Multiwavelength anomalous diffraction at high x-ray intensity

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Center for Free-Electron Laser Science







Overview

- > MAD for phase problem
- > XFEL for femtosecond x-ray nanocrystallography
- Possible to use MAD with XFEL?
- Electronic damage to heavy atoms
- > MAD with XFEL: generalized Karle-Hendrickson equation
- > Conclusion



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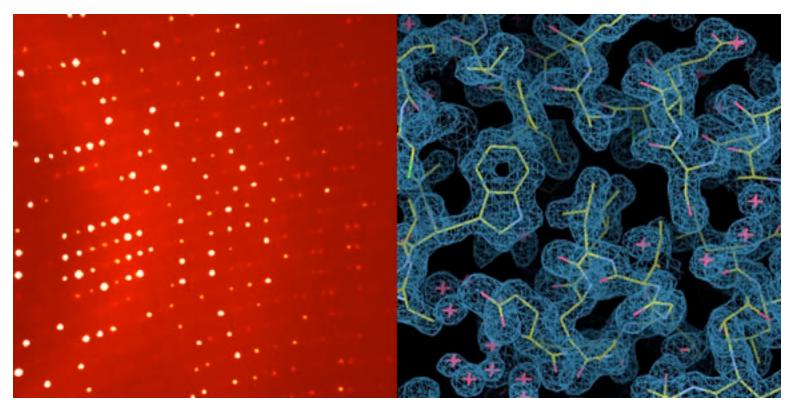


Henry Chapman





Structural determination of macromolecules



Picture taken from http://www.sandiego.edu/cas/chemistry/newsletter/

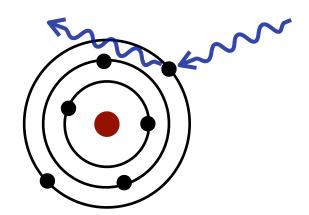




X-ray scattering

Elastic x-ray scattering form factor

$$f^{0}(\mathbf{Q}) = \int d^{3}r \ \rho(\mathbf{r}) \ e^{i\mathbf{Q}\cdot\mathbf{r}}$$
$$\frac{d\sigma(\mathbf{Q})}{d\Omega} = \left(\frac{d\sigma}{d\Omega}\right)_{T} \left|f^{0}(\mathbf{Q})\right|^{2}$$



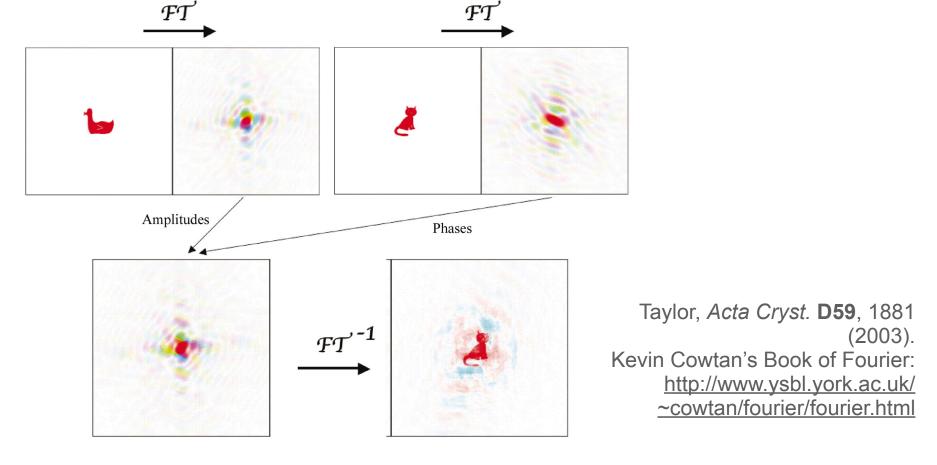
> **Phase problem**: a fundamental obstacle in constructing an electronic density map from x-ray diffraction

$$f^{0}(\mathbf{Q}) = \int d^{3}r \ \rho(\mathbf{r}) \ e^{i\mathbf{Q}\cdot\mathbf{r}} = |f^{0}(\mathbf{Q})| \ e^{i\phi^{0}(\mathbf{Q})}$$





Why are phases important?



cat's phases + duck's amplitudes → reconstruction of the cat





Phasing method

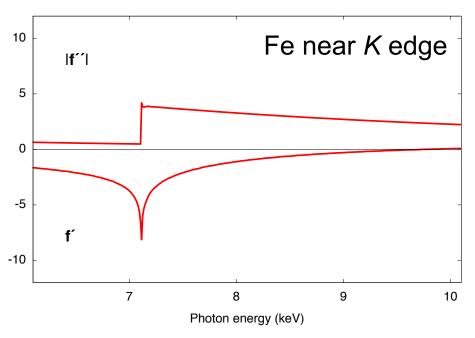
- Many phasing methods have been proposed and applied.
 - Molecular replacement (MR)
 - Single / multiple isomorphous replacement (SIR / MIR)
 - SIR / MIR with anomalous scattering (SIRAS / MIRAS)
 - Multi-wavelength anomalous diffraction (MAD)
 - Single-wavelength anomalous diffraction (SAD)
- molecular replacement: phases borrowed from a similar molecule, but structural determination severely biased
- > ab initio phasing: without knowing a similar structure, mainly employing anomalous scattering of heavy atoms





Multiwavelength Anomalous Diffraction

> Resonant elastic x-ray scattering: $f(\mathbf{Q},\omega) = f^0(\mathbf{Q}) + f'(\omega) + if''(\omega)$



MAD phasing: The Karle-Hendrickson equation provides a simple way for phasing from the contrast at two or more wavelengths.

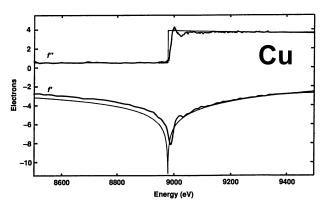
Karle, *Int. J. Quant. Chem.* **18**, Suppl. S7, 357 (1980). Hendrickson, *Trans. Am. Crystalgr. Assoc.* **21**, 11 (1985).





MAD with synchrotron radiation

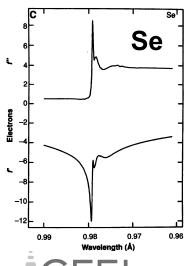
MAD has been a well-established phasing method with synchrotron radiation since 80's.

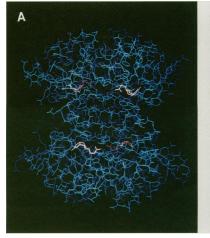


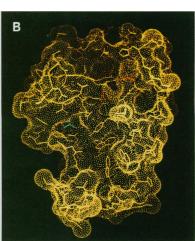
Heptahydrido *bis*(diiopropylphenyl)phosphine **Re** LIII-edge

Arndt et al., Nature 298, 835 (1982).

Cucumber basic blue protein Guss et al., Science **241**, 806 (1988).







A) Streptavidin Hendrickson *et al.*, *PNAS* **86**, 2190 (1989).

B) Ribonuclease H Yang *et al.*, *Science* **249**, 1398 (1990).

Picture taken from Hendrickson, *Science* **254**, 51 (1991).



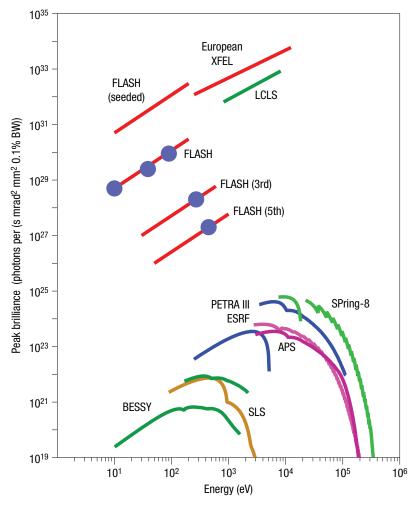


What is XFEL?

- > XFEL: X-ray Free-Electron Laser
- > Ultraintense
 - synchrotron: at most one photon absorbed per pulse
 - XFEL: many photons absorbed per pulse
 - fluence: ~10¹³ photons per μm² per pulse
 - peak intensity: ~10¹⁸ W/cm²

> Ultrafast

pulse duration: femtoseconds or sub-fs



Ackermann et al., Nature Photon. 1, 336 (2007).





Where are XFELs?

- > FLASH at DESY, Germany (2004)
- > LCLS at SLAC, USA (2009)
- SACLA at RIKEN Harima, Japan (2011)
- > PAL XFEL at Pohang, Korea (2015)
- > European XFEL, Germany (2015)











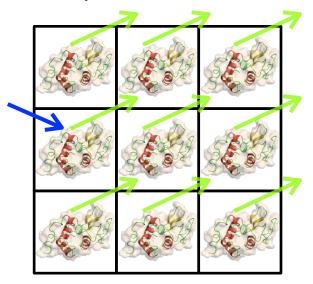


Why need for *ultraintense*

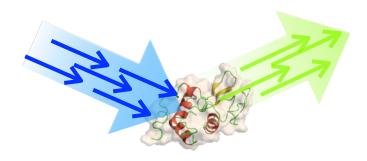
Carbon at synchrotron radiation:
 12 keV, 10⁶ photons
 on 10µm × 10µm



Signal amplification



~108 molecules in a µm-sized crystal

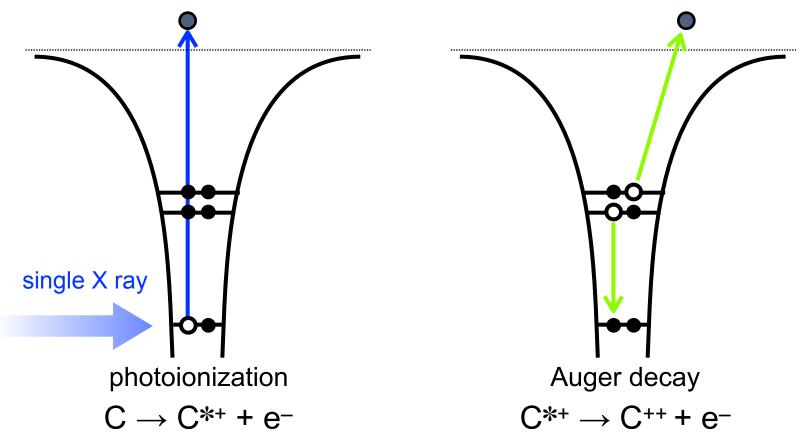


high x-ray fluence from XFEL (×108~10¹⁰ more than synchrotron radiation)





Photoabsorption by X-rays

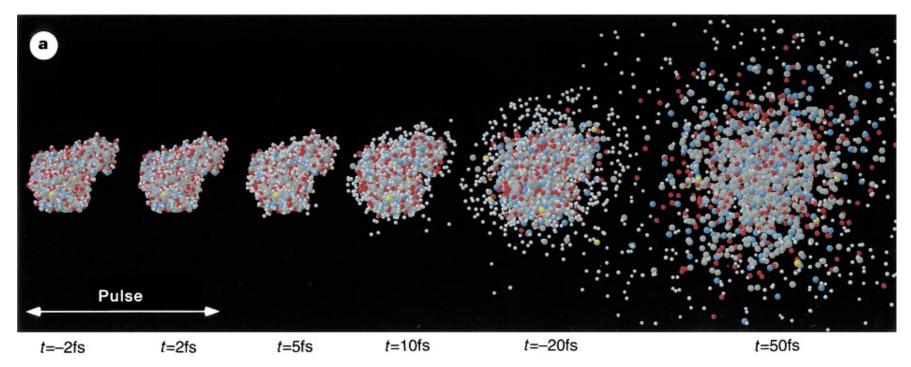


- > Typically absorption cross section is larger than scattering cross section.
- XFEL induces multiphoton multiple ionization dynamics.





Radiation damage (Coulomb explosion)



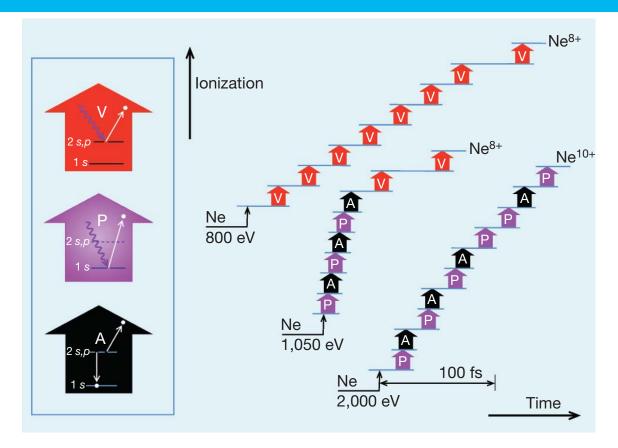
Neutze et al., Nature 406, 752 (2000)

Diffraction-before-destruction: needs ultrafast pulses (~femtosecond)





Electronic damage during XFEL pulses



Young et al., Nature **466**, 56 (2010).

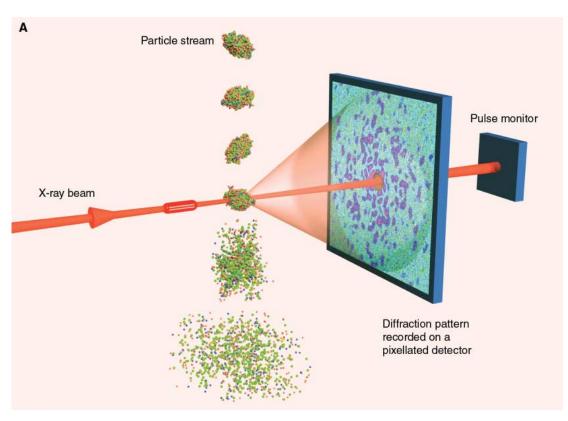
- Multiphoton multiple ionization via a sequence of one-photon ionizations
- Diffraction-during-ionization: unavoidable electronic damage (~attosec.)





Femtosecond X-ray nanocrystallography

- Growing high-quality crystals is one of major bottlenecks in x-ray crystallography.
- Ultraintense and ultrafast pulses from XFEL
- Enough signals from nanosized crystals and single molecules
- Single-shot molecular imaging: revolutionary impact on structural biology



Gaffney & Chapman, Science 316, 1444 (2007).





New theory for new experiment

Can we use the MAD phasing with XFEL?

phase problem → MAD
growing high-quality crystals → XFEL

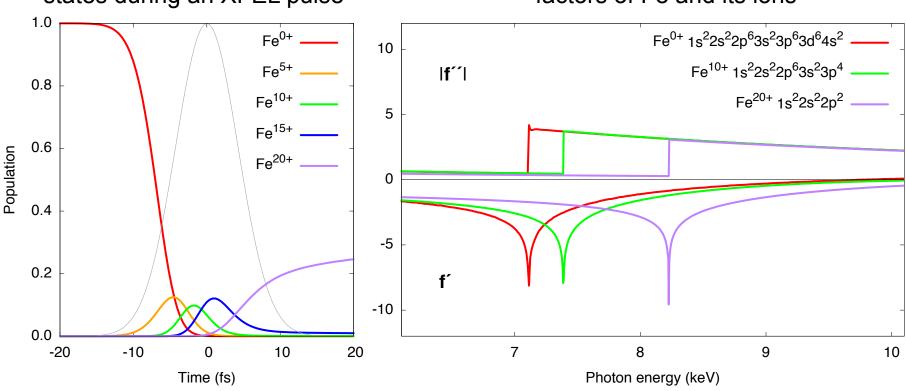




Electronic damage to heavy atoms

Population dynamics of Fe charge states during an XFEL pulse

Dispersion corrections of atomic form factors of Fe and its ions



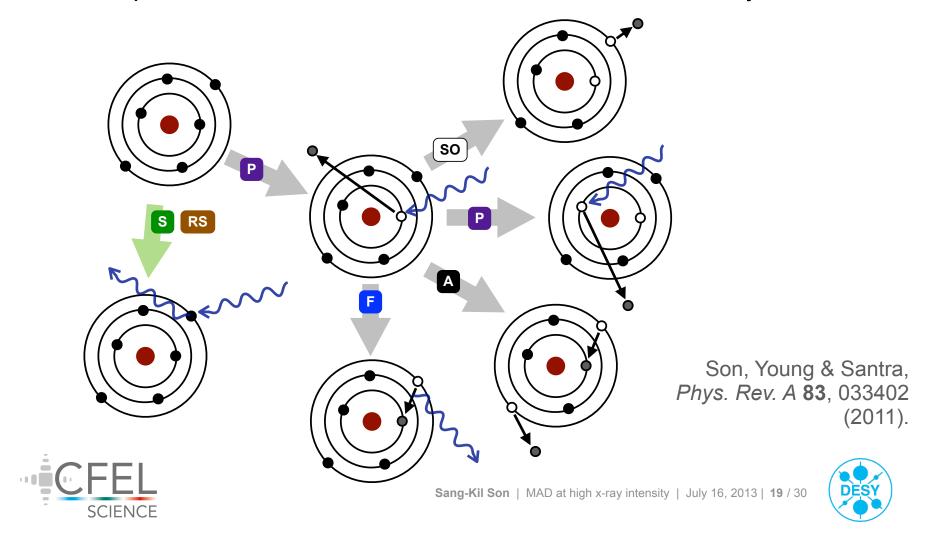
(8 keV, 5×10^{12} photons/ μ m², 10 fs FWHM)



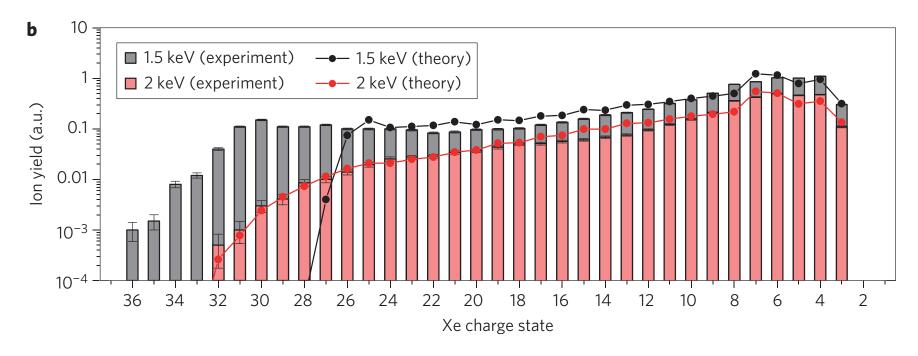


XATOM: x-ray and atomic physics toolkit

- X-ray-induced atomic processes for any given element and configuration
- Rate equation model to simulate ionization and relaxation dynamics



Xe at LCLS



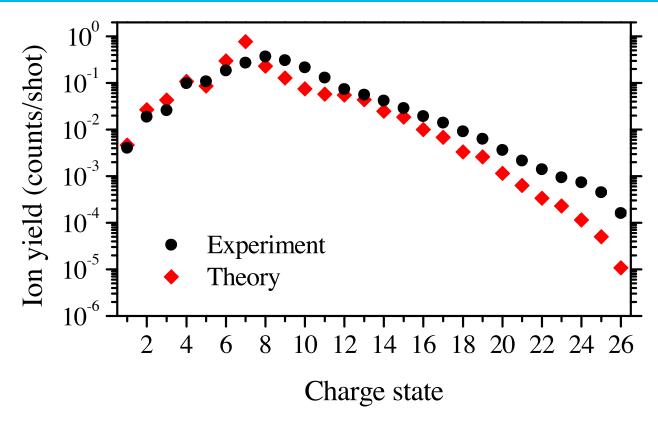
- Charge state distribution of Xe measured at LCLS
- > At 2 keV: good agreement between experiment and theory
- > At 1.5 eV: unprecedented high charge states (up to Xe³⁶⁺) in experiment

Rudek, Son et al., Nature Photon. 6, 858 (2012).





Xe at SACLA



> At 5.5 keV: deep inner-shell (*L*-shell) ionization dynamics good agreement between experiment and theory

Fukuzawa, Son et al., Phys. Rev. Lett. 110, 173005 (2013).





Prior speculations regarding MAD at XFEL

- Unavoidable electronic damage, especially to heavy atoms
- Dramatic change of anomalous scattering for high charge states
- Stochastic electronic damage to heavy atoms would destroy coherent scattering signals in nanocrystals
- MAD would not be an applicable route for phasing at XFEL...?

- We demonstrate the existence of a Karle-Hendrickson-type equation in the high-intensity regime.
- We show that MAD not only works, but also the extensive electronic rearrangements at high x-ray intensity provide a new path to phasing.





Karle-Hendrickson equation

Karle-Hendrickson eq. represents a set of equations at different ω:

$$\begin{aligned} |F_T(\mathbf{Q},\omega)|^2 &= |F_T^0(\mathbf{Q})|^2 + |F_A^0(\mathbf{Q})|^2 a(\mathbf{Q},\omega) \\ &+ |F_T^0(\mathbf{Q})| |F_A^0(\mathbf{Q})| b(\mathbf{Q},\omega) \cos \Delta \phi^0(\mathbf{Q}) \\ &+ |F_T^0(\mathbf{Q})| |F_A^0(\mathbf{Q})| c(\mathbf{Q},\omega) \sin \Delta \phi^0(\mathbf{Q}) \end{aligned}$$

> MAD coefficients (determined theoretically or experimentally):

$$\frac{a(\mathbf{Q},\omega)}{\left\{f_A^0(\mathbf{Q})\right\}^2}, \quad b(\mathbf{Q},\omega) = \frac{2f_A'(\omega)}{f_A^0(\mathbf{Q})}, \quad c(\mathbf{Q},\omega) = \frac{2f_A''(\omega)}{f_A^0(\mathbf{Q})}$$

- > 3 unknowns at every **Q**: $|F_T^0(\mathbf{Q})|$, $|F_A^0(\mathbf{Q})|$, $\Delta \phi^0(\mathbf{Q}) \left[= \phi_T^0(\mathbf{Q}) \phi_A^0(\mathbf{Q}) \right]$
- These 3 unknowns are algebraically solved with 3 measurements.

Karle, Int. J. Quant. Chem. Quant. Bio. Symp. 7, 357 (1980) Hendrickson, Trans. Am. Crystalgr. Assoc. 21, 11 (1985)





Scattering intensity including elec. damage

$$\frac{dI(\mathbf{Q},\omega)}{d\Omega} = \mathcal{F}C(\Omega) \int_{-\infty}^{\infty} dt \, g(t) \sum_{I} P_{I}(t) \left| F_{P}^{0}(\mathbf{Q}) + \sum_{j=1}^{N_{H}} f_{I_{j}}(\mathbf{Q},\omega) e^{i\mathbf{Q} \cdot \mathbf{R}_{j}} \right|^{2}$$

$$I = (I_1, I_2, \dots I_{N_H}), \quad P_I(t) = \prod_{j=1}^{N_H} P_{I_j}(t)$$
$$f_{I_j}(\mathbf{Q}, \omega) = f_{I_j}^0(\mathbf{Q}) + f'_{I_j}(\omega) + i f''_{I_j}(\omega)$$

- > All changes among N_H heavy atoms are included.
- > P: protein, H: heavy atoms; only heavy atoms scatter anomalously and undergo damage dynamics during an x-ray pulse.
- Heavy atoms are ionized independently.
- Only one species of heavy atoms is considered.





Generalized Karle-Hendrickson equation

$$\frac{dI(\mathbf{Q},\omega)}{d\Omega} = \mathcal{F}C(\Omega) \Big[|F_P^0(\mathbf{Q})|^2 + |F_H^0(\mathbf{Q})|^2 \tilde{a}(\mathbf{Q},\omega) \\
+ |F_P^0(\mathbf{Q})| |F_H^0(\mathbf{Q})| b(\mathbf{Q},\omega) \cos \Delta \phi^0(\mathbf{Q}) \\
+ |F_P^0(\mathbf{Q})| |F_H^0(\mathbf{Q})| c(\mathbf{Q},\omega) \sin \Delta \phi^0(\mathbf{Q}) \\
+ N_H |f_H^0(\mathbf{Q})|^2 \{a(\mathbf{Q},\omega) - \tilde{a}(\mathbf{Q},\omega)\} \Big]$$

- > MAD coefficients: $a(\mathbf{Q}, \omega), b(\mathbf{Q}, \omega), c(\mathbf{Q}, \omega), \text{ and } \tilde{a}(\mathbf{Q}, \omega)$
 - → measured or calculated with time evolution of config. populations

e.g.)
$$ilde{a}(\omega) = rac{1}{\left\{f_H^0
ight\}^2} \int_{-\infty}^{\infty}\!\!\!\!dt\,g(t) \left|\sum_{I_H} P_{I_H}(t) f_{I_H}(\omega)\right|^2$$

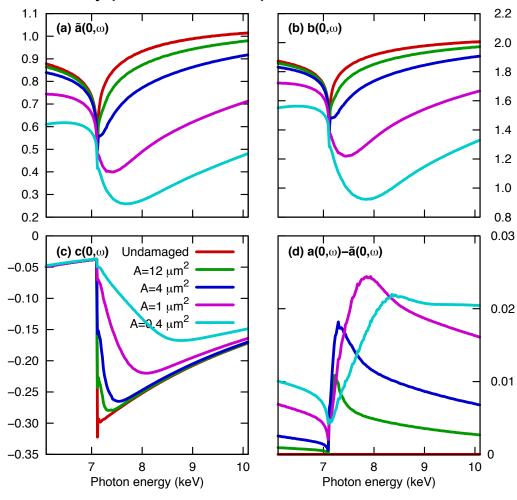
- > 3 unknowns: $|F_P^0(\mathbf{Q})|, |F_H^0(\mathbf{Q})|, \Delta \phi^0(\mathbf{Q}) [= \phi_P^0(\mathbf{Q}) \phi_H^0(\mathbf{Q})]$
 - → solvable with measurements at 3 different wavelengths.





MAD coefficients

Fe in an x-ray pulse of 2×10¹² photons and 10 fs FWHM



- dynamical behavior of heavy atoms calculated by XATOM
- bleaching effect:
 minimum deepened and
 edge broadened
 → easy to choose
 wavelengths
- MAD works: enhanced contrast at different wavelengths
- potential new phasing methods





Brand-new phasing method

- > **SIR** (single isomorphic replacement): atomic replacement in sample preparation; native vs. derivative
- > RIP (radiation-damage induced phasing): chemical rearrangement during the x-ray pulses; S—S bond vs. bond breaking
- **MAD** (multi-wavelength anomalous diffraction): $\Delta F_{\Delta\lambda}$
- > **SAD** (single-wavelength anomalous diffraction): ΔF_{\pm}

New phasing method: neither **SIR** nor **RIP**Fluences rather than wavelengths: neither **MAD** nor **SAD**





Conclusion

- MAD phasing method in extreme conditions of ionizing radiations
- Combination of ultrafast electronic dynamics at the atomic level and imaging of macromolecules by intense x-ray pulses
- Existence of a generalized Karle-Hendrickson equation for the MAD method at high x-ray intensity
- Bleaching effect on the scattering strength to be beneficial to the phasing method
- A new opportunity for solving the phase problem in femtosecond nanocrystallography with XFELs
 - → A breakthrough in structural biology





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Take-home message

XFEL goes MAD.



