Metadata needed for the full exploitation of diffuse scattering data from protein crystals

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Diffuse X-Ray Scattering to Model Protein Motions

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Problems in biology increasingly need models of protein flexibility to understand and control protein function. At the same time, as they improve, crystallographic methods are marching closer to the limits of what can be learned from Bragg data in isolation. It is thus inevitable that mainstream protein crystallography will turn to diffuse scattering to model protein motions and improve crystallographic models. The time is ripe to make it happen.

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Kathleen Lonsdale Diffuse Scattering Pioneer



- 1924, student of Lawrence Bragg
- 1928, Solved benzene structure

 Ended 60 year debate about flat aromatic ring
- 1942, Champion of diffuse X-ray scattering
- 1945, Fellow of Royal Society

 One of first two women (along with Marjory Stophonson)
 - Stephenson)
- 1956, Dame Commander of OBE
- 1966, First woman president of IUCr



$$I_D(\mathbf{q}) = \sum_{\mathbf{g},s} \left[I_0(\mathbf{q} + \mathbf{g}) + I_0(\mathbf{q} - \mathbf{g}) \right] \mathbf{q} \cdot \mathbf{v}_s(\mathbf{g})^2$$

Born & Sarginson, 1941 James, 1948

Thermal Diffuse Scattering (TDS)

$$F = \sum_{n} f_{n} e^{i\mathbf{q}\cdot\mathbf{R}_{n}}$$

$$I(t) = |F|^{2} = \sum_{n} \sum_{m} f_{n} f_{m}^{*} e^{i\mathbf{q}\cdot(\mathbf{R}_{n}-\mathbf{R}_{m})}$$

$$I = \sum_{n} \sum_{m} \left[\left(\left\langle \left| f_{n} \right|^{2} \right\rangle - \left| \left\langle f_{n} \right\rangle \right|^{2} \right) \delta_{nm} + \left| \left\langle f_{n} \right\rangle \right|^{2} \right] e^{i\mathbf{q}\cdot(\mathbf{R}_{n}-\mathbf{R}_{m})}$$

$$I = I_{D} + I_{B}$$

$$I_{B} = \left| \left\langle f \right\rangle \right|^{2} \sum_{n} \sum_{m} e^{i\mathbf{q}\cdot(\mathbf{R}_{n}-\mathbf{R}_{m})}$$

$$I_{D}(\mathbf{q}) = N \left(\left\langle \left| f_{n} - \left\langle f_{n} \right\rangle \right|^{2} \right\rangle_{n} \right) \propto N \int d\mathbf{x} e^{i\mathbf{q}\cdot\mathbf{x}} \int d\mathbf{x}' \left\langle \Delta \rho_{n}\left(\mathbf{x}'\right) \Delta \rho_{n}\left(\mathbf{x}'+\mathbf{x}\right) \right\rangle_{n}$$

Sample using molecular dynamics simulations

Guinier 1956, 1963



Wall, 1996 (Ph.D. Thesis) Wall, Ealick, and Gruner, *PNAS* 1997 Wall, *Methods Mol Biol* 2009 http://github.com/mewall/lunus



Wall, Clarage & Phillips, Structure 1997

$I_D(\vec{q}) = (1 - e^{-\vec{q} \cdot \mathbf{U} \cdot \vec{q}}) e^{-\vec{q} \cdot \mathbf{U} \cdot \vec{q}} I_0(\vec{q}) * \Gamma_{\mathbf{G}}(\vec{q})$

$$\Gamma_{\mathbf{G}}(\vec{q}) = \frac{8\pi \operatorname{det} ||\mathbf{G}||}{\left[1 + |\mathbf{G}\vec{q}|^2\right]^2}$$

Liquid-like motions (Caspar et al)

Snase: γ = 10 Å; σ = 0.36 Å Calmodulin: γ = 4.8 Å; σ = 0.38 Å

 $\Gamma_{\mathbf{G}}(\vec{q}) = \frac{4\pi \operatorname{det} ||\mathbf{G}||}{1 + |\mathbf{G}\vec{q}|^2}$

Acoustic modes

 $\begin{array}{lll} \mbox{Correlations:} & \mbox{Displacements:} \\ \gamma_1 = 50 \mbox{ \AA} & \sigma_1 = 0.0 \mbox{ \AA} \\ \gamma_2 = 135 \mbox{ \AA} & \sigma_2 = 0.4 \mbox{ \AA} \\ \gamma_3 = 85 \mbox{ \AA} & \sigma_3 = 0.0 \mbox{ \AA} \end{array}$



$$L = \frac{1}{2} \sum m_i \dot{\mathbf{R}}_i^2 - U(\mathbf{R}_1, ..., \mathbf{R}_N)$$
$$\frac{d}{dt} \left(\frac{\partial L}{\partial \dot{\mathbf{R}}_k}\right) = -\frac{\partial L}{\partial \mathbf{R}_k}$$
$$\mathbf{f}_i = m_i \ddot{\mathbf{R}}_i = -\frac{\partial U}{\partial \mathbf{R}_i}$$



Clarage et al, PNAS 1995

Meinhold & Smith, Biophys J 2005



Wall et al, PNAS 2014



14,800 independent data points



MD Model 22 Aug 2015

Data

Michael Wall, LA-UR-15-23866

Predicting X-ray diffuse scattering from translationlibration-screw structural ensembles

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22 Aug 2015





Image processing and scripts in Lunus (http://github.com/mewall/lunus)

DIALS indexing methods from Nicholas Sauter and Aaron Brewster, LBNL

Experiments of James Fraser and Others

Raw Images

- No compression
- Simple layout of data
- Shared conventions for *r,c->x,y*
- Human-readable
 header
 - *i.e.* SMV or the like



Beam Metadata

- Beam line
- Wavelength
 - Spectrum
- Polarization
 Evidence
- Beam center
 Evidence



Michael Wall, LA-UR-15-23866

Detector Metadata

- Detector model
- Detailed operating mode
- Relation of ADU to X-ray counts
- Distance
 - Evidence
- Detector face rotation
 Evidence
- Pixel size



Crystal Metadata

- Space group
 - Evidence
- Unit cell
 - Evidence
- Chemical contents



- Light microscopy image for each exposure
 - Distinguish crystal from other scattering sources
 - Tomography model of specimen

Integrated Diffuse Data Deposition

- D(hkl)
 - Fractional hkl possible
- Image processing parameters
 - Beam polarization
 - Solid-angle normalization
 - Bragg peak filtering
- Scale factors

- Frame-by-frame indexing information
- Flexible with respect to future needs for combined integration of Bragg and diffuse data

Diffuse Scattering Model Deposition

- Dynamical parameter values
 - Displacement correlations
 - Displacement amplitudes
 - Dispersion relation
- MD trajectories (large!)
- Calculated diffuse intensities





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 - Nicholas Sauter
 - Aaron Brewster



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Further Reading

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