**Molecular weight determination by ESI-MS**

**Request Form**

Proteomics and Mass Spectrometry Core Facility

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|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **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 |  | Average:  Monoisotopic: |  |  |  |
| 2 |  | Average:  Monoisotopic: |  |  |  |
| 3 |  | Average:  Monoisotopic: |  |  |  |
| 4 |  | Average:  Monoisotopic: |  |  |  |
| 5 |  | Average:  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