Mass Spectrometry Unit

Request Form UniMS



Name: Institution: Address: Department /Group : Number of sample: Delivery date: Analysis date:		De Re	VAT n° Linha*: Phone: e-mail: Delivered by: Report date:		by:	
API-ION TRAP MA			-TOF/TOF	Triple T	Triple TOF 6600	
Small molecule mass measurement		Intact m	Intact mass measurement		Intact mass measurement	
Protein mass measurement		Protein i	Protein identification		Protein characterization	
LC-MS (positive/n	egative)	Nano-LC	Nano-LC-MALDI		Protein quantification	
NanoLC-MS (positive/negative)			<u>ine appropriate fields with x</u>	NanoLC-T	riple TOF	
<u>Mark the appro</u>	ipriate tields with x		Mici		Triple TOF	
Tdentification of samples Additional information						
Samples identification	Estimated molecular m (if appropriate)	ass (Da)	Quality System Solubilization solvent Sample amount & volum Pure sample Contaminants (salts, de Type of compound Reduction & alkylation	ie tergents, etc)	 None Non-Regulated biomedical research** GMP** Yes No Protein Carbohydrate Glycoprotein Organomethalic Polymers Other: 	
Sample preparation			Biological source of sample(s)			
Gel stain O Coomassie		stain	Elimination after analysis		O Yes O No	
O Other: O Trypsin			Observations			
O Photase O Other:						
Cleaning & Concentration O C18 O C8 O Other:						

Report delivery: within two weeks, the time may vary depending on the occupation or unexpected equipment failure. Further questions should be addressed to unims.technicians@itqb.unl.pt/unims.technicians@ibet.pt

Note: Publications that use data obtained by the UniMS - Mass Spectrometry Unit, should include this information in the appropriate section. This should be stated as: "Data provided/obtained by the UniMS - Mass Spectrometry Unit, ITQB/IBET, Oeiras, Portugal".