Biomolecular Chemistry / Mass Spectrometry

Order Form for MS Analysis Name: Group: Institute: Street: Postal Code: Town: Email: Please send the gel and a picture of the gel with the bands/spots of interest labeled How many samples should we analyse: From which organism ist the sample: Human Yeast E. coli Drosophila **Others** Which method should we use: Protein identification by PMF (MALDI-TOF MS) Protein identification by peptide sequencing (LC-ESI-MS/MS) De Novo sequencing by ESI-MS/MS Modifications Mass determination of proteins/peptides Send the gel to:

ZMBH Biomolekulare Chemie Im Neuenheimer Feld 282 69120 Heidelberg