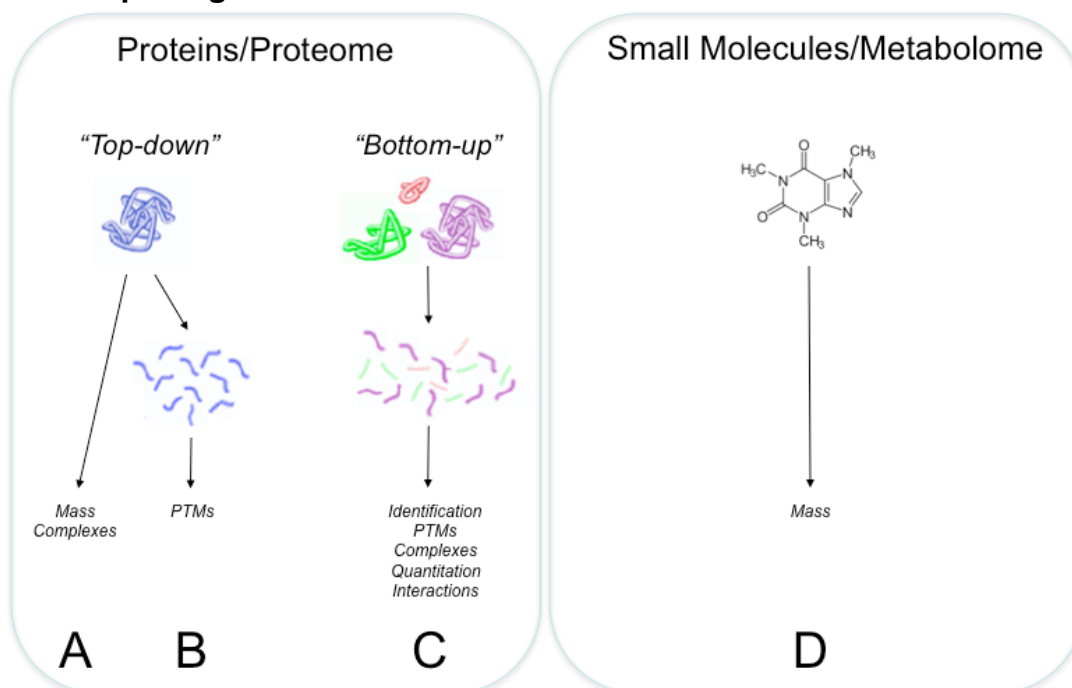


ProFI Terms, Conditions and Fees

By submitting a project to ProFI I agree to abide to the following Terms:

- i. The use, operation and maintenance of facility equipment are performed exclusively by the ProFI staff.
- ii. A submission form is required before the initiation of any research project, in order to design the proteomics approach, to calculate the costs and to evaluate the feasibility of the proposed project by ProFI staff.
- iii. The preparation and isolation of samples is the responsibility of the user, performed in his/her laboratory with guidance provided by the ProFI staff for the proper design and use of methods and consumables compatible with proteomics analysis.
- iv. Sample preparation for mass spectrometric analysis (including the proteolytic digestion of proteins) is performed at the dedicated workspace in ProFI by the user, unless otherwise agreed with ProFI staff.
- v. The costs include consumables/instrument spare parts needed for protein digestion, reversed phase liquid separation (RPLC) of proteins/peptides, mass spectrometric analysis, quality control and the initial processing of the data for protein or metabolite/compound identification.
- vi. Initial data processing and analysis for protein or metabolite/compound identification are performed by ProFI staff (see ProFI site). The users are provided with a viewer version of the software required (e.g. Proteome discoverer, Scaffold) for data visualization. A brief written report is sent by e-mail to the user, including the conditions and results of the analysis.
- vii. In case of required additional separation/purification of samples by methods/techniques available in ProFI (see ProFI site) this is performed by ProFI staff at additional costs.
- viii. In case further involvement in experimental design, data processing and analysis is required (meta-processing, e.g. functional and pathway analysis), this is performed after consultation with ProFI staff on a collaborative basis or at additional cost.
- ix. The MS raw files are archived (saved) by the ProFI staff and are not provided to the users.
- x. In case data produced by ProFI are presented in conferences, abstracts and publications, ProFI staff should be informed in advance and properly referenced, including acknowledgment of the ProFI funding agents.
- xi. Intense and active involvement of ProFI staff in collaborative projects indicates authorship of the involved ProFI members, after agreement with the user.

Services and pricing



Important

- ❖ For routine proteomics experiments “sample” refers to a single slice from a SDS-PAGE or a Native-PAGE gel. In case of liquid samples (e.g. protein solutions after chromatographic separation), “sample” refers to a single chromatographic fraction of low enough complexity to be analyzed by nanoLC-MS.
- ❖ For a single robust determination users should calculate that at least 2-3 triplicate identical samples will be needed (technical repeats). Also, in the case of gels, triplicate samples of an adjacent control lane will be needed. For a complete study of a single sample more than three biological repeats (i.e. different experiments) will be needed.
- ❖ Common experience indicates that even with relative “pure” targeted proteomics experiments (such as those of using immunoprecipitation), several protein bands will appear on a gel.

Price List IMBB-ProFI ¹				
No	Description	IMBB	External Academic	Commercial
A	Biomolecule mass determination by Mass Spectrometry	25 €	50€	90€
B, C	Protein identification and characterization by tryptic digestion and Mass Spectrometry	25 €	50 € (20% discount ≥ 20 samples) ²	90€
D	Accurate Mass Determination (Small Molecules)	15 €	(a) 30 € (15% discount ≥ 20 samples) ²	(a) 60 €
			(b) 50 € ³ (15% discount ≥ 20 samples) ²	(b) 80 € ³
	Extra analysis of results	- ⁴	60 €/hr ⁴	100 €/hr ⁴

¹- All prices are intended per sample submitted

²- Discount on total cost for simultaneous submission of 20 or more samples

³- For samples containing more than one masses

⁴- List of identified proteins is provided with confidence scores for their determination

Additional analysis will have to be discusses on a case by case basis.