IRB Barcelona - Mass Spectrometry & Proteomics 2026

Service	Public and CERCA Institutions (€)
Data Analysis	
Data analysis-scripts/dedicated analysis (hour)	80,59
LCT-Premier	
LC-MS (1h)_ LCTPremier	166,47
LC-MS (short)_ LCTPremier	150,90
LC-MS_LCTPremier_auto	88,27
LC-MS_mABs_ LCTPremier	227,55
MS_LCT Premier	154,41
MS_LCT Premier_auto	113,70
Orbitrap-Eclipse	
LC-MS/MS <60 min run_ECLIPSE	248,71
MS_ECLIPSE	236,61
MS_noreport_ECLIPSE	213,61
LC-MS/MS (88min gradient), ECLIPSE-EVOSEP	310,33
LC-MS/MS (88min gradient), ECLIPSE-EVOSEP_only run	256,61
LC-MS/MS (44min gradient), ECLIPSE-EVOSEP	265,38
LC-MS/MS (44min gradient), ECLIPSE-EVOSEP_only run	206,57
Orbitrap-Lumos	
LC-MS/MS (120min gradient),150 min run_LUMOS	305,35
LC-MS/MS (120min gradient),150 min run_LUMOS_only run	246,54
LC-MS/MS (150min gradient),180 min run_LUMOS	346,44
LC-MS/MS (150min gradient),180 min run_LUMOS_only run	287,63
LC-MS/MS (180min gradient),210 min run_LUMOS	370,56
LC-MS/MS (180min gradient),210 min run_LUMOS_only run	311,75

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Orbitrap-Lumos	
LC-MS/MS (210min gradient),240min run_LUMOS	391,29
LC-MS/MS (210min gradient),240min run_LUMOS_onlyrun	332,48
LC-MS/MS (240min gradient),270min run_LUMOS	426,93
LC-MS/MS (240min gradient),270min run_LUMOS_only run	368,12
LC-MS/MS (270min gradient),300min run_LUMOS	447,66
LC-MS/MS (270min gradient),300min run_LUMOS_only run	388,85
LC-MS/MS (30min gradient), 60 min run_LUMOS	243,16
LC-MS/MS (30min gradient), 60 min run_LUMOS_only run	184,35
LC-MS/MS (60min gradient),90 min run_LUMOS	263,89
LC-MS/MS (60min gradient),90 min run_LUMOS_only run	205,08
LC-MS/MS (90min gradient),120 min run_LUMOS	284,62
LC-MS/MS (90min gradient),120 min run_LUMOS_only run	225,81
LC-MS/MS <60 min run_LUMOS	246,78
MS_LUMOS	218,33
MS_noreport_LUMOS	169,93
Sample prep	
Antibody deglycosilation	140,84
Crosslinking	241,62
Desalting and concentration by reversed phase	47,11
Desalting and concentration by SCX	75,46
Desalting and concentration by ultracentrifugation	47,49
Lysis & Protein Extraction	57,94
GelFree analysis	188,86
In solution Protein Digestion	85,20
In-gel Protein digestion	122,52

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Sample prep	
Isobaric labelling: TMT/iTRAQ (per sample)	871,39
off-line Fractionation kit (per sample/8 fractions)	231,02
Phosphopeptide enrichment	224,10
Protein quantitation (micro BSA)	33,44
Sample depletion (kit)	112,26
Sample prep (biospin)	120,17
Sample prep (FASP digestion)	87,55
Sample enrichment (ENRICH)	138,04
Sample enrichment (ENRICH plus)	165,67
Ultra 2	
LC-MS/MS (6min gradient), 120 SPD, Ultra2-EVOSEP	59,65
LC-MS/MS (16min gradient), 80 SPD, Ultra2-EVOSEP	86,82
LC-MS/MS (33min gradient), 40SPD, Ultra2-EVOSEP	135,39
LC-MS/MS (57min gradient), 20 SPD, Ultra2-EVOSEP	229,77
NeoFlex	
MALDI-MS_Neoflex	117,61
MALDI-MSI_Imaging_Neoflex	177,40

Please be advised that any future increases in consumable prices (e.g. due to customs duties on imported consumables or new tenders) may result in fee adjustments. Should these external costs rise, our pricing might be updated accordingly.

New services prices will be approved retroactively at the subsequent Board Meeting