

**QUESTIONNAIRE TO BE FILLED OUT FOR EACH PROTEIN TO BE BROUGHT
BY PARTICIPANTS OF THE ISBio 2023 COURSE**

Data of the participant	
Participant name:	
Institution:	
Address:	
Phone:	
Fax:	
Email:	

Data of the proposed macromolecule	
Name of the macromolecule	
Source:	
Biological function	
Molecular weight	
Status of oligomerization	
Iso-electric point	
Thermal stability	
Influence of ageing on the stability of the solution.	
Sample data	
Protein buffer:	
Protein concentration:	
Storage conditions:	
Additives:	
Notes:	

Please, select below the preferred methodologies:

Methodologies	
NMR *	
(0,5mL of protein sample with a concentration of 10-20 μ M and 0,5mL of its ligand with a concentration of aprox. 5mM (in DMSO or buffer) are needed to perform ligand-based interaction studies)	
Protein Crystallization **	
(0,5 to 2mg of protein needed for performing a crystallization screening)	
X-ray diffraction **	
(protein single crystals)	
ITC ***	
(2,5mL of protein sample with a concentration of 10-50 μ M and 0,8mL of its ligand with a concentration of 0,1-0,5mM are needed to perform one titration)	
CryoEM ****	

Useful information to attach:

SDS-PAGE, Native or IEF gel of the protein sample, ELISA and other analysis relevant to the study of the interaction

To send samples by mail, please use the following address[§]:

Cristalografia de Raios-X (lab. 6.05 / 6.07)
Departamento de Quimica
Faculdade de Ciências e Tecnologia Universidade Nova de Lisboa
2829-516 Caparica, Portugal

VERY IMPORTANT: Before deciding to ship any samples, please:

* contact Eurico Cabrita (ejc@fct.unl.pt)

** contact Márcia Correia (marcia.correia@fct.unl.pt) and provide information on harvesting and cryo-conditions (or bring pre-prepared solutions) and/or any previous information regarding crystal characterization

*** contact Benedita Pinheiro (b.pinheiro@fct.unl.pt)

**** contact Hartmut Luecke (hudel@fct.unl.pt)

§ inform each responsible of shipping (including tracking number) and if you have any questions