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**Center for Advanced Proteomics Research (CAPR)
 Service Order Form (F004)**
(We do not have service for radioactive samples at present time)

Group: _____ Name: _____
 Address: _____ Date: _____
 _____ Tel: _____
 _____ Fax: _____
 _____ E-mail: _____

Rutgers Project No: _____
 PO Billing Address: _____

Code	Service	Internal Price	Unit	Qty.	Total
S1	Protein Mixture Identification by LC/MS/MS	\$425	each		
S2	Protein Mixture Identification by LC/MS/MS (bulk rate for up to 10 samples submitted together)	\$3,300	each		
S3	Protein Mass Determination	\$300	each		
S4	Full SDS-PAGE Lane (mini-gel) Digestion	\$100	hour		
S5	Sample Clean Up	\$85	each		
S6-8	iTRAQ (8 plex) Proteomics	\$6,050	each		
S7-6	Iodo TMT (6 plex) Proteomics	\$4,500	each		
S8	TMT (16 plex) Proteomics	\$8,000	each		
S9	Data Sharing	\$500	each		
S10	Custom Service	\$150	hour		
PI Signature*		Date			
Print PI Name					

***As per Rutgers guidelines, PI must sign the service order form.**
Additional Information:

Specific objectives of the analysis:

Instructions on remaining sample after analysis:

**Sample buffer composition attached: Yes / No If
No, please specify:**

Sample species (e.g. *Homo sapiens*): _____

Sample Storage Condition: _____

Sample Post-translational modification:

Sample Protein/Peptide known structure:

Sample Protein/Peptide Quantity: _____

Sample Protein/Peptide Concentration: _____

Method used to measure Protein/Peptide Concentration (e.g. BCA, Bradford etc.):

Interested in molecular weight (MW) range:

Interested in pH range: _____

Expected sequence (for MS sequencing only):

Sample ID:

Sample #1: _____

Sample #2: _____

Sample #3: _____

Sample #4: _____

Sample #5: _____

Sample #6: _____

Sample #7: _____

Sample #8: _____

Sample #9: _____