**Proteomics & MS Core Facility**, WHA Room-4303, 750 E. Adams Street, Syracuse, NY 13210

Phone: 315-464-4563 Email: kakhniad@upstate.edu Web: www.upstate.edu/proteomics/

# Sample Submission/Service Request Form

**CUSTOMER / PAYMENT (**Upstate Investigators)

Contact Name PI Name

Department Building/room

Phone Email

State Account #

RF Project/Task/Award #

 / /

\*Signature of PI Date

\*- Samples will be processed when this requisition has an account number and PI signature

**REQUESTED LTQ-Orbitrap Based SERVICES**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Service Type | Service Code | Number of Units | Unit Price,$ | Setup Price,$ | Full Price\*,$ |
| Trypsin Digestion | 100 |  | 15.00 | 35.00 |  |
| MW Determination | 200 |  | 50.00 | -- |  |
| Protein Identification | 300 |  | 60.00 | -- |  |
| 301 |  | 108.00 | -- |  |
| 302 |  | 180.00 | -- |  |
| Mapping of Specified PTMs\*\* | 400 |  |  60.00 | -- |  |
| 401 |  |  108.00 | -- |  |
| 402 |  | 180.00 | -- |  |
| Protein Profiling cICAT, iTRAQ, or SILAC Analysis | 500 |  | 200.00 | -- |  |
| 501 |  | 150.00 | -- |  |
| 502 |  | 350.00 | -- |  |
| 504 |  | 650.00 | -- |  |
| 508 |  | 1200.00 | -- |  |
| Protein Profiling 2-D DIGE Analysis | 600 |  | 600.00 | -- |  |
| 601 |  | 200.00 | -- |  |
| 602 |  | 450.00 | -- |  |
| 603 |  | 123.00 | 35.00 |  |
| Small MoleculeQuantification | 700 |  | 17.00 | -- |  |
| 701 |  | **3**0.00 | -- |  |
| Other | 800 |  |  |  |  |
| Total Cost | -- | -- | -- | -- |  |

\*-Full price = (Number of units) x (Unit Price) + (Setup Price)

\*\*- Add additional $10.00 per PTM to the full price

## SAMPLE DETAILS: see page 2

*Please notify us of any publications and grants utilizing the results of these services*

## Office Use Only

Core Account Detail Code: PROT Account Number: 90020800

Order Accepted (Date): Order Completed (Date):

**SAMPLE DETAILS**

Please indicate the sample origin/species (human, bovine, mouse, rodent, E.coli, yeast, etc), the accurate (or ~estimated) protein amount in the gel slice/liquid samples (pmol, ug, coomassie or Sypro Ruby stain, etc), the protein concentration in the liquid samples (uM, ug/ul, etc), and MW of target proteins when known

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| ## | Sample Name | Sample Origin | Protein Amount  | ProteinConcentration | MW |
| 1 |  |  |  |  |  |
| 2 |  |  |  |  |  |
| 3 |  |  |  |  |  |
| 4 |  |  |  |  |  |
| 5 |  |  |  |  |  |
| 6 |  |  |  |  |  |
| 7 |  |  |  |  |  |
| 8 |  |  |  |  |  |

**Additional Sample Infofmation**

Gel Stain: □ Coomassie □ Sypro Ruby

Is Sample Digested? □ No □ Yes, Enzyme:

Liquid Sample Buffer:

**Sample Comments:**