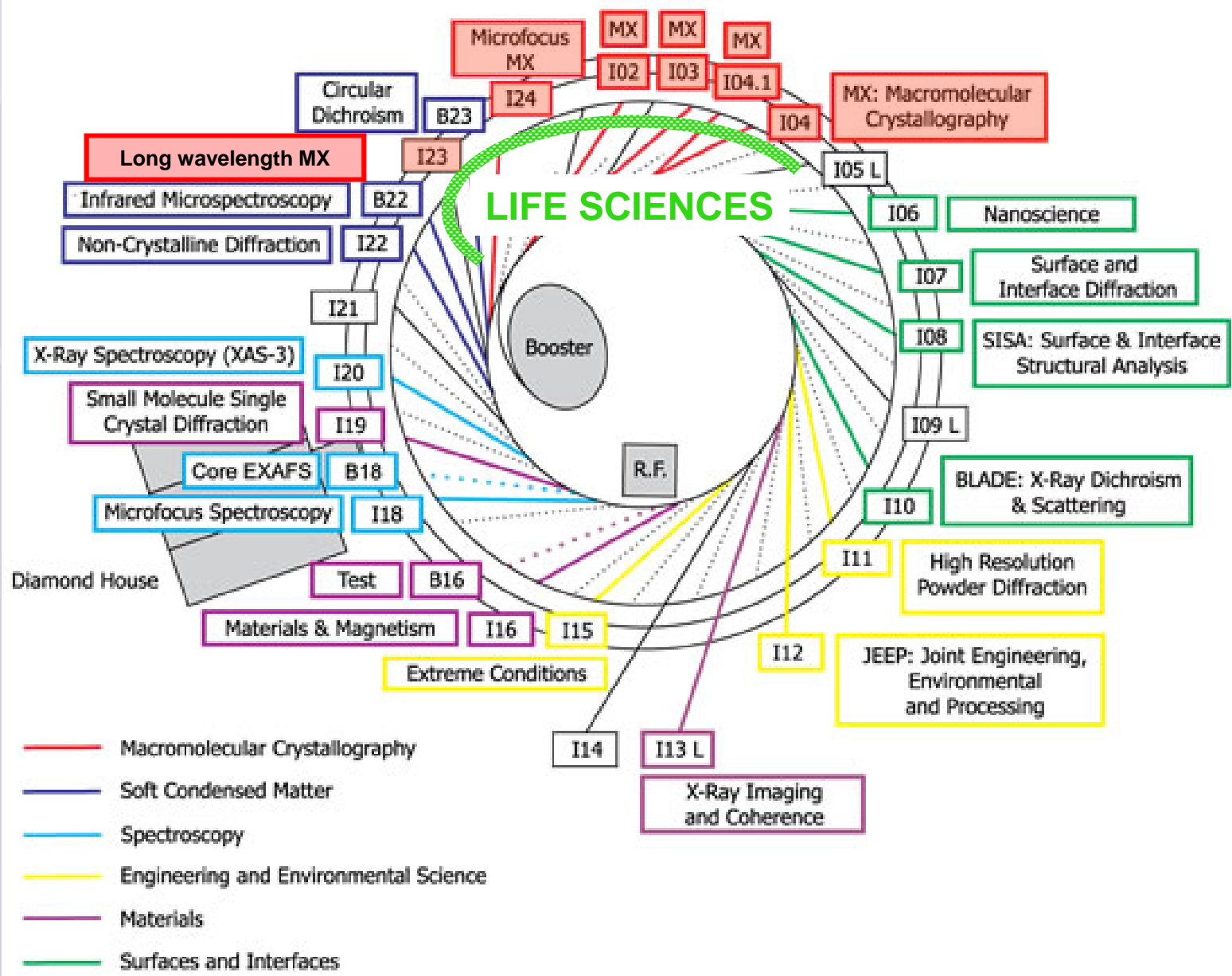




The Long-Wavelength Macromolecular Crystallography Beamline I23 at Diamond Light Source

Armin Wagner
Diamond Light Source



Why long wavelengths?

- Experimental Phasing
 - SAD phasing from native proteins and DNA by using anomalous signal from sulphur and phosphorous.
 - SAD and MAD phasing for large complexes by using enormous anomalous signals from M-edges.
- Element specific analysis

On the routine use of soft X-rays in macromolecular crystallography. Part III. The optimal data-collection wavelength

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Santosh Panjekar, Paul A. Tucker
and Manfred S. Weiss***

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Complete and highly redundant data sets were collected at different wavelengths between 0.80 and 2.65 Å for a total of ten different protein and DNA model systems. The magnitude of the anomalous signal-to-noise ratio as assessed by the quotient $R_{\text{anom}}/R_{\text{r.i.m.}}$ was found to be influenced by the data-collection wavelength and the nature of the anomalously scattering substructure. By utilizing simple empirical correlations, for instance between the estimated $\Delta F/F$ and the expected R_{anom} or the data-collection wavelength and the expected $R_{\text{r.i.m.}}$, the wavelength at which the highest anomalous signal-to-noise ratio can be expected could be estimated even before the experiment. Almost independent of the nature of the anomalously scattering substructure and provided that no elemental X-ray absorption edge is nearby, **this optimal wavelength is 2.1 Å.**

Received 6 April 2005
Accepted 5 July 2005

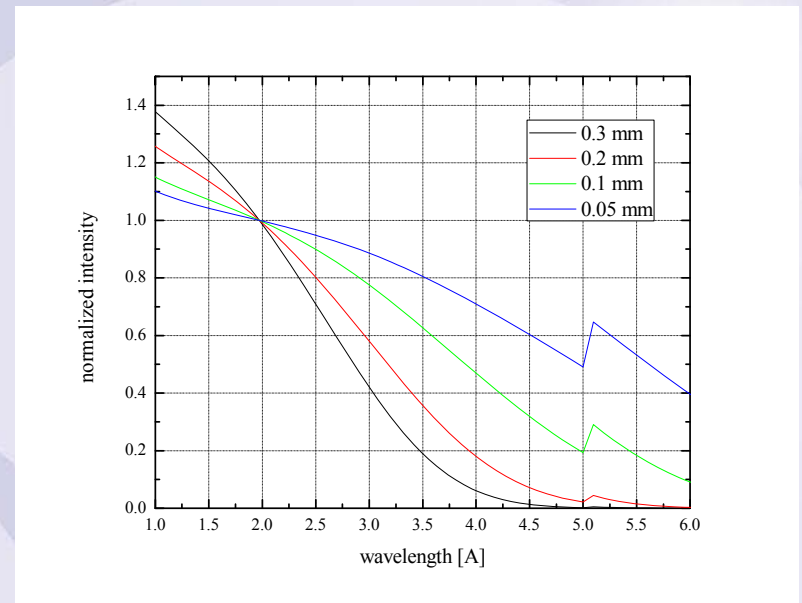
PDB References: ConA-Xe, 2a7a, r2a7asf; adaptin-Xe, 2a7b, r2a7bsf; PPE-Xe, 2a7c, r2a7csf; HEL-Xe, 2a7d, r2a7dsf; DNA, 2a7e, r2a7esf; HEL, 2a7f, r2a7fsf; thermo-lysin, 2a7g, r2a7gsf; trypsin, 2a7h, r2a7hsf; thaumatin, 2a7i, r2a7isf; PPE-Ca, 2a7j, r2a7jsf.

Optimal wavelength

- Radiation damage proportional to absorbed dose

$$I_E \propto \frac{t^3 e^{-\mu t}}{1 - e^{-\mu t}} \lambda^3$$

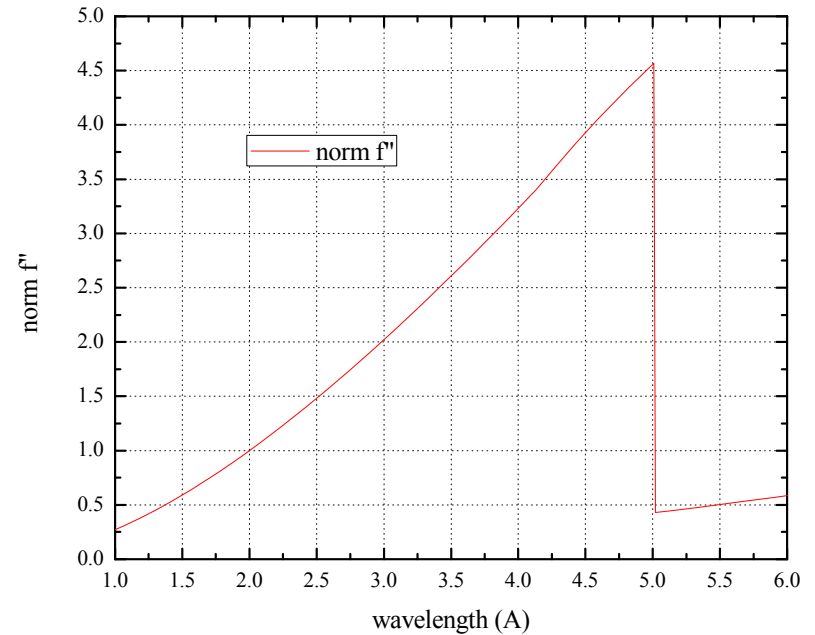
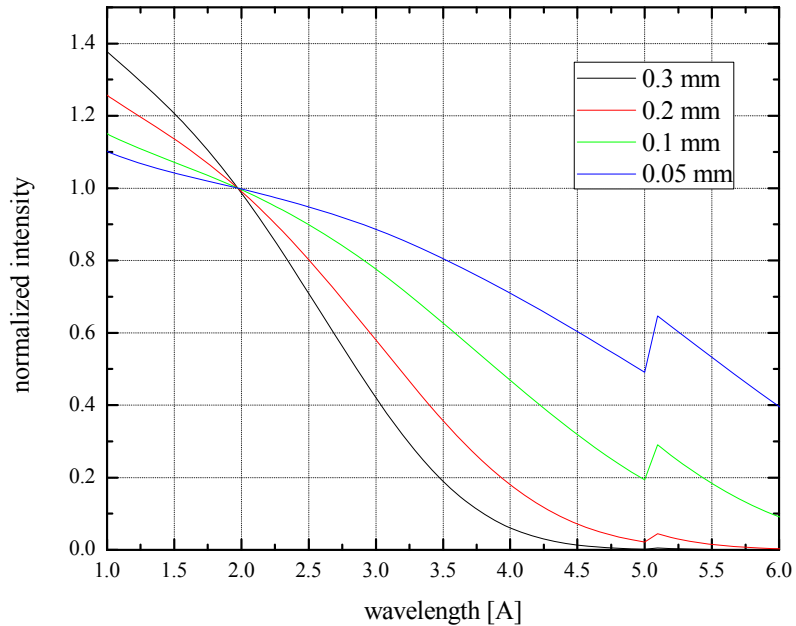
- Arndt (1984)
- Isometric crystal
- Bragg intensity per energy absorbed
- Radiation damage wavelength independent



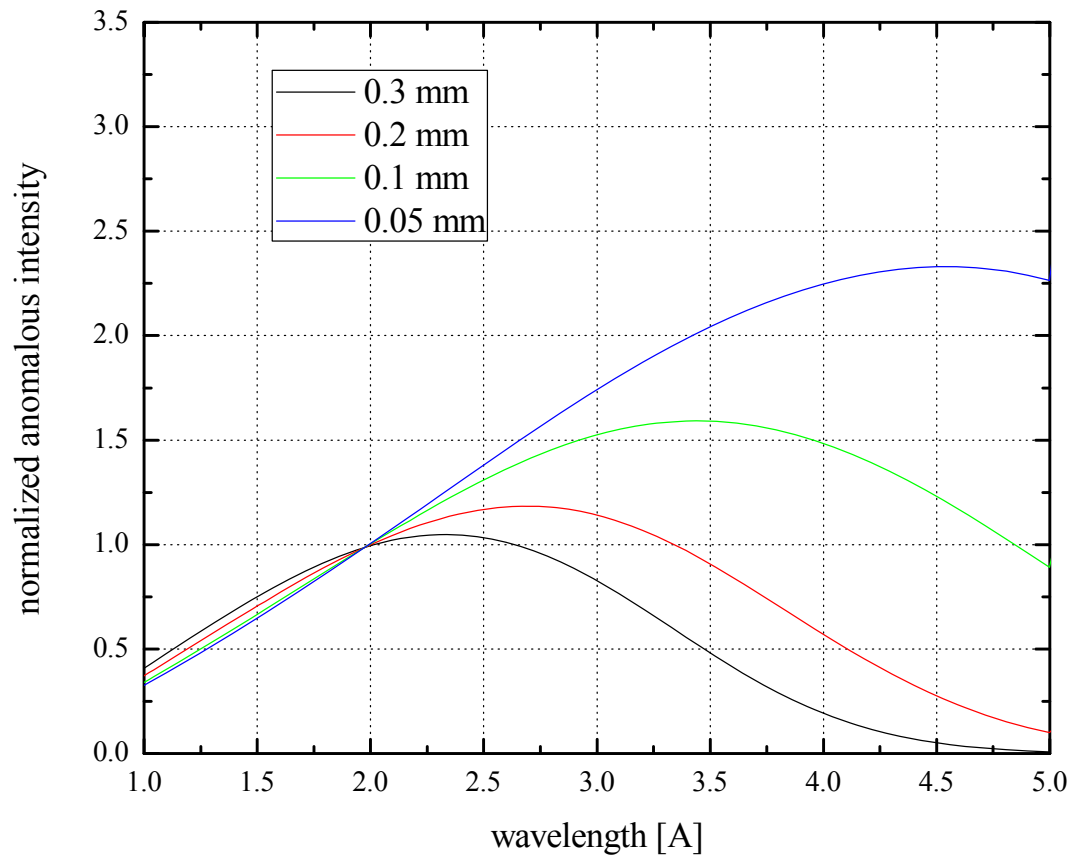
J. Appl. Cryst. (1984). 17, 118–119

Optimum X-ray wavelength for protein crystallography. By U. W. ARNDT, MRC Laboratory of Molecular Biology, Hills Road, Cambridge CB2 2QH, England

Optimal wavelength for S-SAD



Optimal wavelength for S-SAD



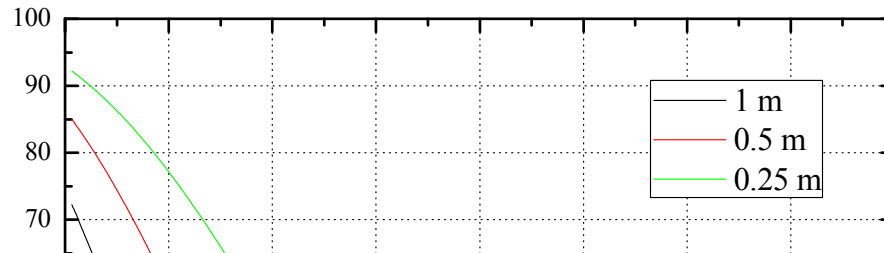
BUT!!!

ABSORPTION

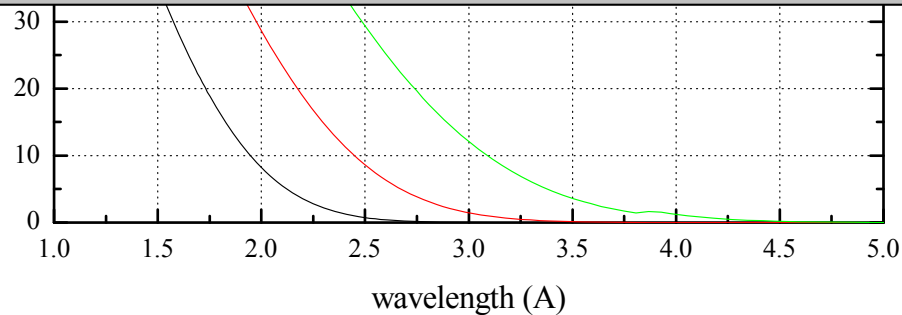
DETECTORS



Air absorption

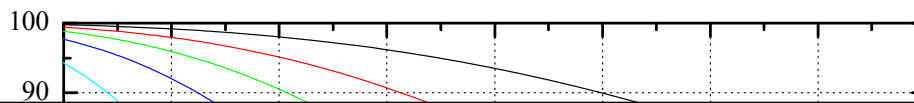


Sample and detector have to be either in He or vacuum!

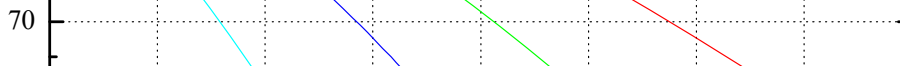


Sample absorption

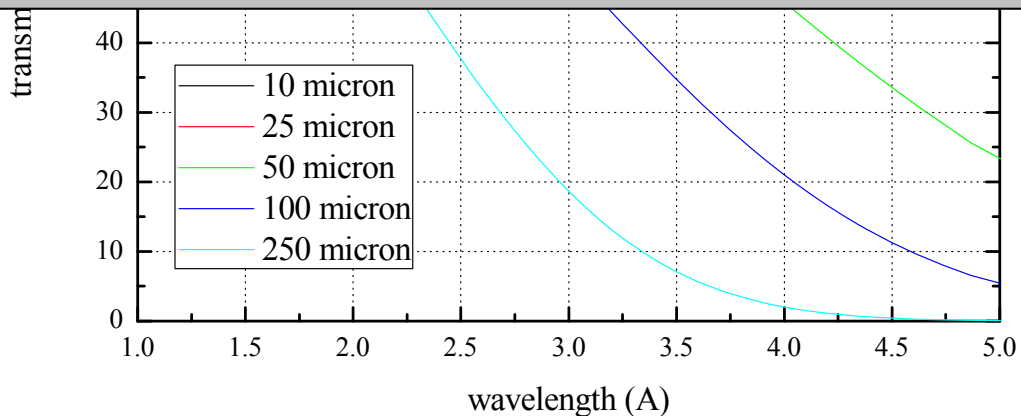
Lysozyme



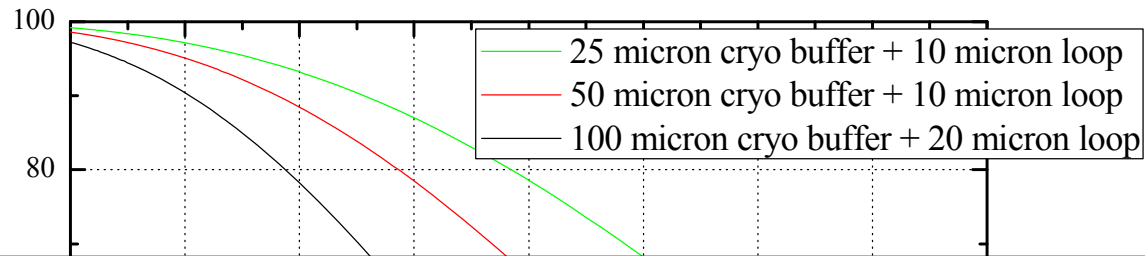
Longest wavelengths only suitable for small crystals!



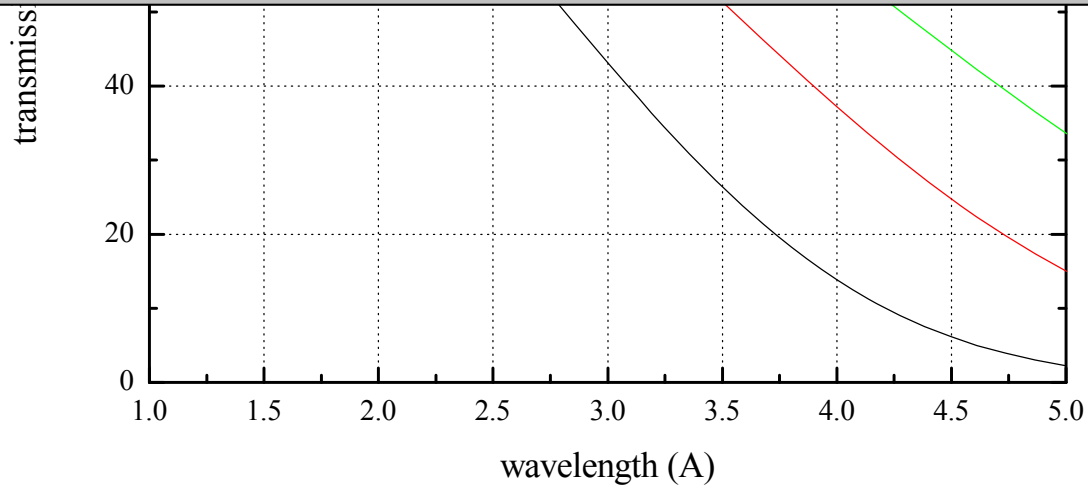
Analytical absorption correction necessary!



Solvent and loop absorption



Reduce amount of buffer around the crystal!



I23 – Long wavelength MX beamline

Energy (wavelength) range:

2.1 - 12 keV (1 - 5.9 Å), optimized for 3 - 8 keV (1.5 - 4 Å)

Band-pass ($\Delta E/E$):

2×10^{-4}

Beam size at sample:

100 - 1000 μm

Photon flux:

5×10^{12} ph/s in 100 x 100 μm @ 4 keV

End station:

Multi-axis goniometer in vacuum with micrometer precision for X-ray diffraction and tomography experiments from frozen macromolecular crystals in the temperature range 30 - 120 K.

Detector:

Area detector with high quantum efficiency to cover large solid angle



B23

I23

B22

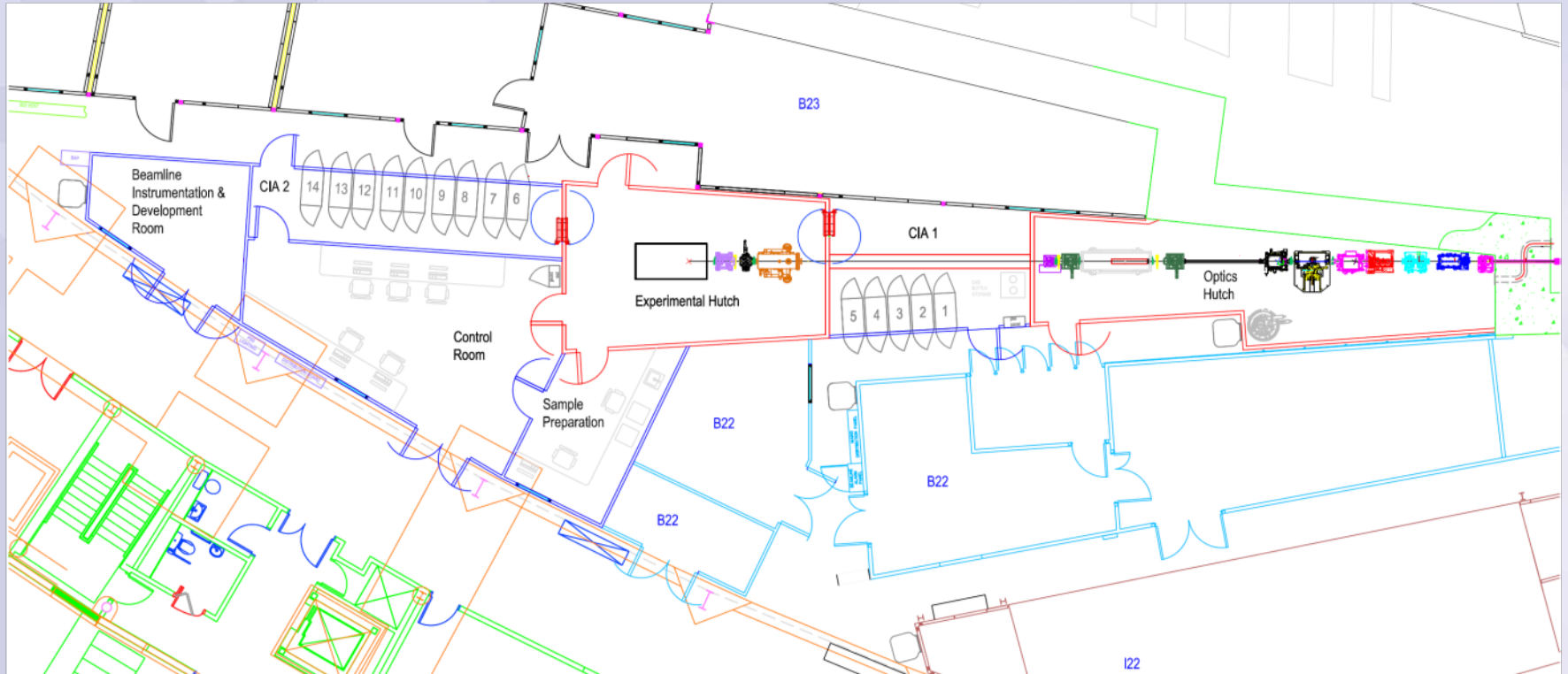
ISCOVER

KONIGDRUMS

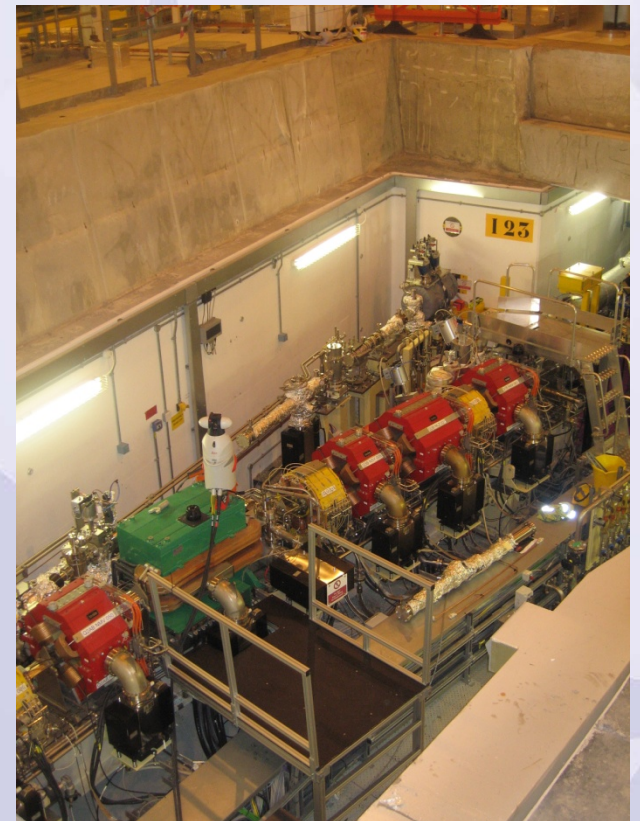
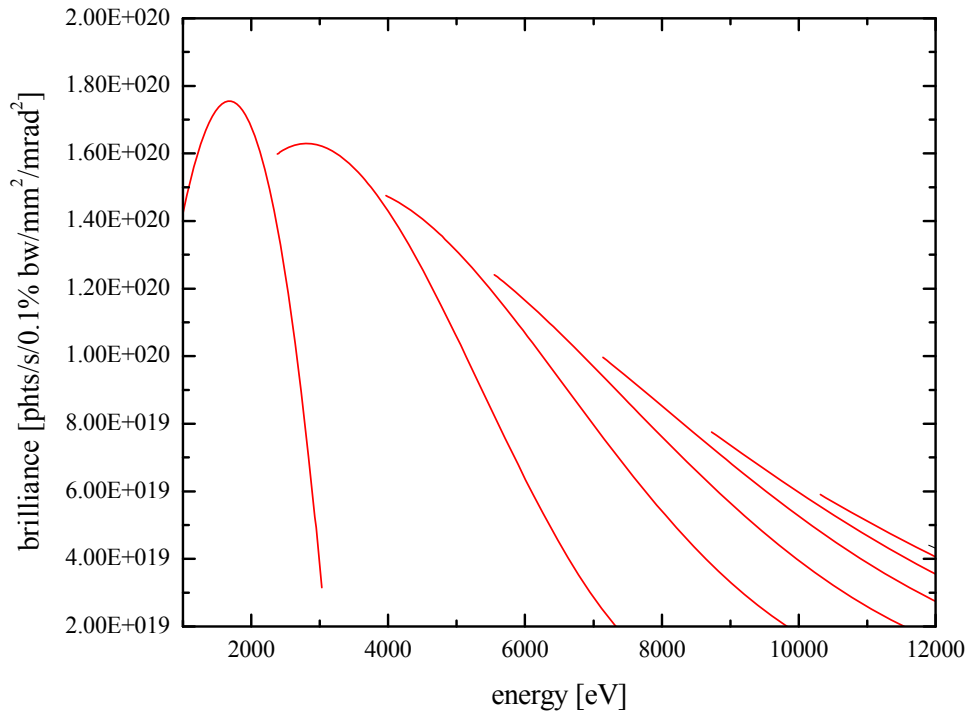
20t

01

Beamline Layout



Insertion Device Front End



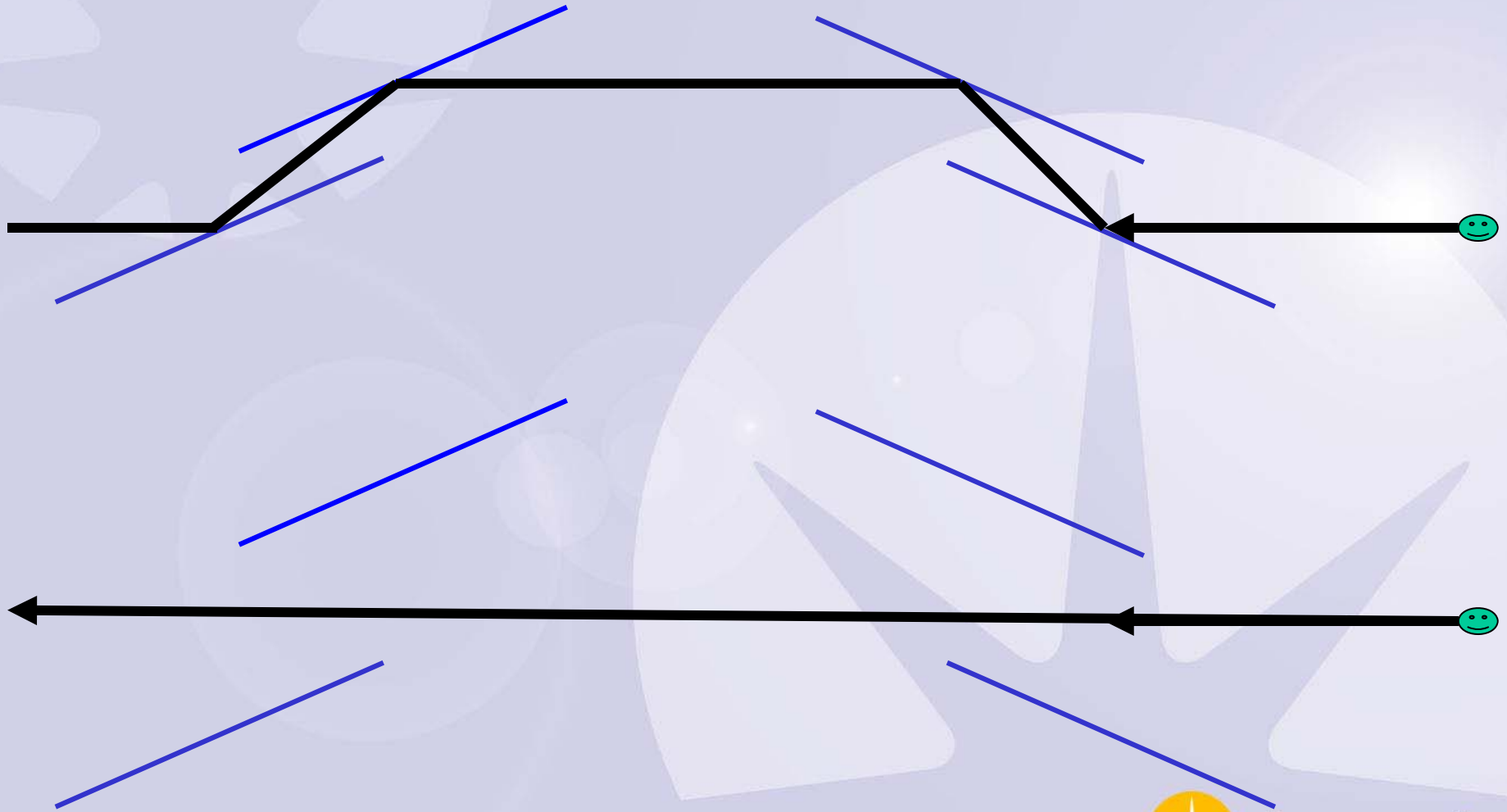
Phase I of FE installation during Christmas shutdown.

Optical layout

- All mirrors horizontally deflecting.
- Elliptical mirror for tangential focussing in horizontal plane.
- Cylindrical mirror for sagittal focussing in vertical plane.
- Two flat mirrors for harmonic rejection.

- Fast switching by moving first and last mirror either vertically or horizontally out of the beam.
- No optical elements for tomography!

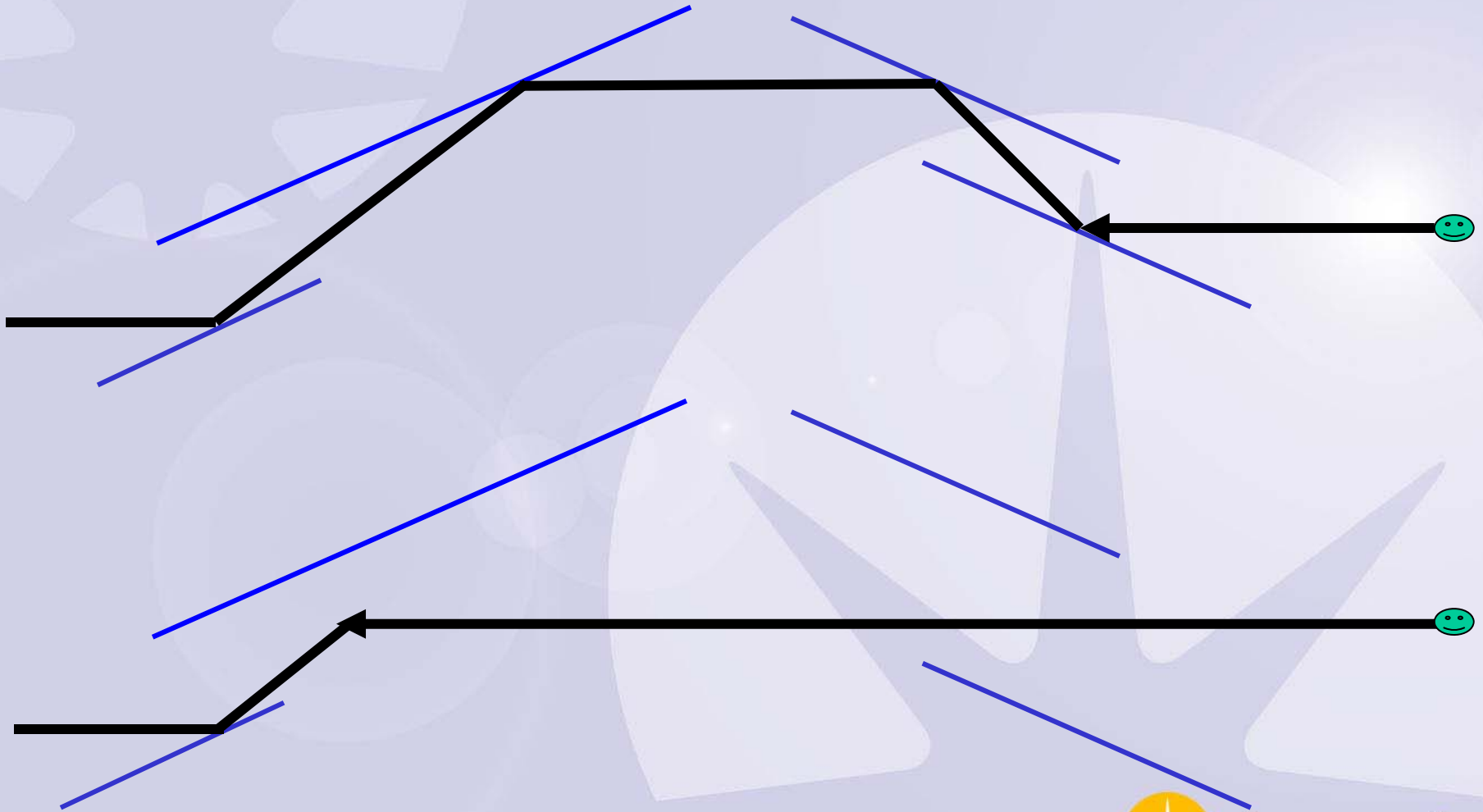
Four-mirror system (case A)



No optical elements, no harmonic rejection for tomography



Four-mirror system (case B)



Two mirrors for harmonic rejection for tomography

Challenges

- Absorption
- Limited resolution
- Radiation damage

Air absorption

Vacuum sample environment

Crystal, solvent and loop absorption

X-ray tomography

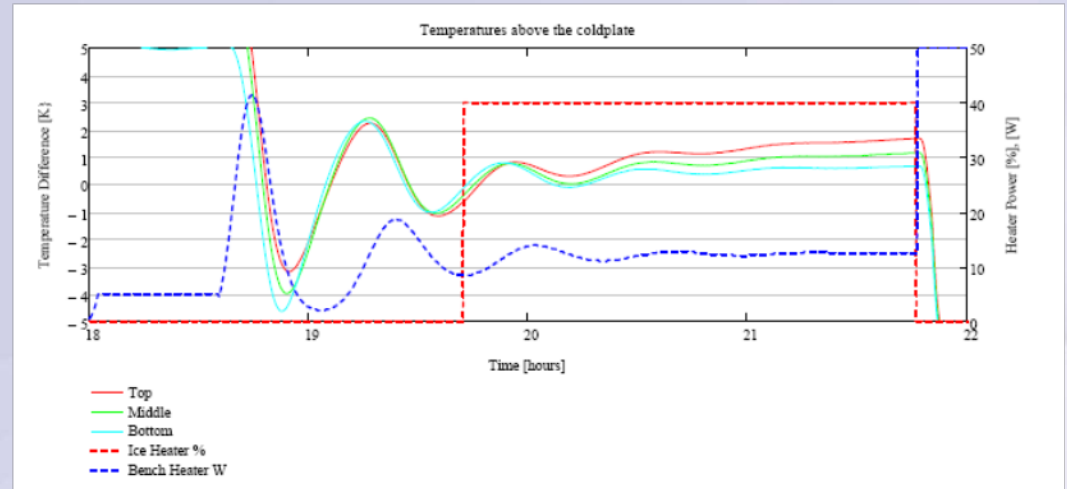
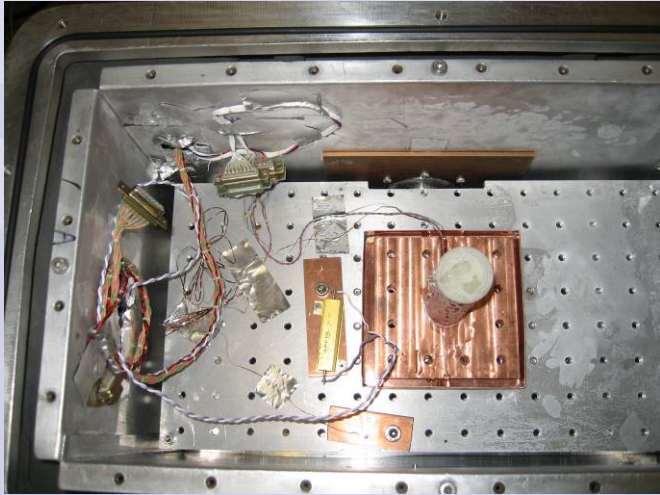
Analytical absorption correction

Conductive sample cooling

- As one main risk the User Working Group identified the efficiency of conductive cooling for protein crystals.
- Main uncertainty, thermal contact conductance to allow sample change and conductivity of vitreous ice.



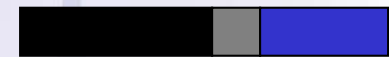
Thermal conductivity of vitreous ice



- Preliminary results
 - $3.5 \text{ Wm}^{-1}\text{K}^{-1}$ from temperature differences between sensors in vitreous ice sample (20% Ethylenglycol)
 - $1.9 \text{ Wm}^{-1}\text{K}^{-1}$ from cooling rate

Crystal Heating

- 20 MGy in 30 s
- Cylindrical crystal 100 μm long, 30 μm diameter
- Protein crystal density (30% solvent)
- 0.059 mW on crystal
- Thermal conductivity of crystal like vitreous ice
- Heating: <5 K



Diamond ice crystal

Sample transfer

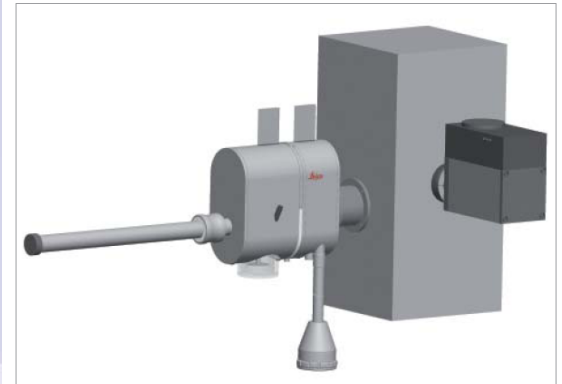


EM VCT100 shuttle with specimen and holder

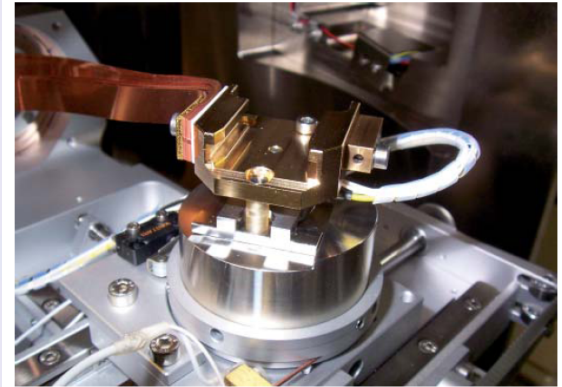


EM VCT100 cryo preparation workstation

- Standard system without customization



EM VCT100 docking station and shuttle with specimen and holder connected to the cryo stage.

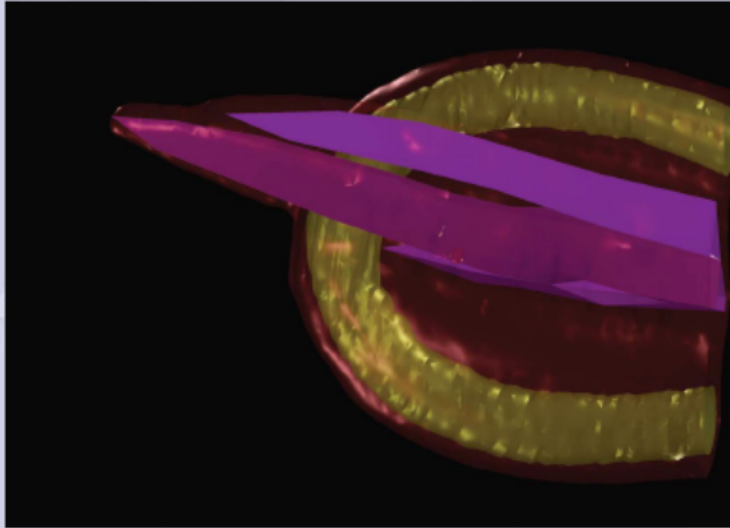


Custom-made cryo stage



The EM VCT100 is compatible with all current SEMs.

X-ray Tomography



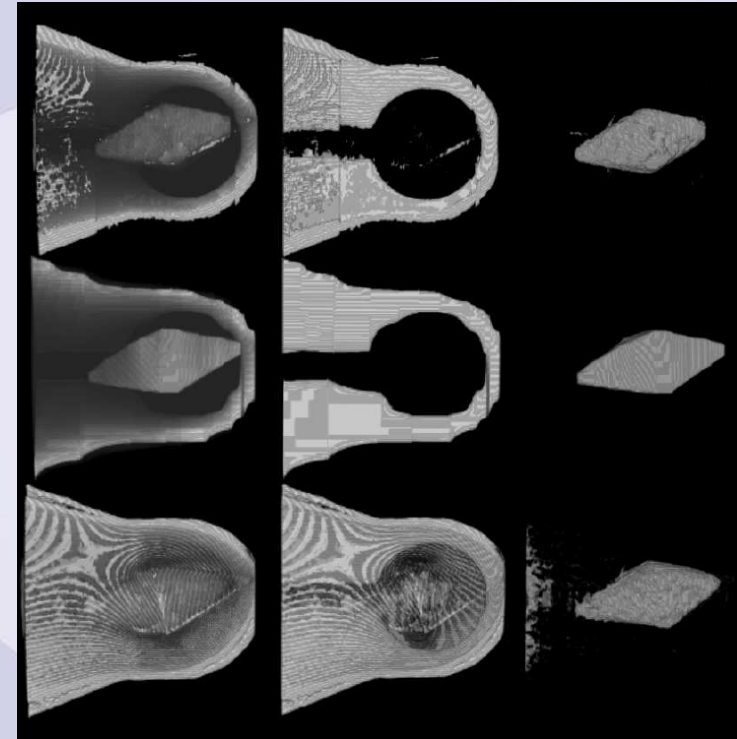
- Both absorption and phase contrast can be exploited
- Resolution: $\sim 1 \mu\text{m}$
- Diamond software development for analytical absorption correction based on tomographic reconstruction

Diamond Developments

Analytical absorption correction based on tomographic reconstructions

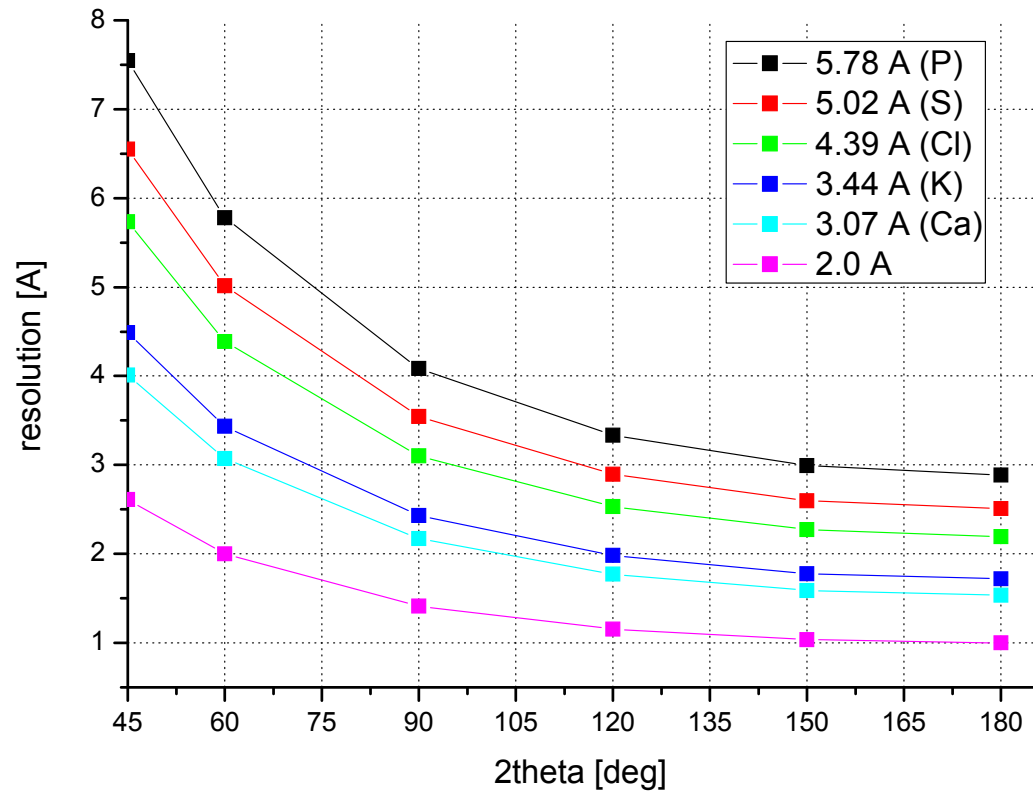
W. Armour et al.

3 publications in preparation.



Challenges

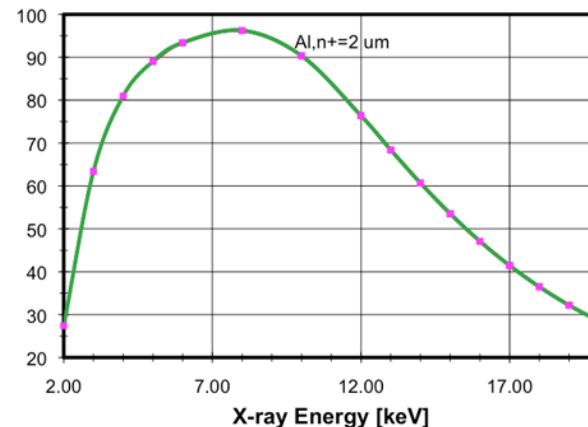
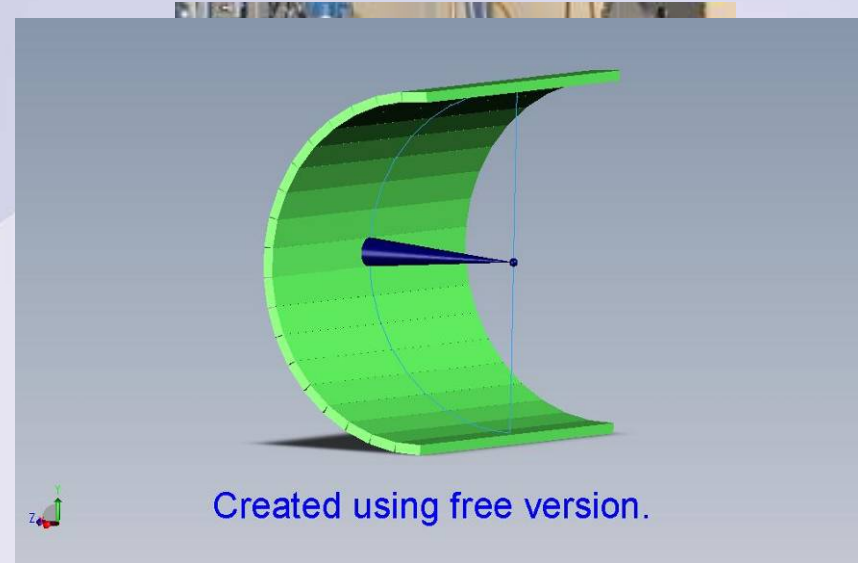
- Abs
- Lim
- Ra



Detector

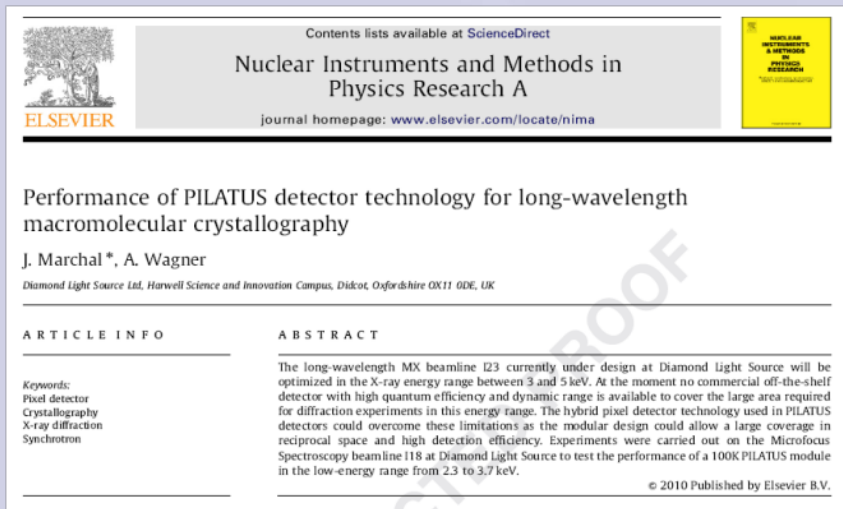
- Pilatus 6M
- Active area 431 x 448 mm²
- Energy range 3–30 keV
- Readout time <3.6 ms
- Framing rate 10 Hz

- No dark current
- No readout noise
- Single photon counter



Detector

- Low-energy experiments on Diamond beamlines I18 and B16 to test Pilatus technology at low energies in vacuum.



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13th INTERNATIONAL WORKSHOP ON RADIATION IMAGING DETECTORS,
3–7 JULY 2011,
ETH ZURICH, SWITZERLAND

Low-energy X-ray detection with an in-vacuum PILATUS detector

Julien Marchal,^{a,1} Benjamin Luethi,^b Catalin Ursachi,^a Vitaliy Mykhaylyk^a and Armin Wagner^a

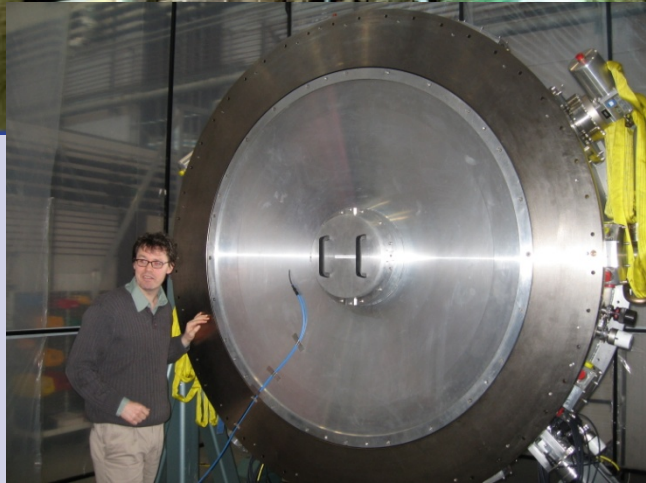
^a*Diamond Light Source,
Harwell Science and Innovation Campus, Didcot, Oxfordshire, OX11 0DE, U.K.*

^b*Dectris Ltd,
Neuenhoferstrasse 107, 5400 Baden, Switzerland*

Pilatus 12M

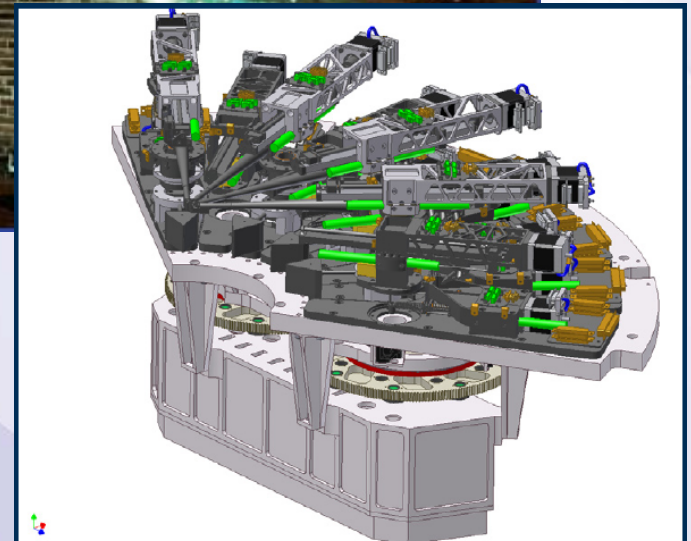
- In-vacuum detector
- Half cylinder $r = 250$ mm
- $2\theta = \pm 90^\circ$

Royal Observatory Edinburgh



KMOS K-band multi-object spectrometer

To be installed at European Southern Observatory in Chile soon.



A section of the instrument showing the pick off arms and calibrator.

In-vacuum goniometer

- Commissioned to Astronomy Technology Centre in Edinburgh
- 3 μm sphere of confusion

Challenges

- Absorption
- Limited resolution
- Radiation damage

Air absorption

Vacuum sample environment

Crystal, solvent and loop absorption

X-ray tomography

Analytical absorption correction

Large area detector

2theta arm

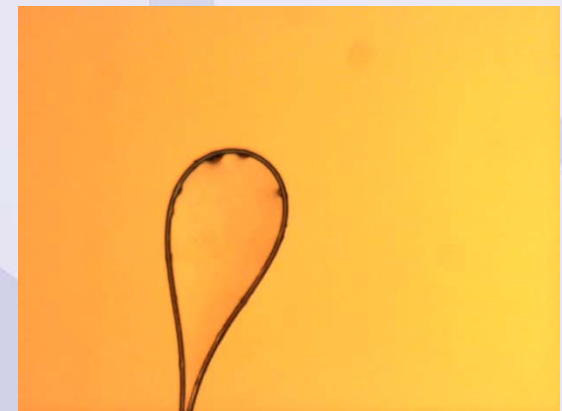
Multi axis goniometer

Radiation Damage

- Most severe limitation on 3rd generation synchrotrons
- Drastic reduction from RT to 100 K
- 95% of experiments at 100 K
- Specific radiation damage alters anomalous substructure

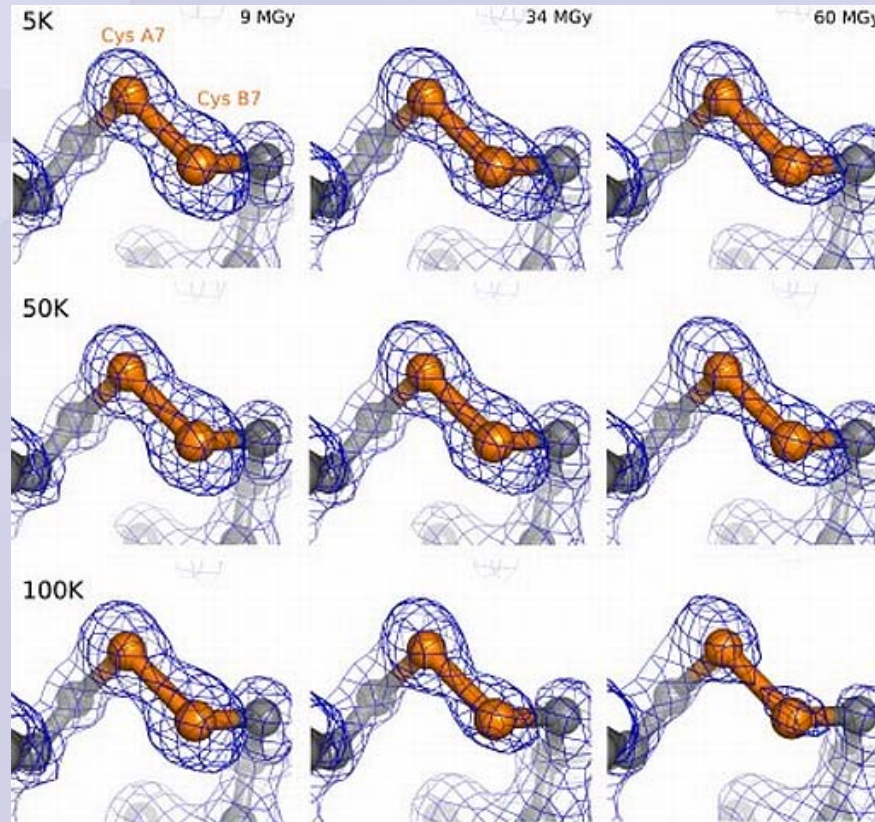


10 s exposure
(protein buffer + 20% ethylenglycol)



200 s exposure (30% ethylenglycol)
Warming up to ~200K

Specific radiation damage

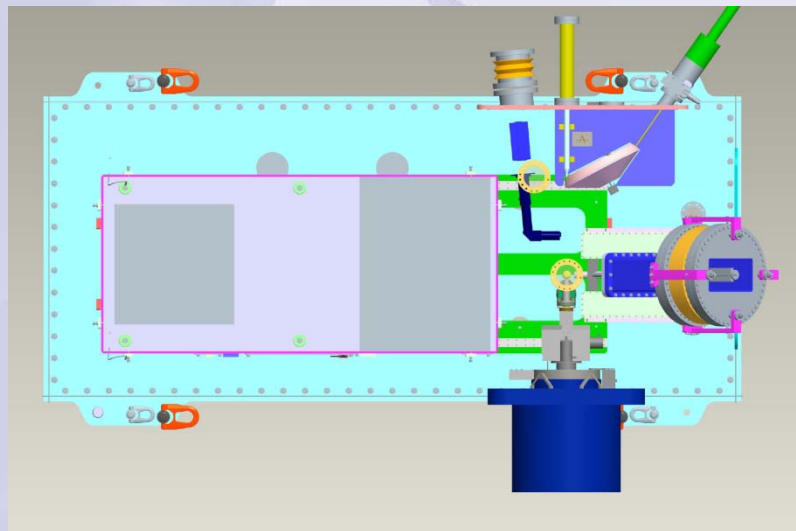


- A. Meents, S. Gutmann, A. Wagner, C. Schulze-Briese (2010) PNAS 107, 1094 – 1099.

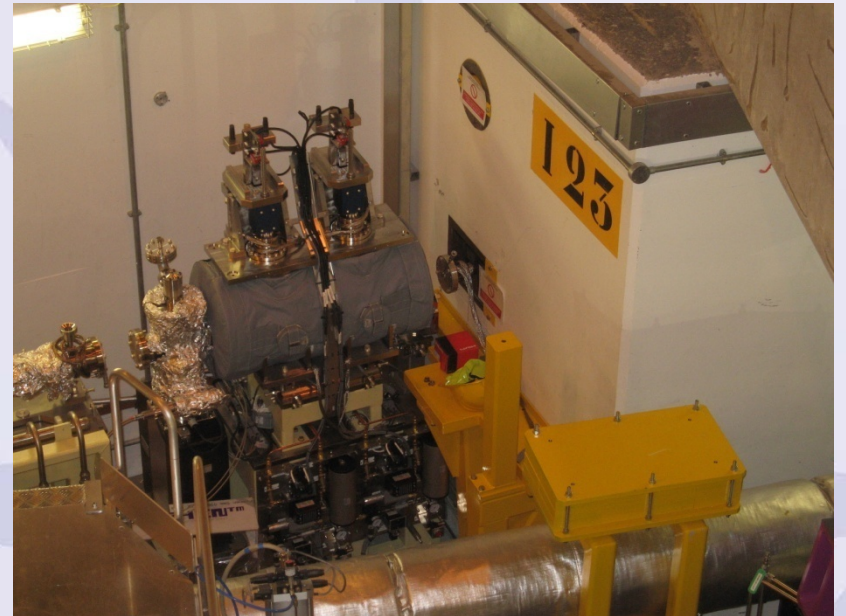
Summary

End station most important aspect of beamline!

- High precision in-vacuum multi-axis goniometer
- In-vacuum detector
- Conductive cooling for temperatures ~ 40 K
- Sample holders compatible with cooling
- Sample transfer of frozen crystals into vacuum chamber
- Integration of tomography system



Milestones	Date
<u>Design</u>	
Conceptual Design Report Review	12/01/09
Technical Design Report Review	01/07/10
<u>Construction</u>	
Hutches Complete	19/07/11
Hutches, Cabins and Services complete	27/03/12
<u>Installation</u>	
Optics Hutch installation complete	04/09/12
Experimental Hutch installation complete	10/04/13
<u>Commissioning</u>	
First Light for beamline I23	06/09/12
I23 First User	04/10/13



Project Team

- Vitaliy Mykhaylyk (Beamline scientist)
- Martin Burt (Project engineer)
- Jon Kelly (Mechanical engineer)
- Kevin Wilkinson (Electrical engineer)
- Gary McIntyre (Design engineer)
- Stephen Green (Design engineer)
- Ronaldo Mercado (Controls)
- Richard Fearn (Data Acquisition)
- Lucia Alianelli (Optics)
- Emily Longhi (ID)
- Julien Marchal (Detectors)