

*Institute of Bioinformatics (IOB) (<http://www.ibioinformatics.org>) is a premier center for Proteomics and Computational Biology. IOB has also established a state-of-the-art proteomics facility to discover and validate candidate biomarkers in human diseases including neurological disorders, infectious diseases and cancers.*

Scientists at IOB will share their expertise in using high-resolution proteomic technologies to highly motivated young researchers in India through a series of workshops. We have successfully conducted four workshops on high resolution mass spectrometry - based proteomics. Several faculties and students from reputed National Institutes and Universities from India participated in these workshops (<http://www.ibioinformatics.org/workshop.php>).

This 4-day long intensive workshop will offer a theoretical and practical understanding of high-resolution mass spectrometry-based proteomic technologies and their application to clinical and biological research. Each participant will get hands on training for sample prep and mass spectrometry-derived proteomic data analysis.

### Eligibility

Open to students and researchers from academia and industry. Applications must include a statement of purpose (<100 words) along with their CV.

### Registration

Visit [http://www.ibioinformatics.org/registration\\_proteomics.php](http://www.ibioinformatics.org/registration_proteomics.php) to register for the workshop. Participants will be selected on first-come, first-served basis. Participants can also reserve their slots by directly contacting the organizers.

Registration fee:

Participants from Academia - Rs. 20,000

Participants from Industry - Rs. 30,000

A discount of Rs.5000/- will be provided to the participants of a previous workshop at IOB.

Refreshments and lunch will be provided throughout the day.

Applicants can contact organizers for accommodation with an additional cost (Single occupancy Rs.1100/day; Deluxe sharing: Rs.800/day; inclusive of taxes).

### Focus of the Workshop

- Principles of mass spectrometry
- Sample preparation strategies
- Fractionation techniques
- Depletion/enrichment strategies
- LC-MS/MS analysis
- Operation of LTQ-Orbitrap Velos mass spectrometer
- Proteomic data analysis
- Quantitative proteomics

### Resource Personnel

- Dr. T. S. Keshava Prasad
- Dr. Nandini A. Sahasrabudde
- Dr. S. K. Shankar
- Dr. Ravi Sirdeshmukh
- Dr. Aditi Chatterjee
- Dr. Ramu Gundimeda
- Dr. Rajesh Raju

## Workshop Program

### Day 1: Tutorial Lectures

Mass spectrometry-based proteomics  
Quantitative proteomics

### Hands on training

Sample prep I: In-gel and in-solution digestion  
Fractionation strategies - SCX/bRPLC

### Day 3: Tutorial Lectures

Analysis of post-translational modifications  
Operation of LC-MS/MS (demo)  
Multiple reaction monitoring (demo)

### Hands on training

Data analysis – quantitative proteomics  
Phosphopeptide enrichment

### Day 2: Tutorial Lectures

Introduction to mass spectrometry data analysis

### Hands on training

Sample prep II - peptide extraction and iTRAQ labeling  
Fractionation: depletion strategies  
Introduction to Mass spectrometry data analysis

### Day 4: Tutorial Lectures

Multipronged proteomic analysis  
Biological interpretation of proteomics data (demo)  
Introduction to metabolomics

### Hands on training

Analysis of phosphoproteomic data  
Biological interpretation of proteomics data

### Contact

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