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Scientific case

X-ray crystallography has emerged as a very effective technique to perform 3D structural studies on biomolecules at the atomic level.

In response to this, one of the phase-I beamlines (**XALOC – BL13**) of the third generation 3-GeV synchrotron Alba will be devoted to Macromolecular Crystallography.

Overall status

Beam commissioning: Oct 2011

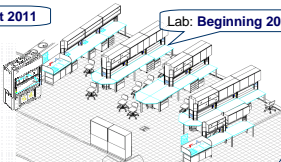
EH commissioning: Jan 2012



Hardware: Oct 2011

Lab: Beginning 2012

TANGO control system/MxCuBE: Jan 2012



XALOC is a versatile beamline

...for many techniques

- All wavelength-dependent techniques can be performed (MAD, SAD, etc) due to the **full-wavelength tunability** and the high resolution ($\Delta\lambda/\lambda \sim 2 \cdot 10^{-4}$) of the beam,
- All common **K and L₃ absorption edges** are reached: 2.4–0.6 Å or 5–21 keV. Covered elements are: V → Mo (K); La → U (L₃),
- An in-vacuum undulator provides a **high-flux beam** ($>2 \times 10^{12}$ ph/s) over the whole energy range

...for many crystal types

- Small crystals** can be studied by **focusing** the beam down to $\sim 50 \times 10 \mu\text{m}^2$ (h×v) with a beam divergence of $0.5 \times 0.1 \text{ mrad}^2$,
- Any medium-sized crystals** can be dealt with by **defocusing** the beam up to $\sim 300 \times 100 \mu\text{m}^2$ (h×v),
- Larger crystals** can be completely exposed to the x-ray beam by **unfocusing** the beam to $\sim 500 \times 500 \mu\text{m}^2$ (h×v). This beam is also highly vertically collimated (0.03 mrad) and hence it can be used for crystals with **very large unit cells** like protein-protein complexes, viruses, etc.

Specifications of the in-vacuum undulator (IU21)

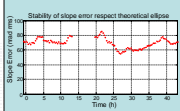
| | |
|--------------------------------|----------------------------|
| Period (number of periods) | 21.6 mm (92) |
| K (at minimum gap, 5.5 mm) | 1.60 |
| Photon source size (h×v, FWHM) | 309 × 18 μm ² |
| Photon source div. (h×v, FWHM) | 112 × 30 μrad ² |



Undulator IU21

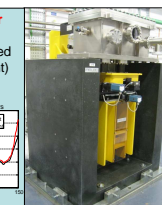
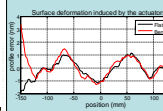
Horizontal Focusing Mirror

- 600 mm optical surface
- Bare Si, Rh, and Ir coated
- 83 nrad slope error (bent)
- 3 Å roughness



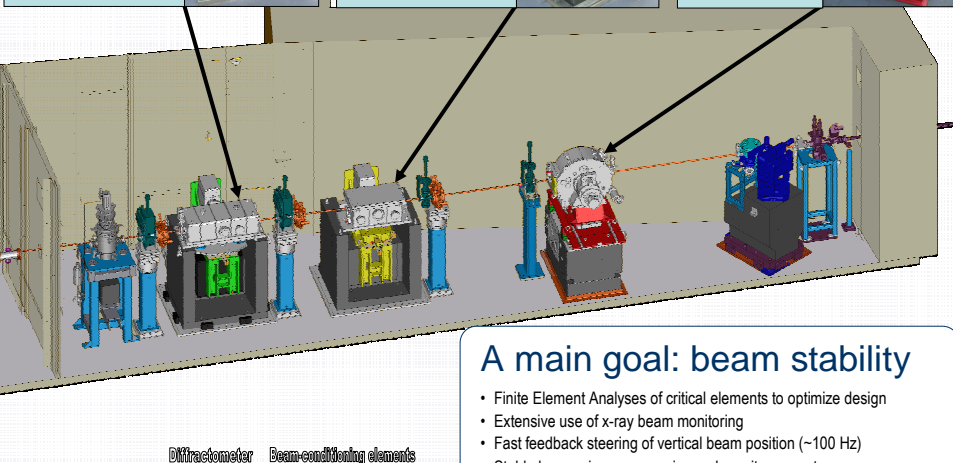
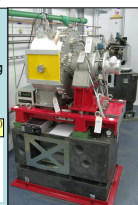
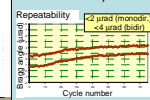
Vertical Focusing Mirror

- 300 mm optical surface
- Bare Si, Rh, and Ir coated
- 70 nrad slope error (bent)
- 3 Å roughness



Monochromator

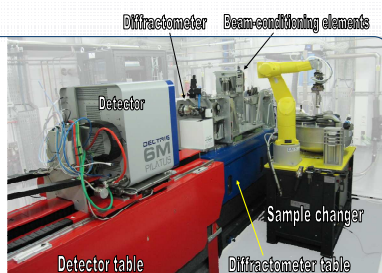
- Si(111) channel-cut
- Liquid N₂ cooled
- Fast V beam steering
- Stability <0.05 eV/h
- Resolution 1 μrad



End Station: the concept

The End station consists of two in-house developed **translation/rotation tables** that support the diffractometer and the detector sitting on a granite base. The tables are adjustable to **1 μm resolution and repeatability**.

An automatic sample changer stands on a nearby table for automatic sample mounting and allows **easy access** to manual mounting.

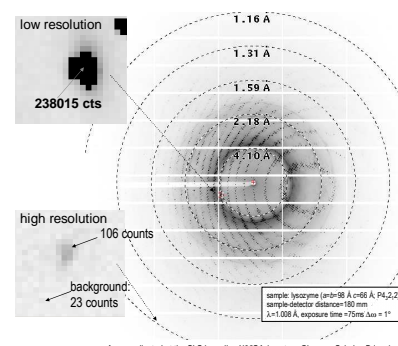


A main goal: beam stability

- Finite Element analyses of critical elements to optimize design
- Extensive use of x-ray beam monitoring
- Fast feedback steering of vertical beam position (~100 Hz)
- Stable base using epoxy resins and granite supports
- Strain gauges in mirror benders to stabilize focusing
- Seismographs close to optical surfaces to deal with vibrations

Detector: Dectris Pilatus 6M

- High dynamic range** (1 million): collection of low and high resolution data on the same frame
- Extremely low background noise**
- Fast read-out**: shutterless collection, minimizing systematic errors
- Thin ω -slicing** which often results in better data in high-resolution data and large unit cell crystals
- Fast frame collection** (80ms/image): a 360° 1.1 Å resolution dataset can be collected in 33 seconds



Diffractometer: Maatel minidiffractometer

Diffractometer with a compact & commercial design, able to deal with crystallization plates

Omega axis

- Air bearing
- Accuracy ±0.7 mdeg
- Run-out ~2 μm

Removable mini-kappa

- Run-out <5 μm
- Improve completeness
- Improve MAD/SAD datasets

Back-lit sample

- Improves visualization

Fluorescence detector

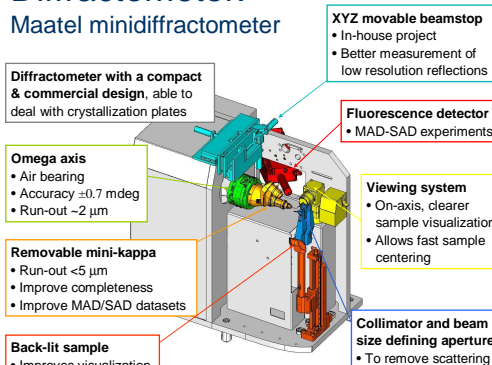
- MAD-SAD experiments

Viewing system

- On-axis, clearer sample visualization
- Allows fast sample centering

Collimator and beam size defining aperture

- To remove scattering



Sample changer: Irelec CATS for SPINE pins & crystallization plates

