

## X-ray structure analysis of macromolecules

## X-ray station user form

In order to schedule experiment, this form must be filled with every user who intent to use the X-ray station.

Submit form to: Jiří Brynda, room 35 (JV, basement), tel.: 220 183 210, brynda@uochb.cas.cz

Name of project:	
Supervisor of project:	
Responsible scientist:	
Contact [email, phone]	
Source of protein:	
Protein MW:	
Ligands, cofactors, etc.	
Toxicity problems (specify if any):	
Infectivity risks (specify if any):	
Protein specific comments:	
Crystallization condition	
(list of all components):	
Crystal dimensions	
[mm x mm x mm]	
Space group:	
Lattice parameters:	
Summary of project	
(max. 3 sentences):	
Purpose of X row structural applying	
Purpose of X-ray structural analysis (max. 3 sentences):	
(max. 5 sentences).	
Analysis you plan to perform:	With which analysis you need assistance:
Measurement of complete dataset	Measurement of complete dataset 1
Data processing	Data processing
Structure solution	Structure solution
	Structure deposition
	Structure analysis

Red items are obligatory.

Green items are only possible on the basis of close collaboration.