

**Supplementary Table 1.** Comparative analysis between the proteomic results obtained with the THP-1 cultured on UnicCa with respect the proteomic results obtained with C samples at 1 and 3 days. Proteins with  $p \leq 0.05$  (yellow) and a ratio higher than 1.5 in either direction were considered as significantly different. Up-regulated proteins in THP-1 cultured on UnicCa are marked in red, while those downregulated are marked in green.

Uniprot entry	Protein name	Unique peptide	Day 1		Day 3	
			p value	Ratio UnicCa/C	p value	Ratio UnicCa/C
CO7A1_HUMAN	Collagen alpha-1(VII) chain	5	8,2E-04	26,36	1,0E-02	8,27
HIBCH_HUMAN	3-hydroxyisobutyryl-CoA hydrolase mitochondrial	7	6,1E-01	1,50	2,1E-03	7,32
MSH2_HUMAN	DNA mismatch repair protein Msh2	3	9,6E-01	0,95	1,1E-02	7,30
CHKB_HUMAN	Choline/ethanolamine kinase	3	1,3E-01	3,72	4,5E-04	5,12
PCH2_HUMAN	Pachytene checkpoint protein 2 homolog	3	2,2E-01	3,23	4,1E-03	5,08
SCRB1_HUMAN	Scavenger receptor class B member 1	2	8,4E-01	0,78	3,7E-02	4,61
CNOT9_HUMAN	CCR4-NOT transcription complex subunit 9	4	2,3E-01	3,33	3,9E-02	4,43
IPP2_HUMAN	Protein phosphatase inhibitor 2	2	3,0E-02	0,04	2,7E-01	4,20
GRWD1_HUMAN	Glutamate-rich WD repeat-containing protein 1	7	8,3E-03	0,15	1,4E-01	3,81
GSLG1_HUMAN	Golgi apparatus protein 1	20	4,0E-01	2,33	2,6E-02	3,74
MEPCE_HUMAN	7SK snRNA methylphosphate capping enzyme	4	5,1E-01	1,71	4,1E-02	3,71
RL27A_HUMAN	60S ribosomal protein L27a	7	3,9E-02	0,39	2,9E-01	3,68
NU214_HUMAN	Nuclear pore complex protein Nup214	8	6,1E-01	1,31	1,6E-02	3,19
NH2L1_HUMAN	NHP2-like protein 1	6	6,5E-01	1,44	1,1E-02	3,15
CBWD1_HUMAN	COBW domain-containing protein 1	3	9,3E-03	5,22	2,4E-01	3,14
HGD_HUMAN	Homogentisate 1 2-dioxygenase	7	2,8E-01	3,79	1,9E-02	3,12
RM28_HUMAN	39S ribosomal protein L28 mitochondrial	2	7,0E-02	2,92	7,1E-03	3,09
PNPT1_HUMAN	Polyribonucleotide nucleotidyltransferase 1 mitochondrial	12	2,0E-02	18,72	2,6E-01	3,08
RM46_HUMAN	39S ribosomal protein L46 mitochondrial	2	8,4E-01	1,15	1,4E-02	2,98
ITA4_HUMAN	Integrin alpha-4	2	3,3E-02	3,96	5,6E-02	2,97
RTRAF_HUMAN	RNA transcription translation and transport factor protein	6	2,8E-02	7,64	1,8E-01	2,96
CBL_HUMAN	E3 ubiquitin-protein ligase CBL	14	9,5E-01	1,06	3,1E-02	2,72
SHIP1_HUMAN	Phosphatidylinositol 3 4 5-trisphosphate 5-phosphatase 1	16	3,0E-02	0,36	1,8E-01	2,60

NQO1_HUMAN	NAD(P)H dehydrogenase [quinone] 1	11	3,0E-01	2,53	3,2E-02	2,60
ODP2_HUMAN	Dihydrolipoyllysine-residue acetyltransferase component of pyruvate dehydrogenase complex mitochondrial	10	2,4E-02	4,14	1,6E-01	2,58
RTN1_HUMAN	Reticulon-1	2	4,2E-01	1,78	4,5E-02	2,51
NDUF4_HUMAN	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 4	3	2,8E-02	2,41	3,8E-01	2,51
EMC10_HUMAN	ER membrane protein complex subunit 10	2	2,2E-02	14,84	5,1E-01	2,46
SYSC_HUMAN	Serine--tRNA ligase cytoplasmic	17	2,2E-02	2,76	9,1E-02	2,41
MIA2_HUMAN	Melanoma inhibitory activity protein 2	2	4,8E-02	5,64	5,1E-01	2,36
TCRG1_HUMAN	Transcription elongation regulator 1	9	3,0E-02	19,88	4,7E-01	2,34
NEK9_HUMAN	Serine/threonine-protein kinase Nek9	7	3,0E-01	1,71	7,6E-03	2,33
UBP2L_HUMAN	Ubiquitin-associated protein 2-like	20	1,9E-01	2,64	4,2E-02	2,28
UFC1_HUMAN	Ubiquitin-fold modifier-conjugating enzyme 1	5	4,7E-02	0,29	3,5E-01	2,23
PDP1_HUMAN	[Pyruvate dehydrogenase [acetyl-transferring]]-ph phatase 1 mitochondrial	5	4,1E-03	33,99	1,3E-01	2,20
RBP56_HUMAN	TATA-binding protein-associated factor 2N	4	7,9E-01	1,19	3,7E-02	2,20
OCAD1_HUMAN	OCIA domain-containing protein 1	7	2,8E-02	4,15	1,5E-01	2,19
PSB6_HUMAN	Proteasome subunit beta type-6	4	7,3E-01	1,39	3,4E-02	2,18
NIPS1_HUMAN	Protein NipSnap homolog 1	5	7,0E-01	0,75	2,5E-02	2,18
SHLB1_HUMAN	Endophilin-B1	10	3,1E-01	2,11	6,3E-03	2,14
MYPT1_HUMAN	Protein phosphatase 1 regulatory subunit 12A	5	4,8E-01	1,83	2,6E-02	2,14
TRPV2_HUMAN	Transient receptor potential cation channel subfamily V member 2	3	4,8E-02	4,59	3,5E-01	2,08
PUR9_HUMAN	Bifunctional purine biosynthesis protein PURH	38	7,2E-01	0,77	3,6E-02	2,07
EST1_HUMAN	Liver carboxylesterase 1	35	5,8E-01	0,79	3,2E-02	2,00
PDLI7_HUMAN	PDZ and LIM domain protein 7	4	1,2E-02	6,83	4,1E-01	1,98
RPIA_HUMAN	Ribose-5-phosphate isomerase	2	4,2E-01	0,59	2,8E-02	1,95
CO4A_HUMAN	Complement C4-A	12	4,8E-01	1,39	3,9E-02	1,91
HS74L_HUMAN	Heat shock 70 kDa protein 4L	9	2,0E-02	0,41	4,0E-01	1,90
RM09_HUMAN	39S ribosomal protein L9 mitochondrial	2	3,9E-02	0,12	2,7E-01	1,83
BICD2_HUMAN	Protein bicaudal D homolog 2	2	3,0E-02	4,19	4,0E-01	1,78

NMI_HUMAN	N-myc-interactor	5	2,0E-01	0,41	1,9E-03	1,76
LTOR3_HUMAN	Ragulator complex protein LAMTOR3	2	4,3E-02	2,37	2,6E-01	1,74
ABHDA_HUMAN	Mycophenolic acid acyl-glucuronide esterase mitochondrial	2	4,8E-02	3,64	4,3E-01	1,71
GRP75_HUMAN	Stress-70 protein mitochondrial	45	6,3E-03	3,65	2,7E-01	1,70
RS6_HUMAN	40S ribosomal protein S6	13	1,4E-04	0,05	7,1E-01	1,68
BGAL_HUMAN	Beta-galactosidase	4	4,2E-02	4,58	2,4E-01	1,67
RS10_HUMAN	40S ribosomal protein S10	13	2,5E-02	1,96	1,2E-01	1,66
TF2B_HUMAN	Transcription initiation factor IIB	3	2,7E-02	10,90	7,1E-01	1,61
NELFB_HUMAN	Negative elongation factor B	2	3,2E-02	6,81	3,8E-01	1,60
RUFY1_HUMAN	RUN and FYVE domain-containing protein 1	10	3,8E-02	0,29	6,5E-01	1,59
RSU1_HUMAN	Ras suppressor protein 1	5	2,3E-02	2,47	3,5E-01	1,59
PSD11_HUMAN	26S proteasome non-ATPase regulatory subunit 11	26	4,0E-02	2,54	2,1E-01	1,59
RUVB1_HUMAN	RuvB-like 1	23	2,6E-02	2,33	2,0E-01	1,59
RUXE_HUMAN	Small nuclear ribonucleoprotein E	4	8,1E-03	5,98	1,1E-01	1,59
HUWE1_HUMAN	E3 ubiquitin-protein ligase HUWE1	27	2,4E-03	2,10	2,0E-01	1,58
PSB1_HUMAN	Proteasome subunit beta type-1	11	1,1E-02	2,31	1,3E-01	1,57
KCAB2_HUMAN	Voltage-gated potassium channel subunit beta-2	13	7,8E-03	5,25	5,2E-01	1,56
TBL3_HUMAN	Transducin beta-like protein 3	7	6,0E-02	4,20	4,1E-02	1,56
GGPPS_HUMAN	Geranylgeranyl pyrophosphate synthase	6	1,7E-02	2,70	3,9E-01	1,54
ODR4_HUMAN	Protein odr-4 homolog	2	2,0E-02	2,78	3,0E-01	1,54
RPP30_HUMAN	Ribonuclease P protein subunit p30	2	2,2E-02	3,71	5,2E-01	1,54
PTSS1_HUMAN	Phosphatidylserine synthase 1	7	9,2E-03	11,53	5,6E-01	1,52
P5CS_HUMAN	Delta-1-pyrroline-5-carboxylate synthase	4	2,0E-02	5,49	4,6E-01	1,51
ESTD_HUMAN	S-formylglutathione hydrolase	11	1,5E-02	4,33	5,2E-01	1,50
LTOR1_HUMAN	Ragulator complex protein LAMTOR1	6	5,6E-04	4,24	2,0E-01	1,48
NSUN2_HUMAN	RNA cytosine C(5)-methyltransferase NSUN2	19	4,2E-02	1,72	2,2E-02	1,47
ITAX_HUMAN	Integrin alpha-X	23	3,6E-02	2,31	3,7E-01	1,45
NLTP_HUMAN	Non-specific lipid-transfer protein	12	4,8E-04	3,35	5,0E-01	1,43

NFKB2_HUMAN	Nuclear factor NF-kappa-B p100 subunit	2	4,5E-02	5,80	5,7E-01	1,42
SF01_HUMAN	Splicing factor 1	6	3,8E-03	2,40	3,7E-01	1,41
NIF3L_HUMAN	NIF3-like protein 1	2	1,8E-02	7,23	3,7E-01	1,39
BNIP2_HUMAN	BCL2/adenovirus E1B 19 kDa protein-interacting protein 2	2	9,3E-03	7,55	6,3E-01	1,39
THIC_HUMAN	Acetyl-CoA acetyltransferase cytosolic	9	1,4E-02	2,77	4,5E-01	1,37
PPIB_HUMAN	Peptidyl-prolyl cis-trans isomerase B	23	1,8E-02	1,91	2,3E-01	1,36
ELAV1_HUMAN	ELAV-like protein 1	15	2,3E-02	2,77	4,8E-01	1,35
EXOS5_HUMAN	Exosome complex component RRP46	5	2,8E-02	3,70	5,3E-01	1,34
H2A1C_HUMAN	Histone H2A type 1-C	2	2,5E-02	0,04	8,3E-01	1,34
MBNL1_HUMAN	Muscleblind-like protein 1	5	3,1E-02	4,07	6,9E-01	1,33
ATP9A_HUMAN	Probable phospholipid-transporting ATPase IIA	2	2,3E-02	3,61	3,6E-01	1,32
USP9X_HUMAN	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	13	4,5E-02	2,39	1,5E-01	1,32
GID8_HUMAN	Glucose-induced degradation protein 8 homolog	2	3,8E-05	11,50	7,1E-01	1,32
TM165_HUMAN	Transmembrane protein 165	2	3,3E-02	1,66	7,8E-01	1,31
NXP20_HUMAN	Protein NOXP20	3	7,1E-03	2,92	7,9E-01	1,30
PCY2_HUMAN	Ethanolamine-phosphate cytidyltransferase	4	6,6E-03	8,09	7,7E-01	1,29
ERAP1_HUMAN	Endoplasmic reticulum aminopeptidase 1	17	2,6E-03	9,42	6,7E-01	1,27
CPPED_HUMAN	Serine/threonine-protein phosphatase CPPED1	6	2,5E-02	0,16	7,7E-01	1,26
PSMD5_HUMAN	26S proteasome non-ATPase regulatory subunit 5	6	4,5E-02	4,25	7,1E-01	1,25
MCU_HUMAN	Calcium uniporter protein mitochondrial	2	6,6E-03	4,57	7,2E-01	1,24
FA98A_HUMAN	Protein FAM98A	2	4,2E-02	4,63	5,8E-01	1,24
GRHPR_HUMAN	Glyoxylate reductase/hydroxypyruvate reductase	12	2,3E-02	2,17	6,7E-01	1,24
PSIP1_HUMAN	PC4 and SFRS1-interacting protein	2	1,4E-02	2,45	7,5E-01	1,23
PFD5_HUMAN	Prefoldin subunit 5	3	4,4E-02	0,20	8,2E-01	1,22
LGUL_HUMAN	Lactoylglutathione lyase	20	3,8E-02	2,87	7,1E-01	1,21
RL11_HUMAN	60S ribosomal protein L11	12	3,6E-02	2,55	5,8E-01	1,20
GCP2_HUMAN	Gamma-tubulin complex component 2	3	2,7E-02	8,42	8,4E-01	1,19
AP2A2_HUMAN	AP-2 complex subunit alpha-2	13	8,2E-03	3,00	7,7E-01	1,19

PGAM1_HUMAN	Phosphoglycerate mutase 1	22	7,4E-03	2,25	6,6E-01	1,18
NEK7_HUMAN	Serine/threonine-protein kinase Nek7	2	3,1E-03	7,00	7,5E-01	1,17
HNRPL_HUMAN	Heterogeneous nuclear ribonucleoprotein L	28	6,7E-03	2,64	7,7E-01	1,17
RU2A_HUMAN	U2 small nuclear ribonucleoprotein A'	12	3,8E-02	1,70	7,7E-01	1,17
PANX1_HUMAN	Pannexin-1	3	4,0E-02	6,94	6,9E-01	1,17
TPC12_HUMAN	Trafficking protein particle complex subunit 12	2	3,8E-02	3,60	7,8E-01	1,17
PRKRA_HUMAN	Interferon-inducible double-stranded RNA-dependent protein kinase activator A	4	1,5E-02	6,53	9,0E-01	1,16
RPN2_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2	27	9,5E-04	5,20	8,3E-01	1,15
H4_HUMAN	Histone H4	18	2,4E-02	0,41	7,4E-01	1,15
SCAM2_HUMAN	Secretory carrier-associated membrane protein 2	4	2,1E-03	1,57	5,2E-01	1,13
MEAK7_HUMAN	MTOR-associated protein MEAK7	2	2,0E-02	4,91	9,0E-01	1,13
BABA2_HUMAN	BRISC and BRCA1-A complex member 2	6	4,0E-02	4,67	8,9E-01	1,13
ATG5_HUMAN	Autophagy protein 5	3	2,6E-02	3,35	8,3E-01	1,13
PIGS_HUMAN	GPI transamidase component PIG-S	7	3,9E-03	8,18	8,3E-01	1,12
PSA5_HUMAN	Proteasome subunit alpha type-5	13	2,6E-02	1,74	7,9E-01	1,12
ATPA_HUMAN	ATP synthase subunit alpha mitochondrial	40	4,3E-02	2,51	7,8E-01	1,12
PSB3_HUMAN	Proteasome subunit beta type-3	11	2,4E-02	2,24	7,1E-01	1,11
MPPB_HUMAN	Mitochondrial-processing peptidase subunit beta	13	3,9E-02	3,90	9,1E-01	1,11
CD14_HUMAN	Monocyte differentiation antigen CD14	4	1,4E-02	10,31	8,6E-01	1,11
IN35_HUMAN	Interferon-induced 35 kDa protein	2	2,5E-02	4,32	9,0E-01	1,11
ARPC2_HUMAN	Actin-related protein 2/3 complex subunit 2	30	3,0E-02	2,78	8,2E-01	1,10
LRC40_HUMAN	Leucine-rich repeat-containing protein 40	2	2,3E-02	8,10	8,9E-01	1,10
VP33B_HUMAN	Vacuolar protein sorting-associated protein 33B	5	7,6E-03	4,98	9,1E-01	1,09
S10A4_HUMAN	Protein S100-A4	6	1,0E-02	4,03	9,1E-01	1,08
DUS3L_HUMAN	tRNA-dihydrouridine(47) synthase [NAD(P)(+)]-like	9	5,0E-02	8,46	9,4E-01	1,07
MAPK3_HUMAN	MAP kinase-activated protein kinase 3	2	2,1E-02	2,71	9,3E-01	1,06
G6PI_HUMAN	Glucose-6-phosphate isomerase	23	1,2E-02	2,13	7,7E-01	1,06

DAB2_HUMAN	Disabled homolog 2	7	1,9E-02	3,88	8,4E-01	1,05
GSTO1_HUMAN	Glutathione S-transferase omega-1	20	1,3E-02	2,53	9,3E-01	1,03
SDCB1_HUMAN	Syntenin-1	10	2,0E-02	3,41	9,6E-01	1,03
ETF_A_HUMAN	Electron transfer flavoprotein subunit alpha mitochondrial	17	3,0E-02	2,18	9,1E-01	1,03
EF1A1_HUMAN	Elongation factor 1-alpha 1	19	3,8E-02	3,41	9,7E-01	1,02
IFT25_HUMAN	Intraflagellar transport protein 25 homolog	3	4,1E-02	9,40	9,8E-01	1,02
THADA_HUMAN	Thyroid adenoma-associated protein	2	2,3E-02	2,50	9,7E-01	1,02
LANC1_HUMAN	Glutathione S-transferase LANCL1	2	4,0E-02	3,90	9,7E-01	1,01
MPI_HUMAN	Mannose-6-phosphate isomerase	5	2,6E-02	4,66	9,9E-01	1,00
TBCE_HUMAN	Tubulin-specific chaperone E	6	1,9E-02	2,30	9,9E-01	1,00
CC88B_HUMAN	Coiled-coil domain-containing protein 88B	3	7,7E-03	0,14	1,0E+00	1,00
RS2_HUMAN	40S ribosomal protein S2	13	3,1E-02	3,06	1,0E+00	1,00
T2FB_HUMAN	General transcription factor IIF subunit 2	3	4,1E-02	0,18	1,0E+00	1,00
AT2C1_HUMAN	Calcium-transporting ATPase type 2C member 1	2	7,4E-03	0,46	9,9E-01	0,99
TETN_HUMAN	Tetranectin	7	1,6E-02	3,43	9,8E-01	0,98
RFC5_HUMAN	Replication factor C subunit 5	3	1,8E-02	7,03	9,5E-01	0,98
ZC3HF_HUMAN	Zinc finger CCCH domain-containing protein 15	2	3,7E-02	0,42	9,7E-01	0,98
COR1C_HUMAN	Coronin-1C	19	2,1E-02	2,43	9,5E-01	0,98
UBA1_HUMAN	Ubiquitin-like modifier-activating enzyme 1	45	4,4E-02	2,20	9,4E-01	0,98
DOP2_HUMAN	Protein dopey-2	5	2,3E-02	2,86	9,4E-01	0,97
ATPD_HUMAN	ATP synthase subunit delta mitochondrial	3	2,2E-02	3,29	9,8E-01	0,97
TOIP2_HUMAN	Torsin-1A-interacting protein 2	3	2,3E-02	3,39	9,8E-01	0,97
IPO7_HUMAN	Importin-7	27	4,2E-02	2,80	9,4E-01	0,97
NOMO2_HUMAN	Nodal modulator 2	34	6,6E-04	4,06	9,5E-01	0,96
AIMP2_HUMAN	Aminoacyl tRNA synthase complex-interacting multifunctional protein 2	6	2,9E-03	2,41	9,4E-01	0,96
ENPL_HUMAN	Endoplasmic reticulum protein	73	4,6E-02	2,03	8,9E-01	0,95
COPG1_HUMAN	Coatomer subunit gamma-1	30	4,9E-02	2,67	9,2E-01	0,95
TB182_HUMAN	182 kDa tankyrase-1-binding protein	11	1,9E-02	0,29	8,5E-01	0,95

H2A2C_HUMAN	Histone H2A type 2-C	3	1,9E-02	0,09	9,1E-01	0,94
EIF3A_HUMAN	Eukaryotic translation initiation factor 3 subunit A	37	2,6E-02	2,44	8,7E-01	0,93
MSTO1_HUMAN	Protein misato homolog 1	2	2,3E-02	3,58	8,7E-01	0,93
P3H2_HUMAN	Prolyl 3-hydroxylase 2	2	1,5E-02	1,84	8,3E-01	0,92
NIP7_HUMAN	60S ribosome subunit biogenesis protein NIP7 homolog	2	3,1E-02	2,53	8,9E-01	0,90
DAZP1_HUMAN	DAZ-associated protein 1	9	3,7E-02	2,15	8,4E-01	0,90
CFAI_HUMAN	Complement factor I	2	1,7E-02	11,59	9,2E-01	0,90
MGST2_HUMAN	Microsomal glutathione S-transferase 2	2	7,9E-04	12,40	9,0E-01	0,90
HGH1_HUMAN	Protein HGH1 homolog	5	4,8E-02	3,86	9,0E-01	0,90
CAN1_HUMAN	Calpain-1 catalytic subunit	18	1,4E-02	2,19	7,3E-01	0,89
MGAT1_HUMAN	Alpha-1 3-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyltransferase	2	1,5E-02	2,97	8,1E-01	0,89
SYAC_HUMAN	Alanine--tRNA ligase cytoplasmic	36	4,4E-02	3,17	7,9E-01	0,88
DC1L2_HUMAN	Cytoplasmic dynein 1 light intermediate chain 2	3	7,6E-03	15,95	8,9E-01	0,87
E2AK2_HUMAN	Interferon-induced double-stranded RNA-activated protein kinase	5	4,4E-02	4,48	8,2E-01	0,87
PPID_HUMAN	Peptidyl-prolyl cis-trans isomerase D	5	2,3E-02	2,16	6,1E-01	0,87
DPP2_HUMAN	Dipeptidyl peptidase 2	14	8,2E-03	3,85	7,5E-01	0,85
PIMT_HUMAN	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	9	3,9E-03	10,23	7,8E-01	0,85
RHG27_HUMAN	Rho GTPase-activating protein 27	2	1,9E-02	2,06	8,1E-01	0,85
GCN1_HUMAN	eIF-2-alpha kinase activator GCN1	26	2,5E-02	3,04	7,3E-01	0,85
HYOU1_HUMAN	Hypoxia up-regulated protein 1	54	3,9E-02	0,52	7,2E-01	0,84
AT1B3_HUMAN	Sodium/potassium-transporting ATPase subunit beta-3	8	1,5E-02	2,94	7,1E-01	0,84
CATG_HUMAN	Cathepsin G	17	1,9E-02	2,09	7,0E-01	0,83
TMM33_HUMAN	Transmembrane protein 33	5	5,0E-03	13,51	8,4E-01	0,83
FA5_HUMAN	Coagulation factor V	8	3,5E-02	0,16	8,3E-01	0,83
FLOT2_HUMAN	Flotillin-2	4	2,7E-02	4,55	8,2E-01	0,83
ORNT1_HUMAN	Mitochondrial ornithine transporter 1	2	3,2E-02	5,65	7,9E-01	0,81
RS12_HUMAN	40S ribosomal protein S12	10	3,4E-02	3,03	6,6E-01	0,81
HXK2_HUMAN	Hexokinase-2	19	2,8E-02	6,51	7,0E-01	0,81

TM10C_HUMAN	tRNA methyltransferase 10 homolog C	2	4,8E-05	13,06	8,2E-01	0,81
GLYM_HUMAN	Serine hydroxymethyltransferase mitochondrial	27	1,9E-02	4,99	7,4E-01	0,81
LACB2_HUMAN	Endoribonuclease LACTB2	8	2,4E-02	1,99	6,6E-01	0,81
TECR_HUMAN	Very-long-chain enoyl-CoA reductase	13	1,7E-02	5,02	7,2E-01	0,79
WDR1_HUMAN	WD repeat-containing protein 1	42	4,2E-02	2,63	5,2E-01	0,79
UBP24_HUMAN	Ubiquitin carboxyl-terminal hydrolase 24	3	2,4E-02	2,18	4,4E-01	0,78
THOC6_HUMAN	THO complex subunit 6 homolog	4	3,0E-02	3,51	7,3E-01	0,78
IF4E_HUMAN	Eukaryotic translation initiation factor 4E	8	2,9E-02	0,42	5,1E-01	0,78
FSCN1_HUMAN	Fascin	39	1,0E-02	1,52	2,7E-01	0,78
CY24B_HUMAN	Cytochrome b-245 heavy chain	5	3,8E-02	3,24	7,4E-01	0,78
MCCA_HUMAN	Methylcrotonoyl-CoA carboxylase subunit alpha mitochondrial	5	7,1E-03	12,35	6,5E-01	0,77
NNRE_HUMAN	NAD(P)H-hydrate epimerase	9	2,9E-02	6,75	7,3E-01	0,77
BST1_HUMAN	ADP-ribosyl cyclase/cyclic ADP-ribose hydrolase 2	12	1,5E-02	4,05	7,6E-01	0,76
EF1G_HUMAN	Elongation factor 1-gamma	24	2,4E-02	2,42	1,5E-01	0,76
5NTD_HUMAN	5'-nucleotidase	13	2,8E-02	8,84	5,1E-01	0,76
ERO1A_HUMAN	ERO1-like protein alpha	11	2,0E-02	7,53	7,5E-01	0,76
DHRS4_HUMAN	Dehydrogenase/reductase SDR family member 4	2	4,4E-02	1,67	3,6E-01	0,75
DNJC9_HUMAN	DnaJ homolog subfamily C member 9	3	2,0E-03	2,73	7,0E-01	0,75
LETM1_HUMAN	Mitochondrial proton/calcium exchanger protein	14	8,0E-03	0,34	5,3E-01	0,75
MBB1A_HUMAN	Myb-binding protein 1A	2	1,6E-02	4,92	7,4E-01	0,75
GELS_HUMAN	Gelsolin	55	3,7E-02	2,46	6,1E-01	0,74
FAN_HUMAN	Protein FAN	3	2,1E-02	5,60	6,9E-01	0,74
PSA3_HUMAN	Proteasome subunit alpha type-3	13	2,1E-02	0,44	5,2E-01	0,74
STX5_HUMAN	Syntaxin-5	2	1,9E-02	5,12	6,5E-01	0,74
ARPC3_HUMAN	Actin-related protein 2/3 complex subunit 3	12	4,0E-02	2,51	3,9E-01	0,74
SF3B6_HUMAN	Splicing factor 3B subunit 6	4	2,9E-02	0,37	2,6E-01	0,74
LRC59_HUMAN	Leucine-rich repeat-containing protein 59	14	5,4E-03	2,46	5,5E-01	0,73
GOSR1_HUMAN	Golgi SNAP receptor complex member 1	2	4,3E-02	5,47	6,7E-01	0,73



RHG17_HUMAN	Rho GTPase-activating protein 17	10	9,6E-03	7,04	7,4E-01	0,72
RAD21_HUMAN	Double-strand-break repair protein rad21 homolog	6	1,7E-02	1,92	5,3E-01	0,72
ST1A1_HUMAN	Sulfotransferase 1A1	6	4,7E-02	3,18	3,5E-01	0,72
PPIF_HUMAN	Peptidyl-prolyl cis-trans isomerase F mitochondrial	11	4,2E-03	0,37	2,6E-01	0,72
KPYM_HUMAN	Pyruvate kinase PKM	56	3,8E-02	1,69	2,3E-01	0,71
HSP74_HUMAN	Heat shock 70 kDa protein 4	49	3,7E-02	0,64	1,6E-01	0,71
HNRPK_HUMAN	Heterogeneous nuclear ribonucleoprotein K	31	4,2E-02	3,91	6,4E-01	0,71
CAH2_HUMAN	Carbonic anhydrase 2	30	6,9E-03	2,45	3,8E-01	0,70
MEP50_HUMAN	Methylosome protein 50	6	1,1E-02	4,56	6,0E-01	0,70
FCL_HUMAN	GDP-L-fucose synthase	7	2,5E-02	3,50	1,8E-01	0,68
FAM3C_HUMAN	Protein FAM3C	5	1,1E-02	3,49	3,1E-01	0,68
IF4E2_HUMAN	Eukaryotic translation initiation factor 4E type 2	2	3,7E-02	2,20	2,4E-01	0,68
COMP_HUMAN	Cartilage oligomeric matrix protein	7	2,3E-02	0,19	6,4E-01	0,67
EF1D_HUMAN	Elongation factor 1-delta	14	2,2E-01	0,20	1,5E-02	0,66
MYLK_HUMAN	Myosin light chain kinase smooth muscle	3	4,2E-02	6,25	5,2E-01	0,65
IF4G3_HUMAN	Eukaryotic translation initiation factor 4 gamma 3	3	3,3E-02	4,26	5,0E-01	0,65
DPOD2_HUMAN	DNA polymerase delta subunit 2	3	1,7E-02	3,88	7,2E-01	0,65
PRS7_HUMAN	26S proteasome regulatory subunit 7	21	1,3E-02	0,35	7,2E-01	0,65
AT2B4_HUMAN	Plasma membrane calcium-transporting ATPase 4	7	9,5E-01	1,06	2,9E-02	0,65
CELF1_HUMAN	CUGBP Elav-like family member 1	4	3,6E-02	0,43	3,9E-01	0,64
RPAP3_HUMAN	RNA polymerase II-associated protein 3	3	3,3E-02	4,23	5,2E-01	0,64
NUCKS_HUMAN	Nuclear ubiquitous casein and cyclin-dependent kinase substrate 1	5	2,1E-02	0,14	6,1E-01	0,64
TT39B_HUMAN	Tetratricopeptide repeat protein 39B	4	1,1E-02	10,13	6,2E-01	0,63
NAA50_HUMAN	N-alpha-acetyltransferase 50	10	3,8E-02	0,13	2,5E-01	0,63
TGO1_HUMAN	Transport and Golgi organization protein 1 homolog	3	4,6E-01	0,53	3,3E-02	0,63
TRAM1_HUMAN	Translocating chain-associated membrane protein 1	2	3,3E-02	3,92	6,1E-01	0,63
C1TM_HUMAN	Monofunctional C1-tetrahydrofolate synthase mitochondrial	5	3,7E-02	3,30	5,1E-01	0,63
NDUS1_HUMAN	NADH-ubiquinone oxidoreductase 75 kDa subunit mitochondrial	15	3,6E-02	10,74	3,8E-01	0,63

COMT_HUMAN	Catechol O-methyltransferase	24	4,9E-02	1,99	7,6E-02	0,62
TCPB_HUMAN	T-complex protein 1 subunit beta	43	8,3E-03	1,72	2,1E-01	0,62
DPYD_HUMAN	Dihydropyrimidine dehydrogenase [NADP(+)]	3	1,7E-02	3,01	1,0E-01	0,62
SPN90_HUMAN	NCK-interacting protein with SH3 domain	6	5,7E-02	0,29	2,9E-02	0,61
H2AY_HUMAN	Core histone macro-H2A.1	15	7,3E-01	0,87	4,6E-02	0,61
ATPG_HUMAN	ATP synthase subunit gamma mitochondrial	9	1,3E-03	2,22	3,2E-01	0,61
HMG1_HUMAN	Non-histone chromosomal protein HMG-14	3	1,2E-03	0,11	7,1E-02	0,60
FINC_HUMAN	Fibronectin	44	2,1E-03	6,93	5,8E-01	0,59
DNJA2_HUMAN	DnaJ homolog subfamily A member 2	7	6,6E-01	0,71	3,1E-02	0,59
RL31_HUMAN	60S ribosomal protein L31	4	1,4E-02	7,96	5,4E-01	0,59
STK10_HUMAN	Serine/threonine-protein kinase 10	31	8,1E-01	1,10	2,9E-02	0,59
CSTOS_HUMAN	Protein CUSTOS	2	7,8E-03	0,25	2,7E-01	0,59
ACTN4_HUMAN	Alpha-actinin-4	41	4,2E-02	5,09	6,0E-01	0,58
DHB12_HUMAN	Very-long-chain 3-oxoacyl-CoA reductase	10	4,6E-01	0,73	3,4E-02	0,58
PAIRB_HUMAN	Plasminogen activator inhibitor 1 RNA-binding protein	18	8,0E-01	0,82	3,9E-02	0,58
AP1B1_HUMAN	AP-1 complex subunit beta-1	13	2,0E-02	0,32	2,7E-01	0,57
CO5_HUMAN	Complement C5	3	4,5E-02	5,73	3,6E-01	0,57
RTCA_HUMAN	RNA 3'-terminal phosphate cyclase	10	3,8E-02	4,97	6,0E-01	0,57
CLPB_HUMAN	Caseinolytic peptidase B protein homolog	7	3,0E-02	4,78	5,5E-01	0,57
PHB_HUMAN	Prohibitin	12	9,8E-01	1,01	4,3E-02	0,57
ISG15_HUMAN	Ubiquitin-like protein ISG15	3	4,7E-02	5,97	5,9E-01	0,57
NT5D2_HUMAN	5'-nucleotidase domain-containing protein 2	9	5,7E-01	0,70	8,4E-04	0,56
AP2S1_HUMAN	AP-2 complex subunit sigma	4	4,5E-02	7,65	4,2E-01	0,56
BCAR1_HUMAN	Breast cancer anti-estrogen resistance protein 1	2	1,4E-02	3,47	3,3E-01	0,56
PGRC1_HUMAN	Membrane-associated progesterone receptor component 1	4	9,9E-03	2,45	6,9E-02	0,56
NIPA_HUMAN	Nuclear-interacting partner of ALK	2	4,1E-02	3,64	2,1E-01	0,56
RL30_HUMAN	60S ribosomal protein L30	6	8,6E-03	1,80	4,0E-01	0,56
COA3_HUMAN	Cytochrome c oxidase assembly factor 3 homolog mitochondrial	2	6,0E-01	0,79	4,6E-02	0,55

EH1L1_HUMAN	EH domain-binding protein 1-like protein 1	2	4,4E-02	0,52	3,0E-01	0,54
IF2A_HUMAN	Eukaryotic translation initiation factor 2 subunit 1	26	7,5E-01	0,86	3,2E-02	0,54
VP26C_HUMAN	Vacuolar protein sorting-associated protein 26C	2	3,0E-02	3,04	2,9E-01	0,54
YIF1A_HUMAN	Protein YIF1A	2	3,1E-02	5,80	4,2E-01	0,54
SIL1_HUMAN	Nucleotide exchange factor SIL1	5	5,0E-02	4,08	3,4E-01	0,54
PRP31_HUMAN	U4/U6 small nuclear ribonucleoprotein Prp31	5	4,4E-02	0,20	4,5E-01	0,53
RNH2B_HUMAN	Ribonuclease H2 subunit B	2	1,2E-01	2,85	4,9E-02	0,52
DCTN4_HUMAN	Dynactin subunit 4	2	4,6E-02	3,40	2,1E-01	0,52
PML_HUMAN	Protein PML	7	7,5E-01	0,76	1,9E-02	0,52
TCTP_HUMAN	Translationally-controlled tumor protein	10	5,6E-01	1,44	3,7E-02	0,51
FLNB_HUMAN	Filamin-B	11	4,7E-01	1,24	4,9E-02	0,50
DPP3_HUMAN	Dipeptidyl peptidase 3	23	3,7E-01	0,30	4,5E-02	0,50
NIBA1_HUMAN	Protein Niban 1	10	3,9E-01	1,60	4,5E-02	0,50
PCY1A_HUMAN	Choline-phosphate cytidyltransferase A	8	4,2E-01	0,55	3,7E-02	0,50
PPCE_HUMAN	Prolyl endopeptidase	33	1,4E-01	2,80	3,7E-02	0,50
IMA3_HUMAN	Importin subunit alpha-3	9	3,2E-02	7,74	5,3E-01	0,50
PLXB2_HUMAN	Plexin-B2	23	3,7E-02	0,24	3,5E-01	0,50
ENOF1_HUMAN	Mitochondrial enolase superfamily member 1	2	2,0E-02	6,19	3,1E-01	0,49
RIC8A_HUMAN	Synembryn-A	5	5,1E-01	1,59	4,4E-02	0,49
ACPH_HUMAN	Acylamino-acid-releasing enzyme	13	3,0E-01	2,51	2,7E-02	0,48
HMCS1_HUMAN	Hydroxymethylglutaryl-CoA synthase cytoplasmic	9	3,5E-02	2,18	3,5E-01	0,48
OGT1_HUMAN	UDP-N-acetylglucosamine--peptide N-acetylglucosaminyltransferase 110 kDa subunit	14	3,2E-02	2,62	4,4E-01	0,48
HS2ST_HUMAN	Heparan sulfate 2-O-sulfotransferase 1	6	9,8E-01	1,03	4,2E-02	0,47
RTN4_HUMAN	Reticulon-4	8	2,4E-02	2,28	2,9E-01	0,47
SYTC_HUMAN	Threonine--tRNA ligase 1 cytoplasmic	30	1,2E-02	3,21	4,0E-01	0,47
KSYK_HUMAN	Tyrosine-protein kinase SYK	14	2,7E-01	1,51	1,6E-02	0,47
S39AE_HUMAN	Metal cation symporter ZIP14	3	2,0E-01	0,41	2,7E-02	0,47

CAR19_HUMAN	Caspase recruitment domain-containing protein 19	2	4,9E-02	2,52	2,3E-01	0,47
MDHC_HUMAN	Malate dehydrogenase cytoplasmic	21	4,6E-02	2,45	2,8E-01	0,46
ICAM1_HUMAN	Intercellular adhesion molecule 1	9	2,9E-01	0,35	2,9E-02	0,46
MOES_HUMAN	Moesin	44	7,3E-01	0,85	2,7E-02	0,46
HNRPD_HUMAN	Heterogeneous nuclear ribonucleoprotein D0	22	1,5E-02	2,58	3,3E-01	0,46
S4A7_HUMAN	Sodium bicarbonate cotransporter 3	17	5,9E-01	2,02	3,8E-02	0,45
SAM50_HUMAN	Sorting and assembly machinery component 50 homolog	12	4,8E-01	1,32	2,6E-02	0,45
ITPA_HUMAN	Inosine triphosphate pyrophosphatase	3	3,3E-02	17,03	5,3E-01	0,45
FL2D_HUMAN	Pre-mRNA-splicing regulator WTAP	4	4,6E-02	0,28	1,9E-01	0,44
CPNS1_HUMAN	Calpain small subunit 1	18	7,2E-01	1,42	8,6E-03	0,44
OAT_HUMAN	Ornithine aminotransferase mitochondrial	6	5,7E-01	1,27	2,7E-02	0,44
ZN622_HUMAN	Zinc finger protein 622	2	9,6E-01	1,02	3,2E-02	0,44
CTNA1_HUMAN	Catenin alpha-1	4	1,5E-01	0,33	2,5E-02	0,44
TKFC_HUMAN	Triokinase/FMN cyclase	8	4,3E-02	3,29	1,9E-01	0,44
SRSF5_HUMAN	Serine/arginine-rich splicing factor 5	6	1,2E-03	0,07	2,7E-01	0,44
SET_HUMAN	Protein SET	16	9,9E-01	1,00	2,2E-02	0,43
SRSF3_HUMAN	Serine/arginine-rich splicing factor 3	7	3,3E-03	0,26	1,7E-01	0,42
GSK3A_HUMAN	Glycogen synthase kinase-3 alpha	3	8,9E-01	0,93	2,1E-02	0,42
SNX5_HUMAN	Sorting nexin-5	9	5,5E-01	0,85	3,6E-02	0,42
PCM1_HUMAN	Pericentriolar material 1 protein	34	2,1E-02	0,09	7,1E-02	0,41
CND1_HUMAN	Condensin complex subunit 1	5	1,2E-02	8,41	2,7E-01	0,41
RALB_HUMAN	Ras-related protein Ral-B	3	7,6E-01	1,25	1,7E-02	0,41
HNRPC_HUMAN	Heterogeneous nuclear ribonucleoproteins C1/C2	10	5,4E-01	0,79	1,1E-02	0,41
CSN7B_HUMAN	COP9 signalosome complex subunit 7b	2	6,7E-01	1,28	1,5E-02	0,40
CHP1_HUMAN	Calcineurin B homologous protein 1	5	5,0E-02	5,64	3,1E-01	0,40
CNPY3_HUMAN	Protein canopy homolog 3	9	3,3E-01	0,83	3,7E-02	0,40
RFA2_HUMAN	Replication protein A 32 kDa subunit	3	2,6E-01	3,05	2,9E-02	0,40
BRE1A_HUMAN	E3 ubiquitin-protein ligase BRE1A	2	8,0E-01	0,87	1,8E-02	0,39

H15_HUMAN	Histone H1.5	8	8,7E-01	1,04	5,0E-02	0,39
PEF1_HUMAN	Peflin	2	3,7E-02	2,93	3,2E-01	0,39
GRN_HUMAN	Progranulin	2	5,0E-01	0,69	4,4E-02	0,39
TGM2_HUMAN	Protein-glutamine gamma-glutamyltransferase 2	23	7,3E-01	0,77	5,5E-03	0,39
NELFE_HUMAN	Negative elongation factor E	2	3,3E-01	0,49	4,5E-02	0,39
BIN2_HUMAN	Bridging integrator 2	9	2,5E-02	10,18	3,3E-01	0,39
WASP_HUMAN	Wiskott-Aldrich syndrome protein	3	3,9E-01	1,84	3,9E-02	0,39
PHAG1_HUMAN	Phosphoprotein associated with glycosphingolipid-enriched microdomains 1	2	1,7E-01	0,34	4,6E-02	0,39
ARFG1_HUMAN	ADP-ribosylation factor GTPase-activating protein 1	2	4,0E-01	0,57	4,5E-02	0,38
T23O_HUMAN	Tryptophan 2 3-dioxygenase	8	1,8E-01	0,36	4,7E-02	0,38
NDKB_HUMAN	Nucleoside diphosphate kinase B	5	2,2E-01	5,92	4,8E-02	0,37
SUMF2_HUMAN	Inactive C-alpha-formylglycine-generating enzyme 2	7	6,0E-01	1,55	5,5E-03	0,37
IL1B_HUMAN	Interleukin-1 beta	14	5,9E-01	1,55	3,4E-02	0,37
SRSF7_HUMAN	Serine/arginine-rich splicing factor 7	10	6,9E-01	1,38	1,2E-02	0,36
NPM_HUMAN	Nucleophosmin	21	4,3E-02	0,47	2,0E-02	0,36
KPCD_HUMAN	Protein kinase C delta type	8	8,8E-01	0,97	3,3E-02	0,35
DJC13_HUMAN	DnaJ homolog subfamily C member 13	3	5,4E-01	1,57	2,2E-02	0,35
RRP8_HUMAN	Ribosomal RNA-processing protein 8	2	9,3E-01	0,93	4,8E-02	0,35
YTDC1_HUMAN	YTH domain-containing protein 1	2	1,1E-02	0,11	2,1E-01	0,35
PDIA5_HUMAN	Protein disulfide-isomerase A5	4	4,2E-02	0,30	2,1E-01	0,34
SLK_HUMAN	STE20-like serine/threonine-protein kinase	2	5,2E-01	0,70	2,1E-02	0,34
RHG01_HUMAN	Rho GTPase-activating protein 1	19	3,7E-02	4,86	4,7E-01	0,33
TMF1_HUMAN	TATA element modulatory factor	2	8,4E-01	0,90	3,4E-03	0,32
FKBP5_HUMAN	Peptidyl-prolyl cis-trans isomerase FKBP5	12	7,4E-01	1,53	2,1E-02	0,32
FIBG_HUMAN	Fibrinogen gamma chain	2	2,1E-02	4,91	5,6E-02	0,32
LCP2_HUMAN	Lymphocyte cytosolic protein 2	3	4,2E-02	0,34	1,5E-01	0,32
FRIL_HUMAN	Ferritin light chain	3	1,9E-02	9,47	3,4E-01	0,32
PRKDC_HUMAN	DNA-dependent protein kinase catalytic subunit	53	4,9E-03	4,10	4,0E-01	0,31

MAK16_HUMAN	Protein MAK16 homolog	5	2,9E-03	0,07	1,2E-01	0,31
BABA1_HUMAN	BRISC and BRCA1-A complex member 1	3	3,8E-01	0,55	4,8E-02	0,31
RAB13_HUMAN	Ras-related protein Rab-13	5	1,4E-02	4,62	7,4E-02	0,30
IF1AX_HUMAN	Eukaryotic translation initiation factor 1A X-chromosomal	6	8,4E-02	0,23	2,7E-03	0,29
SPB10_HUMAN	Serpin B10	9	4,4E-01	0,60	3,3E-02	0,29
RHG30_HUMAN	Rho GTPase-activating protein 30	3	2,2E-02	3,30	5,7E-02	0,29
SKP1_HUMAN	S-phase kinase-associated protein 1	5	8,1E-01	1,20	9,2E-03	0,28
TIAR_HUMAN	Nucleolysin TIAR	2	7,4E-01	0,78	1,6E-02	0,28
HBS1L_HUMAN	HBS1-like protein	5	8,5E-01	0,87	3,7E-03	0,28
CASP8_HUMAN	Caspase-8	3	1,9E-01	1,75	9,2E-03	0,28
NRP1_HUMAN	Neuropilin-1	13	5,0E-03	0,04	1,9E-01	0,28
PRAF3_HUMAN	PRA1 family protein 3	3	1,3E-01	2,71	2,0E-02	0,27
U2AF2_HUMAN	Splicing factor U2AF 65 kDa subunit	8	5,4E-01	1,52	4,4E-02	0,27
RAP1A_HUMAN	Ras-related protein Rap-1A	2	4,1E-02	0,41	1,9E-01	0,27
PSMD8_HUMAN	26S proteasome non-ATPase regulatory subunit 8	3	1,3E-01	2,83	3,5E-02	0,27
SYRC_HUMAN	Arginine--tRNA ligase cytoplasmic	38	2,7E-03	0,24	9,4E-02	0,27
AMPD3_HUMAN	AMP deaminase 3	11	8,6E-03	0,02	2,0E-03	0,27
CBX3_HUMAN	Chromobox protein homolog 3	6	4,2E-01	0,63	4,8E-02	0,27
PPWD1_HUMAN	Peptidylprolyl isomerase domain and WD repeat-containing protein 1	2	7,8E-01	1,13	2,9E-02	0,27
BT3L4_HUMAN	Transcription factor BTF3 homolog 4	6	2,5E-01	0,55	8,2E-03	0,26
ARSB_HUMAN	Arylsulfatase B	2	8,7E-01	0,87	3,9E-02	0,26
RTN3_HUMAN	Reticulon-3	2	5,6E-04	0,28	2,7E-01	0,25
MESD_HUMAN	LRP chaperone MESD	3	7,3E-01	0,72	4,0E-02	0,25
GYS1_HUMAN	Glycogen [starch] synthase muscle	4	9,7E-01	1,03	1,4E-02	0,25
UN93B_HUMAN	Protein unc-93 homolog B1	3	2,9E-01	0,54	3,7E-02	0,25
KIF2A_HUMAN	Kinesin-like protein KIF2A	2	5,5E-01	1,52	7,2E-03	0,25
DAD1_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1	2	4,7E-02	8,93	1,7E-01	0,25
AN32E_HUMAN	Acidic leucine-rich nuclear phosphoprotein 32 family member E	6	7,9E-01	0,90	1,2E-02	0,23

GON7_HUMAN	EKC/KEOPS complex subunit GON7	2	1,1E-01	7,17	3,0E-02	0,23
IL8_HUMAN	Interleukin-8	2	5,2E-04	2,56	7,3E-02	0,23
EBP2_HUMAN	Probable rRNA-processing protein EBP2	5	8,6E-01	0,89	4,9E-02	0,22
NOLC1_HUMAN	Nucleolar and coiled-body phosphoprotein 1	2	1,9E-02	0,21	7,8E-02	0,22
ARRB2_HUMAN	Beta-arrestin-2	5	1,1E-01	0,30	4,7E-02	0,22
SPTC1_HUMAN	Serine palmitoyltransferase 1	7	6,0E-01	1,59	2,2E-02	0,21
AN32B_HUMAN	Acidic leucine-rich nuclear phosphoprotein 32 family member B	6	6,1E-01	0,80	3,6E-02	0,20
NUCL_HUMAN	Nucleolin	49	6,3E-01	0,77	4,9E-02	0,20
UBE2Z_HUMAN	Ubiquitin-conjugating enzyme E2 Z	3	2,1E-02	0,19	4,0E-02	0,19
MAGD2_HUMAN	Melanoma-associated antigen D2	3	6,5E-01	0,77	1,6E-02	0,18
DENR_HUMAN	Density-regulated protein	7	1,4E-01	5,53	1,9E-02	0,18
ABI1_HUMAN	Abl interactor 1	5	9,1E-01	0,91	4,9E-02	0,18
HDAC2_HUMAN	Histone deacetylase 2	4	9,5E-01	0,94	8,7E-03	0,18
TRA2B_HUMAN	Transformer-2 protein homolog beta	11	9,5E-01	0,98	4,0E-02	0,17
CBX5_HUMAN	Chromobox protein homolog 5	7	5,0E-03	0,17	2,0E-03	0,16
JMJD6_HUMAN	Bifunctional arginine demethylase and lysyl-hydroxylase JMJD6	4	2,2E-01	0,52	1,3E-02	0,15
TSP4_HUMAN	Thrombospondin-4	5	1,2E-01	0,33	4,6E-02	0,15
TCP4_HUMAN	Activated RNA polymerase II transcriptional coactivator p15	7	7,4E-02	0,14	3,9E-02	0,15
RL36L_HUMAN	60S ribosomal protein L36a-like	2	2,7E-01	1,30	2,2E-02	0,14
DNJC7_HUMAN	DnaJ homolog subfamily C member 7	16	4,6E-02	3,88	7,4E-02	0,14
RU17_HUMAN	U1 small nuclear ribonucleoprotein 70 kDa	10	3,8E-01	0,53	1,1E-02	0,14
M3K20_HUMAN	Mitogen-activated protein kinase kinase kinase 20	4	7,6E-01	0,87	2,6E-02	0,14
NVL_HUMAN	Nuclear valosin-containing protein-like	2	2,3E-01	0,54	4,1E-02	0,13
TOP1_HUMAN	DNA topoisomerase 1	9	7,2E-01	0,72	1,3E-02	0,13
TBB3_HUMAN	Tubulin beta-3 chain	3	1,7E-02	8,53	1,1E-01	0,13
IF2B_HUMAN	Eukaryotic translation initiation factor 2 subunit 2	17	8,4E-01	0,76	4,6E-02	0,11
PLEK_HUMAN	Pleckstrin	16	4,8E-01	2,02	2,8E-02	0,11
UGGG1_HUMAN	UDP-glucose:glycoprotein glucosyltransferase 1	38	4,0E-01	0,40	1,3E-02	0,10

SRSF6_HUMAN	Serine/arginine-rich splicing factor 6	6	1,6E-03	0,25	2,5E-03	0,10
SMAP_HUMAN	Small acidic protein	3	7,5E-01	0,69	1,6E-02	0,10
ARFG3_HUMAN	ADP-ribosylation factor GTPase-activating protein 3	5	3,5E-01	2,24	1,4E-02	0,09
CHMP5_HUMAN	Charged multivesicular body protein 5	5	5,8E-01	0,79	1,9E-02	0,09
SRSF2_HUMAN	Serine/arginine-rich splicing factor 2	5	7,3E-01	0,67	3,4E-02	0,06
TEBP_HUMAN	Prostaglandin E synthase 3	10	5,6E-01	0,69	1,5E-02	0,03