



Proteomics Core Facility

Price list for 2018

All prices are in HUF and VAT is excluded. In case of commercial requests discounts can be established based on the type of analysis and the number of samples.

The users of the Proteomics Core Facility agree that in their publications or power point presentations using the data acquired in/with the help of Proteomics Core Facility, in the „Materials and Methods” or „Acknowledgment” section they mention that *“The analyses were carried out at the Proteomics Core Facility, Department of Biochemistry and Molecular Biology, University of Debrecen”* and send a pdf of the paper, a copy of the figure or the first and last slides of their presentation to the Proteomics Core Facility as soon as possible but maximum to the end of each year.

Services	Price for BMBI	Price for laboratories from UD Faculty of Medicine	Price for academia	Price for companies
Mass spectrometry analysis				
In-gel or in-solution digestion with trypsin	1500	1500	4000	8000
LC-MS/MS based protein identification	13500	13500	16500	33000
SRM/MRM based protein analysis	13500	13500	16500	33000
C18ZipTip cleaning*	1000	1000	1100	2000
Electrophoresis				
2D electrophoresis on 7 cm gel**	7000	7000	31000	62000
2D electrophoresis on 24 cm gel**	30000	30000	55000	110000
7 cm gel/blot documentation using PharosFXPlus scanner	0	200	500	1000
24 cm gel documentation using PharosFXPlus scanner	0	500	700	1400



Gel band analysis using QuantityOne software for 7 cm gels	0	2000	3000	5000
Sample cleanup for 2D electrophoresis	2000	2000	4500	9000
Ultracentrifugation				
Centrifugation below 100000g***	0	800	1100	1500
Centrifugation with or above 100000g***	0	1100	1500	2000
O/N equilibrium gradient centrifugation	0	19000/day	25000/day	30000/day

* ZipTip cleaning is required in case of in-solution digestion or in case of samples digested by the Customer

** the price include staining with RuBPS

*** the price refers to each started 30 minute. In case of protocols when brake is off or is set to low value the time required for the rotor to stop will be included into the centrifugation time

For further enquires or digestion with other protease please contact the Proteomics Core Facility manager.