SAXS review

- SAXS review
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Reading assignment: Chapter 5.2

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Homework Assignment #04: Chapter 4: 2, 4, 6, 7, 10 due Monday, October 24, 2016

The SAXS scattered intensity from a dilute solution depends on the single particle form factor, $\mathcal{F}(\vec{Q})$, the volume of the particle, V_p , and the density difference from the solvent, $\Delta \rho = (\rho_{sl,p} - \rho_{sl,0})$

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p = 0 p = 10%p = 20%



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Inter-particle interactions

Many interesting problems fall outside the dilute limit.

In these cases, the SAXS modeling must include not only the particle form factor but an additional structure factor, S(Q)

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The book has an example of this and we will look at a couple of others from recent journal articles

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• SAXS of irradiated Zn nanoparticles

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The book has an example of this and we will look at a couple of others from recent journal articles

- SAXS of irradiated Zn nanoparticles
- Nucleation and growth of & glycine crystals

Zn nanoparticles formed in SiO₂ by ion implantation are irradiated with high energy Xe^{+14} ions.

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SAXS is measured using 18 keV x-rays both parallel and perpendicular to the direction of Xe^{+14} irradiation.

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Expt. geometry

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Expt. geometry

Unirradiated

Zn nanoparticles formed in SiO₂ by ion implantation are irradiated with high energy Xe^{+14} ions.

SAXS is measured using 18 keV x-rays both parallel and perpendicular to the direction of Xe^{+14} irradiation.





SAXS intensity for \parallel and \perp x-ray incidence

"Shape elongation of embedded Zn nanoparticles induced by swift heavy ion irradiation: A SAXS study", H. Amekura, K. Kono, N. Okubo, and N. Ishikawa, *Phys. Status Solidi B* **252**, 165-169 (2015).



SAXS intensity for \parallel and \perp x-ray incidence

Interparticle distance as a function of irradiation fluence

"Shape elongation of embedded Zn nanoparticles induced by swift heavy ion irradiation: A SAXS study", H. Amekura, K. Kono, N. Okubo, and N. Ishikawa, *Phys. Status Solidi B* **252**, 165-169 (2015).



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growth of interparticle spacing is due to dissolution and re-agglomeration with fluence

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Nucleation & growth of glycine

Can SAXS help us understand the nucleation and growth of a simple molecule which is the prototype for pharmaceutical compounds?



initial studies at 12 keV show change but no crystallization

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Glycine nucleation



change to 25 keV x-rays study neutral (top) and acidic (bottom) solutions



Glycine nucleation



change to 25 keV x-rays study neutral (top) and acidic (bottom) solutions



Glycine R_g



in aqueous solution, R_g implies dimerization and increases due to aggregation until crystallization

in acidic solution, *Rg* remains small and implies that no dimerization or aggregation occurs before nucleation

"Relationship between Self-Association of Glycine Molecules in Supersaturated Solution and Solid State Outcome", D. Erdemir et al. *Phys. Rev. Lett.* **99**, 115702 (2007)

SAXS of biological molecules is an excellent way of getting some information about the molecules as they exist in solution.

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Obtaining information about R_g and the Porod region, combined with modeling and the known crystallographic structures can give a more complete picture of how these molecules function.

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Matthew, Mirza & Menhart, "liquid-chromatography-coupled SAXS for accurate sizing of aggregating proteins," *J. Synchrotron Rad.* **11**, 314-318 (2004) developed a technique which is now being used routinely in biological SAXS, called Size Exclusion Chromatography SAXS.





2m SAXS camera, 1.03Å(12 keV) x-rays were used



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samples of (1) cytochrome c, (2) plasminogen, (3) mixture of cytochrome c bovine serum albumin, and blue dextran

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SEC-SAXS experimental setup



Matthew, Mirza & Menhart, "liquid-chromatography-coupled SAXS for accurate sizing of aggregating proteins," J. Synchrotron Rad. 11, 314-318 (2004).

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Cytochrome c



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Cytochrome c - Guinier plots



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Plasminogen



Matthew, Mirza & Menhart, "liquid-chromatography-coupled SAXS for accurate sizing of aggregating proteins," J. Synchrotron Rad. 11, 314-318 (2004).

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PHYS 570 - Fall 2016

October 12, 2016 17 / 25

Three component mixture



Matthew, Mirza & Menhart, "liquid-chromatography-coupled SAXS for accurate sizing of aggregating proteins," J. Synchrotron Rad. 11, 314-318 (2004).

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October 12, 2016 18 / 25

$$\vec{R}_n = n_1 \vec{a}_1 + n_2 \vec{a}_2$$



 $\vec{R}_n = n_1 \vec{a}_1 + n_2 \vec{a}_2$

primitive



$$\vec{R}_n = n_1 \vec{a}_1 + n_2 \vec{a}_2$$

primitive

non-primitive



$$\vec{R}_n = n_1 \vec{a}_1 + n_2 \vec{a}_2$$

primitive

non-primitive

non-conventional

More about lattice vectors



sometimes conventional axes...

More about lattice vectors





sometimes conventional axes...

... are not primitive

Miller indices



planes designated (hk), intercept the unit cell axes at

$$\frac{a_1}{h}, \quad \frac{a_2}{k}$$

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Miller indices

(20) a₂ planes designated (hk), intercept the unit cell axes at

 $\frac{a_1}{h}, \quad \frac{a_2}{k}$

for a lattice with orthogonal unit vectors

$$\frac{1}{d_{hk}^2} = \frac{h^2}{a_1^2} + \frac{k^2}{a_2^2}$$

Reciprocal lattice



Reciprocal lattice



$$ec{a}_1^*=rac{2\pi}{V_c}ec{a}_2 imesec{a}_3 \qquad ec{a}_2^*=rac{2\pi}{V_c}ec{a}_3 imesec{a}_1 \qquad ec{a}_3^*=rac{2\pi}{V_c}ec{a}_1 imesec{a}_2$$

Reciprocal lattice



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convoluting the lattice and basis function we write

(

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$$\mathcal{L}(x) = \sum_n \delta(x - na)$$

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$$F^{crystal}(\vec{Q}) = \sum_{l}^{N} f_{l}(\vec{Q}) e^{i\vec{Q}\cdot\vec{r}_{l}}$$

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Since $F^{crystal}(\vec{Q})$ is simply the Fourier Transform of the crystal function, $C(x) = \mathcal{L}(x) \star \mathcal{B}(x)$, it must be the product of the Fourier Transforms of $\mathcal{L}(x)$ and $\mathcal{B}(x)$.

$$F^{crystal}(\vec{Q}) = \sum_{l}^{N} f_{l}(\vec{Q}) e^{i\vec{Q}\cdot\vec{r}_{l}} = \sum_{\vec{R}_{n}+\vec{r}_{j}}^{N} f_{j}(\vec{Q}) e^{i\vec{Q}\cdot(\vec{R}_{n}+\vec{r}_{j})}$$
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$$\vec{Q} \cdot \vec{R}_n = 2\pi m, \quad m = \text{integer}$$

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 $ec{G}_{hkl}=hec{a}_1^*+kec{a}_2^*+lec{a}_3^*$

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$$\vec{Q} \cdot \vec{R}_n = 2\pi m, \quad m = \text{integer}$$

$$\vec{G}_{hkl} = h\vec{a}_1^* + k\vec{a}_2^* + l\vec{a}_3^*$$

$$\vec{G}_{hkl} \cdot \vec{R}_n = (n_1\vec{a}_1 + n_2\vec{a}_2 + n_3\vec{a}_3) \cdot (h\vec{a}_1^* + k\vec{a}_2^* + l\vec{a}_3^*)$$

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$$= 2\pi (hn_{1} + kn_{2} + ln_{3}) = 2\pi m$$

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$$= 2\pi (hn_{1} + kn_{2} + ln_{3}) = 2\pi m$$

$$\therefore \quad \vec{Q} = \vec{G}_{hkl}$$



$$\vec{a}_1 = \frac{a}{2}(\hat{y} + \hat{z}),$$



$$\vec{a}_1 = rac{a}{2}(\hat{y} + \hat{z}), \quad \vec{a}_2 = rac{a}{2}(\hat{z} + \hat{x}),$$



$$ec{a}_1 = rac{a}{2}(\hat{y} + \hat{z}), \quad ec{a}_2 = rac{a}{2}(\hat{z} + \hat{x}), \quad ec{a}_3 = rac{a}{2}(\hat{x} + \hat{y})$$



The primitive lattice vectors of the face-centered cubic lattice are

$$ec{a}_1 = rac{a}{2}(\hat{y} + \hat{z}), \quad ec{a}_2 = rac{a}{2}(\hat{z} + \hat{x}), \quad ec{a}_3 = rac{a}{2}(\hat{x} + \hat{y})$$

The volume of the unit cell is



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The volume of the unit cell is

$$v_c = \vec{a}_1 \cdot \vec{a}_2 \times \vec{a}_3 = \vec{a}_1 \cdot \frac{a^2}{4} (\hat{y} + \hat{z} - \hat{x})$$

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 $= \frac{4\pi}{a} \left(\frac{\hat{y}}{2} + \frac{\hat{z}}{2} - \frac{\hat{x}}{2}\right)$

The primitive lattice vectors of the face-centered cubic lattice are

$$\vec{a}_1 = \frac{a}{2}(\hat{y} + \hat{z}), \quad \vec{a}_2 = \frac{a}{2}(\hat{z} + \hat{x}), \quad \vec{a}_3 = \frac{a}{2}(\hat{x} + \hat{y})$$



The volume of the unit cell is

$$v_{c} = \vec{a}_{1} \cdot \vec{a}_{2} \times \vec{a}_{3} = \vec{a}_{1} \cdot \frac{a^{2}}{4} (\hat{y} + \hat{z} - \hat{x}) = \frac{a^{3}}{4}$$

$$\vec{a}_{1}^{*} = \frac{2\pi}{v_{c}} \vec{a}_{2} \times \vec{a}_{3} = \frac{2\pi}{v_{c}} \frac{a^{2}}{4} (\hat{y} + \hat{z} - \hat{x})$$

$$= \frac{4\pi}{a} \left(\frac{\hat{y}}{2} + \frac{\hat{z}}{2} - \frac{\hat{x}}{2} \right)$$

$$\vec{a}_{2}^{*} = \frac{4\pi}{a} \left(\frac{\hat{z}}{2} + \frac{\hat{x}}{2} - \frac{\hat{y}}{2} \right)$$

$$\vec{a}_{3}^{*} = \frac{4\pi}{a} \left(\frac{\hat{x}}{2} + \frac{\hat{y}}{2} - \frac{\hat{z}}{2} \right)$$

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The primitive lattice vectors of the face-centered cubic lattice are

$$\vec{a}_1 = \frac{a}{2}(\hat{y} + \hat{z}), \quad \vec{a}_2 = \frac{a}{2}(\hat{z} + \hat{x}), \quad \vec{a}_3 = \frac{a}{2}(\hat{x} + \hat{y})$$



which is a body-centered cubic lattice

The volume of the unit cell is

$$\begin{aligned} v_c &= \vec{a}_1 \cdot \vec{a}_2 \times \vec{a}_3 = \vec{a}_1 \cdot \frac{a^2}{4} \left(\hat{y} + \hat{z} - \hat{x} \right) = \frac{a^3}{4} \\ \vec{a}_1^* &= \frac{2\pi}{v_c} \vec{a}_2 \times \vec{a}_3 = \frac{2\pi}{v_c} \frac{a^2}{4} \left(\hat{y} + \hat{z} - \hat{x} \right) \\ &= \frac{4\pi}{a} \left(\frac{\hat{y}}{2} + \frac{\hat{z}}{2} - \frac{\hat{x}}{2} \right) \\ \vec{a}_2^* &= \frac{4\pi}{a} \left(\frac{\hat{z}}{2} + \frac{\hat{x}}{2} - \frac{\hat{y}}{2} \right) \\ \vec{a}_3^* &= \frac{4\pi}{a} \left(\frac{\hat{x}}{2} + \frac{\hat{y}}{2} - \frac{\hat{z}}{2} \right) \end{aligned}$$

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