

SAXS for biology optimized on the beamline SWING



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Abstract

SWING is a beamline devoted to (A)SAXS, WAXS and (A)GISAXS at SOLEIL, opened to all scientific domains with no exclusion. The source is a U20 in-vacuum undulator, thus providing a beam with small size and low divergence. The energy range is 5-17 keV (Si111 Double Crystal Monochromator) with an expected flux of about 10¹³ photons/s. Focusing of the beam onto the detector plane is achieved by bending two perpendicular mirrors (Kirckpatrick Baez configuration), thus being insensitive to a change in energy. The size of the focused beam should be typically 400 µm x 100 µm (HWHM). A large motorized table allows for fine positioning of sample environments. A 17 x 17 cm² CCD detector is positioned on a three axes translation stage within a large chamber under primary vacuum. The sample to detector distance ranges from 0.6 to 8 m. A friendly graphical interface written in Java allows for collection and online data reduction, including automated determination of radius of gyration, peak maximum intensity and width, etc. Absolute intensity measurements will be achieved by using reference samples and calibrated monitors. 4 1000 Despite its scientific versatility, a great effort has been made to provide biologists with optimized experimental conditions. An in vacuum thermostated capillary cell with online measurement of UV-Vis absorption will permit precise and reproducible measurements of proteins in solution. An HPLC system will be proposed for online purification. An automated solution sampler is presently under optimization for high throughput data collection. The complete biochemistry lab of SOLEIL, situated close to the beamline, is opened to those who need it. SWING is at the moment under assembly and commissioning. We expect to welcome our first external users by the end of 2007. The beam was focused at the detector position for the first time on July The white beam was delivered for the first time on 12th, just before the summer shutdown ! March, 15th, in the monochromator optics hutch Detector vacuum chamber (10-2 mbar) KB mirrors vacuum chambe (x,z):4 stops + 2 U20 brilliance of energy v 1001 WING a fixed gap, g=12 m compared nA, to ed with diod otion spectrum of a thin foil of Cu ne dge @ 8.979 keV, with mirrors inser Absorr its K-edge @ The monocl *han 0.2 eV after the b Focusing was straightforward in the vertical direction, while it still requires fine electrode HV tuning in the horizontal direction. The vertical width was checked to remain around 100 μ m whatever the position within the detector vacuum chamber. U A complete system of purification and automated solution sampler will be available for SAXS measurements on proteins in solution

