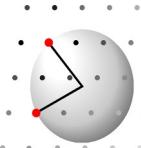

DISCUS Workshop

Single crystal PDF 3D-PDF 3D- Δ -PDF Chemical short range order

Reinhard B. Neder
Crystallography and Structural Physics
Friedrich-Alexander-Universität Erlangen-Nürnberg

reinhard.neder@fau.de



Pair Distribution Function

Intensity: $I(\vec{h}) = F(\vec{h}) \cdot F^*(\vec{h})$

$$\begin{aligned}\mathcal{F}[I(\vec{h})] &= \mathcal{F}[F(\vec{h}) \cdot F^*(\vec{h})] \\ &= \mathcal{F}[F(\vec{h})] \circ \mathcal{F}[F^*(\vec{h})] \\ &= \rho(\vec{r}) \circ \rho^*(-\vec{r}) \\ &= \rho(\vec{r}) \circ \rho(-\vec{r}) = P(\vec{u}) \\ &= \text{Autocorrelation function } P(\vec{u})\end{aligned}$$

Intensity is product of
Structure factor and conj. complex

Fourier transform to go
back to direct space

Convolution of individual
Fourier transforms

Back to electron densities
real valued function of r

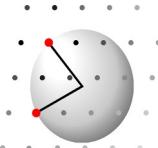
Auto correlation function: peaks at

$$P(\vec{u}) \quad \text{if} \quad \rho(\vec{r}) * \rho(\vec{r} + \vec{u}) \gg 0$$

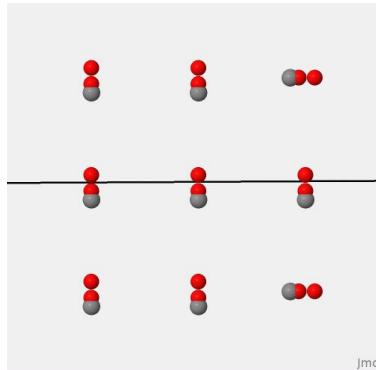
Vectors \mathbf{u} that correspond to interatomic vectors

Powder	Pair distribution function radial distribution function	PDF RDF	$G(\mathbf{r})$, $g(\mathbf{r})$ $RDF(\mathbf{r})$
--------	--	------------	--

Single crystal 3D-PDF / 3D- Δ -PDF



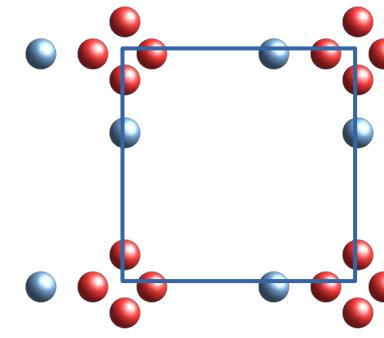
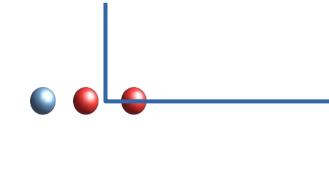
Linear **A-B-B** molecule in two orientations



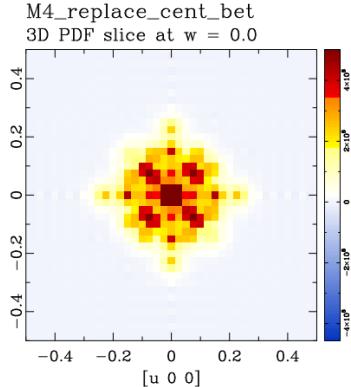
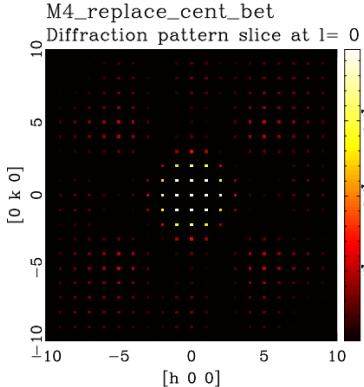
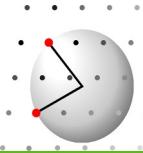
Complete structure

no correlation between molecules

Complete structure



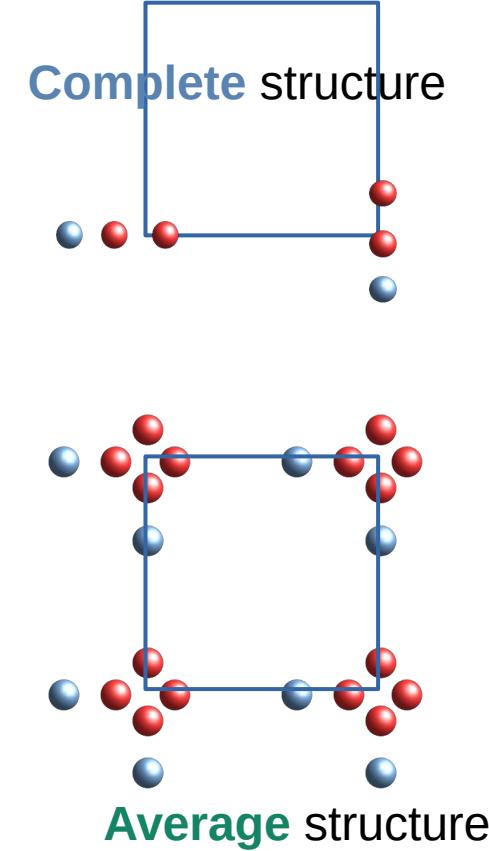
Average structure

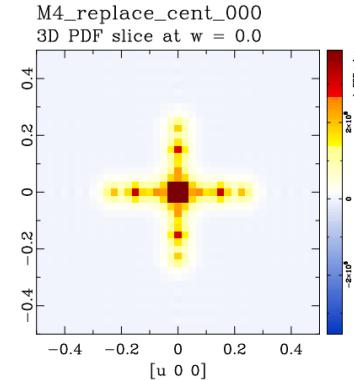
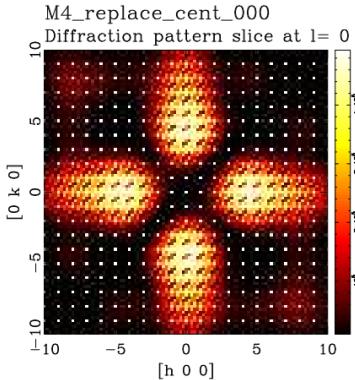
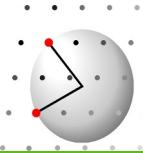


Patterson

Average structure Patterson

Positive peaks
interatomic vectors in average structure



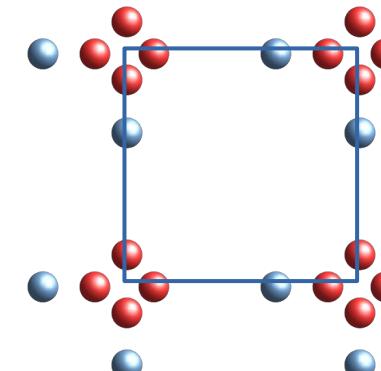
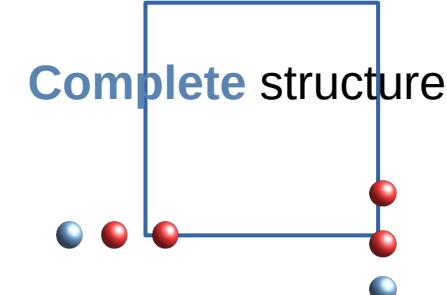


3D PDF

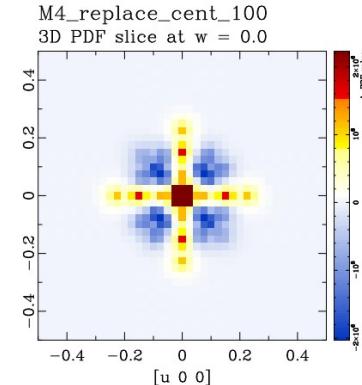
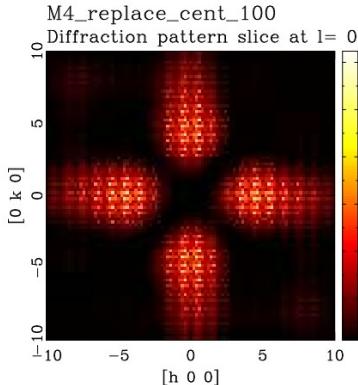
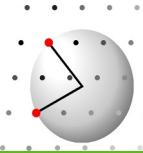
Complete structure Patterson

Positive peaks

interatomic vectors in actual / complete structure



Average structure

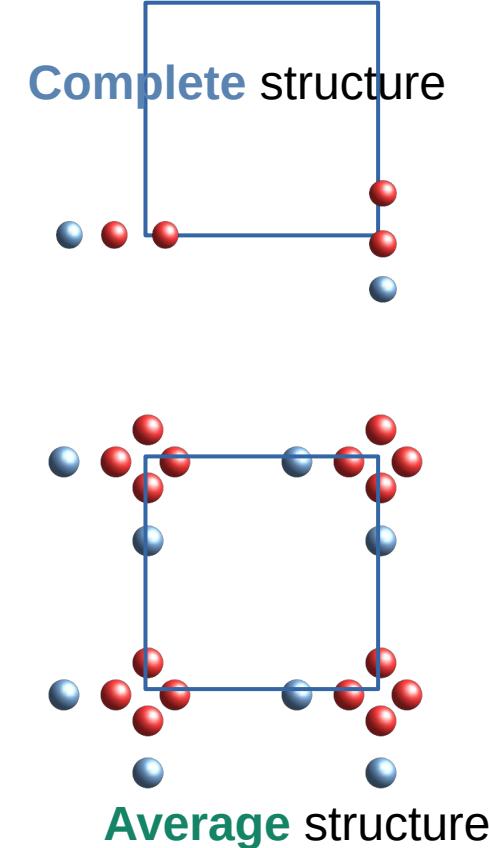


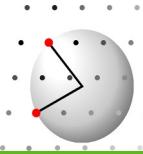
3D - Δ - PDF

„Local“ structure Patterson

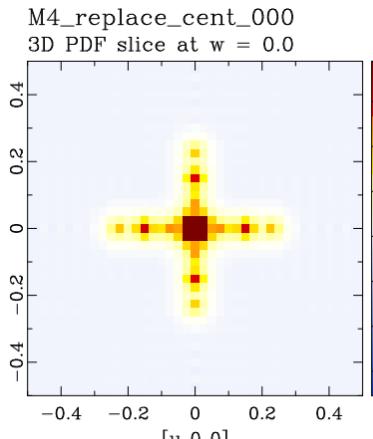
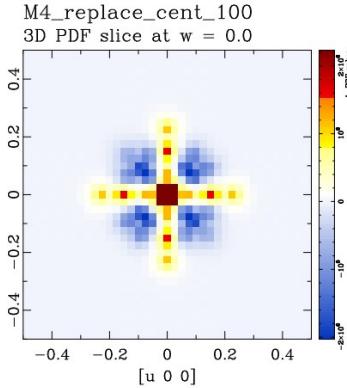
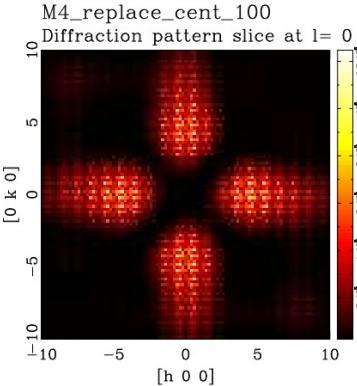
Difference: Complete – Average

Positive and Negative peaks
Deviations of local from average structure

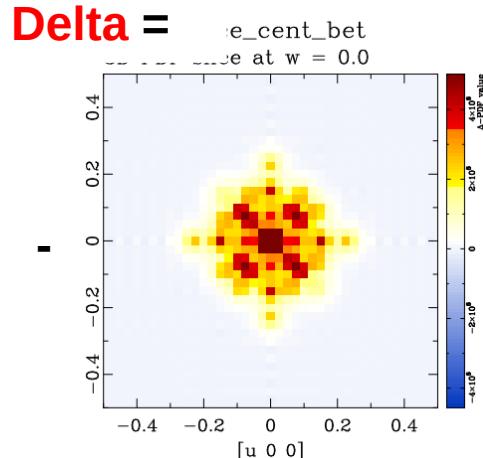




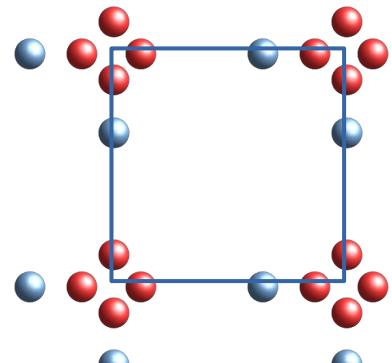
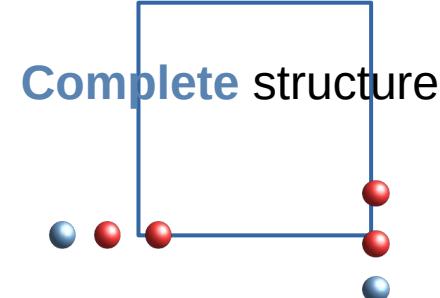
3D-PDF

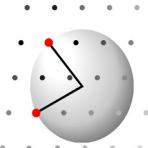


Complete



Average



**Patterson**

3-D inverse Fourier transform
of the **Bragg** scattering

$$P(\mathbf{u}) = F^{-1} I(\mathbf{hkl})$$

$$P(\mathbf{u}) = F^{-1} [F^*(\mathbf{hkl}) \cdot F(\mathbf{hkl})] = \rho(\mathbf{r}) \otimes \rho(-\mathbf{r})$$

$$P(\mathbf{u}) = F^{-1} [F^*(\mathbf{hkl}) \cdot E(\mathbf{hkl})]$$
 Sharpened **Patterson**

$$P(\mathbf{u}) = F^{-1} [E^*(\mathbf{hkl}) \cdot E(\mathbf{hkl})]$$
 Super sharpened **Patterson**

$$PDF(\mathbf{u}) = F^{-1} \frac{I(\mathbf{Q})}{\langle f^2 \rangle}$$

Super sharpened **3D PDF**

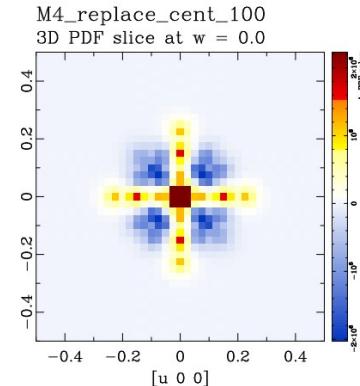
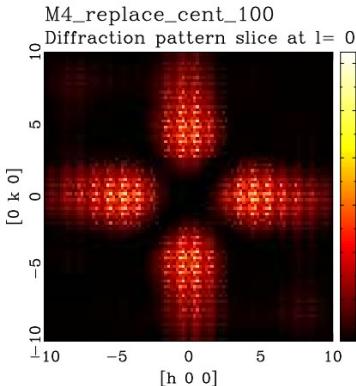
$$E(\mathbf{hkl}) = \frac{F(\mathbf{hkl})}{\sqrt{\langle I(\mathbf{hkl}) \rangle_{\Delta\Theta}}}$$

Interatomic vectors
average structure

periodic



3D-PDF

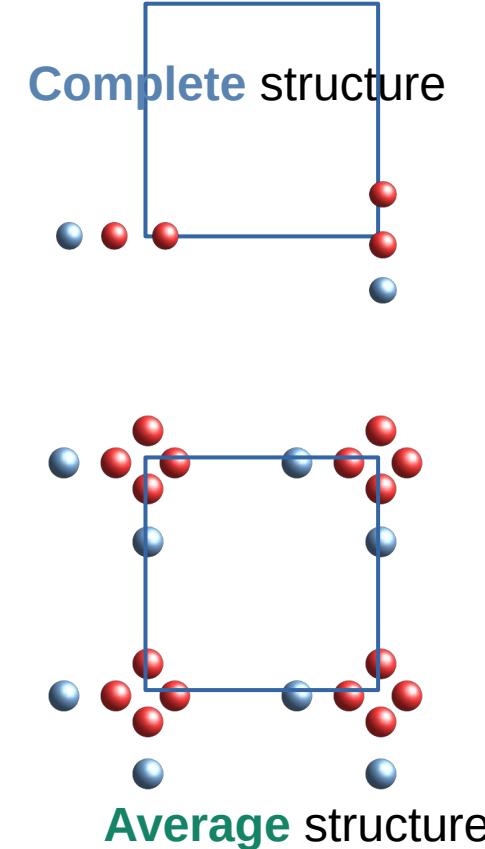


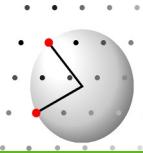
3D - Δ - PDF

„Local“ structure Patterson

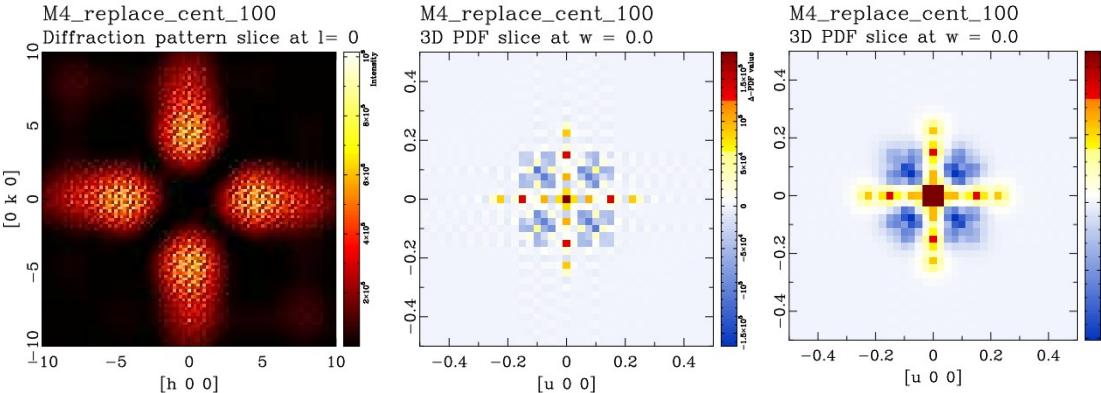
Difference: Complete – Average

Positive and Negative peaks
Deviations of local from average structure





3D-PDF

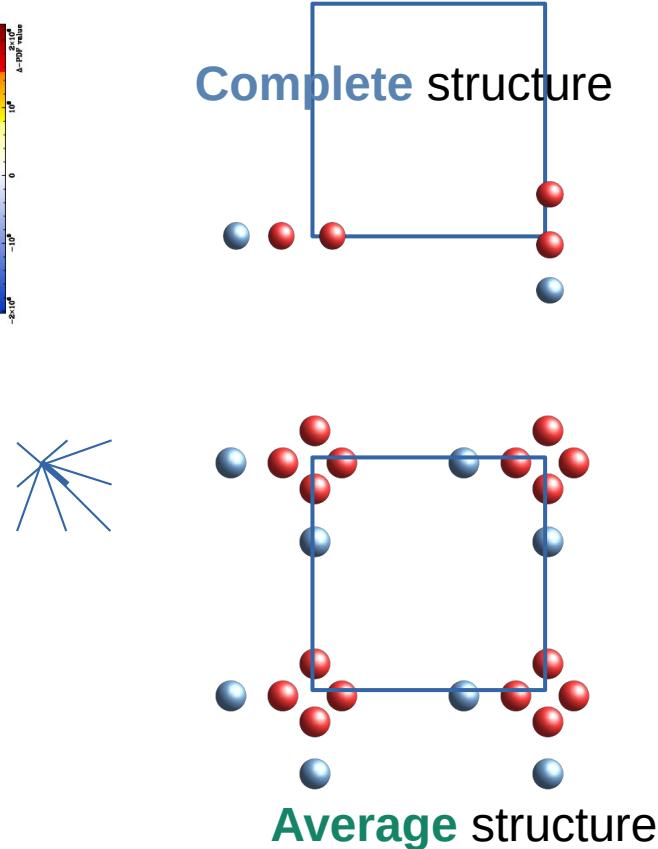


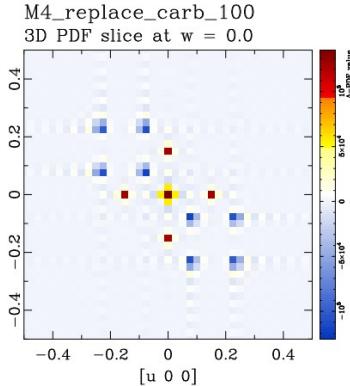
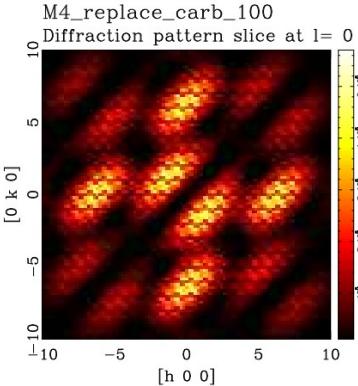
3D - Δ - PDF

„Local“ structure Patterson

Difference: Complete – Average

Positive and Negative peaks
Deviations of local from average structure





3D - Δ - PDF

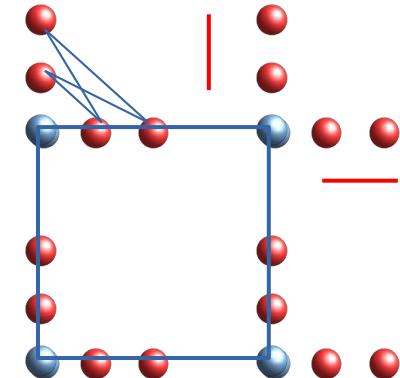
„Local“ structure Patterson

Difference: Complete - Average

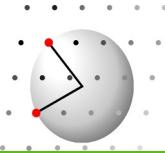
No A..B vectors
in 3D - Δ - PDF

No positive /negative
Although disordered!

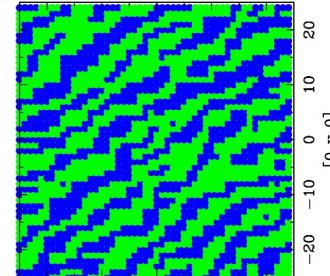
Complete structure

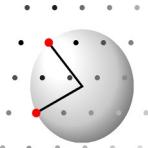


Average structure

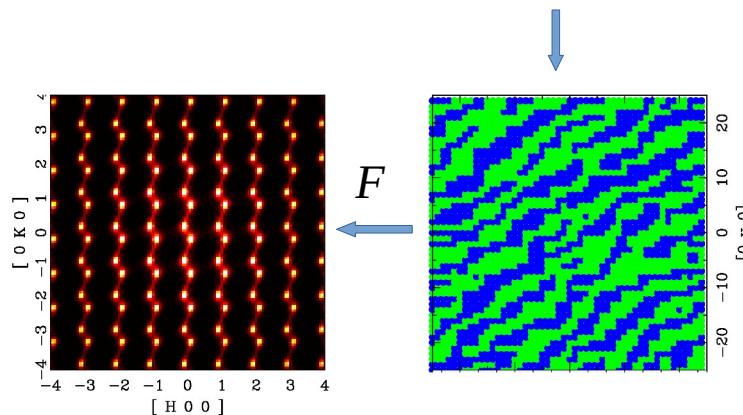


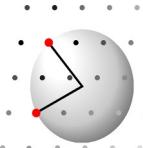
Simulation / refinement of disordered structural model



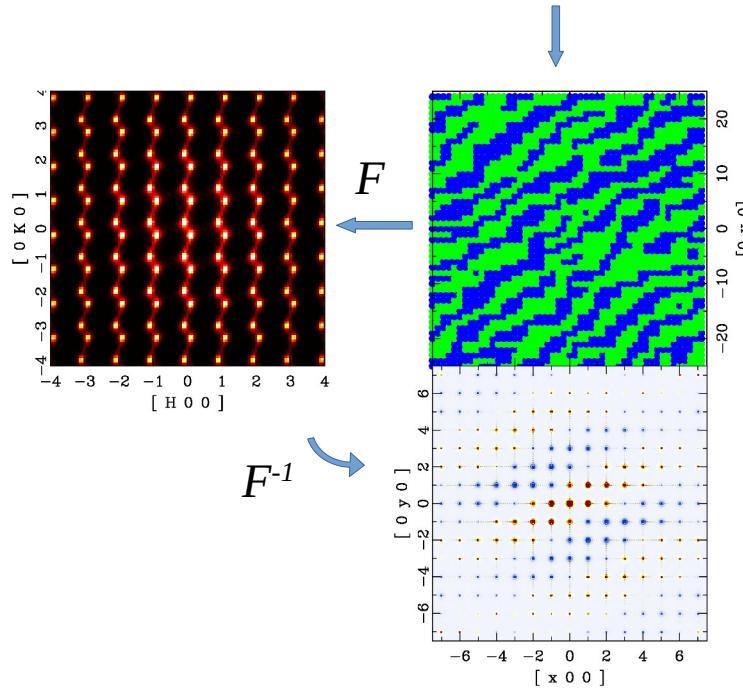


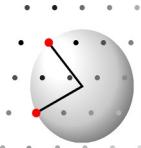
Simulation / refinement of disordered structural model



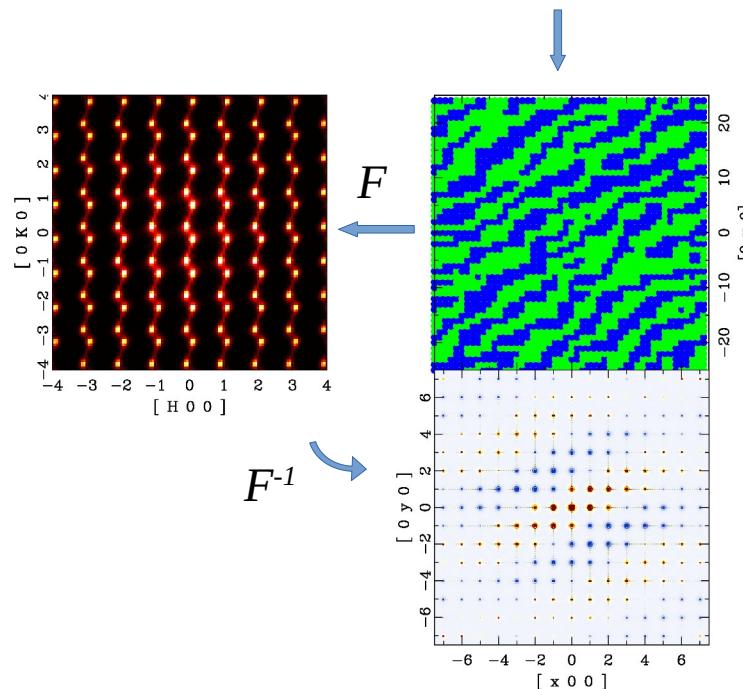


Simulation / refinement of disordered structural model





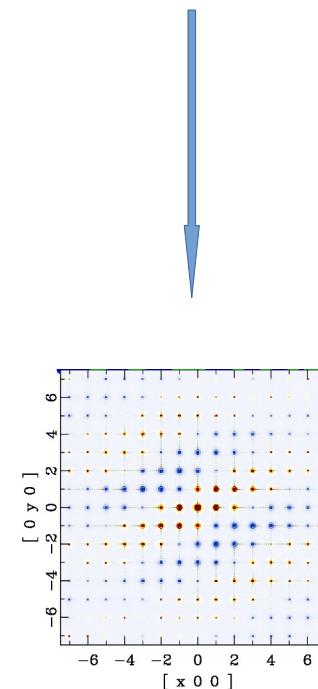
Simulation / refinement of disordered structural model



DISCUS: www.github.com/tproffen/DiffuseCode

Neder & Proffen, Oxford University Press

Simulation / refinement of correlation parameters



Yell: www.github.com/YellProgram

Simonov et al. J.Appl.Cryst. (2014), 47, 1146