

# Order form for mass spectrometry

## Customer address:

Company/Institute:	<input type="text"/>	Contact:	<input type="text"/>
Department:	<input type="text"/>	Phone:	<input type="text"/>
Street:	<input type="text"/>	Fax:	<input type="text"/>
Zip and city:	<input type="text"/>	Country:	<input type="text"/>
E-mail:	<input type="text"/>	Language:	<input type="text"/>

## Billing address (if different):

Accounting office:	<input type="text"/>		
Street:	<input type="text"/>	Shipping date:	<input type="text"/>
Zip and city:	<input type="text"/>	Your order no.:	<input type="text"/>
Country:	<input type="text"/>	Your signature:	<input type="text"/>

## Sample information:

Number of samples:	<input type="text"/>	<input type="text"/>
Sample name(s):	<input type="text"/>	<input type="checkbox"/> External sheet attached
State:	<input type="radio"/> In gel <input type="radio"/> Solution <input type="radio"/> Dry <input type="radio"/> Blot	
Storage condition:	<input type="radio"/> + 4°C <input type="radio"/> - 20°C <input type="radio"/> - 80°C <input type="radio"/> Room temperature	
Protein amount/concentration:	<input type="text"/>	
Protein modification (e.g. alkylation):	<input type="text"/>	
Buffer (conc, pH):	<input type="text"/>	
More information:	<input type="text"/>	
Hazard class:	<input type="radio"/> infectious (e.g. patient material) <input type="radio"/> cytotoxic or hormone	
If other, please specify:	<input type="text"/>	

**Please note, we don't process radioactive samples!**

Species:	<input type="radio"/> Homo sapiens <input type="radio"/> Mus musculus <input type="radio"/> Other	<input type="text"/>
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Sequence of protein sample known?	<input type="radio"/> no <input type="radio"/> yes	<input type="checkbox"/> external sheet attached
If yes, please provide with sequence information:		

Database ref. no. (e.g. UniProt):	<input type="text"/>
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## Requested analysis:

Aim:

e.g.:

- Protein identification
- Mass determination
- Modification analysis
- Sequence analysis etc.

Protein identification

Enzyme:

Trypsin

Glu-C

Lys-C

Asp-N

Arg-C

Other(s)

Instrument:

MALDI-TOF MS (**PMF**)

MALDI-TOF-MS/MS

nanoLC-ESI-MS/MS

Mass determination

MALDI-TOF MS

nanoLC-ESI-MS

ESI-MS

Quantification

label free, emPAI (rough estimate)

iTRAQ

Isotopic labelling

MRM Analysis

ICPL

## Reporting:

Report language:  English  German (**default**)

Type:  digital version only (default)  additional hardcopy by mail

Additional recipients:

E-mail:

Raw data:

To be kept at TOPLAB (standard process). Exceptions only with additional agreement.

## Remaining sample material:

- destroyed 3 months after end of the project (**default**)  returned at customers' expense  stored until further agreement

**Information:** General terms and conditions apply (<http://www.toplab.de/Downloads.html>).