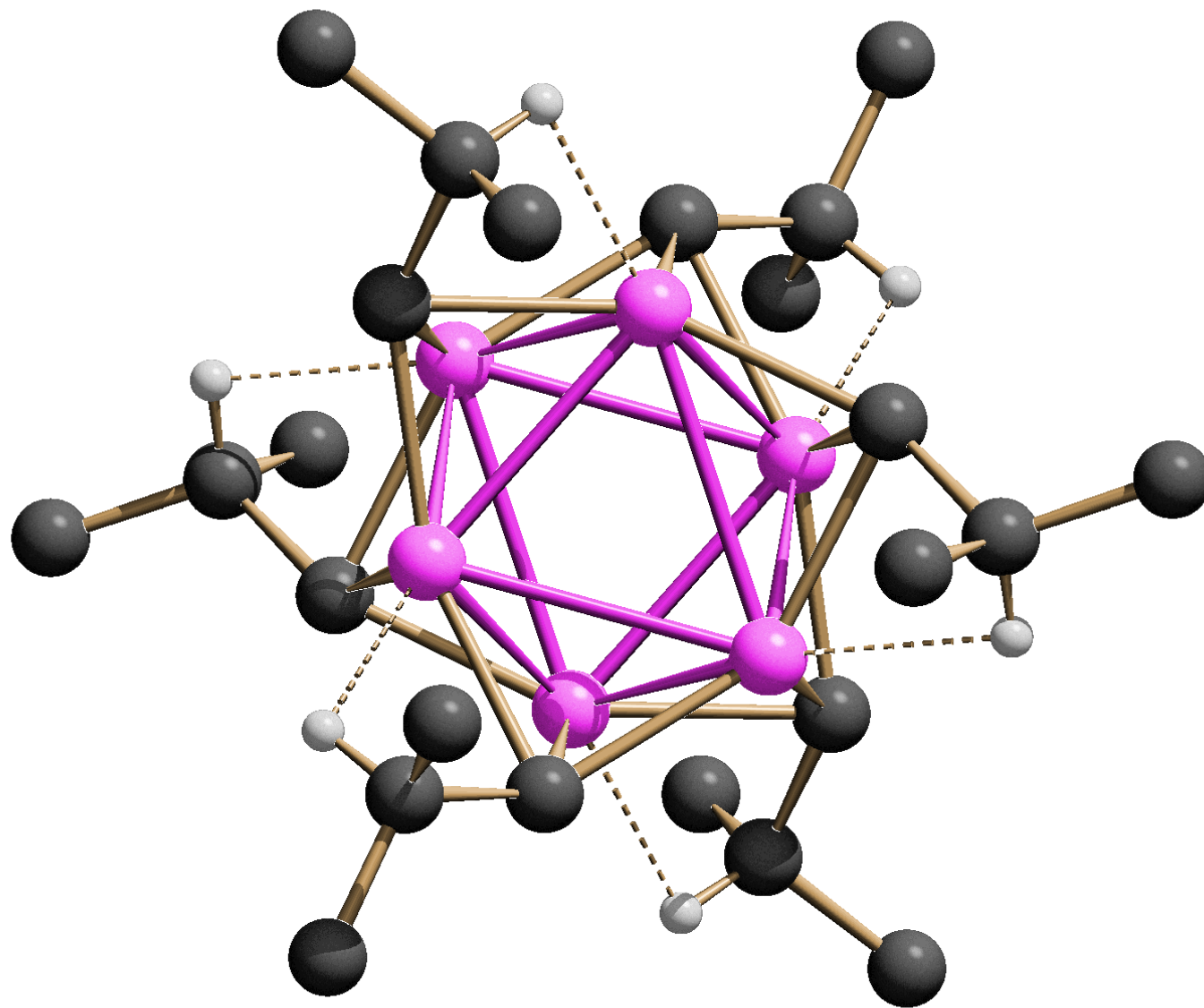
Final solution:

Space group  $P\ \bar{1}$  with two symmetrically independent hexamers, both on inversion centres

# Low-temperature phase ( $\alpha$ -phase)

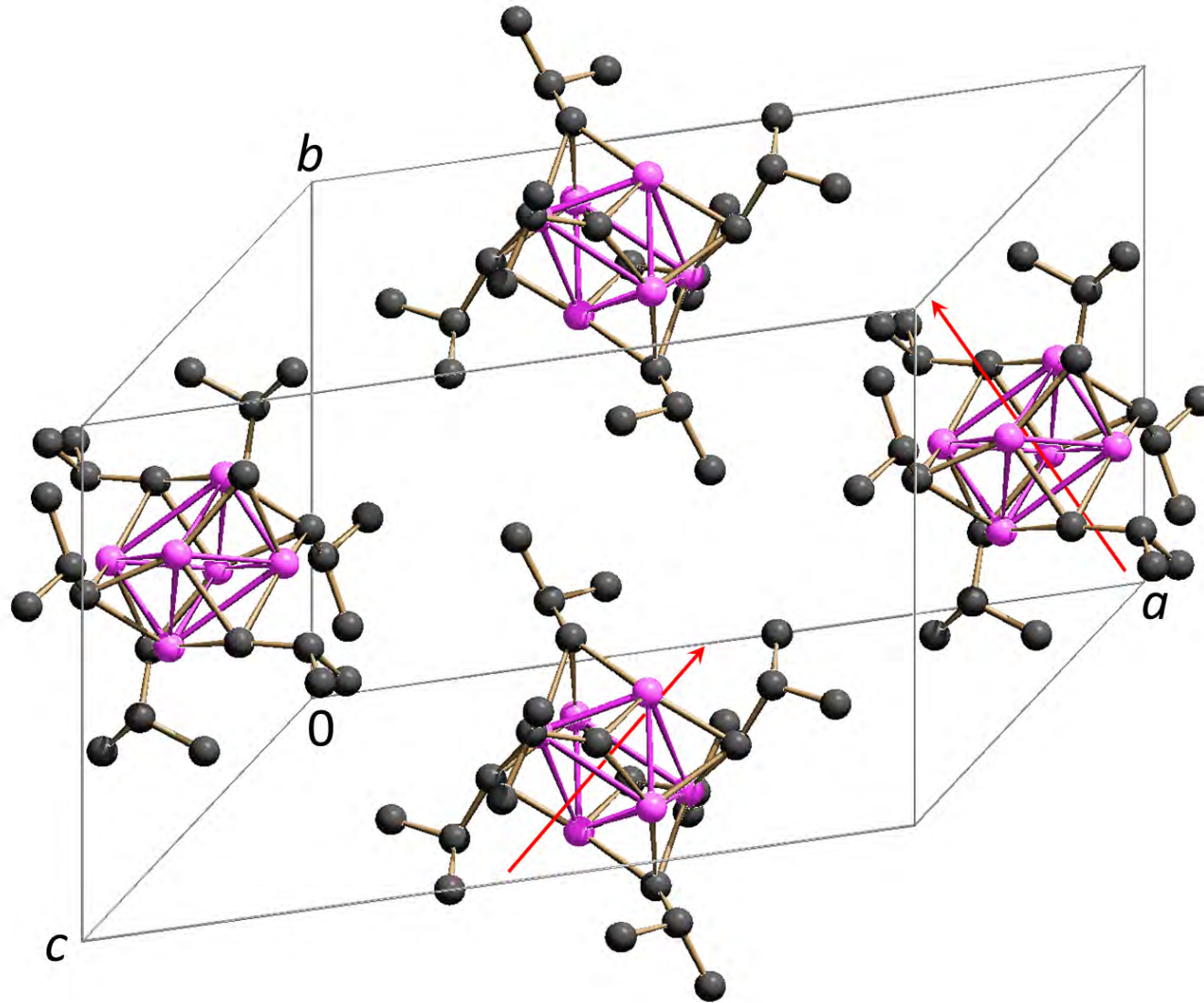


# Low-temperature phase ( $\alpha$ -phase)



**Hexamer**

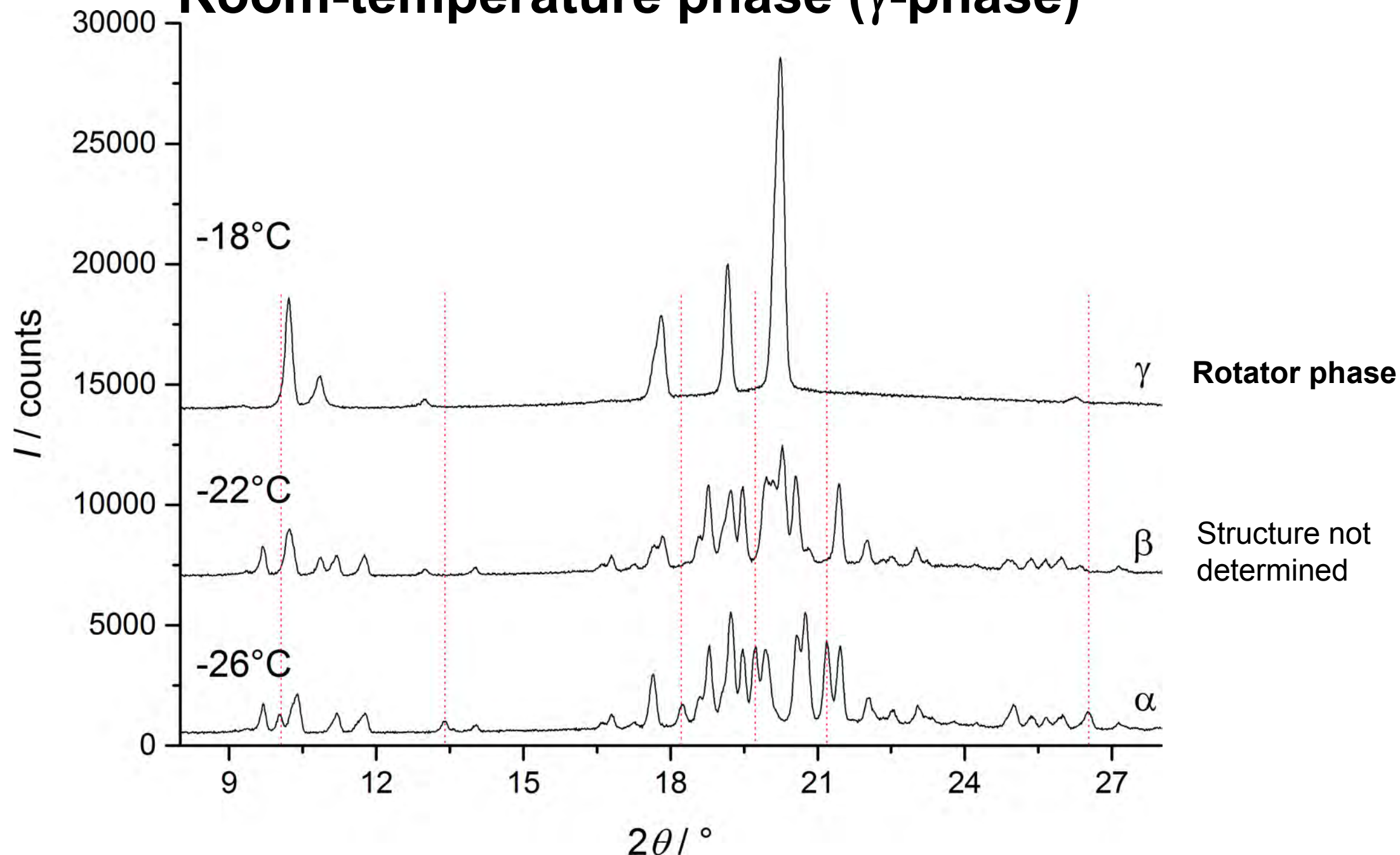
# Low-temperature phase ( $\alpha$ -phase)



$P\bar{1}$ ,  $Z = 2$

Two symmetrically independent hexamers, both on inversion centres

# Room-temperature phase ( $\gamma$ -phase)



# Room-temperature phase ( $\gamma$ -phase)

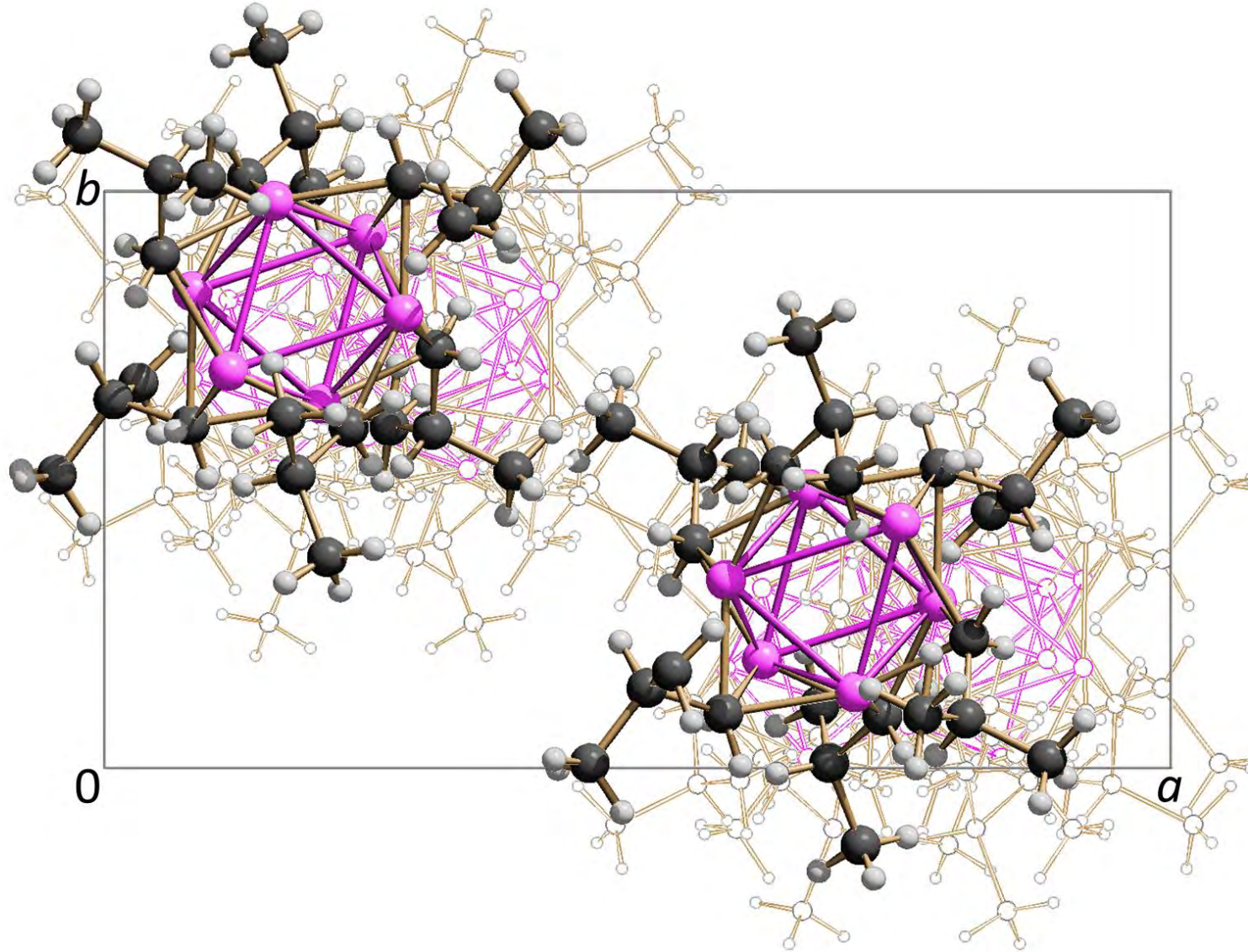
Indexing: orthorhombic,  $Z' = 2$

Space group unclear

Rotator phase (Plastic phase)!

Structure finally solved in Pnnn  
Molecules 4-fold disordered

# Room-temperature phase ( $\gamma$ -phase)



**Rotator phase (Plastic phase)**

**Hexamers rotate in the solid state (Dynamic disorder)**

# **NaOEt**

**First Synthesis: Liebig (1837)**

**180 years later:**

**Crystal structure determined by  
X-ray powder diffraction (2019)**



$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

Vacuum  $\rightarrow$  Solid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

Vacuum  $\rightarrow$  Solid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  Liquid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  Liquid

Vacuum  $\rightarrow$  Solid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt in EtOH (Solution)}$

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Ar  $\rightarrow$  Liquid

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  Solid

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt}$  in EtOH (Solution)

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  Liquid

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  Solid

Mixture of 3-4 phases. Mainly  $\text{NaOEt} \cdot 2 \text{EtOH}$

$\text{Na} + \text{EtOH} \rightarrow \text{NaOEt}$  in EtOH (Solution)

Vacuum  $\rightarrow$  Solid

Ar  $\rightarrow$  liquid

Vacuum  $\rightarrow$  Solid

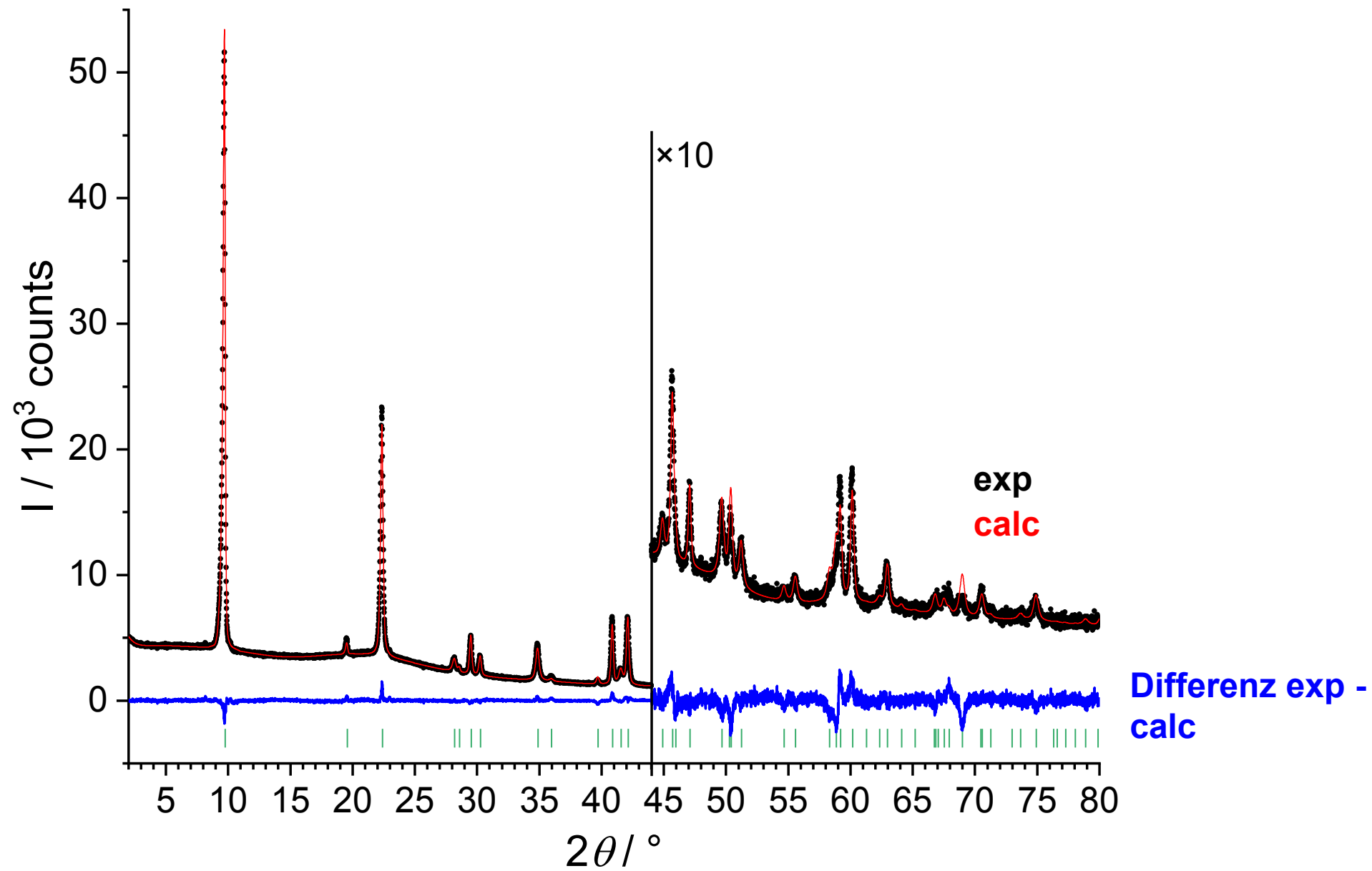
Ar  $\rightarrow$  Liquid

Vacuum  $\rightarrow$  Solid

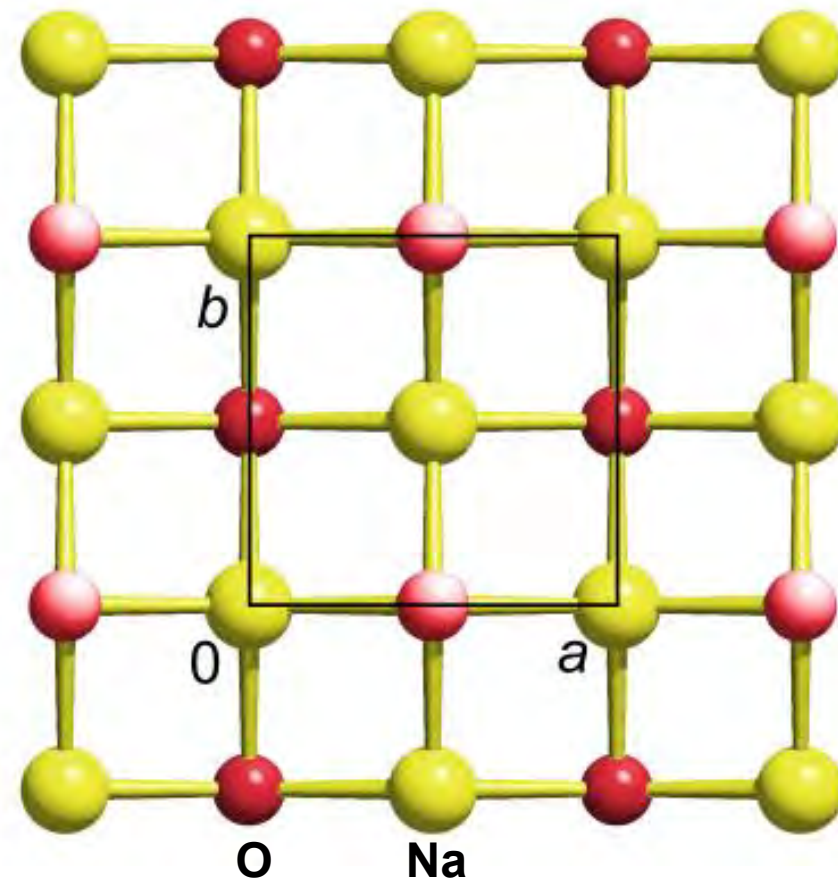
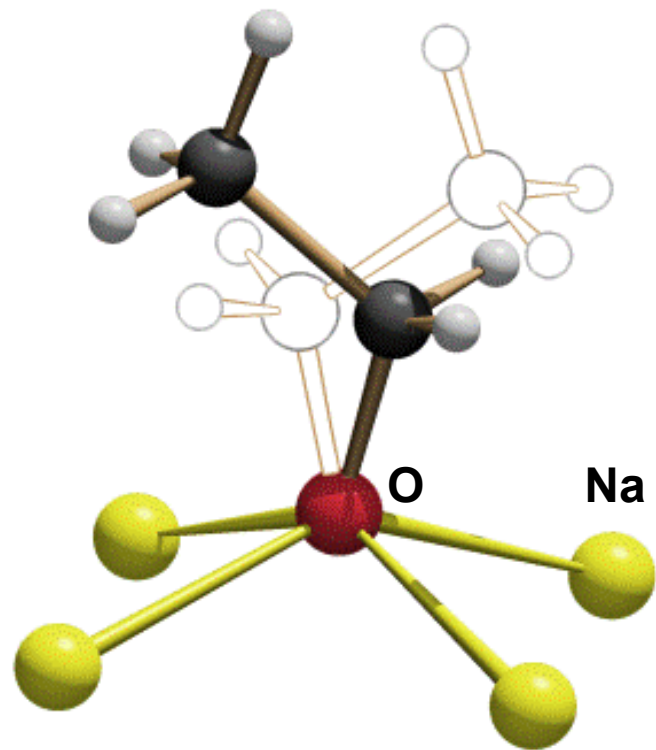
Ar  $\rightarrow$  Solid

Mixture of 3-4 phases. Mainly  $\text{NaOEt} \cdot 2 \text{EtOH}$

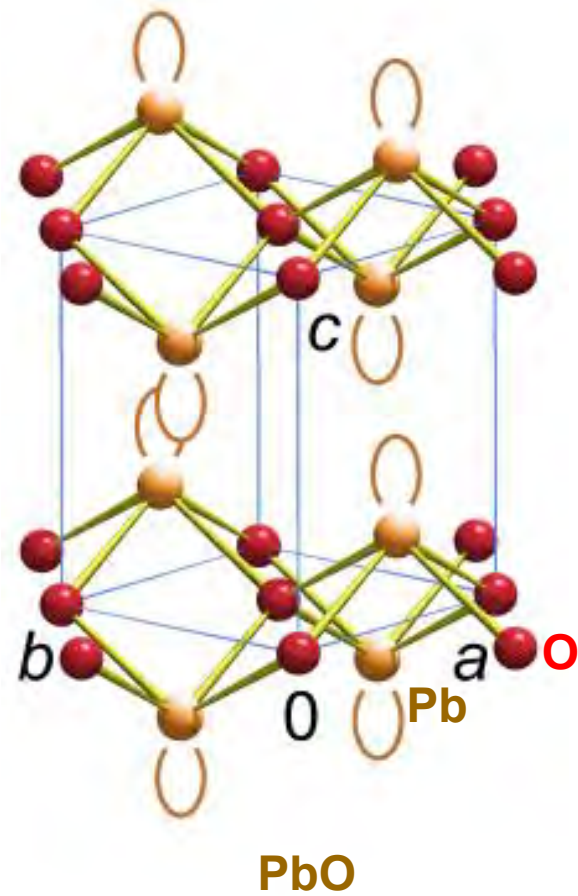
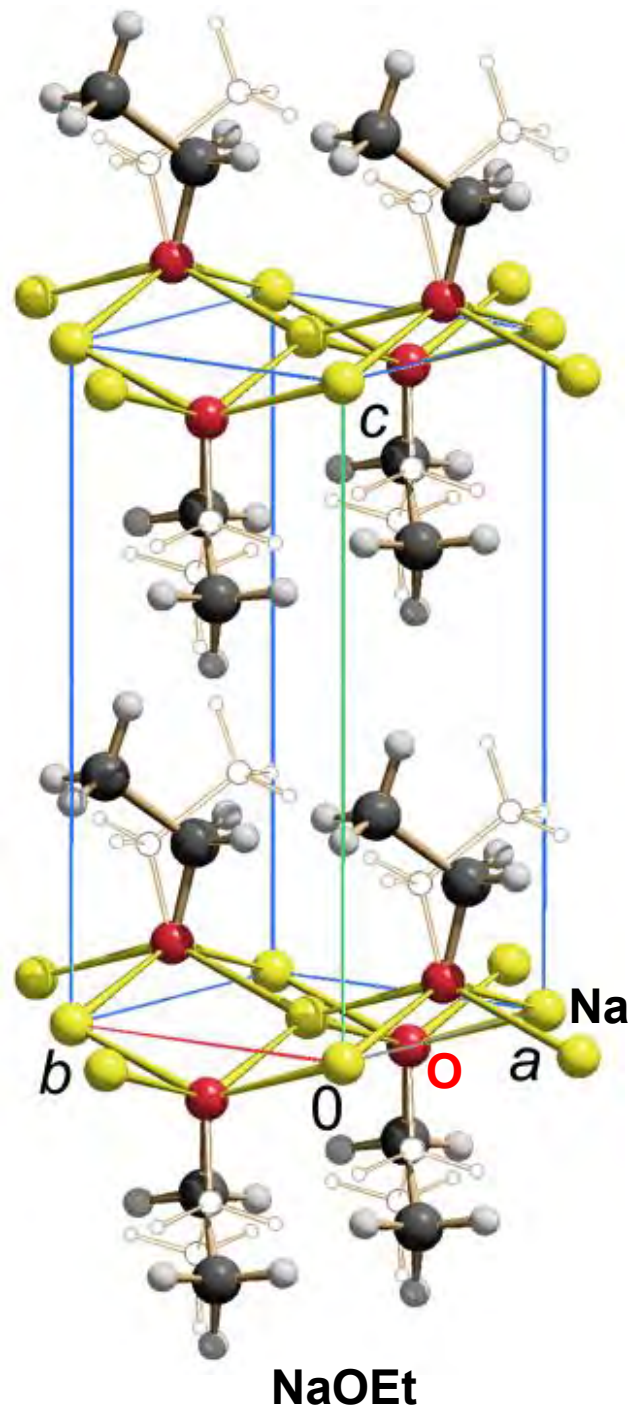
Vacuum,  $50^\circ\text{C}$ , several hours  $\rightarrow$  NaOEt (pure)



**NaOEt: Rietveld refinement**

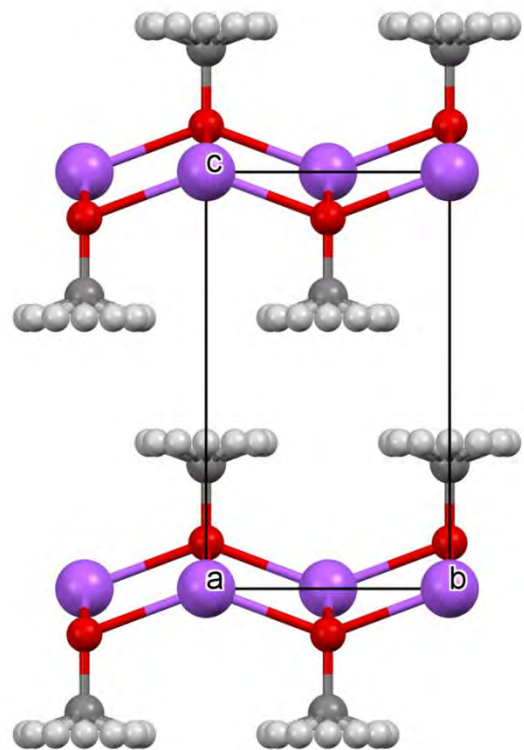


NaOEt

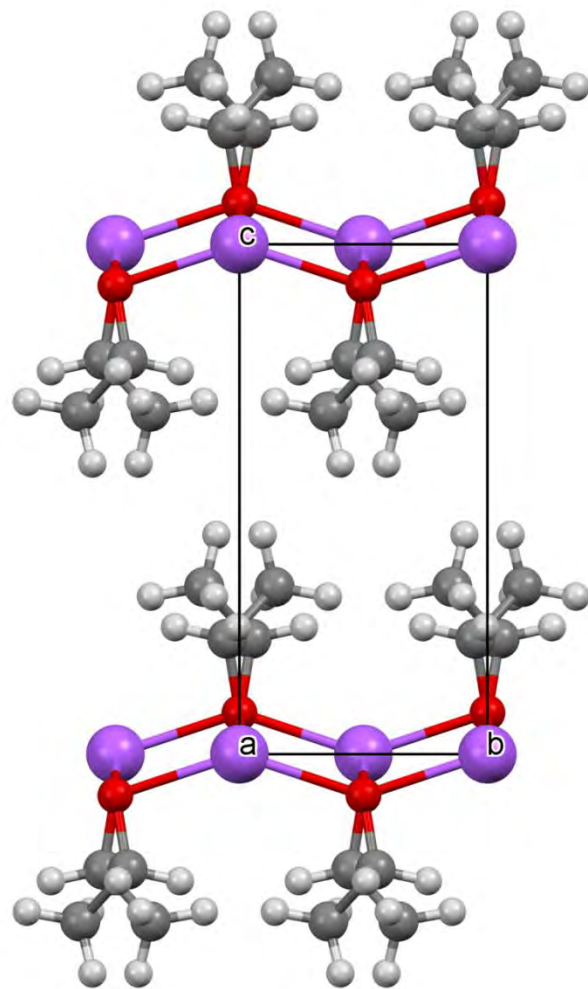


|    |    |    |
|----|----|----|
| Na | O  | Et |
| O  | Pb | LP |

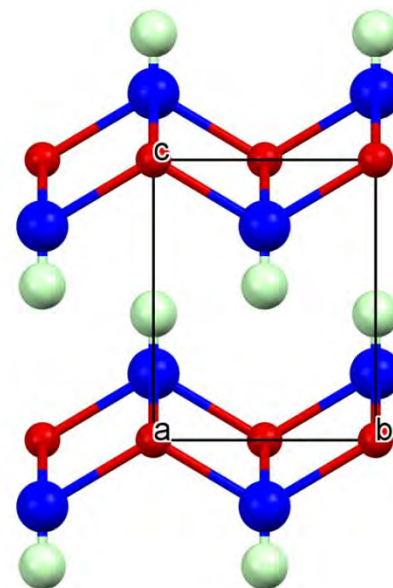
**NaOEt has  
Anti-PbO structure.**



(a)  
NaOMe  
 $P4/nmm$

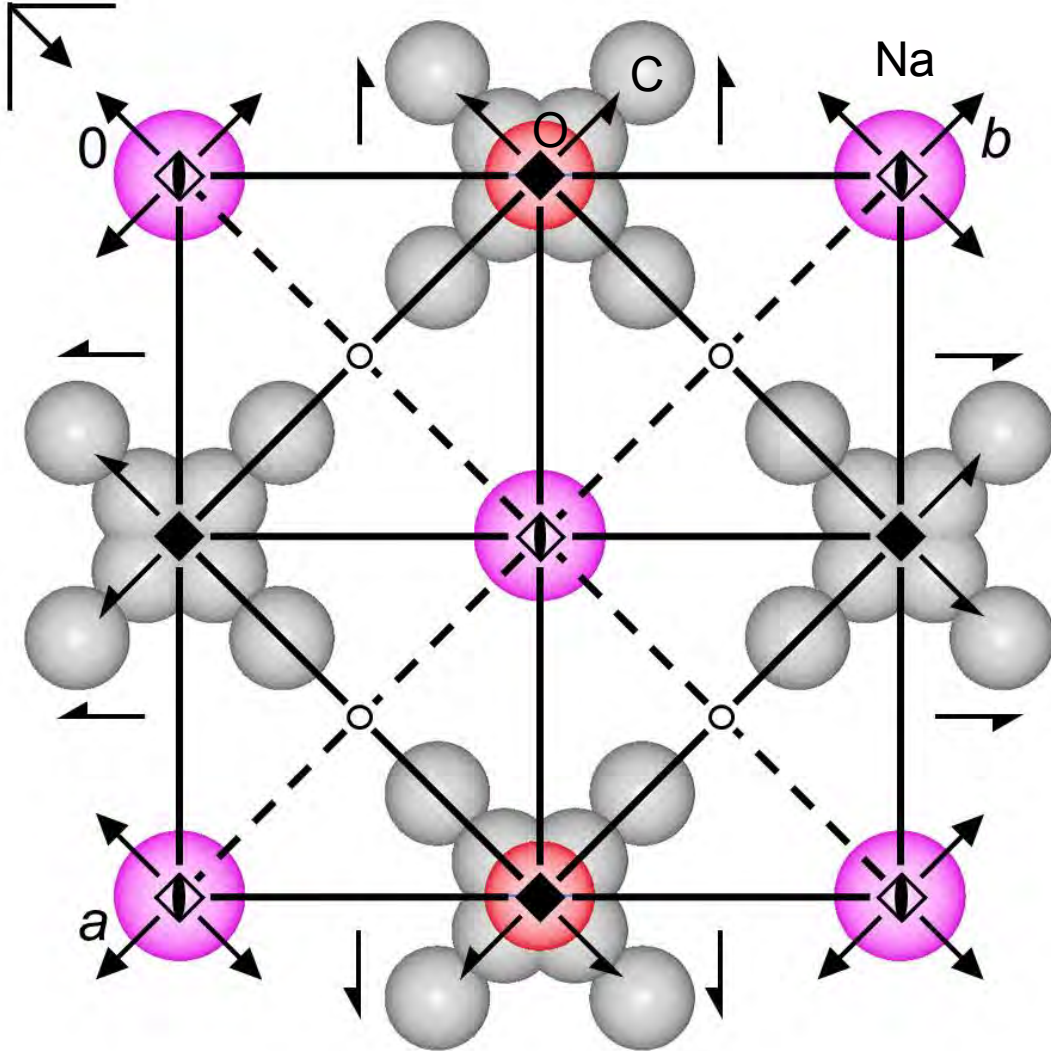


(b)  
NaOEt  
 $P\bar{4}2_1m$

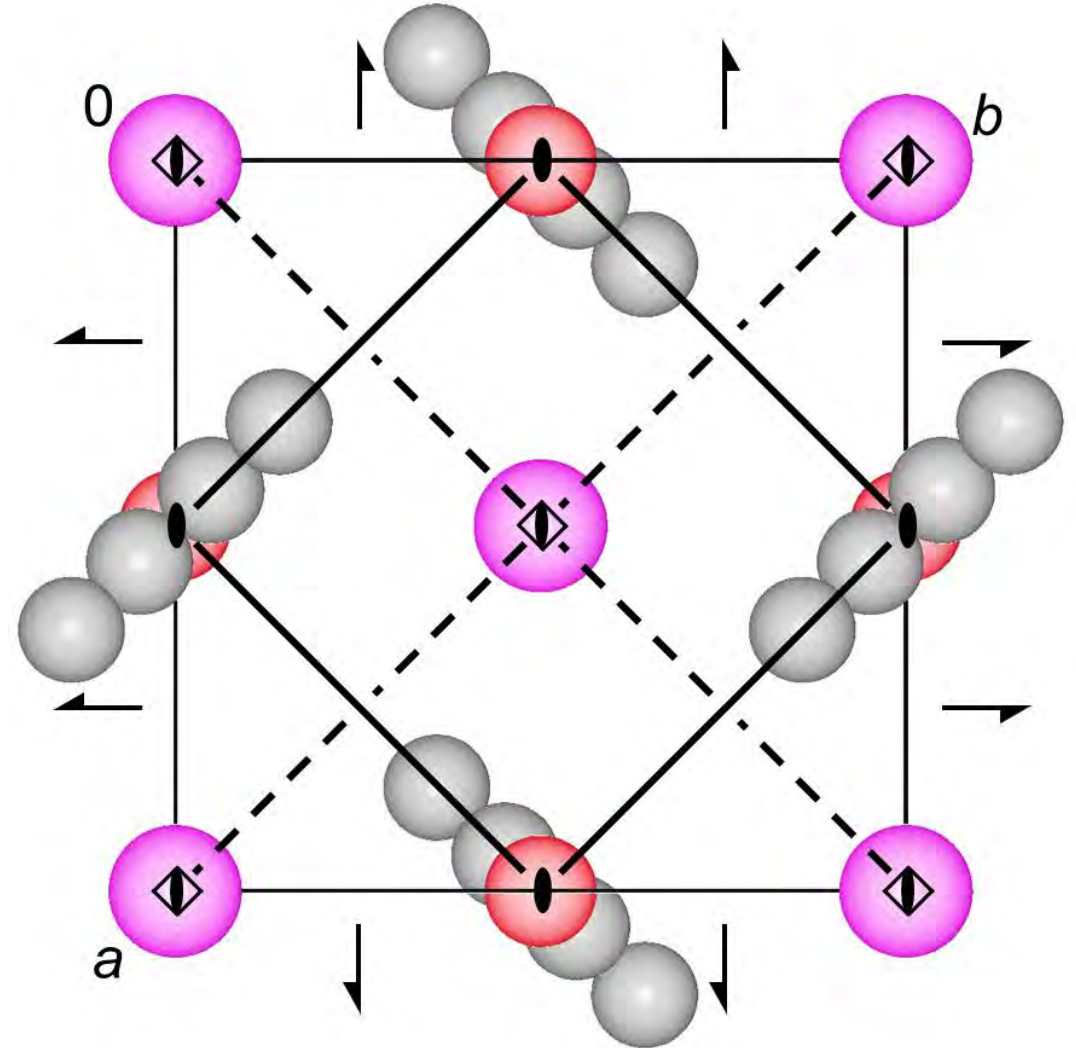


(c)  
PbO  
 $P4/nmm$

# NaOEt: Space group and disorder



$P 4/nmm$ ,  $Z = 2$   
4-fold disorder



$P \bar{4} 2_1 m$ ,  $Z = 2$   
2-fold disorder

**$P\bar{4}$ ,  $Z = 4$**

| Na        | Na        | Na | O  | C  |
|-----------|-----------|----|----|----|
| 1a        | 1c        | 2g | 4h | 4h |
| $\bar{4}$ | $\bar{4}$ | 2  | 1  | 1  |

C<sub>2</sub>H<sub>5</sub> ordered

ordered

**$P2_1 1 1$ ,  $Z = 2$**

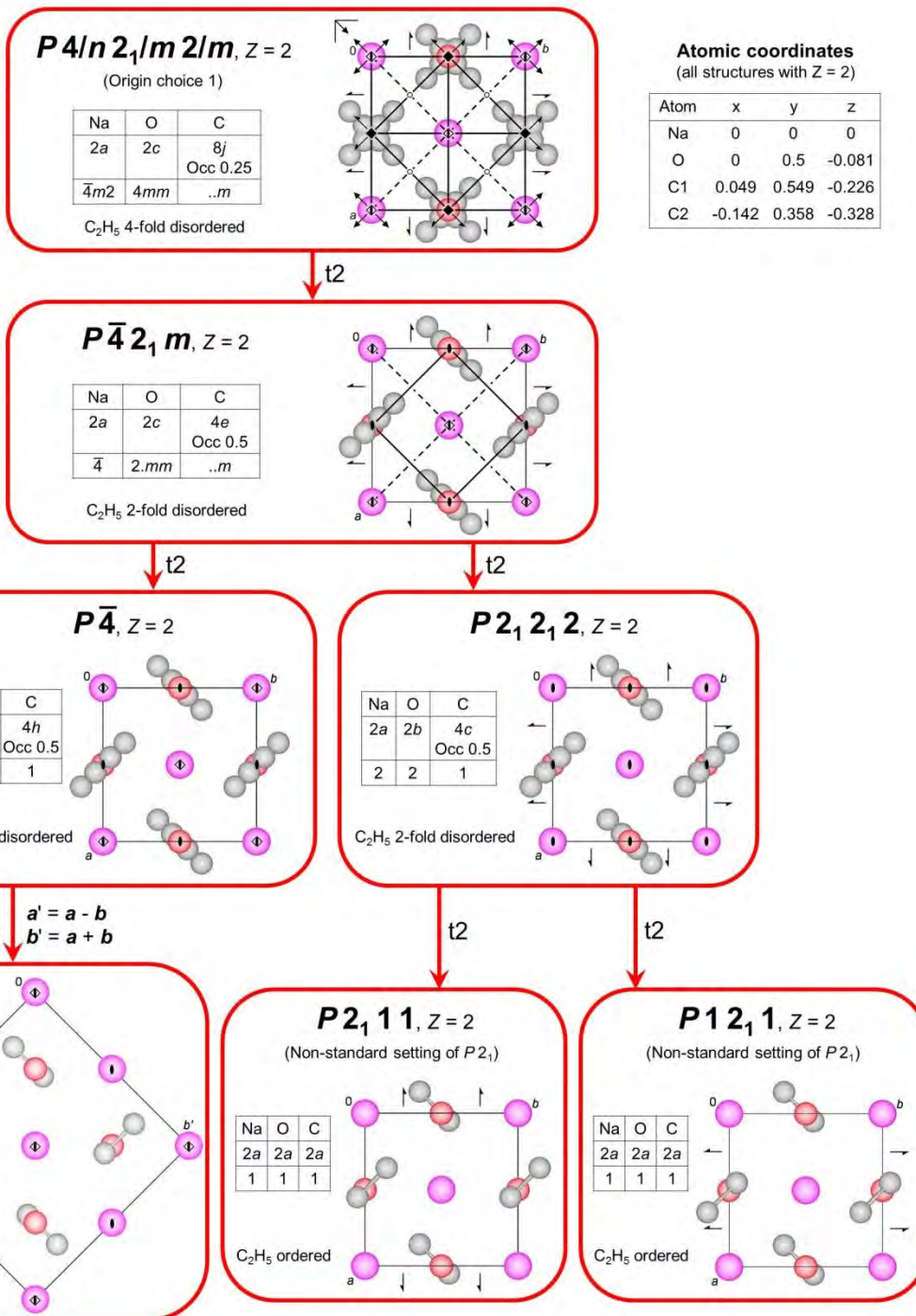
(Non-standard setting of  $P2_1$ )

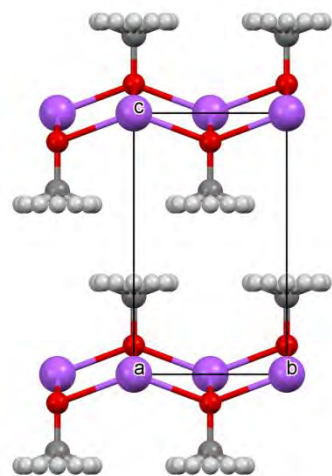
| Na | O  | C  |
|----|----|----|
| 2a | 2a | 2a |
| 1  | 1  | 1  |

C<sub>2</sub>H<sub>5</sub> ordered

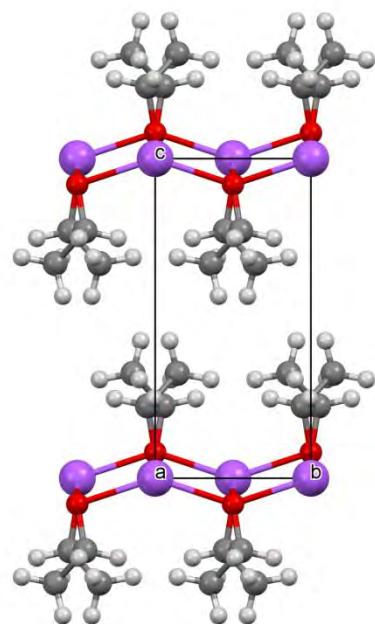
ordered

# Bärnighausen tree

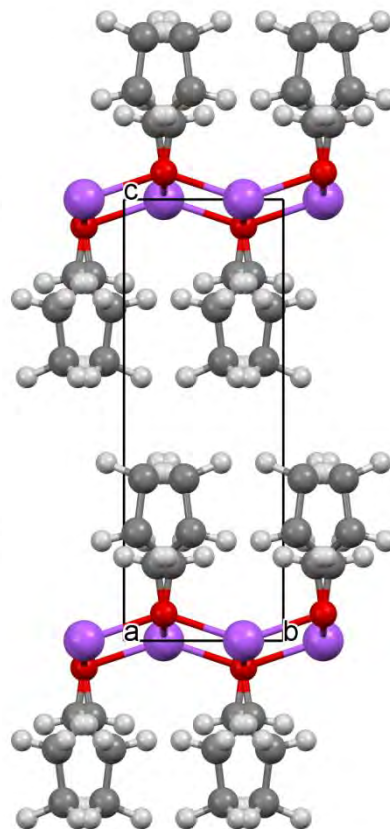




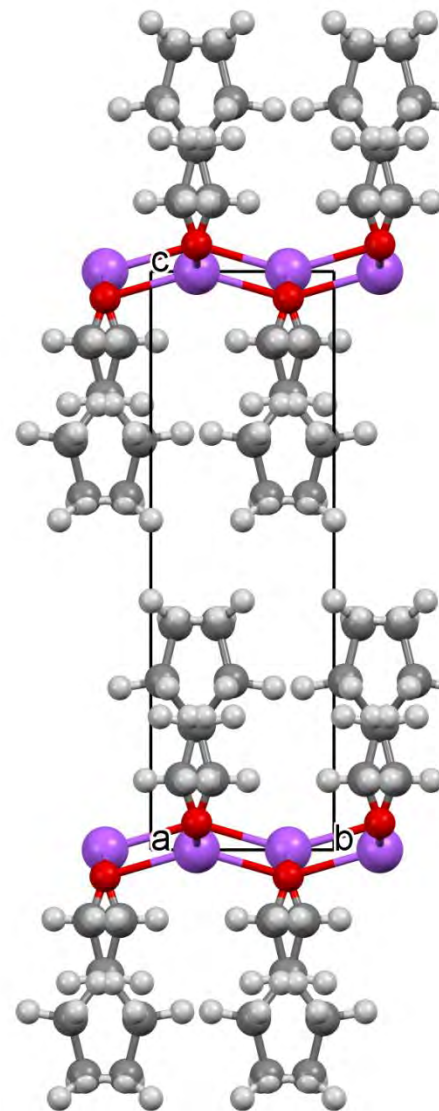
**NaOMe**



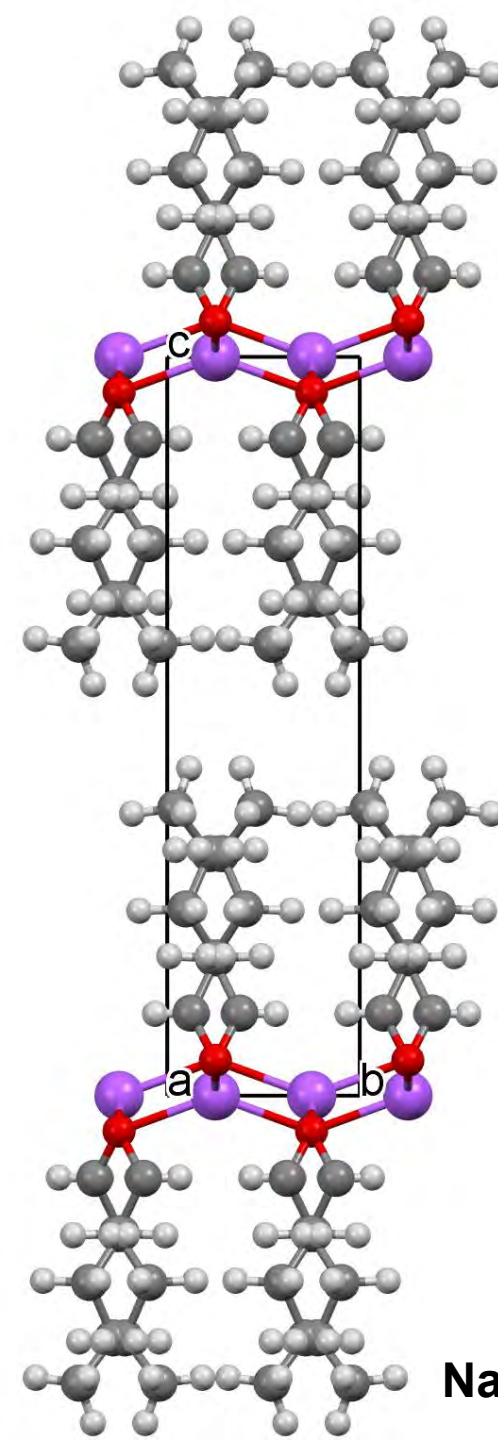
**NaOEt**



**NaOPr**



**NaOBu**



**NaOAm**

Published as:

1) Maurice Beske, Lukas Tapmeyer, Martin U. Schmidt:

"Crystal structure of sodium ethoxide ( $\text{C}_2\text{H}_5\text{ONa}$ ), unravelled after a 180 years"

*Chem. Commun.*, **2020**, 56, 3520-3523

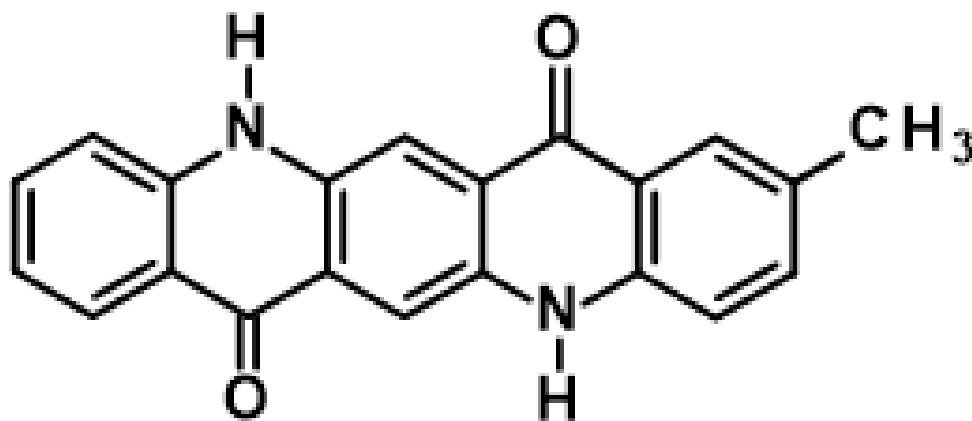
2) Maurice Beske, Stephanie Cronje, Martin U. Schmidt\*, Lukas Tapmeyer:

"Disordered sodium alkoxides from powder data: crystal structures of sodium ethoxide, propoxide, butoxide and pentoxide, and some of their solvates"

*Acta Cryst.*, **2021**, B77, 1-15.

**Orientational disorder  
in monomethyl-quinacridone  
investigated by Rietveld refinement,  
pair-distribution function analysis  
and lattice-energy minimisations**

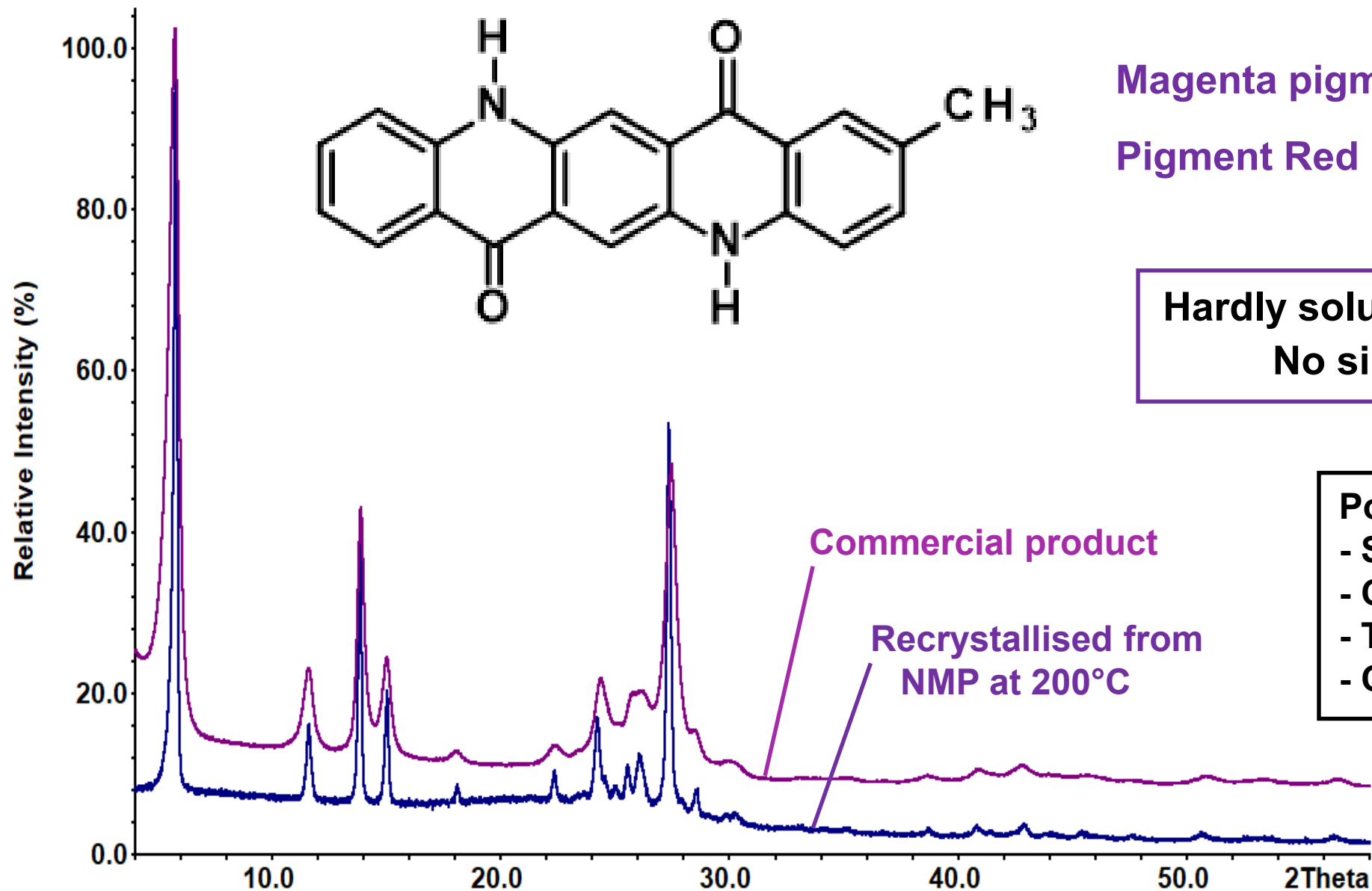
# Monomethyl-quinacridone



Magenta pigment for paints

Pigment Red 192

Hardly soluble in all solvents.  
No single crystals



Powder diffraction:

- STOE STADI-P
- Capillary
- Transmission geometry
- Cu-K $\alpha_1$  radiation

# Structure determination from powder data

- Indexing
- Structure solution by real-space methods with DASH
- "Rietveld" refinement with TOPAS  
Restraints on bond lengths, bond angles and planarity

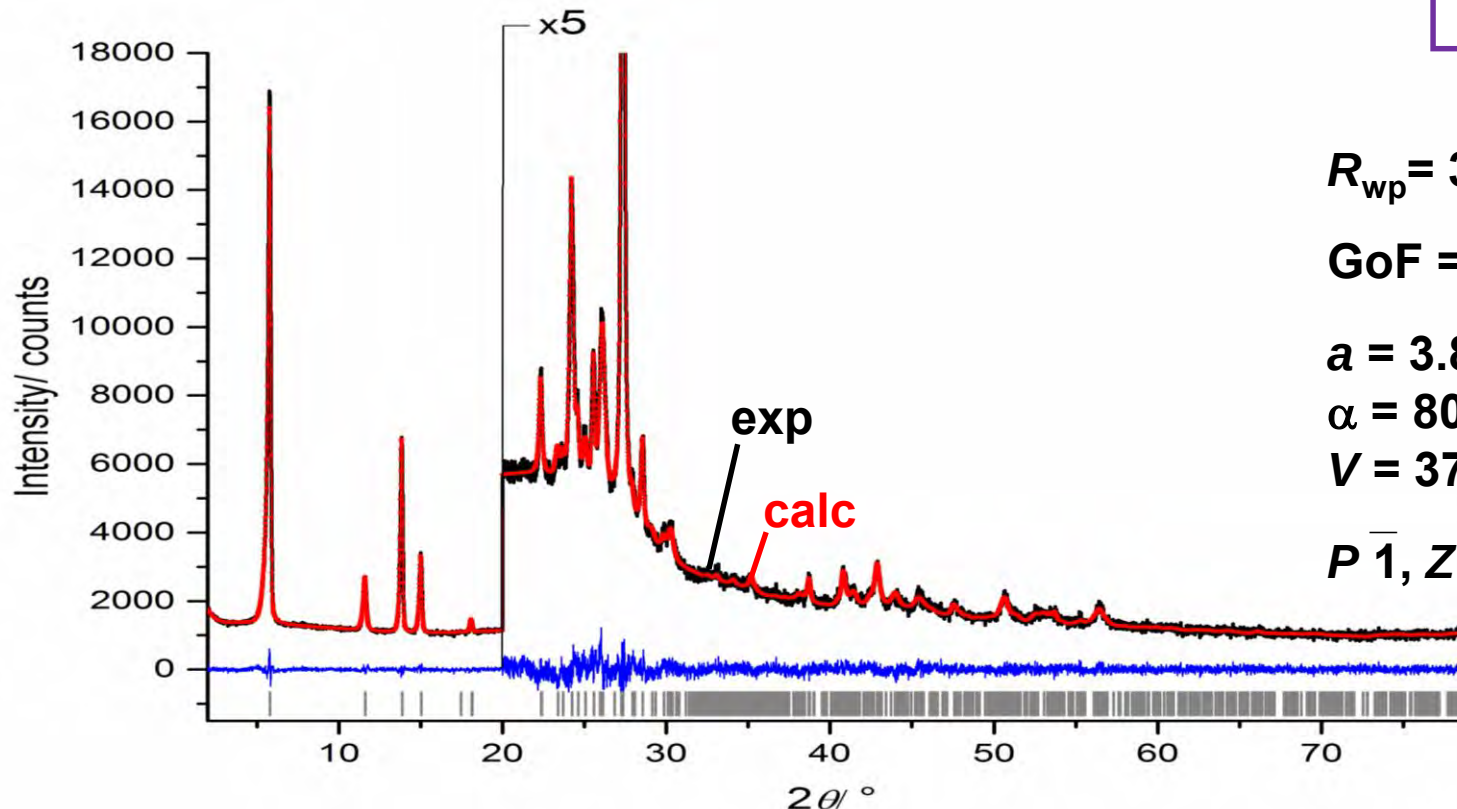
*Comment on the name "Rietveld method":*

*- Idea of Loopstra, 1963*

*- Programmed by Rietveld, 1966*

*⇒ Correct name: "Loopstra method"*

*[Loopstra & Rietveld, Acta Cryst B25, 787 (1969);  
van Laar & Schenk, Acta Cryst A74, 88 (2018)]*



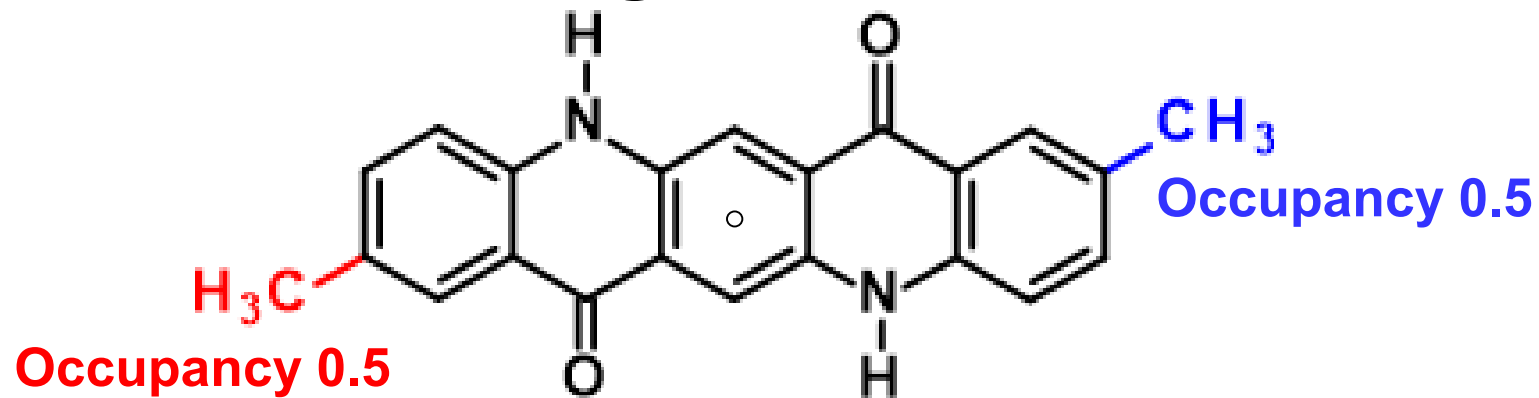
$$R_{wp} = 3.888$$

$$GoF = 1.065$$

$$a = 3.82627 (2), b = 6.87845 (4), c = 16.800 (19) \text{ \AA},$$
$$\alpha = 80.77 (10), \beta = 64.38 (8), \gamma = 68.14 (5)^\circ,$$
$$V = 371.44 \text{ \AA}^3$$

$$P \bar{1}, Z = 1, Z' = 0.5$$

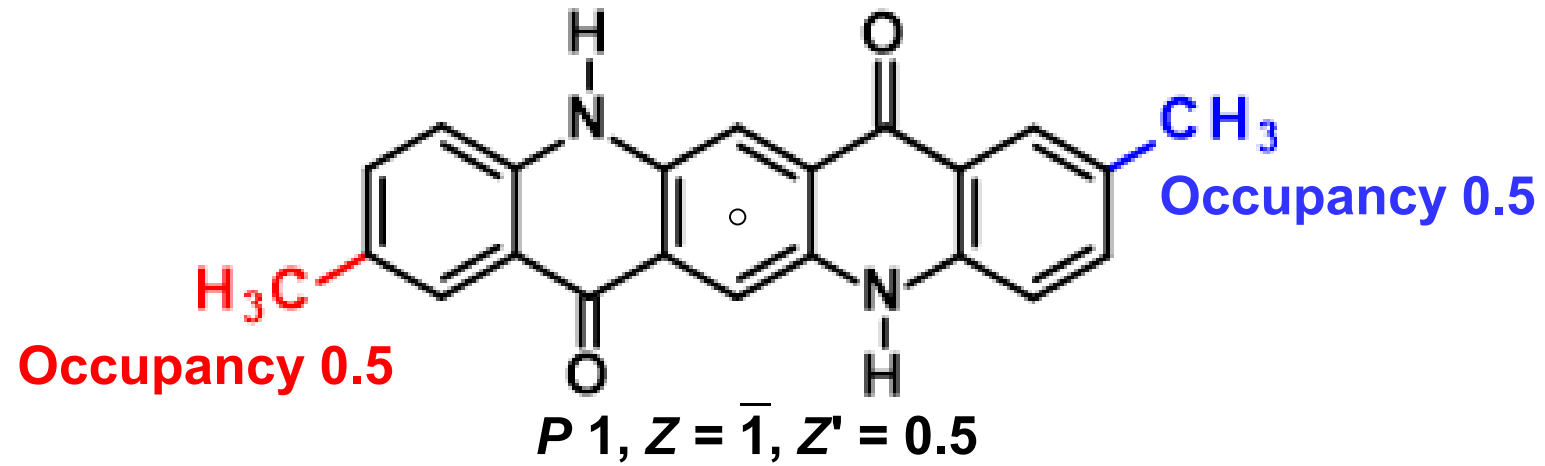
# Average structure



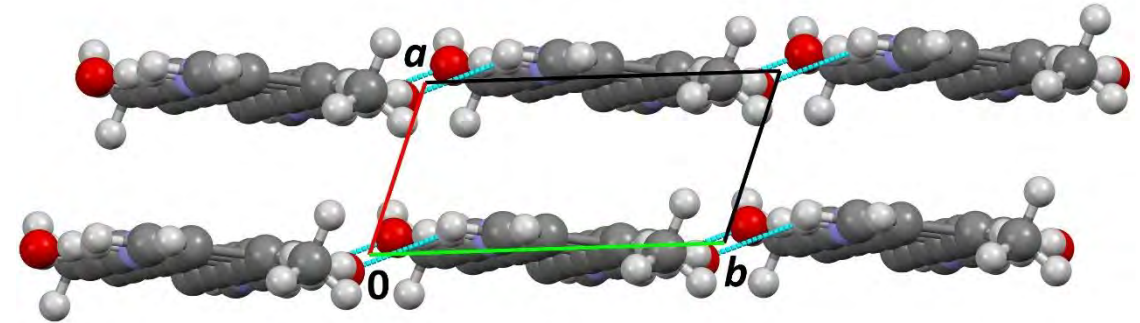
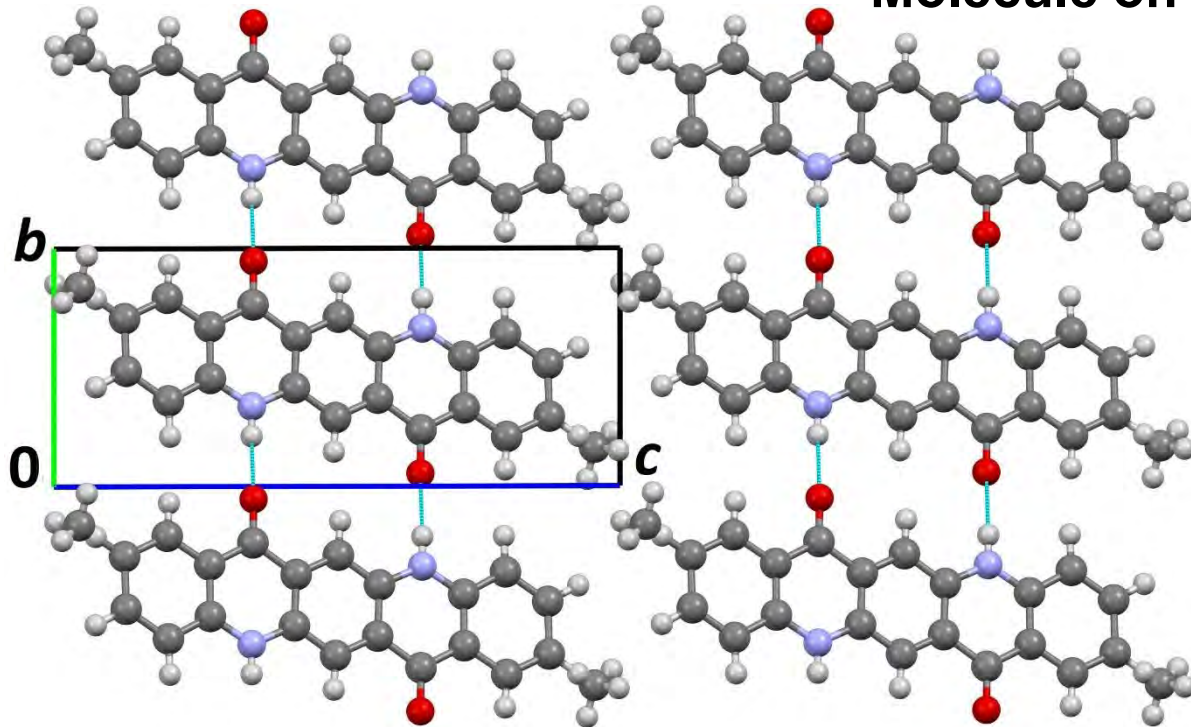
$P\ 1, Z = \bar{1}, Z' = 0.5$

Molecule on inversion centre

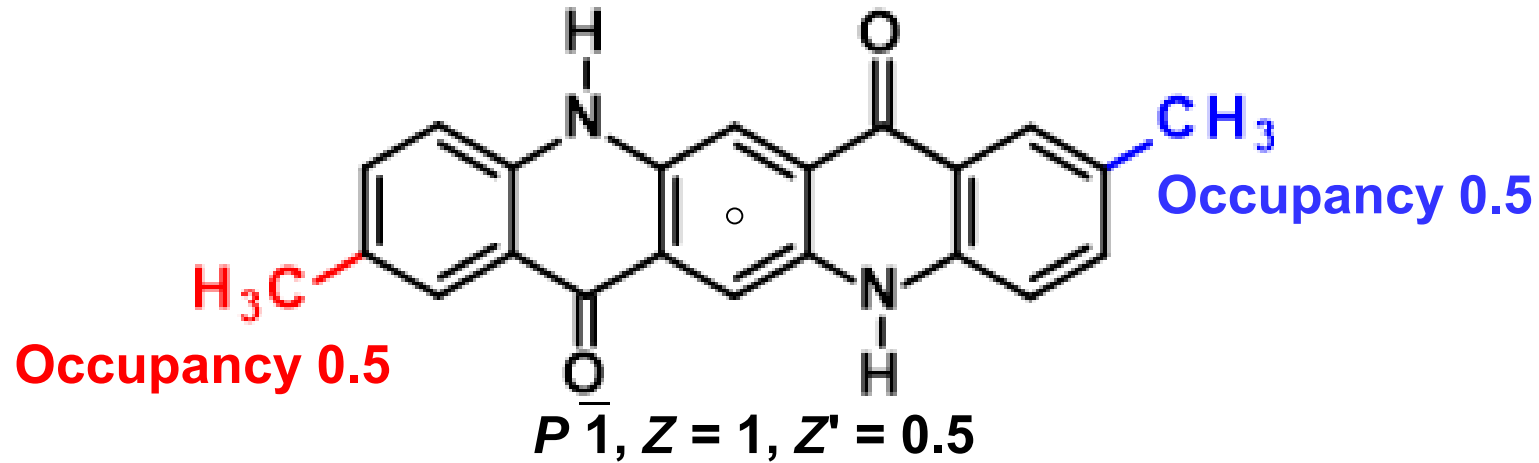
# Average structure



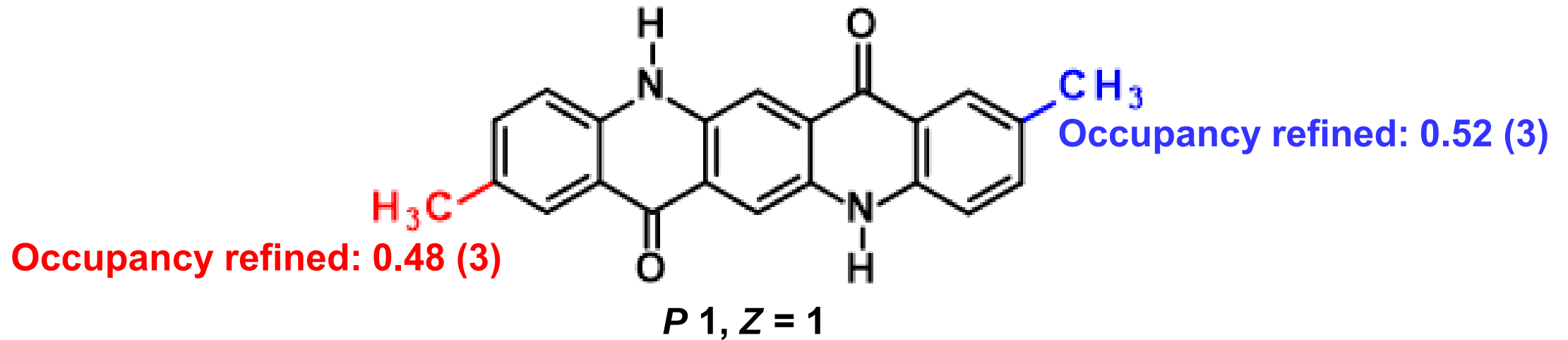
Molecule on inversion centre



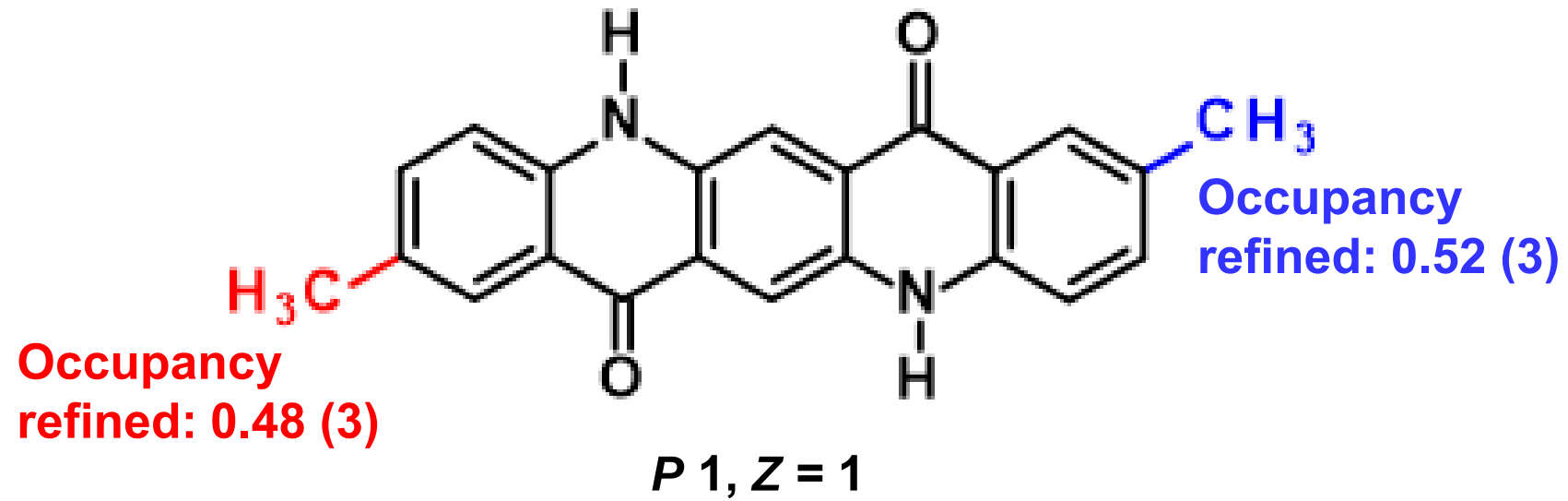
# Average structure



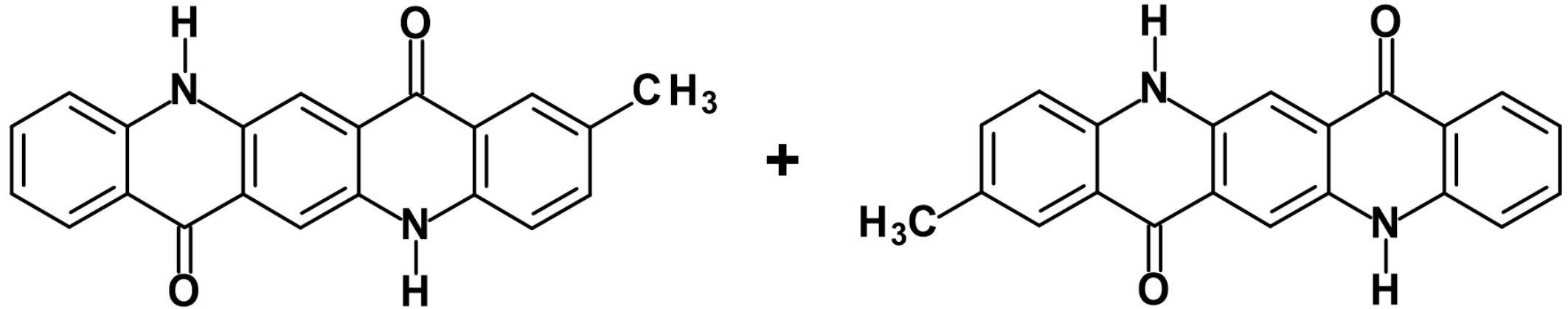
Molecule on inversion centre



# Refinement of occupancies

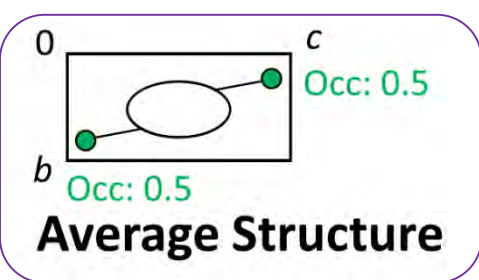


# Head-to-tail disorder

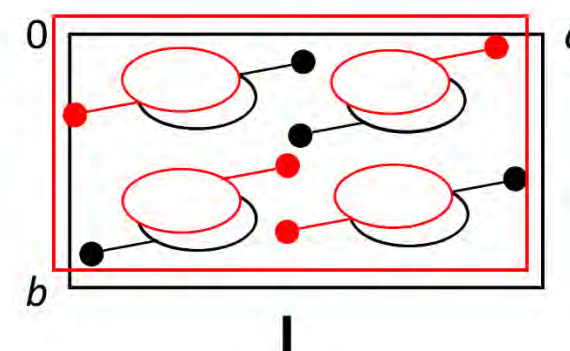
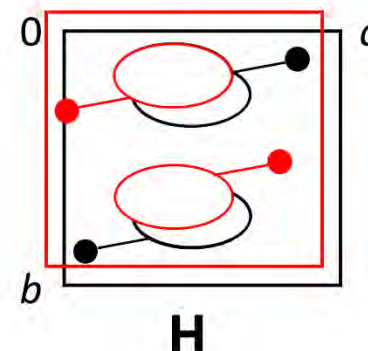
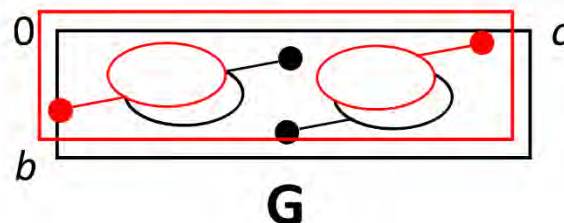
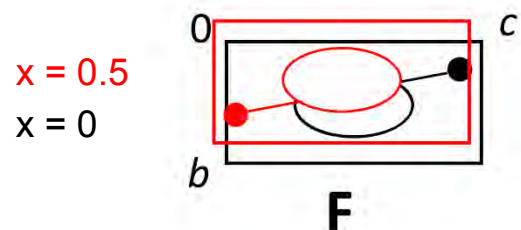
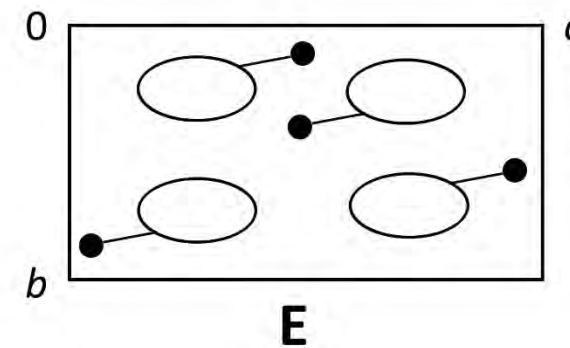
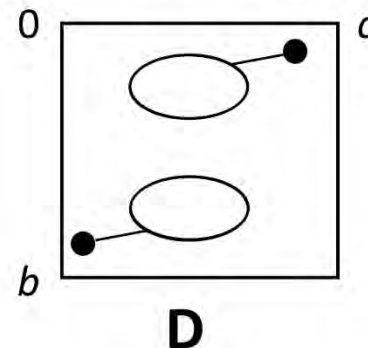
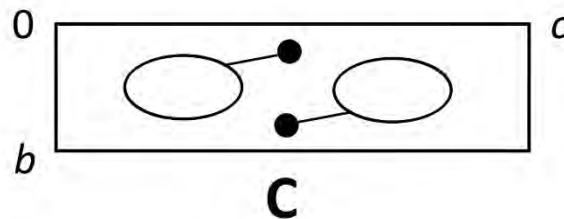
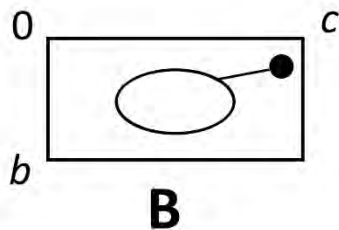


## Local ordering?

- No superstructure reflections visible
- No diffuse scattering visible (Powder data!)

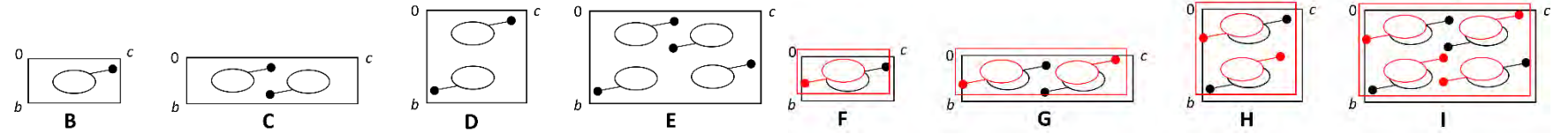


# Construction of ordered models

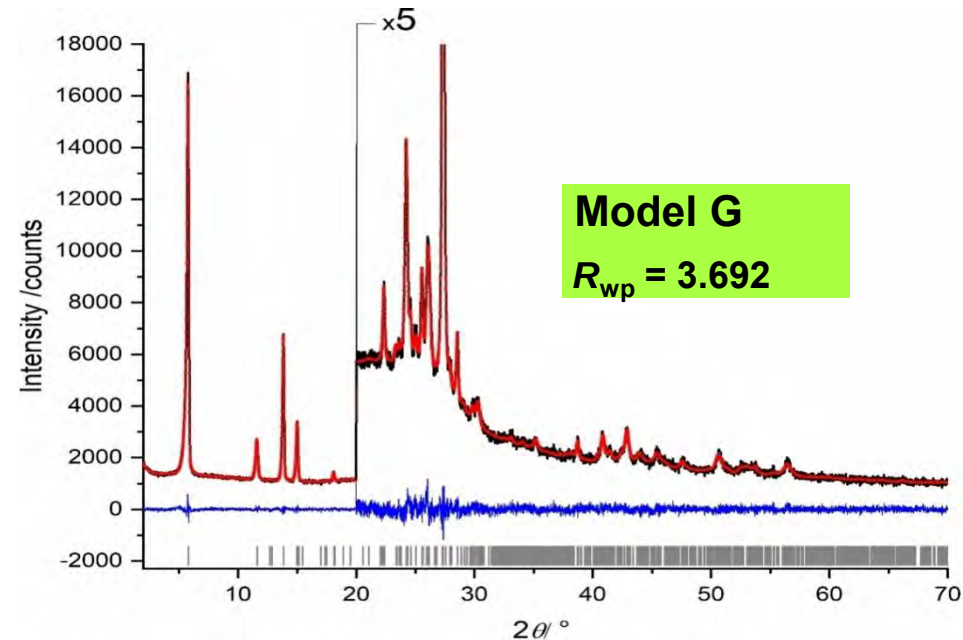
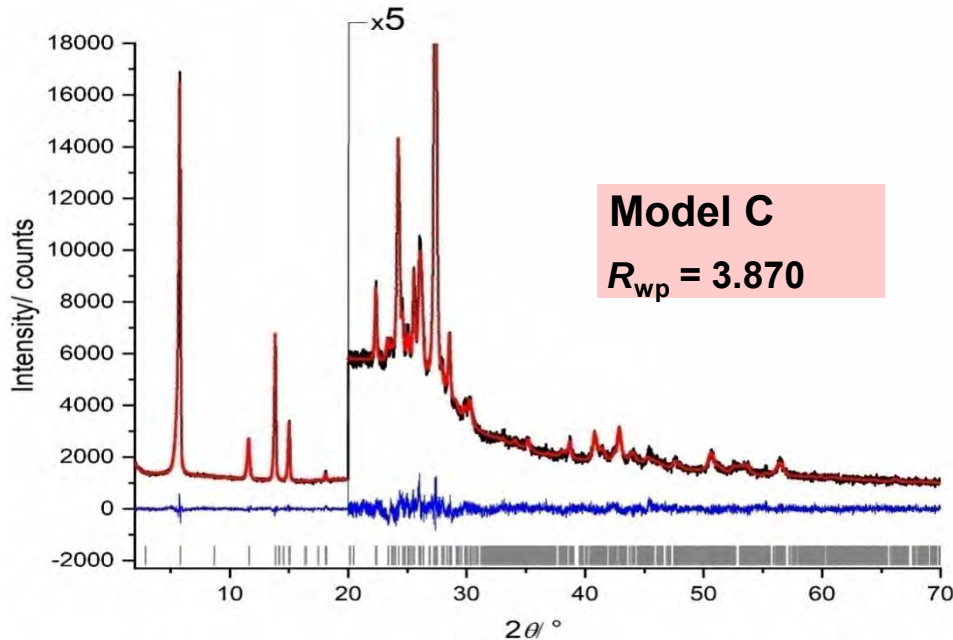


- Rietveld refinements
- Refinement to the Pair-Distribution Function (PDF)
- Lattice-energy minimisations

# Rietveld refinements

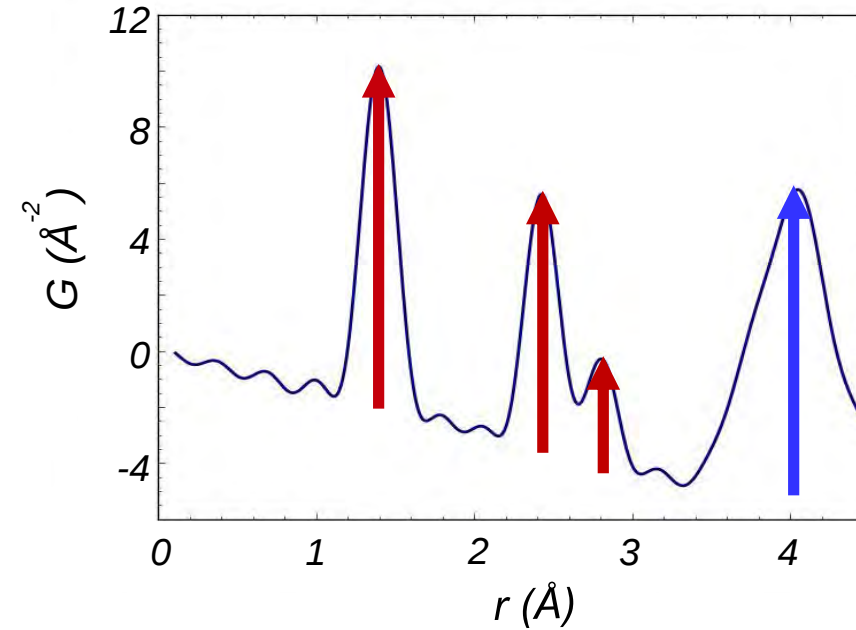
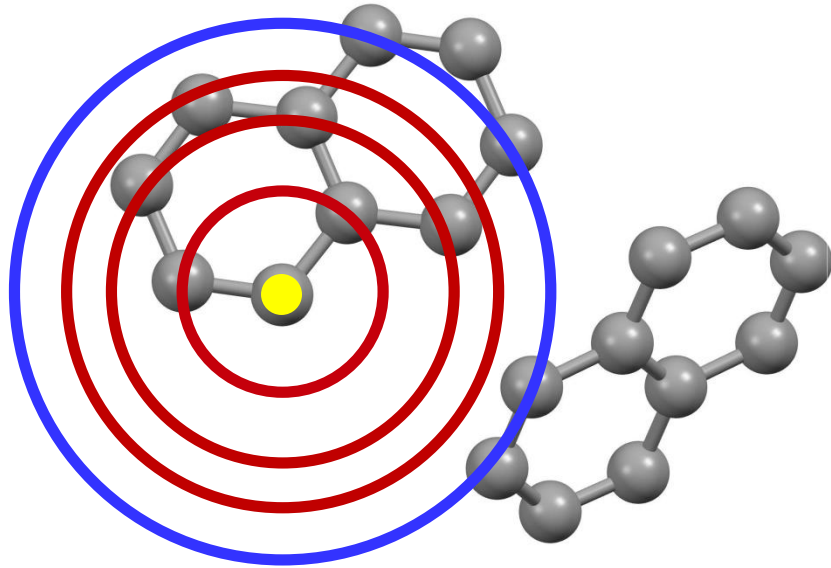


| Structural model | Average structure | B     | C          | D          | E          | F          | G          | H          | I          |
|------------------|-------------------|-------|------------|------------|------------|------------|------------|------------|------------|
| Space group      | $P\bar{1}$        | $P1$  | $P\bar{1}$ | $P\bar{1}$ | $P\bar{1}$ | $P\bar{1}$ | $P\bar{1}$ | $P\bar{1}$ | $P\bar{1}$ |
| Z                | 1                 | 1     | 2          | 2          | 2          | 2          | 2          | 2          | 2          |
| $R_{wp} / \%$    | 3.888             | 3.713 | 3.870      | 3.783      | 3.706      | 3.799      | 3.692      | 3.843      | 3.757      |
| $\chi^2$         | 1.065             | 1.020 | 1.063      | 1.039      | 1.018      | 1.044      | 1.014      | 1.056      | 1.032      |



No unique ordering

# The Pair-Distribution Function (PDF)



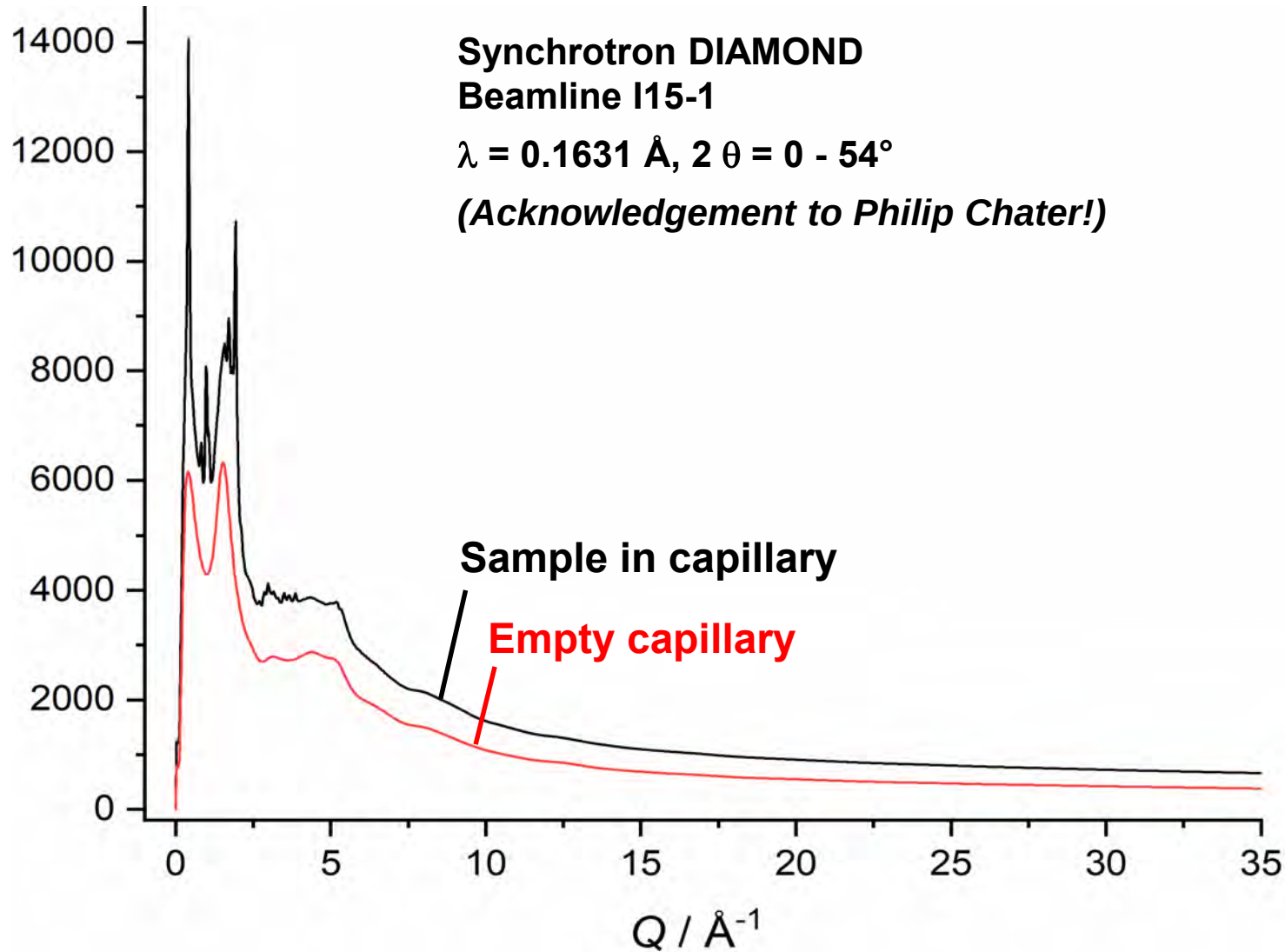
**PDF: Probability to find two atoms separated by a distance  $r$**

- weighted with the scattering power
- summed over all atoms
- normalised to a homogeneous atom density
- Fourier transform of the Intensities of a powder pattern (total scattering)

$$G(r) = 4\pi r[\rho(r) - \rho_0] = \frac{2}{\pi} \int_{\min}^{\max} Q[S(Q) - 1] \sin(Qr) dQ$$

# The Pair-Distribution Function (PDF)

I / counts



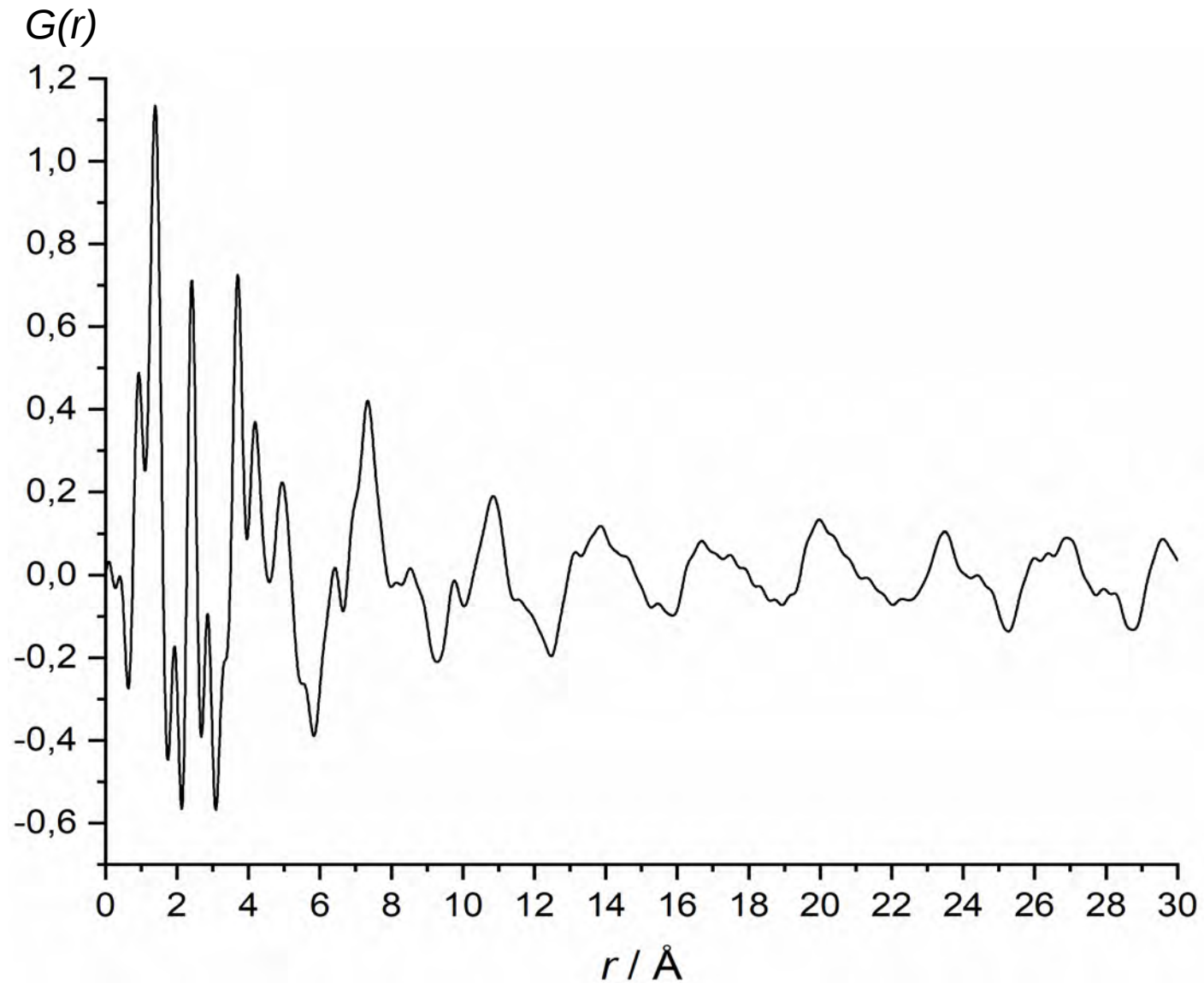
Careful  
background  
correction

Normalisation

Fourier  
transformation

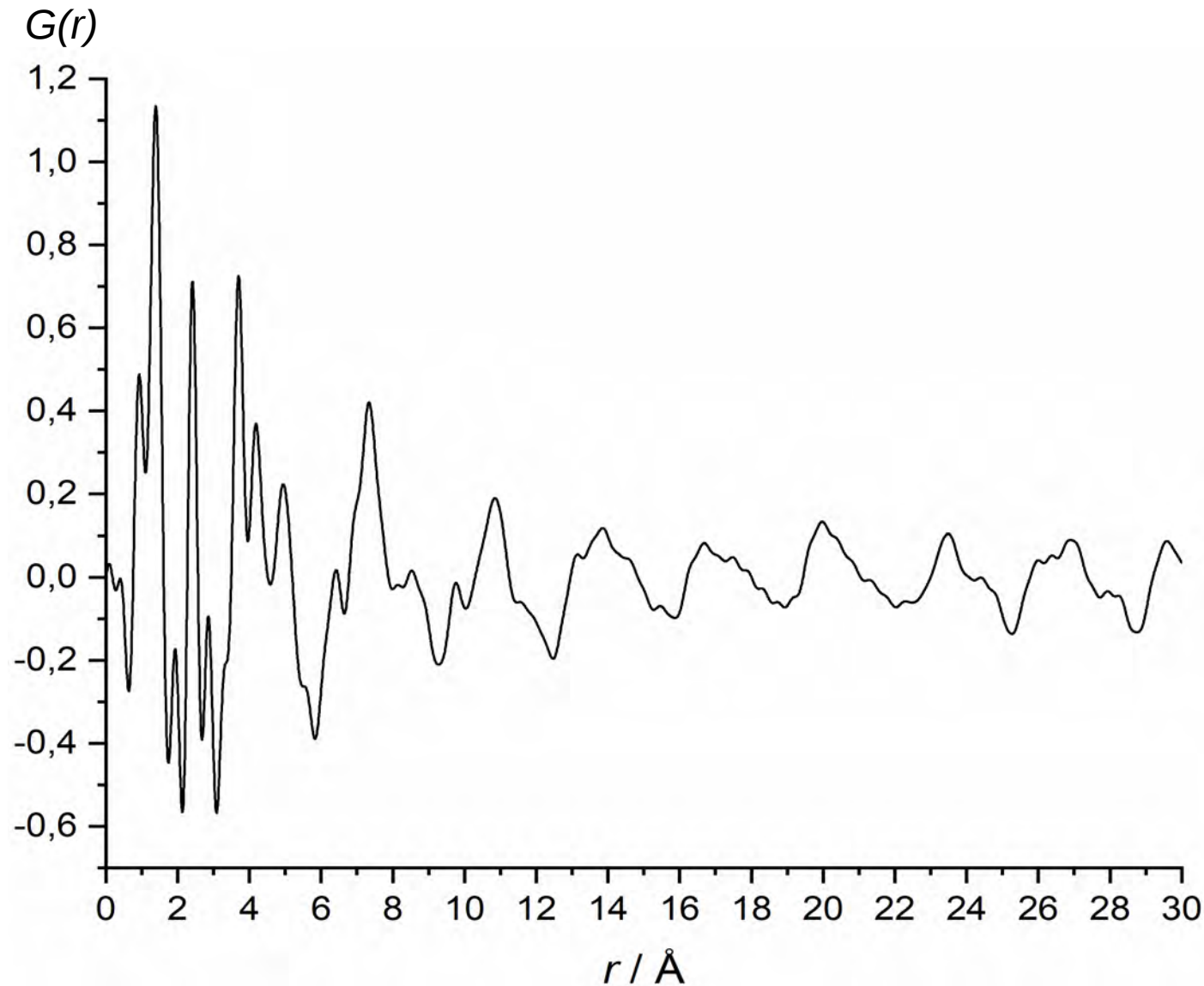
PDF

# Structure refinement by fit to the PDF



**Experimental PDF**

# Structure refinement by fit to the PDF



**Experimental PDF**

**Input: Structural models B - I**

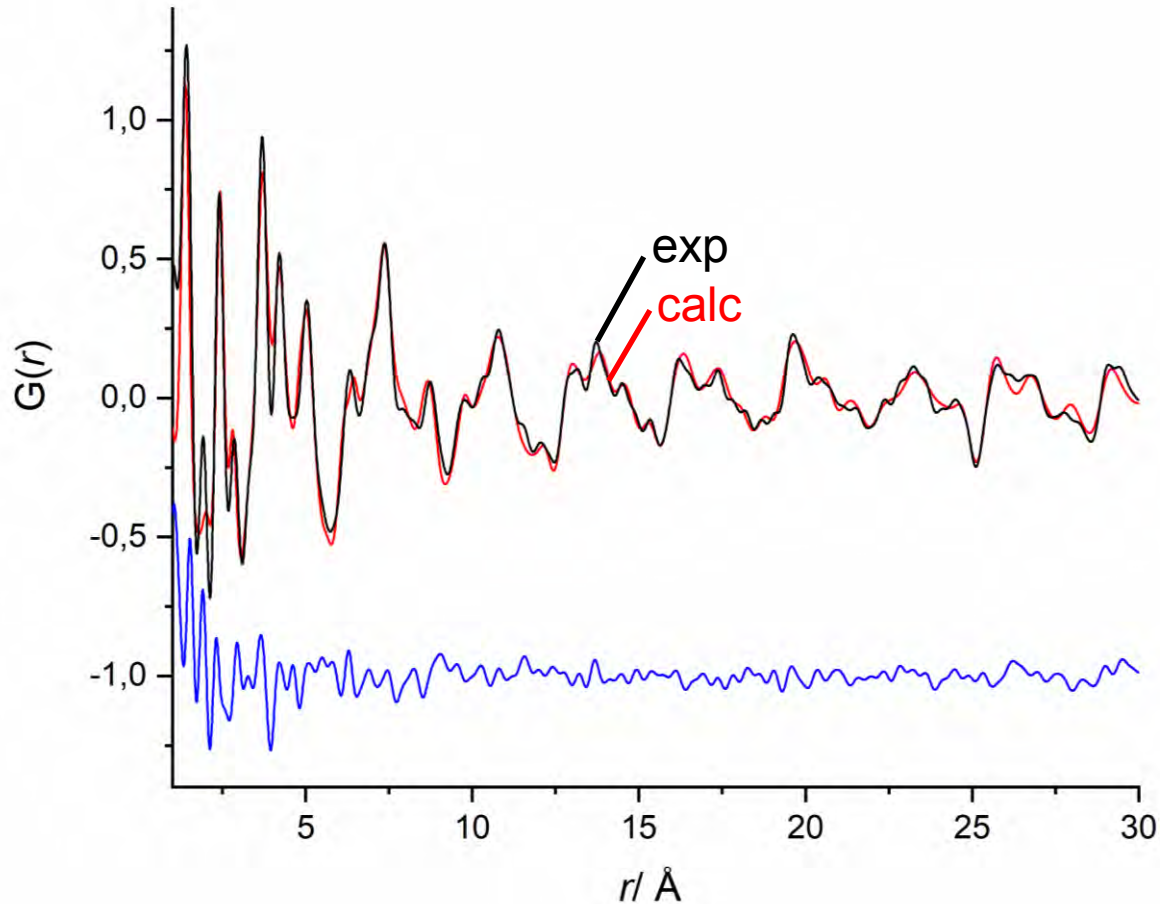
**Refinement of:**

- **Lattice parameters**
- **Atomic coordinates** (Molecules rigid)
- **Scale factor**
- **Peak widths** (described by  $B_{\text{intra}}$  and  $B_{\text{inter}}$ )  
[N. Rademacher et al., *J. Appl. Cryst.* 45, 482-488 (2012).  
D. Prill et al., *J. Appl. Cryst.* 48, 171-178 (2015).  
D. Prill et al., *Acta Cryst. A* 72, 62-72 (2016).]
- **Dampening of PDF at higher  $r$**

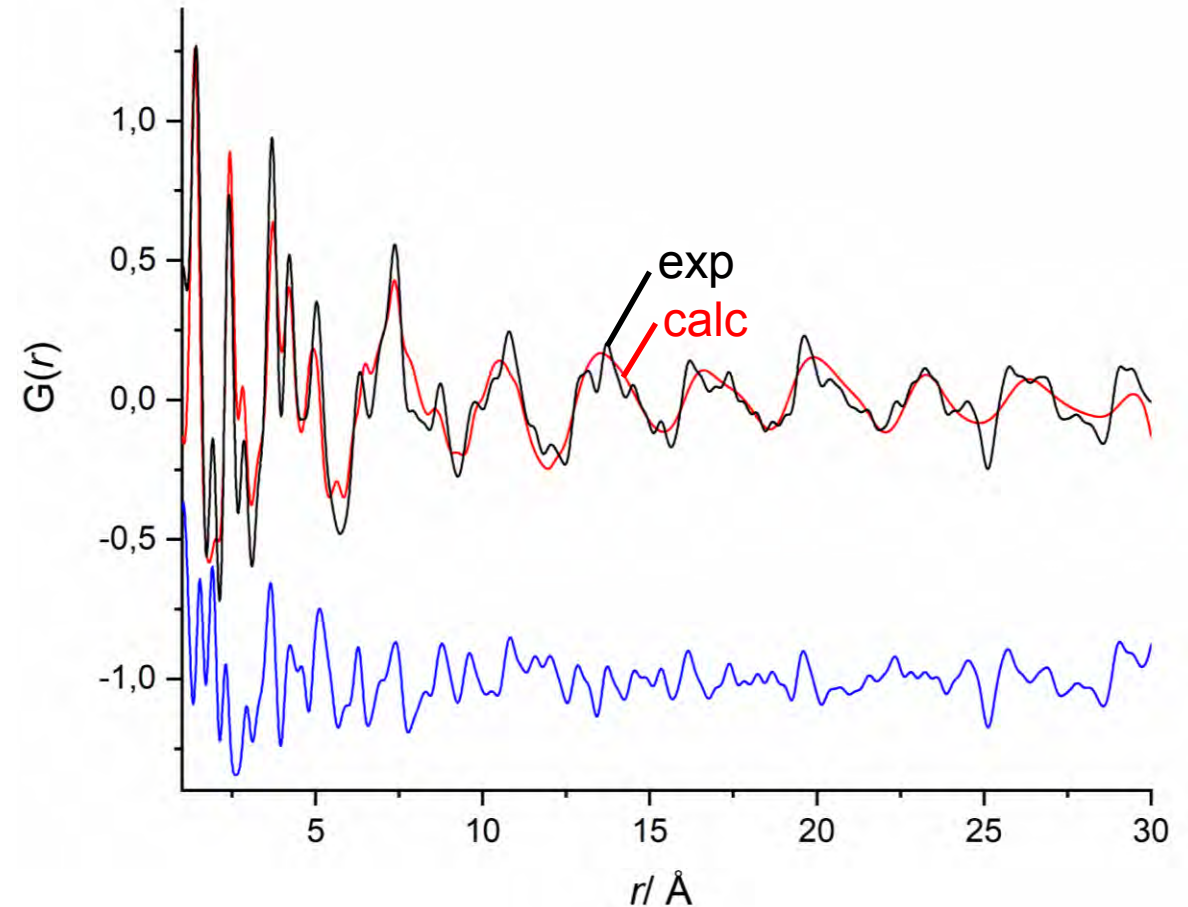
**Programs:**

- **PDFgetX3** (calculation of the PDF curve)
- **TOPAS-6** (for structure fit to the PDF)  
[Acknowledgement to Alan Coelho!]  
Calculation time: 20-30 min per structure

# Structure refinement by fit to the PDF

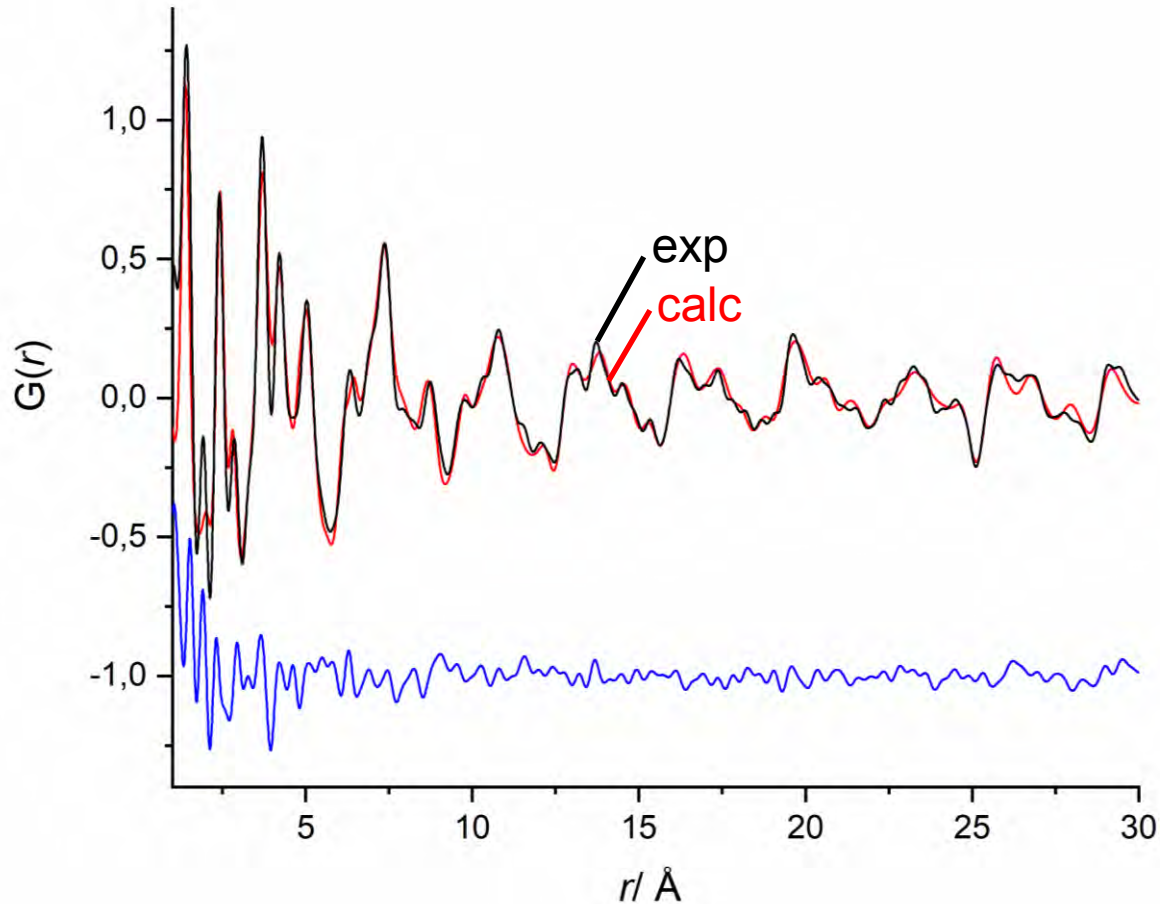


**Model E**  
(best fit to the PDF,  $R_{\text{wp}}^{\text{PDF}} = 0.341$ )

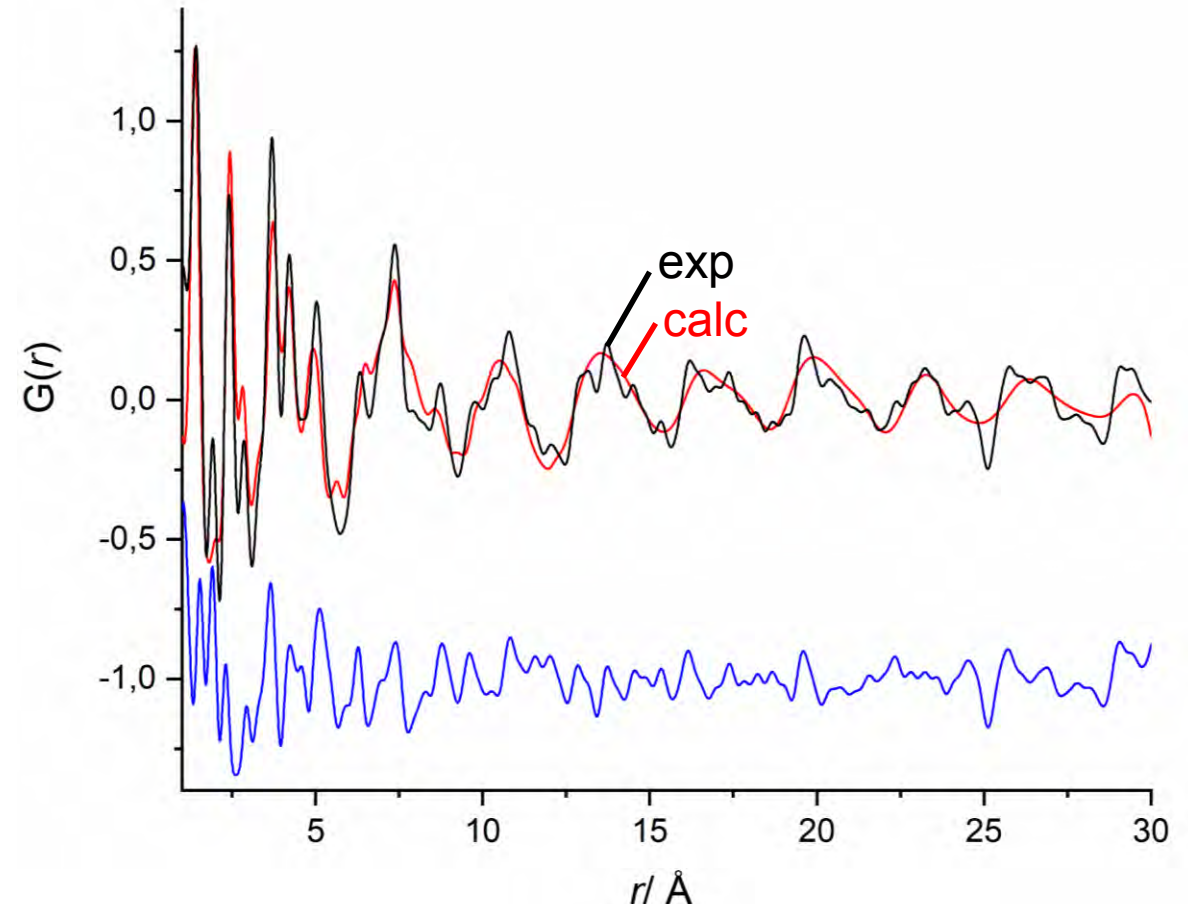


**Model I**  
(worst fit to the PDF,  $R_{\text{wp}}^{\text{PDF}} = 0.469$ )

# Structure refinement by fit to the PDF



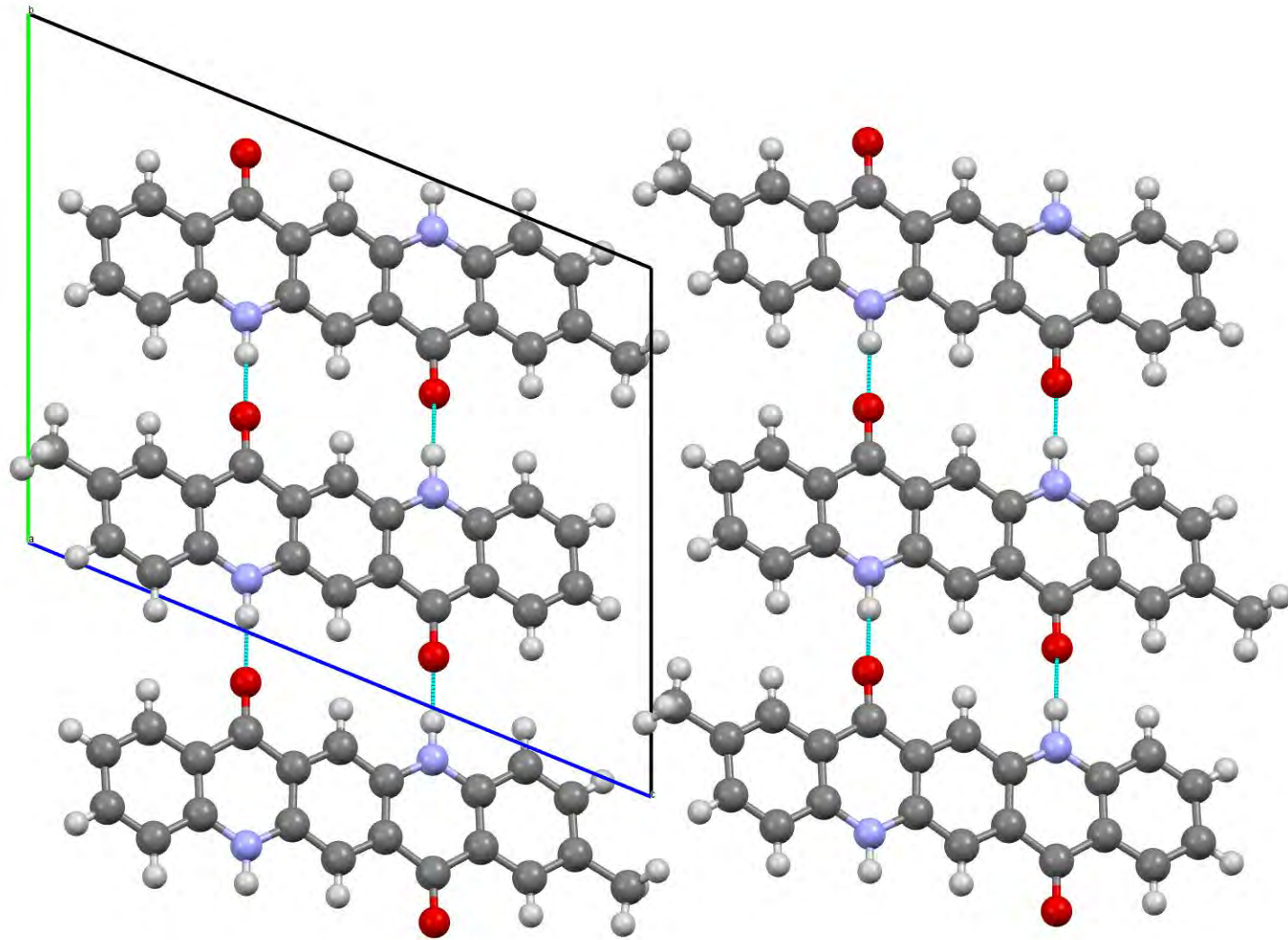
**Model E**  
(best fit to the PDF,  $R_{\text{wp}}^{\text{PDF}} = 0.341$ )



**Model I**  
(worst fit to the PDF,  $R_{\text{wp}}^{\text{PDF}} = 0.469$ )

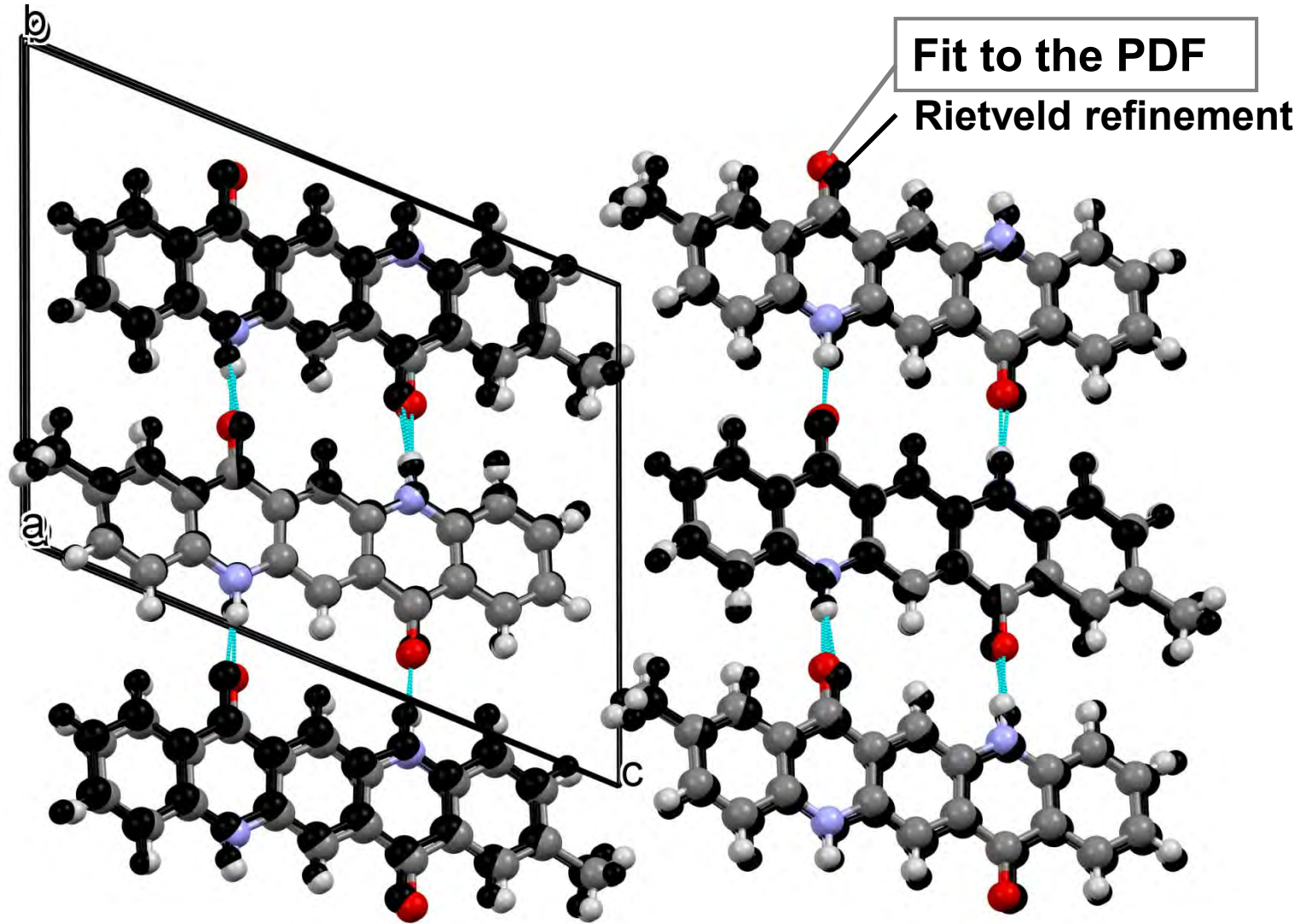
**Refinement of organic structures to PDF data:**  
**- sensitive to small structural deviations**

# Structure refinement by fit to the PDF



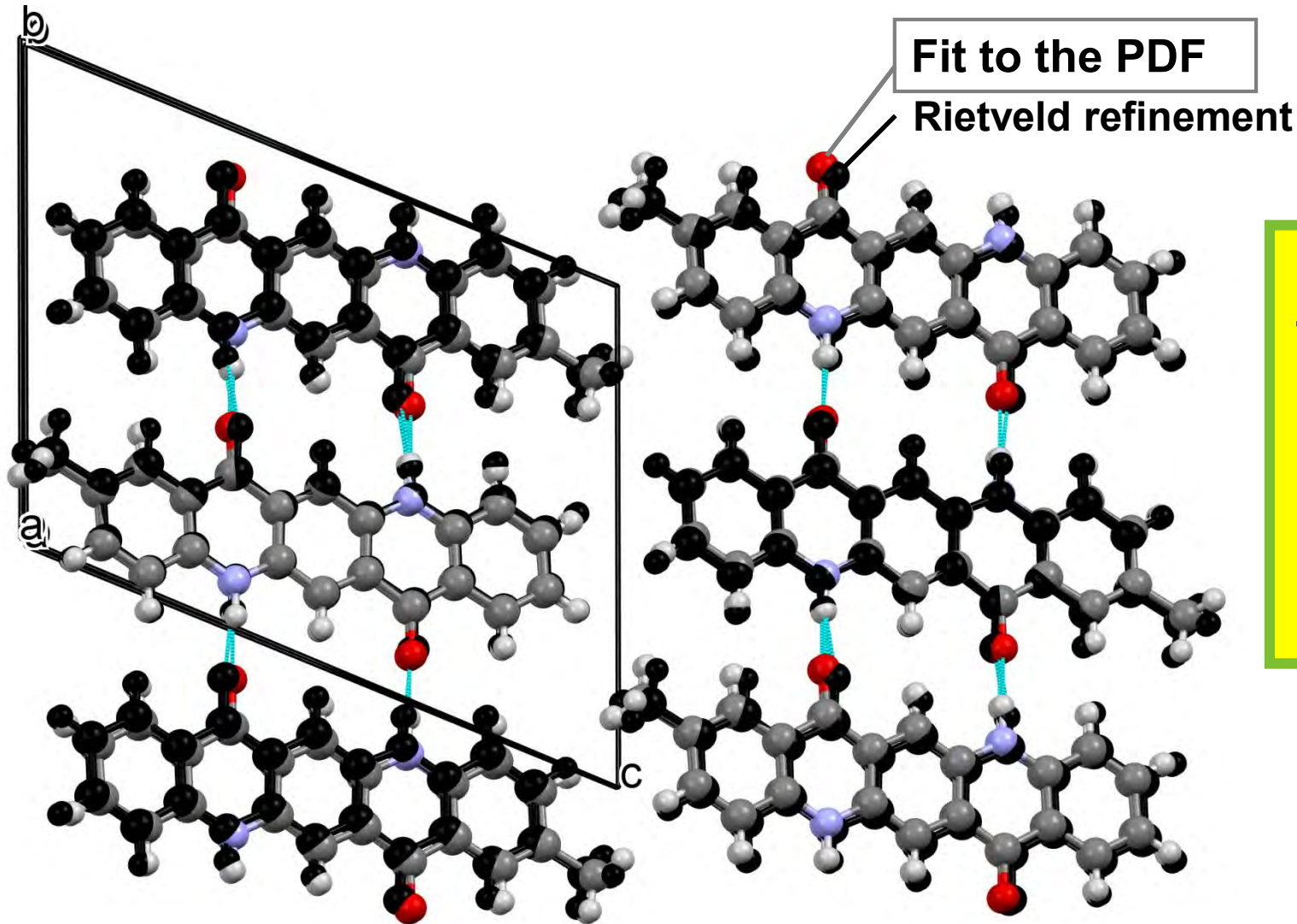
**Model E (best fit to the PDF)**

# Structure refinement by fit to the PDF



Model E (best fit to the PDF)

# Structure refinement by fit to the PDF

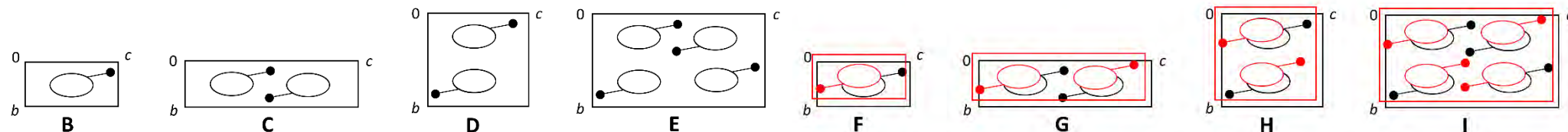


**Refinement of organic structures to PDF data:**

- feasible
- reliable
- sensitive to small structural deviations

**Model E (best fit to the PDF)**

# Structure refinement by fit to the PDF



| Structural model | B     | C     | D     | E     | F     | G     | H     | I     |
|------------------|-------|-------|-------|-------|-------|-------|-------|-------|
| $R_{wp}^{PDF}$   | 0.364 | 0.345 | 0.377 | 0.341 | 0.450 | 0.450 | 0.354 | 0.469 |

Different local structures possible

# Lattice energy minimisations

## Input:

- Models B - I

## Method:

- Dreiding force field
- Charges: HF/6-31G\*\* + ESP

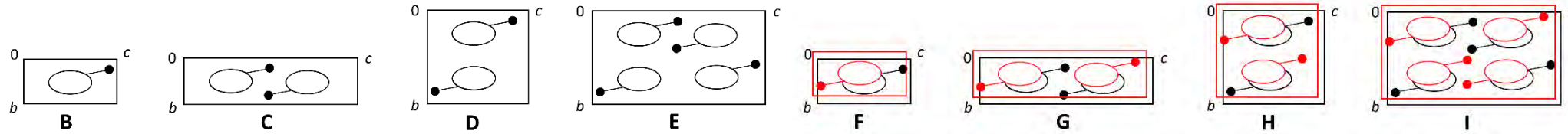
## Refinement of:

- Lattice parameters
- All atomic coordinates (flexible molecules)

## Program for lattice-energy minimisations:

- Materials Studio

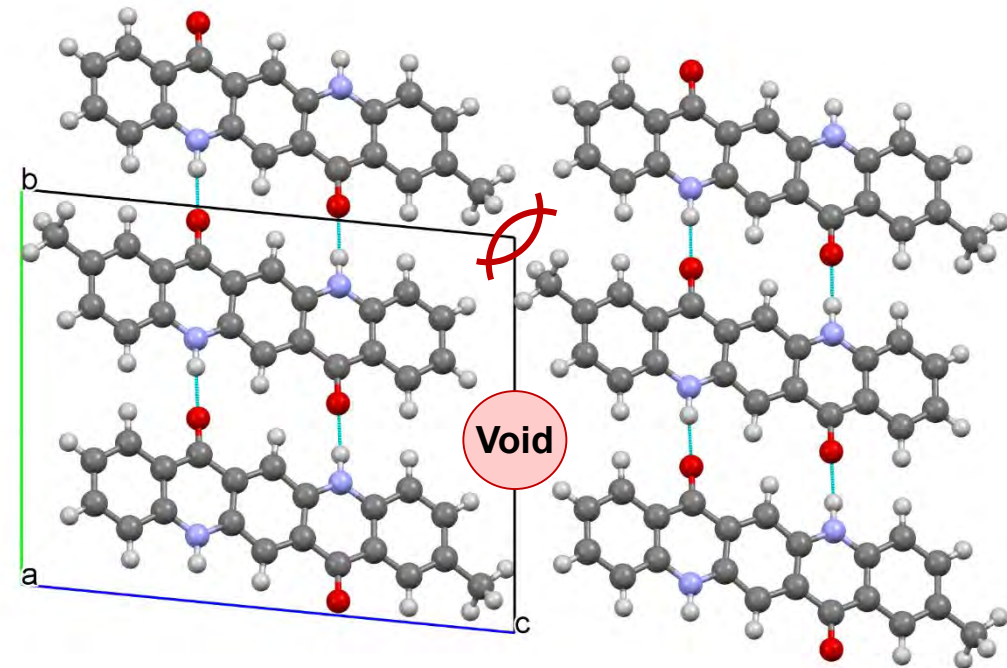
# Lattice energy minimisations



| Structural model                 | B     | C     | D     | E     | F     | G     | H     | I     |
|----------------------------------|-------|-------|-------|-------|-------|-------|-------|-------|
| $E_{\text{rel}} / \text{kJ/mol}$ | 1.84  | 0.5   | 6.48  | 0.63  | 0.04  | 1.76  | 4.89  | 0     |
| $V_{\text{mol}}$                 | 369.2 | 369.7 | 384.1 | 369.1 | 366.7 | 370.7 | 377.2 | 366.4 |

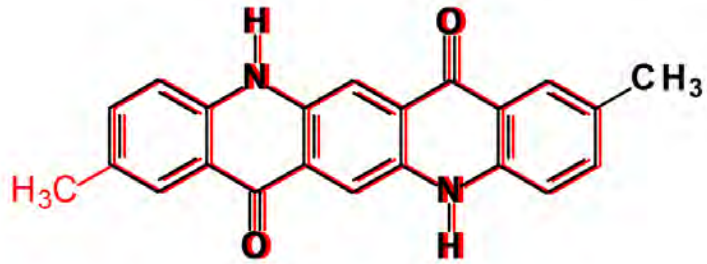
Different local structures possible

Unfavorable local structure  
(Models D + H):



# Local structure

Superposition of molecules:

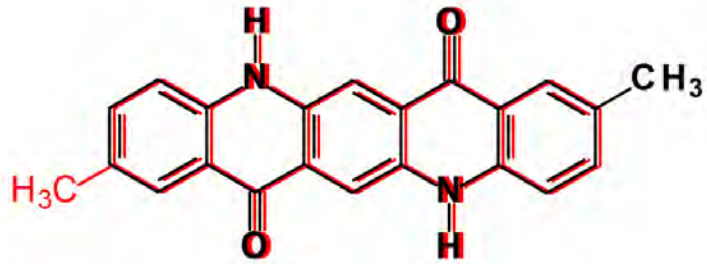


or

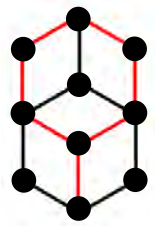
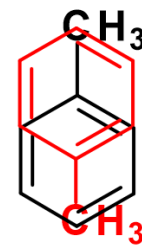
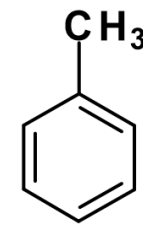


# Local structure

Superposition of molecules:

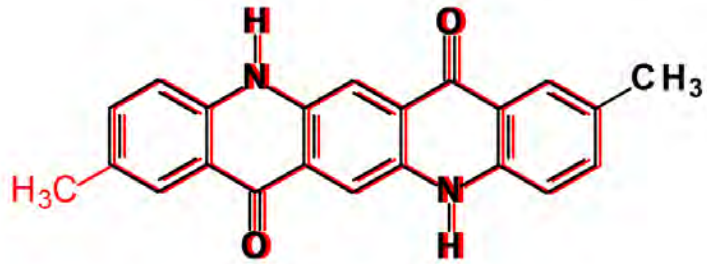


or



# Local structure

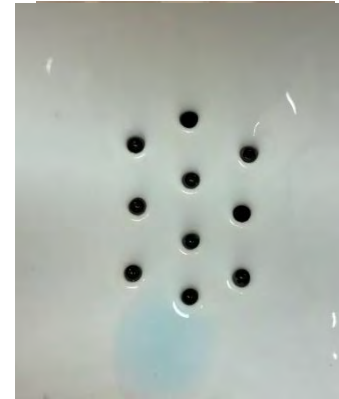
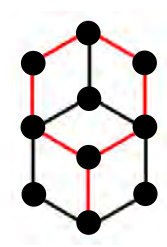
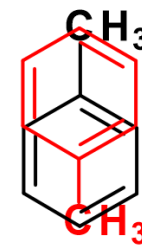
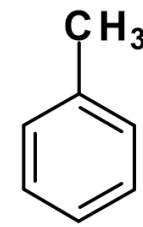
Superposition of molecules:



or



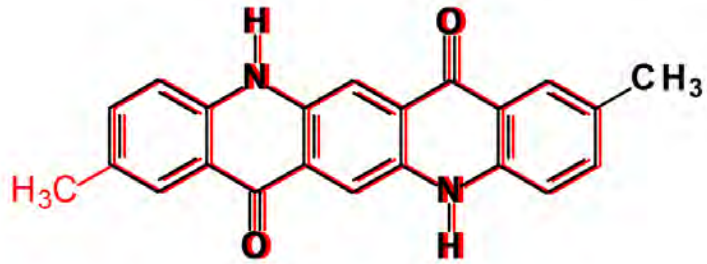
?



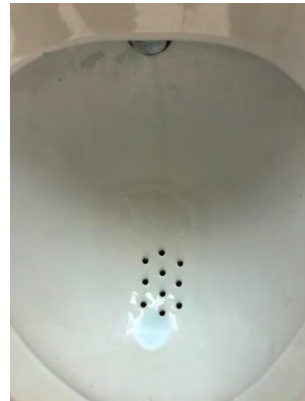
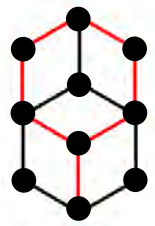
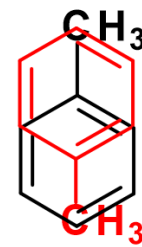
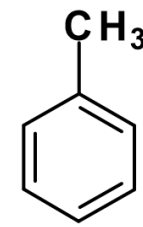
(Idea + Photo:  
Peter Müller)

# Local structure

Superposition of molecules:



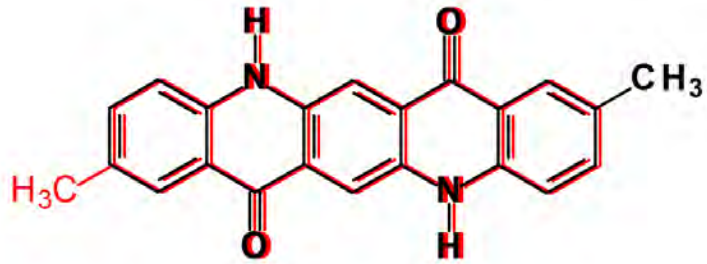
or



(Idea + Photo:  
Peter Müller)

# Local structure

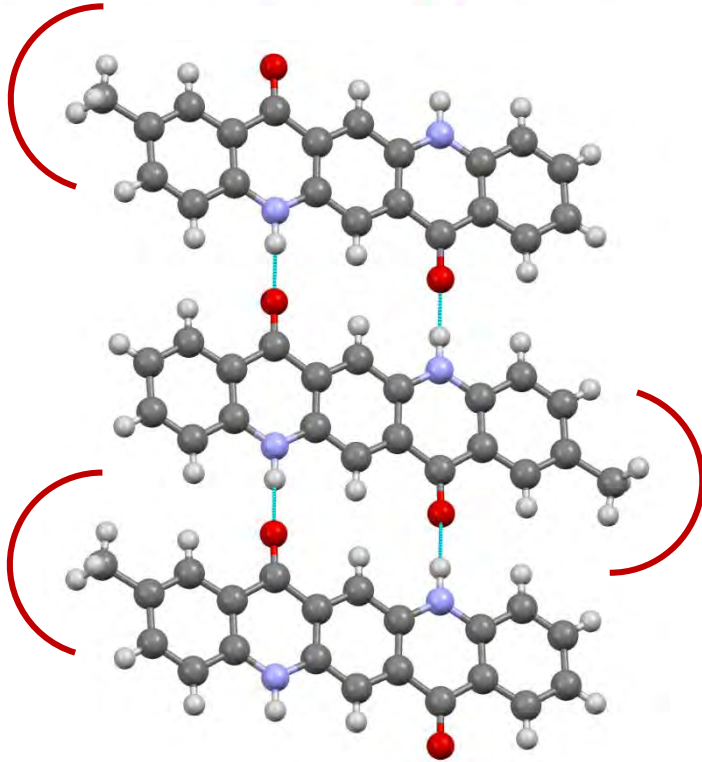
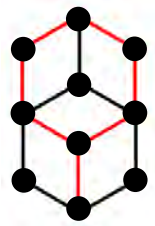
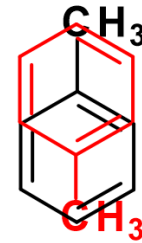
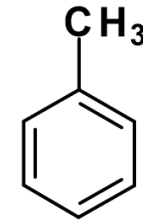
Superposition of molecules:



or

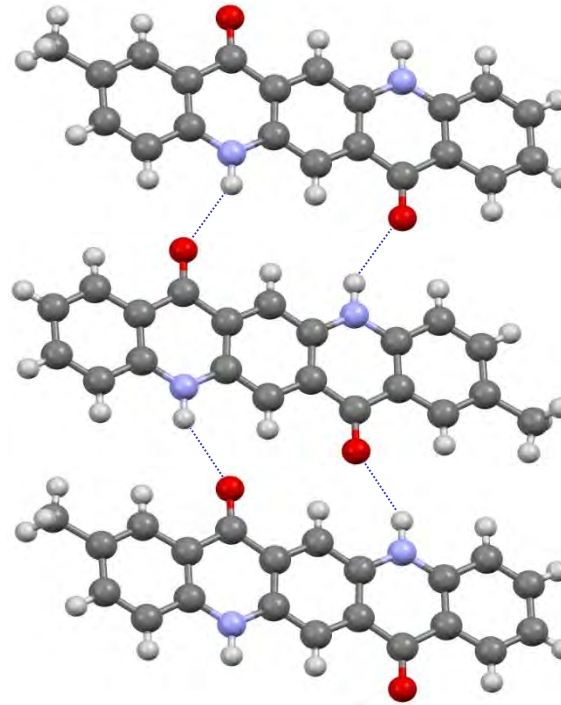


?



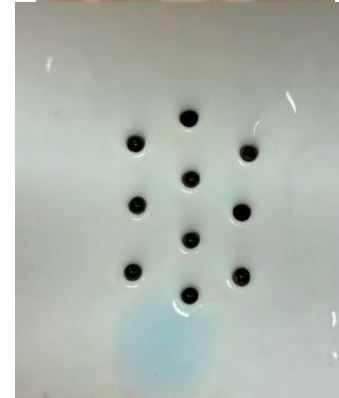
Good H-bonds

Sterically demanding



Worse H-bonds

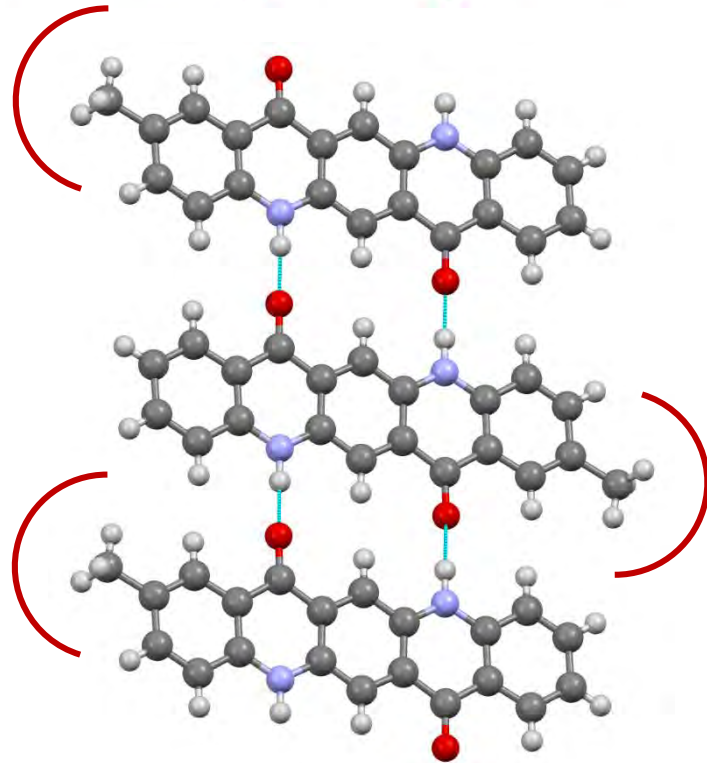
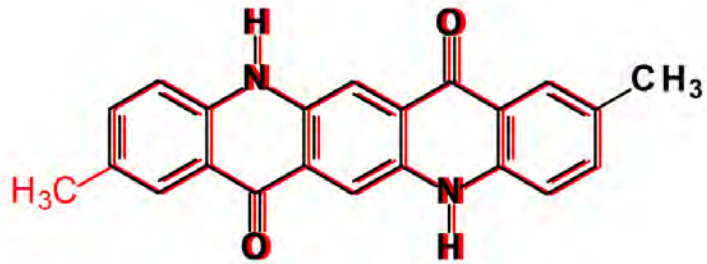
Better space filling



(Idea + Photo:  
Peter Müller)

# Local structure

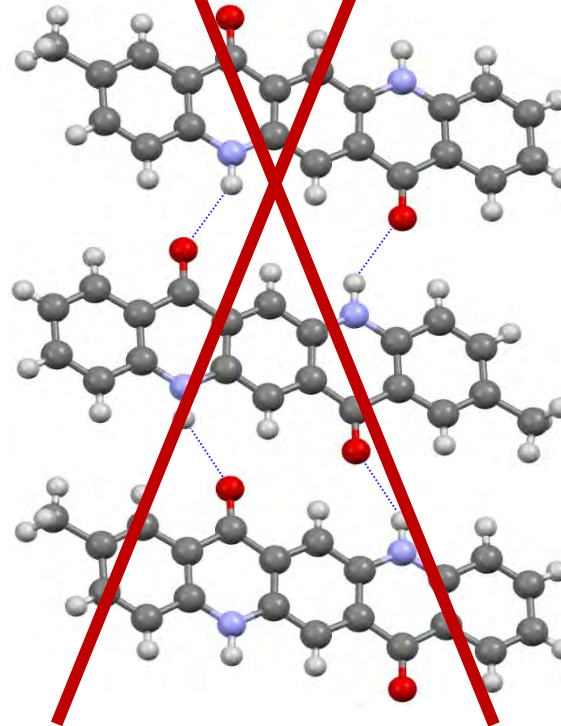
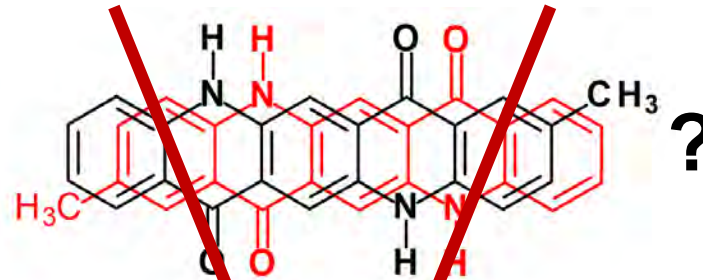
Superposition of molecules:



Good H-bonds

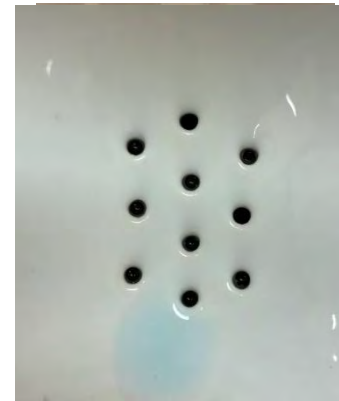
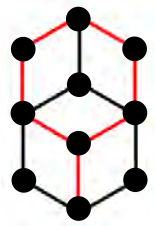
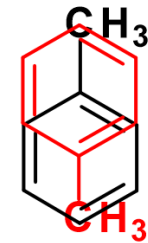
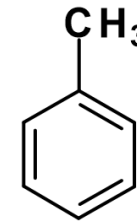
Sterically demanding

or



Worse H-bonds

Better space filling



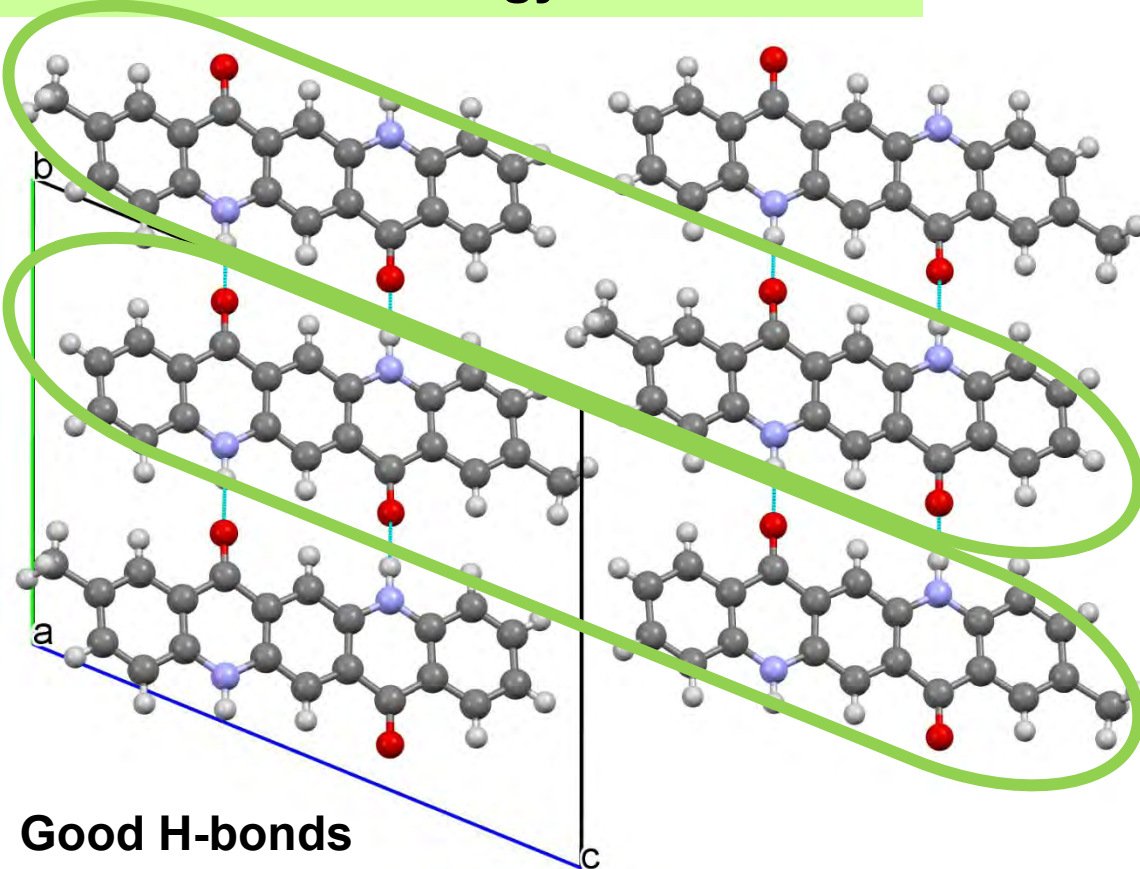
(Idea + Photo:  
Peter Müller)

Never observed

# Local structure

Typical local structure: Model E

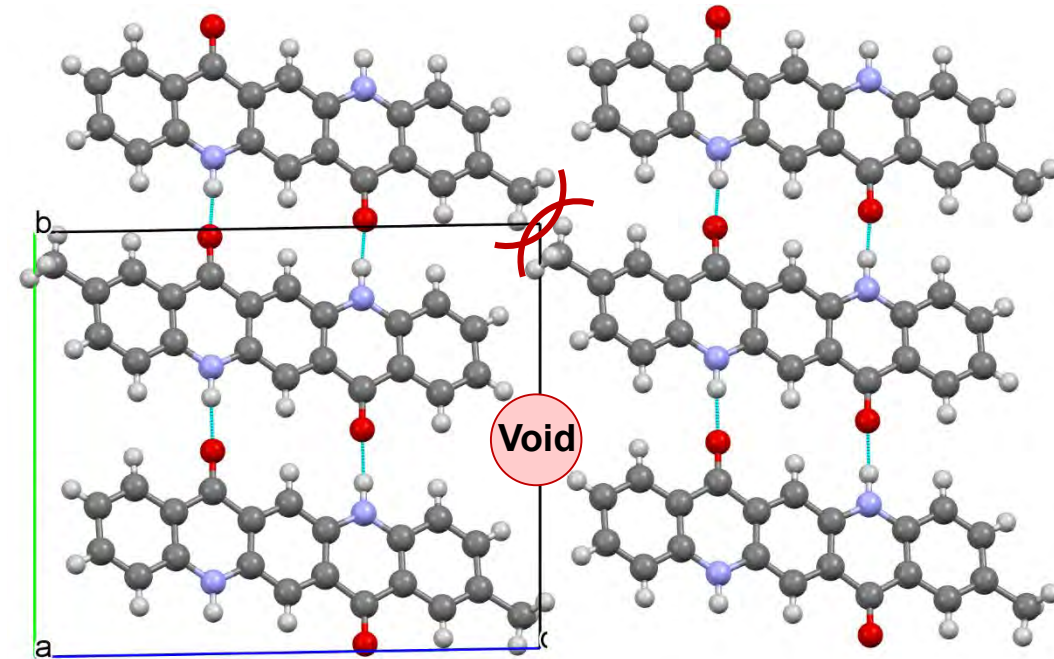
- Good Rietveld fit
- Best PDF fit
- Good lattice energy



Good H-bonds

Good space filling due to local ordering

Unfavorable local structure:

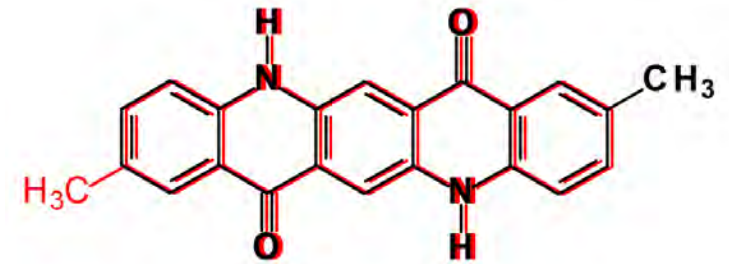


Local ordering  
of neighbouring molecules  
in this direction

No superstructure reflections

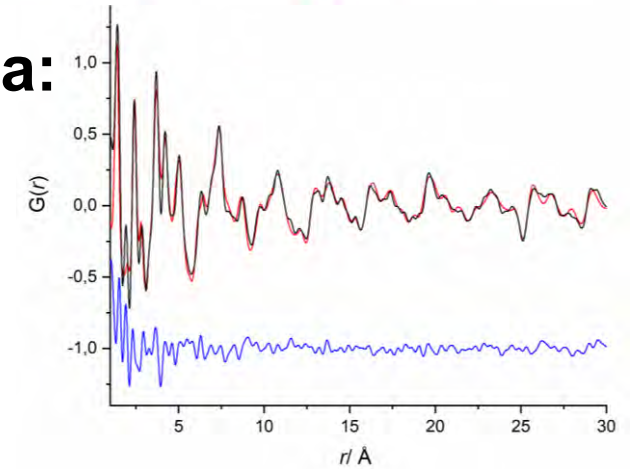
# Conclusion

1. Statistical head-to-tail disorder  
with local ordering of molecules in 1 direction



2. Refinement of organic crystal structures to PDF data:

- feasible
- reliable
- sensitive to small structural deviations



Acknowledgements:

- Carina Schlesinger
- Dragica Prill