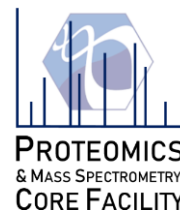


Molecular weight determination by ESI-MS

Request Form

Proteomics and Mass Spectrometry Core Facility
Malopolska Centre of Biotechnology, Jagiellonian University
e-mail: urszula.jankowska@uj.edu.pl lab: 1/40, phone: 12 664 6096



User Information

Name:	Date:
Mail:	Group name (PI):
Institution:	

Sample Details

Purpose of research/experiment:
--

No	Eppendorf mark	Approximate Molecular weight [Da]	Buffer composition	Concentration and volume	Additional information (e.g. purification history, mutations, modifications, sequence)
1		<input type="checkbox"/> Average: <input type="checkbox"/> Monoisotopic:			
2		<input type="checkbox"/> Average: <input type="checkbox"/> Monoisotopic:			
3		<input type="checkbox"/> Average: <input type="checkbox"/> Monoisotopic:			
4		<input type="checkbox"/> Average: <input type="checkbox"/> Monoisotopic:			
5		<input type="checkbox"/> Average: <input type="checkbox"/> Monoisotopic:			
...					

Please remember!

- bring your samples in eppendorf tubes (1,5 mL or 2 mL)
- use high-grade reagents and fresh buffers
- use gloves
- avoid detergents, especially PEG and Triton