User Charges Latest Addition:

Sl. No	Instrument and Software	Total per sample (Rs)	
		Academic	Private
1	Scanning Electron Microscope	2000	3900
2	Ultra-centrifuge	2900	5800
3	RNA sequencing	5900	8260
4	Exome sequencing	5900	8260
5	ChIP sequencing	5900	8260
6	sc-ATAC sequencing	49000	51350
7	sc-RNA sequencing	49000	51350
8	MD simulation	9440	11800
9	Server	9440	11800
10	GC-MS analysis	5000	7000
11	LC-MS/MS analysis, (Q-TRAP)	5000	7000
12	LC-MS/MS proteome analysis (digested sample) [ORBITRAP MS/MS]	9000	12000
13	LC-MS/MS proteome analysis (including sample preparation) [ORBITRAP MS/MS]	12000	15000
14	LC-MS/MS Quantitative proteomics (Labelled samples6 fractions maximum) [ORBITRAP MS/MS]	35000	50000
15	LC-MS/MS PTM analysis (Encriched sample) [ORBITRAP MS/MS]	15000	20000
16	LC-MS/MS Label free Quantitative proteome analysis (2sets of digested sample) [ORBITRAP MS/MS]	20000	30000
17	LC-MS/MS Label free Quantitative proteome analysis (including 2sets of sample preparation) [ORBITRAP MS/MS]	25000	35000
18	LC-MS/MS Label free Quantitative proteome analysis (3 sets of digested sample) [ORBITRAP MS/MS]	30000	40000
19	LC-MS/MS Label free Quantitative proteome analysis (including 3 sets of sample preparation) [ORBITRAP MS/MS]	35000	45000
20	LC-MS/MS Label free Quantitative proteome analysis (4 sets of digested sample) [ORBITRAP MS/MS]	40000	50000
21	LC-MS/MS Label free Quantitative proteome analysis (including 4 sets of sample preparation) [ORBITRAP MS/MS]	45000	55000