

Plectin OS=Homo sapiens GN=PLEC PE=1 SV=3	sp Q15149 PLEC_HU	532 kDa	100%	100%	100%	100%			100%	100%	100%
Triosephosphate isomerase OS=Homo sapiens GN=TP11 PE=1 SV=3	sp P60174 TPIS_HU	31 kDa				100%	67%		100%	100%	100%
Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	sp P11142 HSP7C_HU	71 kDa	100%	98%	100%		98%		14%		100%
Voltage-dependent anion- selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2	sp P21796 VDAC1_HU	31 kDa				100%	100%				88%
Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2	sp P38646 GRP75_HU	74 kDa	68%	100%	100%	100%	68%	100%	100%	79%	100%
Mitochondrial-processing peptidase subunit alpha OS=Homo sapiens GN=PMPCA PE=1 SV=2	sp Q10713 MPPA_HU	58 kDa	100%	100%	100%				16%	79%	
Carbamoyl-phosphate synthase [ammonia], mitochondrial OS=Homo sapiens GN=CPS1 PE=1 SV=2	sp P31327 CPSM_HU	165 kDa	100%	100%	100%	100%	68%	100%	100%	100%	100%
60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2	sp P10809 CH60_HU	61 kDa		100%	100%	100%	100%	100%	100%	77%	100%
Keratin, type I cytoskeletal 14 OS=Homo sapiens GN=KRT14 PE=1 SV=4	sp P02533 K1C14_HU	52 kDa	100%	100%	100%	100%		100%	84%	100%	100%
Keratin, type II cytoskeletal 5 OS=Homo sapiens GN=KRT5 PE=1 SV=3	sp P13647 K2C5_HU	62 kDa	100%	100%	100%			100%	100%	100%	100%
Alkaline phosphatase, placental type OS=Homo sapiens GN=ALPP PE=1 SV=2	sp P05187 PPB1_HU	58 kDa				80%					100%

Aspartate--tRNA ligase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2	sp P14868 SYDC_H	57 kDa	100%	100%	100%				
Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A PE=1 SV=1	sp Q71U36 TBA1A	50 kDa	100%	100%	100%	100%		100%	100%
Pyruvate dehydrogenase protein X component, mitochondrial OS=Homo sapiens GN=PDHX PE=1 SV=3	sp O00330 ODPX_H	54 kDa	100%	100%	100%				
Keratin, type I cytoskeletal 16 OS=Homo sapiens GN=KRT16 PE=1 SV=4	sp P08779 K1C16_H	51 kDa	100%	100%	100%	80%	100%		100%
4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3	sp P49189 AL9A1_H	54 kDa	100%	100%	100%				
L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2	sp P00338 LDHA_H	37 kDa					100%	100%	
Leucine-rich PPR motif- containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3	sp P42704 LRPPRC_H	158 kDa			100%	100%		74%	100%
Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1	sp P54652 HSP72_H	70 kDa	16%	84%	100%	80%		67%	100%
Aldehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH2 PE=1 SV=2	sp P05091 ALDH2_H	56 kDa			100%	100%			
Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2	sp Q99623 PHB2_H	33 kDa					100%	100%	
Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1	sp P60842 IF4A1_H	46 kDa				100%		79%	73%

Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	sp P60709 ACTB_H	42 kDa	100%	100%	100%	100%		69%	100%	100%
Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2	sp P34931 HS71L_H	70 kDa	43%	70%	69%		100%		75%	100%
Alpha-aminoadipic semialdehyde dehydrogenase OS=Homo sapiens GN=ALDH7A1 PE=1 SV=5	sp P49419 AL7A1_H	58 kDa	100%	100%	100%					
Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1	sp P61978 HNRPK_	51 kDa	100%	100%	100%	100%		100%		100%
Actin, aortic smooth muscle OS=Homo sapiens GN=ACTA2 PE=1 SV=1	sp P62736 ACTA_H	42 kDa	100%	100%	100%	100%		74%	100%	100%
Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=2	sp P02786 TFR1_HU	85 kDa	100%	100%	100%	100%		100%	77%	79%
Keratinocyte proline-rich protein OS=Homo sapiens GN=KPRP PE=1 SV=1	sp Q5T749 KPRP_H	64 kDa	100%	100%	100%		100%	100%	100%	
Nucleoprotein TPR OS=Homo sapiens GN=TPR PE=1 SV=3	sp P12270 TPR_HU	267 kDa	100%	100%	100%	100%			79%	100%
GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3	sp P62826 RAN_HU	24 kDa						100%	100%	
Serum albumin precursor (Allergen Bos d 6) (BSA) cRAP	gi 1351907 sp P0276	69 kDa	100%	100%	100%	100%	68%	100%		78%
Keratin, type I cytoskeletal 13 OS=Homo sapiens GN=KRT13 PE=1 SV=4	sp P13646 K1C13_H	50 kDa	100%	100%	100%	99%		100%	100%	100%
ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4	sp Q08211 DHX9_H	141 kDa			69%	100%		100%	15%	100%
Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3	sp P34897 GLYM_H	56 kDa	100%	100%	100%					

Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2	sp P22626 ROA2_H	37 kDa					100%	100%		99%		
Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	sp P06733 ENOA_H	47 kDa	100%	100%	100%	100%				100%		100%
Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondrial OS=Homo sapiens GN=OXCT1 PE=1 SV=1	sp P55809 SCOT1_H	56 kDa	100%	100%	100%							
WD repeat-containing protein 82 OS=Homo sapiens GN=WDR82 PE=1 SV=1	sp Q6UXN9 WDR82_H	35 kDa					100%	100%				
RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1	sp Q9Y265 RUVB1_H	50 kDa	100%	100%	100%							100%
Keratin, type II cytoskeletal 7 OS=Homo sapiens GN=KRT7 PE=1 SV=5	sp P08729 K2C7_HU	51 kDa	68%	100%	100%							
Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	sp P04406 G3P_HU	36 kDa	100%		100%	19%	100%	100%		100%	74%	88%
D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4	sp O43175 SERA_H	57 kDa	100%	100%	100%					100%		
Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3	sp P28838 AMPL_H	56 kDa	100%	100%	100%							
Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 SV=3	sp P31930 QCR1_HU	53 kDa					100%				100%	73%
Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3	sp P30041 PRDX6_H	25 kDa						100%	100%			

Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3	sp P22314 UBA1_H	118 kDa	100%	100%	100%	80%				100%
Intestinal-type alkaline phosphatase OS=Homo sapiens GN=ALPI PE=1 SV=2	sp P09923 PPBI_HU	57 kDa		100%						100%
ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3	sp O75947 ATP5H_H	18 kDa						100%		
Spectrin alpha chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTAN1 PE=1 SV=3	sp Q13813 SPTN1_H	285 kDa			100%	80%				100%
Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3	sp P40926 MDHM_H	36 kDa					100%	100%		
Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4	sp P18206 VINC_HU	124 kDa	100%	100%	100%	80%				
Pre-mRNA-processing factor 19 OS=Homo sapiens GN=PRPF19 PE=1 SV=1	sp Q9UMS4 PRP19_H	55 kDa	100%	100%	100%				79%	
ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3	sp P06576 ATPB_HU	57 kDa			69%				100%	100%
Spectrin beta chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTBN1 PE=1 SV=2	sp Q01082 SPTB2_H	275 kDa			69%	100%				100%
Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5	sp P08107 HSP71_H	70 kDa			69%			100%		100%
Heterogeneous nuclear ribonucleoprotein A1-like 2 OS=Homo sapiens GN=HNRNPA1L2 PE=2 SV=2	sp Q32P51 RA1L2_H	34 kDa					100%	100%		

Keratin, type I cytoskeletal 17 OS=Homo sapiens GN=KRT17 PE=1 SV=2	sp Q04695 K1C17_H	48 kDa	68%		69%	100%		74%	100%	100%
T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3	sp P40227 TCPZ_HU	58 kDa		98%	99%		67%	100%	100%	
Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2	sp P23246 SFPQ_HU	76 kDa	100%	100%	100%			92%		
Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1	sp P26599 PTBP1_H	57 kDa	100%	100%	100%		67%			
Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform OS=Homo sapiens GN=PPP2R2A PE=1 SV=1	sp P63151 2ABA_HU	52 kDa	100%	100%	100%					
Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2	sp O60506 HNRPQ	70 kDa		68%	26%	100%				100%
Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3	sp P52272 HNRPM	78 kDa					100%	100%	56%	
Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=1 SV=3	sp Q16836 HCDH_H	34 kDa					100%	100%		
U2 small nuclear ribonucleoprotein B' OS=Homo sapiens GN=SNRNPB2 PE=1 SV=1	sp P08579 RU2B_HU	25 kDa						100%	100%	
Nuclear mitotic apparatus protein 1 OS=Homo sapiens GN=NUMA1 PE=1 SV=2	sp Q14980 NUMA1	238 kDa			99%					100%
Trypsin-1 OS=Homo sapiens GN=PRSS1 PE=1 SV=1	sp P07477 TRY1_HU	27 kDa	68%	13%	100%		68%	67%	100%	100% 88%
Histone H2A type 1-A OS=Homo sapiens GN=HIST1H2AA PE=1 SV=3	sp Q96QV6 H2A1A	14 kDa		100%		80%		99%	99%	99%

Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4	sp P08670 VIME_HU	54 kDa		70%	100%		100%	77%	100%
Isoleucine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=IARS PE=1 SV=2	sp P41252 SYIC_HU	145 kDa	100%	100%	98%	100%			
Bifunctional glutamate/proline-- tRNA ligase OS=Homo sapiens GN=EPRS PE=1 SV=5	sp P07814 SYEP_HU	171 kDa	100%	100%	99%				
Putative heat shock protein HSP 90-beta-3 OS=Homo sapiens GN=HSP90AB3P PE=5 SV=1	sp Q58FF7 H90B3_H	68 kDa				100%			100%
Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70A PE=1 SV=1	sp O94826 TOM70_H	67 kDa					79%		100%
Splicing factor 45 OS=Homo sapiens GN=RBM17 PE=1 SV=1	sp Q96125 SPF45_H	45 kDa				100%			17%
Actin-like protein 6A OS=Homo sapiens GN=ACTL6A PE=1 SV=1	sp O96019 ACL6A_H	47 kDa					100%		
26S protease regulatory subunit 7 OS=Homo sapiens GN=PSMC2 PE=1 SV=3	sp P35998 PRS7_HU	49 kDa					100%		
Sorting nexin-1 OS=Homo sapiens GN=SNX1 PE=1 SV=3	sp Q13596 SNX1_H	59 kDa							100%
Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2	sp P00367 DHE3_HU	61 kDa	100%	70%	100%		100%		
Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1	sp Q5D862 FILA2_H	248 kDa		70%	69%		100%	100%	
Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3	sp P49327 FAS_HU	273 kDa		100%	100%	100%			
Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 PE=1 SV=3	sp Q12906 ILF3_HU	95 kDa	98%	100%	100%				

Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3	sp P63244 GBLP_HU	35 kDa			100%	100%		74%				
Pyrroline-5-carboxylate reductase 2 OS=Homo sapiens GN=PYCR2 PE=1 SV=1	sp Q96C36 P5CR2_HU	34 kDa			100%	100%						
AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2	sp Q9Y4W6 AFG32_HU	89 kDa	100%	100%								
Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondrial OS=Homo sapiens GN=CHCHD3 PE=1 SV=1	sp Q9NX63 CHCH3_HU	26 kDa						100%	77%			
Kinectin OS=Homo sapiens GN=KTN1 PE=1 SV=1	sp Q86UP2 KTN1_HU	156 kDa	69%									100%
Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	sp O43707 ACTN4_HU	105 kDa										100%
Myosin-10 OS=Homo sapiens GN=MYH10 PE=1 SV=3	sp P35580 MYH10_HU	229 kDa										100%
Chromobox protein homolog 3 OS=Homo sapiens GN=CBX3 PE=1 SV=4	sp Q13185 CBX3_HU	21 kDa									100%	
Evolutionarily conserved signaling intermediate in Toll pathway, mitochondrial OS=Homo sapiens GN=ECSIT PE=1 SV=1	sp Q9BQ95 ECSIT_HU	49 kDa				100%						
Cationic trypsin precursor (Beta-trypsin) [Contains: Alpha-trypsin chain 1; Alpha-trypsin chain 2] cRAP	gij 2507249 sp P0076	25 kDa	69%		68%	67%	84%	74%	100%	79%	100%	
Keratin, type II cytoskeletal 6B OS=Homo sapiens GN=KRT6B PE=1 SV=5	sp P04259 K2C6B_HU	60 kDa	100%	100%	100%			100%	77%			
Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	sp P07437 TBB5_HU	50 kDa	68%	59%	100%	99%						100%

Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4	sp P55072 TERA_H	89 kDa	68%	70%	100%			100%
Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1	sp P50995 ANX11_H	54 kDa		100%	100%		100%	
Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3	sp P37802 TAGL2_H	22 kDa					100%	
V-type proton ATPase catalytic subunit A OS=Homo sapiens GN=ATP6V1A PE=1 SV=2	sp P38606 VATA_HU	68 kDa						100%
Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2	sp Q07065 CKAP4_H	66 kDa						100%
UDP-glucose:glycoprotein glucosyltransferase 1 OS=Homo sapiens GN=UGGT1 PE=1 SV=3	sp Q9NYU2 UGGG1_H	177 kDa						100%
Tyrosine-protein phosphatase non-receptor type 1 OS=Homo sapiens GN=PTPN1 PE=1 SV=1	sp P18031 PTN1_HU	50 kDa	68%	100%	100%	80%		
Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2	sp P54886 P5CS_HU	87 kDa	68%	70%			98%	100%
Mitochondrial inner membrane protein OS=Homo sapiens GN=IMMT PE=1 SV=1	sp Q16891 IMMT_H	84 kDa		40%	98%	100%		79%
Lamina-associated polypeptide 2, isoform alpha OS=Homo sapiens GN=TMPO PE=1 SV=2	sp P42166 LAP2A_H	75 kDa	68%	100%	100%		34%	
Dihydrolipoyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2	sp P09622 DLDH_H	54 kDa	100%	100%	100%			
Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3 PE=1 SV=3	sp P30048 PRDX3_H	28 kDa	100%	100%				100%

Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3	sp Q16658 FSCN1_H	55 kDa	100%	100%	100%	
Phosphatidylinositol 4-kinase type 2-alpha OS=Homo sapiens GN=PI4K2A PE=1 SV=1	sp Q9BTU6 P4K2A_H	54 kDa	68%	100%	100%	
Heterogeneous nuclear ribonucleoprotein F OS=Homo sapiens GN=HNRNPF PE=1 SV=3	sp P52597 HNRNPF_H	46 kDa		47%		100%
Citrate synthase, mitochondrial OS=Homo sapiens GN=CS PE=1 SV=2	sp O75390 CISY_HU	52 kDa			80%	100%
Acylpyruvase FAHD1, mitochondrial OS=Homo sapiens GN=FAHD1 PE=1 SV=2	sp Q6P587 FAHD1_H	25 kDa				100%
Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	sp P08238 HS90B_H	83 kDa			100%	100%
Eukaryotic translation initiation factor 3 subunit F OS=Homo sapiens GN=EIF3F PE=1 SV=1	sp O00303 EIF3F_H	38 kDa			100%	100%
Cob(I)yrinic acid a,c-diamide adenosyltransferase, mitochondrial OS=Homo sapiens GN=MMAB PE=1 SV=1	sp Q96EY8 MMAB_H	27 kDa				100%
Cathepsin D OS=Homo sapiens GN=CTSD PE=1 SV=1	sp P07339 CATD_H	45 kDa			100%	
Chromodomain-helicase-DNA- binding protein 4 OS=Homo sapiens GN=CHD4 PE=1 SV=2	sp Q14839 CHD4_H	218 kDa				100%
[Pyruvate dehydrogenase [acetyl-transferring]]- phosphatase 1, mitochondrial OS=Homo sapiens GN=PDP1 PE=1 SV=3	sp Q9P0J1 PDP1_HU	61 kDa			100%	

Putative beta-actin-like protein 3 OS=Homo sapiens GN=POTEKP PE=5 SV=1	sp Q9BYX7 ACTBM	42 kDa			69%	80%		100%	100%
Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3	sp P26641 EF1G_HU	50 kDa	60%	70%	69%	100%			
Inosine-5'-monophosphate dehydrogenase 2 OS=Homo sapiens GN=IMPDH2 PE=1 SV=2	sp P12268 IMDH2_H	56 kDa	99%	100%	100%				
Nodal modulator 1 OS=Homo sapiens GN=NOMO1 PE=1 SV=5	sp Q15155 NOMO1_H	134 kDa		24%	69%	100%			100%
Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 PE=1 SV=2	sp Q8NC51 PAIRB_H	45 kDa	68%	100%	100%				
Adenylosuccinate lyase OS=Homo sapiens GN=ADSL PE=1 SV=2	sp P30566 PUR8_HU	55 kDa	68%	100%	100%				
Septin-8 OS=Homo sapiens GN=SEPT8 PE=1 SV=4	sp Q92599 SEPT8_H	56 kDa		100%	100%				
X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=2	sp P12956 XRCC6_H	70 kDa					100%		100%
Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=2	sp P43243 MATR3_H	95 kDa					74%		100%
SPRY domain-containing protein 4 OS=Homo sapiens GN=SPRYD4 PE=1 SV=2	sp Q8WW59 SPRY4_H	23 kDa				58%		100%	
U5 small nuclear ribonucleoprotein 200 kDa helicase OS=Homo sapiens GN=SNRNP200 PE=1 SV=2	sp O75643 U520_HU	245 kDa							100%
Electron transfer flavoprotein subunit beta OS=Homo sapiens GN=ETFB PE=1 SV=3	sp P38117 ETFB_HU	28 kDa					100%		
Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	sp Q06830 PRDX1_H	22 kDa		55%	69%		67%	74%	100%

HLA class I histocompatibility antigen, A-1 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=1	sp P30443 1A01_HU	41 kDa	70%	100%	100%	74%	
ATP-dependent RNA helicase DDX39A OS=Homo sapiens GN=DDX39A PE=1 SV=2	sp O00148 DX39A_H	49 kDa		69%	100%		100% 73%
Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3	sp Q14697 GANAB_H	107 kDa	98%	70%	100%		73%
Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2	sp P12814 ACTN1_H	103 kDa	68%		69%		79% 100%
Carnitine O-palmitoyltransferase 1, liver isoform OS=Homo sapiens GN=CPT1A PE=1 SV=2	sp P50416 CPT1A_H	88 kDa		100%	100%	100%	
Lysosomal protective protein OS=Homo sapiens GN=CTSA PE=1 SV=2	sp P10619 PPGB_HU	54 kDa	100%	70%	100%		
Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM PE=1 SV=4	sp P14618 KPYM_H	58 kDa			69%	100%	100%
Serine hydroxymethyltransferase, cytosolic OS=Homo sapiens GN=SHMT1 PE=1 SV=1	sp P34896 GLYC_H	53 kDa	100%	100%	69%		
Rho GTPase-activating protein 1 OS=Homo sapiens GN=ARHGAP1 PE=1 SV=1	sp Q07960 RHG01_H	50 kDa	68%	100%	100%		
Basigin OS=Homo sapiens GN=BSG PE=1 SV=2	sp P35613 BASI_HU	42 kDa			69%	95%	100%
L-lactate dehydrogenase A-like 6A OS=Homo sapiens GN=LDHAL6A PE=2 SV=1	sp Q6ZMR3 LDH6A_H	37 kDa				100% 100%	16%
Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	sp P07900 HS90A_H	85 kDa			100%		100%
Myoferlin OS=Homo sapiens GN=MYOF PE=1 SV=1	sp Q9NZM1 MYOF_H	235 kDa			100%		100%

2-oxoglutarate dehydrogenase, mitochondrial OS=Homo sapiens GN=OGDH PE=1 SV=3	sp Q02218 ODO1_H	116 kDa	100%	100%			
Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2	sp P49411 EFTU_HU	50 kDa		100%		74%	
Guanine nucleotide-binding protein-like 3 OS=Homo sapiens GN=GNL3 PE=1 SV=2	sp Q9BVP2 GNL3_H	62 kDa		69%			100%
L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2	sp P07195 LDHB_H	37 kDa	100%		29%	17%	
Flotillin-1 OS=Homo sapiens GN=FLOT1 PE=1 SV=3	sp O75955 FLOT1_H	47 kDa		100%			
Ornithine aminotransferase, mitochondrial OS=Homo sapiens GN=OAT PE=1 SV=1	sp P04181 OAT_HU	49 kDa		100%			
Zinc finger CCCH domain-containing protein 14 OS=Homo sapiens GN=ZC3H14 PE=1 SV=1	sp Q6PJT7 ZC3HE_H	83 kDa		100%			
Elongation factor G, mitochondrial OS=Homo sapiens GN=GFM1 PE=1 SV=2	sp Q96RP9 EFGM_H	83 kDa		100%			
Structural maintenance of chromosomes protein 3 OS=Homo sapiens GN=SMC3 PE=1 SV=2	sp Q9UQE7 SMC3_H	142 kDa		100%			
Guanine deaminase OS=Homo sapiens GN=GDA PE=1 SV=1	sp Q9Y2T3 GUAD_H	51 kDa		100%			
DNA polymerase alpha subunit B OS=Homo sapiens GN=POLA2 PE=1 SV=2	sp Q14181 DPOA2_H	66 kDa					100%
Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2	sp P14866 HNRPL_H	64 kDa			97%	100%	67% 73%

Lon protease homolog, mitochondrial OS=Homo sapiens GN=LONP1 PE=1 SV=2	sp P36776 LONM_H	106 kDa	68%	70%	100%	80%		
Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	sp P81605 DCD_HU	11 kDa		70%	69%		40%	100%
Ankyrin repeat and MYND domain-containing protein 2 OS=Homo sapiens GN=ANKMY2 PE=1 SV=1	sp Q8IV38 ANKY2_	49 kDa	99%	100%	69%			
Nuclear pore membrane glycoprotein 210 OS=Homo sapiens GN=NUP210 PE=1 SV=3	sp Q8TEM1 PO210_	205 kDa	68%		100%	100%		
Exosome component 10 OS=Homo sapiens GN=EXOSC10 PE=1 SV=2	sp Q01780 EXOSX_	101 kDa	38%	100%	69%			
Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2	sp P13667 PDIA4_H	73 kDa					100%	100%
Metaxin-3 OS=Homo sapiens GN=MTX3 PE=1 SV=2	sp Q5HYI7 MTX3_I	35 kDa					100%	
Ribose-5-phosphate isomerase OS=Homo sapiens GN=RPIA PE=1 SV=3	sp P49247 RPIA_HU	33 kDa					98%	100%
Ribonucleoside-diphosphate reductase large subunit OS=Homo sapiens GN=RRM1 PE=1 SV=1	sp P23921 RIR1_HU	90 kDa		100%	69%			
DBIRD complex subunit ZNF326 OS=Homo sapiens GN=ZNF326 PE=1 SV=2	sp Q5BKZ1 ZN326_	66 kDa						100%
TAR DNA-binding protein 43 OS=Homo sapiens GN=TARDBP PE=1 SV=1	sp Q13148 TADBP_	45 kDa				100%		
Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	sp Q08380 LG3BP_I	65 kDa						100%
40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1	sp P62081 RS7_HUM	22 kDa						100%

Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	sp P32119 PRDX2_H	22 kDa							100%
Protein Hikeshi OS=Homo sapiens GN=C11orf73 PE=1 SV=2	sp Q53FT3 HIKES_H	22 kDa							100%
26S proteasome non-ATPase regulatory subunit 8 OS=Homo sapiens GN=PSMD8 PE=1 SV=2	sp P48556 PSMD8_H	40 kDa						100%	
Serine/arginine-rich splicing factor 6 OS=Homo sapiens GN=SRSF6 PE=1 SV=2	sp Q13247 SRSF6_H	40 kDa		100%					
Histone H2B type 1-C/E/F/G/I OS=Homo sapiens GN=HIST1H2BC PE=1 SV=4	sp P62807 H2B1C_H	14 kDa			98%	100%		74%	
Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLST PE=1 SV=4	sp P36957 ODO2_H	49 kDa		100%	100%	70%			
Keratin, type II cytoskeletal 3 OS=Homo sapiens GN=KRT3 PE=1 SV=3	sp P12035 K2C3_HU	64 kDa		70%	100%			100%	
tRNA (cytosine(34)-C(5))-methyltransferase OS=Homo sapiens GN=NSUN2 PE=1 SV=2	sp Q08J23 NSUN2_H	86 kDa	68%	100%	100%				
Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	sp P68363 TBA1B_H	50 kDa		70%				79%	100%
Filamin-B OS=Homo sapiens GN=FLNB PE=1 SV=2	sp O75369 FLNB_H	278 kDa		70%	100%				73%
Septin-6 OS=Homo sapiens GN=SEPT6 PE=1 SV=4	sp Q14141 SEPT6_H	50 kDa	100%	100%					
2,4-dienoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=DECR1 PE=1 SV=1	sp Q16698 DECR_H	36 kDa				100%	100%		

Programmed cell death protein 10 OS=Homo sapiens GN=PDCD10 PE=1 SV=1	sp Q9BUL8 PDC10_	25 kDa			100%	100%	
Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2	sp P25787 PSA2_HU	26 kDa				100%	77%
Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1	sp Q8WUM4 PDC6I	96 kDa	100%			74%	
Mitochondrial import receptor subunit TOM34 OS=Homo sapiens GN=TOMM34 PE=1 SV=2	sp Q15785 TOM34_	35 kDa			68%	100%	
Mammalian ependymin-related protein 1 OS=Homo sapiens GN=EPDR1 PE=1 SV=2	sp Q9UM22 EPDR1_	25 kDa				100%	62%
Nuclear RNA export factor 1 OS=Homo sapiens GN=NXF1 PE=1 SV=1	sp Q9UBU9 NXF1_	70 kDa	45%	12%			100%
Putative heat shock protein HSP90-beta 2 OS=Homo sapiens GN=HSP90AB2P PE=1 SV=2	sp Q58FF8 H90B2_	44 kDa			13%	35%	100%
SWI/SNF complex subunit SMARCC1 OS=Homo sapiens GN=SMARCC1 PE=1 SV=3	sp Q92922 SMRC1_	123 kDa					100%
39S ribosomal protein L15, mitochondrial OS=Homo sapiens GN=MRPL15 PE=1 SV=1	sp Q9P015 RM15_H	33 kDa				100%	
Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2	sp P07741 APT_HU	20 kDa					100%
Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5	sp Q13263 TIF1B_H	89 kDa					100%

DnaJ homolog subfamily A member 3, mitochondrial OS=Homo sapiens GN=DNAJA3 PE=1 SV=2	sp Q96EY1 DNJA3	52 kDa		100%				
5-formyltetrahydrofolate cyclo- ligase OS=Homo sapiens GN=MTHFS PE=1 SV=2	sp P49914 MTHFS_	23 kDa				100%		
Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1	sp Q15084 PDIA6_H	48 kDa		80%			79%	100%
Large neutral amino acids transporter small subunit 1 OS=Homo sapiens GN=SLC7A5 PE=1 SV=2	sp Q01650 LAT1_HU	55 kDa	69%	100%				73%
Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1	sp Q9BVC6 TM109	26 kDa	69%				100%	73%
Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 PE=1 SV=2	sp P11940 PABP1_H	71 kDa				100%		88% 49%
Sideroflexin-1 OS=Homo sapiens GN=SFXN1 PE=1 SV=4	sp Q9H9B4 SFXN1	36 kDa			100%	100%		
Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	sp P04075 ALDOA	39 kDa				99%	100%	
Inosine-5'-monophosphate dehydrogenase 1 OS=Homo sapiens GN=IMPDH1 PE=1 SV=2	sp P20839 IMDH1_H	55 kDa	100%	98%				
Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=1 SV=4	sp P07910 HNRPC_	34 kDa				100%		79% 18%
Enoyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=ECHS1 PE=1 SV=4	sp P30084 ECHM_H	31 kDa				100%		79%
Thioredoxin reductase 1, cytoplasmic OS=Homo sapiens GN=TXNRD1 PE=1 SV=3	sp Q16881 TRXR1_	71 kDa		70%	100%			

Keratin, type II cytoskeletal 78 OS=Homo sapiens GN=KRT78 PE=2 SV=2	sp Q8N1N4 K2C78_	57 kDa		70%		100%
Beta-arrestin-1 OS=Homo sapiens GN=ARRB1 PE=1 SV=2	sp P49407 ARRB1_1	47 kDa	100%	70%		
Probable ATP-dependent RNA helicase DDX46 OS=Homo sapiens GN=DDX46 PE=1 SV=2	sp Q7L014 DDX46_1	117 kDa	100%	70%		
Keratin, type II cytoskeletal 1b OS=Homo sapiens GN=KRT77 PE=2 SV=3	sp Q7Z794 K2C1B_1	62 kDa			69%	100%
Inhibitor of growth protein 3 OS=Homo sapiens GN=ING3 PE=1 SV=2	sp Q9NXR8 ING3_H1	47 kDa		100%	69%	
Peptidyl-prolyl cis-trans isomerase FKBP5 OS=Homo sapiens GN=FKBP5 PE=1 SV=2	sp Q13451 FKBP5_1	51 kDa		65%	100%	
Cell division cycle 5-like protein OS=Homo sapiens GN=CDC5L PE=1 SV=2	sp Q99459 CDC5L_1	92 kDa	100%	32%		
Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	sp P09211 GSTP1_H1	23 kDa				100%
Cysteine and glycine-rich protein 1 OS=Homo sapiens GN=CSRP1 PE=1 SV=3	sp P21291 CSRP1_H1	21 kDa				100%
Exosome complex component RRP4 OS=Homo sapiens GN=EXOSC2 PE=1 SV=2	sp Q13868 EXOS2_1	33 kDa			100%	
116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 PE=1 SV=1	sp Q15029 U5S1_HU	109 kDa			100%	
Keratin, type I cuticular Ha1 OS=Homo sapiens GN=KRT31 PE=2 SV=3	sp Q15323 K1H1_H1	47 kDa				100%

Vacuolar protein sorting-associated protein 37A OS=Homo sapiens GN=VPS37A PE=1 SV=1	sp Q8NEZ2 VP37A_	44 kDa	100%	
G patch domain and KOW motifs-containing protein OS=Homo sapiens GN=GPKOW PE=1 SV=2	sp Q92917 GPKOW_	52 kDa	100%	
Cell division cycle-associated 7-like protein OS=Homo sapiens GN=CDCA7L PE=1 SV=2	sp Q96GN5 CDA7L_	52 kDa		100%
RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3	sp Q9Y230 RUVB2_	51 kDa	100%	
Plasma alpha-L-fucosidase OS=Homo sapiens GN=FUCA2 PE=1 SV=2	sp Q9BTY2 FUCO2_	54 kDa	100%	
Sideroflexin-3 OS=Homo sapiens GN=SFXN3 PE=1 SV=2	sp Q9BWM7 SFXN3_	36 kDa		100%
DNA mismatch repair protein Msh2 OS=Homo sapiens GN=MSH2 PE=1 SV=1	sp P43246 MSH2_H_	105 kDa	100%	
Ribosome biogenesis protein WDR12 OS=Homo sapiens GN=WDR12 PE=1 SV=2	sp Q9GZL7 WDR12_	48 kDa	100%	
Dihydropteridine reductase OS=Homo sapiens GN=QDPR PE=1 SV=2	sp P09417 DHPR_H_	26 kDa		100%
Leucine-rich repeat-containing protein 47 OS=Homo sapiens GN=LRR47 PE=1 SV=1	sp Q8N1G4 LRC47_	63 kDa		100%
Serine--tRNA ligase, mitochondrial OS=Homo sapiens GN=SARS2 PE=1 SV=1	sp Q9NP81 SYSM_F_	58 kDa	100%	
Translation initiation factor eIF-2B subunit gamma OS=Homo sapiens GN=EIF2B3 PE=1 SV=1	sp Q9NR50 EI2BG_	50 kDa	100%	

Thiosulfate sulfurtransferase OS=Homo sapiens GN=TST PE=1 SV=4	sp Q16762 THTR_H	33 kDa			100%	
Dynactin subunit 3 OS=Homo sapiens GN=DCTN3 PE=1 SV=1	sp O75935 DCTN3_	21 kDa				100%
Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1	sp P26447 S10A4_H	12 kDa				100%
4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3	sp P08195 4F2_HUM	68 kDa				100%
Double-stranded RNA-specific adenosine deaminase OS=Homo sapiens GN=ADAR PE=1 SV=4	sp P55265 DSRAD_	136 kDa	100%			
Ferritin heavy chain OS=Homo sapiens GN=FTH1 PE=1 SV=2	sp P02794 FRIH_HU	21 kDa				100%
Exosome complex component RRP46 OS=Homo sapiens GN=EXOSC5 PE=1 SV=1	sp Q9NQT4 EXOS5_	25 kDa			100%	
Pre-mRNA 3'-end-processing factor FIP1 OS=Homo sapiens GN=FIP1L1 PE=1 SV=1	sp Q6UN15 FIP1_HU	67 kDa				100%
Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Homo sapiens GN=PDHB PE=1 SV=3	sp P11177 ODPB_HU	39 kDa				100%
Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2	sp Q12905 ILF2_HU	43 kDa		100%		
Procollagen-lysine,2- oxoglutarate 5-dioxygenase 3 OS=Homo sapiens GN=PLOD3 PE=1 SV=1	sp O60568 PLOD3_I	85 kDa	100%	80%		
RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX PE=1 SV=3	sp P38159 RBMX_H	42 kDa		80%		100%

Fumarate hydratase, mitochondrial OS=Homo sapiens GN=FH PE=1 SV=3	sp P07954 FUMH_H	55 kDa			100%	74%	
Poly(U)-binding-splicing factor PUF60 OS=Homo sapiens GN=PUF60 PE=1 SV=1	sp Q9UHX1 PUF60	60 kDa			100%		73%
Putative eukaryotic translation initiation factor 2 subunit 3-like protein OS=Homo sapiens GN=EIF2S3L PE=5 SV=2	sp Q2VIR3 IF2GL_H	51 kDa	70%	100%			
Torsin-1A-interacting protein 1 OS=Homo sapiens GN=TOR1AIP1 PE=1 SV=2	sp Q5JTV8 TOIP1_H	66 kDa				69%	100%
Serine/threonine-protein phosphatase PGAM5, mitochondrial OS=Homo sapiens GN=PGAM5 PE=1 SV=2	sp Q96HS1 PGAM5	32 kDa			100%	67%	
Regulator of chromosome condensation OS=Homo sapiens GN=RCC1 PE=1 SV=1	sp P18754 RCC1_HU	45 kDa			100%		66%
Tyrosine-protein kinase Fyn OS=Homo sapiens GN=FYN PE=1 SV=3	sp P06241 FYN_HU	61 kDa	65%	100%			
RNA-binding protein 10 OS=Homo sapiens GN=RBM10 PE=1 SV=3	sp P98175 RBM10_H	104 kDa				36%	100%
Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21 PE=1 SV=5	sp Q9NR30 DDX21	87 kDa		17%	100%		
Beta-galactosidase OS=Homo sapiens GN=GLB1 PE=1 SV=2	sp P16278 BGAL_H	76 kDa					100%
Cytochrome b-c1 complex subunit 2, mitochondrial OS=Homo sapiens GN=UQCRC2 PE=1 SV=3	sp P22695 QCR2_HU	48 kDa				100%	

ES1 protein homolog, mitochondrial OS=Homo sapiens GN=C21orf33 PE=1 SV=3	sp P30042 ES1_HUN	28 kDa		100%	
ATP synthase subunit gamma, mitochondrial OS=Homo sapiens GN=ATP5C1 PE=1 SV=1	sp P36542 ATPG_HU	33 kDa		100%	
Dolichyl- diphosphooligosaccharide-- protein glycosyltransferase 48 kDa subunit OS=Homo sapiens GN=DDOST PE=1 SV=4	sp P39656 OST48_H	51 kDa	100%		
Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4	sp P40925 MDHC_H	36 kDa		100%	
Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2	sp P49720 PSB3_HU	23 kDa			100%
Peroxisomal multifunctional enzyme type 2 OS=Homo sapiens GN=HSD17B4 PE=1 SV=3	sp P51659 DHB4_H	80 kDa		100%	
Serine/arginine-rich splicing factor 3 OS=Homo sapiens GN=SRSF3 PE=1 SV=1	sp P84103 SRSF3_H	19 kDa			100%
Serine/arginine-rich splicing factor 4 OS=Homo sapiens GN=SRSF4 PE=1 SV=2	sp Q08170 SRSF4_H	57 kDa	100%		
Cytoplasmic dynein 1 intermediate chain 2 OS=Homo sapiens GN=DYNC112 PE=1 SV=3	sp Q13409 DC112_H	71 kDa			100%
Rab-like protein 3 OS=Homo sapiens GN=RABL3 PE=1 SV=1	sp Q5HYI8 RABL3_	26 kDa		100%	
Calponin-2 OS=Homo sapiens GN=CNN2 PE=1 SV=4	sp Q99439 CNN2_H	34 kDa		100%	

COMM domain-containing protein 3 OS=Homo sapiens GN=COMMD3 PE=1 SV=1	sp Q9UBI1 COMD3	22 kDa		100%	
Ubiquilin-2 OS=Homo sapiens GN=UBQLN2 PE=1 SV=2	sp Q9UHD9 UBQL2	66 kDa			100%
Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=3	sp Q9UQ80 PA2G4	44 kDa	100%		
Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2	sp P31949 S10AB_H	12 kDa			100%
Wiskott-Aldrich syndrome protein family member 2 OS=Homo sapiens GN=WASF2 PE=1 SV=3	sp Q9Y6W5 WASF2	54 kDa			100%
Protein regulator of cytokinesis 1 OS=Homo sapiens GN=PRC1 PE=1 SV=2	sp O43663 PRC1_HU	72 kDa			100%
BH3-interacting domain death agonist OS=Homo sapiens GN=BID PE=1 SV=1	sp P55957 BID_HUM	22 kDa		100%	
Mitochondrial carrier homolog 2 OS=Homo sapiens GN=MTCH2 PE=1 SV=1	sp Q9Y6C9 MTCH2	33 kDa		100%	
Sorting nexin-2 OS=Homo sapiens GN=SNX2 PE=1 SV=2	sp O60749 SNX2_H	58 kDa			100%
Eukaryotic translation initiation factor 2 subunit 2 OS=Homo sapiens GN=EIF2S2 PE=1 SV=2	sp P20042 IF2B_HU	38 kDa	100%		
Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1	sp O14818 PSA7_HU	28 kDa		100%	
Calcium-binding mitochondrial carrier protein SCaMC-1 OS=Homo sapiens GN=SLC25A24 PE=1 SV=2	sp Q6NUK1 SCMC1	53 kDa	100%		

Receptor-type tyrosine-protein phosphatase F OS=Homo sapiens GN=PTPRF PE=1 SV=2	sp P10586 PTPRF_H	213 kDa			100%
Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1	sp Q15363 TMED2	23 kDa			100%
Alanine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2	sp P49588 SYAC_H	107 kDa	100%		
THO complex subunit 7 homolog OS=Homo sapiens GN=THOC7 PE=1 SV=3	sp Q619Y2 THOC7	24 kDa			100%
Probable RNA-binding protein 23 OS=Homo sapiens GN=RBM23 PE=1 SV=1	sp Q86U06 RBM23	49 kDa			100%
Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1 PE=1 SV=2	sp O14773 TPP1_H	61 kDa	100%		
Protein phosphatase 1 regulatory subunit 12A OS=Homo sapiens GN=PPP1R12A PE=1 SV=1	sp O14974 MYPT1	115 kDa			100%
Gamma-glutamylcyclotransferase OS=Homo sapiens GN=GGCT PE=1 SV=1	sp O75223 GGCT_H	21 kDa		100%	
NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial OS=Homo sapiens GN=NDUFS2 PE=1 SV=2	sp O75306 NDUS2	53 kDa	100%		
Pre-mRNA-processing factor 40 homolog A OS=Homo sapiens GN=PRPF40A PE=1 SV=2	sp O75400 PR40A_H	109 kDa			100%
Heat shock 70 kDa protein 4L OS=Homo sapiens GN=HSPA4L PE=1 SV=3	sp O95757 HS74L_H	95 kDa			100%

Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1	sp P00403 COX2_HU	26 kDa			100%
Salivary acidic proline-rich phosphoprotein 1/2 OS=Homo sapiens GN=PRH1 PE=1 SV=2	sp P02810 PRPC_HU	17 kDa			100%
Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2	sp P04792 HSPB1_HU	23 kDa		100%	
Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4	sp P06744 G6PI_HU	63 kDa		100%	
Dihydrolipoyllysine-residue acetyltransferase component of pyruvate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLAT PE=1 SV=3	sp P10515 ODP2_HU	69 kDa			100%
Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase, mitochondrial OS=Homo sapiens GN=MTHFD2 PE=1 SV=2	sp P13995 MTDC_HU	38 kDa		100%	
Zinc finger protein RFP OS=Homo sapiens GN=TRIM27 PE=1 SV=1	sp P14373 TRI27_HU	58 kDa	100%		
Small nuclear ribonucleoprotein-associated proteins B and B' OS=Homo sapiens GN=SNRNPB PE=1 SV=2	sp P14678 RSMB_HU	25 kDa		100%	
V-type proton ATPase subunit B, kidney isoform OS=Homo sapiens GN=ATP6V1B1 PE=1 SV=3	sp P15313 VATB1_HU	57 kDa	100%		
60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3	sp P18621 RL17_HU	21 kDa			100%

V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3	sp P21281 VATB2_H	57 kDa	100%	
DNA replication licensing factor MCM7 OS=Homo sapiens GN=MCM7 PE=1 SV=4	sp P33993 MCM7_H	81 kDa	100%	
Flap endonuclease 1 OS=Homo sapiens GN=FEN1 PE=1 SV=1	sp P39748 FEN1_HU	43 kDa		100%
Adenylyl cyclase-associated protein 2 OS=Homo sapiens GN=CAP2 PE=1 SV=1	sp P40123 CAP2_HU	53 kDa	100%	
Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1	sp P46060 RAGP1_H	64 kDa		100%
Serpin B8 OS=Homo sapiens GN=SERPINB8 PE=1 SV=2	sp P50452 SPB8_HU	43 kDa	100%	
Transcription activator BRG1 OS=Homo sapiens GN=SMARCA4 PE=1 SV=2	sp P51532 SMCA4_H	185 kDa		100%
Translocon-associated protein subunit delta OS=Homo sapiens GN=SSR4 PE=1 SV=1	sp P51571 SSRD_HU	19 kDa		100%
Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5	sp Q00610 CLH1_HU	192 kDa		100%
Kinesin heavy chain isoform 5A OS=Homo sapiens GN=KIF5A PE=1 SV=2	sp Q12840 KIF5A_H	117 kDa		100%
Plastin-1 OS=Homo sapiens GN=PLS1 PE=1 SV=2	sp Q14651 PLSI_HU	70 kDa		100%
Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2	sp Q15365 PCBP1_H	37 kDa	100%	
DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1	sp Q16531 DDB1_HU	127 kDa		100%

Cleavage and polyadenylation specificity factor subunit 6 OS=Homo sapiens GN=CPSF6 PE=1 SV=2	sp Q16630 CPSF6_H	59 kDa			100%
Proteasome subunit alpha type-7-like OS=Homo sapiens GN=PSMA8 PE=1 SV=3	sp Q8TAA3 PSA7L	29 kDa		100%	
Phosphatidylglycerophosphatase and protein-tyrosine phosphatase 1 OS=Homo sapiens GN=PTPMT1 PE=1 SV=1	sp Q8WUK0 PTPM1	23 kDa			100%
DnaJ homolog subfamily B member 3 OS=Homo sapiens GN=DNAJB3 PE=1 SV=1	sp Q8WWF6 DNJB3	17 kDa		100%	
Protein NDRG1 OS=Homo sapiens GN=NDRG1 PE=1 SV=1	sp Q92597 NDRG1	43 kDa	100%		
Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2	sp Q92973 TNPO1_I	102 kDa	100%		
Serpin B12 OS=Homo sapiens GN=SERPINB12 PE=1 SV=1	sp Q96P63 SPB12_H	46 kDa		100%	
Intraflagellar transport protein 27 homolog OS=Homo sapiens GN=IFT27 PE=1 SV=1	sp Q9BW83 IFT27_I	20 kDa			100%
Inosine triphosphate pyrophosphatase OS=Homo sapiens GN=ITPA PE=1 SV=2	sp Q9BY32 ITPA_H	21 kDa			100%
Ubiquitin-conjugating enzyme E2 O OS=Homo sapiens GN=UBE2O PE=1 SV=3	sp Q9C0C9 UBE2O	141 kDa			100%
39S ribosomal protein L44, mitochondrial OS=Homo sapiens GN=MRPL44 PE=1 SV=1	sp Q9H9J2 RM44_H	38 kDa		100%	
Probable Xaa-Pro aminopeptidase 3 OS=Homo sapiens GN=XPNPEP3 PE=1 SV=1	sp Q9NQH7 XPP3_I	57 kDa	100%		

GTP-binding protein SAR1a OS=Homo sapiens GN=SAR1A PE=1 SV=1	sp Q9NR31 SAR1A_	22 kDa		100%	
Thyroid hormone receptor- associated protein 3 OS=Homo sapiens GN=THRAP3 PE=1 SV=2	sp Q9Y2W1 TR150_	109 kDa			100%
Non-structural maintenance of chromosomes element 1 homolog OS=Homo sapiens GN=NSMCE1 PE=1 SV=5	sp Q8WV22 NSE1_I	31 kDa	100%		