UF Scripps | **Biomedical Research**
UNIVERSITY *of* FLORIDA

PROTEOMICS CORE
130 Scripps Way, B147
Jupiter, Florida 33458
(561) 228-2360

PROTEOMICS SAMPLE SUMBISSION FORM

Job No. _____
(provided by Core personnel)

Proteomics Core Sample Information Instructions

In order to serve you better, please fill out as much information as possible regarding your sample(s). Also note any special sample instructions or analysis instructions on the form. The more information we have, the better the results and the faster we can return data. A completed sample submission form will be required before we can start working on samples. If you are a new user of the Core, we advise that you schedule a meeting with staff to discuss your project, goals, expectations, sample preparation and data analysis. Email the completed form to George Tsaprailis (Gtsaprailis@ufl.edu) and to Cathy Scharager (c.scharagertapia@ufl.edu), or print it and drop it off with your samples.

Date Submitted: _____

User Name: _____

User Email: _____

PI Name: _____

PI Account #: _____ (Required to initiate service)

Estimated Cost: _____ (Based on service units below)

PI/Approval Signature _____   Date: _____
(Required to initiate service)

The PI agrees to the transfer of the final cost from the account provide above to the Proteomics Core account by providing an electronic (or signed) signature. The PI may also designate members of their laboratory to approve invoices. Charges will be made once analysis is completed and the results are provided. Results will be provided via a Dropbox link to a secure folder.

_____

PROTEOMICS CORE INVOICE

Job No. _____

Final Cost: _____

Date Submitted: _____

Date Completed: _____

Core personnel Signature: _____

A copy of this invoice will be provided to Ms. Mayra Caramazana (mcaramazana@ufl.edu) for internal billing.

PROTEOMICS SAMPLE INFORMATION

Goal of experiment and comments (ex., general protein ID, PTMs, quantification, 1D SDS-PAGE, protein determination) – to be filled out by user

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Sample type: _____

No of samples: _____

Sample name(s):   _____   _____
                      _____   _____
                      _____   _____
                      _____   _____

Concentration
and volume: _____   _____
(N/A if unknown)   _____   _____
                      _____   _____
                      _____   _____

Target Protein MW   _____   _____
(if applicable)   _____   _____
                      _____   _____
                      _____   _____

Gel Picture: Yes (attach gel picture hardcopy to request when dropping off samples)
                      No

Organism type: _____

Job No. _____

Proteomics Services Requested

Services Requested	Quantity	Rate	Total
Basic Protein Services			
Mini gel electrophoresis			
Mini gel Coomassie staining			
Protein/Peptide assay			
Protein precipitation			
LC-MS/MS Analysis on Fusion (incl. digestion, LC-MS/MS and database search)			
High Resolution 1HR (eg. simple protein)			
High Resolution 2HR (eg. IP)			
High Resolution 3HR (eg. IP)			
High Resolution 4HR (eg. complex proteome)			
LC-MS/MS Analysis on Q Exactive (incl. digestion, LC-MS/MS and database search)			
High Resolution 1HR (eg. simple protein)			
High Resolution 2HR (eg. IP)			
High Resolution 3HR (eg. IP)			
High Resolution 4HR (eg. complex proteome)			
LC-MS/MS Analysis on Fusion of self-digested and cleaned-up sample (LC-MS/MS and database search)			
High Resolution 1HR (eg. simple protein)			
High Resolution 2HR (eg. IP)			
High Resolution 3HR (eg. IP)			
High Resolution 4HR (eg. complex proteome)			
LC-MS/MS Analysis on Q Exactive of self-digested and cleaned-up sample (LC-MS/MS and database search)			
High Resolution 1HR (eg. simple protein)			
High Resolution 2HR (eg. IP)			
High Resolution 3HR (eg. IP)			
High Resolution 4HR (eg. complex proteome)			
Other Services			
Sample clean-up (each LC-MS/MS unit requires 1)			
TMT protein quantification (quote to be provided)		NA	
HR MS (or MS/MS) (Q Exactive)			
Other (cost to be determined by Core personnel)			
TOTAL			

Additional data mining/bioinformatics provided by Dr. Gogce Crynen at hourly rates (inquire at GCrynen@scripps.edu)

Disclaimers:

Long term storage of the RAW files and any data mining results is the sole responsibility of the user. The Proteomics Core will however, backup files to a Scripps Dropbox account as long as the institution maintains a license for Dropbox.

Job No. _____

PROTEOMICS CORE NOTES

Job No. <mark>_____</mark>