

Institute of Bioinformatics

Discoverer Building, 7th Floor International Tech Park

Whitefield, Bangalore - 560 066

Karnataka, India.

Ph: +91 8028416140

We received a quotation from one company for the following instrumentation. If there are any other vendors who would like to provide quotations, please do so by sending the quotation in a sealed envelope on or before November 10, 2015 at 4 PM IST

No. IOB/001/2015

Date: 01.11.2015

Quotations are invited for the following equipment.

Sl. no	Particulars	Description
1.	Name of the equipment	GelFree Fractionation system
2.	Quantity	One
3.	Specifications	Details enclosed as Annexure I

The last date for submission of the quotation is 10.11.2015,

to <u>keshav@ibioinformatics.org</u>; santosh.shiri@ibioinformatics.org

Sd/-

Keshava Prasad, Ph.D.

Faculty Scientist

Institute of Bioinformatics

Annexure I

GelFree Fractionation system		
Instrument specifications	The instrument should enable molecular weight-based fractionation of intact proteins with liquid phase recovery	
Interface	The instrument should contain a programmable control module that will enable analysis of eight samples in parallel within 90 minutes	
Analytical capability	The instrument should have the capability to separate analytes using gel free methods to isolate and enrich user-selected molecular weight fractions for targeted protein quantification using LC-MS/MS, isolate intact proteins to analyze variants, posttranslational modifications, alterations, separate protein pull-down components for target protein purification	
System performance		
	The instrument should contain eight independent channels for molecular weight fractionation and liquid phase recovery. The channels should consist of a precision-cast gel column surrounded by pipette-accessible sample loading and fraction collection chambers	
Loading capacity	The instrument should enable high sample loading volume (>5X more than a 1D gel)	
Multiplexing	The system should provide robust fractionation over the mass range 3.5 kDa -500 kDa with high loading capacity, reproducibility, and recovery	
Protein Recovery	The instrument should enable high protein recovery (>80%)	
Reproducibility	The instrument should provide high reproducibility (<15% CV)	
Data System	The software must be a robust instrument control software: It should contain workflow based method editor	