RUTGERS
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Center for Advanced Proteomics Research (CAPR)
Service Order Form (F003)

(We do not have service for radioactive samples at present time)

Group:______________________________ Name:______________________
Address:____________________________ Date:_______________________
____________________________________ Tel:________________________
____________________________________ Fax:________________________
____________________________________ E-mail:_____________________

Rutgers Index No.:__________________ Non-Rutgers PO No.:__________
PO Billing Address:____
____
____

Code	**Service**	**Unit Price**	**Unit**	**Qty.**	**Total**
S002	Low Level Protein Identification by MALDI-TOF/TOF	$150.00	each		
S003	Protein Mixture Identification by LC/MS/MS(LTQ Orbitrap Velos)	$275.00	each		
S003-B	Protein Mixture Identification by LC/MS/MS(LTQ Orbitrap Velos) (bulk rate for up to 10 samples)	$2000.00	each		
S005	Extra Protein Identification Data Analysis Time (over the default 2 hours)	$100.00	hour		
S006	Peptide Mass Determination by MALDI-TOF	$50.00	each		
S007	Protein Mass Determination by LTQ Orbitrap Velos	$150.00	each		
S012	BioRad Criterion 2D Gel (11cm)	$160.00	gel		
S013	Sypro Ruby Staining and Typhoon Gel Imaging	$50.00	gel		
S014	Extra Gel Data Analysis Time (over the default 1 hour/gel)	$75.00	hour		
S015	Sample Clean Up	$50.00	each		
S016	Custom Service	$100.00	hour		
S017	Protein Identification by MALDI (bulk rate up to 100)	$5000.00	each		
S018-4	iTRAQ (4 plex) Proteomics	$4000.00	each		
S018-8	iTRAQ (8 plex) Proteomics	$5000.00	each		
S019	DIGE Gel and Imaging (3 plex)	$1500.00	each		
S021	Consultation with CAPR staff	$75.00	hour		
S022	Instrument Training	$100.00	hour		
S023	Consultation with CAPR director	$100.00	each		
S025	User Operation of Typhoon Imager	$40.00	hour		
S026-1	ICAT (One heavy/light pair)	$1500.00	each		
S026-3	ICAT (Three heavy/light pair) with data analysis	$4500.00	each		
S028-1	Protein Mixture Identification by 2D LC/MS/MS (MALDI)	$3000.00	each		
S028-2	Protein Mixture Identification by 2D LC/MS/MS (LTQ Orbitrap Velos)	$3000.00	each		
PI Signature*	**Date**	**Total Charge**			
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:____

Sample #10:____