

MS44-2-8 The BM07-FIP2 beamline for macromolecular crystallography at ESRF
#MS44-2-8

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Abstract

BM07-FIP2 is available to users since October 1st, 2021. It is the follow-up of beamline FIP-BM30A [1] which ended operation in 2018 upon ESRF-EBS upgrade. FIP2 (French beamline for the Investigation of Proteins) is located on the Bending Magnet section 07 (BM07) of the ESRF. It is especially dedicated to crystallography of biological macromolecules. Its optics can deliver a focused beam on a fixed sample position in a large energy range (7-17 keV, to be extended to 5-25 keV). The beam size can be adjusted between 50 x 50 and 250 x 250 μm^2 . Data collection is controlled using the browser-based MXCuBE3 user interface, and sample (meta)data are handled and stored by ISPyB. FIP offers the following capabilities:

- High-throughput screening of cryocooled crystals
- (Multiwavelength) anomalous diffraction experiments
- In situ (in plate) crystal screening and data collection [2]
- On-line UV-Vis absorption microspectrophotometry [3]
- Humidity-controlled experiments [4]

References

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