**FIRST ATTEMPT CRYOEM SESSION**

* Name of the PI and the scientist involved:
* Sample name and project identifier:
* Sample components (protein/nucleic acids/ligands):
* Molecular weight (kDa):
* Sample preparation date and storage conditions:
* Concentration (µM) and method of measuring concentration:
* Sample volume and buffer composition:

Biophysical characterization

SDS-PAGE with coomassie stain Gel Filtration profile Mass Photometry profile

INFORMATION

* Sample size:

Please bear in mind that those samples with molecular weights below 150 kDa would require significant efforts for structure determination, in particular, when the goal is to achieve a high-resolution structure.