

The in crystallo optical spectroscopy (icOS) Lab

Towards time-resolved icOS (TR-icOS)

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Complementary spectroscopy techniques

The *ic*OS Lab, an IBS/ESRF platform, located at the ESRF (formerly known as the Cryobench)

• UV-visible abs/fluorescence/Raman spectra directly measured on crystals at cryogenic or room temperature





Why performing optical spectroscopy on crystals?

Optical spectroscopy used in complement to MX:

(1) To determine the **functional state** of the crystalline protein

(2) To evaluate the extent of specific radiation damage effects

(3) To perform kinetic crystallography experiments (structure determination of unstable species, in time or dose)

When and where?

- Before or after the diffraction experiment: **Offline setup**
- During the diffraction experiment: **Online setup**

Offline setup: the icOS Lab – Control Cabin and Experimental Hutch Now in Chartreuse Hall between new ID29 and ID30B



New automated spectroscopy setup



- Standard MX beamline equipment (minidiffractometer MD2M with 3-click sample centring, back/ front lights, on-axis viewer)
- Motorized optical objectives

on-line UV-vis absorption spectroscopy on BM07-FIP2 (large crystals, > 50-100 um)

- FIP2 = reconstruction of BM30A-FIP on BM07
- FIP2: 3x to 10x higher flux + Pilatus detector -> faster data collection -> faster time resolution
- Cryogenic (still very) low dose spectroscopic characterization -> complementary to other MX beamlines
- Importance of keeping a TopHat beam vs. Gaussian beam for radiation damage studies





McGeehan et al., J. Synchrotron Rad. (2009)

Future: on-line UV-vis absorption spectroscopy on ID30A-3 (smaller crystals, 15-20 um)

- Brand new design (O. Hignette/P. Theveneau, ESRF) use of parabolic mirrors
- Developed by David von Stetten and now Igor Melnikov





Future: on-line Raman spectroscopy on ID30B

660

640

- Online Raman not suitable for ID29 any more (beam size)
- Similar setup as ID29 (6tec/ESRF design), but re-designed to accommodate the MD2S (P. Jacquet, IBS)
- Radiation damage studies monitoring specific bond breakage or deformation of groups (or of secondary structures)







von Stetten et al., J. Struct. Biol. (2017)

Recent *ic*OS highlights

- Carpentier et al., Nucleic Acids Res. (2020) Structural, biochemical and functional analyses of the tRNAmonooxygenase enzyme MiaE
- Kovalev et al., Nat. Commun. (2020) Molecular mechanism of light-driven sodium pumping
- Kovalev et al., PNAS (2020) High-resolution structural insights into the heliorhodopsin family
- De Zitter et al., JACS (2020) Mechanistic Investigations of Green mEos4b Reveal a Dynamic Long-Lived Dark State
- Xu *et al., PNAS* (2020) Structural elements regulating the photochromicity in a cyanobacteriochrome



Microsecond time-resolved UV-vis absorption spectroscopy on *ic*OS

- Use with 10-15 um crystals
- Actinic pulse: nanoseconds (from a tunable ns laser)
- Time resolution of probe: microseconds (from a Xe flashlamp)



Feasibility experiment on bacteriorhodopsin crystals



Building the TR-icOS setup – hopefully working end of 2021



Preliminary results on bacteriorhodopsin crystals







Summary: various available spectroscopies, or soon-to-be

Spectroscopy	Off-line	On-line
UV-vis absorption	icOS	BM07-FIP2 (large beam > 50-100 um) ID30A-3 (MASSIF-3) (small beam ~15-20 um)
Fluorescence	icOS	
Raman	icOS	ID30B
Time-resolved UV-vis absorption (microsecond)	icOS	-

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Example of a time-resolved crystallography experiment on ID30A-3 aided by off-line time-resolved optical spectroscopy

Time-resolved crystallography setup on beamline ID30A-3



Crystals preservation at RT

HC1 humidity controller

Populate the crystal in light state



1 <u>100 µm</u>

Record oscillation data sets as fast as possible

MD2 Microdiffractometer

Eiger X 4M pixel detector





470 nm LED

Photoadduct build-up



Spectroscopic characterization of photoadduct build-up

Time-resolved UV-vis absorption spectroscopy at the *ic*OS Lab (offline mode), with **41 ms time resolution**



X-ray data collection principle



Molecular movie of the photoadduct population build-up in a LOV2 crystal

66 time points Cys426 One frame every 63 ms Slowed x2 **GIn489** FMN Phe470 0000 ms Aumonier et al., IUCrJ (2020)

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Diffraction-based calculation of the build-up time constant



Aumonier et al., IUCrJ (2020)



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