# **National Institute of Plant Genome Research**

Aruna Asaf Ali Marg, P.O. Box No. 10531, New Delhi – 110067.

#### **Indent Form for Proteomics Facility (Internal Users)**

ndent No						Date:	
S.No.	Type of analysis	Service code pricelist)	e (in R	nte	Quantity	Amount	
1							
2							
3							
4							
	•	GR / Project			Total		
1. 2. 3.	Proteomic Facility publication ofdata	abide by the safety , NIPGR shall not t ds are available und	take responsibil	ity on the ana			
1)	Indentor Name, Signature)		denting PI e, Signature)		Proteomic (Name, Sig		
Rupees	i						
•		c.) Rs					
Total E	xpenditure	•••••		Payı	ment terms		
Recom	mendation of the St	anding Purchase Co	ommittee				
	Dr. Manoj Maj Chairman	iee	Dr. Saloni M Member		Dr. Jyothi	ilakshmi Vadasser Member	
		Vineeta Sharma Member		udhir Patwal Member			
comme	endation of the Stand	ding Purchase Com	mittee is submi	ted for appro	val.		

P.S.O. COA Director

# **National Institute of Plant Genome Research**

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#### **Sample Submission Form for Proteomics Facility (Internal Users)**

Name of the Use	er: Date:	
Name of the Fac	culty/Scientist:	
Lab No:	Contact No:	
Email ID:		
Service Code: _		
	Sample Information	
Organism (speci	ries):	
Source of sampl	le (Tissue/Organ/Organelle):	
Sample complex	xity: 1D digest/ 2D digest/ Pull down/ whole cell digest	
Protein:	Pure protein ☐ Mixture of proteins ☐	
Gels stained wit	th: Coomassie □ Silver □	
Chemicals used	for reduction/ alkylation:	
Enzyme used: _		
Number of Sam	ples:	
	red (Please tick):	
	ation/ Peptide Mass Fingerprinting (PMF)/ Quantitation (iTr Label Free)/ PTM analysis/ Molecular weight confirmation.	raq / SILAC /
For PTM analys	sis:	
Known and poss	sible protein modifications:	
Site of PTM (if l	known):	
Additional infor		

### Sample details (Please fill all the required sections):

	wher of samples, places attach an extra sheet is similar

<sup>\*</sup>For more number of samples, please attach an extra sheet in similar format.

Pre-submission Checklist: (Please tick)					
☐ The stain used is compatible with mass spectrometry.					
☐ Photo of the stained gel attached.					
☐ The samples are free of particles or debris.					
☐ The samples are free of detergents or other chemical interferences.					
☐ High-abundant proteins, such as RuBisco: Depleted ☐ Not depleted ☐					
Please indicate composition of solution (including any salts, etc.):					
<b>Undertaking</b>					
<ol> <li>I understand that all Core Facility services are for research purposes only.</li> <li>I have read and understood the sample digestion guidelines provided by the facility and prepared the samples accordingly.</li> </ol>					
3. I understand that the presence of high-abundance proteins (e.g. RuBisco) and/or protein reagents (e.g. antibody, protein A/G) in the sample will affect the results.					
4. I understand that limitations apply to the detection of post-translational modifications (PTMs), and I have indicated on this form all modifications that are relevant to this analysis.					
5. I/We shall not claim for any damage/harm to my samples submitted for the analysis by Facility's equipment.					
6. I/We shall give due acknowledgement of Proteomics Facility, NIPGR in publications as mentioned below and also inform the facility about the publications coming out of the work done at the facility. Proteomics Facility shall not take any responsibility about the analysis, interpretation and publication of data acquired using equipment at this facility.					
7. I/We undertake to mention in any publication using data generated in the facility by stating in the Acknowledgment. Statement for Acknowledgement: "We thank the Proteomics Facility, NIPGR funded by Department of Biotechnology, Government of India for protein identification and mass spectrometry analysis used in the manuscript".					
8. I/ We take responsibility to pay user charges as per norms of facility/institute.  (Signature of the User) (Signature of the Scientist with Stamp)					
(FOR INTERNAL USE ONLY)					
Purchase Order No:					