

# MASS SPECTROMETRY RESEARCH FACILITY

CRL Department of Chemistry  
University Of Oxford  
Mansfield Road, Oxford  
Telephone: (01865 275942)  
e-mail: [elisabete.pires@chem.ox.ac.uk](mailto:elisabete.pires@chem.ox.ac.uk)

## Proteomics Submission Form

### Contact Information

Name: \_\_\_\_\_ Department: \_\_\_\_\_  
Research Group: \_\_\_\_\_ Email: \_\_\_\_\_ Telephone: \_\_\_\_\_  
Project Code/Charge Account \_\_\_\_\_ Date: \_\_\_\_\_

### Sample Information

Number of samples: \_\_\_\_\_  
Sample ID( multiple samples can be submitted here) \_\_\_\_\_  
Sample type: Tryptic digest\*  Intact Protein  Peptide   
Sample Volume supplied: \_\_\_\_\_ Sample concentration: \_\_\_\_\_  
Sample molecular weight: \_\_\_\_\_  
Sample buffer/solvent: \_\_\_\_\_  
Sample buffer concentration: \_\_\_\_\_  
Sample Purification: ZipTip  C18 spin Columns  SepPak   
Amino acid sequence (if known): \_\_\_\_\_  
Specific modifications (if known): \_\_\_\_\_  
Organism\*: \_\_\_\_\_  
Enzyme used for digestion\*: \_\_\_\_\_  
Protein name\* / Uniprot code\*: \_\_\_\_\_

### Experiment Information for Triptic Digestion performed in the CRL-Mass Spectrometry Lab

Sample Submitted as: Entire Gel  Gel Band  Solution  Pellet   
Gel Stain Type: \_\_\_\_\_ Protein Concentration: \_\_\_\_\_  
Sample buffer/solvent: \_\_\_\_\_

### MS Analysis required

nanoLC-ELITE (CID)  nanoLC-ELITE (ETD)  MALDI -TOF/TOF  LC-MS  Native MS/Synapt

\*Just in case of Tryptic digest samples

**Note:** For collaborative proteomics projects we would normally expect to analyse samples and perform data processing and analysis followed by a meeting to discuss results. We will provide the raw data files upon request. We aim to send results back to you within two - three weeks after sample submission but technical issues can cause delays. Please contact Elisabete directly for an update if results have not been communicated by three weeks.