

# On dose related issues in XFELs vs. ERLs

## Outline:

- Motivation
- Overview of processes involved
- “Conventional” protein damage in crystallography
- Cryoprotected X-ray microscopy

# Motivation

Biological and Environmental Research Advisory Committee (BERAC) at Department of Energy (DOE) meeting on April 30 - May 1 (2003) Washington, DC aims at addressing the following question from the Director of Office of Science

<http://www.science.doe.gov/ober/berac/synchrotron.html>

*“What characteristics of the next generation x-ray light sources (e.g., their extremely short femtosecond time scale x-ray pulses; high average or peak brightness; coherence) are most important in enabling science, from determination of physical structures to biological functions, for the biological community in the coming 10-20 years?”*

The response suggests (see <http://www.science.doe.gov/ober/berac/StructBio.pdf>):

... superiority of X-FELs over ERLs in “greater potential for **breakdown** science by biological community” (page 4). The reasoning behind this view is most succinctly recapped by the following phrase that “ERLs will be far less powerful than X-FELs due to limited number of photons per ERL pulse” (page 16).

# Recap of radiation properties

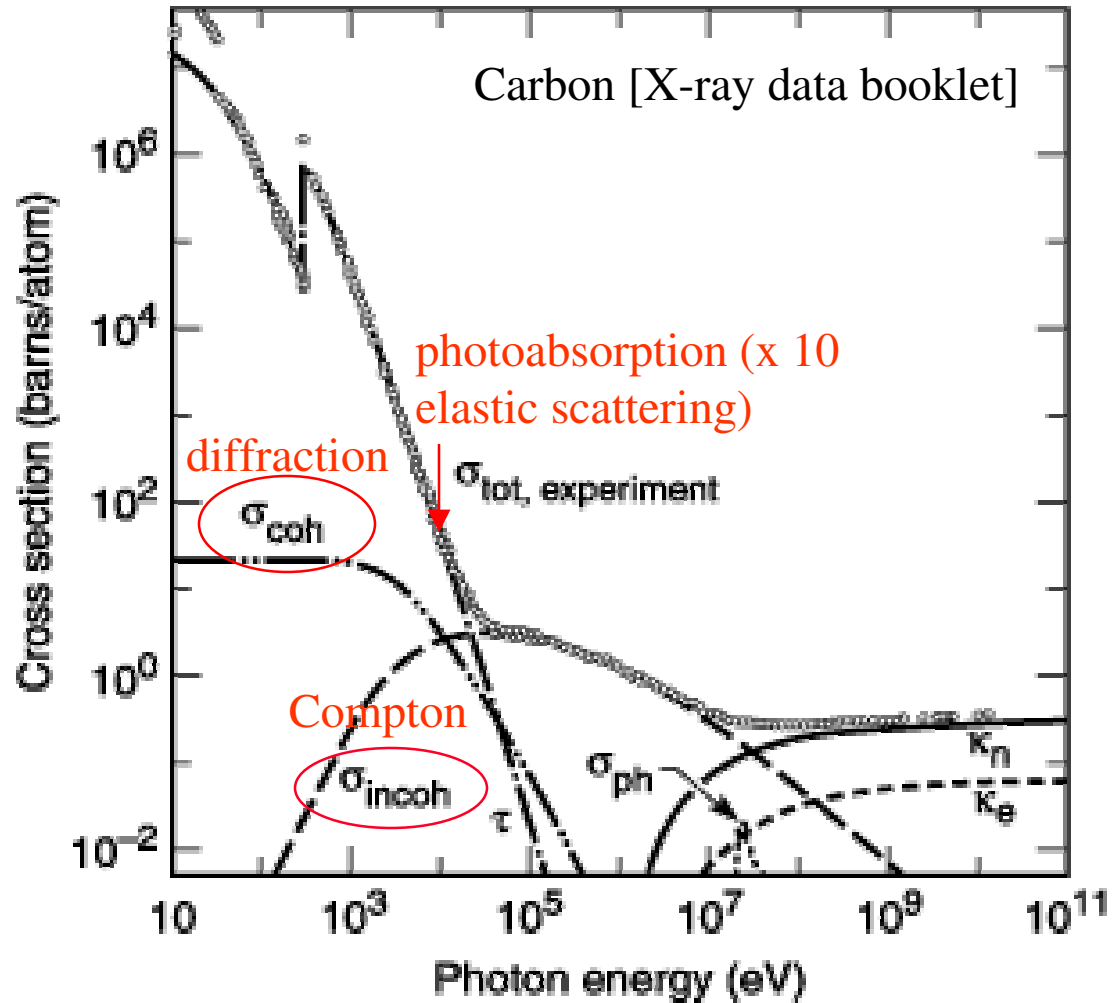
	XFEL	ERL
Photons/pulse	$10^{12}$	$10^{6-7}$
Rep. Rate [Hz]	$10^2$	$10^9$
Pulse duration [ps]	0.1	0.1 – 2
Source size [ $\mu\text{m}$ ]	30	10

# The scare of XFELs

- If focused to 10  $\mu\text{m}$  spot, the peak power density is  $10^{16}$  W/cm<sup>2</sup>
- 200 kiloton nuclear weapon where 6% of the energy is emitted in X and  $\gamma$  rays over a time period of 100 ns creates peak photon density of  $10^{17}$  W/cm<sup>2</sup> within the bomb casting
- $E \sim \sqrt{Z_0 I}$ ,  $Z_0 = 377 \Omega$ , i.e. electric field  $\sim 10^{11}$  V/m
- Coulomb field acting on an outer electron  $10 \text{ V} / 1 \text{ \AA} \sim 10^{11}$  V/m
- NOT a strong field regime:  $U_p = e^2 E^2 (\lambda^2 / 16 \pi^2) / mc^2$ , e.g. average kinetic energy of wiggling electron is only  $\sim 1 \mu\text{eV}$
- OR amplitude of the wiggling motion is  $a_w = 4U_p / eE \sim 5 \times 10^{-7}$  Angstrom

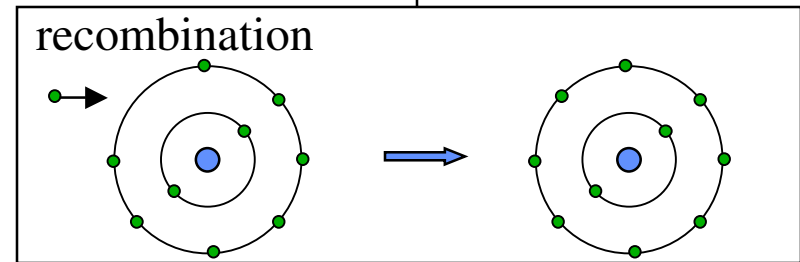
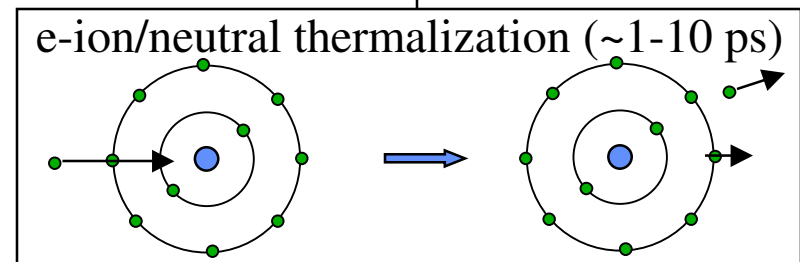
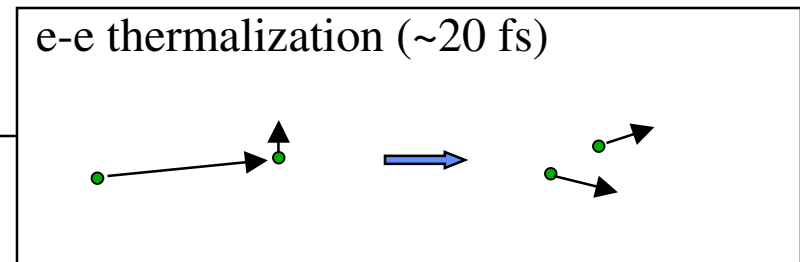
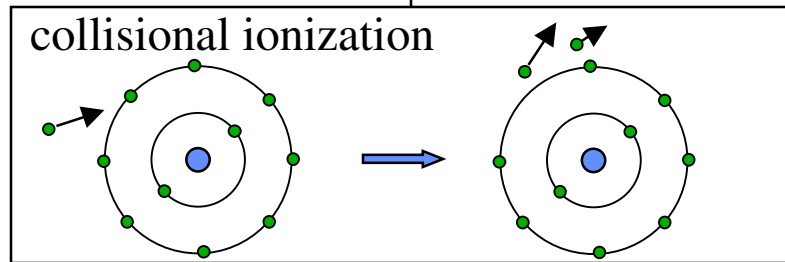
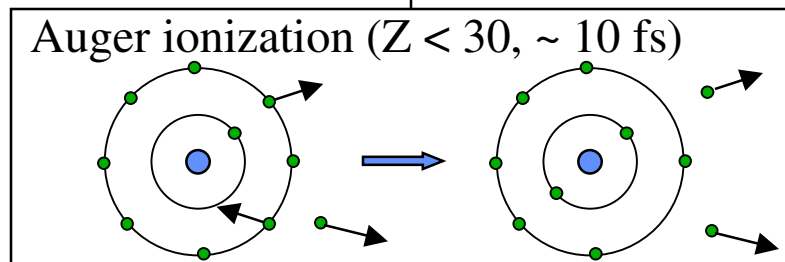
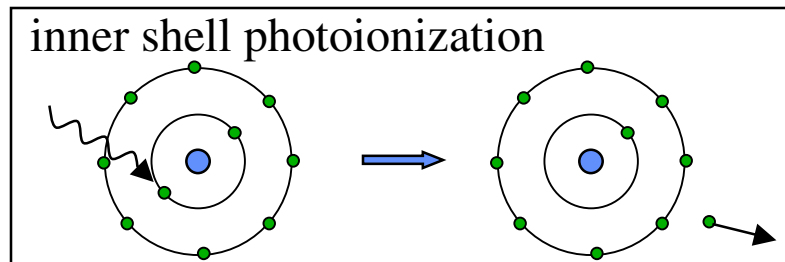
*1  $\text{\AA}$  light oscillation takes only 0.3 attosecond*

# Processes involved

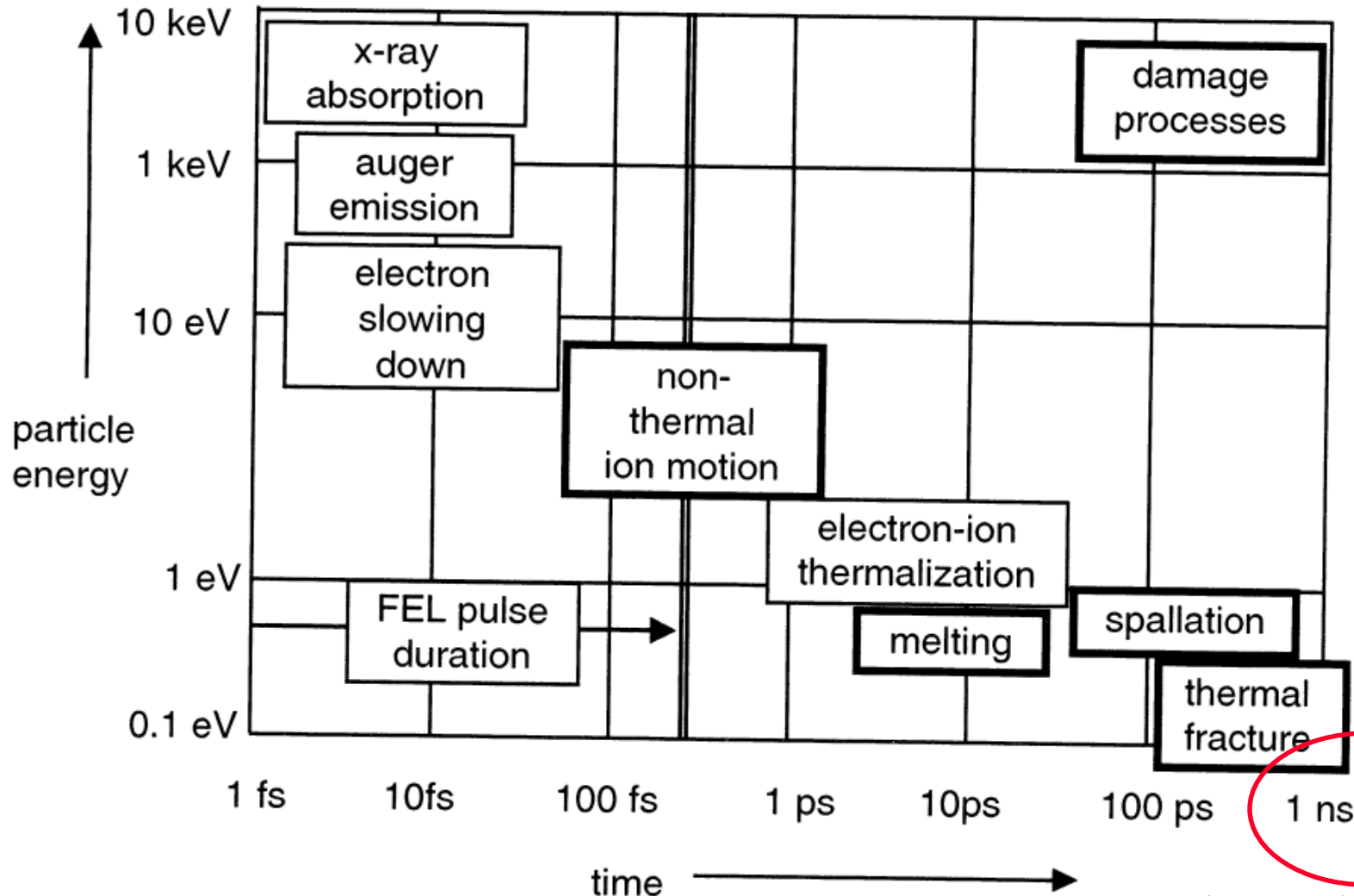


# X-ray and electron processes

R.A. London et al, "Computational studies of high intensity X-ray matter interaction",  
Optics for Fourth Generation X-ray Sources, SPIE Proc (2001), **4500**, p. 51



# Energy-time plot for x-ray-matter interaction



temperature and density relax. time

## Fluence [ $\text{J}/\text{cm}^2$ ] that matters

- for very short pulses (e.g. both XFELs and ERLs) it's fluence that matters
- XFEL:  $10^{12} \times 10 \text{keV} \sim 1 \text{ mJ} / \text{pulse}$ ; ERL:  $10^6\text{-}10^7 \times 10 \text{keV} \sim 10 \text{ nJ} / \text{pulse}$
- tolerable dose can be estimated as following (e.g. Si):
  - specific heat times  $1700 - 300 \text{ K}$  temperature difference
  - plus fusion heat, =  $78 \text{ kJ/mole}$
  - normalized per atom  $\sim 0.8 \text{ eV/atom}$
- most elements have melting dose between  $1/3$  to  $1 \text{ eV/atom}$
- $< 0.1 \text{ eV/atom}$  considered safe



## Damage onset with instantaneous dose

0.1 eV/atom  
threshold for permanent structural changes



1 eV/atom  
most materials are melted



10 eV/atom  
ablation begins



higher dose eventually leads to Coulomb explosion

# Tolerable spot size for melting

Use protein density  $0.8 \text{ Da}/\text{\AA}^3$

carbon photo-absorption cross-section  $\sigma_a = 85 \text{ barn/atom}$

dose per atom:  $E_{\text{pulse}} \sigma_a / \text{area}$

For tolerable dose  $0.1 - 1 \text{ eV/atom}$

XFEL  
 $10 - 30 \mu\text{m}$

ERL  
 $6^* - 60 \text{ nm}$

For smaller spot sizes, one moves into a “single shot” regime

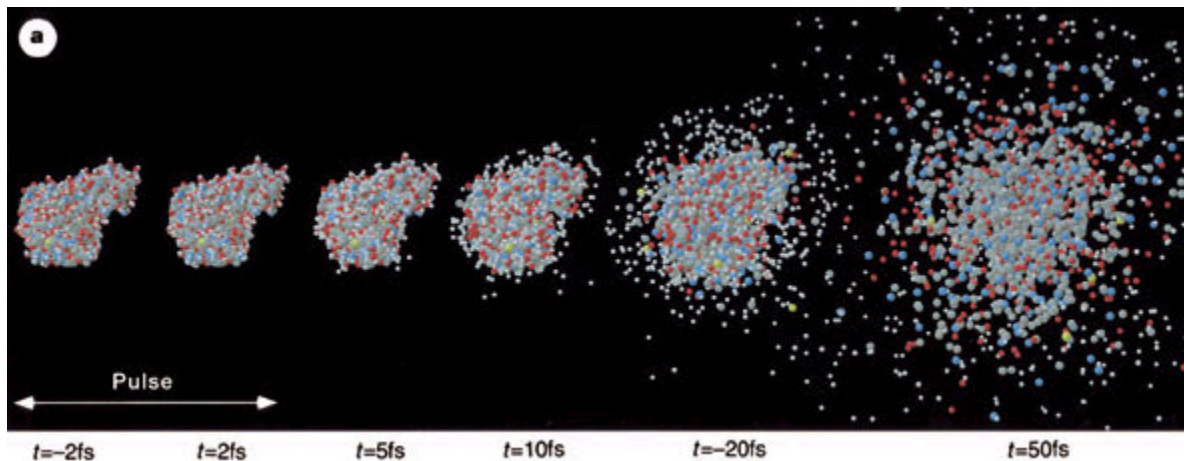
For high-Z materials this number is worse

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\* low flux regime

# Explosive proteins

R. Neutze, et al., *Nature*, **406**, 752-757, January 17, 2000



Very briefly: calculations were done for T4 lysozyme (diameter  $32 \text{ \AA}$ ,  $N_C \sim 1000$ ); flux  $4 \times 10^6 \text{ X-rays/\AA}^2$  with  $\sim 2000$  primary ionization events; elastically scattered  $\sim 200$  photons. The claim is that if pulse is sufficiently short (much shorter than the LCLS spec),  $5 \times 5 \times 5$  lysozyme nanocrystal will scatter to  $< 2 \text{ \AA}$  resolution.

# Conventional damage to proteins

- Primary: breaking of chemical bonds
- ~~• Secondary: chemical damage by free radicals~~
- ~~• Tertiary: crystal lattice destabilized in absence of further chemical damage (domino effect)~~

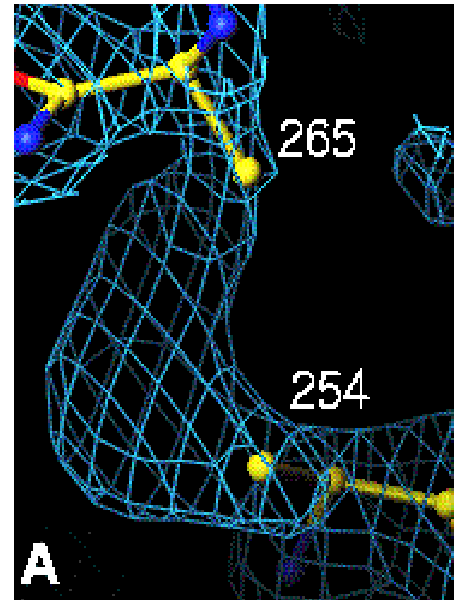
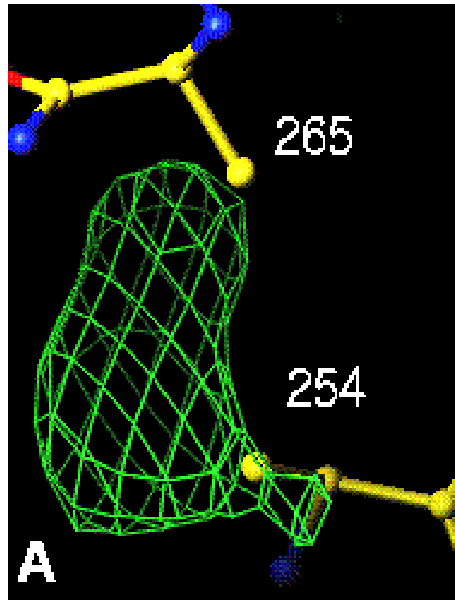
cryoprotection helps with these two (prevents mass loss)

- Primary radiation dose  $10^7$  Gy or  $\sim 200$  X-rays/Angstrom<sup>2</sup> (Henderson's limit)
- It's accumulated dose that matters (unlike "fast melting")
- Despite the very different mechanisms, the damage dose is 1.4 eV/atom, very similar to the "single shot melting"
- Coulomb explosion requires much greater dose (delivered in a single pulse)

# Disulphide bonds go first

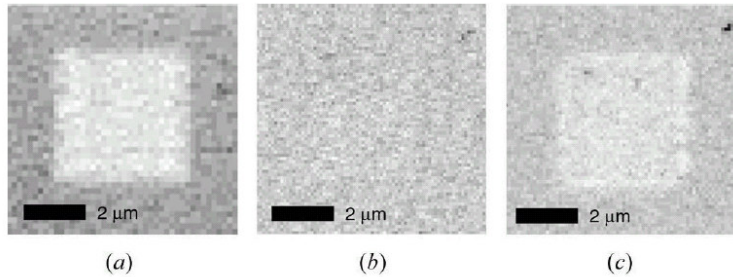
M. Weik, et al., *PNAS*, Vol. 97, Issue 2, 623-628, January 18, 2000

Each frame is a complete data-set collected in about 3.5 minutes, each data-set is a 15 minutes time point. The left panel is the  $3F_o-2F_c$  map and model of the 254-265 disulfide bridge, and the right panel is the  $F_o-F_c$  map of the same S-S bond.



# Cryoprotection prevents mass loss

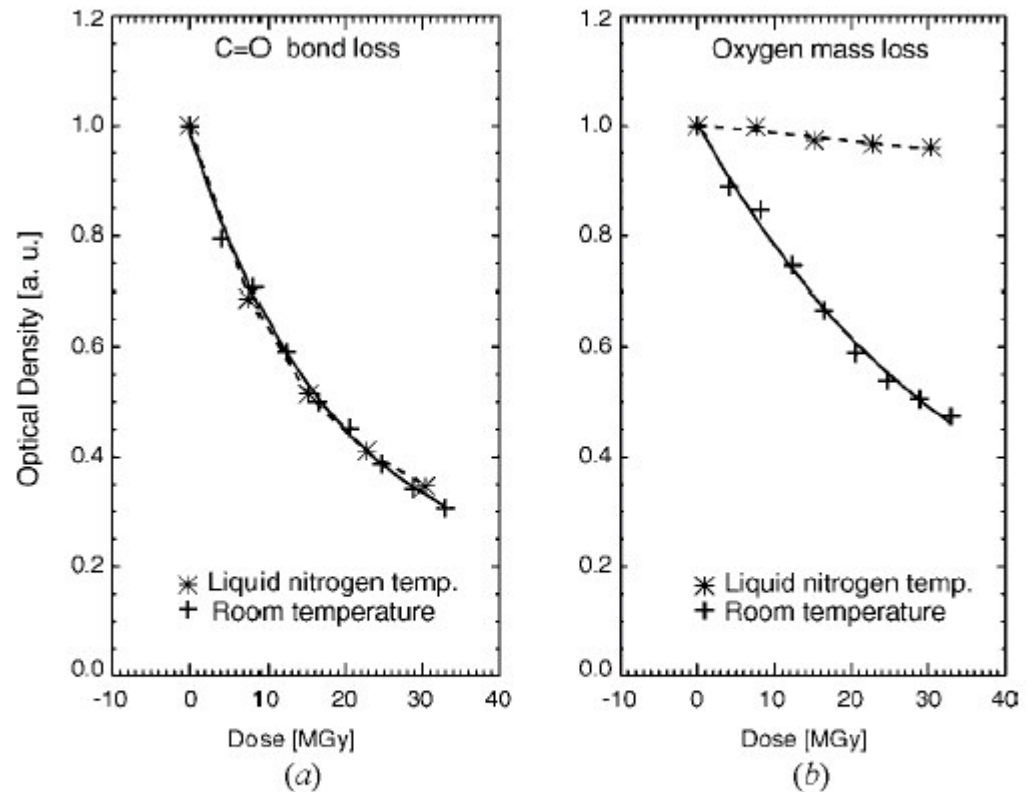
T. Beetz and C. Jacobsen, *J. Synchrotron Rad.* (2003). **10**, 280–283



**Figure 2**

(a) Coarse area scan at room temperature of a region that was dosed at room temperature. The square in the middle indicates the mass loss of the dosed region. (b) Coarse area scan at liquid-nitrogen temperature of the region that was dosed at liquid-nitrogen temperature. The image shows no visible mass loss at the dosed region. (c) Same area as that in image (b) but after warming up the sample; the dosed region becomes visible.

Radiation damage for  $\sim 10$  nm resolution in cryoprotected X-ray microscopy is estimated to be at  $\sim 10^{10}$  Gy ( $\sim 1$  keV/atom ac. dose) [J. Maser et al., *J. Microscopy* (2000), **197**, p. 68]



**Figure 4**

(a) Loss of C=O peak intensity and (b) mass loss as a function of dose at liquid-nitrogen and room temperatures. The data were fitted using equations (2) and (3), where the fitting coefficients are summarized in Table 1.

## Some preliminary conclusions

- Cryoprotection is likely to be ineffective for “single shot melting”, i.e. X-ray microscopy is better off with c.w. source like an ERL
- Applications requiring multiple exposures of the same sample (e.g. tomography) with good resolution will prefer c.w. source over XFEL
- “single shot” experiments are for XFELs
- ...