QconCAT inquiry form

Dear valued customer,

thank you very much for your request.

To receive an appropriate quotation, please answer the following questions to the best of your knowledge and return the order inquiry form to PolyQuant via E-mail (info@polyquant.com) or fax (+49-9405-96999-28). Any information you provide will be treated with the highest confidentiality independent of a CDA[3]. If you have any questions, please do not hesitate to contact us.

For internal use:	Name	
	Position	
	Company/Institute	
	Street Address	
	Zip Code	
	City	
	Country	
	Telephone/Fax	
	E-mail	
	VAT-ID	

1) General information:

Source (organism/matrix):

(e.g. bacteria, yeast, cultured cells,
 mouse/kidney, human/plasma, plant/leaves etc)

Number of proteins to quantify:

Total number of peptides for quantification:

MS-data for peptides available[1]: ☐ Yes ☐ No   [1](the peptides have been detected previously e.g. by the customer)

Peptide selection by: ☐ Customer ☐ PolyQuant ☐ Customer/PolyQuant

MTA[2] necessary: ☐ Yes ☐ No   [2]Material transfer agreement

CDA[3] necessary: ☐ Yes ☐ No   [3]Confidentiality disclosure agreement

The QconCAT(s) will be used for: ☐ Exploratory/Discovery Analysis

☐ High-throughput analysis

2) Gene construct information

Gene construct exists: ☐ Yes   ☐ No

Select yes if gene construct exists and continue with 3)
Select no if gene construct should be designed by PolyQuant

Protein Optimization: ☐ Yes   ☐ No

Select yes if PolyQuant shall shuffle peptides to optimize the properties of the QconCAT
Select no if the order of the peptides needs to remain as provided by the customer

3) Expression specific information

Labeling option: ☐ unlabeled ☐ 13C Arg/Lys ☐ 13C, 15N Arg/Lys ☐ 15N

other (please specify): [   ]

Purity: ☐ crude extract

☐ IMAC purified (6 M GdnHCl)

☐ IMAC purified (8 M Urea)

☐ IMAC purified, dialysed   requires additional buffer screen to identify suitable storage buffer

4) Other services

☐ Absolute protein quantification to be performed by PolyQuant

☐ Discovery proteomics to identify peptides suitable for quantification by PolyQuant

☐ Development of protein extraction protocol by PolyQuant

Comment: [   ]