**Customer Details:**

Name:Click or tap here to enter text.

Group:Click or tap here to enter text.

Email:Click or tap here to enter text.

Phone:Click or tap here to enter text.

**Details on sample(s):**

Nature of sample: Choose an item.

Organism (e.g. Human, Mouse, etc.): Click or tap here to enter text.

Expression host (e.g. E.coli, S.cerevisae, Human, etc.): Click or tap here to enter text.

Molecular weight (kDa): Click or tap here to enter text.

Estimated amount/concentration (µg/µl): Click or tap here to enter text.

Are there any detergents (even low amounts count) in your sample?

Yes (Please indicate which in buffer composition)

No!

Buffer composition: Click or tap here to enter text.

pH of buffer: acidic neutral basic

Do you have a gel image? (if so, please send it to us) Yes No

If yes, have all bands for analysis been assigned in the gel image!

(Tick for 'Yes', if not, please do so!)  Yes

**Additional information:** (e.g. expected contaminants, purification history, fusion proteins, known PTMs, truncation, mutation, etc., if you have nothing to mention here, please write NA)

Click or tap here to enter text.

Please paste here the protein sequence(s) incl. all tags, mutations, etc. in FASTA format [Uniprot] (If not applicable, write NA):

Click or tap here to enter text.

**Type of analysis:** Choose an item.

**Other details:**

Describe what you want to know about your sample: (default protease is Trypsin (K,R, if you need a different one, please write this here):

Click or tap here to enter text.

**Data analysis:**

For quantitative experiments (e.g. TMT, iBAQ, LFQ, Dimethyl, SILAC): Please indicate here which ratio comparisons you would like to have in the end and in which order you want to have them in the plots. State the controls, please. For IP’s, name the bait protein here.

Click or tap here to enter text.