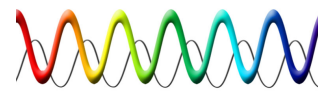


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Order form - SAXS measurements

Name of user

Address (institute, working group)

Phone-no., email

I (budget responsible person) recognized the „terms of use“ and „charges and fees“ and agree on these (<https://www.css.hhu.de/en/translate-to-english-kontakt-und-service>).

All users of the CSS must acknowledge the DFG in their publications:

"The Center for Structural Studies is part of StrukturaLINK Rhein-Ruhr which is funded by the Deutsche Forschungsgemeinschaft (DFG Grant number 573727698) ." For X-ray diffraction data please mention also „INST 208/740-1 FUGG“ (X-ray diffractometer), for SAXS data „INST 208/761-1 FUGG“ (SAXS machine).

Date , signature (budget responsible person)

Name and/or acronym of the protein:

Source organism: Safety level: S1 S2

Expression organism: Safety level: S1 S2

soluble protein

membrane protein

Date of purification:

Is there a tag fused to the protein?

no

yes (which tag, N- or C-terminal?)

signal peptide?

modifications? (e. g. phosphorylation):

Total no. of residues (incl. tag):

Store/stable at [°C]:

Exact size of the protein [kDa]:

Extinction coefficient:

Oligomeric state, symmetry:

Protein concentration [mg/ml]:

Total volume of protein [μl]:

Composition of protein buffer:

SEC column:

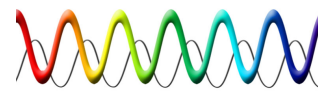
Enclosed substrate(s) (name, concentration, K_d value):

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Short sample description of the protein (function, catalyzed reaction, ...):

Enclosed files/analyses/solutions (**obligatory**):

3 ml protein buffer (original)
protein sequence incl. tag (electronical file to be send
via email)

Is the above mentioned project (protein) **funded
by third party**? If so, please give the grant and
the corresponding grant number (**obligatory**):

Additional files/analyses/info:

SDS-Page
gel filtration
structure or homology model
other