

Price - Proteomics Platform - Protein identification by mass spectrometry

Code	Proteomics - Price per sample <i>Valid from January 1st 2015</i>	UdeS \$CAN	Academic (outside UdeS) \$CAN without indirect fees	Academic (outside UdeS) \$CAN with indirect fees (15%)	Non-academic \$CAN
Reagents					
<i>Amino acids: the price indicate de amount required for 500 ml of DMEM</i>					
ARG0	Arginine R0	5	6	6,9	10
ARG6	Arginine R6	130	160	184	210
ARG10	Arginin R10	90	100	115	150
LYS0	Lysine K0	5	6	6,9	10
LYS4	Lysine K4	120	150	172,5	200
LYS8	Lysine K8	180	210	241,5	300
FBSDIAL	Dialyzed FBS (50 ml)	90	60	69	120
DMEMRK	DMEM -R-K (500 ml)	50	60	69	120
TRYP	Trypsin Mass Spec Grade (10 ug)	15	20	23	30
SPINCOL	Microcon Spin Column	5	6	6,9	10
ZIPTIP	ZipTips	3,5	4	4,6	8
VIAL	Injection vial + cap	1,5	2	2,3	4
R0K0	DMEM with R0K0 + dialyzed serum	170	180	207	250
R6K4	DMEM with R6K4 + dialyzed serum	410	475	546,25	750
R10K8	DMEM with R10K8 + dialyzed serum	430	490	563,5	775
GFPTRAP	Sepharose + GFP Ac (per microliter)	0,3	0,5	0,575	2
LOBIND	Lo-Bind Eppendorf (per box)	35	40	46	56
PRECAST	4-15% precast gel + buffer	15	20	23	30
SBLUE	SimplyBlue Safestain (20ml)	10	15	17,25	30
FILTER	Filter 500 ml	20	25	28,75	50
Sample Preparation					
LIQDIG	Liquid digestion	100	120	138	200
1DGEL	1D Gel (gradient) + coomassie safestain	50	60	69	150
INGELDIG	In-gel digestion + peptide extraction	50	60	69	100
PHOSPHO	Phospho peptide enrichment (excluding digestion)	100	120	138	200
Mass Spectrometry					
LCLONG	LC-MS/MS (240min gradient)	50	60	69	210
ANALYSIS	Analysis (/hour)	40	48	55,2	100