

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)

In-gel digestion

In-solution digestion

Relative protein quantification or expression (label-free)

Analysis Method:

ESI (MS or MS/MS)

MALDI (MS or MS/MS)

Others (specify)

Project Investigator (Name & Signature):

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

- Samples without completed submission forms & without the signature of the PI will not be accepted.
- Include a gel picture with proper numbering of bands/spots, if the samples are from gels
- 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 long term for clients.
- At the conclusion of a project, data given to clients will include all raw data files along with processed data files and report. It will be the sole responsibility of the client to keep and store these files in a safe place.