Suppl. Table 2A: Total number of proteins detected in THP-1 cells infected with only DV2 (score calculated from 2 biological replicates and 3 technical replicates)

Entry	MASCOT Score	Mass	Sequences coverage	Description
CH60_HUMAN	1645	61187	24 (16)	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2
ACTB_HUMAN	1622	42052	21 (18)	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1
ACTC_HUMAN	584	42334	14 (10)	Actin, alpha cardiac muscle 1 OS=Homo sapiens GN=ACTC1 PE=1 SV=1
ATPB_HUMAN	937	56525	19 (16)	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3
PDIA1_HUMAN	876	57480	20 (14)	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3
ATPA_HUMAN	743	59828	17 (13)	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1
CAP1_HUMAN	640	52222	20 (14)	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=4
TBA1B_HUMAN	563	50804	12 (8)	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1
GRP75_HUMAN	560	73920	13 (7)	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2
K2C1_HUMAN	529	66170	21 (15)	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6
K22E_HUMAN	122	65678	8 (5)	Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2
TBB5_HUMAN	513	50095	17 (11)	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2
ENOA_HUMAN	507	47481	19 (12)	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2
K1C10_HUMAN	478	59020	17 (13)	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6
K1C9_HUMAN	370	62255	12 (11)	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3
EF1A1_HUMAN	470	50451	12 (9)	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1
H4_HUMAN	441	11360	5 (5)	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2
ROA2_HUMAN	433	37464	14 (10)	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2
ROA1_HUMAN	421	38837	10 (7)	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5
HSP7C_HUMAN	428	71082	12 (9)	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1
GRP78_HUMAN	386	72402	18 (10)	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2
PDIA6_HUMAN	311	48490	6 (4)	Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1
HNRPU_HUMAN	290	91269	12 (8)	Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6
VDAC1_HUMAN	285	30868	11 (7)	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2
H2B1B_HUMAN	285	13942	5 (3)	Histone H2B type 1-B OS=Homo sapiens GN=HIST1H2BB PE=1 SV=2
PROF1_HUMAN	277	15216	6 (4)	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2
G3P_HUMAN	272	36201	7 (5)	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3
EF2_HUMAN	253	96246	8 (3)	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4
CALX_HUMAN	252	67982	11 (6)	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2
ENPL_HUMAN	250	92696	13 (8)	Endoplasmin OS=Homo sapiens GN=HSP90B1 PE=1 SV=1
HS90B_HUMAN	224	83554	16 (8)	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4
HS90A_HUMAN	104	85006	13 (5)	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5
TRAP1_HUMAN	45	80345	3 (2)	Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3
ECH1_HUMAN	237	36136	6 (4)	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH1 PE=1 SV=2
PDIA3_HUMAN	220	57146	13 (9)	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4
PPIA_HUMAN	215	18229	7 (6)	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2
IMB1_HUMAN	210	98420	6 (2)	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2
CALR_HUMAN	201	48283	8 (5)	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1
KAD2_HUMAN	200	26689	6 (3)	Adenylate kinase 2, mitochondrial OS=Homo sapiens GN=AK2 PE=1 SV=2
ALBU_HUMAN	190	71317	5 (4)	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2
KPYM_HUMAN	190	58470	9 (4)	Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM2 PE=1 SV=4
ALDOA_HUMAN	189	39851	13 (6)	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2
MDHM_HUMAN	186	35937	10 (3)	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3
LDHB_HUMAN	185	36900	8 (4)	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2

LDHA_HUMAN	136	36950	5 (4)	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2
TCPD_HUMAN	178	58401	5 (4)	T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4
CAH2_HUMAN	175	29285	6 (2)	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2
TKT_HUMAN	170	68519	5 (4)	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3
HNRPK_HUMAN	169	51230	5 (3)	Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1
PGK1_HUMAN	163	44985	5 (3)	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3
PPT1_HUMAN	162	34627	6 (4)	Palmitoyl-protein thioesterase 1 OS=Homo sapiens GN=PPT1 PE=1 SV=1
LMAN2_HUMAN	156	40545	7 (3)	Vesicular integral-membrane protein VIP36 OS=Homo sapiens GN=LMAN2 PE=1 SV=1
GANAB_HUMAN	151	107263	7 (4)	Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3
RNT2_HUMAN	148	30089	7 (4)	Ribonuclease T2 OS=Homo sapiens GN=RNASET2 PE=1 SV=2
1433Z_HUMAN	144	27899	6 (4)	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1
1433E_HUMAN	48	29326	4 (2)	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1
1433G_HUMAN	45	28456	3 (2)	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2
H2A1D_HUMAN	141	14099	5 (3)	Histone H2A type 1-D OS=Homo sapiens GN=HIST1H2AD PE=1 SV=2
SERA_HUMAN	139	57356	5 (3)	D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4
RL14_HUMAN	132	23531	2 (2)	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4
KU70_HUMAN	130	70084	9 (4)	ATP-dependent DNA helicase 2 subunit 1 OS=Homo sapiens GN=XRCC6 PE=1 SV=2
ANXA5_HUMAN	130	35971	6 (3)	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2
PSME2_HUMAN	129	27515	3 (1)	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=3
TCPE_HUMAN	129	60089	5 (3)	T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=1 SV=1
ADT2_HUMAN	126	33102	5 (3)	ADP/ATP translocase 2 OS=Homo sapiens GN=SLC25A5 PE=1 SV=6
MOES_HUMAN	124	67892	13 (6)	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3
PLSL_HUMAN	123	70815	6 (3)	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=5
DHSA_HUMAN	117	73672	2 (2)	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=2
TCPA_HUMAN	117	60819	8 (4)	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1
NPM_HUMAN	113	32726	2 (1)	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2
LMNB1_HUMAN	111	66653	3 (3)	Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2
ATPG_HUMAN	109	33032	5 (3)	ATP synthase subunit gamma, mitochondrial OS=Homo sapiens GN=ATP5C1 PE=1 SV=1
RPN1_HUMAN	109	68641	5 (3)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1
IDH3A_HUMAN	107	40022	2 (1)	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Homo sapiens GN=IDH3A PE=1 SV=1
H32_HUMAN	107	15436	8 (4)	Histone H3.2 OS=Homo sapiens GN=HIST2H3A PE=1 SV=3
VDAC2_HUMAN	105	32060	5 (2)	Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1 SV=2
PTBP1_HUMAN	105	57357	5 (3)	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1
ARPC4_HUMAN	104	19768	3 (1)	Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3
HSPB1_HUMAN	101	22826	7 (3)	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2
PRDX1_HUMAN	98	22324	8 (3)	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1
6PGD_HUMAN	97	53619	3 (1)	6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=1 SV=3
UBIQ_HUMAN	96	8560	3 (2)	Ubiquitin OS=Homo sapiens GN=RPS27A PE=1 SV=1
DDX5_HUMAN	94	69618	5 (2)	Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1
HNRPM_HUMAN	92	77749	8 (3)	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3
RL6_HUMAN	91	32765	3 (3)	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3
PRDX3_HUMAN	91	28017	3 (1)	Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3 PE=1 SV=3
1A02_HUMAN	90	41181	3 (2)	HLA class I histocompatibility antigen, A-2 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=1
COMT_HUMAN	90	30474	3 (3)	Catechol O-methyltransferase OS=Homo sapiens GN=COMT PE=1 SV=2
ACTN1_HUMAN	88	103563	7 (3)	Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2
TPP1_HUMAN	87	61723	4 (2)	Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1 PE=1 SV=2
GLYM_HUMAN	87	56414	7 (2)	Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3
SFPQ_HUMAN	87	76216	5 (3)	Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2

COR1A_HUMAN	87	51678	3 (2)	Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4
RL7_HUMAN	87	29264	4 (2)	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1
RS24_HUMAN	84	15413	1 (1)	40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=1 SV=1
CCD47_HUMAN	83	56123	1 (1)	Coiled-coil domain-containing protein 47 OS=Homo sapiens GN=CCDC47 PE=1 SV=1
RL12_HUMAN	82	17979	4 (2)	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1
PRTN3_HUMAN	82	28245	3 (3)	Myeloblastin OS=Homo sapiens GN=PRTN3 PE=1 SV=3
COF1_HUMAN	78	18719	5 (3)	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3
IF4A1_HUMAN	77	46353	3 (2)	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1
FUBP1_HUMAN	77	67690	1 (1)	Far upstream element-binding protein 1 OS=Homo sapiens GN=FUBP1 PE=1 SV=3
NUCL_HUMAN	76	76625	6 (3)	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3
STML2_HUMAN	76	38624	2 (1)	Stomatin-like protein 2 OS=Homo sapiens GN=STOML2 PE=1 SV=1
CH10_HUMAN	74	10925	2 (2)	10 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPE1 PE=1 SV=2
NONO_HUMAN	73	54311	5 (1)	Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV=4
CY1_HUMAN	71	35709	3 (1)	Cytochrome c1, heme protein, mitochondrial OS=Homo sapiens GN=CYC1 PE=1 SV=2
LAMP1_HUMAN	71	45367	5 (2)	Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3
OST48_HUMAN	68	50940	6 (4)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Homo sapiens GN=DDOST PE=1 SV
LAMP2_HUMAN	68	45503	3 (1)	Lysosome-associated membrane glycoprotein 2 OS=Homo sapiens GN=LAMP2 PE=1 SV=2
RAP1B_HUMAN	68	21040	5 (1)	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1
RL27A_HUMAN	67	16665	1 (1)	60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=1 SV=2
DLDH_HUMAN	66	54713	4 (2)	Dihydrolipoyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2
PHB2_HUMAN	66	33276	4 (2)	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2
ETFA_HUMAN	65	35400	2 (1)	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Homo sapiens GN=ETFA PE=1 SV=1
LPPRC_HUMAN	65	159003	5 (2)	Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3
HEXB_HUMAN	65	63527	5 (2)	Beta-hexosaminidase subunit beta OS=Homo sapiens GN=HEXB PE=1 SV=3
RAB10_HUMAN	65	22755	3 (3)	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1
ANXA6_HUMAN	64	76168	3 (2)	Annexin A6 OS=Homo sapiens GN=ANXA6 PE=1 SV=3
MPPA_HUMAN	58	58729	3 (1)	Mitochondrial-processing peptidase subunit alpha OS=Homo sapiens GN=PMPCA PE=1 SV=2
EFTU_HUMAN	57	49852	6 (2)	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2
GBLP_HUMAN	56	35511	6 (2)	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3
SYAC_HUMAN	56	107484	4 (2)	Alanyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2
DCXR_HUMAN	56	26182	2 (1)	L-xylulose reductase OS=Homo sapiens GN=DCXR PE=1 SV=2
ARP3_HUMAN	55	47797	5 (2)	Actin-related protein 3 OS=Homo sapiens GN=ACTR3 PE=1 SV=3
TCPB_HUMAN	54	57794	6 (1)	T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4
PABP2_HUMAN	53	32843	2 (1)	Polyadenylate-binding protein 2 OS=Homo sapiens GN=PABPN1 PE=1 SV=3
IF5A1_HUMAN	53	17049	2 (1)	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2
RL18_HUMAN	53	21735	3 (1)	60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2
CLH1_HUMAN	52	193260	4 (2)	Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5
VAT1_HUMAN	52	42122	3 (1)	Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2
ECHB_HUMAN	52	51547	5 (1)	Trifunctional enzyme subunit beta, mitochondrial OS=Homo sapiens GN=HADHB PE=1 SV=3
FUBP2_HUMAN	52	73443	1 (1)	Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=3
RL13_HUMAN	52	24304	4 (1)	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4
ILF2_HUMAN	51	43263	4 (3)	Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2
PGAM1_HUMAN	51	28900	2 (1)	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2
ARPC2_HUMAN	51	34426	4 (3)	Actin-related protein 2/3 complex subunit 2 OS=Homo sapiens GN=ARPC2 PE=1 SV=1
KU86_HUMAN	51	83222	6 (1)	ATP-dependent DNA helicase 2 subunit 2 OS=Homo sapiens GN=XRCC5 PE=1 SV=3
RSSA_HUMAN	51	32947	1 (1)	40S ribosomal protein SA OS=Homo sapiens GN=RPSA PE=1 SV=4
GDIB_HUMAN	50	51087	5 (2)	Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GDI2 PE=1 SV=2
RS13_HUMAN	50	17212	3 (2)	40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2

PHB_HUMAN	50	29843	4 (2)	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1
RL22_HUMAN	50	14835	3 (1)	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2
TLN1_HUMAN	50	271766	6 (1)	Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3
CISY_HUMAN	50	51908	2 (1)	Citrate synthase, mitochondrial OS=Homo sapiens GN=CS PE=1 SV=2
GNAI2_HUMAN	49	40995	4 (2)	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3
RL3_HUMAN	49	46365	2 (1)	60S ribosomal protein L3 OS=Homo sapiens GN=RPL3 PE=1 SV=2
TMED2_HUMAN	48	22860	1 (1)	Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1
SET_HUMAN	48	33469	3 (1)	Protein SET OS=Homo sapiens GN=SET PE=1 SV=3
RS3_HUMAN	48	26842	3 (2)	40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2
DHB4_HUMAN	48	80092	5 (3)	Peroxisomal multifunctional enzyme type 2 OS=Homo sapiens GN=HSD17B4 PE=1 SV=3
RB27A_HUMAN	47	25137	3 (1)	Ras-related protein Rab-27A OS=Homo sapiens GN=RAB27A PE=1 SV=3
TCPZ_HUMAN	47	58444	4 (2)	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3
PPIB_HUMAN	46	23785	2 (1)	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2
HNRPD_HUMAN	45	38581	4 (2)	Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD PE=1 SV=1
RS6_HUMAN	44	28834	2 (1)	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1
SYTC_HUMAN	44	84294	4 (1)	Threonyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=TARS PE=1 SV=3
FAS_HUMAN	44	275877	5 (1)	Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3
CPT2_HUMAN	43	74244	3 (1)	Carnitine O-palmitoyltransferase 2, mitochondrial OS=Homo sapiens GN=CPT2 PE=1 SV=2
CATG_HUMAN	43	29161	3 (2)	Cathepsin G OS=Homo sapiens GN=CTSG PE=1 SV=2
EF1G_HUMAN	43	50429	5 (2)	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3
ASSY_HUMAN	42	46786	1 (1)	Argininosuccinate synthase OS=Homo sapiens GN=ASS1 PE=1 SV=2
RL30_HUMAN	41	12947	2 (1)	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2
ROA0_HUMAN	41	30993	1 (1)	Heterogeneous nuclear ribonucleoprotein A0 OS=Homo sapiens GN=HNRNPA0 PE=1 SV=1
PCBP1_HUMAN	40	37987	1 (1)	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2
AMPB_HUMAN	40	73234	2 (2)	Aminopeptidase B OS=Homo sapiens GN=RNPEP PE=1 SV=2
PSB3_HUMAN	40	23219	1 (1)	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2
U2AF2_HUMAN	39	53809	2 (1)	Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2 PE=1 SV=4
RHOC_HUMAN	39	22334	2 (1)	Rho-related GTP-binding protein RhoC OS=Homo sapiens GN=RHOC PE=1 SV=1
HNRH1_HUMAN	38	49484	3 (1)	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4
ANXA2_HUMAN	38	38808	3 (2)	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2
SSRA_HUMAN	38	32215	1 (1)	Translocon-associated protein subunit alpha OS=Homo sapiens GN=SSR1 PE=1 SV=3
HNRPC_HUMAN	38	33707	4 (1)	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=1 SV=4
PYRG1_HUMAN	38	67332	1 (1)	CTP synthase 1 OS=Homo sapiens GN=CTPS PE=1 SV=2
ECHM_HUMAN	37	31823	2 (1)	Enoyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=ECHS1 PE=1 SV=4
LONM_HUMAN	37	106936	3 (1)	Lon protease homolog, mitochondrial OS=Homo sapiens GN=LONP1 PE=1 SV=2
TOM70_HUMAN	37	68096	2 (1)	Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70A PE=1 SV=1
DHE3_HUMAN	37	61701	2 (1)	Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2
HBB_HUMAN	37	16102	1 (1)	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2
GBB1_HUMAN	37	38151	4 (1)	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3
NDUS2_HUMAN	37	52911	2 (1)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial OS=Homo sapiens GN=NDUFS2 PE=1 SV=2
APT_HUMAN	37	19766	1 (1)	Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2
UAP56_HUMAN	37	49416	4 (1)	Spliceosome RNA helicase BAT1 OS=Homo sapiens GN=BAT1 PE=1 SV=1
RL7A_HUMAN	36	30148	1 (1)	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2
LMNA_HUMAN	36	74380	6 (1)	Lamin-A/C OS=Homo sapiens GN=LMNA PE=1 SV=1
CH3L1_HUMAN	36	42998	3 (2)	Chitinase-3-like protein 1 OS=Homo sapiens GN=CHI3L1 PE=1 SV=2
CLIC1_HUMAN	36	27248	2 (1)	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4
RL15_HUMAN	35	24245	2 (1)	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2
IPYR2_HUMAN	35	38409	3 (1)	Inorganic pyrophosphatase 2, mitochondrial OS=Homo sapiens GN=PPA2 PE=1 SV=2

RL10L_HUMAN	35	24959	1 (1)	60S ribosomal protein L10-like OS=Homo sapiens GN=RPL10L PE=1 SV=3
RS10_HUMAN	34	18886	1 (1)	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1
C1QBP_HUMAN	34	31742	3 (1)	Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QBP PE=1 SV=1
SYPL1_HUMAN	34	28889	1 (1)	Synaptophysin-like protein 1 OS=Homo sapiens GN=SYPL1 PE=1 SV=1
GLP2R_HUMAN	34	63873	3 (1)	Glucagon-like peptide 2 receptor OS=Homo sapiens GN=GLP2R PE=1 SV=1
RS4X_HUMAN	34	29807	2 (1)	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2
RS11_HUMAN	34	18590	1 (1)	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3
FABP5_HUMAN	34	15497	1 (1)	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3
PRKDC_HUMAN	34	473749	12 (1)	DNA-dependent protein kinase catalytic subunit OS=Homo sapiens GN=PRKDC PE=1 SV=3
ANXA4_HUMAN	33	36088	5 (1)	Annexin A4 OS=Homo sapiens GN=ANXA4 PE=1 SV=4
ECHA_HUMAN	33	83688	8 (2)	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2
ERP29_HUMAN	33	29032	2 (1)	Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=1 SV=4
G6PI_HUMAN	33	63335	1 (1)	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4
RLA0_HUMAN	33	34423	8 (1)	60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1
COX2_HUMAN	32	25719	2 (1)	Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1
GGH_HUMAN	32	36340	2 (1)	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2
SMD1_HUMAN	32	13273	2 (1)	Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1
PECA1_HUMAN	32	83396	1 (1)	Platelet endothelial cell adhesion molecule OS=Homo sapiens GN=PECAM1 PE=1 SV=1
ALDH2_HUMAN	32	56859	5 (1)	Aldehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH2 PE=1 SV=2
4F2_HUMAN	32	68180	2 (1)	4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3
RS7_HUMAN	32	22113	3 (1)	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1
ACLY_HUMAN	32	121674	3 (1)	ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3
MA2B1_HUMAN	31	114357	2 (1)	Lysosomal alpha-mannosidase OS=Homo sapiens GN=MAN2B1 PE=1 SV=3
IDHP_HUMAN	31	51333	3 (1)	Isocitrate dehydrogenase [NADP], mitochondrial OS=Homo sapiens GN=IDH2 PE=1 SV=2
RPN2_HUMAN	30	69355	3 (1)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo sapiens GN=RPN2 PE=1 SV=3
MCM6_HUMAN	30	93801	2 (1)	DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1
HCD2_HUMAN	30	27134	1 (1)	3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3
RS18_HUMAN	30	17708	4 (1)	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3
PGM2_HUMAN	30	68754	3 (1)	Phosphoglucomutase-2 OS=Homo sapiens GN=PGM2 PE=1 SV=4
PNPH_HUMAN	30	32325	3 (1)	Purine nucleoside phosphorylase OS=Homo sapiens GN=NP PE=1 SV=2
HBA_HUMAN	30	15305	1 (1)	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2
THIL_HUMAN	29	45456	5 (2)	Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1
RS3A_HUMAN	29	30154	5 (2)	40S ribosomal protein S3a OS=Homo sapiens GN=RPS3A PE=1 SV=2
H15_HUMAN	28	22566	1 (1)	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3
DHX9_HUMAN	28	142181	6 (1)	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4
M2OM_HUMAN	28	34211	2 (1)	Mitochondrial 2-oxoglutarate/malate carrier protein OS=Homo sapiens GN=SLC25A11 PE=1 SV=3
RAB5C_HUMAN	28	23696	4 (1)	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2
CAC1E_HUMAN	28	263617	6 (1)	Voltage-dependent R-type calcium channel subunit alpha-1E OS=Homo sapiens GN=CACNA1E PE=1 SV=3
COTL1_HUMAN	27	16049	2 (1)	Coactosin-like protein OS=Homo sapiens GN=COTL1 PE=1 SV=3
O2T35_HUMAN	27	36704	1 (1)	Olfactory receptor 2T35 OS=Homo sapiens GN=OR2T35 PE=2 SV=1
TSP1_HUMAN	27	133291	5 (1)	Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2
ARF1_HUMAN	27	20741	4 (1)	ADP-ribosylation factor 1 OS=Homo sapiens GN=ARF1 PE=1 SV=2
SCOT1_HUMAN	26	56578	1 (1)	Succinyl-CoA:3-ketoacid-coenzyme A transferase 1, mitochondrial OS=Homo sapiens GN=OXCT1 PE=1 SV=1
TMX1_HUMAN	26	32170	1 (1)	Thioredoxin-related transmembrane protein 1 OS=Homo sapiens GN=TMX1 PE=1 SV=1
SYEP_HUMAN	26	172080	2 (1)	Bifunctional aminoacyl-tRNA synthetase OS=Homo sapiens GN=EPRS PE=1 SV=5
MYH9_HUMAN	25	227646	5 (2)	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4
HCDH_HUMAN	25	34313	2 (1)	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=1 SV=2
TCPG_HUMAN	25	61066	3 (1)	T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4

ASAP1_HUMAN	24	126363	3 (1)	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 1 OS=Homo sapiens GN=ASAP1 PE=1 SV=3
IF4G1_HUMAN	24	176168	5 (1)	Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1 PE=1 SV=3
ROAA_HUMAN	23	36316	2 (1)	Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=1 SV=2
PUR6_HUMAN	23	47790	2 (1)	Multifunctional protein ADE2 OS=Homo sapiens GN=PAICS PE=1 SV=3
LYSC_HUMAN	23	16982	3 (1)	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1
SPB8_HUMAN	22	43309	3 (1)	Serpin B8 OS=Homo sapiens GN=SERPINB8 PE=1 SV=2
FBRL_HUMAN	22	33877	2 (1)	rRNA 2'-O-methyltransferase fibrillarin OS=Homo sapiens GN=FBL PE=1 SV=2
RS25_HUMAN	22	13791	2 (1)	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1
PSA7_HUMAN	21	28041	1 (1)	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1
HNRPQ_HUMAN	19	69788	7 (2)	Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2
TCF15_HUMAN	21	21146	1 (1)	Transcription factor 15 OS=Homo sapiens GN=TCF15 PE=2 SV=3
H13_HUMAN	20	22336	4 (1)	Histone H1.3 OS=Homo sapiens GN=HIST1H1D PE=1 SV=2
SODM_HUMAN	19	24878	1 (1)	Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2
MPCP_HUMAN	19	40525	2 (1)	Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2
TCPH_HUMAN	18	59842	4 (1)	T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2
AMPN_HUMAN	18	109870	2 (1)	Aminopeptidase N OS=Homo sapiens GN=ANPEP PE=1 SV=4
ITB2_HUMAN	18	87976	5 (1)	Integrin beta-2 OS=Homo sapiens GN=ITGB2 PE=1 SV=2
HNRPG_HUMAN	17	42306	5 (1)	Heterogeneous nuclear ribonucleoprotein G OS=Homo sapiens GN=RBMX PE=1 SV=3
VIME_HUMAN	17	53676	2 (1)	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4
UT2_HUMAN	16	102213	1 (1)	Urea transporter 2 OS=Homo sapiens GN=SLC14A2 PE=1 SV=2
MCCB_HUMAN	14	61808	2 (1)	Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=MCCC2 PE=1 SV=1
S22AO_HUMAN	14	36650	1 (1)	Solute carrier family 22 member 24 OS=Homo sapiens GN=SLC22A24 PE=2 SV=1
PSA4_HUMAN	14	29750	1 (1)	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1

Suppl. Table 2B: Total number of proteins detected in THP-1 cells treated with DV2-activated platelets (score calculated from 2 biological replicates and 3 technical replicates)

Entry	MASCOT Score	Mass	Sequences coverage	Description
ACTB_HUMAN	1666	42052	24 (21)	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1
ACTC_HUMAN	908	42334	14 (13)	Actin, alpha cardiac muscle 1 OS=Homo sapiens GN=ACTC1 PE=1 SV=1
PPIA_HUMAN	1542	18229	11 (9)	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2
CH60_HUMAN	1527	61187	26 (21)	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2
H2B1D_HUMAN	1359	13928	9 (6)	Histone H2B type 1-D OS=Homo sapiens GN=HIST1H2BD PE=1 SV=2
H2B1B_HUMAN	1356	13942	7 (6)	Histone H2B type 1-B OS=Homo sapiens GN=HIST1H2BB PE=1 SV=2
ATPB_HUMAN	1069	56525	19 (16)	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3
K1C10_HUMAN	1065	59020	21 (17)	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6
K2C1_HUMAN	1058	66170	24 (19)	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6
K22E_HUMAN	333	65678	12 (9)	Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2
H4_HUMAN	993	11360	13 (11)	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2
H2A2A_HUMAN	909	14087	10 (7)	Histone H2A type 2-A OS=Homo sapiens GN=HIST2H2AA3 PE=1 SV=3
H2A2B_HUMAN	656	13987	4 (3)	Histone H2A type 2-B OS=Homo sapiens GN=HIST2H2AB PE=1 SV=3
H2A1B_HUMAN	643	14127	6 (4)	Histone H2A type 1-B/E OS=Homo sapiens GN=HIST1H2AB PE=1 SV=2
H2AV_HUMAN	557	13501	7 (4)	Histone H2A.V OS=Homo sapiens GN=H2AFV PE=1 SV=3
CH10_HUMAN	903	10925	9 (7)	10 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPE1 PE=1 SV=2
TAGL2_HUMAN	868	22548	14 (12)	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3
NPM_HUMAN	815	32726	9 (6)	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2
PDIA1_HUMAN	787	57480	15 (10)	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3
RS5_HUMAN	768	23033	11 (9)	40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4
TBB5_HUMAN	729	50095	18 (14)	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2
TBB2C_HUMAN	507	50255	15 (12)	Tubulin beta-2C chain OS=Homo sapiens GN=TUBB2C PE=1 SV=1
1433Z_HUMAN	705	27899	15 (10)	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1
1433B_HUMAN	370	28179	10 (7)	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3
1433T_HUMAN	245	28032	8 (5)	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1
1433G_HUMAN	123	28456	6 (3)	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2
1433F_HUMAN	110	28372	4 (3)	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4
HSPB1_HUMAN	659	22826	14 (13)	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2
CAP1_HUMAN	636	52222	18 (13)	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=4
KAD2_HUMAN	614	26689	11 (9)	Adenylate kinase 2, mitochondrial OS=Homo sapiens GN=AK2 PE=1 SV=2
PROF1_HUMAN	612	15216	8 (8)	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2
ATPA_HUMAN	610	59828	13 (10)	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1
TBA1B_HUMAN	591	50804	12 (9)	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1
NDKB_HUMAN	518	17401	9 (9)	Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2 PE=1 SV=1
NDKA_HUMAN	500	17309	9 (7)	Nucleoside diphosphate kinase A OS=Homo sapiens GN=NME1 PE=1 SV=1
ATPO_HUMAN	500	23377	10 (8)	ATP synthase subunit O, mitochondrial OS=Homo sapiens GN=ATP5O PE=1 SV=1
CALX_HUMAN	500	67982	10 (9)	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2
EF1A3_HUMAN	481	50495	12 (9)	Putative elongation factor 1-alpha-like 3 OS=Homo sapiens GN=EEF1AL3 PE=5 SV=1
HMGB1_HUMAN	462	25049	10 (6)	High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3

HMGB2_HUMAN	359	24190	8 (5)	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2
LEG1_HUMAN	441	15048	9 (7)	Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2
K1C9_HUMAN	422	62255	13 (7)	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3
RL12_HUMAN	420	17979	5 (5)	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1
RNT2_HUMAN	384	30089	7 (4)	Ribonuclease T2 OS=Homo sapiens GN=RNASET2 PE=1 SV=2
RAP1A_HUMAN	383	21316	6 (5)	Ras-related protein Rap-1A OS=Homo sapiens GN=RAP1A PE=1 SV=1
G3P_HUMAN	379	36201	10 (6)	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3
TCP4_HUMAN	371	14386	7 (6)	Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SUB1 PE=1 SV=3
COX5A_HUMAN	359	16923	10 (7)	Cytochrome c oxidase subunit 5A, mitochondrial OS=Homo sapiens GN=COX5A PE=1 SV=2
GRP75_HUMAN	337	73920	12 (8)	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2
MYL6_HUMAN	331	17090	8 (8)	Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2
H12_HUMAN	325	21352	10 (5)	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2
ALBU_HUMAN	321	71317	7 (5)	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2
COF1_HUMAN	318	18719	7 (7)	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3
HBA_HUMAN	315	15305	7 (7)	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2
RLA2_HUMAN	313	11658	4 (4)	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1
CYC_HUMAN	306	11855	7 (5)	Cytochrome c OS=Homo sapiens GN=CYCS PE=1 SV=2
H32_HUMAN	305	15436	7 (5)	Histone H3.2 OS=Homo sapiens GN=HIST2H3A PE=1 SV=3
H33_HUMAN	186	15376	7 (5)	Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2
H31_HUMAN	173	15509	7 (5)	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2
RS18_HUMAN	304	17708	9 (7)	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3
ATP5H_HUMAN	295	18537	11 (8)	ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3
HS90B_HUMAN	294	83554	11 (7)	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4
ENPL_HUMAN	154	92696	3 (3)	Endoplasmin OS=Homo sapiens GN=HSP90B1 PE=1 SV=1
ML12A_HUMAN	288	19839	8 (7)	Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=1 SV=2
MYL9_HUMAN	131	19871	6 (5)	Myosin regulatory light polypeptide 9 OS=Homo sapiens GN=MYL9 PE=1 SV=4
VDAC1_HUMAN	286	30868	7 (5)	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2
VDAC3_HUMAN	145	30981	2 (2)	Voltage-dependent anion-selective channel protein 3 OS=Homo sapiens GN=VDAC3 PE=1 SV=1
HBB_HUMAN	282	16102	6 (6)	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2
HBG1_HUMAN	193	16187	2 (2)	Hemoglobin subunit gamma-1 OS=Homo sapiens GN=HBG1 PE=1 SV=2
ENOA_HUMAN	281	47481	11 (4)	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2
RS25_HUMAN	279	13791	5 (4)	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1
RL30_HUMAN	276	12947	3 (3)	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2
PEBP1_HUMAN	275	21158	6 (4)	Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3
CXCL7_HUMAN	273	14171	5 (5)	Platelet basic protein OS=Homo sapiens GN=PPBP PE=1 SV=3
ATP5L_HUMAN	272	11421	4 (4)	ATP synthase subunit g, mitochondrial OS=Homo sapiens GN=ATP5L PE=1 SV=3
IF5A1_HUMAN	269	17049	8 (4)	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2
RS19_HUMAN	264	16051	9 (7)	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2
RL27A_HUMAN	260	16665	3 (3)	60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=1 SV=2
PDIA6_HUMAN	259	48490	6 (5)	Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1
RL18_HUMAN	256	21735	5 (4)	60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2
RL35_HUMAN	252	14543	4 (4)	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2
RL22_HUMAN	248	14835	4 (3)	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2

PPIB_HUMAN	246	23785	8 (7)	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2
CUTA_HUMAN	242	19218	3 (2)	Protein CutA OS=Homo sapiens GN=CUTA PE=1 SV=2
RS14_HUMAN	242	16434	4 (4)	40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3
LYSC_HUMAN	240	16982	7 (3)	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1
ERP29_HUMAN	239	29032	8 (4)	Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=1 SV=4
TFIP8_HUMAN	233	23160	2 (1)	Tumor necrosis factor alpha-induced protein 8 OS=Homo sapiens GN=TNFAIP8 PE=1 SV=1
SRP14_HUMAN	233	14675	4 (4)	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2
SC61B_HUMAN	232	10025	3 (2)	Protein transport protein Sec61 subunit beta OS=Homo sapiens GN=SEC61B PE=1 SV=2
MDHM_HUMAN	229	35937	11 (9)	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3
RS13_HUMAN	225	17212	9 (7)	40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2
NDUA4_HUMAN	221	9421	5 (4)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=Homo sapiens GN=NDUFA4 PE=1 SV=1
CNPY2_HUMAN	220	20981	4 (3)	Protein canopy homolog 2 OS=Homo sapiens GN=CNPY2 PE=1 SV=1
FRIL_HUMAN	217	20064	1 (1)	Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2
RL38_HUMAN	216	8270	4 (3)	60S ribosomal protein L38 OS=Homo sapiens GN=RPL38 PE=1 SV=2
HSP7C_HUMAN	210	71082	8 (4)	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1
CAH2_HUMAN	209	29285	7 (5)	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2
RAB5C_HUMAN	206	23696	7 (5)	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2
CY1_HUMAN	202	35709	3 (2)	Cytochrome c1, heme protein, mitochondrial OS=Homo sapiens GN=CYC1 PE=1 SV=2
ATPK_HUMAN	202	11025	2 (2)	ATP synthase subunit f, mitochondrial OS=Homo sapiens GN=ATP5J2 PE=1 SV=3
GANAB_HUMAN	202	107263	5 (4)	Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3
UBIQ_HUMAN	198	8560	4 (3)	Ubiquitin OS=Homo sapiens GN=RPS27A PE=1 SV=1
RNAS2_HUMAN	196	18855	3 (3)	Non-secretory ribonuclease OS=Homo sapiens GN=RNASE2 PE=1 SV=2
PRDX5_HUMAN	192	22298	8 (6)	Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=3
SODM_HUMAN	190	24878	5 (3)	Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2
RL29_HUMAN	190	17798	2 (1)	60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2
LAP2B_HUMAN	188	50696	6 (4)	Lamina-associated polypeptide 2, isoforms beta/gamma OS=Homo sapiens GN=TMPO PE=1 SV=2
SSBP_HUMAN	187	17249	6 (3)	Single-stranded DNA-binding protein, mitochondrial OS=Homo sapiens GN=SSBP1 PE=1 SV=1
BID_HUMAN	185	22152	4 (2)	BH3-interacting domain death agonist OS=Homo sapiens GN=BID PE=1 SV=1
ROA2_HUMAN	185	37464	6 (4)	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2
ALDOA_HUMAN	185	39851	6 (3)	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2
SLIRP_HUMAN	184	12398	7 (6)	SRA stem-loop-interacting RNA-binding protein, mitochondrial OS=Homo sapiens GN=SLIRP PE=1 SV=1
COX2_HUMAN	181	25719	4 (4)	Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1
RS17_HUMAN	180	15597	6 (4)	40S ribosomal protein S17 OS=Homo sapiens GN=RPS17 PE=1 SV=2
HNRPK_HUMAN	180	51230	7 (5)	Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1
CLIC1_HUMAN	174	27248	7 (4)	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4
PPT1_HUMAN	173	34627	3 (2)	Palmitoyl-protein thioesterase 1 OS=Homo sapiens GN=PPT1 PE=1 SV=1
RS28_HUMAN	171	7893	2 (2)	40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1
RS20_HUMAN	171	13478	4 (3)	40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1
LGUL_HUMAN	171	20992	10 (6)	Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 PE=1 SV=4
NDUBA_HUMAN	170	21048	4 (3)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3
PDCD6_HUMAN	169	21912	5 (5)	Programmed cell death protein 6 OS=Homo sapiens GN=PDCD6 PE=1 SV=1
TLN1_HUMAN	166	271766	7 (2)	Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3
UB2V1_HUMAN	165	16598	4 (3)	Ubiquitin-conjugating enzyme E2 variant 1 OS=Homo sapiens GN=UBE2V1 PE=1 SV=2

RS15_HUMAN	165	17029	4 (2)	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2
RL14_HUMAN	162	23531	2 (2)	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4
ARPC5_HUMAN	161	16367	2 (2)	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3
NDUB4_HUMAN	160	15256	5 (4)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 OS=Homo sapiens GN=NDUFB4 PE=1 SV=3
ECH1_HUMAN	157	36136	5 (3)	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH1 PE=1 SV=2
RL36_HUMAN	157	12303	4 (4)	60S ribosomal protein L36 OS=Homo sapiens GN=RPL36 PE=1 SV=3
CYB5B_HUMAN	156	16436	2 (2)	Cytochrome b5 type B OS=Homo sapiens GN=CYB5B PE=1 SV=2
TPIS_HUMAN	155	26938	5 (4)	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=2
TMEDA_HUMAN	155	25131	5 (4)	Transmembrane emp24 domain-containing protein 10 OS=Homo sapiens GN=TMED10 PE=1 SV=2
ATPD_HUMAN	155	17479	2 (1)	ATP synthase subunit delta, mitochondrial OS=Homo sapiens GN=ATP5D PE=1 SV=2
RL19_HUMAN	155	23565	3 (2)	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1
MIF_HUMAN	154	12639	2 (2)	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4
SFRS3_HUMAN	154	19546	5 (3)	Splicing factor, arginine/serine-rich 3 OS=Homo sapiens GN=SFRS3 PE=1 SV=1
PLF4_HUMAN	154	11123	4 (4)	Platelet factor 4 OS=Homo sapiens GN=PF4 PE=1 SV=2
CX7A2_HUMAN	152	9390	3 (3)	Cytochrome c oxidase subunit 7A2, mitochondrial OS=Homo sapiens GN=COX7A2 PE=1 SV=1
CALR_HUMAN	152	48283	7 (4)	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1
RLA1_HUMAN	150	11621	2 (2)	60S acidic ribosomal protein P1 OS=Homo sapiens GN=RPLP1 PE=1 SV=1
C1QBP_HUMAN	149	31742	3 (3)	Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QBP PE=1 SV=1
RS16_HUMAN	148	16549	5 (3)	40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2
RL23_HUMAN	147	14970	4 (2)	60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1
S10AB_HUMAN	145	11847	6 (4)	Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2
1B15_HUMAN	145	40648	3 (3)	HLA class I histocompatibility antigen, B-15 alpha chain OS=Homo sapiens GN=HLA-B PE=1 SV=2
PDIA3_HUMAN	144	57146	6 (4)	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4
S10A8_HUMAN	144	10885	2 (2)	Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1
UE2NL_HUMAN	143	17366	3 (2)	Putative ubiquitin-conjugating enzyme E2 N-like OS=Homo sapiens GN=UBE2NL PE=5 SV=1
CHCH2_HUMAN	143	15731	2 (1)	Coiled-coil-helix-coiled-coil-helix domain-containing protein 2, mitochondrial OS=Homo sapiens GN=CHCHD2 PE=1 SV=
RS24_HUMAN	141	15413	3 (2)	40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=1 SV=1
PRDX3_HUMAN	139	28017	6 (4)	Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3 PE=1 SV=3
NDUA7_HUMAN	137	12601	4 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=Homo sapiens GN=NDUFA7 PE=1 SV=3
RL10_HUMAN	136	25044	6 (2)	60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=1 SV=4
ATP5I_HUMAN	135	7928	3 (3)	ATP synthase subunit e, mitochondrial OS=Homo sapiens GN=ATP5I PE=1 SV=2
CISD2_HUMAN	135	15497	4 (2)	CDGSH iron sulfur domain-containing protein 2 OS=Homo sapiens GN=CISD2 PE=1 SV=1
DOPD_HUMAN	134	12818	2 (2)	D-dopachrome decarboxylase OS=Homo sapiens GN=DDT PE=1 SV=3
RS10_HUMAN	134	18886	3 (3)	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1
USMG5_HUMAN	133	6510	2 (2)	Up-regulated during skeletal muscle growth protein 5 OS=Homo sapiens GN=USMG5 PE=1 SV=1
HINT2_HUMAN	133	17208	5 (3)	Histidine triad nucleotide-binding protein 2, mitochondrial OS=Homo sapiens GN=HINT2 PE=1 SV=1
CX6B1_HUMAN	132	10414	5 (3)	Cytochrome c oxidase subunit 6B1 OS=Homo sapiens GN=COX6B1 PE=1 SV=2
PRDX1_HUMAN	132	22324	10 (4)	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1
PRDX2_HUMAN	64	22049	2 (2)	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5
ARPC4_HUMAN	128	19768	4 (4)	Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3
LYPA1_HUMAN	127	24996	2 (1)	Acyl-protein thioesterase 1 OS=Homo sapiens GN=LYPLA1 PE=1 SV=1
S10A4_HUMAN	127	11949	3 (2)	Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1
CK059_HUMAN	126	17848	2 (2)	RhoA activator C11orf59 OS=Homo sapiens GN=C11orf59 PE=1 SV=2

RL24_HUMAN	125	17882	3 (3)	60S ribosomal protein L24 OS=Homo sapiens GN=RPL24 PE=1 SV=1
RS12_HUMAN	125	14905	6 (5)	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3
KAD3_HUMAN	124	25550	4 (2)	GTP:AMP phosphotransferase mitochondrial OS=Homo sapiens GN=AK3 PE=1 SV=4
PRDX6_HUMAN	124	25133	6 (4)	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3
RAB7A_HUMAN	123	23760	7 (2)	Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1
MA2B1_HUMAN	122	114357	5 (3)	Lysosomal alpha-mannosidase OS=Homo sapiens GN=MAN2B1 PE=1 SV=3
H15_HUMAN	120	22566	4 (3)	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3
PGAM1_HUMAN	120	28900	3 (2)	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2
LYRIC_HUMAN	119	63856	2 (2)	Protein LYRIC OS=Homo sapiens GN=MTDH PE=1 SV=2
VATG1_HUMAN	119	13863	3 (3)	V-type proton ATPase subunit G 1 OS=Homo sapiens GN=ATP6V1G1 PE=1 SV=3
RL11_HUMAN	118	20468	1 (1)	60S ribosomal protein L11 OS=Homo sapiens GN=RPL11 PE=1 SV=2
QCR9_HUMAN	118	7304	2 (2)	Cytochrome b-c1 complex subunit 9 OS=Homo sapiens GN=UQCR10 PE=1 SV=3
SODC_HUMAN	118	16154	3 (1)	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2
NDUF4_HUMAN	117	20311	2 (1)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 4 OS=Homo sapiens GN=NDUFAF4 PE=1 SV=1
RAN_HUMAN	117	24579	3 (3)	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3
NDUAD_HUMAN	117	16688	4 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Homo sapiens GN=NDUFA13 PE=1 SV=3
TOM20_HUMAN	117	16459	2 (2)	Mitochondrial import receptor subunit TOM20 homolog OS=Homo sapiens GN=TOMM20 PE=1 SV=1
COX5B_HUMAN	116	13915	4 (4)	Cytochrome c oxidase subunit 5B, mitochondrial OS=Homo sapiens GN=COX5B PE=1 SV=2
VAMP8_HUMAN	116	11488	4 (3)	Vesicle-associated membrane protein 8 OS=Homo sapiens GN=VAMP8 PE=1 SV=1
GP1BB_HUMAN	116	22274	4 (3)	Platelet glycoprotein Ib beta chain OS=Homo sapiens GN=GP1BB PE=1 SV=1
CREG1_HUMAN	115	24173	1 (1)	Protein CREG1 OS=Homo sapiens GN=CREG1 PE=1 SV=1
HNRPG_HUMAN	114	42306	5 (3)	Heterogeneous nuclear ribonucleoprotein G OS=Homo sapiens GN=RBMX PE=1 SV=3
KPYM_HUMAN	114	58470	3 (2)	Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM2 PE=1 SV=4
NLTP_HUMAN	114	59640	7 (5)	Non-specific lipid-transfer protein OS=Homo sapiens GN=SCP2 PE=1 SV=2
SCPDH_HUMAN	114	47464	2 (1)	Probable saccharopine dehydrogenase OS=Homo sapiens GN=SCCPDH PE=1 SV=1
ARL8B_HUMAN	113	21753	4 (2)	ADP-ribosylation factor-like protein 8B OS=Homo sapiens GN=ARL8B PE=1 SV=1
HNRPC_HUMAN	112	33707	4 (3)	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=1 SV=4
TBCA_HUMAN	112	12904	5 (4)	Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=1 SV=3
RAB1B_HUMAN	111	22328	4 (3)	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1
RAB10_HUMAN	97	22755	5 (3)	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1
THIO_HUMAN	111	12015	3 (1)	Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3
HEXB_HUMAN	111	63527	7 (5)	Beta-hexosaminidase subunit beta OS=Homo sapiens GN=HEXB PE=1 SV=3
PSME2_HUMAN	110	27515	4 (2)	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=3
PGRC2_HUMAN	110	23861	2 (2)	Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRMC2 PE=1 SV=1
COX41_HUMAN	110	19621	4 (3)	Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Homo sapiens GN=COX4I1 PE=1 SV=1
6PGL_HUMAN	109	27815	4 (3)	6-phosphogluconolactonase OS=Homo sapiens GN=PGLS PE=1 SV=2
PSB1_HUMAN	109	26700	2 (1)	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2
NDUB9_HUMAN	108	22045	2 (2)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 9 OS=Homo sapiens GN=NDUFB9 PE=1 SV=3
EF2_HUMAN	107	96246	5 (3)	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4
PA1B2_HUMAN	105	25724	3 (2)	Platelet-activating factor acetylhydrolase IB subunit beta OS=Homo sapiens GN=PAFAH1B2 PE=1 SV=1
NDUV2_HUMAN	105	27659	3 (2)	NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Homo sapiens GN=NDUFV2 PE=1 SV=2
PHB_HUMAN	102	29843	3 (1)	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1
RS15A_HUMAN	101	14944	7 (3)	40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2

RCL_HUMAN	101	19211	2 (2)	Deoxyribonucleoside 5'-monophosphate N-glycosidase OS=Homo sapiens GN=RCL PE=1 SV=1
ANXA2_HUMAN	99	38808	6 (2)	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2
TFAM_HUMAN	99	29306	5 (2)	Transcription factor A, mitochondrial OS=Homo sapiens GN=TFAM PE=1 SV=1
ARF1_HUMAN	99	20741	6 (3)	ADP-ribosylation factor 1 OS=Homo sapiens GN=ARF1 PE=1 SV=2
ADT2_HUMAN	98	33102	9 (6)	ADP/ATP translocase 2 OS=Homo sapiens GN=SLC25A5 PE=1 SV=6
NDUA5_HUMAN	98	13507	3 (2)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 OS=Homo sapiens GN=NDUFA5 PE=1 SV=3
PRTN3_HUMAN	98	28245	3 (2)	Myeloblastin OS=Homo sapiens GN=PRTN3 PE=1 SV=3
RS23_HUMAN	97	15969	2 (2)	40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=1 SV=3
MTPN_HUMAN	97	13058	2 (1)	Myotrophin OS=Homo sapiens GN=MTPN PE=1 SV=2
TPD54_HUMAN	97	22281	1 (1)	Tumor protein D54 OS=Homo sapiens GN=TPD52L2 PE=1 SV=2
RM12_HUMAN	96	21563	4 (3)	39S ribosomal protein L12, mitochondrial OS=Homo sapiens GN=MRPL12 PE=1 SV=2
LDHA_HUMAN	96	36950	5 (2)	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2
GPX1_HUMAN	96	22218	6 (5)	Glutathione peroxidase 1 OS=Homo sapiens GN=GPX1 PE=1 SV=3
LYAG_HUMAN	95	106107	3 (2)	Lysosomal alpha-glucosidase OS=Homo sapiens GN=GAA PE=1 SV=3
ACTN1_HUMAN	94	103563	8 (2)	Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2
QCR7_HUMAN	93	13522	5 (4)	Cytochrome b-c1 complex subunit 7 OS=Homo sapiens GN=UQCRB PE=1 SV=2
DUT_HUMAN	92	26975	4 (2)	Deoxyuridine 5'-triphosphate nucleotidohydrolase, mitochondrial OS=Homo sapiens GN=DUT PE=1 SV=3
RL31_HUMAN	92	14454	4 (4)	60S ribosomal protein L31 OS=Homo sapiens GN=RPL31 PE=1 SV=1
COTL1_HUMAN	91	16049	3 (2)	Coactosin-like protein OS=Homo sapiens GN=COTL1 PE=1 SV=3
TPP1_HUMAN	91	61723	2 (1)	Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1 PE=1 SV=2
TM109_HUMAN	90	26194	1 (1)	Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1
DHSB_HUMAN	90	32407	5 (4)	Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Homo sapiens GN=SDHB PE=1 SV=3
PLSL_HUMAN	90	70815	3 (2)	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=5
LONM_HUMAN	89	106936	4 (1)	Lon protease homolog, mitochondrial OS=Homo sapiens GN=LONP1 PE=1 SV=2
SMD3_HUMAN	88	14021	3 (2)	Small nuclear ribonucleoprotein Sm D3 OS=Homo sapiens GN=SNRPD3 PE=1 SV=1
BAX_HUMAN	88	21285	1 (1)	Apoptosis regulator BAX OS=Homo sapiens GN=BAX PE=1 SV=1
TCPZ_HUMAN	88	58444	3 (1)	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3
RL23A_HUMAN	85	17684	4 (1)	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1
NDUS8_HUMAN	85	24203	2 (2)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Homo sapiens GN=NDUFS8 PE=1 SV=1
NUCL_HUMAN	84	76625	5 (1)	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3
VDAC2_HUMAN	84	32060	2 (1)	Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1 SV=2
RLA0_HUMAN	82	34423	4 (1)	60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1
GSTP1_HUMAN	82	23569	2 (2)	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2
RAB2A_HUMAN	81	23702	1 (1)	Ras-related protein Rab-2A OS=Homo sapiens GN=RAB2A PE=1 SV=1
TMCO1_HUMAN	81	21389	3 (1)	Transmembrane and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=TMCO1 PE=1 SV=1
RA1L3_HUMAN	80	34373	5 (1)	Putative heterogeneous nuclear ribonucleoprotein A1-like 3 OS=Homo sapiens GN=HNRPA1L3 PE=5 SV=1
RL17_HUMAN	79	21611	6 (4)	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3
COMT_HUMAN	78	30474	5 (2)	Catechol O-methyltransferase OS=Homo sapiens GN=COMT PE=1 SV=2
QCR8_HUMAN	78	9900	2 (1)	Cytochrome b-c1 complex subunit 8 OS=Homo sapiens GN=UQCRQ PE=1 SV=4
MGST3_HUMAN	77	16734	1 (1)	Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1
OCAD1_HUMAN	77	27780	4 (1)	OCIA domain-containing protein 1 OS=Homo sapiens GN=OCIAD1 PE=1 SV=1
FKBP2_HUMAN	76	15810	1 (1)	Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Homo sapiens GN=FKBP2 PE=1 SV=2
PSB6_HUMAN	76	25570	1 (1)	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4

ARP5L_HUMAN	76	16931	2 (2)	Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5L PE=1 SV=1
NDUA8_HUMAN	76	20548	4 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=Homo sapiens GN=NDUFA8 PE=1 SV=3
RL27_HUMAN	75	15788	3 (2)	60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2
PARK7_HUMAN	75	20050	6 (4)	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2
RM49_HUMAN	74	19243	2 (1)	39S ribosomal protein L49, mitochondrial OS=Homo sapiens GN=MRPL49 PE=1 SV=1
TOM22_HUMAN	74	15512	2 (1)	Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=1 SV=3
ITA2B_HUMAN	73	114446	1 (1)	Integrin alpha-IIb OS=Homo sapiens GN=ITGA2B PE=1 SV=3
RL7_HUMAN	72	29264	6 (1)	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1
GLRX5_HUMAN	72	16732	1 (1)	Glutaredoxin-related protein 5 OS=Homo sapiens GN=GLRX5 PE=1 SV=2
UB2L3_HUMAN	72	18021	2 (2)	Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1
SNX3_HUMAN	72	18808	2 (2)	Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3
ALDH2_HUMAN	71	56859	1 (1)	Aldehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH2 PE=1 SV=2
TEBP_HUMAN	70	18971	2 (1)	Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1
DHSA_HUMAN	70	73672	3 (1)	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=2
TKT_HUMAN	70	68519	3 (2)	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3
GRPE1_HUMAN	69	24492	5 (4)	GrpE protein homolog 1, mitochondrial OS=Homo sapiens GN=GRPEL1 PE=1 SV=2
SMD1_HUMAN	68	13273	3 (1)	Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1
SH3L3_HUMAN	68	10488	1 (1)	SH3 domain-binding glutamic acid-rich-like protein 3 OS=Homo sapiens GN=SH3BGRL3 PE=1 SV=1
RL13_HUMAN	68	24304	6 (2)	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4
BAK_HUMAN	68	23508	2 (2)	Bcl-2 homologous antagonist/killer OS=Homo sapiens GN=BAK1 PE=1 SV=1
CATA_HUMAN	67	59947	4 (1)	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3
SSRD_HUMAN	66	19158	1 (1)	Translocon-associated protein subunit delta OS=Homo sapiens GN=SSR4 PE=1 SV=1
LYRM7_HUMAN	65	12004	1 (1)	LYR motif-containing protein 7 OS=Homo sapiens GN=LYRM7 PE=1 SV=1
BTF3_HUMAN	65	22211	2 (1)	Transcription factor BTF3 OS=Homo sapiens GN=BTF3 PE=1 SV=1
RS7_HUMAN	65	22113	2 (2)	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1
SSRG_HUMAN	64	21067	2 (1)	Translocon-associated protein subunit gamma OS=Homo sapiens GN=SSR3 PE=1 SV=1
ATPG_HUMAN	64	33032	3 (2)	ATP synthase subunit gamma, mitochondrial OS=Homo sapiens GN=ATP5C1 PE=1 SV=1
VMA21_HUMAN	64	11347	1 (1)	Vacuolar ATPase assembly integral membrane protein VMA21 OS=Homo sapiens GN=VMA21 PE=1 SV=1
NDUS3_HUMAN	63	30337	5 (2)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Homo sapiens GN=NDUFS3 PE=1 SV=1
RUXE_HUMAN	63	10854	1 (1)	Small nuclear ribonucleoprotein E OS=Homo sapiens GN=SNRPE PE=1 SV=1
DAD1_HUMAN	62	12660	2 (1)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 OS=Homo sapiens GN=DAD1 PE=1 SV=3
TMED2_HUMAN	62	22860	1 (1)	Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1
ARPC3_HUMAN	62	20761	4 (2)	Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3
AN32A_HUMAN	62	28682	5 (3)	Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Homo sapiens GN=ANP32A PE=1 SV=1
RL7A_HUMAN	61	30148	4 (1)	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2
CISY_HUMAN	61	51908	2 (2)	Citrate synthase, mitochondrial OS=Homo sapiens GN=CS PE=1 SV=2
HNRPU_HUMAN	60	91269	5 (1)	Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6
NWD1_HUMAN	60	176808	5 (1)	NACHT and WD repeat domain-containing protein 1 OS=Homo sapiens GN=NWD1 PE=2 SV=2
EZRI_HUMAN	59	69484	7 (1)	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4
LDHB_HUMAN	59	36900	4 (2)	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2
STMN1_HUMAN	59	17292	3 (1)	Stathmin OS=Homo sapiens GN=STMN1 PE=1 SV=3
APT_HUMAN	59	19766	3 (2)	Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2
PGK1_HUMAN	58	44985	2 (2)	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3

SC22B_HUMAN	58	24896	3 (1)	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=3
FKB1A_HUMAN	57	12000	1 (1)	Peptidyl-prolyl cis-trans isomerase FKBP1A OS=Homo sapiens GN=FKBP1A PE=1 SV=2
PLP2_HUMAN	57	17022	1 (1)	Proteolipid protein 2 OS=Homo sapiens GN=PLP2 PE=1 SV=1
PGRC1_HUMAN	56	21772	1 (1)	Membrane-associated progesterone receptor component 1 OS=Homo sapiens GN=PGRMC1 PE=1 SV=3
GRN_HUMAN	55	68518	2 (1)	Granulins OS=Homo sapiens GN=GRN PE=1 SV=2
ASPG_HUMAN	55	37812	2 (1)	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase OS=Homo sapiens GN=AGA PE=1 SV=2
TXD12_HUMAN	55	19365	4 (2)	Thioredoxin domain-containing protein 12 OS=Homo sapiens GN=TXNDC12 PE=1 SV=1
ITB2_HUMAN	55	87976	2 (1)	Integrin beta-2 OS=Homo sapiens GN=ITGB2 PE=1 SV=2
RAC2_HUMAN	55	21814	5 (3)	Ras-related C3 botulinum toxin substrate 2 OS=Homo sapiens GN=RAC2 PE=1 SV=1
RHOG_HUMAN	29	21751	3 (2)	Rho-related GTP-binding protein RhoG OS=Homo sapiens GN=RHOG PE=1 SV=1
RS21_HUMAN	55	9220	4 (1)	40S ribosomal protein S21 OS=Homo sapiens GN=RPS21 PE=1 SV=1
HIG2A_HUMAN	55	11578	1 (1)	HIG1 domain family member 2A OS=Homo sapiens GN=HIGD2A PE=1 SV=1
RL40_HUMAN	55	6462	2 (1)	60S ribosomal protein L40 OS=Homo sapiens GN=UBA52 PE=1 SV=1
RM23_HUMAN	55	17770	3 (1)	39S ribosomal protein L23, mitochondrial OS=Homo sapiens GN=MRPL23 PE=1 SV=1
PM14_HUMAN	54	14690	1 (1)	Pre-mRNA branch site protein p14 OS=Homo sapiens GN=SF3B14 PE=1 SV=1
PFD2_HUMAN	53	16695	2 (2)	Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1
NONO_HUMAN	53	54311	2 (1)	Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV=4
RL6_HUMAN	53	32765	2 (1)	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3
B2MG_HUMAN	53	13820	3 (3)	Beta-2-microglobulin OS=Homo sapiens GN=B2M PE=1 SV=1
S10A6_HUMAN	52	10230	2 (2)	Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1
H2AY_HUMAN	52	39764	3 (1)	Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY PE=1 SV=4
SSRA_HUMAN	52	32215	3 (1)	Translocon-associated protein subunit alpha OS=Homo sapiens GN=SSR1 PE=1 SV=3
GBLP_HUMAN	52	35511	1 (1)	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3
BR44L_HUMAN	52	12510	1 (1)	Brain protein 44-like protein OS=Homo sapiens GN=BRP44L PE=1 SV=1
ACPM_HUMAN	51	17577	3 (2)	Acyl carrier protein, mitochondrial OS=Homo sapiens GN=NDUFAB1 PE=1 SV=3
THOC4_HUMAN	51	26872	2 (1)	THO complex subunit 4 OS=Homo sapiens GN=THOC4 PE=1 SV=3
CHSP1_HUMAN	51	16110	1 (1)	Calcium-regulated heat stable protein 1 OS=Homo sapiens GN=CARHSP1 PE=1 SV=2
GBB1_HUMAN	51	38151	2 (2)	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3
RS30_HUMAN	50	6644	2 (1)	40S ribosomal protein S30 OS=Homo sapiens GN=FAU PE=1 SV=1
NDUAC_HUMAN	50	17104	2 (2)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Homo sapiens GN=NDUFA12 PE=1 SV=1
GAPR1_HUMAN	50	17322	1 (1)	Golgi-associated plant pathogenesis-related protein 1 OS=Homo sapiens GN=GLIPR2 PE=1 SV=3
CF115_HUMAN	50	9108	2 (1)	UPF0727 protein C6orf115 OS=Homo sapiens GN=C6orf115 PE=1 SV=1
RM41_HUMAN	50	15430	1 (1)	39S ribosomal protein L41, mitochondrial OS=Homo sapiens GN=MRPL41 PE=1 SV=1
CNN2_HUMAN	50	34074	3 (2)	Calponin-2 OS=Homo sapiens GN=CNN2 PE=1 SV=4
SFRS2_HUMAN	50	25461	2 (1)	Splicing factor, arginine/serine-rich 2 OS=Homo sapiens GN=SFRS2 PE=1 SV=4
NH2L1_HUMAN	50	14393	2 (1)	NHP2-like protein 1 OS=Homo sapiens GN=NHP2L1 PE=1 SV=3
AT1B3_HUMAN	49	31834	2 (1)	Sodium/potassium-transporting ATPase subunit beta-3 OS=Homo sapiens GN=ATP1B3 PE=1 SV=1
RL15_HUMAN	49	24245	2 (1)	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2
RM43_HUMAN	49	23701	4 (2)	39S ribosomal protein L43, mitochondrial OS=Homo sapiens GN=MRPL43 PE=1 SV=1
PPIF_HUMAN	48	22368	2 (1)	Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Homo sapiens GN=PPIF PE=1 SV=1
SNG2_HUMAN	47	25079	2 (1)	Synaptogyrin-2 OS=Homo sapiens GN=SYNGR2 PE=1 SV=1
RL36A_HUMAN	47	12718	3 (2)	60S ribosomal protein L36a OS=Homo sapiens GN=RPL36A PE=1 SV=2
FCERG_HUMAN	47	9775	2 (1)	High affinity immunoglobulin epsilon receptor subunit gamma OS=Homo sapiens GN=FCER1G PE=1 SV=1

HNRH1_HUMAN	46	49484	3 (2)	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4
PIMT_HUMAN	46	24806	2 (2)	Protein-L-isoaspartate(D-aspartate) O-methyltransferase OS=Homo sapiens GN=PCMT1 PE=1 SV=3
RM14_HUMAN	46	16165	2 (1)	39S ribosomal protein L14, mitochondrial OS=Homo sapiens GN=MRPL14 PE=1 SV=1
ANXA5_HUMAN	46	35971	4 (3)	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2
ANXA4_HUMAN	45	36088	1 (1)	Annexin A4 OS=Homo sapiens GN=ANXA4 PE=1 SV=4
AT5F1_HUMAN	45	28947	4 (1)	ATP synthase subunit b, mitochondrial OS=Homo sapiens GN=ATP5F1 PE=1 SV=2
RBM3_HUMAN	45	17160	1 (1)	Putative RNA-binding protein 3 OS=Homo sapiens GN=RBM3 PE=1 SV=1
PSA7_HUMAN	45	28041	2 (1)	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1
SERA_HUMAN	45	57356	5 (2)	D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4
NDUBB_HUMAN	45	17363	1 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 11, mitochondrial OS=Homo sapiens GN=NDUFB11 PE=
AIBP_HUMAN	44	31996	3 (2)	Apolipoprotein A-I-binding protein OS=Homo sapiens GN=APOA1BP PE=1 SV=2
CATG_HUMAN	44	29161	2 (1)	Cathepsin G OS=Homo sapiens GN=CTSG PE=1 SV=2
RS29_HUMAN	44	6900	2 (1)	40S ribosomal protein S29 OS=Homo sapiens GN=RPS29 PE=1 SV=2
SNP23_HUMAN	43	23682	2 (1)	Synaptosomal-associated protein 23 OS=Homo sapiens GN=SNAP23 PE=1 SV=1
TCPG_HUMAN	43	61066	3 (1)	T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4
BAG2_HUMAN	43	23928	3 (1)	BAG family molecular chaperone regulator 2 OS=Homo sapiens GN=BAG2 PE=1 SV=1
PPIH_HUMAN	43	19481	3 (1)	Peptidyl-prolyl cis-trans isomerase H OS=Homo sapiens GN=PPIH PE=1 SV=1
FABP5_HUMAN	43	15497	2 (1)	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3
RS11_HUMAN	42	18590	2 (1)	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3
RS27_HUMAN	42	9797	3 (1)	40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3
GDIR1_HUMAN	42	23250	2 (1)	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDIA PE=1 SV=3
RT36_HUMAN	41	11459	2 (2)	28S ribosomal protein S36, mitochondrial OS=Homo sapiens GN=MRPS36 PE=1 SV=2
LSM4_HUMAN	41	15511	3 (1)	U6 snRNA-associated Sm-like protein LSm4 OS=Homo sapiens GN=LSM4 PE=1 SV=1
ECHM_HUMAN	41	31823	3 (1)	Enoyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=ECHS1 PE=1 SV=4
RL9_HUMAN	41	21964	3 (1)	60S ribosomal protein L9 OS=Homo sapiens GN=RPL9 PE=1 SV=1
CI142_HUMAN	41	21968	1 (1)	Uncharacterized protein C9orf142 OS=Homo sapiens GN=C9orf142 PE=1 SV=2
DECR_HUMAN	40	36330	2 (1)	2,4-dienoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=DECR1 PE=1 SV=1
NPC2_HUMAN	40	16902	3 (1)	Epididymal secretory protein E1 OS=Homo sapiens GN=NPC2 PE=1 SV=1
RL39_HUMAN	40	6403	2 (1)	60S ribosomal protein L39 OS=Homo sapiens GN=RPL39 PE=2 SV=2
HNRPM_HUMAN	40	77749	5 (1)	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3
RB11A_HUMAN	40	24492	3 (1)	Ras-related protein Rab-11A OS=Homo sapiens GN=RAB11A PE=1 SV=3
MCA3_HUMAN	40	19855	2 (1)	Eukaryotic translation elongation factor 1 epsilon-1 OS=Homo sapiens GN=EEF1E1 PE=1 SV=1
RL26_HUMAN	40	17248	3 (2)	60S ribosomal protein L26 OS=Homo sapiens GN=RPL26 PE=1 SV=1
FA36A_HUMAN	40	13511	1 (1)	Protein FAM36A OS=Homo sapiens GN=FAM36A PE=1 SV=2
SORCN_HUMAN	39	21947	4 (1)	Sorcin OS=Homo sapiens GN=SRI PE=1 SV=1
TCTP_HUMAN	39	19697	4 (2)	Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1
IMB1_HUMAN	39	98420	4 (1)	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2
MPKS1_HUMAN	39	13671	1 (1)	Mitogen-activated protein kinase scaffold protein 1 OS=Homo sapiens GN=MAPKSP1 PE=1 SV=1
GRB2_HUMAN	39	25304	1 (1)	Growth factor receptor-bound protein 2 OS=Homo sapiens GN=GRB2 PE=1 SV=1
IDHP_HUMAN	39	51333	2 (1)	Isocitrate dehydrogenase [NADP], mitochondrial OS=Homo sapiens GN=IDH2 PE=1 SV=2
SET_HUMAN	39	33469	1 (1)	Protein SET OS=Homo sapiens GN=SET PE=1 SV=3
AT5EL_HUMAN	38	5860	1 (1)	ATP synthase subunit epsilon-like protein, mitochondrial OS=Homo sapiens GN=ATP5EP2 PE=1 SV=1
RS26_HUMAN	38	13292	4 (1)	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3

S10A9_HUMAN	38	13291	3 (2)	Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1
RT23_HUMAN	37	21814	2 (1)	28S ribosomal protein S23, mitochondrial OS=Homo sapiens GN=MRPS23 PE=1 SV=2
MANF_HUMAN	37	20700	4 (3)	Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=2
FUMH_HUMAN	36	54773	1 (1)	Fumarate hydratase, mitochondrial OS=Homo sapiens GN=FH PE=1 SV=3
CPSF6_HUMAN	36	59344	2 (1)	Cleavage and polyadenylation specificity factor subunit 6 OS=Homo sapiens GN=CPSF6 PE=1 SV=2
RSU1_HUMAN	36	31521	1 (1)	Ras suppressor protein 1 OS=Homo sapiens GN=RSU1 PE=1 SV=3
TMED9_HUMAN	36	27374	2 (1)	Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=2
LRC59_HUMAN	36	35308	6 (2)	Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRRC59 PE=1 SV=1
IR3IP_HUMAN	36	9020	1 (1)	Immediate early response 3-interacting protein 1 OS=Homo sapiens GN=IER3IP1 PE=1 SV=1
ROAA_HUMAN	35	36316	3 (1)	Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=1 SV=2
RS6_HUMAN	34	28834	3 (1)	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1
NDUB1_HUMAN	34	7014	1 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 1 OS=Homo sapiens GN=NDUFB1 PE=1 SV=1
VPS25_HUMAN	34	20849	2 (1)	Vacuolar protein-sorting-associated protein 25 OS=Homo sapiens GN=VPS25 PE=1 SV=1
CA151_HUMAN	34	8916	1 (1)	UPF0327 protein C1orf151 OS=Homo sapiens GN=C1orf151 PE=1 SV=1
EFTU_HUMAN	34	49852	3 (1)	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2
ICT1_HUMAN	34	23843	2 (1)	Immature colon carcinoma transcript 1 protein OS=Homo sapiens GN=ICT1 PE=1 SV=1
PTBP1_HUMAN	34	57357	2 (1)	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1
IL25_HUMAN	34	20887	1 (1)	Interleukin-25 OS=Homo sapiens GN=IL25 PE=1 SV=1
ASC_HUMAN	34	21670	1 (1)	Apoptosis-associated speck-like protein containing a CARD OS=Homo sapiens GN=PYCARD PE=1 SV=2
SFRS9_HUMAN	33	25640	3 (1)	Splicing factor, arginine/serine-rich 9 OS=Homo sapiens GN=SFRS9 PE=1 SV=1
DLDH_HUMAN	33	54713	2 (1)	Dihydrolipoyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2
NDUF3_HUMAN	33	20566	2 (1)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 3 OS=Homo sapiens GN=NDUFAF3 PE=1 SV=1
IPYR2_HUMAN	33	38409	3 (1)	Inorganic pyrophosphatase 2, mitochondrial OS=Homo sapiens GN=PPA2 PE=1 SV=2
IF4H_HUMAN	33	27425	2 (1)	Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H PE=1 SV=5
RL32_HUMAN	33	15964	2 (1)	60S ribosomal protein L32 OS=Homo sapiens GN=RPL32 PE=1 SV=2
HNRPD_HUMAN	33	38581	4 (1)	Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD PE=1 SV=1
FBRL_HUMAN	33	33877	2 (1)	rRNA 2'-O-methyltransferase fibrillarin OS=Homo sapiens GN=FBL PE=1 SV=2
NDUB8_HUMAN	32	21865	2 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrial OS=Homo sapiens GN=NDUFB8 PE=1
ANXA1_HUMAN	32	38918	2 (1)	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2
ETFA_HUMAN	32	35400	2 (2)	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Homo sapiens GN=ETFA PE=1 SV=1
MPCP_HUMAN	32	40525	2 (1)	Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2
PSME1_HUMAN	32	28876	3 (1)	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1
IF4G1_HUMAN	32	176168	5 (1)	Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1 PE=1 SV=3
TCPB_HUMAN	31	57794	4 (2)	T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4
DYL1_HUMAN	31	10530	1 (1)	Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1
GDIR2_HUMAN	31	23031	4 (1)	Rho GDP-dissociation inhibitor 2 OS=Homo sapiens GN=ARHGDIB PE=1 SV=3
UB2D3_HUMAN	30	16904	2 (1)	Ubiquitin-conjugating enzyme E2 D3 OS=Homo sapiens GN=UBE2D3 PE=1 SV=1
RL18A_HUMAN	30	21034	4 (1)	60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A PE=1 SV=2
TMX1_HUMAN	30	32170	3 (1)	Thioredoxin-related transmembrane protein 1 OS=Homo sapiens GN=TMX1 PE=1 SV=1
RM50_HUMAN	30	18484	3 (1)	39S ribosomal protein L50, mitochondrial OS=Homo sapiens GN=MRPL50 PE=1 SV=2
LPPRC_HUMAN	30	159003	2 (1)	Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3
RL8_HUMAN	30	28235	1 (1)	60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=1 SV=2
QIL1_HUMAN	30	13136	1 (1)	Protein QIL1 OS=Homo sapiens GN=QIL1 PE=1 SV=1

HCDH_HUMAN	30	34313	2 (2)	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=1 SV=2
NUCB2_HUMAN	30	50278	1 (1)	Nucleobindin-2 OS=Homo sapiens GN=NUCB2 PE=1 SV=2
PCBP2_HUMAN	29	38955	2 (1)	Poly(rC)-binding protein 2 OS=Homo sapiens GN=PCBP2 PE=1 SV=1
NDUA2_HUMAN	28	11029	2 (1)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 2 OS=Homo sapiens GN=NDUFA2 PE=1 SV=3
CO4A4_HUMAN	28	165759	3 (1)	Collagen alpha-4(IV) chain OS=Homo sapiens GN=COL4A4 PE=1 SV=3
MMAB_HUMAN	28	27713	1 (1)	Cob(I)yrinic acid a,c-diamide adenosyltransferase, mitochondrial OS=Homo sapiens GN=MMAB PE=1 SV=1
RSSA_HUMAN	28	32947	5 (1)	40S ribosomal protein SA OS=Homo sapiens GN=RPSA PE=1 SV=4
RPN1_HUMAN	28	68641	3 (1)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1
NRAP_HUMAN	28	197920	5 (1)	Nebulin-related-anchoring protein OS=Homo sapiens GN=NRAP PE=2 SV=2
BRI3B_HUMAN	28	27932	1 (1)	BRI3-binding protein OS=Homo sapiens GN=BRI3BP PE=1 SV=1
RINI_HUMAN	28	51766	2 (1)	Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2
RHOA_HUMAN	28	22096	1 (1)	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1
THIL_HUMAN	28	45456	3 (1)	Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1
OSTF1_HUMAN	27	23943	1 (1)	Osteoclast-stimulating factor 1 OS=Homo sapiens GN=OSTF1 PE=1 SV=2
GSTK1_HUMAN	27	25594	2 (1)	Glutathione S-transferase kappa 1 OS=Homo sapiens GN=GSTK1 PE=1 SV=3
O2T35_HUMAN	27	36704	1 (1)	Olfactory receptor 2T35 OS=Homo sapiens GN=OR2T35 PE=2 SV=1
KLOTB_HUMAN	27	120473	2 (1)	Beta-klotho OS=Homo sapiens GN=KLB PE=1 SV=1
SDPR_HUMAN	27	47202	2 (1)	Serum deprivation-response protein OS=Homo sapiens GN=SDPR PE=1 SV=3
HMGA1_HUMAN	26	11669	2 (1)	High mobility group protein HMG-I/HMG-Y OS=Homo sapiens GN=HMGA1 PE=1 SV=3
UCRIL_HUMAN	26	31081	1 (1)	Cytochrome b-c1 complex subunit Rieske-like protein 1 OS=Homo sapiens GN=UQCRFSL1 PE=2 SV=1
GGT1_HUMAN	26	61714	2 (1)	Gamma-glutamyltranspeptidase 1 OS=Homo sapiens GN=GGT1 PE=1 SV=2
COX6C_HUMAN	26	8776	3 (2)	Cytochrome c oxidase subunit 6C OS=Homo sapiens GN=COX6C PE=1 SV=2
HCD2_HUMAN	26	27134	3 (1)	3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3
CSK2B_HUMAN	26	25268	1 (1)	Casein kinase II subunit beta OS=Homo sapiens GN=CSNK2B PE=1 SV=1
PSA5_HUMAN	26	26565	3 (1)	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3
QCR6_HUMAN	26	11017	2 (1)	Cytochrome b-c1 complex subunit 6, mitochondrial OS=Homo sapiens GN=UQCRH PE=1 SV=2
NDUV3_HUMAN	25	11990	1 (1)	NADH dehydrogenase [ubiquinone] flavoprotein 3, mitochondrial OS=Homo sapiens GN=NDUFV3 PE=1 SV=2
FIS1_HUMAN	25	16984	2 (1)	Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2
ARP3B_HUMAN	24	48090	1 (1)	Actin-related protein 3B OS=Homo sapiens GN=ACTR3B PE=1 SV=1
EXD2_HUMAN	24	57165	4 (1)	Exonuclease 3'-5' domain-containing protein 2 OS=Homo sapiens GN=EXD2 PE=1 SV=1
CS010_HUMAN	24	18897	1 (1)	UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1
PPAP_HUMAN	24	44880	2 (1)	Prostatic acid phosphatase OS=Homo sapiens GN=ACPP PE=1 SV=3
PFD5_HUMAN	24	17374	1 (1)	Prefoldin subunit 5 OS=Homo sapiens GN=PFDN5 PE=1 SV=2
PPAC_HUMAN	23	18487	3 (1)	Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1 PE=1 SV=3
MAPIP_HUMAN	23	13613	1 (1)	Mitogen-activated protein-binding protein-interacting protein OS=Homo sapiens GN=ROBLD3 PE=1 SV=1
PSA2_HUMAN	22	25996	2 (1)	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2
GLI3_HUMAN	22	171410	2 (1)	Zinc finger protein GLI3 OS=Homo sapiens GN=GLI3 PE=1 SV=6
MYH9_HUMAN	21	227646	7 (1)	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4
RISC_HUMAN	21	51083	1 (1)	Retinoid-inducible serine carboxypeptidase OS=Homo sapiens GN=SCPEP1 PE=1 SV=1
PDCD5_HUMAN	21	14276	1 (1)	Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3
TCPE_HUMAN	21	60089	2 (1)	T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=1 SV=1
RM13_HUMAN	21	20736	1 (1)	39S ribosomal protein L13, mitochondrial OS=Homo sapiens GN=MRPL13 PE=1 SV=1
OSTC_HUMAN	21	16932	1 (1)	Oligosaccharyltransferase complex subunit OSTC OS=Homo sapiens GN=OSTC PE=1 SV=1

RFA3_HUMAN	20	13674	2 (1)	Replication protein A 14 kDa subunit OS=Homo sapiens GN=RPA3 PE=1 SV=1
TIM14_HUMAN	20	12491	1 (1)	Mitochondrial import inner membrane translocase subunit TIM14 OS=Homo sapiens GN=DNAJC19 PE=1 SV=3
NDUS6_HUMAN	20	14045	1 (1)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Homo sapiens GN=NDUFS6 PE=1 SV=1
S14L4_HUMAN	20	47070	1 (1)	SEC14-like protein 4 OS=Homo sapiens GN=SEC14L4 PE=2 SV=1
PSB4_HUMAN	19	29242	1 (1)	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4
NR4A3_HUMAN	19	69327	2 (1)	Nuclear receptor subfamily 4 group A member 3 OS=Homo sapiens GN=NR4A3 PE=2 SV=3
CDN2C_HUMAN	19	18173	1 (1)	Cyclin-dependent kinase 4 inhibitor C OS=Homo sapiens GN=CDKN2C PE=1 SV=1
ECHA_HUMAN	19	83688	10 (1)	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2
PDLI7_HUMAN	19	50896	1 (1)	PDZ and LIM domain protein 7 OS=Homo sapiens GN=PDLIM7 PE=1 SV=1
VAPA_HUMAN	19	28103	5 (1)	Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA PE=1 SV=3
RL13A_HUMAN	18	23619	5 (1)	60S ribosomal protein L13a OS=Homo sapiens GN=RPL13A PE=1 SV=2
TXD17_HUMAN	18	14217	1 (1)	Thioredoxin domain-containing protein 17 OS=Homo sapiens GN=TXNDC17 PE=1 SV=1
PSA4_HUMAN	18	29750	1 (1)	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1
IGSF3_HUMAN	18	136252	2 (1)	Immunoglobulin superfamily member 3 OS=Homo sapiens GN=IGSF3 PE=1 SV=3
RS2_HUMAN	18	31590	1 (1)	40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2
EF1B_HUMAN	18	24919	1 (1)	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3
NDUS5_HUMAN	18	12737	1 (1)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 5 OS=Homo sapiens GN=NDUFS5 PE=1 SV=3
RT33_HUMAN	17	12621	1 (1)	28S ribosomal protein S33, mitochondrial OS=Homo sapiens GN=MRPS33 PE=1 SV=1

Suppl. Table 2C: Total number of proteins detected in THP-1 incubated with untreated platelets (score calculated from 2 biological replicates and 3 technical replicates)

Entry	MASCOT Score	Mass	Sequences coverage	Description
CH60_HUMAN	1434	61187	17 (14)	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2
K1C10_HUMAN	1373	59020	30 (22)	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6
K1C14_HUMAN	366	51872	9 (7)	Keratin, type I cytoskeletal 14 OS=Homo sapiens GN=KRT14 PE=1 SV=4
K1C16_HUMAN	186	51578	9 (6)	Keratin, type I cytoskeletal 16 OS=Homo sapiens GN=KRT16 PE=1 SV=4
PPIA_HUMAN	1342	18229	12 (11)	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2
K2C1_HUMAN	1291	66170	26 (22)	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6
K22E_HUMAN	415	65678	14 (10)	Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2
K2C6A_HUMAN	232	60293	10 (6)	Keratin, type II cytoskeletal 6A OS=Homo sapiens GN=KRT6A PE=1 SV=3
K2C5_HUMAN	124	62568	9 (5)	Keratin, type II cytoskeletal 5 OS=Homo sapiens GN=KRT5 PE=1 SV=3
ACTB_HUMAN	1235	42052	18 (17)	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1
ACTC_HUMAN	772	42334	11 (11)	Actin, alpha cardiac muscle 1 OS=Homo sapiens GN=ACTC1 PE=1 SV=1
H2B1D_HUMAN	1040	13928	8 (5)	Histone H2B type 1-D OS=Homo sapiens GN=HIST1H2BD PE=1 SV=2
H2B1B_HUMAN	1003	13942	5 (5)	Histone H2B type 1-B OS=Homo sapiens GN=HIST1H2BB PE=1 SV=2
TAGL2_HUMAN	1000	22548	14 (12)	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3
ATPB_HUMAN	965	56525	19 (16)	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3
KAD2_HUMAN	954	26689	16 (13)	Adenylate kinase 2, mitochondrial OS=Homo sapiens GN=AK2 PE=1 SV=2
K1C9_HUMAN	903	62255	20 (15)	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3
HMGB1_HUMAN	733	25049	10 (10)	High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3
HMGB2_HUMAN	462	24190	10 (7)	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2
PROF1_HUMAN	687	15216	8 (8)	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2
NPM_HUMAN	676	32726	8 (6)	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2
PDIA1_HUMAN	656	57480	16 (12)	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3
CAP1_HUMAN	641	52222	16 (10)	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=4
TBB5_HUMAN	595	50095	18 (14)	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2
TBB2C_HUMAN	402	50255	15 (12)	Tubulin beta-2C chain OS=Homo sapiens GN=TUBB2C PE=1 SV=1
H4_HUMAN	592	11360	9 (9)	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2
RNT2_HUMAN	583	30089	12 (9)	Ribonuclease T2 OS=Homo sapiens GN=RNASET2 PE=1 SV=2
HSPB1_HUMAN	574	22826	15 (14)	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2
COF1_HUMAN	567	18719	9 (8)	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3
CH10_HUMAN	562	10925	7 (6)	10 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPE1 PE=1 SV=2
NDKB_HUMAN	537	17401	10 (7)	Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2 PE=1 SV=1
NDKA_HUMAN	525	17309	10 (7)	Nucleoside diphosphate kinase A OS=Homo sapiens GN=NME1 PE=1 SV=1
ATPA_HUMAN	529	59828	11 (7)	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1
TBA1B_HUMAN	527	50804	13 (9)	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1
EF1A1_HUMAN	516	50451	9 (8)	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1
G3P_HUMAN	505	36201	10 (7)	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3
C1QBP_HUMAN	491	31742	5 (5)	Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QBP PE=1 SV=1
RAP1B_HUMAN	475	21040	9 (8)	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1

CLIC1_HUMAN	470	27248	10 (6)	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4
AN32A_HUMAN	464	28682	11 (9)	Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Homo sapiens GN=ANP32A PE=1 SV=1
AN32B_HUMAN	403	28941	9 (8)	Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Homo sapiens GN=ANP32B PE=1 SV=1
PPIB_HUMAN	453	23785	12 (8)	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2
1433Z_HUMAN	439	27899	11 (8)	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1
1433B_HUMAN	254	28179	10 (5)	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3
1433T_HUMAN	226	28032	7 (3)	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1
1433E_HUMAN	124	29326	9 (3)	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1
1433G_HUMAN	88	28456	8 (3)	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2
PEBP1_HUMAN	437	21158	7 (7)	Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3
TPD54_HUMAN	408	22281	6 (5)	Tumor protein D54 OS=Homo sapiens GN=TPD52L2 PE=1 SV=2
RS5_HUMAN	380	23033	8 (6)	40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4
CYC_HUMAN	366	11855	8 (5)	Cytochrome c OS=Homo sapiens GN=CYCS PE=1 SV=2
BID_HUMAN	362	22152	7 (2)	BH3-interacting domain death agonist OS=Homo sapiens GN=BID PE=1 SV=1
PARK7_HUMAN	361	20050	8 (6)	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2
CY1_HUMAN	357	35709	7 (4)	Cytochrome c1, heme protein, mitochondrial OS=Homo sapiens GN=CYC1 PE=1 SV=2
RS14_HUMAN	354	16434	4 (4)	40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3
RLA2_HUMAN	354	11658	6 (5)	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1
ML12A_HUMAN	351	19839	7 (7)	Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=1 SV=2
MYL9_HUMAN	159	19871	5 (5)	Myosin regulatory light polypeptide 9 OS=Homo sapiens GN=MYL9 PE=1 SV=4
RL12_HUMAN	351	17979	6 (5)	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1
IF5A1_HUMAN	349	17049	7 (4)	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2
AN32E_HUMAN	345	30902	5 (5)	Acidic leucine-rich nuclear phosphoprotein 32 family member E OS=Homo sapiens GN=ANP32E PE=1 SV=1
CALX_HUMAN	332	67982	8 (5)	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2
ENOA_HUMAN	327	47481	12 (7)	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2
PRDX6_HUMAN	322	25133	7 (5)	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3
ROA2_HUMAN	322	37464	9 (7)	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2
ROA1_HUMAN	101	38837	7 (3)	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5
CYB5B_HUMAN	316	16436	3 (3)	Cytochrome b5 type B OS=Homo sapiens GN=CYB5B PE=1 SV=2
CNPY2_HUMAN	304	20981	6 (5)	Protein canopy homolog 2 OS=Homo sapiens GN=CNPY2 PE=1 SV=1
ATP5H_HUMAN	296	18537	12 (8)	ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3
VDAC1_HUMAN	296	30868	10 (6)	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2
VDAC3_HUMAN	228	30981	3 (3)	Voltage-dependent anion-selective channel protein 3 OS=Homo sapiens GN=VDAC3 PE=1 SV=1
COX5A_HUMAN	295	16923	11 (6)	Cytochrome c oxidase subunit 5A, mitochondrial OS=Homo sapiens GN=COX5A PE=1 SV=2
SODC_HUMAN	290	16154	3 (2)	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2
H12_HUMAN	286	21352	8 (6)	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2
LEG1_HUMAN	284	15048	7 (6)	Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2
HNRPK_HUMAN	283	51230	9 (6)	Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1
GRP75_HUMAN	281	73920	10 (8)	Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2
MDHM_HUMAN	280	35937	13 (10)	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3
ATPO_HUMAN	277	23377	8 (7)	ATP synthase subunit O, mitochondrial OS=Homo sapiens GN=ATP5O PE=1 SV=1
ERP29_HUMAN	275	29032	9 (7)	Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=1 SV=4

TCP4_HUMAN	265	14386	7 (4)	Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SUB1 PE=1 SV=3
RL30_HUMAN	261	12947	3 (3)	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2
LYRIC_HUMAN	257	63856	5 (2)	Protein LYRIC OS=Homo sapiens GN=MTDH PE=1 SV=2
RS13_HUMAN	255	17212	9 (7)	40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2
RU2A_HUMAN	254	28512	5 (5)	U2 small nuclear ribonucleoprotein A' OS=Homo sapiens GN=SNRPA1 PE=1 SV=2
PGRC2_HUMAN	252	23861	5 (5)	Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRMC2 PE=1 SV=1
PGRC1_HUMAN	134	21772	7 (3)	Membrane-associated progesterone receptor component 1 OS=Homo sapiens GN=PGRMC1 PE=1 SV=3
HBA_HUMAN	251	15305	7 (5)	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2
RNAS2_HUMAN	250	18855	3 (3)	Non-secretory ribonuclease OS=Homo sapiens GN=RNASE2 PE=1 SV=2
RS25_HUMAN	246	13791	6 (5)	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1
RS19_HUMAN	246	16051	8 (6)	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2
NDUA5_HUMAN	244	13507	5 (4)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 OS=Homo sapiens GN=NDUFA5 PE=1 SV=3
PRDX3_HUMAN	241	28017	6 (5)	Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3 PE=1 SV=3
RS18_HUMAN	240	17708	7 (6)	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3
CAH2_HUMAN	239	29285	8 (5)	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2
RL11_HUMAN	239	20468	3 (2)	60S ribosomal protein L11 OS=Homo sapiens GN=RPL11 PE=1 SV=2
RL18_HUMAN	237	21735	3 (3)	60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2
ALDOA_HUMAN	231	39851	7 (6)	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2
ARF1_HUMAN	230	20741	10 (6)	ADP-ribosylation factor 1 OS=Homo sapiens GN=ARF1 PE=1 SV=2
HBB_HUMAN	229	16102	5 (4)	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2
HBG1_HUMAN	166	16187	3 (2)	Hemoglobin subunit gamma-1 OS=Homo sapiens GN=HBG1 PE=1 SV=2
HSP7C_HUMAN	228	71082	8 (6)	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1
SC61B_HUMAN	225	10025	3 (2)	Protein transport protein Sec61 subunit beta OS=Homo sapiens GN=SEC61B PE=1 SV=2
ECH1_HUMAN	224	36136	7 (5)	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH1 PE=1 SV=2
PRDX5_HUMAN	223	22298	9 (6)	Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=3
H2A1D_HUMAN	223	14099	7 (4)	Histone H2A type 1-D OS=Homo sapiens GN=HIST1H2AD PE=1 SV=2
H2A1A_HUMAN	142	14225	4 (2)	Histone H2A type 1-A OS=Homo sapiens GN=HIST1H2AA PE=1 SV=3
RSSA_HUMAN	220	32947	6 (4)	40S ribosomal protein SA OS=Homo sapiens GN=RPSA PE=1 SV=4
CUTA_HUMAN	219	19218	3 (3)	Protein CutA OS=Homo sapiens GN=CUTA PE=1 SV=2
SFRS1_HUMAN	219	27842	10 (6)	Splicing factor, arginine/serine-rich 1 OS=Homo sapiens GN=SFRS1 PE=1 SV=2
ATP5L_HUMAN	217	11421	3 (3)	ATP synthase subunit g, mitochondrial OS=Homo sapiens GN=ATP5L PE=1 SV=3
TMEDA_HUMAN	216	25131	3 (3)	Transmembrane emp24 domain-containing protein 10 OS=Homo sapiens GN=TMED10 PE=1 SV=2
RAB1B_HUMAN	216	22328	6 (5)	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1
RAB10_HUMAN	191	22755	5 (4)	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1
RAB8A_HUMAN	135	23824	4 (3)	Ras-related protein Rab-8A OS=Homo sapiens GN=RAB8A PE=1 SV=1
RAB6A_HUMAN	106	23692	3 (2)	Ras-related protein Rab-6A OS=Homo sapiens GN=RAB6A PE=1 SV=3
LGUL_HUMAN	215	20992	11 (6)	Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 PE=1 SV=4
RS20_HUMAN	214	13478	5 (4)	40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1
RS28_HUMAN	212	7893	2 (2)	40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1
LYSC_HUMAN	211	16982	6 (4)	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1
MYL6_HUMAN	209	17090	6 (5)	Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2
SFRS3_HUMAN	208	19546	5 (4)	Splicing factor, arginine/serine-rich 3 OS=Homo sapiens GN=SFRS3 PE=1 SV=1

SFRS7_HUMAN	48	27578	5 (3)	Splicing factor, arginine/serine-rich 7 OS=Homo sapiens GN=SFRS7 PE=1 SV=1
TBCA_HUMAN	206	12904	5 (3)	Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=1 SV=3
COTL1_HUMAN	204	16049	5 (2)	Coactosin-like protein OS=Homo sapiens GN=COTL1 PE=1 SV=3
UBIQ_HUMAN	203	8560	4 (3)	Ubiquitin OS=Homo sapiens GN=RPS27A PE=1 SV=1
COX2_HUMAN	202	25719	5 (4)	Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1
GSTP1_HUMAN	199	23569	4 (4)	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2
NDUS3_HUMAN	198	30337	6 (3)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Homo sapiens GN=NDUFS3 PE=1 SV=1
RAB5C_HUMAN	197	23696	6 (6)	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2
RAN_HUMAN	194	24579	5 (3)	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3
SSRD_HUMAN	193	19158	4 (3)	Translocon-associated protein subunit delta OS=Homo sapiens GN=SSR4 PE=1 SV=1
UB2L3_HUMAN	192	18021	5 (2)	Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1
PPT1_HUMAN	192	34627	2 (2)	Palmitoyl-protein thioesterase 1 OS=Homo sapiens GN=PPT1 PE=1 SV=1
RL14_HUMAN	191	23531	3 (2)	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4
SSBP_HUMAN	190	17249	3 (3)	Single-stranded DNA-binding protein, mitochondrial OS=Homo sapiens GN=SSBP1 PE=1 SV=1
RS17_HUMAN	187	15597	6 (6)	40S ribosomal protein S17 OS=Homo sapiens GN=RPS17 PE=1 SV=2
COMT_HUMAN	185	30474	7 (3)	Catechol O-methyltransferase OS=Homo sapiens GN=COMT PE=1 SV=2
RL38_HUMAN	185	8270	4 (3)	60S ribosomal protein L38 OS=Homo sapiens GN=RPL38 PE=1 SV=2
ATPK_HUMAN	183	11025	2 (2)	ATP synthase subunit f, mitochondrial OS=Homo sapiens GN=ATP5J2 PE=1 SV=3
H31_HUMAN	182	15509	8 (3)	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2
H32_HUMAN	170	15436	8 (3)	Histone H3.2 OS=Homo sapiens GN=HIST2H3A PE=1 SV=3
H33_HUMAN	118	15376	8 (3)	Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2
HS90B_HUMAN	180	83554	12 (6)	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4
ENPL_HUMAN	159	92696	8 (2)	Endoplasmin OS=Homo sapiens GN=HSP90B1 PE=1 SV=1
PDIA6_HUMAN	180	48490	3 (3)	Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1
NDUA7_HUMAN	179	12601	4 (2)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=Homo sapiens GN=NDUFA7 PE=1 SV=3
ARPC4_HUMAN	177	19768	5 (4)	Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3
SRP14_HUMAN	176	14675	4 (3)	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2
ATPD_HUMAN	175	17479	2 (1)	ATP synthase subunit delta, mitochondrial OS=Homo sapiens GN=ATP5D PE=1 SV=2
GP1BB_HUMAN	175	22274	4 (2)	Platelet glycoprotein Ib beta chain OS=Homo sapiens GN=GP1BB PE=1 SV=1
PRDX1_HUMAN	171	22324	10 (8)	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1
PRDX2_HUMAN	108	22049	4 (4)	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5
PRDX4_HUMAN	61	30749	2 (2)	Peroxiredoxin-4 OS=Homo sapiens GN=PRDX4 PE=1 SV=1
CXCL7_HUMAN	169	14171	6 (5)	Platelet basic protein OS=Homo sapiens GN=PPBP PE=1 SV=3
MA2B1_HUMAN	169	114357	5 (3)	Lysosomal alpha-mannosidase OS=Homo sapiens GN=MAN2B1 PE=1 SV=3
UB2V2_HUMAN	168	16409	5 (4)	Ubiquitin-conjugating enzyme E2 variant 2 OS=Homo sapiens GN=UBE2V2 PE=1 SV=4
PHB_HUMAN	167	29843	6 (4)	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1
VAPA_HUMAN	166	28103	5 (5)	Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA PE=1 SV=3
PA1B2_HUMAN	166	25724	3 (2)	Platelet-activating factor acetylhydrolase IB subunit beta OS=Homo sapiens GN=PAFAH1B2 PE=1 SV=1
CX6B1_HUMAN	165	10414	4 (3)	Cytochrome c oxidase subunit 6B1 OS=Homo sapiens GN=COX6B1 PE=1 SV=2
APT_HUMAN	163	19766	4 (4)	Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2
GDIR1_HUMAN	163	23250	2 (2)	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDIA PE=1 SV=3
RAB2A_HUMAN	160	23702	5 (5)	Ras-related protein Rab-2A OS=Homo sapiens GN=RAB2A PE=1 SV=1

S10A4_HUMAN	158	11949	3 (1)	Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1
TMCO1_HUMAN	157	21389	3 (1)	Transmembrane and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=TMCO1 PE=1 SV=1
QCR7_HUMAN	156	13522	8 (6)	Cytochrome b-c1 complex subunit 7 OS=Homo sapiens GN=UQCRB PE=1 SV=2
CISD2_HUMAN	153	15497	3 (2)	CDGSH iron sulfur domain-containing protein 2 OS=Homo sapiens GN=CISD2 PE=1 SV=1
RHOA_HUMAN	152	22096	7 (4)	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1
VAMP8_HUMAN	151	11488	4 (4)	Vesicle-associated membrane protein 8 OS=Homo sapiens GN=VAMP8 PE=1 SV=1
PSB1_HUMAN	150	26700	1 (1)	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2
RAB7A_HUMAN	149	23760	9 (6)	Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1
ARPC5_HUMAN	149	16367	4 (4)	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3
USMG5_HUMAN	147	6510	2 (1)	Up-regulated during skeletal muscle growth protein 5 OS=Homo sapiens GN=USMG5 PE=1 SV=1
UBE2N_HUMAN	147	17184	3 (2)	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1
TFAM_HUMAN	146	29306	6 (3)	Transcription factor A, mitochondrial OS=Homo sapiens GN=TFAM PE=1 SV=1
SSRA_HUMAN	146	32215	3 (2)	Translocon-associated protein subunit alpha OS=Homo sapiens GN=SSR1 PE=1 SV=3
BAX_HUMAN	144	21285	2 (2)	Apoptosis regulator BAX OS=Homo sapiens GN=BAX PE=1 SV=1
MIF_HUMAN	143	12639	1 (1)	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4
RS12_HUMAN	143	14905	5 (5)	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3
SLIRP_HUMAN	142	12398	6 (5)	SRA stem-loop-interacting RNA-binding protein, mitochondrial OS=Homo sapiens GN=SLIRP PE=1 SV=1
VDAC2_HUMAN	138	32060	6 (3)	Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1 SV=2
ALBU_HUMAN	137	71317	7 (4)	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2
ITA2B_HUMAN	135	114446	3 (2)	Integrin alpha-IIb OS=Homo sapiens GN=ITGA2B PE=1 SV=3
H15_HUMAN	134	22566	4 (3)	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3
GANAB_HUMAN	134	107263	4 (3)	Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3
ATP5I_HUMAN	132	7928	3 (2)	ATP synthase subunit e, mitochondrial OS=Homo sapiens GN=ATP5I PE=1 SV=2
SMD1_HUMAN	132	13273	4 (2)	Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1
LAP2B_HUMAN	132	50696	5 (4)	Lamina-associated polypeptide 2, isoforms beta/gamma OS=Homo sapiens GN=TMPO PE=1 SV=2
RS16_HUMAN	130	16549	3 (3)	40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2
SODM_HUMAN	128	24878	3 (2)	Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2
EF1B_HUMAN	127	24919	3 (3)	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3
PDIA3_HUMAN	127	57146	7 (5)	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4
RL27A_HUMAN	127	16665	4 (3)	60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=1 SV=2
NDUA4_HUMAN	126	9421	4 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=Homo sapiens GN=NDUFA4 PE=1 SV=1
RL36_HUMAN	126	12303	4 (3)	60S ribosomal protein L36 OS=Homo sapiens GN=RPL36 PE=1 SV=3
CALR_HUMAN	125	48283	7 (4)	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1
TFIP8_HUMAN	124	23160	1 (1)	Tumor necrosis factor alpha-induced protein 8 OS=Homo sapiens GN=TNFAIP8 PE=1 SV=1
TPP1_HUMAN	123	61723	2 (2)	Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1 PE=1 SV=2
RL35_HUMAN	121	14543	3 (3)	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2
OCAD1_HUMAN	121	27780	2 (2)	OCIA domain-containing protein 1 OS=Homo sapiens GN=OCIAD1 PE=1 SV=1
TM109_HUMAN	121	26194	1 (1)	Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1
NDUB4_HUMAN	121	15256	3 (3)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 OS=Homo sapiens GN=NDUFB4 PE=1 SV=3
RL22_HUMAN	120	14835	4 (2)	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2
RS15A_HUMAN	120	14944	7 (4)	40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2
RS10_HUMAN	120	18886	4 (4)	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1

CREG1_HUMAN	120	24173	1 (1)	Protein CREG1 OS=Homo sapiens GN=CREG1 PE=1 SV=1
ARL8B_HUMAN	119	21753	4 (3)	ADP-ribosylation factor-like protein 8B OS=Homo sapiens GN=ARL8B PE=1 SV=1
SERA_HUMAN	118	57356	4 (2)	D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4
APOA1_HUMAN	117	30759	2 (1)	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1
RLA1_HUMAN	116	11621	2 (2)	60S acidic ribosomal protein P1 OS=Homo sapiens GN=RPLP1 PE=1 SV=1
1A01_HUMAN	116	41105	2 (2)	HLA class I histocompatibility antigen, A-1 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=1
SMD3_HUMAN	115	14021	4 (3)	Small nuclear ribonucleoprotein Sm D3 OS=Homo sapiens GN=SNRPD3 PE=1 SV=1
KAD3_HUMAN	114	25550	5 (4)	GTP:AMP phosphotransferase mitochondrial OS=Homo sapiens GN=AK3 PE=1 SV=4
HIG2A_HUMAN	113	11578	1 (1)	HIG1 domain family member 2A OS=Homo sapiens GN=HIGD2A PE=1 SV=1
SFRS9_HUMAN	111	25640	4 (3)	Splicing factor, arginine/serine-rich 9 OS=Homo sapiens GN=SFRS9 PE=1 SV=1
GPX1_HUMAN	111	22218	5 (2)	Glutathione peroxidase 1 OS=Homo sapiens GN=GPX1 PE=1 SV=3
NDUBA_HUMAN	110	21048	5 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3
LDHB_HUMAN	110	36900	8 (4)	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2
LDHA_HUMAN	87	36950	5 (3)	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2
NDUAD_HUMAN	109	16688	5 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Homo sapiens GN=NDUFA13 PE=1 SV=3
TEBP_HUMAN	109	18971	3 (1)	Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1
ARP5L_HUMAN	109	16931	3 (2)	Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5L PE=1 SV=1
RS7_HUMAN	108	22113	4 (3)	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1
HNRH1_HUMAN	107	49484	6 (3)	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4
NDUA8_HUMAN	106	20548	4 (3)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=Homo sapiens GN=NDUFA8 PE=1 SV=3
CE032_HUMAN	106	10966	1 (1)	UPF0467 protein C5orf32 OS=Homo sapiens GN=C5orf32 PE=2 SV=1
RL23_HUMAN	104	14970	3 (2)	60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1
PSB3_HUMAN	104	23219	2 (1)	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2
RM23_HUMAN	103	17770	3 (3)	39S ribosomal protein L23, mitochondrial OS=Homo sapiens GN=MRPL23 PE=1 SV=1
PFD2_HUMAN	102	16695	2 (2)	Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1
RB11A_HUMAN	102	24492	5 (3)	Ras-related protein Rab-11A OS=Homo sapiens GN=RAB11A PE=1 SV=3
LRC59_HUMAN	102	35308	8 (6)	Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRRC59 PE=1 SV=1
HCD2_HUMAN	101	27134	4 (3)	3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3
RS24_HUMAN	101	15413	3 (2)	40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=1 SV=1
RL24_HUMAN	101	17882	5 (3)	60S ribosomal protein L24 OS=Homo sapiens GN=RPL24 PE=1 SV=1
ADT3_HUMAN	99	33073	9 (4)	ADP/ATP translocase 3 OS=Homo sapiens GN=SLC25A6 PE=1 SV=4
PDCD6_HUMAN	99	21912	4 (2)	Programmed cell death protein 6 OS=Homo sapiens GN=PDCD6 PE=1 SV=1
HEXB_HUMAN	98	63527	4 (2)	Beta-hexosaminidase subunit beta OS=Homo sapiens GN=HEXB PE=1 SV=3
HNRPC_HUMAN	98	33707	6 (4)	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=1 SV=4
ARPC3_HUMAN	97	20761	4 (2)	Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3
SNP23_HUMAN	95	23682	3 (2)	Synaptosomal-associated protein 23 OS=Homo sapiens GN=SNAP23 PE=1 SV=1
RS23_HUMAN	94	15969	2 (2)	40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=1 SV=3
RL31_HUMAN	94	14454	2 (2)	60S ribosomal protein L31 OS=Homo sapiens GN=RPL31 PE=1 SV=1
HNRPG_HUMAN	93	42306	6 (3)	Heterogeneous nuclear ribonucleoprotein G OS=Homo sapiens GN=RBMX PE=1 SV=3
RAC2_HUMAN	93	21814	4 (3)	Ras-related C3 botulinum toxin substrate 2 OS=Homo sapiens GN=RAC2 PE=1 SV=1
NDUB9_HUMAN	91	22045	2 (2)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 9 OS=Homo sapiens GN=NDUFB9 PE=1 SV=3
RM17_HUMAN	91	20095	6 (1)	39S ribosomal protein L17, mitochondrial OS=Homo sapiens GN=MRPL17 PE=1 SV=1

TOM22_HUMAN	91	15512	3 (1)	Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=1 SV=3
PSB6_HUMAN	91	25570	2 (1)	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4
HMGB3_HUMAN	88	23137	4 (2)	High mobility group protein B3 OS=Homo sapiens GN=HMGB3 PE=1 SV=4
EFTU_HUMAN	88	49852	2 (1)	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2
ACPM_HUMAN	88	17577	3 (1)	Acyl carrier protein, mitochondrial OS=Homo sapiens GN=NDUFAB1 PE=1 SV=3
RUXE_HUMAN	88	10854	2 (2)	Small nuclear ribonucleoprotein E OS=Homo sapiens GN=SNRPE PE=1 SV=1
COX41_HUMAN	88	19621	4 (3)	Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Homo sapiens GN=COX4I1 PE=1 SV=1
RLA0L_HUMAN	87	34514	4 (2)	60S acidic ribosomal protein P0-like OS=Homo sapiens PE=1 SV=1
CX7A2_HUMAN	87	9390	3 (1)	Cytochrome c oxidase subunit 7A2, mitochondrial OS=Homo sapiens GN=COX7A2 PE=1 SV=1
RL7_HUMAN	86	29264	4 (1)	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1
TPIS_HUMAN	86	26938	4 (1)	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=2
VATG1_HUMAN	86	13863	3 (3)	V-type proton ATPase subunit G 1 OS=Homo sapiens GN=ATP6V1G1 PE=1 SV=3
NDUS5_HUMAN	86	12737	3 (3)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 5 OS=Homo sapiens GN=NDUFS5 PE=1 SV=3
MTPN_HUMAN	86	13058	1 (1)	Myotrophin OS=Homo sapiens GN=MTPN PE=1 SV=2
RANG_HUMAN	85	23467	5 (3)	Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1
RM12_HUMAN	85	21563	4 (3)	39S ribosomal protein L12, mitochondrial OS=Homo sapiens GN=MRPL12 PE=1 SV=2
FKBP3_HUMAN	84	25218	3 (2)	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1
TCPH_HUMAN	83	59842	3 (1)	T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2
SFR2B_HUMAN	83	32382	2 (1)	Splicing factor, arginine/serine-rich 2B OS=Homo sapiens GN=SFRS2B PE=1 SV=1
LYAG_HUMAN	83	106107	3 (2)	Lysosomal alpha-glucosidase OS=Homo sapiens GN=GAA PE=1 SV=3
COX5B_HUMAN	82	13915	4 (2)	Cytochrome c oxidase subunit 5B, mitochondrial OS=Homo sapiens GN=COX5B PE=1 SV=2
QCR9_HUMAN	82	7304	2 (1)	Cytochrome b-c1 complex subunit 9 OS=Homo sapiens GN=UQCR10 PE=1 SV=3
MGST3_HUMAN	82	16734	1 (1)	Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1
PSA6_HUMAN	81	27838	2 (2)	Proteasome subunit alpha type-6 OS=Homo sapiens GN=PSMA6 PE=1 SV=1
CHSP1_HUMAN	80	16110	1 (1)	Calcium-regulated heat stable protein 1 OS=Homo sapiens GN=CARHSP1 PE=1 SV=2
ANXA5_HUMAN	80	35971	7 (4)	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2
MIMIT_HUMAN	79	19844	4 (1)	Mimitin, mitochondrial OS=Homo sapiens GN=NDUFAF2 PE=1 SV=1
RL17_HUMAN	79	21611	4 (2)	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3
NDUS8_HUMAN	78	24203	4 (3)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Homo sapiens GN=NDUFS8 PE=1 SV=1
RS15_HUMAN	78	17029	4 (2)	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2
QCR1_HUMAN	77	53297	2 (1)	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 SV=3
PLF4_HUMAN	77	11123	4 (4)	Platelet factor 4 OS=Homo sapiens GN=PF4 PE=1 SV=2
CPSF6_HUMAN	77	59344	3 (1)	Cleavage and polyadenylation specificity factor subunit 6 OS=Homo sapiens GN=CPSF6 PE=1 SV=2
FABP5_HUMAN	77	15497	2 (2)	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3
ROAA_HUMAN	77	36316	4 (3)	Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=1 SV=2
HNRDL_HUMAN	34	46580	2 (2)	Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRPDL PE=1 SV=3
PCBP2_HUMAN	77	38955	3 (2)	Poly(rC)-binding protein 2 OS=Homo sapiens GN=PCBP2 PE=1 SV=1
PRTN3_HUMAN	76	28245	2 (2)	Myeloblastin OS=Homo sapiens GN=PRTN3 PE=1 SV=3
CHP1_HUMAN	75	22442	2 (2)	Calcium-binding protein p22 OS=Homo sapiens GN=CHP PE=1 SV=3
DECR_HUMAN	75	36330	2 (2)	2,4-dienoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=DECR1 PE=1 SV=1
HP1B3_HUMAN	75	61454	4 (1)	Heterochromatin protein 1-binding protein 3 OS=Homo sapiens GN=HP1BP3 PE=1 SV=1
ECHM_HUMAN	75	31823	3 (1)	Enoyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=ECHS1 PE=1 SV=4

TMED2_HUMAN	75	22860	2 (1)	Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1
DHSA_HUMAN	73	73672	2 (1)	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=2
ITB2_HUMAN	73	87976	2 (2)	Integrin beta-2 OS=Homo sapiens GN=ITGB2 PE=1 SV=2
PLSL_HUMAN	73	70815	5 (2)	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=5
NLTP_HUMAN	73	59640	4 (3)	Non-specific lipid-transfer protein OS=Homo sapiens GN=SCP2 PE=1 SV=2
DUT_HUMAN	72	26975	4 (3)	Deoxyuridine 5'-triphosphate nucleotidohydrolase, mitochondrial OS=Homo sapiens GN=DUT PE=1 SV=3
PGK1_HUMAN	72	44985	4 (1)	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3
ACTN1_HUMAN	71	103563	4 (1)	Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2
RL19_HUMAN	70	23565	2 (2)	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1
BR44L_HUMAN	70	12510	2 (1)	Brain protein 44-like protein OS=Homo sapiens GN=BRP44L PE=1 SV=1
RL13_HUMAN	70	24304	5 (1)	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4
CBX3_HUMAN	69	20969	3 (1)	Chromobox protein homolog 3 OS=Homo sapiens GN=CBX3 PE=1 SV=4
RL27_HUMAN	69	15788	2 (2)	60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2
RM41_HUMAN	68	15430	6 (2)	39S ribosomal protein L41, mitochondrial OS=Homo sapiens GN=MRPL41 PE=1 SV=1
GBB1_HUMAN	67	38151	4 (2)	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3
COR1A_HUMAN	66	51678	3 (1)	Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4
MATR3_HUMAN	66	95078	2 (1)	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=2
NUCL_HUMAN	66	76625	3 (1)	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3
DOPD_HUMAN	65	12818	2 (2)	D-dopachrome decarboxylase OS=Homo sapiens GN=DDT PE=1 SV=3
NDUF4_HUMAN	65	20311	1 (1)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 4 OS=Homo sapiens GN=NDUFAF4 PE=1 SV=1
TXD12_HUMAN	64	19365	3 (2)	Thioredoxin domain-containing protein 12 OS=Homo sapiens GN=TXNDC12 PE=1 SV=1
G6PI_HUMAN	64	63335	1 (1)	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4
S10AB_HUMAN	64	11847	5 (1)	Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2
LPPRC_HUMAN	63	159003	7 (2)	Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3
RL9_HUMAN	63	21964	5 (2)	60S ribosomal protein L9 OS=Homo sapiens GN=RPL9 PE=1 SV=1
RL23A_HUMAN	63	17684	6 (2)	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1
RL6_HUMAN	63	32765	3 (2)	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3
PGAM2_HUMAN	62	28919	3 (2)	Phosphoglycerate mutase 2 OS=Homo sapiens GN=PGAM2 PE=1 SV=3
PPIH_HUMAN	61	19481	1 (1)	Peptidyl-prolyl cis-trans isomerase H OS=Homo sapiens GN=PPIH PE=1 SV=1
CNN2_HUMAN	60	34074	3 (1)	Calponin-2 OS=Homo sapiens GN=CNN2 PE=1 SV=4
TMED9_HUMAN	60	27374	2 (2)	Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=2
RL18A_HUMAN	60	21034	3 (2)	60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A PE=1 SV=2
ETFA_HUMAN	60	35400	2 (1)	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Homo sapiens GN=ETFA PE=1 SV=1
TCPQ_HUMAN	60	60153	2 (1)	T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4
MOES_HUMAN	59	67892	10 (1)	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3
NDUB1_HUMAN	58	7014	1 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 1 OS=Homo sapiens GN=NDUFB1 PE=1 SV=1
BAG2_HUMAN	58	23928	2 (1)	BAG family molecular chaperone regulator 2 OS=Homo sapiens GN=BAG2 PE=1 SV=1
RS27_HUMAN	58	9797	3 (2)	40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3
PSA7_HUMAN	57	28041	2 (1)	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1
GRN_HUMAN	57	68518	2 (1)	Granulins OS=Homo sapiens GN=GRN PE=1 SV=2
TCTP_HUMAN	57	19697	3 (1)	Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1
FKB1A_HUMAN	56	12000	1 (1)	Peptidyl-prolyl cis-trans isomerase FKBP1A OS=Homo sapiens GN=FKBP1A PE=1 SV=2

PPIF_HUMAN	56	22368	3 (3)	Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Homo sapiens GN=PPIF PE=1 SV=1
MPCP_HUMAN	56	40525	3 (2)	Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2
GDIR2_HUMAN	56	23031	4 (3)	Rho GDP-dissociation inhibitor 2 OS=Homo sapiens GN=ARHGDIB PE=1 SV=3
IDHP_HUMAN	55	51333	4 (2)	Isocitrate dehydrogenase [NADP], mitochondrial OS=Homo sapiens GN=IDH2 PE=1 SV=2
RL29_HUMAN	55	17798	1 (1)	60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2
NDUBB_HUMAN	54	17363	3 (2)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 11, mitochondrial OS=Homo sapiens GN=NDUFB11 PE=
TKT_HUMAN	54	68519	1 (1)	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3
STMN1_HUMAN	54	17292	5 (1)	Stathmin OS=Homo sapiens GN=STMN1 PE=1 SV=3
RL10A_HUMAN	54	24987	4 (2)	60S ribosomal protein L10a OS=Homo sapiens GN=RPL10A PE=1 SV=2
LSM4_HUMAN	53	15511	5 (1)	U6 snRNA-associated Sm-like protein LSm4 OS=Homo sapiens GN=LSM4 PE=1 SV=1
ASC_HUMAN	53	21670	1 (1)	Apoptosis-associated speck-like protein containing a CARD OS=Homo sapiens GN=PYCARD PE=1 SV=2
ASPG_HUMAN	53	37812	2 (2)	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase OS=Homo sapiens GN=AGA PE=1 SV=2
SORCN_HUMAN	53	21947	5 (1)	Sorcin OS=Homo sapiens GN=SRI PE=1 SV=1
SH3L3_HUMAN	53	10488	1 (1)	SH3 domain-binding glutamic acid-rich-like protein 3 OS=Homo sapiens GN=SH3BGRL3 PE=1 SV=1
NDUV2_HUMAN	53	27659	3 (2)	NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Homo sapiens GN=NDUFV2 PE=1 SV=2
AT5EL_HUMAN	52	5860	1 (1)	ATP synthase subunit epsilon-like protein, mitochondrial OS=Homo sapiens GN=ATP5EP2 PE=1 SV=1
HINT1_HUMAN	52	13907	2 (1)	Histidine triad nucleotide-binding protein 1 OS=Homo sapiens GN=HINT1 PE=1 SV=2
PLIN3_HUMAN	52	47189	4 (2)	Perilipin-3 OS=Homo sapiens GN=PLIN3 PE=1 SV=2
NH2L1_HUMAN	52	14393	1 (1)	NHP2-like protein 1 OS=Homo sapiens GN=NHP2L1 PE=1 SV=3
PSME2_HUMAN	52	27515	2 (2)	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=3
MMAB_HUMAN	52	27713	1 (1)	Cob(I)yrinic acid a,c-diamide adenosyltransferase, mitochondrial OS=Homo sapiens GN=MMAB PE=1 SV=1
TCPG_HUMAN	51	61066	6 (2)	T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4
QCR6_HUMAN	51	11017	2 (2)	Cytochrome b-c1 complex subunit 6, mitochondrial OS=Homo sapiens GN=UQCRH PE=1 SV=2
RL36A_HUMAN	51	12718	3 (1)	60S ribosomal protein L36a OS=Homo sapiens GN=RPL36A PE=1 SV=2
IDH3A_HUMAN	51	40022	6 (1)	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Homo sapiens GN=IDH3A PE=1 SV=1
GBLP_HUMAN	50	35511	3 (2)	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3
ATPG_HUMAN	50	33032	3 (1)	ATP synthase subunit gamma, mitochondrial OS=Homo sapiens GN=ATP5C1 PE=1 SV=1
FKBP2_HUMAN	50	15810	1 (1)	Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Homo sapiens GN=FKBP2 PE=1 SV=2
LYRM7_HUMAN	49	12004	2 (1)	LYR motif-containing protein 7 OS=Homo sapiens GN=LYRM7 PE=1 SV=1
RM14_HUMAN	49	16165	1 (1)	39S ribosomal protein L14, mitochondrial OS=Homo sapiens GN=MRPL14 PE=1 SV=1
RS6_HUMAN	48	28834	3 (1)	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1
LMAN2_HUMAN	48	40545	3 (2)	Vesicular integral-membrane protein VIP36 OS=Homo sapiens GN=LMAN2 PE=1 SV=1
TMX1_HUMAN	48	32170	2 (1)	Thioredoxin-related transmembrane protein 1 OS=Homo sapiens GN=TMX1 PE=1 SV=1
EF2_HUMAN	48	96246	9 (2)	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4
RL40_HUMAN	48	6462	2 (1)	60S ribosomal protein L40 OS=Homo sapiens GN=UBA52 PE=1 SV=1
SC11A_HUMAN	47	20612	3 (1)	Signal peptidase complex catalytic subunit SEC11A OS=Homo sapiens GN=SEC11A PE=1 SV=1
SF13A_HUMAN	47	31339	4 (1)	Splicing factor, arginine/serine-rich 13A OS=Homo sapiens GN=SFRS13A PE=1 SV=1
HCDH_HUMAN	47	34313	2 (1)	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=1 SV=2
RM47_HUMAN	47	29603	2 (1)	39S ribosomal protein L47, mitochondrial OS=Homo sapiens GN=MRPL47 PE=1 SV=2
DYL1_HUMAN	47	10530	1 (1)	Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1
CATA_HUMAN	47	59947	4 (1)	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3
RL26_HUMAN	46	17248	4 (1)	60S ribosomal protein L26 OS=Homo sapiens GN=RPL26 PE=1 SV=1

MPKS1_HUMAN	46	13671	1 (1)	Mitogen-activated protein kinase scaffold protein 1 OS=Homo sapiens GN=MAPKSP1 PE=1 SV=1
RM50_HUMAN	46	18484	2 (1)	39S ribosomal protein L50, mitochondrial OS=Homo sapiens GN=MRPL50 PE=1 SV=2
SMD2_HUMAN	46	13632	2 (1)	Small nuclear ribonucleoprotein Sm D2 OS=Homo sapiens GN=SNRPD2 PE=1 SV=1
CALM_HUMAN	46	16827	1 (1)	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2
RL32_HUMAN	46	15964	3 (2)	60S ribosomal protein L32 OS=Homo sapiens GN=RPL32 PE=1 SV=2
IF4H_HUMAN	45	27425	2 (1)	Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H PE=1 SV=5
HNRPU_HUMAN	45	91269	4 (2)	Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6
CO024_HUMAN	45	26454	1 (1)	UPF0480 protein C15orf24 OS=Homo sapiens GN=C15orf24 PE=1 SV=1
NDUB8_HUMAN	45	21865	3 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrial OS=Homo sapiens GN=NDUFB8 PE=1
SNX3_HUMAN	44	18808	3 (3)	Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3
FCERG_HUMAN	44	9775	2 (1)	High affinity immunoglobulin epsilon receptor subunit gamma OS=Homo sapiens GN=FCER1G PE=1 SV=1
CATG_HUMAN	44	29161	3 (1)	Cathepsin G OS=Homo sapiens GN=CTSG PE=1 SV=2
GSTM3_HUMAN	44	26998	4 (1)	Glutathione S-transferase Mu 3 OS=Homo sapiens GN=GSTM3 PE=1 SV=3
PTBP1_HUMAN	43	57357	2 (1)	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1
IMB1_HUMAN	43	98420	4 (1)	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2
S10A6_HUMAN	43	10230	2 (1)	Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1
RB27A_HUMAN	43	25137	3 (3)	Ras-related protein Rab-27A OS=Homo sapiens GN=RAB27A PE=1 SV=3
VATE1_HUMAN	42	26186	3 (1)	V-type proton ATPase subunit E 1 OS=Homo sapiens GN=ATP6V1E1 PE=1 SV=1
GRPE1_HUMAN	42	24492	5 (2)	GrpE protein homolog 1, mitochondrial OS=Homo sapiens GN=GRPEL1 PE=1 SV=2
TCPZ_HUMAN	42	58444	1 (1)	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3
RL15_HUMAN	42	24245	1 (1)	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2
SARNP_HUMAN	41	23713	2 (2)	SAP domain-containing ribonucleoprotein OS=Homo sapiens GN=SARNP PE=1 SV=3
MCA3_HUMAN	41	19855	2 (1)	Eukaryotic translation elongation factor 1 epsilon-1 OS=Homo sapiens GN=EEF1E1 PE=1 SV=1
HPCA_HUMAN	41	22527	2 (1)	Neuron-specific calcium-binding protein hippocalcin OS=Homo sapiens GN=HPCA PE=1 SV=2
PSA5_HUMAN	41	26565	4 (2)	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3
KU70_HUMAN	40	70084	4 (1)	ATP-dependent DNA helicase 2 subunit 1 OS=Homo sapiens GN=XRCC6 PE=1 SV=2
RS2_HUMAN	40	31590	2 (1)	40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2
RS26_HUMAN	40	13292	3 (2)	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3
RINI_HUMAN	40	51766	3 (1)	Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2
PSB2_HUMAN	40	22993	4 (1)	Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1
ARP3B_HUMAN	40	48090	1 (1)	Actin-related protein 3B OS=Homo sapiens GN=ACTR3B PE=1 SV=1
THOC4_HUMAN	39	26872	2 (1)	THO complex subunit 4 OS=Homo sapiens GN=THOC4 PE=1 SV=3
IF4A1_HUMAN	39	46353	2 (1)	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1
BAK_HUMAN	39	23508	1 (1)	Bcl-2 homologous antagonist/killer OS=Homo sapiens GN=BAK1 PE=1 SV=1
RPN1_HUMAN	39	68641	3 (1)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1
RM10_HUMAN	39	29492	1 (1)	39S ribosomal protein L10, mitochondrial OS=Homo sapiens GN=MRPL10 PE=1 SV=3
LAMP2_HUMAN	39	45503	1 (1)	Lysosome-associated membrane glycoprotein 2 OS=Homo sapiens GN=LAMP2 PE=1 SV=2
RM54_HUMAN	39	15866	2 (1)	39S ribosomal protein L54, mitochondrial OS=Homo sapiens GN=MRPL54 PE=1 SV=1
RM43_HUMAN	39	23701	4 (3)	39S ribosomal protein L43, mitochondrial OS=Homo sapiens GN=MRPL43 PE=1 SV=1
SPCS3_HUMAN	39	20358	2 (1)	Signal peptidase complex subunit 3 OS=Homo sapiens GN=SPCS3 PE=1 SV=1
NDUS4_HUMAN	38	20095	3 (1)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial OS=Homo sapiens GN=NDUFS4 PE=1 SV=1
HINT2_HUMAN	38	17208	3 (1)	Histidine triad nucleotide-binding protein 2, mitochondrial OS=Homo sapiens GN=HINT2 PE=1 SV=1

RL10_HUMAN	38	25044	4 (1)	60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=1 SV=4
PFD5_HUMAN	38	17374	2 (1)	Prefoldin subunit 5 OS=Homo sapiens GN=PFDN5 PE=1 SV=2
MDHC_HUMAN	37	36631	2 (1)	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4
STAG3_HUMAN	37	140257	1 (1)	Cohesin subunit SA-3 OS=Homo sapiens GN=STAG3 PE=1 SV=2
RS30_HUMAN	37	6644	2 (1)	40S ribosomal protein S30 OS=Homo sapiens GN=FAU PE=1 SV=1
OSTF1_HUMAN	36	23943	1 (1)	Osteoclast-stimulating factor 1 OS=Homo sapiens GN=OSTF1 PE=1 SV=2
CX4NB_HUMAN	36	24214	1 (1)	Neighbor of COX4 OS=Homo sapiens GN=COX4NB PE=1 SV=1
RRFM_HUMAN	36	29373	2 (1)	Ribosome-recycling factor, mitochondrial OS=Homo sapiens GN=MRRF PE=1 SV=1
GLRX5_HUMAN	36	16732	1 (1)	Glutaredoxin-related protein 5 OS=Homo sapiens GN=GLRX5 PE=1 SV=2
RRS1_HUMAN	35	41225	4 (1)	Ribosome biogenesis regulatory protein homolog OS=Homo sapiens GN=RRS1 PE=1 SV=2
PSA4_HUMAN	35	29750	2 (1)	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1
KU86_HUMAN	35	83222	4 (1)	ATP-dependent DNA helicase 2 subunit 2 OS=Homo sapiens GN=XRCC5 PE=1 SV=3
GLYM_HUMAN	35	56414	4 (1)	Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3
RS4X_HUMAN	35	29807	3 (1)	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2
PFD4_HUMAN	34	15419	1 (1)	Prefoldin subunit 4 OS=Homo sapiens GN=PFDN4 PE=1 SV=1
RL7A_HUMAN	34	30148	3 (1)	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2
MFSD8_HUMAN	34	58046	1 (1)	Major facilitator superfamily domain-containing protein 8 OS=Homo sapiens GN=MFSD8 PE=1 SV=1
TM173_HUMAN	33	42736	1 (1)	Transmembrane protein 173 OS=Homo sapiens GN=TMEM173 PE=1 SV=1
IF4G1_HUMAN	33	176168	2 (1)	Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1 PE=1 SV=3
MANF_HUMAN	33	20700	3 (1)	Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=2
TXTP_HUMAN	33	34333	4 (1)	Tricarboxylate transport protein, mitochondrial OS=Homo sapiens GN=SLC25A1 PE=1 SV=2
RM49_HUMAN	33	19243	1 (1)	39S ribosomal protein L49, mitochondrial OS=Homo sapiens GN=MRPL49 PE=1 SV=1
SKP1_HUMAN	32	18817	3 (1)	S-phase kinase-associated protein 1 OS=Homo sapiens GN=SKP1 PE=1 SV=2
RS11_HUMAN	32	18590	1 (1)	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3
COX6C_HUMAN	32	8776	2 (2)	Cytochrome c oxidase subunit 6C OS=Homo sapiens GN=COX6C PE=1 SV=2
BASP1_HUMAN	32	22680	2 (1)	Brain acid soluble protein 1 OS=Homo sapiens GN=BASP1 PE=1 SV=2
PSB4_HUMAN	31	29242	1 (1)	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4
TSPOA_HUMAN	31	18881	1 (1)	Translocator protein OS=Homo sapiens GN=TSPO PE=1 SV=2
UBC9_HUMAN	31	18223	3 (1)	SUMO-conjugating enzyme UBC9 OS=Homo sapiens GN=UBE2I PE=1 SV=1
NTAL_HUMAN	30	26819	4 (1)	Linker for activation of T-cells family member 2 OS=Homo sapiens GN=LAT2 PE=1 SV=1
RPN2_HUMAN	30	69355	2 (1)	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo sapiens GN=RPN2 PE=1 SV=3
RL21_HUMAN	30	18610	3 (1)	60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2
B2MG_HUMAN	30	13820	3 (1)	Beta-2-microglobulin OS=Homo sapiens GN=B2M PE=1 SV=1
CS010_HUMAN	30	18897	1 (1)	UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1
PLP2_HUMAN	29	17022	1 (1)	Proteolipid protein 2 OS=Homo sapiens GN=PLP2 PE=1 SV=1
IL25_HUMAN	29	20887	1 (1)	Interleukin-25 OS=Homo sapiens GN=IL25 PE=1 SV=1
TOM20_HUMAN	29	16459	4 (1)	Mitochondrial import receptor subunit TOM20 homolog OS=Homo sapiens GN=TOMM20 PE=1 SV=1
DHSB_HUMAN	29	32407	3 (1)	Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Homo sapiens GN=SDHB PE=1 SV=3
G45IP_HUMAN	28	25425	4 (1)	Growth arrest and DNA damage-inducible proteins-interacting protein 1 OS=Homo sapiens GN=GADD45GIP1 PE=1 SV=
HNRPM_HUMAN	28	77749	8 (2)	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3
NDUA6_HUMAN	28	15127	1 (1)	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=Homo sapiens GN=NDUFA6 PE=1 SV=2
PLEK_HUMAN	28	40471	1 (1)	Pleckstrin OS=Homo sapiens GN=PLEK PE=1 SV=2

ARF6_HUMAN	28	20183	3 (1)	ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2
SC22B_HUMAN	28	24896	1 (1)	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=3
NDUS6_HUMAN	27	14045	1 (1)	NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Homo sapiens GN=NDUFS6 PE=1 SV=1
CK059_HUMAN	26	17848	1 (1)	RhoA activator C11orf59 OS=Homo sapiens GN=C11orf59 PE=1 SV=2
NDUC2_HUMAN	25	14235	2 (1)	NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Homo sapiens GN=NDUFC2 PE=1 SV=1
THIL_HUMAN	25	45456	4 (1)	Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1
CISY_HUMAN	25	51908	2 (1)	Citrate synthase, mitochondrial OS=Homo sapiens GN=CS PE=1 SV=2
RS9_HUMAN	24	22635	7 (1)	40S ribosomal protein S9 OS=Homo sapiens GN=RPS9 PE=1 SV=3
UCRIL_HUMAN	24	31081	2 (1)	Cytochrome b-c1 complex subunit Rieske-like protein 1 OS=Homo sapiens GN=UQCRFSL1 PE=2 SV=1
RT23_HUMAN	24	21814	2 (1)	28S ribosomal protein S23, mitochondrial OS=Homo sapiens GN=MRPS23 PE=1 SV=2
CLIC4_HUMAN	23	28982	2 (1)	Chloride intracellular channel protein 4 OS=Homo sapiens GN=CLIC4 PE=1 SV=4
AT5F1_HUMAN	23	28947	5 (2)	ATP synthase subunit b, mitochondrial OS=Homo sapiens GN=ATP5F1 PE=1 SV=2
SAP_HUMAN	23	59899	2 (1)	Proactivator polypeptide OS=Homo sapiens GN=PSAP PE=1 SV=2
HDGF_HUMAN	22	26886	1 (1)	Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1
NUCB2_HUMAN	22	50278	1 (1)	Nucleobindin-2 OS=Homo sapiens GN=NUCB2 PE=1 SV=2
LASP1_HUMAN	22	30097	3 (1)	LIM and SH3 domain protein 1 OS=Homo sapiens GN=LASP1 PE=1 SV=2
RBM3_HUMAN	22	17160	2 (1)	Putative RNA-binding protein 3 OS=Homo sapiens GN=RBM3 PE=1 SV=1
FIS1_HUMAN	22	16984	1 (1)	Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2
PLCA_HUMAN	22	32038	2 (1)	1-acyl-sn-glycerol-3-phosphate acyltransferase alpha OS=Homo sapiens GN=AGPAT1 PE=2 SV=2
PSA2_HUMAN	22	25996	2 (1)	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2
FRIL_HUMAN	21	20064	1 (1)	Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2
PHB2_HUMAN	21	33276	1 (1)	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2
SRPR_HUMAN	21	70223	2 (1)	Signal recognition particle receptor subunit alpha OS=Homo sapiens GN=SRPR PE=1 SV=2
PTMA_HUMAN	20	12196	1 (1)	Prothymosin alpha OS=Homo sapiens GN=PTMA PE=1 SV=2
CYB5_HUMAN	20	15321	1 (1)	Cytochrome b5 OS=Homo sapiens GN=CYB5A PE=1 SV=2
CN166_HUMAN	20	28165	3 (1)	UPF0568 protein C14orf166 OS=Homo sapiens GN=C14orf166 PE=1 SV=1
GRB2_HUMAN	20	25304	1 (1)	Growth factor receptor-bound protein 2 OS=Homo sapiens GN=GRB2 PE=1 SV=1
PAIRB_HUMAN	20	44995	2 (1)	Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 PE=1 SV=2
PSA3_HUMAN	20	28643	1 (1)	Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 PE=1 SV=2
NDUB6_HUMAN	19	15479	3 (1)	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 6 OS=Homo sapiens GN=NDUFB6 PE=1 SV=3
COPD_HUMAN	19	57630	2 (1)	Coatomer subunit delta OS=Homo sapiens GN=ARCN1 PE=1 SV=1
SCO1_HUMAN	19	34021	2 (1)	Protein SCO1 homolog, mitochondrial OS=Homo sapiens GN=SCO1 PE=1 SV=1
C2C2L_HUMAN	19	76590	1 (1)	C2 domain-containing protein 2-like OS=Homo sapiens GN=C2CD2L PE=1 SV=3
PNPH_HUMAN	19	32325	3 (1)	Purine nucleoside phosphorylase OS=Homo sapiens GN=NP PE=1 SV=2
FBRL_HUMAN	18	33877	2 (1)	rRNA 2'-O-methyltransferase fibrillarin OS=Homo sapiens GN=FBL PE=1 SV=2
RM11_HUMAN	18	20727	2 (1)	39S ribosomal protein L11, mitochondrial OS=Homo sapiens GN=MRPL11 PE=1 SV=1
PDCD5_HUMAN	18	14276	2 (1)	Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3
NFU1_HUMAN	18	28615	4 (1)	NFU1 iron-sulfur cluster scaffold homolog, mitochondrial OS=Homo sapiens GN=NFU1 PE=1 SV=2
CAP7_HUMAN	18	27325	1 (1)	Azurocidin OS=Homo sapiens GN=AZU1 PE=1 SV=3
RNH2C_HUMAN	17	17943	3 (1)	Ribonuclease H2 subunit C OS=Homo sapiens GN=RNASEH2C PE=1 SV=1
D3D2_HUMAN	17	33080	1 (1)	3,2-trans-enoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=DCI PE=1 SV=1
LONF1_HUMAN	15	88494	3 (1)	LON peptidase N-terminal domain and RING finger protein 1 OS=Homo sapiens GN=LONRF1 PE=2 SV=2

EMAL4_HUMAN	14	109919	3 (1)	Echinoderm microtubule-associated protein-like 4 OS=Homo sapiens GN=EML4 PE=1 SV=2