

Protein & Peptide Sample Submission Form
Mass Spectrometry & Proteomic Core Facility, RGCB

Submitted By:

Submission Date:

Email:

Phone:

Sample details:

Number of samples:

Sample ID(s):

Concentration:

Solvents/buffer:

Expected mass (Da):

Molecular formula:

Storage condition (RT, 4°C, -20°C):

Hazards/toxicity:

Organism (species):

Source of sample:

Nature of samples:

Protein

Peptide

Others (specify):

Gels stained with: Coomassie

Silver

Others (specify)

Protein:

Pure protein

Mixture of proteins

Fusion protein

Analysis Requirement(s):

Molecular weight confirmation / Accurate Mass determination

peptide sequencing (MS or MS/MS)

Peptide Mass Fingerprinting (MS)

Protein Identification (MS/MS)

Relative protein quantification or expression (label-free)

Analysis Method:

ESI (MS or MS/MS)

MALDI (MS or MS/MS)

Others (specify)

Funding source for payment (Analyses cost for RGCB staff):

Project Investigator (Name & Signature):

Date Received:	Date Completed:	Analyst:
Time spent	Result Delivered	

- Samples without filled submission forms & without the signature of the PI will not be accepted.
- At the conclusion of analysis, all extra samples will be returned to the clients only if such request is made at the time of sample submission. The proteomics facility will not store samples more than 2 weeks once the results are given to the clients.
- At the conclusion of a project, data given to clients will include all raw data files along with processed data files and report. Proteomics facility will not keep the results and raw data files.