

Beamline BL13–XALOC: Macromolecular Crystallography

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www.cells.es/beamlines/XALOC

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Scientific case

The need for 3D atomic information of biological molecules in Biology, Biomedicine, and related disciplines has increased exponentially in the past few decades. X-ray crystallography has emerged as a very effective technique to perform these 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.



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 ($\Delta\lambda/\lambda\sim 2$ 10⁻⁴) of the beam,
- All common K and L₃ absorption edges can be reached due to the large wavelength span: 2.4–0.6 Å or 5–21 keV. Covered elements are: V→Mo (K edge); La→ U (L₃ edge).,
- An in-vacuum undulator (IVU21) feeds the beamline with a high-flux beam (>10¹² ph/s in 100×100 μm²) over the whole wavelength range.

Specifications of the in-vacuum undulator (IVU21)

Period (number of periods)	21.6 mm (92)
K (at minimum gap, 5.5 mm)	1.60
Photon source size (h \times v, FWHM)	$309\times18~\mu\text{m}^2$
Photon source div $(h \times v \in WHM)$	$112 \times 30 \mu rad^2$

...for many crystal types

- Small crystals can be studied by focusing the beam down to ~50×10 μm² (h×v) where the beam divergence is 0.5 ×0.1 mrad², allowing the study of macromolecules that crystallize only in small crystals,
- Any medium-sized crystals can be dealt with by defocusing the beam up to ~300×100 μm² (h×v),
 - Larger crystals can be completely exposed to the x-ray beam by unfocusing the beam to ~500×500 μm² (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.





The End station consists of two in-house developed translation/rotation tables that support the diffractometer and the detector, and to guarantee stability they sit on a granite base. The tables are adjustable in X,Y,Z to 1 μ m resolution and repeatability. An automatic sample changer stands on a nearby table for automatic sample mounting and allows easy access to manual sample mounting.





Automatic Sample changer



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 Status Call for tender issued Delivery End of 2010

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

