

Supplemental Table 2A. Proteins detected in the membrane fraction of Human RBCs.

Accession	Description
P31946	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3 - [1433B_HUMAN]
P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1 - [1433E_HUMAN]
Q04917	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4 - [1433F_HUMAN]
P61981	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2 - [1433G_HUMAN]
P27348	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1 - [1433T_HUMAN]
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1 - [1433Z_HUMAN]
Q9NUQ2	1-acyl-sn-glycerol-3-phosphate acyltransferase epsilon OS=Homo sapiens GN=AGPAT5 PE=1 SV=3 - [PLCE_HUMAN]
Q4AC99	1-aminocyclopropane-1-carboxylate synthase-like protein 2 OS=Homo sapiens GN=ACCSL PE=2 SV=3 - [1A1L2_HUMAN]
P09543	2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Homo sapiens GN=CNP PE=1 SV=2 - [CN37_HUMAN]
P62333	26S protease regulatory subunit 10B OS=Homo sapiens GN=PSMC6 PE=1 SV=1 - [PRS10_HUMAN]
P62191	26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1 - [PRS4_HUMAN]
P17980	26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=1 SV=3 - [PRS6A_HUMAN]
P43686	26S protease regulatory subunit 6B OS=Homo sapiens GN=PSMC4 PE=1 SV=2 - [PRS6B_HUMAN]
P35998	26S protease regulatory subunit 7 OS=Homo sapiens GN=PSMC2 PE=1 SV=3 - [PRS7_HUMAN]
Q99460	26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1 PE=1 SV=2 - [PSMD1_HUMAN]
O00231	26S proteasome non-ATPase regulatory subunit 11 OS=Homo sapiens GN=PSMD11 PE=1 SV=3 - [PSD11_HUMAN]
O00232	26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV=3 - [PSD12_HUMAN]
Q9UNM6	26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV=2 - [PSD13_HUMAN]
O00487	26S proteasome non-ATPase regulatory subunit 14 OS=Homo sapiens GN=PSMD14 PE=1 SV=1 - [PSDE_HUMAN]
P55036	26S proteasome non-ATPase regulatory subunit 4 OS=Homo sapiens GN=PSMD4 PE=1 SV=1 - [PSMD4_HUMAN]
Q15008	26S proteasome non-ATPase regulatory subunit 6 OS=Homo sapiens GN=PSMD6 PE=1 SV=1 - [PSMD6_HUMAN]
P48556	26S proteasome non-ATPase regulatory subunit 8 OS=Homo sapiens GN=PSMD8 PE=1 SV=2 - [PSMD8_HUMAN]
Q13442	28 kDa heat- and acid-stable phosphoprotein OS=Homo sapiens GN=PDAP1 PE=1 SV=1 - [HAP28_HUMAN]
P82909	28S ribosomal protein S36, mitochondrial OS=Homo sapiens GN=MRPS36 PE=1 SV=2 - [RT36_HUMAN]
P82933	28S ribosomal protein S9, mitochondrial OS=Homo sapiens GN=MRPS9 PE=1 SV=2 - [RT09_HUMAN]
Q96RE3	36 kDa muscleblind protein EXP36 OS=Homo sapiens GN=MBNL1 PE=2 SV=1 - [Q96RE3_HUMAN]
P52815	39S ribosomal protein L12, mitochondrial OS=Homo sapiens GN=MRPL12 PE=1 SV=2 - [RM12_HUMAN]
Q9P035	3-hydroxyacyl-CoA dehydratase 3 OS=Homo sapiens GN=PTPLAD1 PE=1 SV=2 - [HACD3_HUMAN]
Q99714	3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3 - [HCD2_HUMAN]
P42765	3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2 PE=1 SV=2 - [THIM_HUMAN]
P46783	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1 - [RS10_HUMAN]
P62280	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3 - [RS11_HUMAN]
P25398	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3 - [RS12_HUMAN]
P62277	40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2 - [RS13_HUMAN]
P62263	40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3 - [RS14_HUMAN]
P62841	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2 - [RS15_HUMAN]
P62244	40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2 - [RS15A_HUMAN]
P62249	40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2 - [RS16_HUMAN]
P0CW22	40S ribosomal protein S17-like OS=Homo sapiens GN=RPS17L PE=3 SV=1 - [RS17L_HUMAN]
P62269	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3 - [RS18_HUMAN]
P39019	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2 - [RS19_HUMAN]
P15880	40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2 - [RS2_HUMAN]
P60866	40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1 - [RS20_HUMAN]

Q8WVC2	40S ribosomal protein S21 OS=Homo sapiens GN=RPS21 PE=2 SV=1 - [Q8WVC2_HUMAN]
P62266	40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=1 SV=3 - [RS23_HUMAN]
Q5TOP8	40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=2 SV=1 - [Q5TOP8_HUMAN]
P62851	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1 - [RS25_HUMAN]
P62854	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3 - [RS26_HUMAN]
P42677	40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3 - [RS27_HUMAN]
P62857	40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1 - [RS28_HUMAN]
P62273	40S ribosomal protein S29 OS=Homo sapiens GN=RPS29 PE=1 SV=2 - [RS29_HUMAN]
P23396	40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2 - [RS3_HUMAN]
P62861	40S ribosomal protein S30 OS=Homo sapiens GN=FAU PE=1 SV=1 - [RS30_HUMAN]
P61247	40S ribosomal protein S3a OS=Homo sapiens GN=RPS3A PE=1 SV=2 - [RS3A_HUMAN]
P62701	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2 - [RS4X_HUMAN]
P46782	40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4 - [RS5_HUMAN]
P62753	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1 - [RS6_HUMAN]
P62081	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1 - [RS7_HUMAN]
P46781	40S ribosomal protein S9 OS=Homo sapiens GN=RPS9 PE=1 SV=3 - [RS9_HUMAN]
P49189	4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3 - [AL9A1_HUMAN]
Q8IZH2	5'-3' exoribonuclease 1 OS=Homo sapiens GN=XRN1 PE=1 SV=1 - [XRN1_HUMAN]
Q13131	5'-AMP-activated protein kinase catalytic subunit alpha-1 OS=Homo sapiens GN=PRKAA1 PE=1 SV=4 - [AAPK1_HUMAN]
P54619	5'-AMP-activated protein kinase subunit gamma-1 OS=Homo sapiens GN=PRKAG1 PE=1 SV=1 - [AAKG1_HUMAN]
Q86UY8	5'-nucleotidase domain-containing protein 3 OS=Homo sapiens GN=NT5DC3 PE=2 SV=1 - [NT5D3_HUMAN]
P10809	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2 - [CH60_HUMAN]
P05387	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1 - [RLA2_HUMAN]
P27635	60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=1 SV=4 - [RL10_HUMAN]
P62906	60S ribosomal protein L10a OS=Homo sapiens GN=RPL10A PE=1 SV=2 - [RL10A_HUMAN]
P30050	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1 - [RL12_HUMAN]
P26373	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4 - [RL13_HUMAN]
P40429	60S ribosomal protein L13a OS=Homo sapiens GN=RPL13A PE=1 SV=2 - [RL13A_HUMAN]
P50914	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4 - [RL14_HUMAN]
P18621	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3 - [RL17_HUMAN]
Q07020	60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2 - [RL18_HUMAN]
P84098	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1 - [RL19_HUMAN]
P46778	60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2 - [RL21_HUMAN]
P35268	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2 - [RL22_HUMAN]
P62829	60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1 - [RL23_HUMAN]
P62750	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1 - [RL23A_HUMAN]
P61254	60S ribosomal protein L26 OS=Homo sapiens GN=RPL26 PE=1 SV=1 - [RL26_HUMAN]
P61353	60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2 - [RL27_HUMAN]
P46779	60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=1 SV=3 - [RL28_HUMAN]
P47914	60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2 - [RL29_HUMAN]
P39023	60S ribosomal protein L3 OS=Homo sapiens GN=RPL3 PE=1 SV=2 - [RL3_HUMAN]
P62888	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2 - [RL30_HUMAN]
P49207	60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3 - [RL34_HUMAN]

P42766	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2 - [RL35_HUMAN]
Q9Y3U8	60S ribosomal protein L36 OS=Homo sapiens GN=RPL36 PE=1 SV=3 - [RL36_HUMAN]
P83881	60S ribosomal protein L36a OS=Homo sapiens GN=RPL36A PE=1 SV=2 - [RL36A_HUMAN]
P63173	60S ribosomal protein L38 OS=Homo sapiens GN=RPL38 PE=1 SV=2 - [RL38_HUMAN]
P36578	60S ribosomal protein L4 OS=Homo sapiens GN=RPL4 PE=1 SV=5 - [RL4_HUMAN]
Q02878	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3 - [RL6_HUMAN]
P18124	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1 - [RL7_HUMAN]
P62424	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2 - [RL7A_HUMAN]
P17858	6-phosphofructokinase, liver type OS=Homo sapiens GN=PFKL PE=1 SV=6 - [K6PL_HUMAN]
P08237	6-phosphofructokinase, muscle type OS=Homo sapiens GN=PFKM PE=1 SV=2 - [K6PF_HUMAN]
F5H7U0	6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=3 SV=1 - [F5H7U0_HUMAN]
P11021	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]
D2K8Q1	AAA domain containing 3A protein OS=Homo sapiens GN=ATAD3A PE=2 SV=1 - [D2K8Q1_HUMAN]
Q2L6I2	ABC50 protein OS=Homo sapiens GN=ABCF1 PE=2 SV=1 - [Q2L6I2_HUMAN]
P24752	Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1 - [THIL_HUMAN]
Q92688	Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Homo sapiens GN=ANP32B PE=1 SV=1 - [AN32B_HUMAN]
Q5VZA6	Aconitase 1, soluble OS=Homo sapiens GN=ACO1 PE=2 SV=1 - [Q5VZA6_HUMAN]
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 - [ACTB_HUMAN]
P63267	Actin, gamma-enteric smooth muscle OS=Homo sapiens GN=ACTG2 PE=1 SV=1 - [ACTH_HUMAN]
P61160	Actin-related protein 2 OS=Homo sapiens GN=ACTR2 PE=1 SV=1 - [ARP2_HUMAN]
O15143	Actin-related protein 2/3 complex subunit 1B OS=Homo sapiens GN=ARPC1B PE=1 SV=3 - [ARC1B_HUMAN]
O15144	Actin-related protein 2/3 complex subunit 2 OS=Homo sapiens GN=ARPC2 PE=1 SV=1 - [ARPC2_HUMAN]
P59998	Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3 - [ARPC4_HUMAN]
B1ALC0	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=3 SV=1 - [B1ALC0_HUMAN]
Q9BPX5	Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5L PE=1 SV=1 - [ARP5L_HUMAN]
Q9H9F9	Actin-related protein 5 OS=Homo sapiens GN=ACTR5 PE=1 SV=2 - [ARP5_HUMAN]
Q7LC44	Activity-regulated cytoskeleton-associated protein OS=Homo sapiens GN=ARC PE=1 SV=1 - [ARC_HUMAN]
O14561	Acyl carrier protein, mitochondrial OS=Homo sapiens GN=NDUFAB1 PE=1 SV=3 - [ACPM_HUMAN]
P13798	Acylamino-acid-releasing enzyme OS=Homo sapiens GN=APEH PE=1 SV=4 - [ACPH_HUMAN]
Q9H845	Acyl-CoA dehydrogenase family member 9, mitochondrial OS=Homo sapiens GN=ACAD9 PE=1 SV=1 - [ACAD9_HUMAN]
Q5JWV8	Acyl-CoA synthetase long-chain family member 4 OS=Homo sapiens GN=ACSL4 PE=2 SV=1 - [Q5JWV8_HUMAN]
Q6E213	Acyl-CoA wax alcohol acyltransferase 2 OS=Homo sapiens GN=AWAT2 PE=1 SV=1 - [AWAT2_HUMAN]
Q92604	Acyl-CoA:lysophosphatidylglycerol acyltransferase 1 OS=Homo sapiens GN=LPGAT1 PE=2 SV=1 - [LGAT1_HUMAN]
P82987	ADAMTS-like protein 3 OS=Homo sapiens GN=ADAMTSL3 PE=2 SV=4 - [ATL3_HUMAN]
Q86XM2	ADD1 protein OS=Homo sapiens GN=ADD1 PE=2 SV=1 - [Q86XM2_HUMAN]
A2A3N8	Adducin 1 (Alpha) OS=Homo sapiens GN=ADD1 PE=4 SV=1 - [A2A3N8_HUMAN]
F5H737	Adenosylhomocysteinase OS=Homo sapiens GN=AHCY PE=3 SV=1 - [F5H737_HUMAN]
P00568	Adenylate kinase isoenzyme 1 OS=Homo sapiens GN=AK1 PE=1 SV=3 - [KAD1_HUMAN]
P30566	Adenylosuccinate lyase OS=Homo sapiens GN=ADSL PE=1 SV=2 - [PUR8_HUMAN]
F5H6Z3	Adenylyl cyclase-associated protein OS=Homo sapiens GN=CAP1 PE=3 SV=1 - [F5H6Z3_HUMAN]
Q9HDC9	Adipocyte plasma membrane-associated protein OS=Homo sapiens GN=APMAP PE=1 SV=2 - [APMAP_HUMAN]
P05141	ADP/ATP translocase 2 OS=Homo sapiens GN=SLC25A5 PE=1 SV=7 - [ADT2_HUMAN]
P12236	ADP/ATP translocase 3 OS=Homo sapiens GN=SLC25A6 PE=1 SV=4 - [ADT3_HUMAN]

P62330	ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2 - [ARF6_HUMAN]
Q9NVJ2	ADP-ribosylation factor-like protein 8B OS=Homo sapiens GN=ARL8B PE=1 SV=1 - [ARL8B_HUMAN]
Q9Y4W6	AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2 - [AFG32_HUMAN]
P49588	Alanyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2 - [SYAC_HUMAN]
Q8IZ83	Aldehyde dehydrogenase family 16 member A1 OS=Homo sapiens GN=ALDH16A1 PE=1 SV=2 - [A16A1_HUMAN]
Q9H6U8	Alpha-1,2-mannosyltransferase ALG9 OS=Homo sapiens GN=ALG9 PE=1 SV=2 - [ALG9_HUMAN]
Q9H553	Alpha-1,3/1,6-mannosyltransferase ALG2 OS=Homo sapiens GN=ALG2 PE=2 SV=1 - [ALG2_HUMAN]
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3 - [A2MG_HUMAN]
P61163	Alpha-centractin OS=Homo sapiens GN=ACTR1A PE=1 SV=1 - [ACTZ_HUMAN]
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2 - [ENOA_HUMAN]
Q9NZD4	Alpha-hemoglobin-stabilizing protein OS=Homo sapiens GN=AHSP PE=1 SV=1 - [AHSP_HUMAN]
P54920	Alpha-soluble NSF attachment protein OS=Homo sapiens GN=NAPA PE=1 SV=3 - [SNAA_HUMAN]
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sapiens GN=AIMP1 PE=1 SV=2 - [AIMP1_HUMAN]
A2RU49	Aminoglycoside phosphotransferase domain-containing protein 1 OS=Homo sapiens GN=AGPHD1 PE=2 SV=2 - [AGPD1_HUMAN]
A8K6C4	Aminolevulinate, delta-, synthase 2 OS=Homo sapiens GN=ALAS2 PE=2 SV=1 - [A8K6C4_HUMAN]
P04920	Anion exchange protein 2 OS=Homo sapiens GN=SLC4A2 PE=1 SV=4 - [B3A2_HUMAN]
Q6PK32	ANK1 protein OS=Homo sapiens GN=ANK1 PE=2 SV=1 - [Q6PK32_HUMAN]
Q53ER1	Ankyrin 1 isoform 5 variant (Fragment) OS=Homo sapiens GN=ANK1 PE=2 SV=1 - [Q53ER1_HUMAN]
Q5T457	Ankyrin repeat domain 2 (Stretch responsive muscle) OS=Homo sapiens GN=ANKRD2 PE=2 SV=1 - [Q5T457_HUMAN]
Q6UB99	Ankyrin repeat domain-containing protein 11 OS=Homo sapiens GN=ANKRD11 PE=1 SV=3 - [ANR11_HUMAN]
Q4UJ75	Ankyrin repeat domain-containing protein 20A4 OS=Homo sapiens GN=ANKRD20A4 PE=2 SV=1 - [A20A4_HUMAN]
Q96NW4	Ankyrin repeat domain-containing protein 27 OS=Homo sapiens GN=ANKRD27 PE=1 SV=2 - [ANR27_HUMAN]
Q9BQI6	Ankyrin repeat domain-containing protein 32 OS=Homo sapiens GN=ANKRD32 PE=1 SV=2 - [ANR32_HUMAN]
Q12955	Ankyrin-3 OS=Homo sapiens GN=ANK3 PE=1 SV=3 - [ANK3_HUMAN]
Q10567	AP-1 complex subunit beta-1 OS=Homo sapiens GN=AP1B1 PE=1 SV=2 - [AP1B1_HUMAN]
O43747	AP-1 complex subunit gamma-1 OS=Homo sapiens GN=AP1G1 PE=1 SV=5 - [AP1G1_HUMAN]
O95782	AP-2 complex subunit alpha-1 OS=Homo sapiens GN=AP2A1 PE=1 SV=3 - [AP2A1_HUMAN]
P63010	AP-2 complex subunit beta OS=Homo sapiens GN=AP2B1 PE=1 SV=1 - [AP2B1_HUMAN]
P53680	AP-2 complex subunit sigma OS=Homo sapiens GN=AP2S1 PE=1 SV=2 - [AP2S1_HUMAN]
Q2M2I8	AP2-associated protein kinase 1 OS=Homo sapiens GN=AAK1 PE=1 SV=3 - [AAK1_HUMAN]
O95236	Apolipoprotein L3 OS=Homo sapiens GN=APOL3 PE=2 SV=3 - [APOL3_HUMAN]
O95831	Apoptosis-inducing factor 1, mitochondrial OS=Homo sapiens GN=AIFM1 PE=1 SV=1 - [AIFM1_HUMAN]
P18054	Arachidonate 12-lipoxygenase, 12S-type OS=Homo sapiens GN=ALOX12 PE=1 SV=4 - [LOX12_HUMAN]
P52594	Arf-GAP domain and FG repeats-containing protein 1 OS=Homo sapiens GN=AGFG1 PE=1 SV=2 - [AGFG1_HUMAN]
O95081	Arf-GAP domain and FG repeats-containing protein 2 OS=Homo sapiens GN=AGFG2 PE=1 SV=2 - [AGFG2_HUMAN]
Q15057	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing protein 2 OS=Homo sapiens GN=ACAP2 PE=1 SV=3 - [ACAP2_HUMAN]
Q99490	Arf-GAP with GTPase, ANK repeat and PH domain-containing protein 2 OS=Homo sapiens GN=AGAP2 PE=1 SV=2 - [AGAP2_HUMAN]
Q9ULH1	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 1 OS=Homo sapiens GN=ASAP1 PE=1 SV=4 - [ASAP1_HUMAN]
P05089	Arginase-1 OS=Homo sapiens GN=ARG1 PE=1 SV=2 - [ARGI1_HUMAN]
P78540	Arginase-2, mitochondrial OS=Homo sapiens GN=ARG2 PE=1 SV=1 - [ARGI2_HUMAN]
B7Z637	Armadillo repeat containing 8, isoform CRA_g OS=Homo sapiens GN=ARMC8 PE=2 SV=1 - [B7Z637_HUMAN]
O43776	Asparaginyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=NARS PE=1 SV=1 - [SYNC_HUMAN]
P00505	Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3 - [AATM_HUMAN]

P14868	Aspartyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2 - [SYDC_HUMAN]
Q6PI48	Aspartyl-tRNA synthetase, mitochondrial OS=Homo sapiens GN=DARS2 PE=1 SV=1 - [SYDM_HUMAN]
O43313	ATM interactor OS=Homo sapiens GN=ATMIN PE=1 SV=2 - [ATMIN_HUMAN]
B4DL14	ATP synthase gamma chain OS=Homo sapiens GN=ATP5C1 PE=2 SV=1 - [B4DL14_HUMAN]
P25705	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1 - [ATPA_HUMAN]
P06576	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3 - [ATPB_HUMAN]
O75947	ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3 - [ATP5H_HUMAN]
P30049	ATP synthase subunit delta, mitochondrial OS=Homo sapiens GN=ATP5D PE=1 SV=2 - [ATPD_HUMAN]
P56385	ATP synthase subunit e, mitochondrial OS=Homo sapiens GN=ATP5I PE=1 SV=2 - [ATP5I_HUMAN]
P56381	ATP synthase subunit epsilon, mitochondrial OS=Homo sapiens GN=ATP5E PE=1 SV=2 - [ATP5E_HUMAN]
P48047	ATP synthase subunit O, mitochondrial OS=Homo sapiens GN=ATP5O PE=1 SV=1 - [ATPO_HUMAN]
Q5QNZ2	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit B1 OS=Homo sapiens GN=ATP5F1 PE=2 SV=1 - [Q5QNZ2_HUMAN]
P18859	ATP synthase-coupling factor 6, mitochondrial OS=Homo sapiens GN=ATP5J PE=1 SV=1 - [ATP5J_HUMAN]
Q96QE3	ATPase family AAA domain-containing protein 5 OS=Homo sapiens GN=ATAD5 PE=1 SV=4 - [ATAD5_HUMAN]
Q9UII2	ATPase inhibitor, mitochondrial OS=Homo sapiens GN=ATPIF1 PE=1 SV=1 - [ATIF1_HUMAN]
B1AKY9	ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide OS=Homo sapiens GN=ATP1A2 PE=3 SV=1 - [B1AKY9_HUMAN]
Q86UK0	ATP-binding cassette sub-family A member 12 OS=Homo sapiens GN=ABCA12 PE=1 SV=3 - [ABCAC_HUMAN]
O94911	ATP-binding cassette sub-family A member 8 OS=Homo sapiens GN=ABCA8 PE=1 SV=3 - [ABCA8_HUMAN]
Q9NRK6	ATP-binding cassette sub-family B member 10, mitochondrial OS=Homo sapiens GN=ABCB10 PE=1 SV=2 - [ABCBA_HUMAN]
Q9NP58	ATP-binding cassette sub-family B member 6, mitochondrial OS=Homo sapiens GN=ABCB6 PE=1 SV=1 - [ABCB6_HUMAN]
P61221	ATP-binding cassette sub-family E member 1 OS=Homo sapiens GN=ABCE1 PE=1 SV=1 - [ABCE1_HUMAN]
Q5STZ8	ATP-binding cassette sub-family F (GCN20) member 1 (Fragment) OS=Homo sapiens GN=ABCF1 PE=2 SV=2 - [Q5STZ8_HUMAN]
Q9UNQ0	ATP-binding cassette sub-family G member 2 OS=Homo sapiens GN=ABCG2 PE=1 SV=3 - [ABCG2_HUMAN]
A4D1D5	ATP-binding cassette, sub-family B (MDR/TAP), member 4 OS=Homo sapiens GN=ABCB4 PE=3 SV=1 - [A4D1D5_HUMAN]
P53396	ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3 - [ACLY_HUMAN]
Q08211	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4 - [DHX9_HUMAN]
Q7Z478	ATP-dependent RNA helicase DHX29 OS=Homo sapiens GN=DHX29 PE=1 SV=2 - [DHX29_HUMAN]
Q4LE39	AT-rich interactive domain-containing protein 4B OS=Homo sapiens GN=ARID4B PE=1 SV=2 - [ARI4B_HUMAN]
P02730	Band 3 anion transport protein OS=Homo sapiens GN=SLC4A1 PE=1 SV=3 - [B3AT_HUMAN]
P50895	Basal cell adhesion molecule OS=Homo sapiens GN=BCAM PE=1 SV=2 - [BCAM_HUMAN]
Q54A51	Basigin (OK blood group), isoform CRA_a OS=Homo sapiens GN=hEMMPRIN PE=2 SV=1 - [Q54A51_HUMAN]
P51572	B-cell receptor-associated protein 31 OS=Homo sapiens GN=BCAP31 PE=1 SV=3 - [BAP31_HUMAN]
Q5TE63	BCL2-like 1 OS=Homo sapiens GN=BCL2L1 PE=2 SV=1 - [Q5TE63_HUMAN]
P35612	Beta-adducin OS=Homo sapiens GN=ADD2 PE=1 SV=3 - [ADDB_HUMAN]
P07686	Beta-hexosaminidase subunit beta OS=Homo sapiens GN=HEXB PE=1 SV=3 - [HEXB_HUMAN]
A1A5D9	Bicaudal D-related protein 2 OS=Homo sapiens GN=CCDC64B PE=2 SV=2 - [BICR2_HUMAN]
P07814	Bifunctional aminoacyl-tRNA synthetase OS=Homo sapiens GN=EPRS PE=1 SV=5 - [SYEP_HUMAN]
P53004	Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2 - [BIEA_HUMAN]
P78537	Biogenesis of lysosome-related organelles complex 1 subunit 1 OS=Homo sapiens GN=BLOC1S1 PE=1 SV=2 - [BL1S1_HUMAN]
P07738	Bisphosphoglycerate mutase OS=Homo sapiens GN=BPGM PE=1 SV=2 - [PMGE_HUMAN]
Q13867	Bleomycin hydrolase OS=Homo sapiens GN=BLMH PE=1 SV=1 - [BLMH_HUMAN]
Q02161	Blood group Rh(D) polypeptide OS=Homo sapiens GN=RHD PE=1 SV=3 - [RHD_HUMAN]
Q9NSY1	BMP-2-inducible protein kinase OS=Homo sapiens GN=BMP2K PE=1 SV=2 - [BMP2K_HUMAN]

Q8TDC3	BR serine/threonine-protein kinase 1 OS=Homo sapiens GN=BRSK1 PE=1 SV=1 - [BRSK1_HUMAN]
Q5R3B4	Brain protein 44 (Fragment) OS=Homo sapiens GN=BRP44 PE=2 SV=1 - [Q5R3B4_HUMAN]
Q9NWW8	BRISC and BRCA1-A complex member 1 OS=Homo sapiens GN=BABAM1 PE=1 SV=1 - [BABA1_HUMAN]
Q15018	BRISC complex subunit Abro1 OS=Homo sapiens GN=FAM175B PE=1 SV=2 - [F175B_HUMAN]
Q14681	BTB/POZ domain-containing protein KCTD2 OS=Homo sapiens GN=KCTD2 PE=1 SV=3 - [KCTD2_HUMAN]
P11586	C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3 - [C1TC_HUMAN]
O14523	C2 domain-containing protein 2-like OS=Homo sapiens GN=C2CD2L PE=1 SV=3 - [C2C2L_HUMAN]
A6H8M8	C4A protein OS=Homo sapiens GN=C4A PE=2 SV=1 - [A6H8M8_HUMAN]
P27708	CAD protein OS=Homo sapiens GN=CAD PE=1 SV=3 - [PYR1_HUMAN]
Q99828	Calcium and integrin-binding protein 1 OS=Homo sapiens GN=CIB1 PE=1 SV=4 - [CIB1_HUMAN]
Q9Y376	Calcium-binding protein 39 OS=Homo sapiens GN=CAB39 PE=1 SV=1 - [CAB39_HUMAN]
B2ZWH1	Calcyclin binding protein (Fragment) OS=Homo sapiens GN=CACYBP PE=4 SV=1 - [B2ZWH1_HUMAN]
P62158	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2 - [CALM_HUMAN]
Q5T5Y3	Calmodulin-regulated spectrin-associated protein 1 OS=Homo sapiens GN=CAMSAP1 PE=1 SV=2 - [CAMP1_HUMAN]
P27824	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2 - [CALX_HUMAN]
P04632	Calpain small subunit 1 OS=Homo sapiens GN=CAPNS1 PE=1 SV=1 - [CPNS1_HUMAN]
P07384	Calpain-1 catalytic subunit OS=Homo sapiens GN=CAPN1 PE=1 SV=1 - [CAN1_HUMAN]
O15484	Calpain-5 OS=Homo sapiens GN=CAPN5 PE=1 SV=2 - [CAN5_HUMAN]
P27797	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1 - [CALR_HUMAN]
O43852	Calumenin OS=Homo sapiens GN=CALU PE=1 SV=2 - [CALU_HUMAN]
P17612	cAMP-dependent protein kinase catalytic subunit alpha OS=Homo sapiens GN=PRKACA PE=1 SV=2 - [KAPCA_HUMAN]
P10644	cAMP-dependent protein kinase type I-alpha regulatory subunit OS=Homo sapiens GN=PRKAR1A PE=1 SV=1 - [KAPO_HUMAN]
B1AK87	Capping protein (Actin filament) muscle Z-line, beta OS=Homo sapiens GN=CAPZB PE=4 SV=1 - [B1AK87_HUMAN]
D3YHP0	CAR4/6 OS=Homo sapiens PE=2 SV=1 - [D3YHP0_HUMAN]
Q7L1S5	Carbohydrate sulfotransferase 9 OS=Homo sapiens GN=CHST9 PE=2 SV=2 - [CHST9_HUMAN]
P00918	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2 - [CAH2_HUMAN]
A3EY21	Cardiac sodium channel alpha subunit (Fragment) OS=Homo sapiens GN=SCN5A PE=2 SV=1 - [A3EY21_HUMAN]
P50416	Carnitine O-palmitoyltransferase 1, liver isoform OS=Homo sapiens GN=CPT1A PE=1 SV=2 - [CPT1A_HUMAN]
P48729	Casein kinase I isoform alpha OS=Homo sapiens GN=CSNK1A1 PE=1 SV=2 - [KC1A_HUMAN]
P67870	Casein kinase II subunit beta OS=Homo sapiens GN=CSNK2B PE=1 SV=1 - [CSK2B_HUMAN]
P04040	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3 - [CATA_HUMAN]
P21964	Catechol O-methyltransferase OS=Homo sapiens GN=COMT PE=1 SV=2 - [COMT_HUMAN]
Q5JZG9	Cathepsin A (Fragment) OS=Homo sapiens GN=CTSA PE=2 SV=1 - [Q5JZG9_HUMAN]
P07339	Cathepsin D OS=Homo sapiens GN=CTSD PE=1 SV=1 - [CATD_HUMAN]
P14091	Cathepsin E OS=Homo sapiens GN=CTSE PE=1 SV=2 - [CATE_HUMAN]
O95628	CCR4-NOT transcription complex subunit 4 OS=Homo sapiens GN=CNOT4 PE=1 SV=3 - [CNOT4_HUMAN]
Q9Y5K6	CD2-associated protein OS=Homo sapiens GN=CD2AP PE=1 SV=1 - [CD2AP_HUMAN]
B6EAT9	CD44 OS=Homo sapiens PE=2 SV=1 - [B6EAT9_HUMAN]
B1AMW1	CD58 antigen, (Lymphocyte function-associated antigen 3), isoform CRA_c OS=Homo sapiens GN=CD58 PE=4 SV=1 - [B1AMW1_HUMAN]
A6NIW1	CD99 antigen, isoform CRA_b OS=Homo sapiens GN=CD99 PE=4 SV=1 - [A6NIW1_HUMAN]
Q8N5K1	CDGSH iron-sulfur domain-containing protein 2 OS=Homo sapiens GN=CISD2 PE=1 SV=1 - [CISD2_HUMAN]
B3KW52	cDNA FLJ42145 fis, clone TESTI4000228, highly similar to Mus musculus ubiquitin family domain containing 1 (Ubf1), mRNA OS=Homo sapiens PE=2 SV=1 - [B3KW52_HUMAN]
B3KXX3	cDNA FLJ46254 fis, clone TESTI4023096, highly similar to Solute carrier family 12 member 6 OS=Homo sapiens PE=2 SV=1 - [B3KXX3_HUMAN]

B7Z3E2	cDNA FLJ50777, highly similar to Serine/threonine-protein phosphatase 6 (EC 3.1.3.16) OS=Homo sapiens PE=2 SV=1 - [B7Z3E2_HUMAN]
B4DM63	cDNA FLJ51245, highly similar to Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens PE=2 SV=1 - [B4DM63_HUMAN]
B4DEW9	cDNA FLJ52478, highly similar to Eukaryotic translation initiation factor 3 subunit 5 OS=Homo sapiens PE=2 SV=1 - [B4DEW9_HUMAN]
B7Z3Q0	cDNA FLJ52550 OS=Homo sapiens PE=2 SV=1 - [B7Z3Q0_HUMAN]
B4DS57	cDNA FLJ53176, highly similar to Nuclear autoantigenic sperm protein OS=Homo sapiens PE=2 SV=1 - [B4DS57_HUMAN]
B4DPJ6	cDNA FLJ53635, highly similar to Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 6, mRNA OS=Homo sapiens PE=2 SV=1 - [B4DPJ6_HUMAN]
B4DW31	cDNA FLJ54186, highly similar to Nuclear receptor-binding protein OS=Homo sapiens PE=2 SV=1 - [B4DW31_HUMAN]
B7Z934	cDNA FLJ55302, highly similar to Transcription factor COE2 OS=Homo sapiens PE=2 SV=1 - [B7Z934_HUMAN]
B7Z4B8	cDNA FLJ56481, highly similar to Heterogeneous nuclear ribonucleoprotein U-like protein 1 OS=Homo sapiens PE=2 SV=1 - [B7Z4B8_HUMAN]
B4DLT2	cDNA FLJ56637, highly similar to Nuclear pore complex protein Nup155 OS=Homo sapiens PE=2 SV=1 - [B4DLT2_HUMAN]
B4DK60	cDNA FLJ57515, highly similar to CDK5 and ABL1 enzyme substrate 1 OS=Homo sapiens PE=2 SV=1 - [B4DK60_HUMAN]
B4DZV5	cDNA FLJ58820, highly similar to 55 kDa erythrocyte membrane protein (p55) OS=Homo sapiens PE=2 SV=1 - [B4DZV5_HUMAN]
B4DH26	cDNA FLJ60517, highly similar to DCN1-like protein 4 OS=Homo sapiens PE=2 SV=1 - [B4DH26_HUMAN]
B4DN00	cDNA FLJ60715, highly similar to Pyruvate carboxylase, mitochondrial (EC 6.4.1.1) OS=Homo sapiens PE=2 SV=1 - [B4DN00_HUMAN]
B4DQQ8	cDNA FLJ60806, highly similar to RalBP1-associated Eps domain-containing protein 2 OS=Homo sapiens PE=2 SV=1 - [B4DQQ8_HUMAN]
B7Z1T7	cDNA FLJ61691, highly similar to Homo sapiens myosin head domain containing 1 (MYOHD1), transcript variant 2, mRNA OS=Homo sapiens PE=2 SV=1 - [B7Z1T7_HUMAN]
Q8NBU8	cDNA FLJ90737 fis, clone PLACE1010827, moderately similar to Membrane protein p24B OS=Homo sapiens PE=2 SV=1 - [Q8NBU8_HUMAN]
Q9NV96	Cell cycle control protein 50A OS=Homo sapiens GN=TMEM30A PE=1 SV=1 - [CC50A_HUMAN]
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2 - [CDC42_HUMAN]
Q99741	Cell division control protein 6 homolog OS=Homo sapiens GN=CDC6 PE=1 SV=1 - [CDC6_HUMAN]
Q9NX58	Cell growth-regulating nucleolar protein OS=Homo sapiens GN=LYAR PE=1 SV=2 - [LYAR_HUMAN]
P49454	Centromere protein F OS=Homo sapiens GN=CENPF PE=1 SV=2 - [CENPF_HUMAN]
Q02224	Centromere-associated protein E OS=Homo sapiens GN=CENPE PE=1 SV=2 - [CENPE_HUMAN]
Q5JVD6	Centrosomal protein 110kDa (Fragment) OS=Homo sapiens GN=CEP110 PE=2 SV=1 - [Q5JVD6_HUMAN]
O15078	Centrosomal protein of 290 kDa OS=Homo sapiens GN=CEP290 PE=1 SV=2 - [CE290_HUMAN]
B7ZAT2	Chaperonin containing TCP1, subunit 2 (Beta), isoform CRA_c OS=Homo sapiens GN=CCT2 PE=2 SV=1 - [B7ZAT2_HUMAN]
B7Z4T9	Chaperonin containing TCP1, subunit 7 (Eta), isoform CRA_c OS=Homo sapiens GN=CCT7 PE=2 SV=1 - [B7Z4T9_HUMAN]
Q9HD42	Charged multivesicular body protein 1a OS=Homo sapiens GN=CHMP1A PE=1 SV=1 - [CHM1A_HUMAN]
Q7LBR1	Charged multivesicular body protein 1b OS=Homo sapiens GN=CHMP1B PE=1 SV=1 - [CHM1B_HUMAN]
O43633	Charged multivesicular body protein 2a OS=Homo sapiens GN=CHMP2A PE=1 SV=1 - [CHM2A_HUMAN]
Q9H444	Charged multivesicular body protein 4b OS=Homo sapiens GN=CHMP4B PE=1 SV=1 - [CHM4B_HUMAN]
Q96FZ7	Charged multivesicular body protein 6 OS=Homo sapiens GN=CHMP6 PE=1 SV=3 - [CHMP6_HUMAN]
Q9BWS9	Chitinase domain-containing protein 1 OS=Homo sapiens GN=CHID1 PE=1 SV=1 - [CHID1_HUMAN]
P06276	Cholinesterase OS=Homo sapiens GN=BCHE PE=1 SV=1 - [CHLE_HUMAN]
C9J5P1	Chromosome 6 open reading frame 130, isoform CRA_a OS=Homo sapiens GN=C6orf130 PE=4 SV=1 - [C9J5P1_HUMAN]
Q5JVC0	Citrate lyase beta like (Fragment) OS=Homo sapiens GN=CLYBL PE=2 SV=1 - [Q5JVC0_HUMAN]
B4DJV2	Citrate synthase OS=Homo sapiens GN=CS PE=2 SV=1 - [B4DJV2_HUMAN]
Q00610	Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5 - [CLH1_HUMAN]
Q14677	Clathrin interactor 1 OS=Homo sapiens GN=CLINT1 PE=1 SV=1 - [EPN4_HUMAN]
P09497	Clathrin light chain B OS=Homo sapiens GN=CLTB PE=1 SV=1 - [CLCB_HUMAN]
Q10570	Cleavage and polyadenylation specificity factor subunit 1 OS=Homo sapiens GN=CPSF1 PE=1 SV=2 - [CPSF1_HUMAN]
P53621	Coatomer subunit alpha OS=Homo sapiens GN=COPA PE=1 SV=2 - [COPA_HUMAN]
P53618	Coatomer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3 - [COPB_HUMAN]

P48444	Coatomer subunit delta OS=Homo sapiens GN=ARCN1 PE=1 SV=1 - [COPD_HUMAN]
Q9UBF2	Coatomer subunit gamma-2 OS=Homo sapiens GN=COPG2 PE=1 SV=1 - [COPG2_HUMAN]
Q6P1N0	Coiled-coil and C2 domain-containing protein 1A OS=Homo sapiens GN=CC2D1A PE=1 SV=1 - [C2D1A_HUMAN]
Q96CT7	Coiled-coil domain-containing protein 124 OS=Homo sapiens GN=CCDC124 PE=1 SV=1 - [CC124_HUMAN]
Q2TAC2	Coiled-coil domain-containing protein 57 OS=Homo sapiens GN=CCDC57 PE=2 SV=2 - [CCD57_HUMAN]
Q16204	Coiled-coil domain-containing protein 6 OS=Homo sapiens GN=CCDC6 PE=1 SV=2 - [CCDC6_HUMAN]
Q6P9F0	Coiled-coil domain-containing protein 62 OS=Homo sapiens GN=CCDC62 PE=1 SV=2 - [CCD62_HUMAN]
Q8NA47	Coiled-coil domain-containing protein 63 OS=Homo sapiens GN=CCDC63 PE=1 SV=1 - [CCD63_HUMAN]
Q9Y2S6	Coiled-coil domain-containing protein 72 OS=Homo sapiens GN=CCDC72 PE=1 SV=1 - [CCD72_HUMAN]
Q8IWF9	Coiled-coil domain-containing protein 83 OS=Homo sapiens GN=CCDC83 PE=2 SV=2 - [CCD83_HUMAN]
Q8NCU4	Coiled-coil domain-containing protein KIAA1407 OS=Homo sapiens GN=KIAA1407 PE=2 SV=1 - [K1407_HUMAN]
Q9NX63	Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondrial OS=Homo sapiens GN=CHCHD3 PE=1 SV=1 - [CHCH3_HUMAN]
Q8IZC6	Collagen alpha-1(XVII) chain OS=Homo sapiens GN=COL27A1 PE=1 SV=1 - [CORA1_HUMAN]
A2A2Y8	Collagen, type XVII, alpha 1 OS=Homo sapiens GN=COL17A1 PE=4 SV=1 - [A2A2Y8_HUMAN]
Q9BXJ0	Complement C1q tumor necrosis factor-related protein 5 OS=Homo sapiens GN=C1QTNF5 PE=1 SV=1 - [C1QT5_HUMAN]
Q07021	Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QBP PE=1 SV=1 - [C1QBP_HUMAN]
Q9BPX3	Condensin complex subunit 3 OS=Homo sapiens GN=NCAPG PE=1 SV=1 - [CND3_HUMAN]
A8K1H6	COP9 constitutive photomorphogenic homolog subunit 8 (Arabidopsis), isoform CRA_b OS=Homo sapiens GN=COPS8 PE=2 SV=1 - [A8K1H6_HUMAN]
Q13098	COP9 signalosome complex subunit 1 OS=Homo sapiens GN=GPS1 PE=1 SV=4 - [CSN1_HUMAN]
P61201	COP9 signalosome complex subunit 2 OS=Homo sapiens GN=COPS2 PE=1 SV=1 - [CSN2_HUMAN]
Q9BT78	COP9 signalosome complex subunit 4 OS=Homo sapiens GN=COPS4 PE=1 SV=1 - [CSN4_HUMAN]
Q92905	COP9 signalosome complex subunit 5 OS=Homo sapiens GN=COPS5 PE=1 SV=4 - [CSN5_HUMAN]
Q9H9Q2	COP9 signalosome complex subunit 7b OS=Homo sapiens GN=COPS7B PE=1 SV=1 - [CSN7B_HUMAN]
Q04656	Copper-transporting ATPase 1 OS=Homo sapiens GN=ATP7A PE=1 SV=3 - [ATP7A_HUMAN]
P36551	Coproporphyrinogen-III oxidase, mitochondrial OS=Homo sapiens GN=CPOX PE=1 SV=3 - [HEM6_HUMAN]
P31146	Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4 - [COR1A_HUMAN]
Q9ULV4	Coronin-1C OS=Homo sapiens GN=CORO1C PE=1 SV=1 - [COR1C_HUMAN]
Q9NRF8	CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1 - [PYRG2_HUMAN]
Q2HIY8	CTSC protein OS=Homo sapiens GN=CTSC PE=2 SV=1 - [Q2HIY8_HUMAN]
Q5T2B5	Cullin 2 OS=Homo sapiens GN=CUL2 PE=2 SV=1 - [Q5T2B5_HUMAN]
Q13618	Cullin-3 OS=Homo sapiens GN=CUL3 PE=1 SV=2 - [CUL3_HUMAN]
Q13620	Cullin-4B OS=Homo sapiens GN=CUL4B PE=1 SV=4 - [CUL4B_HUMAN]
Q93034	Cullin-5 OS=Homo sapiens GN=CUL5 PE=1 SV=4 - [CUL5_HUMAN]
Q86VP6	Cullin-associated NEDD8-dissociated protein 1 OS=Homo sapiens GN=CAND1 PE=1 SV=2 - [CAND1_HUMAN]
Q14028	Cyclic nucleotide-gated cation channel beta-1 OS=Homo sapiens GN=CNGB1 PE=1 SV=2 - [CNGB1_HUMAN]
Q15078	Cyclin-dependent kinase 5 activator 1 OS=Homo sapiens GN=CDK5R1 PE=1 SV=1 - [CD5R1_HUMAN]
Q9NZV1	Cysteine-rich motor neuron 1 protein OS=Homo sapiens GN=CRIM1 PE=1 SV=1 - [CRIM1_HUMAN]
A8MVQ3	Cysteinyl-tRNA synthetase, isoform CRA_b OS=Homo sapiens GN=CARS PE=2 SV=1 - [A8MVQ3_HUMAN]
P00167	Cytochrome b5 OS=Homo sapiens GN=CYB5A PE=1 SV=2 - [CYB5_HUMAN]
P31930	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 SV=3 - [QCR1_HUMAN]
P22695	Cytochrome b-c1 complex subunit 2, mitochondrial OS=Homo sapiens GN=UQCRC2 PE=1 SV=3 - [QCR2_HUMAN]
P07919	Cytochrome b-c1 complex subunit 6, mitochondrial OS=Homo sapiens GN=UQCRH PE=1 SV=2 - [QCR6_HUMAN]
O14949	Cytochrome b-c1 complex subunit 8 OS=Homo sapiens GN=UQCRQ PE=1 SV=4 - [QCR8_HUMAN]

P47985	Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Homo sapiens GN=UQCRFS1 PE=1 SV=2 - [UCRI_HUMAN]
Q14061	Cytochrome c oxidase copper chaperone OS=Homo sapiens GN=COX17 PE=1 SV=2 - [COX17_HUMAN]
P00403	Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1 - [COX2_HUMAN]
P13073	Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Homo sapiens GN=COX4I1 PE=1 SV=1 - [COX41_HUMAN]
P20674	Cytochrome c oxidase subunit 5A, mitochondrial OS=Homo sapiens GN=COX5A PE=1 SV=2 - [COX5A_HUMAN]
P10606	Cytochrome c oxidase subunit 5B, mitochondrial OS=Homo sapiens GN=COX5B PE=1 SV=2 - [COX5B_HUMAN]
P14854	Cytochrome c oxidase subunit 6B1 OS=Homo sapiens GN=COX6B1 PE=1 SV=2 - [CX6B1_HUMAN]
P09669	Cytochrome c oxidase subunit 6C OS=Homo sapiens GN=COX6C PE=1 SV=2 - [COX6C_HUMAN]
P08574	Cytochrome c1, heme protein, mitochondrial OS=Homo sapiens GN=CYC1 PE=1 SV=3 - [CY1_HUMAN]
P53701	Cytochrome c-type heme lyase OS=Homo sapiens GN=HCCS PE=1 SV=1 - [CCHL_HUMAN]
Q5VXH9	Cytochrome P450, family 26, subfamily A, polypeptide 1 OS=Homo sapiens GN=CYP26A1 PE=2 SV=1 - [Q5VXH9_HUMAN]
Q8IUI8	Cytokine receptor-like factor 3 OS=Homo sapiens GN=CRLF3 PE=1 SV=2 - [CRLF3_HUMAN]
Q14008	Cytoskeleton-associated protein 5 OS=Homo sapiens GN=CKAP5 PE=1 SV=3 - [CKAP5_HUMAN]
P28838	Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3 - [AMPL_HUMAN]
Q10586	D site-binding protein OS=Homo sapiens GN=DBP PE=2 SV=1 - [DBP_HUMAN]
Q58WW2	DDB1- and CUL4-associated factor 6 OS=Homo sapiens GN=DCAF6 PE=1 SV=1 - [DCAF6_HUMAN]
Q96HY6	DDRKG domain-containing protein 1 OS=Homo sapiens GN=DDRKG1 PE=1 SV=2 - [DDRKG_HUMAN]
D6RJA6	DEAD (Asp-Glu-Ala-Asp) box polypeptide 46, isoform CRA_b OS=Homo sapiens GN=DDX46 PE=4 SV=1 - [D6RJA6_HUMAN]
Q14UF5	Decay-accelerating factor splicing variant 2 OS=Homo sapiens GN=DAF PE=2 SV=1 - [Q14UF5_HUMAN]
Q8IZD9	Dedicator of cytokinesis protein 3 OS=Homo sapiens GN=DOCK3 PE=1 SV=1 - [DOCK3_HUMAN]
Q9Y238	Deleted in lung and esophageal cancer protein 1 OS=Homo sapiens GN=DLEC1 PE=2 SV=2 - [DLEC1_HUMAN]
Q13011	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH1 PE=1 SV=2 - [ECH1_HUMAN]
P54886	Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2 - [P5CS_HUMAN]
B7Z3I9	Delta-aminolevulinic acid dehydratase OS=Homo sapiens PE=2 SV=1 - [B7Z3I9_HUMAN]
Q08495	Dematin OS=Homo sapiens GN=EPB49 PE=1 SV=3 - [DEMA_HUMAN]
O43583	Density-regulated protein OS=Homo sapiens GN=DENR PE=1 SV=2 - [DENR_HUMAN]
Q5QJE6	Deoxynucleotidyltransferase terminal-interacting protein 2 OS=Homo sapiens GN=DNTTIP2 PE=1 SV=2 - [TDIF2_HUMAN]
P15924	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3 - [DESP_HUMAN]
Q96JH7	Deubiquitinating protein VCIP135 OS=Homo sapiens GN=VCPIP1 PE=1 SV=2 - [VCIP1_HUMAN]
F5H4I0	Dihydrolipoyl dehydrogenase OS=Homo sapiens GN=DLD PE=3 SV=1 - [F5H4I0_HUMAN]
P36957	Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLST PE=1 SV=4 - [ODO2_HUMAN]
P53602	Diphosphomevalonate decarboxylase OS=Homo sapiens GN=MVD PE=1 SV=1 - [MVD1_HUMAN]
Q9P265	Disco-interacting protein 2 homolog B OS=Homo sapiens GN=DIP2B PE=1 SV=3 - [DIP2B_HUMAN]
Q9Y4D1	Disheveled-associated activator of morphogenesis 1 OS=Homo sapiens GN=DAAM1 PE=1 SV=2 - [DAAM1_HUMAN]
Q16531	DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1 - [DDB1_HUMAN]
P04053	DNA nucleotidyltransferase OS=Homo sapiens GN=DNTT PE=1 SV=3 - [TDT_HUMAN]
Q92878	DNA repair protein RAD50 OS=Homo sapiens GN=RAD50 PE=1 SV=1 - [RAD50_HUMAN]
P16989	DNA-binding protein A OS=Homo sapiens GN=CSDA PE=1 SV=4 - [DBPA_HUMAN]
F5H643	DNA-directed RNA polymerase OS=Homo sapiens GN=POLR1B PE=3 SV=1 - [F5H643_HUMAN]
D3DQ66	DnaJ (Hsp40) homolog, subfamily C, member 6, isoform CRA_c OS=Homo sapiens GN=DNAJC6 PE=4 SV=1 - [D3DQ66_HUMAN]
P25685	DnaJ homolog subfamily B member 1 OS=Homo sapiens GN=DNAJB1 PE=1 SV=4 - [DNJB1_HUMAN]
Q9UBS4	DnaJ homolog subfamily B member 11 OS=Homo sapiens GN=DNAJB11 PE=1 SV=1 - [DJB11_HUMAN]
Q9UDY4	DnaJ homolog subfamily B member 4 OS=Homo sapiens GN=DNAJB4 PE=1 SV=1 - [DNJB4_HUMAN]

Q96KC8	DnaJ homolog subfamily C member 1 OS=Homo sapiens GN=DNAJC1 PE=1 SV=1 - [DNJC1_HUMAN]
Q9NVH1	DnaJ homolog subfamily C member 11 OS=Homo sapiens GN=DNAJC11 PE=1 SV=2 - [DJC11_HUMAN]
O75165	DnaJ homolog subfamily C member 13 OS=Homo sapiens GN=DNAJC13 PE=1 SV=5 - [DJC13_HUMAN]
Q9NVM6	DnaJ homolog subfamily C member 17 OS=Homo sapiens GN=DNAJC17 PE=1 SV=1 - [DJC17_HUMAN]
Q9H3Z4	DnaJ homolog subfamily C member 5 OS=Homo sapiens GN=DNAJC5 PE=1 SV=1 - [DNJC5_HUMAN]
O75937	DnaJ homolog subfamily C member 8 OS=Homo sapiens GN=DNAJC8 PE=1 SV=2 - [DNJC8_HUMAN]
Q8WXX5	DnaJ homolog subfamily C member 9 OS=Homo sapiens GN=DNAJC9 PE=1 SV=1 - [DNJC9_HUMAN]
O60762	Dolichol-phosphate mannosyltransferase OS=Homo sapiens GN=DPM1 PE=1 SV=1 - [DPM1_HUMAN]
Q9P2X0	Dolichol-phosphate mannosyltransferase subunit 3 OS=Homo sapiens GN=DPM3 PE=1 SV=2 - [DPM3_HUMAN]
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1 - [RPN1_HUMAN]
P49959	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=3 - [MRE11_HUMAN]
P46734	Dual specificity mitogen-activated protein kinase kinase 3 OS=Homo sapiens GN=MAP2K3 PE=1 SV=2 - [MP2K3_HUMAN]
P49761	Dual specificity protein kinase CLK3 OS=Homo sapiens GN=CLK3 PE=1 SV=3 - [CLK3_HUMAN]
Q9NR20	Dual specificity tyrosine-phosphorylation-regulated kinase 4 OS=Homo sapiens GN=DYRK4 PE=1 SV=2 - [DYRK4_HUMAN]
Q5T1I6	Dynactin 3 (P22) (Fragment) OS=Homo sapiens GN=DCTN3 PE=2 SV=1 - [Q5T1I6_HUMAN]
Q13561	Dynactin subunit 2 OS=Homo sapiens GN=DCTN2 PE=1 SV=4 - [DCTN2_HUMAN]
P50570	Dynamin-2 OS=Homo sapiens GN=DNM2 PE=1 SV=2 - [DYN2_HUMAN]
O60313	Dynamin-like 120 kDa protein, mitochondrial OS=Homo sapiens GN=OPA1 PE=1 SV=3 - [OPA1_HUMAN]
Q8IVF4	Dynein heavy chain 10, axonemal OS=Homo sapiens GN=DNAH10 PE=1 SV=4 - [DYH10_HUMAN]
Q8TD57	Dynein heavy chain 3, axonemal OS=Homo sapiens GN=DNAH3 PE=2 SV=1 - [DYH3_HUMAN]
P63167	Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1 - [DYL1_HUMAN]
Q96FJ2	Dynein light chain 2, cytoplasmic OS=Homo sapiens GN=DYNLL2 PE=1 SV=1 - [DYL2_HUMAN]
Q14258	E3 ubiquitin/ISG15 ligase TRIM25 OS=Homo sapiens GN=TRIM25 PE=1 SV=2 - [TRI25_HUMAN]
O95376	E3 ubiquitin-protein ligase ARIH2 OS=Homo sapiens GN=ARIH2 PE=1 SV=1 - [ARI2_HUMAN]
Q9UNE7	E3 ubiquitin-protein ligase CHIP OS=Homo sapiens GN=STUB1 PE=1 SV=2 - [CHIP_HUMAN]
Q9ULT8	E3 ubiquitin-protein ligase HECTD1 OS=Homo sapiens GN=HECTD1 PE=1 SV=3 - [HECD1_HUMAN]
O95714	E3 ubiquitin-protein ligase HERC2 OS=Homo sapiens GN=HERC2 PE=1 SV=2 - [HERC2_HUMAN]
P62877	E3 ubiquitin-protein ligase RBX1 OS=Homo sapiens GN=RBX1 PE=1 SV=1 - [RBX1_HUMAN]
Q5XPI4	E3 ubiquitin-protein ligase RNF123 OS=Homo sapiens GN=RNF123 PE=1 SV=1 - [RN123_HUMAN]
Q8IWW8	E3 ubiquitin-protein ligase UBR2 OS=Homo sapiens GN=UBR2 PE=1 SV=1 - [UBR2_HUMAN]
Q5T4S7	E3 ubiquitin-protein ligase UBR4 OS=Homo sapiens GN=UBR4 PE=1 SV=1 - [UBR4_HUMAN]
Q96JP5	E3 ubiquitin-protein ligase ZFP91 OS=Homo sapiens GN=ZFP91 PE=1 SV=1 - [ZFP91_HUMAN]
Q15075	Early endosome antigen 1 OS=Homo sapiens GN=EEA1 PE=1 SV=2 - [EEA1_HUMAN]
Q93070	Ecto-ADP-ribosyltransferase 4 OS=Homo sapiens GN=ART4 PE=2 SV=2 - [NAR4_HUMAN]
Q4VBZ6	EEF1D protein OS=Homo sapiens GN=EEF1D PE=1 SV=1 - [Q4VBZ6_HUMAN]
Q8N3D4	EH domain-binding protein 1-like protein 1 OS=Homo sapiens GN=EHBP1L1 PE=1 SV=2 - [EH1L1_HUMAN]
Q9H4M9	EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2 - [EHD1_HUMAN]
P38117	Electron transfer flavoprotein subunit beta OS=Homo sapiens GN=ETFB PE=1 SV=3 - [ETFB_HUMAN]
Q2Y0W8	Electroneutral sodium bicarbonate exchanger 1 OS=Homo sapiens GN=SLC4A8 PE=1 SV=1 - [S4A8_HUMAN]
P68104	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]
P24534	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3 - [EF1B_HUMAN]
P26641	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3 - [EF1G_HUMAN]
P13639	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4 - [EF2_HUMAN]

P49411	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2 - [EFTU_HUMAN]
P50402	Emerin OS=Homo sapiens GN=EMD PE=1 SV=1 - [EMD_HUMAN]
Q9BXX0	EMILIN-2 OS=Homo sapiens GN=EMILIN2 PE=1 SV=3 - [EMIL2_HUMAN]
O94919	Endonuclease domain-containing 1 protein OS=Homo sapiens GN=ENDOD1 PE=1 SV=2 - [ENDD1_HUMAN]
Q9NR46	Endophilin-B2 OS=Homo sapiens GN=SH3GLB2 PE=1 SV=1 - [SHLB2_HUMAN]
Q9NZ08	Endoplasmic reticulum aminopeptidase 1 OS=Homo sapiens GN=ERAP1 PE=1 SV=3 - [ERAP1_HUMAN]
P30040	Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=1 SV=4 - [ERP29_HUMAN]
Q9BS26	Endoplasmic reticulum resident protein 44 OS=Homo sapiens GN=ERP44 PE=1 SV=1 - [ERP44_HUMAN]
Q96RQ1	Endoplasmic reticulum-Golgi intermediate compartment protein 2 OS=Homo sapiens GN=ERGIC2 PE=1 SV=2 - [ERGI2_HUMAN]
Q9Y282	Endoplasmic reticulum-Golgi intermediate compartment protein 3 OS=Homo sapiens GN=ERGIC3 PE=1 SV=1 - [ERGI3_HUMAN]
P14625	Endoplasmic OS=Homo sapiens GN=HSP90B1 PE=1 SV=1 - [ENPL_HUMAN]
O60869	Endothelial differentiation-related factor 1 OS=Homo sapiens GN=EDF1 PE=1 SV=1 - [EDF1_HUMAN]
F5H0C8	Enolase OS=Homo sapiens GN=ENO2 PE=3 SV=1 - [F5H0C8_HUMAN]
Q5W0J6	Enoyl Coenzyme A hydratase domain containing 3 (Fragment) OS=Homo sapiens GN=ECHDC3 PE=2 SV=1 - [Q5W0J6_HUMAN]
P42126	Enoyl-CoA delta isomerase 1, mitochondrial OS=Homo sapiens GN=ECI1 PE=1 SV=1 - [ECI1_HUMAN]
Q4KKX0	EPB42 protein OS=Homo sapiens GN=EPB42 PE=2 SV=1 - [Q4KKX0_HUMAN]
B1AUU8	Epidermal growth factor receptor pathway substrate 15 OS=Homo sapiens GN=EPS15 PE=4 SV=1 - [B1AUU8_HUMAN]
Q9NRG7	Epimerase family protein SDR39U1 OS=Homo sapiens GN=SDR39U1 PE=2 SV=2 - [D39U1_HUMAN]
P07099	Epoxide hydrolase 1 OS=Homo sapiens GN=EPHX1 PE=1 SV=1 - [HYEP_HUMAN]
A2RRF3	EPS15L1 protein OS=Homo sapiens GN=EPS15L1 PE=2 SV=1 - [A2RRF3_HUMAN]
Q9Y6I3	Epsin-1 OS=Homo sapiens GN=EPN1 PE=1 SV=2 - [EPN1_HUMAN]
Q99808	Equilibrative nucleoside transporter 1 OS=Homo sapiens GN=SLC29A1 PE=1 SV=3 - [S29A1_HUMAN]
O75477	Erlin-1 OS=Homo sapiens GN=ERLIN1 PE=1 SV=1 - [ERLN1_HUMAN]
Q96PL5	Erythroid membrane-associated protein OS=Homo sapiens GN=ERMAP PE=1 SV=1 - [ERMAP_HUMAN]
P30042	ES1 protein homolog, mitochondrial OS=Homo sapiens GN=C21orf33 PE=1 SV=3 - [ES1_HUMAN]
Q6ZVH7	Espin-like protein OS=Homo sapiens GN=ESPNL PE=1 SV=3 - [ESPNL_HUMAN]
Q53GQ0	Estradiol 17-beta-dehydrogenase 12 OS=Homo sapiens GN=HSD17B12 PE=1 SV=2 - [DHB12_HUMAN]
P60842	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1 - [IF4A1_HUMAN]
P38919	Eukaryotic initiation factor 4A-III OS=Homo sapiens GN=EIF4A3 PE=1 SV=4 - [IF4A3_HUMAN]
P62495	Eukaryotic peptide chain release factor subunit 1 OS=Homo sapiens GN=ETF1 PE=1 SV=3 - [ERF1_HUMAN]
P05198	Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens GN=EIF2S1 PE=1 SV=3 - [IF2A_HUMAN]
P20042	Eukaryotic translation initiation factor 2 subunit 2 OS=Homo sapiens GN=EIF2S2 PE=1 SV=2 - [IF2B_HUMAN]
P41091	Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=3 - [IF2G_HUMAN]
Q5QP88	Eukaryotic translation initiation factor 2B, subunit 3 gamma, 58kDa (Fragment) OS=Homo sapiens GN=EIF2B3 PE=2 SV=1 - [Q5QP88_HUMAN]
B1ALI0	Eukaryotic translation initiation factor 2C, 3 OS=Homo sapiens GN=EIF2C3 PE=4 SV=1 - [B1ALI0_HUMAN]
Q14152	Eukaryotic translation initiation factor 3 subunit A OS=Homo sapiens GN=EIF3A PE=1 SV=1 - [EIF3A_HUMAN]
P60228	Eukaryotic translation initiation factor 3 subunit E OS=Homo sapiens GN=EIF3E PE=1 SV=1 - [EIF3E_HUMAN]
O75821	Eukaryotic translation initiation factor 3 subunit G OS=Homo sapiens GN=EIF3G PE=1 SV=2 - [EIF3G_HUMAN]
O15372	Eukaryotic translation initiation factor 3 subunit H OS=Homo sapiens GN=EIF3H PE=1 SV=1 - [EIF3H_HUMAN]
Q13347	Eukaryotic translation initiation factor 3 subunit I OS=Homo sapiens GN=EIF3I PE=1 SV=1 - [EIF3I_HUMAN]
O75822	Eukaryotic translation initiation factor 3 subunit J OS=Homo sapiens GN=EIF3J PE=1 SV=2 - [EIF3J_HUMAN]
Q9Y262	Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3L PE=1 SV=1 - [EIF3L_HUMAN]
P06730	Eukaryotic translation initiation factor 4E OS=Homo sapiens GN=EIF4E PE=1 SV=2 - [IF4E_HUMAN]

P55010	Eukaryotic translation initiation factor 5 OS=Homo sapiens GN=EIF5 PE=1 SV=2 - [IF5_HUMAN]
P63241	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2 - [IF5A1_HUMAN]
O60841	Eukaryotic translation initiation factor 5B OS=Homo sapiens GN=EIF5B PE=1 SV=4 - [IF2P_HUMAN]
C9JGE3	Ewing sarcoma breakpoint region 1, isoform CRA_e OS=Homo sapiens GN=EWSR1 PE=4 SV=1 - [C9JGE3_HUMAN]
O00471	Exocyst complex component 5 OS=Homo sapiens GN=EXOC5 PE=1 SV=1 - [EXOC5_HUMAN]
Q9BSJ8	Extended synaptotagmin-1 OS=Homo sapiens GN=ESYT1 PE=1 SV=1 - [ESYT1_HUMAN]
A0FGR8	Extended synaptotagmin-2 OS=Homo sapiens GN=ESYT2 PE=1 SV=1 - [ESYT2_HUMAN]
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4 - [EZRI_HUMAN]
P52907	F-actin-capping protein subunit alpha-1 OS=Homo sapiens GN=CAPZA1 PE=1 SV=3 - [CAZA1_HUMAN]
B1AJQ5	Family with sequence similarity 76, member A (Fragment) OS=Homo sapiens GN=FAM76A PE=4 SV=1 - [B1AJQ5_HUMAN]
Q92945	Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=4 - [FUBP2_HUMAN]
P49327	Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3 - [FAS_HUMAN]
Q6PIJ6	F-box only protein 38 OS=Homo sapiens GN=FBXO38 PE=1 SV=3 - [FBX38_HUMAN]
Q5TI86	F-box protein 7 OS=Homo sapiens GN=FBXO7 PE=2 SV=1 - [Q5TI86_HUMAN]
Q96IG2	F-box/LRR-repeat protein 20 OS=Homo sapiens GN=FBXL20 PE=1 SV=2 - [FXL20_HUMAN]
Q5JV73	FERM and PDZ domain-containing protein 3 OS=Homo sapiens GN=FRMPD3 PE=2 SV=2 - [FRPD3_HUMAN]
Q86UX7	Fermitin family homolog 3 OS=Homo sapiens GN=FERMT3 PE=1 SV=1 - [URP2_HUMAN]
P02794	Ferritin heavy chain OS=Homo sapiens GN=FTH1 PE=1 SV=2 - [FRIH_HUMAN]
Q9BXU8	Ferritin heavy polypeptide-like 17 OS=Homo sapiens GN=FTHL17 PE=2 SV=1 - [FHL17_HUMAN]
P02792	Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2 - [FRIL_HUMAN]
P22830	Ferrochelatase, mitochondrial OS=Homo sapiens GN=FECH PE=1 SV=2 - [HEMH_HUMAN]
P55075	Fibroblast growth factor 8 OS=Homo sapiens GN=FGF8 PE=1 SV=1 - [FGF8_HUMAN]
Q5HY54	Filamin A, alpha (Actin binding protein 280) OS=Homo sapiens GN=FLNA PE=2 SV=1 - [Q5HY54_HUMAN]
Q14315	Filamin-C OS=Homo sapiens GN=FLNC PE=1 SV=3 - [FLNC_HUMAN]
Q5T1M5	FK506-binding protein 15 OS=Homo sapiens GN=FKBP15 PE=1 SV=2 - [FKB15_HUMAN]
P30043	Flavin reductase OS=Homo sapiens GN=BLVRB PE=1 SV=3 - [BLVRB_HUMAN]
O75955	Flotillin-1 OS=Homo sapiens GN=FLOT1 PE=1 SV=3 - [FLOT1_HUMAN]
Q14254	Flotillin-2 OS=Homo sapiens GN=FLOT2 PE=1 SV=2 - [FLOT2_HUMAN]
Q9H479	Fructosamine-3-kinase OS=Homo sapiens GN=FN3K PE=1 SV=1 - [FN3K_HUMAN]
P04075	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2 - [ALDOA_HUMAN]
B7Z1N6	Fructose-bisphosphate aldolase OS=Homo sapiens GN=ALDOC PE=2 SV=1 - [B7Z1N6_HUMAN]
Q86TT8	Full-length cDNA clone CS0DC011YL17 of Neuroblastoma of Homo sapiens (human) OS=Homo sapiens GN=MARK3 PE=2 SV=1 - [Q86TT8_HUMAN]
P07954	Fumarate hydratase, mitochondrial OS=Homo sapiens GN=FH PE=1 SV=3 - [FUMH_HUMAN]
Q9BWH2	FUN14 domain-containing protein 2 OS=Homo sapiens GN=FUNDC2 PE=1 SV=2 - [FUND2_HUMAN]
P48544	G protein-activated inward rectifier potassium channel 4 OS=Homo sapiens GN=KCNJ5 PE=1 SV=2 - [IRK5_HUMAN]
P43250	G protein-coupled receptor kinase 6 OS=Homo sapiens GN=GRK6 PE=1 SV=2 - [GRK6_HUMAN]
Q8WWL7	G2/mitotic-specific cyclin-B3 OS=Homo sapiens GN=CCNB3 PE=1 SV=2 - [CCNB3_HUMAN]
P17931	Galectin-3 OS=Homo sapiens GN=LGALS3 PE=1 SV=5 - [LEG3_HUMAN]
Q9UEY8	Gamma-adducin OS=Homo sapiens GN=ADD3 PE=1 SV=1 - [ADDG_HUMAN]
O95166	Gamma-aminobutyric acid receptor-associated protein OS=Homo sapiens GN=GABARAP PE=1 SV=1 - [GBRAP_HUMAN]
P60520	Gamma-aminobutyric acid receptor-associated protein-like 2 OS=Homo sapiens GN=GABARAPL2 PE=1 SV=1 - [GBRL2_HUMAN]
Q92820	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2 - [GGH_HUMAN]
Q99747	Gamma-soluble NSF attachment protein OS=Homo sapiens GN=NAPG PE=1 SV=1 - [SNAG_HUMAN]

Q92696	Geranylgeranyl transferase type-2 subunit alpha OS=Homo sapiens GN=RABGGTA PE=1 SV=2 - [PGTA_HUMAN]
C9IYJ3	Glioblastoma amplified sequence, isoform CRA_a OS=Homo sapiens GN=GBAS PE=4 SV=1 - [C9IYJ3_HUMAN]
Q6UWM5	GLIPR1-like protein 1 OS=Homo sapiens GN=GLIPR1L1 PE=1 SV=2 - [GPRL1_HUMAN]
E9PDV8	Glucose-6-phosphate 1-dehydrogenase OS=Homo sapiens GN=G6PD PE=3 SV=1 - [E9PDV8_HUMAN]
O43826	Glucose-6-phosphate translocase OS=Homo sapiens GN=SLC37A4 PE=1 SV=1 - [G6PT1_HUMAN]
P15104	Glutamine synthetase OS=Homo sapiens GN=GLUL PE=1 SV=4 - [GLNA_HUMAN]
Q86SX6	Glutaredoxin-related protein 5, mitochondrial OS=Homo sapiens GN=GLRX5 PE=1 SV=2 - [GLRX5_HUMAN]
Q92947	Glutaryl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=GCDH PE=1 SV=1 - [GCDH_HUMAN]
C8KIM0	Glutathion reductase delta8+9 alternative splicing variant OS=Homo sapiens GN=GSR PE=2 SV=1 - [C8KIM0_HUMAN]
P07203	Glutathione peroxidase 1 OS=Homo sapiens GN=GPX1 PE=1 SV=4 - [GPX1_HUMAN]
P78417	Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2 - [GSTO1_HUMAN]
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3 - [G3P_HUMAN]
Q9NZC3	Glycerophosphodiester phosphodiesterase 1 OS=Homo sapiens GN=GDE1 PE=1 SV=1 - [GDE1_HUMAN]
B8Q185	Glycophorin A MNS blood group OS=Homo sapiens GN=GYP A PE=4 SV=1 - [B8Q185_HUMAN]
P04921	Glycophorin-C OS=Homo sapiens GN=GYP C PE=1 SV=1 - [GLPC_HUMAN]
P41250	Glycyl-tRNA synthetase OS=Homo sapiens GN=GARS PE=1 SV=3 - [SYG_HUMAN]
P36959	GMP reductase 1 OS=Homo sapiens GN=GMPR PE=1 SV=1 - [GMPR1_HUMAN]
Q5T5I6	Golgi phosphoprotein 3-like (Fragment) OS=Homo sapiens GN=GOLPH3L PE=2 SV=1 - [Q5T5I6_HUMAN]
Q9H4G4	Golgi-associated plant pathogenesis-related protein 1 OS=Homo sapiens GN=GLIPR2 PE=1 SV=3 - [GAPR1_HUMAN]
Q7Z5G4	Golgin subfamily A member 7 OS=Homo sapiens GN=GOLGA7 PE=1 SV=2 - [GOGA7_HUMAN]
Q92538	Golgi-specific brefeldin A-resistance guanine nucleotide exchange factor 1 OS=Homo sapiens GN=GBF1 PE=1 SV=2 - [GBF1_HUMAN]
B3KUR5	GRAM domain containing 1C, isoform CRA_d OS=Homo sapiens GN=GRAMD1C PE=2 SV=1 - [B3KUR5_HUMAN]
P28799	Granulins OS=Homo sapiens GN=GRN PE=1 SV=2 - [GRN_HUMAN]
Q8IWI2	GRIP and coiled-coil domain-containing protein 2 OS=Homo sapiens GN=GCC2 PE=1 SV=4 - [GCC2_HUMAN]
B1B0M1	GRIP1 associated protein 1 OS=Homo sapiens GN=GRIPAP1 PE=4 SV=1 - [B1B0M1_HUMAN]
B4DWV5	GrpE protein homolog OS=Homo sapiens GN=GRPEL1 PE=2 SV=1 - [B4DWV5_HUMAN]
P01112	GTPase HRas OS=Homo sapiens GN=HRAS PE=1 SV=1 - [RASH_HUMAN]
P01116	GTPase KRas OS=Homo sapiens GN=KRAS PE=1 SV=1 - [RASK_HUMAN]
P01111	GTPase NRas OS=Homo sapiens GN=NRAS PE=1 SV=1 - [RASN_HUMAN]
Q14C86	GTPase-activating protein and VPS9 domain-containing protein 1 OS=Homo sapiens GN=GAPVD1 PE=1 SV=2 - [GAPD1_HUMAN]
P62826	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3 - [RAN_HUMAN]
O00178	GTP-binding protein 1 OS=Homo sapiens GN=GTPBP1 PE=1 SV=3 - [GTPB1_HUMAN]
Q9BX10	GTP-binding protein 2 OS=Homo sapiens GN=GTPBP2 PE=2 SV=1 - [GTPB2_HUMAN]
O75628	GTP-binding protein REM 1 OS=Homo sapiens GN=REM1 PE=1 SV=2 - [REM1_HUMAN]
P04899	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3 - [GNAI2_HUMAN]
Q9P2W3	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-13 OS=Homo sapiens GN=GNG13 PE=2 SV=1 - [GBG13_HUMAN]
P59768	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-2 OS=Homo sapiens GN=GNG2 PE=1 SV=2 - [GBG2_HUMAN]
P63218	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-5 OS=Homo sapiens GN=GNG5 PE=1 SV=3 - [GBG5_HUMAN]
Q9UK08	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-8 OS=Homo sapiens GN=GNG8 PE=2 SV=1 - [GBG8_HUMAN]
P62873	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3 - [GBB1_HUMAN]
P62879	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Homo sapiens GN=GNB2 PE=1 SV=3 - [GBB2_HUMAN]
P08754	Guanine nucleotide-binding protein G(k) subunit alpha OS=Homo sapiens GN=GNAI3 PE=1 SV=3 - [GNAI3_HUMAN]
P50148	Guanine nucleotide-binding protein G(q) subunit alpha OS=Homo sapiens GN=GNAQ PE=1 SV=4 - [GNAQ_HUMAN]

P63092	Guanine nucleotide-binding protein G(s) subunit alpha isoforms short OS=Homo sapiens GN=GNAS PE=1 SV=1 - [GNAS2_HUMAN]
P29992	Guanine nucleotide-binding protein subunit alpha-11 OS=Homo sapiens GN=GNA11 PE=1 SV=2 - [GNA11_HUMAN]
Q14344	Guanine nucleotide-binding protein subunit alpha-13 OS=Homo sapiens GN=GNA13 PE=1 SV=2 - [GNA13_HUMAN]
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3 - [GBLP_HUMAN]
B3KSY4	HCG1787779, isoform CRA_b OS=Homo sapiens GN=CORO7 PE=2 SV=1 - [B3KSY4_HUMAN]
B2RDM2	HCG1811539, isoform CRA_b OS=Homo sapiens GN=TXNDC5 PE=2 SV=1 - [B2RDM2_HUMAN]
D3DTT2	HCG1820468, isoform CRA_a OS=Homo sapiens GN=FAM18B2 PE=4 SV=1 - [D3DTT2_HUMAN]
Q6PIK3	HCG1995540, isoform CRA_b (Hcg2044074, isoform cra_d) OS=Homo sapiens GN=RAB4B PE=2 SV=1 - [Q6PIK3_HUMAN]
B4DP52	HCG2005638, isoform CRA_c OS=Homo sapiens GN=DDX39B PE=2 SV=1 - [B4DP52_HUMAN]
A6PVK0	HCG2019817, isoform CRA_b OS=Homo sapiens GN=RBCK1 PE=4 SV=1 - [A6PVK0_HUMAN]
B4DHA4	HCG2031650, isoform CRA_f OS=Homo sapiens GN=NSUN4 PE=2 SV=1 - [B4DHA4_HUMAN]
A2A3U5	HCG20471, isoform CRA_d OS=Homo sapiens GN=OPRS1 PE=2 SV=1 - [A2A3U5_HUMAN]
Q4VXL4	HCG41426, isoform CRA_c OS=Homo sapiens GN=TACC2 PE=2 SV=1 - [Q4VXL4_HUMAN]
Q5T3Q7	HEAT repeat containing 1 OS=Homo sapiens GN=HEATR1 PE=2 SV=1 - [Q5T3Q7_HUMAN]
B8ZZL8	Heat shock 10kDa protein 1 (Chaperonin 10), isoform CRA_b OS=Homo sapiens GN=HSPE1 PE=3 SV=1 - [B8ZZL8_HUMAN]
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5 - [HSP71_HUMAN]
P34931	Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2 - [HS71L_HUMAN]
P34932	Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4 - [HSP74_HUMAN]
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]
P04792	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2 - [HSPB1_HUMAN]
P07900	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5 - [HS90A_HUMAN]
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [HS90B_HUMAN]
Q14527	Helicase-like transcription factor OS=Homo sapiens GN=HLTF PE=1 SV=2 - [HLTF_HUMAN]
P09601	Heme oxygenase 1 OS=Homo sapiens GN=HMOX1 PE=1 SV=1 - [HMOX1_HUMAN]
Q9NRV9	Heme-binding protein 1 OS=Homo sapiens GN=HEBP1 PE=1 SV=1 - [HEBP1_HUMAN]
Q9Y5Z4	Heme-binding protein 2 OS=Homo sapiens GN=HEBP2 PE=1 SV=1 - [HEBP2_HUMAN]
Q9BXL5	Hemogen OS=Homo sapiens GN=HEMGN PE=1 SV=1 - [HEMGN_HUMAN]
P69905	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2 - [HBA_HUMAN]
P68871	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2 - [HBB_HUMAN]
P02042	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2 - [HBD_HUMAN]
P02100	Hemoglobin subunit epsilon OS=Homo sapiens GN=HBE1 PE=1 SV=2 - [HBE_HUMAN]
P69891	Hemoglobin subunit gamma-1 OS=Homo sapiens GN=HBG1 PE=1 SV=2 - [HBG1_HUMAN]
P69892	Hemoglobin subunit gamma-2 OS=Homo sapiens GN=HBG2 PE=1 SV=2 - [HBG2_HUMAN]
Q6B0K9	Hemoglobin subunit mu OS=Homo sapiens GN=HBM PE=2 SV=1 - [HBM_HUMAN]
P02008	Hemoglobin subunit zeta OS=Homo sapiens GN=HBZ PE=1 SV=2 - [HBAZ_HUMAN]
D9YZU8	Hemoglobin, gamma A OS=Homo sapiens GN=HBG1 PE=3 SV=1 - [D9YZU8_HUMAN]
P51858	Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1 - [HDGF_HUMAN]
P09651	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5 - [ROA1_HUMAN]
P31942	Heterogeneous nuclear ribonucleoprotein H3 OS=Homo sapiens GN=HNRNPH3 PE=1 SV=2 - [HNRH3_HUMAN]
Q5T6W5	Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=2 SV=1 - [Q5T6W5_HUMAN]
P14866	Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2 - [HNRPL_HUMAN]
P52272	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3 - [HNRPM_HUMAN]
O60506	Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2 - [HNRPQ_HUMAN]

Q2L7G6	Heterogeneous nuclear ribonucleoprotein-R2 OS=Homo sapiens GN=HNRNPR PE=2 SV=1 - [Q2L7G6_HUMAN]
P22626	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2 - [ROA2_HUMAN]
P26583	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2 - [HMGB2_HUMAN]
Q5T7C0	High-mobility group box 1 (Fragment) OS=Homo sapiens GN=HMGB1 PE=2 SV=1 - [Q5T7C0_HUMAN]
Q9BX68	Histidine triad nucleotide-binding protein 2, mitochondrial OS=Homo sapiens GN=HINT2 PE=1 SV=1 - [HINT2_HUMAN]
P16403	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]
P16402	Histone H1.3 OS=Homo sapiens GN=HIST1H1D PE=1 SV=2 - [H13_HUMAN]
P10412	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2 - [H14_HUMAN]
P16401	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3 - [H15_HUMAN]
Q96KK5	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]
P16104	Histone H2A.x OS=Homo sapiens GN=H2AFX PE=1 SV=2 - [H2AX_HUMAN]
P0C0S5	Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2 - [H2AZ_HUMAN]
P58876	Histone H2B type 1-D OS=Homo sapiens GN=HIST1H2BD PE=1 SV=2 - [H2B1D_HUMAN]
P68431	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2 - [H31_HUMAN]
P62805	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]
B0UX83	HLA-B associated transcript 3 OS=Homo sapiens GN=BAT3 PE=4 SV=1 - [B0UX83_HUMAN]
P51610	Host cell factor 1 OS=Homo sapiens GN=HCFC1 PE=1 SV=2 - [HCFC1_HUMAN]
P50502	Hsc70-interacting protein OS=Homo sapiens GN=ST13 PE=1 SV=2 - [F10A1_HUMAN]
Q16543	Hsp90 co-chaperone Cdc37 OS=Homo sapiens GN=CDC37 PE=1 SV=1 - [CDC37_HUMAN]
O75146	Huntingtin-interacting protein 1-related protein OS=Homo sapiens GN=HIP1R PE=1 SV=2 - [HIP1R_HUMAN]
Q16836	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=1 SV=3 - [HCDH_HUMAN]
Q16775	Hydroxyacylglutathione hydrolase, mitochondrial OS=Homo sapiens GN=HAGH PE=1 SV=2 - [GLO2_HUMAN]
Q6YN16	Hydroxysteroid dehydrogenase-like protein 2 OS=Homo sapiens GN=HSDL2 PE=1 SV=1 - [HSDL2_HUMAN]
P00492	Hypoxanthine-guanine phosphoribosyltransferase OS=Homo sapiens GN=HPRT1 PE=1 SV=2 - [HPRT_HUMAN]
Q9Y4L1	Hypoxia up-regulated protein 1 OS=Homo sapiens GN=HYOU1 PE=1 SV=1 - [HYOU1_HUMAN]
P01880	Ig delta chain C region OS=Homo sapiens GN=IGHD PE=1 SV=2 - [IGHD_HUMAN]
P01854	Ig epsilon chain C region OS=Homo sapiens GN=IGHE PE=1 SV=1 - [IGHE_HUMAN]
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 - [IGHG1_HUMAN]
P01773	Ig heavy chain V-III region BUR OS=Homo sapiens PE=1 SV=1 - [HV312_HUMAN]
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1 - [LAC2_HUMAN]
Q9Y5U9	Immediate early response 3-interacting protein 1 OS=Homo sapiens GN=IER3IP1 PE=1 SV=1 - [IR3IP_HUMAN]
Q6WRI0	Immunoglobulin superfamily member 10 OS=Homo sapiens GN=IGSF10 PE=1 SV=1 - [IGS10_HUMAN]
P78318	Immunoglobulin-binding protein 1 OS=Homo sapiens GN=IGBP1 PE=1 SV=1 - [IGBP1_HUMAN]
Q14974	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2 - [IMB1_HUMAN]
O00410	Importin-5 OS=Homo sapiens GN=IPO5 PE=1 SV=4 - [IPO5_HUMAN]
O95373	Importin-7 OS=Homo sapiens GN=IPO7 PE=1 SV=1 - [IPO7_HUMAN]
Q96P70	Importin-9 OS=Homo sapiens GN=IPO9 PE=1 SV=3 - [IPO9_HUMAN]
Q14164	Inhibitor of nuclear factor kappa-B kinase subunit epsilon OS=Homo sapiens GN=IKBKE PE=1 SV=1 - [IKKE_HUMAN]
O43314	Inositol hexakisphosphate and diphosphoinositol-pentakisphosphate kinase 2 OS=Homo sapiens GN=PPIP5K2 PE=1 SV=3 - [VIP2_HUMAN]
Q9Y6M1	Insulin-like growth factor 2 mRNA-binding protein 2 OS=Homo sapiens GN=IGF2BP2 PE=1 SV=2 - [IF2B2_HUMAN]
P05556	Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2 - [ITB1_HUMAN]
Q14773	Intercellular adhesion molecule 4 OS=Homo sapiens GN=ICAM4 PE=2 SV=1 - [ICAM4_HUMAN]
P80217	Interferon-induced 35 kDa protein OS=Homo sapiens GN=IFI35 PE=1 SV=5 - [IN35_HUMAN]

Q722Y8	Interferon-induced very large GTPase 1 OS=Homo sapiens GN=GVINP1 PE=2 SV=2 - [GVIN1_HUMAN]
Q12905	Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2 - [ILF2_HUMAN]
Q5JU85	IQ motif and SEC7 domain-containing protein 2 OS=Homo sapiens GN=IQSEC2 PE=1 SV=1 - [IQEC2_HUMAN]
P48735	Isocitrate dehydrogenase [NADP], mitochondrial OS=Homo sapiens GN=IDH2 PE=1 SV=2 - [IDHP_HUMAN]
P53990	IST1 homolog OS=Homo sapiens GN=KIAA0174 PE=1 SV=1 - [IST1_HUMAN]
A0T124	Jumonji domain-containing 1 C splice variant OS=Homo sapiens GN=JMJD1C PE=2 SV=1 - [A0T124_HUMAN]
Q9Y624	Junctional adhesion molecule A OS=Homo sapiens GN=F11R PE=1 SV=1 - [JAM1_HUMAN]
Q5TFJ7	Karyopherin alpha 6 (Importin alpha 7) (Fragment) OS=Homo sapiens GN=KPNA6 PE=2 SV=1 - [Q5TFJ7_HUMAN]
P23276	Kell blood group glycoprotein OS=Homo sapiens GN=KEL PE=1 SV=2 - [KELL_HUMAN]
P13645	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 - [K1C10_HUMAN]
P08779	Keratin, type I cytoskeletal 16 OS=Homo sapiens GN=KRT16 PE=1 SV=4 - [K1C16_HUMAN]
Q04695	Keratin, type I cytoskeletal 17 OS=Homo sapiens GN=KRT17 PE=1 SV=2 - [K1C17_HUMAN]
P35527	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3 - [K1C9_HUMAN]
P04264	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6 - [K2C1_HUMAN]
P12035	Keratin, type II cytoskeletal 3 OS=Homo sapiens GN=KRT3 PE=1 SV=3 - [K2C3_HUMAN]
Q3SY84	Keratin, type II cytoskeletal 71 OS=Homo sapiens GN=KRT71 PE=1 SV=3 - [K2C71_HUMAN]
Q8N1N4	Keratin, type II cytoskeletal 78 OS=Homo sapiens GN=KRT78 PE=1 SV=2 - [K2C78_HUMAN]
Q17RZ5	Kinectin 1 (Kinesin receptor) OS=Homo sapiens GN=KTN1 PE=2 SV=1 - [Q17RZ5_HUMAN]
P33176	Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1 - [KINH_HUMAN]
Q9P2E2	Kinesin-like protein KIF17 OS=Homo sapiens GN=KIF17 PE=1 SV=3 - [KIF17_HUMAN]
Q9BW19	Kinesin-like protein KIFC1 OS=Homo sapiens GN=KIFC1 PE=1 SV=2 - [KIFC1_HUMAN]
Q9UEF7	Klotho OS=Homo sapiens GN=KL PE=1 SV=2 - [KLOT_HUMAN]
Q04760	Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 PE=1 SV=4 - [LGUL_HUMAN]
Q03252	Lamin-B2 OS=Homo sapiens GN=LMNB2 PE=1 SV=3 - [LMNB2_HUMAN]
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2 - [LAMB1_HUMAN]
Q13753	Laminin subunit gamma-2 OS=Homo sapiens GN=LAMC2 PE=1 SV=2 - [LAMC2_HUMAN]
Q9NS86	LanC-like protein 2 OS=Homo sapiens GN=LANCL2 PE=1 SV=1 - [LANC2_HUMAN]
Q6PKG0	La-related protein 1 OS=Homo sapiens GN=LARP1 PE=1 SV=2 - [LARP1_HUMAN]
Q01650	Large neutral amino acids transporter small subunit 1 OS=Homo sapiens GN=SLC7A5 PE=1 SV=2 - [LAT1_HUMAN]
O75387	Large neutral amino acids transporter small subunit 3 OS=Homo sapiens GN=SLC43A1 PE=2 SV=1 - [LAT3_HUMAN]
Q14696	LDLR chaperone MESD OS=Homo sapiens GN=MESDC2 PE=1 SV=2 - [MESD_HUMAN]
Q0VAK6	Leiomodin-3 OS=Homo sapiens GN=LMOD3 PE=2 SV=1 - [LMOD3_HUMAN]
O15243	Leptin receptor gene-related protein OS=Homo sapiens GN=LEPROT PE=1 SV=1 - [OBRG_HUMAN]
O95202	LETM1 and EF-hand domain-containing protein 1, mitochondrial OS=Homo sapiens GN=LETM1 PE=1 SV=1 - [LETM1_HUMAN]
Q8N9N7	Leucine-rich repeat-containing protein 57 OS=Homo sapiens GN=LRRC57 PE=1 SV=1 - [LRC57_HUMAN]
Q96AG4	Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRRC59 PE=1 SV=1 - [LRC59_HUMAN]
P07195	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2 - [LDHB_HUMAN]
E9PH51	L-lactate dehydrogenase OS=Homo sapiens GN=LDHA PE=3 SV=2 - [E9PH51_HUMAN]
Q496Y0	LON peptidase N-terminal domain and RING finger protein 3 OS=Homo sapiens GN=LONRF3 PE=1 SV=1 - [LONF3_HUMAN]
O95573	Long-chain-fatty-acid--CoA ligase 3 OS=Homo sapiens GN=ACSL3 PE=1 SV=3 - [ACSL3_HUMAN]
Q969J3	Loss of heterozygosity 12 chromosomal region 1 protein OS=Homo sapiens GN=LOH12CR1 PE=1 SV=1 - [L12R1_HUMAN]
O75096	Low-density lipoprotein receptor-related protein 4 OS=Homo sapiens GN=LRP4 PE=1 SV=4 - [LRP4_HUMAN]
P05455	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2 - [LA_HUMAN]

Q7Z4W1	L-xylulose reductase OS=Homo sapiens GN=DCXR PE=1 SV=2 - [DCXR_HUMAN]
Q6ZWT7	Lysophospholipid acyltransferase 2 OS=Homo sapiens GN=MBOAT2 PE=2 SV=2 - [MBOA2_HUMAN]
Q14108	Lysosome membrane protein 2 OS=Homo sapiens GN=SCARB2 PE=1 SV=2 - [SCRB2_HUMAN]
P11279	Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3 - [LAMP1_HUMAN]
P61626	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1 - [LYSC_HUMAN]
A6NFX1	Major facilitator superfamily domain-containing protein 2B OS=Homo sapiens GN=MFS2B PE=2 SV=3 - [MFS2B_HUMAN]
F5GY30	Major prion protein OS=Homo sapiens GN=PRNP PE=3 SV=1 - [F5GY30_HUMAN]
P40925	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4 - [MDHC_HUMAN]
P40926	Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3 - [MDHM_HUMAN]
Q9UM22	Mammalian ependymin-related protein 1 OS=Homo sapiens GN=EPDR1 PE=1 SV=2 - [EPDR1_HUMAN]
B8ZZG1	Membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6), isoform CRA_a OS=Homo sapiens GN=MPP6 PE=4 SV=1 - [B8ZZG1_HUMAN]
P51811	Membrane transport protein XK OS=Homo sapiens GN=XK PE=1 SV=5 - [XK_HUMAN]
O15173	Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRMC2 PE=1 SV=1 - [PGRC2_HUMAN]
P35240	Merlin OS=Homo sapiens GN=NF2 PE=1 SV=1 - [MERL_HUMAN]
P55145	Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=3 - [MANF_HUMAN]
P50579	Methionine aminopeptidase 2 OS=Homo sapiens GN=METAP2 PE=1 SV=1 - [AMPM2_HUMAN]
F5GXS7	Methionine aminopeptidase OS=Homo sapiens GN=METAP1 PE=3 SV=1 - [F5GXS7_HUMAN]
P22033	Methylmalonyl-CoA mutase, mitochondrial OS=Homo sapiens GN=MUT PE=1 SV=4 - [MUTA_HUMAN]
Q9H8H3	Methyltransferase-like protein 7A OS=Homo sapiens GN=METTL7A PE=1 SV=1 - [MET7A_HUMAN]
Q6NT16	MFS-type transporter C6orf192 OS=Homo sapiens GN=C6orf192 PE=2 SV=1 - [CF192_HUMAN]
Q8IY33	MICAL-like protein 2 OS=Homo sapiens GN=MICALL2 PE=1 SV=1 - [MILK2_HUMAN]
O14880	Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1 - [MGST3_HUMAN]
Q96PK2	Microtubule-actin cross-linking factor 1, isoform 4 OS=Homo sapiens GN=MACF1 PE=1 SV=2 - [MACF4_HUMAN]
Q15691	Microtubule-associated protein RP/EB family member 1 OS=Homo sapiens GN=MAPRE1 PE=1 SV=3 - [MARE1_HUMAN]
Q7Z6P5	Minichromosome maintenance complex component 3 (Fragment) OS=Homo sapiens GN=MCM3 PE=2 SV=1 - [Q7Z6P5_HUMAN]
Q8TCT9	Minor histocompatibility antigen H13 OS=Homo sapiens GN=HM13 PE=1 SV=1 - [HM13_HUMAN]
Q9BQT8	Mitochondrial 2-oxodicarboxylate carrier OS=Homo sapiens GN=SLC25A21 PE=1 SV=1 - [ODC_HUMAN]
Q02978	Mitochondrial 2-oxoglutarate/malate carrier protein OS=Homo sapiens GN=SLC25A11 PE=1 SV=3 - [M2OM_HUMAN]
Q9Y3D6	Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2 - [FIS1_HUMAN]
O43615	Mitochondrial import inner membrane translocase subunit TIM44 OS=Homo sapiens GN=TIMM44 PE=1 SV=2 - [TIM44_HUMAN]
O60220	Mitochondrial import inner membrane translocase subunit Tim8 A OS=Homo sapiens GN=TIMM8A PE=1 SV=1 - [TIM8A_HUMAN]
Q9NS69	Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=1 SV=3 - [TOM22_HUMAN]
O96008	Mitochondrial import receptor subunit TOM40 homolog OS=Homo sapiens GN=TOMM40 PE=1 SV=1 - [TOM40_HUMAN]
Q96T52	Mitochondrial inner membrane protease subunit 2 OS=Homo sapiens GN=IMMP2L PE=2 SV=1 - [IMP2L_HUMAN]
Q99797	Mitochondrial intermediate peptidase OS=Homo sapiens GN=MIPEP PE=1 SV=2 - [MIPEP_HUMAN]
Q9NYZ2	Mitoferrin-1 OS=Homo sapiens GN=SLC25A37 PE=1 SV=2 - [MFRN1_HUMAN]
Q9ULH7	MKL/myocardin-like protein 2 OS=Homo sapiens GN=MKL2 PE=1 SV=3 - [MKL2_HUMAN]
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3 - [MOES_HUMAN]
P53985	Monocarboxylate transporter 1 OS=Homo sapiens GN=SLC16A1 PE=1 SV=3 - [MOT1_HUMAN]
Q9Y3A3	Mps one binder kinase activator-like 3 OS=Homo sapiens GN=MOBK3 PE=1 SV=1 - [MOBL3_HUMAN]
Q96T58	Msx2-interacting protein OS=Homo sapiens GN=SPEN PE=1 SV=1 - [MINT_HUMAN]
O15439	Multidrug resistance-associated protein 4 OS=Homo sapiens GN=ABCC4 PE=1 SV=3 - [MRP4_HUMAN]
Q7Z7M0	Multiple epidermal growth factor-like domains protein 8 OS=Homo sapiens GN=MEGF8 PE=1 SV=2 - [MEGF8_HUMAN]

Q9UNW1	Multiple inositol polyphosphate phosphatase 1 OS=Homo sapiens GN=MINPP1 PE=1 SV=1 - [MINP1_HUMAN]
P05976	Myosin light chain 1/3, skeletal muscle isoform OS=Homo sapiens GN=MYL1 PE=1 SV=3 - [MYL1_HUMAN]
P12829	Myosin light chain 4 OS=Homo sapiens GN=MYL4 PE=1 SV=3 - [MYL4_HUMAN]
P60660	Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2 - [MYL6_HUMAN]
P19105	Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=1 SV=2 - [ML12A_HUMAN]
Q3MIV8	Myosin, heavy chain 11, smooth muscle OS=Homo sapiens GN=MYH11 PE=2 SV=1 - [Q3MIV8_HUMAN]
P35580	Myosin-10 OS=Homo sapiens GN=MYH10 PE=1 SV=3 - [MYH10_HUMAN]
Q7Z406	Myosin-14 OS=Homo sapiens GN=MYH14 PE=1 SV=2 - [MYH14_HUMAN]
P35579	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4 - [MYH9_HUMAN]
O94832	Myosin-Id OS=Homo sapiens GN=MYO1D PE=1 SV=2 - [MYO1D_HUMAN]
Q13459	Myosin-IXb OS=Homo sapiens GN=MYO9B PE=1 SV=3 - [MYO9B_HUMAN]
Q9NQX4	Myosin-Vc OS=Homo sapiens GN=MYO5C PE=1 SV=2 - [MYO5C_HUMAN]
Q5FWF5	N-acetyltransferase ESCO1 OS=Homo sapiens GN=ESCO1 PE=1 SV=3 - [ESCO1_HUMAN]
Q13423	NAD(P) transhydrogenase, mitochondrial OS=Homo sapiens GN=NNT PE=1 SV=3 - [NNTM_HUMAN]
B2RUY3	NADH dehydrogenase (Ubiquinone) 1 beta subcomplex, 4, 15kDa OS=Homo sapiens GN=NDUFB4 PE=4 SV=1 - [B2RUY3_HUMAN]
Q9P0J0	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Homo sapiens GN=NDUFA13 PE=1 SV=3 - [NDUAD_HUMAN]
O00483	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=Homo sapiens GN=NDUFA4 PE=1 SV=1 - [NDUA4_HUMAN]
P56556	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=Homo sapiens GN=NDUFA6 PE=1 SV=3 - [NDUA6_HUMAN]
O95182	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=Homo sapiens GN=NDUFA7 PE=1 SV=3 - [NDUA7_HUMAN]
O96000	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3 - [NDUBA_HUMAN]
Q9NX14	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 11, mitochondrial OS=Homo sapiens GN=NDUFB11 PE=1 SV=1 - [NDUBB_HUMAN]
P17568	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Homo sapiens GN=NDUFB7 PE=1 SV=4 - [NDUB7_HUMAN]
E9PRB2	NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Homo sapiens GN=NDUFC2 PE=3 SV=1 - [E9PRB2_HUMAN]
P19404	NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Homo sapiens GN=NDUFV2 PE=1 SV=2 - [NDUV2_HUMAN]
P56181	NADH dehydrogenase [ubiquinone] flavoprotein 3, mitochondrial OS=Homo sapiens GN=NDUFV3 PE=1 SV=2 - [NDUV3_HUMAN]
O75489	NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Homo sapiens GN=NDUFS3 PE=1 SV=1 - [NDUS3_HUMAN]
O43181	NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial OS=Homo sapiens GN=NDUFS4 PE=1 SV=1 - [NDUS4_HUMAN]
O75380	NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Homo sapiens GN=NDUFS6 PE=1 SV=1 - [NDUS6_HUMAN]
Q9UHQ9	NADH-cytochrome b5 reductase 1 OS=Homo sapiens GN=CYB5R1 PE=1 SV=1 - [NB5R1_HUMAN]
P00387	NADH-cytochrome b5 reductase 3 OS=Homo sapiens GN=CYB5R3 PE=1 SV=3 - [NB5R3_HUMAN]
Q13765	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=1 SV=1 - [NACA_HUMAN]
P20929	Nebulin OS=Homo sapiens GN=NEB PE=1 SV=4 - [NEBU_HUMAN]
Q15843	NEDD8 OS=Homo sapiens GN=NEDD8 PE=1 SV=1 - [NEDD8_HUMAN]
Q8WXH0	Nesprin-2 OS=Homo sapiens GN=SYNE2 PE=1 SV=3 - [SYNE2_HUMAN]
Q9UMX5	Neudesin OS=Homo sapiens GN=NENF PE=1 SV=1 - [NENF_HUMAN]
A2RRP1	Neuroblastoma-amplified sequence OS=Homo sapiens GN=NBAS PE=1 SV=2 - [NBAS_HUMAN]
Q9HB89	Neuromedin-U receptor 1 OS=Homo sapiens GN=NMUR1 PE=2 SV=1 - [NMUR1_HUMAN]
B2RAL7	Neuroplastin OS=Homo sapiens GN=NPTN PE=2 SV=1 - [B2RAL7_HUMAN]
Q14697	Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3 - [GANAB_HUMAN]
Q15758	Neutral amino acid transporter B(0) OS=Homo sapiens GN=SLC1A5 PE=1 SV=2 - [AAAT_HUMAN]
Q6PIU2	Neutral cholesterol ester hydrolase 1 OS=Homo sapiens GN=NCEH1 PE=1 SV=3 - [NCEH1_HUMAN]
P59665	Neutrophil defensin 1 OS=Homo sapiens GN=DEFA1 PE=1 SV=1 - [DEF1_HUMAN]
Q5BKU3	NIN protein OS=Homo sapiens GN=NIN PE=2 SV=1 - [Q5BKU3_HUMAN]

Q6KC79	Nipped-B-like protein OS=Homo sapiens GN=NIPBL PE=1 SV=2 - [NIPBL_HUMAN]
Q9NPP4	NLR family CARD domain-containing protein 4 OS=Homo sapiens GN=NLRC4 PE=1 SV=2 - [NLRC4_HUMAN]
Q49AL4	NMNAT3 protein OS=Homo sapiens GN=NMNAT3 PE=2 SV=1 - [Q49AL4_HUMAN]
P69849	Nodal modulator 3 OS=Homo sapiens GN=NOMO3 PE=2 SV=2 - [NOMO3_HUMAN]
B7ZBA8	Novel protein OS=Homo sapiens GN=CCDC154 PE=4 SV=1 - [B7ZBA8_HUMAN]
Q5JQQ4	Novel protein OS=Homo sapiens GN=RP11-529I10.4 PE=2 SV=1 - [Q5JQQ4_HUMAN]
Q9UNZ2	NSFL1 cofactor p47 OS=Homo sapiens GN=NSFL1C PE=1 SV=2 - [NSF1C_HUMAN]
B1AKN7	Nuclear factor 1 OS=Homo sapiens GN=NFA1 PE=3 SV=1 - [B1AKN7_HUMAN]
Q5VW31	Nuclear factor I/B OS=Homo sapiens GN=NFIB PE=2 SV=1 - [Q5VW31_HUMAN]
Q16649	Nuclear factor interleukin-3-regulated protein OS=Homo sapiens GN=NFIL3 PE=1 SV=2 - [NFIL3_HUMAN]
B5B2M6	Nuclear factor of activated T-cells c1 isoform IA-deltaIX OS=Homo sapiens GN=NFATC1 PE=2 SV=1 - [B5B2M6_HUMAN]
Q9Y266	Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1 - [NUDC_HUMAN]
Q14980	Nuclear mitotic apparatus protein 1 OS=Homo sapiens GN=NUMA1 PE=1 SV=2 - [NUMA1_HUMAN]
Q14686	Nuclear receptor coactivator 6 OS=Homo sapiens GN=NCOA6 PE=1 SV=3 - [NCOA6_HUMAN]
P61970	Nuclear transport factor 2 OS=Homo sapiens GN=NUTF2 PE=1 SV=1 - [NUTF2_HUMAN]
Q9H1E3	Nuclear ubiquitous casein and cyclin-dependent kinases substrate OS=Homo sapiens GN=NUCKS1 PE=1 SV=1 - [NUCKS_HUMAN]
P67809	Nuclease-sensitive element-binding protein 1 OS=Homo sapiens GN=YBX1 PE=1 SV=3 - [YBOX1_HUMAN]
Q02818	Nucleobindin-1 OS=Homo sapiens GN=NUCB1 PE=1 SV=4 - [NUCB1_HUMAN]
P19338	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3 - [NUCL_HUMAN]
Q9BYG9	Nucleophosmin (Nucleolar phosphoprotein B23, numatrin) OS=Homo sapiens GN=NPM1 PE=2 SV=1 - [Q9BYG9_HUMAN]
P12270	Nucleoprotein TPR OS=Homo sapiens GN=TPR PE=1 SV=3 - [TPR_HUMAN]
B3KV44	Nucleosome assembly protein 1-like 1, isoform CRA_c OS=Homo sapiens GN=NAP1L1 PE=2 SV=1 - [B3KV44_HUMAN]
Q4LE47	NUP153 variant protein (Fragment) OS=Homo sapiens GN=NUP153 variant protein PE=2 SV=1 - [Q4LE47_HUMAN]
Q9NTK5	Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=1 SV=2 - [OLA1_HUMAN]
Q8NH19	Olfactory receptor 10AG1 OS=Homo sapiens GN=OR10AG1 PE=2 SV=1 - [O10AG_HUMAN]
Q8NH49	Olfactory receptor 4X1 OS=Homo sapiens GN=OR4X1 PE=2 SV=1 - [OR4X1_HUMAN]
O95007	Olfactory receptor 6B1 OS=Homo sapiens GN=OR6B1 PE=2 SV=1 - [OR6B1_HUMAN]
Q96CV9	Optineurin OS=Homo sapiens GN=OPTN PE=1 SV=2 - [OPTN_HUMAN]
P04181	Ornithine aminotransferase, mitochondrial OS=Homo sapiens GN=OAT PE=1 SV=1 - [OAT_HUMAN]
Q8N6M0	OTU domain-containing protein 6B OS=Homo sapiens GN=OTUD6B PE=1 SV=1 - [OTU6B_HUMAN]
Q8TE49	OTU domain-containing protein 7A OS=Homo sapiens GN=OTUD7A PE=1 SV=1 - [OTU7A_HUMAN]
B4DTR3	Oxysterol-binding protein OS=Homo sapiens GN=OSBP2 PE=2 SV=1 - [B4DTR3_HUMAN]
F5H7T2	Oxysterol-binding protein OS=Homo sapiens GN=OSBPL8 PE=3 SV=1 - [F5H7T2_HUMAN]
P51582	P2Y purinoceptor 4 OS=Homo sapiens GN=P2RY4 PE=1 SV=1 - [P2RY4_HUMAN]
Q5T0S4	Palmitoyl-protein thioesterase 1 (Fragment) OS=Homo sapiens GN=PPT1 PE=2 SV=1 - [Q5T0S4_HUMAN]
Q96RD7	Pannexin-1 OS=Homo sapiens GN=PANX1 PE=1 SV=4 - [PANX1_HUMAN]
Q8TD49	PAP1 OS=Homo sapiens GN=PAP1 PE=2 SV=1 - [Q8TD49_HUMAN]
Q9BRP8	Partner of Y14 and mago OS=Homo sapiens GN=WIBG PE=1 SV=1 - [WIBG_HUMAN]
Q8NEN9	PDZ domain-containing protein 8 OS=Homo sapiens GN=PDZD8 PE=1 SV=1 - [PDZD8_HUMAN]
O75127	Pentatricopeptide repeat-containing protein 1 OS=Homo sapiens GN=PTCD1 PE=1 SV=2 - [PTCD1_HUMAN]
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2 - [PPIA_HUMAN]
P23284	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2 - [PPIB_HUMAN]
Q00688	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1 - [FKBP3_HUMAN]

Q02790	Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3 - [FKBP4_HUMAN]
Q14318	Peptidyl-prolyl cis-trans isomerase FKBP8 OS=Homo sapiens GN=FKBP8 PE=1 SV=2 - [FKBP8_HUMAN]
Q9Y237	Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 OS=Homo sapiens GN=PIN4 PE=1 SV=1 - [PIN4_HUMAN]
Q9Y3E5	Peptidyl-tRNA hydrolase 2, mitochondrial OS=Homo sapiens GN=PTRH2 PE=1 SV=1 - [PTH2_HUMAN]
Q9BXM0	Periaxin OS=Homo sapiens GN=PRX PE=1 SV=2 - [PRAX_HUMAN]
O60664	Perilipin-3 OS=Homo sapiens GN=PLIN3 PE=1 SV=3 - [PLIN3_HUMAN]
O60437	Periplakin OS=Homo sapiens GN=PPL PE=1 SV=4 - [PEPL_HUMAN]
Q06830	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1 - [PRDX1_HUMAN]
P32119	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5 - [PRDX2_HUMAN]
P30044	Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=4 - [PRDX5_HUMAN]
P30041	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3 - [PRDX6_HUMAN]
O75192	Peroxisomal membrane protein 11A OS=Homo sapiens GN=PEX11A PE=1 SV=1 - [PX11A_HUMAN]
Q8WW12	PEST proteolytic signal-containing nuclear protein OS=Homo sapiens GN=PCNP PE=1 SV=2 - [PCNP_HUMAN]
Q5JWZ0	PHD finger protein 20 (Fragment) OS=Homo sapiens GN=PHF20 PE=2 SV=1 - [Q5JWZ0_HUMAN]
Q9Y285	Phenylalanyl-tRNA synthetase alpha chain OS=Homo sapiens GN=FARSA PE=1 SV=3 - [SYFA_HUMAN]
Q9NSD9	Phenylalanyl-tRNA synthetase beta chain OS=Homo sapiens GN=FARSB PE=1 SV=3 - [SYFB_HUMAN]
B7ZL07	PHKA1 protein OS=Homo sapiens GN=PHKA1 PE=2 SV=1 - [B7ZL07_HUMAN]
Q13371	Phosducin-like protein OS=Homo sapiens GN=PDCL PE=1 SV=3 - [PHLP_HUMAN]
Q00325	Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2 - [MPCP_HUMAN]
Q92539	Phosphatidate phosphatase LPIN2 OS=Homo sapiens GN=LPIN2 PE=1 SV=1 - [LPIN2_HUMAN]
Q9BTU6	Phosphatidylinositol 4-kinase type 2-alpha OS=Homo sapiens GN=PI4K2A PE=1 SV=1 - [P4K2A_HUMAN]
Q92835	Phosphatidylinositol-3,4,5-trisphosphate 5-phosphatase 1 OS=Homo sapiens GN=INPP5D PE=1 SV=2 - [SHIP1_HUMAN]
P48426	Phosphatidylinositol-5-phosphate 4-kinase type-2 alpha OS=Homo sapiens GN=PIP4K2A PE=1 SV=2 - [PI42A_HUMAN]
Q13492	Phosphatidylinositol-binding clathrin assembly protein OS=Homo sapiens GN=PICALM PE=1 SV=2 - [PICAL_HUMAN]
Q16822	Phosphoenolpyruvate carboxykinase [GTP], mitochondrial OS=Homo sapiens GN=PCK2 PE=1 SV=3 - [PCKGM_HUMAN]
B7Z7A9	Phosphoglycerate kinase OS=Homo sapiens GN=PGK1 PE=2 SV=1 - [B7Z7A9_HUMAN]
P15259	Phosphoglycerate mutase 2 OS=Homo sapiens GN=PGAM2 PE=1 SV=3 - [PGAM2_HUMAN]
Q99570	Phosphoinositide 3-kinase regulatory subunit 4 OS=Homo sapiens GN=PIK3R4 PE=1 SV=3 - [PI3R4_HUMAN]
Q9Y263	Phospholipase A-2-activating protein OS=Homo sapiens GN=PLAA PE=1 SV=2 - [PLAP_HUMAN]
P36969	Phospholipid hydroperoxide glutathione peroxidase, mitochondrial OS=Homo sapiens GN=GPX4 PE=1 SV=3 - [GPX4_HUMAN]
Q6ZR73	Phospholipid scramblase 4, isoform CRA_a OS=Homo sapiens GN=PLSCR4 PE=2 SV=1 - [Q6ZR73_HUMAN]
O60256	Phosphoribosyl pyrophosphate synthase-associated protein 2 OS=Homo sapiens GN=PRPSAP2 PE=1 SV=1 - [KPRB_HUMAN]
Q93100	Phosphorylase b kinase regulatory subunit beta OS=Homo sapiens GN=PHKB PE=1 SV=3 - [KPBB_HUMAN]
Q9GZP4	PITH domain-containing protein 1 OS=Homo sapiens GN=PITHD1 PE=1 SV=1 - [PITH1_HUMAN]
P20020	Plasma membrane calcium-transporting ATPase 1 OS=Homo sapiens GN=ATP2B1 PE=1 SV=3 - [AT2B1_HUMAN]
Q16720	Plasma membrane calcium-transporting ATPase 3 OS=Homo sapiens GN=ATP2B3 PE=1 SV=3 - [AT2B3_HUMAN]
P23634	Plasma membrane calcium-transporting ATPase 4 OS=Homo sapiens GN=ATP2B4 PE=1 SV=2 - [AT2B4_HUMAN]
Q8NC51	Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 PE=1 SV=2 - [PAIRB_HUMAN]
Q14651	Plastin-1 OS=Homo sapiens GN=PLS1 PE=1 SV=2 - [PLSI_HUMAN]
P13796	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6 - [PLSL_HUMAN]
Q8IVE3	Pleckstrin homology domain-containing family H member 2 OS=Homo sapiens GN=PLEKHH2 PE=2 SV=2 - [PKHH2_HUMAN]
Q9NYT0	Pleckstrin-2 OS=Homo sapiens GN=PLEK2 PE=1 SV=1 - [PLEK2_HUMAN]
B1ANR0	Poly(A) binding protein, cytoplasmic 4 (Inducible form) OS=Homo sapiens GN=PABPC4 PE=4 SV=1 - [B1ANR0_HUMAN]

Q68Y55	Poly(RC) binding protein 2 OS=Homo sapiens GN=PCBP2 PE=2 SV=1 - [Q68Y55_HUMAN]
Q15365	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]
P11940	Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 PE=1 SV=2 - [PABP1_HUMAN]
Q9Y257	Polymerase delta-interacting protein 2 OS=Homo sapiens GN=POLDIP2 PE=1 SV=1 - [PDIP2_HUMAN]
P26599	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1 - [PTBP1_HUMAN]
B2RU33	POTE ankyrin domain family member C OS=Homo sapiens GN=POTEC PE=2 SV=2 - [POTEC_HUMAN]
Q9UHV9	Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1 - [PFD2_HUMAN]
Q9NQP4	Prefoldin subunit 4 OS=Homo sapiens GN=PFDN4 PE=1 SV=1 - [PFD4_HUMAN]
Q8N945	PRELI domain-containing protein 2 OS=Homo sapiens GN=PRELID2 PE=2 SV=1 - [PRLD2_HUMAN]
Q5VTL8	Pre-mRNA-splicing factor 38B OS=Homo sapiens GN=PRPF38B PE=1 SV=1 - [PR38B_HUMAN]
Q9BUB1	PRKAR2A protein OS=Homo sapiens GN=PRKAR2A PE=2 SV=1 - [Q9BUB1_HUMAN]
P07602	Proactivator polypeptide OS=Homo sapiens GN=PSAP PE=1 SV=2 - [SAP_HUMAN]
Q9BUQ8	Probable ATP-dependent RNA helicase DDX23 OS=Homo sapiens GN=DDX23 PE=1 SV=3 - [DDX23_HUMAN]
Q9Y235	Probable C->U-editing enzyme APOBEC-2 OS=Homo sapiens GN=APOBEC2 PE=1 SV=1 - [ABEC2_HUMAN]
Q9HD20	Probable cation-transporting ATPase 13A1 OS=Homo sapiens GN=ATP13A1 PE=1 SV=2 - [AT131_HUMAN]
Q9Y4D8	Probable E3 ubiquitin-protein ligase C12orf51 OS=Homo sapiens GN=C12orf51 PE=1 SV=5 - [K0614_HUMAN]
P51531	Probable global transcription activator SNF2L2 OS=Homo sapiens GN=SMARCA2 PE=1 SV=2 - [SMCA2_HUMAN]
Q13395	Probable methyltransferase TARBP1 OS=Homo sapiens GN=TARBP1 PE=1 SV=1 - [TARB1_HUMAN]
Q9UHF3	Probable N-acetyltransferase 8B OS=Homo sapiens GN=NAT8B PE=2 SV=1 - [NAT8B_HUMAN]
Q9H3G5	Probable serine carboxypeptidase CPVL OS=Homo sapiens GN=CPVL PE=1 SV=2 - [CPVL_HUMAN]
Q93008	Probable ubiquitin carboxyl-terminal hydrolase FAF-X OS=Homo sapiens GN=USP9X PE=1 SV=3 - [USP9X_HUMAN]
O00507	Probable ubiquitin carboxyl-terminal hydrolase FAF-Y OS=Homo sapiens GN=USP9Y PE=1 SV=2 - [USP9Y_HUMAN]
Q8NBJ5	Procollagen galactosyltransferase 1 OS=Homo sapiens GN=GLT25D1 PE=1 SV=1 - [GT251_HUMAN]
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2 - [PROF1_HUMAN]
Q6ZVX9	Progesterin and adipoQ receptor family member 9 OS=Homo sapiens GN=PAQR9 PE=2 SV=1 - [PAQR9_HUMAN]
O14737	Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3 - [PDCD5_HUMAN]
Q99623	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2 - [PHB2_HUMAN]
Q9UQ80	Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=3 - [PA2G4_HUMAN]
Q9BRP4	Proteasomal ATPase-associated factor 1 OS=Homo sapiens GN=PAAF1 PE=1 SV=2 - [PAAF1_HUMAN]
Q16186	Proteasomal ubiquitin receptor ADRM1 OS=Homo sapiens GN=ADRM1 PE=1 SV=2 - [ADRM1_HUMAN]
A8K3Z3	Proteasome (Prosome, macropain) 26S subunit, ATPase, 5, isoform CRA_b OS=Homo sapiens GN=PSMC5 PE=2 SV=1 - [A8K3Z3_HUMAN]
B1AJY5	Proteasome (Prosome, macropain) 26S subunit, non-ATPase, 10 OS=Homo sapiens GN=PSMD10 PE=4 SV=1 - [B1AJY5_HUMAN]
Q06323	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1 - [PSME1_HUMAN]
Q9UL46	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=4 - [PSME2_HUMAN]
P25786	Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1 PE=1 SV=1 - [PSA1_HUMAN]
P25787	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2 - [PSA2_HUMAN]
P25788	Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 PE=1 SV=2 - [PSA3_HUMAN]
P25789	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1 - [PSA4_HUMAN]
P28066	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3 - [PSA5_HUMAN]
P60900	Proteasome subunit alpha type-6 OS=Homo sapiens GN=PSMA6 PE=1 SV=1 - [PSA6_HUMAN]
O14818	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1 - [PSA7_HUMAN]
P20618	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2 - [PSB1_HUMAN]
P49721	Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1 - [PSB2_HUMAN]

P49720	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2 - [PSB3_HUMAN]
P28070	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4 - [PSB4_HUMAN]
P28074	Proteasome subunit beta type-5 OS=Homo sapiens GN=PSMB5 PE=1 SV=3 - [PSB5_HUMAN]
P28072	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4 - [PSB6_HUMAN]
Q99436	Proteasome subunit beta type-7 OS=Homo sapiens GN=PSMB7 PE=1 SV=1 - [PSB7_HUMAN]
Q5VYK3	Proteasome-associated protein ECM29 homolog OS=Homo sapiens GN=ECM29 PE=1 SV=2 - [ECM29_HUMAN]
P11171	Protein 4.1 OS=Homo sapiens GN=EPB41 PE=1 SV=4 - [41_HUMAN]
O14744	Protein arginine N-methyltransferase 5 OS=Homo sapiens GN=PRMT5 PE=1 SV=4 - [ANM5_HUMAN]
Q9UKV8	Protein argonaute-2 OS=Homo sapiens GN=EIF2C2 PE=1 SV=3 - [AGO2_HUMAN]
Q9Y2B0	Protein canopy homolog 2 OS=Homo sapiens GN=CNPY2 PE=1 SV=1 - [CNPY2_HUMAN]
Q9BT09	Protein canopy homolog 3 OS=Homo sapiens GN=CNPY3 PE=1 SV=1 - [CNPY3_HUMAN]
Q9UKY7	Protein CDV3 homolog OS=Homo sapiens GN=CDV3 PE=1 SV=1 - [CDV3_HUMAN]
Q8TCG1	Protein CIP2A OS=Homo sapiens GN=KIAA1524 PE=1 SV=2 - [CIP2A_HUMAN]
Q5TDH0	Protein DDI1 homolog 2 OS=Homo sapiens GN=DDI2 PE=1 SV=1 - [DDI2_HUMAN]
P13667	Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2 - [PDIA4_HUMAN]
P07237	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3 - [PDIA1_HUMAN]
Q99497	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2 - [PARK7_HUMAN]
Q8WYP5	Protein ELYS OS=Homo sapiens GN=AHCTF1 PE=1 SV=3 - [ELYS_HUMAN]
P49257	Protein ERGIC-53 OS=Homo sapiens GN=LMAN1 PE=1 SV=2 - [LMAN1_HUMAN]
Q9NRY5	Protein FAM114A2 OS=Homo sapiens GN=FAM114A2 PE=1 SV=4 - [F1142_HUMAN]
Q9BUT9	Protein FAM195A OS=Homo sapiens GN=FAM195A PE=1 SV=2 - [F195A_HUMAN]
Q6UWF7	Protein FAM55D OS=Homo sapiens GN=FAM55D PE=2 SV=1 - [FA55D_HUMAN]
Q96KN4	Protein FAM84A OS=Homo sapiens GN=FAM84A PE=1 SV=2 - [FA84A_HUMAN]
Q68CZ1	Protein fantom OS=Homo sapiens GN=RPGRIP1L PE=1 SV=2 - [FTM_HUMAN]
O94992	Protein HEXIM1 OS=Homo sapiens GN=HEXIM1 PE=1 SV=1 - [HEXI1_HUMAN]
B0QYG7	Protein kinase C and casein kinase substrate in neurons 2 (Fragment) OS=Homo sapiens GN=PAC SIN2 PE=4 SV=1 - [B0QYG7_HUMAN]
Q96RT1	Protein LAP2 OS=Homo sapiens GN=ERBB2IP PE=1 SV=2 - [LAP2_HUMAN]
Q9NUP9	Protein lin-7 homolog C OS=Homo sapiens GN=LIN7C PE=1 SV=1 - [LIN7C_HUMAN]
P12524	Protein L-Myc-1 OS=Homo sapiens GN=MYCL1 PE=1 SV=2 - [MYCL1_HUMAN]
Q9BZQ8	Protein Niban OS=Homo sapiens GN=FAM129A PE=1 SV=1 - [NIBAN_HUMAN]
Q9BRX2	Protein pelota homolog OS=Homo sapiens GN=PELO PE=1 SV=2 - [PELO_HUMAN]
O75688	Protein phosphatase 1B OS=Homo sapiens GN=PPM1B PE=1 SV=1 - [PPM1B_HUMAN]
Q59GB2	Protein phosphatase 1G variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q59GB2_HUMAN]
Q9Y570	Protein phosphatase methylesterase 1 OS=Homo sapiens GN=PPME1 PE=1 SV=3 - [PPME1_HUMAN]
Q92508	Protein PIEZO1 OS=Homo sapiens GN=FAM38A PE=1 SV=4 - [PIEZ1_HUMAN]
Q9P258	Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2 - [RCC2_HUMAN]
O15258	Protein RER1 OS=Homo sapiens GN=RER1 PE=1 SV=1 - [RER1_HUMAN]
P06703	Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1 - [S10A6_HUMAN]
P05109	Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1 - [S10A8_HUMAN]
A2A3K4	Protein tyrosine phosphatase domain-containing protein 1 OS=Homo sapiens GN=PTPDC1 PE=2 SV=1 - [PTPC1_HUMAN]
Q8IWX7	Protein unc-45 homolog B OS=Homo sapiens GN=UNC45B PE=2 SV=1 - [UN45B_HUMAN]
Q08AM6	Protein VAC14 homolog OS=Homo sapiens GN=VAC14 PE=1 SV=1 - [VAC14_HUMAN]
O75695	Protein XRP2 OS=Homo sapiens GN=RP2 PE=1 SV=4 - [XRP2_HUMAN]

Q9Y5H8	Protocadherin alpha-3 OS=Homo sapiens GN=PCDHA3 PE=1 SV=1 - [PCDA3_HUMAN]
Q9Y5G9	Protocadherin gamma-A4 OS=Homo sapiens GN=PCDHGA4 PE=2 SV=1 - [PCDG4_HUMAN]
P61457	Pterin-4-alpha-carbinolamine dehydratase OS=Homo sapiens GN=PCBD1 PE=1 SV=2 - [PHS_HUMAN]
A2VCM6	PTMS protein (Fragment) OS=Homo sapiens GN=PTMS PE=2 SV=1 - [A2VCM6_HUMAN]
P00491	Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2 - [PNPH_HUMAN]
Q5NE16	Putative cathepsin L-like protein 3 OS=Homo sapiens GN=CTSL3 PE=5 SV=1 - [CATL3_HUMAN]
Q9Y315	Putative deoxyribose-phosphate aldolase OS=Homo sapiens GN=DERA PE=1 SV=2 - [DEOC_HUMAN]
A6NDK9	Putative golgin subfamily A member 6C OS=Homo sapiens GN=GOLGA6C PE=5 SV=1 - [GOG6C_HUMAN]
Q5TCQ7	Putative homeodomain transcription factor 1 (Fragment) OS=Homo sapiens GN=PHTF1 PE=2 SV=1 - [Q5TCQ7_HUMAN]
A6ND91	Putative L-aspartate dehydrogenase OS=Homo sapiens GN=ASPDH PE=2 SV=2 - [ASPD_HUMAN]
P59922	Putative olfactory receptor 2B8 OS=Homo sapiens GN=OR2B8P PE=5 SV=1 - [OR2B8_HUMAN]
A2A3N6	Putative PIP5K1A and PSMD4-like protein OS=Homo sapiens GN=PIPSL PE=5 SV=1 - [PIPSL_HUMAN]
Q495Y7	Putative speedy protein E7 OS=Homo sapiens GN=SPDYE7P PE=5 SV=1 - [SPDE7_HUMAN]
Q5QP74	Putative uncharacterized protein C10orf115 OS=Homo sapiens GN=C10orf115 PE=2 SV=2 - [CJ115_HUMAN]
Q6ZS72	Putative uncharacterized protein C19orf35 OS=Homo sapiens GN=C19orf35 PE=2 SV=1 - [CS035_HUMAN]
Q5SQ13	Putative uncharacterized protein C9orf141 OS=Homo sapiens GN=C9orf141 PE=2 SV=1 - [CI141_HUMAN]
Q5HYI5	Putative uncharacterized protein DKFZp313C1541 OS=Homo sapiens GN=DKFZp313C1541 PE=2 SV=1 - [Q5HYI5_HUMAN]
Q9Y427	Putative uncharacterized protein DKFZp586K2222 OS=Homo sapiens GN=DKFZp586K2222 PE=2 SV=2 - [Q9Y427_HUMAN]
Q6MZR2	Putative uncharacterized protein DKFZp779J2370 OS=Homo sapiens GN=DKFZp779J2370 PE=2 SV=1 - [Q6MZR2_HUMAN]
Q8N9K3	Putative uncharacterized protein FLJ37004 OS=Homo sapiens PE=2 SV=1 - [YA015_HUMAN]
Q6ZTK2	Putative uncharacterized protein LOC400499 OS=Homo sapiens PE=2 SV=1 - [YP015_HUMAN]
Q53R32	Putative uncharacterized protein RPIA (Fragment) OS=Homo sapiens GN=RPIA PE=2 SV=1 - [Q53R32_HUMAN]
Q4ZG20	Putative uncharacterized protein TTN OS=Homo sapiens GN=TTN PE=2 SV=1 - [Q4ZG20_HUMAN]
P30613	Pyruvate kinase isozymes R/L OS=Homo sapiens GN=PKLR PE=1 SV=2 - [KPYR_HUMAN]
Q9Y4A3	R33590_1 OS=Homo sapiens GN=CREB3L3 PE=2 SV=1 - [Q9Y4A3_HUMAN]
P50395	Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GDI2 PE=1 SV=2 - [GDIB_HUMAN]
Q8WVF3	RAB28 protein OS=Homo sapiens GN=RAB28 PE=2 SV=1 - [Q8WVF3_HUMAN]
Q9H2M9	Rab3 GTPase-activating protein non-catalytic subunit OS=Homo sapiens GN=RAB3GAP2 PE=1 SV=1 - [RBGPR_HUMAN]
Q96NA2	Rab-interacting lysosomal protein OS=Homo sapiens GN=RILP PE=1 SV=1 - [RILP_HUMAN]
C6K8I4	Rab-like GTP-binding protein 1C OS=Homo sapiens GN=RBEL1 PE=2 SV=1 - [C6K8I4_HUMAN]
Q9H0H5	Rac GTPase-activating protein 1 OS=Homo sapiens GN=RACGAP1 PE=1 SV=1 - [RGAP1_HUMAN]
P35241	Radixin OS=Homo sapiens GN=RDX PE=1 SV=1 - [RADI_HUMAN]
Q9UHA4	Ragulator complex protein LAMTOR3 OS=Homo sapiens GN=LAMTOR3 PE=1 SV=1 - [LTOR3_HUMAN]
Q96D71	RalBP1-associated Eps domain-containing protein 1 OS=Homo sapiens GN=REPS1 PE=1 SV=3 - [REPS1_HUMAN]
P46060	Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1 - [RAGP1_HUMAN]
P43487	Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1 - [RANG_HUMAN]
P63000	Ras-related C3 botulinum toxin substrate 1 OS=Homo sapiens GN=RAC1 PE=1 SV=1 - [RAC1_HUMAN]
P61026	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1 - [RAB10_HUMAN]
P62491	Ras-related protein Rab-11A OS=Homo sapiens GN=RAB11A PE=1 SV=3 - [RB11A_HUMAN]
P51153	Ras-related protein Rab-13 OS=Homo sapiens GN=RAB13 PE=1 SV=1 - [RAB13_HUMAN]
P61106	Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4 - [RAB14_HUMAN]
Q9NP72	Ras-related protein Rab-18 OS=Homo sapiens GN=RAB18 PE=1 SV=1 - [RAB18_HUMAN]
P62820	Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=1 SV=3 - [RAB1A_HUMAN]

Q9H0U4	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1 - [RAB1B_HUMAN]
Q9UL25	Ras-related protein Rab-21 OS=Homo sapiens GN=RAB21 PE=1 SV=3 - [RAB21_HUMAN]
Q9UL26	Ras-related protein Rab-22A OS=Homo sapiens GN=RAB22A PE=1 SV=2 - [RB22A_HUMAN]
P61019	Ras-related protein Rab-2A OS=Homo sapiens GN=RAB2A PE=1 SV=1 - [RAB2A_HUMAN]
Q9H082	Ras-related protein Rab-33B OS=Homo sapiens GN=RAB33B PE=1 SV=1 - [RB33B_HUMAN]
Q15286	Ras-related protein Rab-35 OS=Homo sapiens GN=RAB35 PE=1 SV=1 - [RAB35_HUMAN]
O95716	Ras-related protein Rab-3D OS=Homo sapiens GN=RAB3D PE=1 SV=1 - [RAB3D_HUMAN]
Q8WXH6	Ras-related protein Rab-40A OS=Homo sapiens GN=RAB40A PE=2 SV=2 - [RB40A_HUMAN]
P61020	Ras-related protein Rab-5B OS=Homo sapiens GN=RAB5B PE=1 SV=1 - [RAB5B_HUMAN]
P51148	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2 - [RAB5C_HUMAN]
P51149	Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1 - [RAB7A_HUMAN]
P61006	Ras-related protein Rab-8A OS=Homo sapiens GN=RAB8A PE=1 SV=1 - [RAB8A_HUMAN]
Q92930	Ras-related protein Rab-8B OS=Homo sapiens GN=RAB8B PE=1 SV=2 - [RAB8B_HUMAN]
P11233	Ras-related protein Ral-A OS=Homo sapiens GN=RALA PE=1 SV=1 - [RALA_HUMAN]
P62834	Ras-related protein Rap-1A OS=Homo sapiens GN=RAP1A PE=1 SV=1 - [RAP1A_HUMAN]
P10114	Ras-related protein Rap-2a OS=Homo sapiens GN=RAP2A PE=1 SV=1 - [RAP2A_HUMAN]
P61225	Ras-related protein Rap-2b OS=Homo sapiens GN=RAP2B PE=1 SV=1 - [RAP2B_HUMAN]
Q00765	Receptor expression-enhancing protein 5 OS=Homo sapiens GN=REEP5 PE=1 SV=3 - [REEP5_HUMAN]
Q96TC7	Regulator of microtubule dynamics protein 3 OS=Homo sapiens GN=FAM82A2 PE=1 SV=2 - [RMD3_HUMAN]
Q9BZ17	Regulator of nonsense transcripts 3B OS=Homo sapiens GN=UPF3B PE=1 SV=1 - [REN3B_HUMAN]
Q9HB40	Retinoid-inducible serine carboxypeptidase OS=Homo sapiens GN=SCPEP1 PE=1 SV=1 - [RISC_HUMAN]
Q8TC12	Retinol dehydrogenase 11 OS=Homo sapiens GN=RDH11 PE=1 SV=2 - [RDH11_HUMAN]
Q6ZS25	Rho guanine nucleotide exchange factor 18 OS=Homo sapiens GN=ARHGEF18 PE=1 SV=3 - [ARHGI_HUMAN]
Q13464	Rho-associated protein kinase 1 OS=Homo sapiens GN=ROCK1 PE=1 SV=1 - [ROCK1_HUMAN]
O75116	Rho-associated protein kinase 2 OS=Homo sapiens GN=ROCK2 PE=1 SV=4 - [ROCK2_HUMAN]
B4DFK3	Rhomboid, veinlet-like 1 (Drosophila), isoform CRA_b OS=Homo sapiens GN=RHBDL1 PE=2 SV=1 - [B4DFK3_HUMAN]
P84095	Rho-related GTP-binding protein RhoG OS=Homo sapiens GN=RHOG PE=1 SV=1 - [RHOG_HUMAN]
O75792	Ribonuclease H2 subunit A OS=Homo sapiens GN=RNASEH2A PE=1 SV=2 - [RNH2A_HUMAN]
P60891	Ribose-phosphate pyrophosphokinase 1 OS=Homo sapiens GN=PRPS1 PE=1 SV=2 - [PRPS1_HUMAN]
P11908	Ribose-phosphate pyrophosphokinase 2 OS=Homo sapiens GN=PRPS2 PE=1 SV=2 - [PRPS2_HUMAN]
Q5VVC8	Ribosomal protein L11 (Fragment) OS=Homo sapiens GN=RPL11 PE=2 SV=1 - [Q5VVC8_HUMAN]
E7EQV9	Ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=3 SV=1 - [E7EQV9_HUMAN]
B3KTM6	Ribosomal protein L5, isoform CRA_b OS=Homo sapiens GN=RPL5 PE=2 SV=1 - [B3KTM6_HUMAN]
Q5JR95	Ribosomal protein S8 OS=Homo sapiens GN=RPS8 PE=4 SV=1 - [Q5JR95_HUMAN]
Q9Y3A4	Ribosomal RNA-processing protein 7 homolog A OS=Homo sapiens GN=RRP7A PE=1 SV=2 - [RRP7A_HUMAN]
Q9Y3A5	Ribosome maturation protein SBDS OS=Homo sapiens GN=SBDS PE=1 SV=4 - [SBDS_HUMAN]
Q9P2E9	Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4 - [RRBP1_HUMAN]
Q5EBL4	RILP-like protein 1 OS=Homo sapiens GN=RILPL1 PE=1 SV=1 - [RIPL1_HUMAN]
O00442	RNA 3'-terminal phosphate cyclase OS=Homo sapiens GN=RTCD1 PE=1 SV=1 - [RTC1_HUMAN]
Q9HC36	RNA methyltransferase-like protein 1 OS=Homo sapiens GN=RNMTL1 PE=1 SV=2 - [RMTL1_HUMAN]
Q92541	RNA polymerase-associated protein RTF1 homolog OS=Homo sapiens GN=RTF1 PE=1 SV=4 - [RTF1_HUMAN]
Q9NTZ6	RNA-binding protein 12 OS=Homo sapiens GN=RBM12 PE=1 SV=1 - [RBM12_HUMAN]
Q96PK6	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]

Q3B7A4	RPLP0 protein OS=Homo sapiens GN=RPLP0 PE=2 SV=1 - [Q3B7A4_HUMAN]
Q7L7Q6	RTN4 OS=Homo sapiens GN=RTN4 PE=2 SV=1 - [Q7L7Q6_HUMAN]
Q96T51	RUN and FYVE domain-containing protein 1 OS=Homo sapiens GN=RUFY1 PE=1 SV=2 - [RUFY1_HUMAN]
Q96C34	RUN domain-containing protein 1 OS=Homo sapiens GN=RUNDC1 PE=1 SV=3 - [RUND1_HUMAN]
Q9Y265	RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1 - [RUVB1_HUMAN]
Q9Y230	RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3 - [RUVB2_HUMAN]
P0C7P3	Schlafen family member 14 OS=Homo sapiens GN=SLFN14 PE=2 SV=2 - [SLN14_HUMAN]
O95171	Sciellin OS=Homo sapiens GN=SCEL PE=1 SV=2 - [SCEL_HUMAN]
Q8WU76	Sec1 family domain-containing protein 2 OS=Homo sapiens GN=SCFD2 PE=1 SV=2 - [SCFD2_HUMAN]
B7ZL00	SEC31A protein OS=Homo sapiens GN=SEC31A PE=2 SV=1 - [B7ZL00_HUMAN]
Q969E2	Secretory carrier-associated membrane protein 4 OS=Homo sapiens GN=SCAMP4 PE=2 SV=1 - [SCAM4_HUMAN]
Q13228	Selenium-binding protein 1 OS=Homo sapiens GN=SELENBP1 PE=1 SV=2 - [SBP1_HUMAN]
Q9H2E6	Semaphorin-6A OS=Homo sapiens GN=SEMA6A PE=1 SV=2 - [SEM6A_HUMAN]
O75326	Semaphorin-7A OS=Homo sapiens GN=SEMA7A PE=1 SV=1 - [SEM7A_HUMAN]
D6RGI3	Septin 11, isoform CRA_b OS=Homo sapiens GN=SEPT11 PE=3 SV=1 - [D6RGI3_HUMAN]
A8MYA6	Serine hydroxymethyltransferase OS=Homo sapiens GN=SHMT1 PE=3 SV=1 - [A8MYA6_HUMAN]
B7Z9F1	Serine hydroxymethyltransferase OS=Homo sapiens GN=SHMT2 PE=2 SV=1 - [B7Z9F1_HUMAN]
O15270	Serine palmitoyltransferase 2 OS=Homo sapiens GN=SPTLC2 PE=1 SV=1 - [SPTC2_HUMAN]
Q9Y5S2	Serine/threonine-protein kinase MRCK beta OS=Homo sapiens GN=CDC42BPB PE=1 SV=2 - [MRCKB_HUMAN]
O95747	Serine/threonine-protein kinase OSR1 OS=Homo sapiens GN=OXSR1 PE=1 SV=1 - [OXSR1_HUMAN]
Q9H2K8	Serine/threonine-protein kinase TAO3 OS=Homo sapiens GN=TAOK3 PE=1 SV=2 - [TAOK3_HUMAN]
Q13362	Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit gamma isoform OS=Homo sapiens GN=PPP2R5C PE=1 SV=3 - [2A5G_HUMAN]
P67775	Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform OS=Homo sapiens GN=PPP2CA PE=1 SV=1 - [PP2AA_HUMAN]
P48454	Serine/threonine-protein phosphatase 2B catalytic subunit gamma isoform OS=Homo sapiens GN=PPP3CC PE=1 SV=3 - [PP2BC_HUMAN]
Q9UPN7	Serine/threonine-protein phosphatase 6 regulatory subunit 1 OS=Homo sapiens GN=PPP6R1 PE=1 SV=5 - [PP6R1_HUMAN]
E9PMD7	Serine/threonine-protein phosphatase OS=Homo sapiens GN=PPP1CA PE=3 SV=1 - [E9PMD7_HUMAN]
B4DDZ8	Serine/threonine-protein phosphatase OS=Homo sapiens GN=PPP5C PE=2 SV=1 - [B4DDZ8_HUMAN]
Q96HS1	Serine/threonine-protein phosphatase PGAM5, mitochondrial OS=Homo sapiens GN=PGAM5 PE=1 SV=2 - [PGAM5_HUMAN]
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3 - [TRFE_HUMAN]
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2 - [ALBU_HUMAN]
P49591	Seryl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=SARS PE=1 SV=3 - [SYSC_HUMAN]
Q8N228	Sex comb on midleg-like protein 4 OS=Homo sapiens GN=SCML4 PE=2 SV=2 - [SCML4_HUMAN]
Q5TIH2	SFT2 domain containing 2 OS=Homo sapiens GN=SFT2D2 PE=2 SV=1 - [Q5TIH2_HUMAN]
Q5T123	SH3 domain binding glutamic acid-rich protein like 3 OS=Homo sapiens GN=SH3BGRL3 PE=2 SV=1 - [Q5T123_HUMAN]
Q9UJC5	SH3 domain-binding glutamic acid-rich-like protein 2 OS=Homo sapiens GN=SH3BGRL2 PE=1 SV=2 - [SH3L2_HUMAN]
O75368	SH3 domain-binding glutamic acid-rich-like protein OS=Homo sapiens GN=SH3BGRL PE=1 SV=1 - [SH3L1_HUMAN]
Q9BZZ2	Sialoadhesin OS=Homo sapiens GN=SIGLEC1 PE=1 SV=2 - [SN_HUMAN]
Q9H9B4	Sideroflexin-1 OS=Homo sapiens GN=SFXN1 PE=1 SV=4 - [SFXN1_HUMAN]
P61009	Signal peptidase complex subunit 3 OS=Homo sapiens GN=SPCS3 PE=1 SV=1 - [SPCS3_HUMAN]
P37108	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2 - [SRP14_HUMAN]
O76094	Signal recognition particle 72 kDa protein OS=Homo sapiens GN=SRP72 PE=1 SV=3 - [SRP72_HUMAN]
P49458	Signal recognition particle 9 kDa protein OS=Homo sapiens GN=SRP9 PE=1 SV=2 - [SRP09_HUMAN]
Q9Y5M8	Signal recognition particle receptor subunit beta OS=Homo sapiens GN=SRPRB PE=1 SV=3 - [SRPRB_HUMAN]

P40763	Signal transducer and activator of transcription 3 OS=Homo sapiens GN=STAT3 PE=1 SV=2 - [STAT3_HUMAN]
O00241	Signal-regulatory protein beta-1 OS=Homo sapiens GN=SIRPB1 PE=1 SV=5 - [SIRB1_HUMAN]
Q8N3X5	Similar to ATPase, Ca++ transporting, cardiac muscle, fast twitch 1 (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q8N3X5_HUMAN]
A7XYQ1	Sine oculis-binding protein homolog OS=Homo sapiens GN=SOBP PE=2 SV=2 - [SOBP_HUMAN]
Q14190	Single-minded homolog 2 OS=Homo sapiens GN=SIM2 PE=1 SV=2 - [SIM2_HUMAN]
Q96G53	SLC27A4 protein OS=Homo sapiens GN=SLC27A4 PE=2 SV=1 - [Q96G53_HUMAN]
O43295	SLIT-ROBO Rho GTPase-activating protein 3 OS=Homo sapiens GN=SRGAP3 PE=1 SV=3 - [SRGP2_HUMAN]
Q8NHG7	Small VCP/p97-interacting protein OS=Homo sapiens GN=SVIP PE=2 SV=1 - [SVIP_HUMAN]
C0H5W9	SNX6 protein OS=Homo sapiens GN=SNX6 PE=2 SV=1 - [C0H5W9_HUMAN]
Q9UQD0	Sodium channel protein type 8 subunit alpha OS=Homo sapiens GN=SCN8A PE=1 SV=1 - [SCN8A_HUMAN]
P14415	Sodium/potassium-transporting ATPase subunit beta-2 OS=Homo sapiens GN=ATP1B2 PE=1 SV=3 - [AT1B2_HUMAN]
P54709	Sodium/potassium-transporting ATPase subunit beta-3 OS=Homo sapiens GN=ATP1B3 PE=1 SV=1 - [AT1B3_HUMAN]
B5A923	Soluble VEGFR1 variant 21 OS=Homo sapiens GN=VEGFR1 PE=2 SV=1 - [B5A923_HUMAN]
P11166	Solute carrier family 2, facilitated glucose transporter member 1 OS=Homo sapiens GN=SLC2A1 PE=1 SV=2 - [GTR1_HUMAN]
Q8TD20	Solute carrier family 2, facilitated glucose transporter member 12 OS=Homo sapiens GN=SLC2A12 PE=2 SV=1 - [GTR12_HUMAN]
P14672	Solute carrier family 2, facilitated glucose transporter member 4 OS=Homo sapiens GN=SLC2A4 PE=1 SV=1 - [GTR4_HUMAN]
Q5VTS8	Solute carrier family 25, member 27 (Fragment) OS=Homo sapiens GN=SLC25A27 PE=2 SV=1 - [Q5VTS8_HUMAN]
Q9NP59	Solute carrier family 40 member 1 OS=Homo sapiens GN=SLC40A1 PE=1 SV=1 - [S40A1_HUMAN]
Q9BX66	Sorbin and SH3 domain-containing protein 1 OS=Homo sapiens GN=SORBS1 PE=1 SV=3 - [SRBS1_HUMAN]
Q00796	Sorbitol dehydrogenase OS=Homo sapiens GN=SORD PE=1 SV=4 - [DHSO_HUMAN]
Q9Y512	Sorting and assembly machinery component 50 homolog OS=Homo sapiens GN=SAMM50 PE=1 SV=3 - [SAM50_HUMAN]
P02549	Spectrin alpha chain, erythrocyte OS=Homo sapiens GN=SPTA1 PE=1 SV=5 - [SPTA1_HUMAN]
P11277	Spectrin beta chain, erythrocyte OS=Homo sapiens GN=SPTB PE=1 SV=5 - [SPTB1_HUMAN]
Q5MJ07	Sperm protein associated with the nucleus on the X chromosome N5 OS=Homo sapiens GN=SPANXN5 PE=2 SV=1 - [SPXN5_HUMAN]
B3KUY1	Splicing factor, arginine/serine-rich 2, isoform CRA_d OS=Homo sapiens GN=SFRS2 PE=2 SV=1 - [B3KUY1_HUMAN]
B4E241	Splicing factor, arginine/serine-rich 3, isoform CRA_a OS=Homo sapiens GN=SRSF3 PE=2 SV=1 - [B4E241_HUMAN]
P23246	Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2 - [SFPQ_HUMAN]
Q9H2G2	STE20-like serine/threonine-protein kinase OS=Homo sapiens GN=SLK PE=1 SV=1 - [SLK_HUMAN]
Q9C0K7	STE20-related kinase adapter protein beta OS=Homo sapiens GN=STRADB PE=1 SV=1 - [STRAB_HUMAN]
P31948	Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=1 SV=1 - [STIP1_HUMAN]
O95347	Structural maintenance of chromosomes protein 2 OS=Homo sapiens GN=SMC2 PE=1 SV=2 - [SMC2_HUMAN]
P21912	Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Homo sapiens GN=SDHB PE=1 SV=3 - [DHSB_HUMAN]
P53597	Succinyl-CoA ligase [GDP-forming] subunit alpha, mitochondrial OS=Homo sapiens GN=SUCLG1 PE=1 SV=4 - [SUCA_HUMAN]
Q96199	Succinyl-CoA ligase [GDP-forming] subunit beta, mitochondrial OS=Homo sapiens GN=SUCLG2 PE=1 SV=2 - [SUCB2_HUMAN]
P55809	Succinyl-CoA:3-ketoacid-coenzyme A transferase 1, mitochondrial OS=Homo sapiens GN=OXCT1 PE=1 SV=1 - [SCOT1_HUMAN]
Q9Y6N5	Sulfide:quinone oxidoreductase, mitochondrial OS=Homo sapiens GN=SQRDL PE=1 SV=1 - [SQRD_HUMAN]
P63279	SUMO-conjugating enzyme UBC9 OS=Homo sapiens GN=UBE2I PE=1 SV=1 - [UBC9_HUMAN]
P00441	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2 - [SODC_HUMAN]
O60284	Suppression of tumorigenicity 18 protein OS=Homo sapiens GN=ST18 PE=1 SV=1 - [ST18_HUMAN]
Q99536	Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2 - [VAT1_HUMAN]
O00161	Synaptosomal-associated protein 23 OS=Homo sapiens GN=SNAP23 PE=1 SV=1 - [SNP23_HUMAN]
Q96C24	Synaptotagmin-like protein 4 OS=Homo sapiens GN=SYTL4 PE=1 SV=2 - [SYTL4_HUMAN]
B1AJQ6	Syntaxin 12 (Fragment) OS=Homo sapiens GN=STX12 PE=4 SV=1 - [B1AJQ6_HUMAN]

A8MX0	Syntaxin 4A (Placental), isoform CRA_c OS=Homo sapiens GN=STX4 PE=2 SV=1 - [A8MX0_HUMAN]
B4DKR0	Syntaxin 5A, isoform CRA_c OS=Homo sapiens GN=STX5 PE=2 SV=1 - [B4DKR0_HUMAN]
O43752	Syntaxin-6 OS=Homo sapiens GN=STX6 PE=1 SV=1 - [STX6_HUMAN]
O15400	Syntaxin-7 OS=Homo sapiens GN=STX7 PE=1 SV=4 - [STX7_HUMAN]
Q9UNK0	Syntaxin-8 OS=Homo sapiens GN=STX8 PE=1 SV=2 - [STX8_HUMAN]
O00186	Syntaxin-binding protein 3 OS=Homo sapiens GN=STXBP3 PE=1 SV=2 - [STXB3_HUMAN]
Q5T5C0	Syntaxin-binding protein 5 OS=Homo sapiens GN=STXBP5 PE=1 SV=1 - [STXB5_HUMAN]
Q5TCU6	Talin 1 OS=Homo sapiens GN=TLN1 PE=2 SV=1 - [Q5TCU6_HUMAN]
Q9Y4G6	Talin-2 OS=Homo sapiens GN=TLN2 PE=1 SV=4 - [TLN2_HUMAN]
B1AKP7	TAR DNA binding protein OS=Homo sapiens GN=TARDBP PE=4 SV=1 - [B1AKP7_HUMAN]
Q9ULP9	TBC1 domain family member 24 OS=Homo sapiens GN=TBC1D24 PE=1 SV=2 - [TBC24_HUMAN]
O95759	TBC1 domain family member 8 OS=Homo sapiens GN=TBC1D8 PE=1 SV=3 - [TBCD8_HUMAN]
P17987	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1 - [TCPA_HUMAN]
P50991	T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4 - [TCPD_HUMAN]
P48643	T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=1 SV=1 - [TCPE_HUMAN]
P50990	T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4 - [TCPQ_HUMAN]
P40227	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3 - [TCPZ_HUMAN]
Q92526	T-complex protein 1 subunit zeta-2 OS=Homo sapiens GN=CCT6B PE=1 SV=5 - [TCPW_HUMAN]
Q9NYB0	Telomeric repeat-binding factor 2-interacting protein 1 OS=Homo sapiens GN=TERF2IP PE=1 SV=1 - [TE2IP_HUMAN]
Q59G71	Tensin variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q59G71_HUMAN]
Q9BXA7	Testis-specific serine/threonine-protein kinase 1 OS=Homo sapiens GN=TSSK1B PE=1 SV=1 - [TSSK1_HUMAN]
B1ALW1	Thioredoxin OS=Homo sapiens GN=TXN PE=4 SV=1 - [B1ALW1_HUMAN]
O43396	Thioredoxin-like protein 1 OS=Homo sapiens GN=TXNL1 PE=1 SV=3 - [TXNL1_HUMAN]
Q9H3N1	Thioredoxin-related transmembrane protein 1 OS=Homo sapiens GN=TMX1 PE=1 SV=1 - [TMX1_HUMAN]
P26639	Threonyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=TARS PE=1 SV=3 - [SYTC_HUMAN]
P23919	Thymidylate kinase OS=Homo sapiens GN=DTYMK PE=1 SV=4 - [KTHY_HUMAN]
Q9Y2W1	Thyroid hormone receptor-associated protein 3 OS=Homo sapiens GN=THRAP3 PE=1 SV=2 - [TR150_HUMAN]
Q17RP2	Tigger transposable element-derived protein 6 OS=Homo sapiens GN=TIGD6 PE=2 SV=2 - [TIGD6_HUMAN]
Q8WZ42	Titin OS=Homo sapiens GN=TTN PE=1 SV=2 - [TITIN_HUMAN]
Q7Z649	TMEM9 domain family, member B, isoform CRA_b OS=Homo sapiens GN=TMEM9B PE=2 SV=1 - [Q7Z649_HUMAN]
Q8NFAQ8	Torsin-1A-interacting protein 2 OS=Homo sapiens GN=TOR1AIP2 PE=1 SV=1 - [TOIP2_HUMAN]
O14804	Trace amine-associated receptor 5 OS=Homo sapiens GN=TAAR5 PE=2 SV=2 - [TAAR5_HUMAN]
O75865	Trafficking protein particle complex subunit 6A OS=Homo sapiens GN=TRAPPC6A PE=1 SV=2 - [TPC6A_HUMAN]
Q9NZ01	Trans-2,3-enoyl-CoA reductase OS=Homo sapiens GN=TECR PE=1 SV=1 - [TECR_HUMAN]
Q9BV79	Trans-2-enoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=MECR PE=1 SV=2 - [MECR_HUMAN]
P37837	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2 - [TALDO_HUMAN]
P23193	Transcription elongation factor A protein 1 OS=Homo sapiens GN=TCEA1 PE=1 SV=2 - [TCEA1_HUMAN]
B8ZZU8	Transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B), isoform CRA_b OS=Homo sapiens GN=TCEB2 PE=4 SV=1 - [B8ZZU8_HUMAN]
Q7KZ85	Transcription elongation factor SPT6 OS=Homo sapiens GN=SUPT6H PE=1 SV=2 - [SPT6H_HUMAN]
O14776	Transcription elongation regulator 1 OS=Homo sapiens GN=TCERG1 PE=1 SV=2 - [TCRG1_HUMAN]
Q9UGU0	Transcription factor 20 OS=Homo sapiens GN=TCF20 PE=1 SV=3 - [TCF20_HUMAN]
Q00059	Transcription factor A, mitochondrial OS=Homo sapiens GN=TFAM PE=1 SV=1 - [TFAM_HUMAN]
Q6VUC0	Transcription factor AP-2-epsilon OS=Homo sapiens GN=TFAP2E PE=2 SV=1 - [AP2E_HUMAN]

A0AVK6	Transcription factor E2F8 OS=Homo sapiens GN=E2F8 PE=1 SV=1 - [E2F8_HUMAN]
Q00577	Transcriptional activator protein Pur-alpha OS=Homo sapiens GN=PURA PE=1 SV=2 - [PURA_HUMAN]
Q96QR8	Transcriptional activator protein Pur-beta OS=Homo sapiens GN=PURB PE=1 SV=3 - [PURB_HUMAN]
P02786	Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=2 - [TFR1_HUMAN]
B9ZVW9	Transforming growth factor, beta-induced, 68kDa, isoform CRA_a OS=Homo sapiens GN=TGFBI PE=4 SV=1 - [B9ZVW9_HUMAN]
P61586	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1 - [RHOA_HUMAN]
P55072	Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4 - [TERA_HUMAN]
P49770	Translation initiation factor eIF-2B subunit beta OS=Homo sapiens GN=EIF2B2 PE=1 SV=3 - [EI2BB_HUMAN]
Q9UI10	Translation initiation factor eIF-2B subunit delta OS=Homo sapiens GN=EIF2B4 PE=1 SV=2 - [EI2BD_HUMAN]
Q92616	Translational activator GCN1 OS=Homo sapiens GN=GCN1L1 PE=1 SV=6 - [GCN1L_HUMAN]
Q99598	Translin-associated protein X OS=Homo sapiens GN=TSNAX PE=1 SV=1 - [TSNAX_HUMAN]
Q99442	Translocation protein SEC62 OS=Homo sapiens GN=SEC62 PE=1 SV=1 - [SEC62_HUMAN]
B1AH87	Translocator protein (18kDa) (Fragment) OS=Homo sapiens GN=TSPO PE=4 SV=1 - [B1AH87_HUMAN]
Q9HD45	Transmembrane 9 superfamily member 3 OS=Homo sapiens GN=TM9SF3 PE=1 SV=2 - [TM9S3_HUMAN]
B7Z1P7	Transmembrane and coiled-coil domain family 2, isoform CRA_d OS=Homo sapiens GN=TMCC2 PE=2 SV=1 - [B7Z1P7_HUMAN]
Q13445	Transmembrane emp24 domain-containing protein 1 OS=Homo sapiens GN=TMED1 PE=1 SV=1 - [TMED1_HUMAN]
P49755	Transmembrane emp24 domain-containing protein 10 OS=Homo sapiens GN=TMED10 PE=1 SV=2 - [TMEDA_HUMAN]
Q9BVK6	Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=2 - [TMED9_HUMAN]
Q9NUM4	Transmembrane protein 106B OS=Homo sapiens GN=TMEM106B PE=1 SV=2 - [T106B_HUMAN]
Q9BVC6	Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1 - [TM109_HUMAN]
Q7Z7N9	Transmembrane protein 179B OS=Homo sapiens GN=TMEM179B PE=1 SV=1 - [T179B_HUMAN]
Q6UW68	Transmembrane protein 205 OS=Homo sapiens GN=TMEM205 PE=1 SV=1 - [TM205_HUMAN]
Q5T3F8	Transmembrane protein 63B OS=Homo sapiens GN=TMEM63B PE=1 SV=1 - [TM63B_HUMAN]
Q96KR6	Transmembrane protein C20orf108 OS=Homo sapiens GN=C20orf108 PE=1 SV=2 - [CT108_HUMAN]
Q6IN77	Transportin 2 OS=Homo sapiens GN=TNPO2 PE=2 SV=1 - [Q6IN77_HUMAN]
Q92973	Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2 - [TNPO1_HUMAN]
P53007	Tricarboxylate transport protein, mitochondrial OS=Homo sapiens GN=SLC25A1 PE=1 SV=2 - [TXTP_HUMAN]
P40939	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2 - [ECHA_HUMAN]
P55084	Trifunctional enzyme subunit beta, mitochondrial OS=Homo sapiens GN=HADHB PE=1 SV=3 - [ECHB_HUMAN]
P60174	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=2 - [TPI1_HUMAN]
Q5SRJ5	Tripartite motif-containing 10 OS=Homo sapiens GN=TRIM10 PE=2 SV=1 - [Q5SRJ5_HUMAN]
Q8NG06	Tripartite motif-containing protein 58 OS=Homo sapiens GN=TRIM58 PE=2 SV=2 - [TRI58_HUMAN]
P29144	Tripeptidyl-peptidase 2 OS=Homo sapiens GN=TPP2 PE=1 SV=4 - [TPP2_HUMAN]
Q5JR82	Tropomodulin 4 (Muscle) (Fragment) OS=Homo sapiens GN=TMOD4 PE=2 SV=1 - [Q5JR82_HUMAN]
P28289	Tropomodulin-1 OS=Homo sapiens GN=TMOD1 PE=1 SV=1 - [TMOD1_HUMAN]
D9YZV5	Tropomyosin 1 (Alpha) isoform 4 OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [D9YZV5_HUMAN]
D9YZV7	Tropomyosin 1 (Alpha) isoform 6 OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [D9YZV7_HUMAN]
Q5VU58	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU58_HUMAN]
Q5VU59	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU59_HUMAN]
Q5VU61	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU61_HUMAN]
Q5VU66	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU66_HUMAN]
P09493	Tropomyosin alpha-1 chain OS=Homo sapiens GN=TPM1 PE=1 SV=2 - [TPM1_HUMAN]
P06753	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=1 SV=1 - [TPM3_HUMAN]

Q5LJ98	TROVE domain family, member 2 OS=Homo sapiens GN=TROVE2 PE=2 SV=1 - [Q5LJ98_HUMAN]
P48775	Tryptophan 2,3-dioxygenase OS=Homo sapiens GN=TDO2 PE=1 SV=1 - [T23O_HUMAN]
P23381	Tryptophanyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=WARS PE=1 SV=2 - [SYWC_HUMAN]
Q9Y3Q8	TSC22 domain family protein 4 OS=Homo sapiens GN=TSC22D4 PE=1 SV=2 - [T22D4_HUMAN]
Q9BQE3	Tubulin alpha-1C chain OS=Homo sapiens GN=TUBA1C PE=1 SV=1 - [TBA1C_HUMAN]
P07437	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2 - [TBB5_HUMAN]
P68371	Tubulin beta-2C chain OS=Homo sapiens GN=TUBB2C PE=1 SV=1 - [TBB2C_HUMAN]
B1AH89	Tubulin tyrosine ligase-like family, member 12 OS=Homo sapiens GN=TTLL12 PE=4 SV=1 - [B1AH89_HUMAN]
A8MUB1	Tubulin, alpha 1 (Testis specific), isoform CRA_a OS=Homo sapiens GN=TUBA4A PE=2 SV=1 - [A8MUB1_HUMAN]
Q8NHU6	Tudor domain-containing protein 7 OS=Homo sapiens GN=TDRD7 PE=1 SV=2 - [TDRD7_HUMAN]
Q6UWW7	TUFT1 OS=Homo sapiens GN=TUFT1 PE=2 SV=1 - [Q6UWW7_HUMAN]
P25445	Tumor necrosis factor receptor superfamily member 6 OS=Homo sapiens GN=FAS PE=1 SV=1 - [TNR6_HUMAN]
Q6RW13	Type-1 angiotensin II receptor-associated protein OS=Homo sapiens GN=AGTRAP PE=1 SV=1 - [ATRAP_HUMAN]
Q9UIG0	Tyrosine-protein kinase BAZ1B OS=Homo sapiens GN=BAZ1B PE=1 SV=2 - [BAZ1B_HUMAN]
Q06124	Tyrosine-protein phosphatase non-receptor type 11 OS=Homo sapiens GN=PTPN11 PE=1 SV=2 - [PTN11_HUMAN]
P54577	Tyrosyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=YARS PE=1 SV=4 - [SYYC_HUMAN]
O43290	U4/U6.U5 tri-snRNP-associated protein 1 OS=Homo sapiens GN=SART1 PE=1 SV=1 - [SNUT1_HUMAN]
Q9Y333	U6 snRNA-associated Sm-like protein LSM2 OS=Homo sapiens GN=LSM2 PE=1 SV=1 - [LSM2_HUMAN]
Q9Y4Z0	U6 snRNA-associated Sm-like protein LSM4 OS=Homo sapiens GN=LSM4 PE=1 SV=1 - [LSM4_HUMAN]
Q70EL4	Ubiquitin carboxyl-terminal hydrolase 43 OS=Homo sapiens GN=USP43 PE=1 SV=2 - [UBP43_HUMAN]
P45974	Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5 PE=1 SV=2 - [UBP5_HUMAN]
A6NJA2	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP14 PE=3 SV=1 - [A6NJA2_HUMAN]
E9PCQ3	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP15 PE=3 SV=2 - [E9PCQ3_HUMAN]
B7WPF4	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP24 PE=3 SV=2 - [B7WPF4_HUMAN]
D6RAW0	Ubiquitin carrier protein OS=Homo sapiens GN=UBE2D3 PE=3 SV=1 - [D6RAW0_HUMAN]
Q9BSL1	Ubiquitin-associated domain-containing protein 1 OS=Homo sapiens GN=UBAC1 PE=1 SV=1 - [UBAC1_HUMAN]
P61088	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1 - [UBE2N_HUMAN]
Q9C0C9	Ubiquitin-conjugating enzyme E2 O OS=Homo sapiens GN=UBE2O PE=1 SV=3 - [UBE2O_HUMAN]
Q8WVY7	Ubiquitin-like domain-containing CTD phosphatase 1 OS=Homo sapiens GN=UBLCP1 PE=1 SV=2 - [UBCP1_HUMAN]
P22314	Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3 - [UBA1_HUMAN]
A0AVT1	Ubiquitin-like modifier-activating enzyme 6 OS=Homo sapiens GN=UBA6 PE=1 SV=1 - [UBA6_HUMAN]
P11441	Ubiquitin-like protein 4A OS=Homo sapiens GN=UBL4A PE=1 SV=1 - [UBL4A_HUMAN]
Q9NYU2	UDP-glucose:glycoprotein glucosyltransferase 1 OS=Homo sapiens GN=UGGT1 PE=1 SV=3 - [UGGG1_HUMAN]
Q9NTN3	UDP-glucuronic acid/UDP-N-acetylgalactosamine transporter OS=Homo sapiens GN=SLC35D1 PE=1 SV=1 - [S35D1_HUMAN]
O15294	UDP-N-acetylglucosamine--peptide N-acetylglucosaminyltransferase 110 kDa subunit OS=Homo sapiens GN=OGT PE=1 SV=3 - [OGT1_HUMAN]
A8K6F7	Unc-45 homolog A (C. elegans), isoform CRA_a OS=Homo sapiens GN=UNC45A PE=2 SV=1 - [A8K6F7_HUMAN]
B0QY64	Unc-84 homolog B (C. elegans) (Fragment) OS=Homo sapiens GN=UNC84B PE=4 SV=1 - [B0QY64_HUMAN]
B4DGL8	Uncharacterized protein (Fragment) OS=Homo sapiens GN=ABCB7 PE=2 SV=1 - [B4DGL8_HUMAN]
Q5T446	Uncharacterized protein (Fragment) OS=Homo sapiens GN=UROD PE=2 SV=1 - [Q5T446_HUMAN]
Q5SY21	Uncharacterized protein (Fragment) OS=Homo sapiens GN=ZBTB48 PE=2 SV=1 - [Q5SY21_HUMAN]
Q96GS4	Uncharacterized protein C17orf59 OS=Homo sapiens GN=C17orf59 PE=1 SV=2 - [CQ059_HUMAN]
Q86X18	Uncharacterized protein C19orf68 OS=Homo sapiens GN=C19orf68 PE=1 SV=2 - [CS068_HUMAN]
A0AUZ9	Uncharacterized protein C2orf67 OS=Homo sapiens GN=C2orf67 PE=1 SV=2 - [CB067_HUMAN]

Q9BZA4	Uncharacterized protein CYorf15B OS=Homo sapiens GN=CYorf15B PE=1 SV=1 - [CY15B_HUMAN]
Q9UF83	Uncharacterized protein DKFZp434B061 OS=Homo sapiens PE=2 SV=2 - [YM012_HUMAN]
Q9Y2F5	Uncharacterized protein KIAA0947 OS=Homo sapiens GN=KIAA0947 PE=1 SV=5 - [K0947_HUMAN]
B2RUZ4	Uncharacterized protein LOC388588 OS=Homo sapiens PE=1 SV=1 - [YA047_HUMAN]
F5GXL7	Uncharacterized protein OS=Homo sapiens GN=ABCC1 PE=3 SV=1 - [F5GXL7_HUMAN]
E9PBW0	Uncharacterized protein OS=Homo sapiens GN=ABCC5 PE=3 SV=1 - [E9PBW0_HUMAN]
F5GYC1	Uncharacterized protein OS=Homo sapiens GN=ABCD3 PE=3 SV=1 - [F5GYC1_HUMAN]
C9IZX5	Uncharacterized protein OS=Homo sapiens GN=ABHD10 PE=4 SV=1 - [C9IZX5_HUMAN]
B7Z9I1	Uncharacterized protein OS=Homo sapiens GN=ACADM PE=2 SV=1 - [B7Z9I1_HUMAN]
E9PE82	Uncharacterized protein OS=Homo sapiens GN=ACADS PE=4 SV=1 - [E9PE82_HUMAN]
C9JGQ9	Uncharacterized protein OS=Homo sapiens GN=ACHE PE=4 SV=1 - [C9JGQ9_HUMAN]
B4E3P0	Uncharacterized protein OS=Homo sapiens GN=ACLY PE=2 SV=1 - [B4E3P0_HUMAN]
F5H2A5	Uncharacterized protein OS=Homo sapiens GN=ACO2 PE=4 SV=1 - [F5H2A5_HUMAN]
F5H2L4	Uncharacterized protein OS=Homo sapiens GN=ACOT13 PE=4 SV=1 - [F5H2L4_HUMAN]
F5GWW7	Uncharacterized protein OS=Homo sapiens GN=ACP2 PE=4 SV=1 - [F5GWW7_HUMAN]
E7ERD7	Uncharacterized protein OS=Homo sapiens GN=ACSL6 PE=4 SV=1 - [E7ERD7_HUMAN]
F5H6F4	Uncharacterized protein OS=Homo sapiens GN=ACSS1 PE=4 SV=1 - [F5H6F4_HUMAN]
E7EWU3	Uncharacterized protein OS=Homo sapiens GN=ACSS2 PE=4 SV=1 - [E7EWU3_HUMAN]
F5GXS2	Uncharacterized protein OS=Homo sapiens GN=ACTN4 PE=4 SV=1 - [F5GXS2_HUMAN]
F5H3P5	Uncharacterized protein OS=Homo sapiens GN=ACTR3 PE=3 SV=1 - [F5H3P5_HUMAN]
C9JTM0	Uncharacterized protein OS=Homo sapiens GN=ADD2 PE=4 SV=2 - [C9JTM0_HUMAN]
E7EQY8	Uncharacterized protein OS=Homo sapiens GN=ADPGK PE=4 SV=2 - [E7EQY8_HUMAN]
E7EWH4	Uncharacterized protein OS=Homo sapiens GN=AK2 PE=3 SV=1 - [E7EWH4_HUMAN]
E7ET30	Uncharacterized protein OS=Homo sapiens GN=AK3 PE=3 SV=1 - [E7ET30_HUMAN]
B7Z5R1	Uncharacterized protein OS=Homo sapiens GN=AKT1 PE=2 SV=1 - [B7Z5R1_HUMAN]
B4DLJ0	Uncharacterized protein OS=Homo sapiens GN=ALDH1B1 PE=2 SV=1 - [B4DLJ0_HUMAN]
F5H328	Uncharacterized protein OS=Homo sapiens GN=ALDH5A1 PE=3 SV=1 - [F5H328_HUMAN]
E9PPG2	Uncharacterized protein OS=Homo sapiens GN=AMPD3 PE=4 SV=1 - [E9PPG2_HUMAN]
E7EVE3	Uncharacterized protein OS=Homo sapiens GN=ANK1 PE=4 SV=1 - [E7EVE3_HUMAN]
E9PHW9	Uncharacterized protein OS=Homo sapiens GN=ANK2 PE=4 SV=1 - [E9PHW9_HUMAN]
F5H754	Uncharacterized protein OS=Homo sapiens GN=ANKFY1 PE=4 SV=1 - [F5H754_HUMAN]
B5MEE3	Uncharacterized protein OS=Homo sapiens GN=ANKRD17 PE=4 SV=2 - [B5MEE3_HUMAN]
B4DIW9	Uncharacterized protein OS=Homo sapiens GN=ANKRD28 PE=2 SV=1 - [B4DIW9_HUMAN]
F2Z3H3	Uncharacterized protein OS=Homo sapiens GN=ANP32A PE=4 SV=2 - [F2Z3H3_HUMAN]
E9PPH5	Uncharacterized protein OS=Homo sapiens GN=ANP32E PE=4 SV=1 - [E9PPH5_HUMAN]
E9PHT9	Uncharacterized protein OS=Homo sapiens GN=ANXA5 PE=3 SV=1 - [E9PHT9_HUMAN]
F5H0H0	Uncharacterized protein OS=Homo sapiens GN=AP2M1 PE=4 SV=1 - [F5H0H0_HUMAN]
F5H093	Uncharacterized protein OS=Homo sapiens GN=AP3B1 PE=4 SV=1 - [F5H093_HUMAN]
C9JX71	Uncharacterized protein OS=Homo sapiens GN=APOD PE=4 SV=1 - [C9JX71_HUMAN]
E9PC21	Uncharacterized protein OS=Homo sapiens GN=AQP1 PE=3 SV=1 - [E9PC21_HUMAN]
B7ZB63	Uncharacterized protein OS=Homo sapiens GN=ARF3 PE=2 SV=1 - [B7ZB63_HUMAN]
E9PNT2	Uncharacterized protein OS=Homo sapiens GN=ARHGAP27 PE=4 SV=1 - [E9PNT2_HUMAN]
F5GY14	Uncharacterized protein OS=Homo sapiens GN=ARL6IP1 PE=4 SV=1 - [F5GY14_HUMAN]

E9PER8	Uncharacterized protein OS=Homo sapiens GN=ARL6IP5 PE=4 SV=1 - [E9PER8_HUMAN]
E5RHK3	Uncharacterized protein OS=Homo sapiens GN=ARMC1 PE=4 SV=1 - [E5RHK3_HUMAN]
B7Z7I1	Uncharacterized protein OS=Homo sapiens GN=ARMC4 PE=2 SV=1 - [B7Z7I1_HUMAN]
B4DLQ7	Uncharacterized protein OS=Homo sapiens GN=ARPC1A PE=2 SV=1 - [B4DLQ7_HUMAN]
E7ERV9	Uncharacterized protein OS=Homo sapiens GN=ASAH1 PE=4 SV=1 - [E7ERV9_HUMAN]
E7EVN0	Uncharacterized protein OS=Homo sapiens GN=ASNA1 PE=4 SV=1 - [E7EVN0_HUMAN]
C9JAL9	Uncharacterized protein OS=Homo sapiens GN=ASPSCR1 PE=4 SV=1 - [C9JAL9_HUMAN]
F5GWY2	Uncharacterized protein OS=Homo sapiens GN=ATIC PE=4 SV=1 - [F5GWY2_HUMAN]
F5H6I7	Uncharacterized protein OS=Homo sapiens GN=ATL3 PE=4 SV=1 - [F5H6I7_HUMAN]
F5H3A1	Uncharacterized protein OS=Homo sapiens GN=ATP1A1 PE=3 SV=1 - [F5H3A1_HUMAN]
D6R9H7	Uncharacterized protein OS=Homo sapiens GN=ATP5G1 PE=3 SV=1 - [D6R9H7_HUMAN]
E9PN17	Uncharacterized protein OS=Homo sapiens GN=ATP5L PE=4 SV=1 - [E9PN17_HUMAN]
F5H569	Uncharacterized protein OS=Homo sapiens GN=ATP6V0A1 PE=4 SV=1 - [F5H569_HUMAN]
B7Z1R5	Uncharacterized protein OS=Homo sapiens GN=ATP6V1A PE=2 SV=1 - [B7Z1R5_HUMAN]
A8MQ60	Uncharacterized protein OS=Homo sapiens GN=ATP6V1C1 PE=4 SV=1 - [A8MQ60_HUMAN]
A8MUN4	Uncharacterized protein OS=Homo sapiens GN=ATP6V1E1 PE=2 SV=1 - [A8MUN4_HUMAN]
E9PGE6	Uncharacterized protein OS=Homo sapiens GN=ATP6V1H PE=4 SV=1 - [E9PGE6_HUMAN]
E7EU18	Uncharacterized protein OS=Homo sapiens GN=AUP1 PE=4 SV=1 - [E7EU18_HUMAN]
C9JAB1	Uncharacterized protein OS=Homo sapiens GN=AZGP1 PE=4 SV=1 - [C9JAB1_HUMAN]
B4DXE2	Uncharacterized protein OS=Homo sapiens GN=BAG2 PE=2 SV=1 - [B4DXE2_HUMAN]
E7EX43	Uncharacterized protein OS=Homo sapiens GN=BBS9 PE=4 SV=1 - [E7EX43_HUMAN]
A6NME1	Uncharacterized protein OS=Homo sapiens GN=BCAP29 PE=2 SV=2 - [A6NME1_HUMAN]
A8MVZ6	Uncharacterized protein OS=Homo sapiens GN=BICD1 PE=4 SV=2 - [A8MVZ6_HUMAN]
F5H7R6	Uncharacterized protein OS=Homo sapiens GN=BLM PE=4 SV=1 - [F5H7R6_HUMAN]
B0AZS9	Uncharacterized protein OS=Homo sapiens GN=BNIP3L PE=2 SV=1 - [B0AZS9_HUMAN]
E9PH68	Uncharacterized protein OS=Homo sapiens GN=BRCA1 PE=4 SV=1 - [E9PH68_HUMAN]
A6NF68	Uncharacterized protein OS=Homo sapiens GN=BRDT PE=4 SV=1 - [A6NF68_HUMAN]
B4DM89	Uncharacterized protein OS=Homo sapiens GN=BSDC1 PE=2 SV=1 - [B4DM89_HUMAN]
E9PL10	Uncharacterized protein OS=Homo sapiens GN=BTF3L4 PE=4 SV=1 - [E9PL10_HUMAN]
B4DDM6	Uncharacterized protein OS=Homo sapiens GN=BUB3 PE=2 SV=1 - [B4DDM6_HUMAN]
E9PBZ7	Uncharacterized protein OS=Homo sapiens GN=C13orf38 PE=4 SV=1 - [E9PBZ7_HUMAN]
E5RK14	Uncharacterized protein OS=Homo sapiens GN=C14orf159 PE=4 SV=1 - [E5RK14_HUMAN]
F5H2W0	Uncharacterized protein OS=Homo sapiens GN=C18orf8 PE=4 SV=1 - [F5H2W0_HUMAN]
E5RI90	Uncharacterized protein OS=Homo sapiens GN=C1orf198 PE=4 SV=1 - [E5RI90_HUMAN]
E9PCW3	Uncharacterized protein OS=Homo sapiens GN=C22orf28 PE=4 SV=1 - [E9PCW3_HUMAN]
C9J384	Uncharacterized protein OS=Homo sapiens GN=C3orf26 PE=4 SV=1 - [C9J384_HUMAN]
E7ETQ0	Uncharacterized protein OS=Homo sapiens GN=C4orf26 PE=4 SV=1 - [E7ETQ0_HUMAN]
F5H200	Uncharacterized protein OS=Homo sapiens GN=C6orf146 PE=4 SV=1 - [F5H200_HUMAN]
C9JGX5	Uncharacterized protein OS=Homo sapiens GN=C9orf174 PE=4 SV=1 - [C9JGX5_HUMAN]
C9K005	Uncharacterized protein OS=Homo sapiens GN=C9orf5 PE=4 SV=2 - [C9K005_HUMAN]
E5RJF6	Uncharacterized protein OS=Homo sapiens GN=CA1 PE=4 SV=1 - [E5RJF6_HUMAN]
E7EWT6	Uncharacterized protein OS=Homo sapiens GN=CALCOCO1 PE=4 SV=2 - [E7EWT6_HUMAN]
C9JUG7	Uncharacterized protein OS=Homo sapiens GN=CAPZA2 PE=4 SV=1 - [C9JUG7_HUMAN]

D6RC54	Uncharacterized protein OS=Homo sapiens GN=CAST PE=4 SV=1 - [D6RC54_HUMAN]
E7EVY3	Uncharacterized protein OS=Homo sapiens GN=CAST PE=4 SV=1 - [E7EVY3_HUMAN]
E9PCH5	Uncharacterized protein OS=Homo sapiens GN=CAST PE=4 SV=1 - [E9PCH5_HUMAN]
C9JMM0	Uncharacterized protein OS=Homo sapiens GN=CBX3 PE=4 SV=1 - [C9JMM0_HUMAN]
F5GWI9	Uncharacterized protein OS=Homo sapiens GN=CCDC53 PE=4 SV=1 - [F5GWI9_HUMAN]
C9JPL0	Uncharacterized protein OS=Homo sapiens GN=CCNL1 PE=3 SV=1 - [C9JPL0_HUMAN]
E9PAQ6	Uncharacterized protein OS=Homo sapiens GN=CCT3 PE=3 SV=1 - [E9PAQ6_HUMAN]
E9PB22	Uncharacterized protein OS=Homo sapiens GN=CD47 PE=4 SV=1 - [E9PB22_HUMAN]
E9PNW4	Uncharacterized protein OS=Homo sapiens GN=CD59 PE=4 SV=1 - [E9PNW4_HUMAN]
E9PC70	Uncharacterized protein OS=Homo sapiens GN=CD82 PE=4 SV=1 - [E9PC70_HUMAN]
E7ESI2	Uncharacterized protein OS=Homo sapiens GN=CDK2 PE=3 SV=1 - [E7ESI2_HUMAN]
F5H5A6	Uncharacterized protein OS=Homo sapiens GN=CENPT PE=4 SV=1 - [F5H5A6_HUMAN]
F5GYE8	Uncharacterized protein OS=Homo sapiens GN=CEP112 PE=4 SV=1 - [F5GYE8_HUMAN]
E9PD62	Uncharacterized protein OS=Homo sapiens GN=CEP250 PE=4 SV=1 - [E9PD62_HUMAN]
E9PP50	Uncharacterized protein OS=Homo sapiens GN=CFL1 PE=4 SV=1 - [E9PP50_HUMAN]
C9J0A7	Uncharacterized protein OS=Homo sapiens GN=CHMP2B PE=4 SV=1 - [C9J0A7_HUMAN]
B7Z5Z4	Uncharacterized protein OS=Homo sapiens GN=CLCN4 PE=2 SV=1 - [B7Z5Z4_HUMAN]
E9PDB9	Uncharacterized protein OS=Homo sapiens GN=CLCN7 PE=4 SV=1 - [E9PDB9_HUMAN]
E7EQ42	Uncharacterized protein OS=Homo sapiens GN=CLIP1 PE=4 SV=2 - [E7EQ42_HUMAN]
B4DIN1	Uncharacterized protein OS=Homo sapiens GN=CLTA PE=2 SV=1 - [B4DIN1_HUMAN]
F5H6N3	Uncharacterized protein OS=Homo sapiens GN=CLTA PE=4 SV=1 - [F5H6N3_HUMAN]
A6NCT7	Uncharacterized protein OS=Homo sapiens GN=COL16A1 PE=4 SV=1 - [A6NCT7_HUMAN]
F5GX37	Uncharacterized protein OS=Homo sapiens GN=COL5A2 PE=4 SV=1 - [F5GX37_HUMAN]
F5GXH1	Uncharacterized protein OS=Homo sapiens GN=COPG PE=4 SV=1 - [F5GXH1_HUMAN]
B4DY81	Uncharacterized protein OS=Homo sapiens GN=COPS3 PE=2 SV=1 - [B4DY81_HUMAN]
E7EM64	Uncharacterized protein OS=Homo sapiens GN=COPS6 PE=4 SV=1 - [E7EM64_HUMAN]
F5H4Z0	Uncharacterized protein OS=Homo sapiens GN=COPS7A PE=4 SV=1 - [F5H4Z0_HUMAN]
B4DHZ0	Uncharacterized protein OS=Homo sapiens GN=COPZ1 PE=2 SV=1 - [B4DHZ0_HUMAN]
A8MZ71	Uncharacterized protein OS=Homo sapiens GN=COX7A2 PE=4 SV=1 - [A8MZ71_HUMAN]
E5RJZ1	Uncharacterized protein OS=Homo sapiens GN=COX7A2L PE=4 SV=1 - [E5RJZ1_HUMAN]
D6R9Z7	Uncharacterized protein OS=Homo sapiens GN=COX7C PE=4 SV=1 - [D6R9Z7_HUMAN]
E9PDY4	Uncharacterized protein OS=Homo sapiens GN=CR1 PE=4 SV=1 - [E9PDY4_HUMAN]
E7EU96	Uncharacterized protein OS=Homo sapiens GN=CSNK2A1 PE=4 SV=1 - [E7EU96_HUMAN]
E9PP21	Uncharacterized protein OS=Homo sapiens GN=CSR1 PE=4 SV=1 - [E9PP21_HUMAN]
C9IYG0	Uncharacterized protein OS=Homo sapiens GN=CTDSP1 PE=4 SV=1 - [C9IYG0_HUMAN]
C9IZ88	Uncharacterized protein OS=Homo sapiens GN=CTNNA2 PE=4 SV=1 - [C9IZ88_HUMAN]
F5GWW0	Uncharacterized protein OS=Homo sapiens GN=CTSB PE=3 SV=1 - [F5GWW0_HUMAN]
E9PI30	Uncharacterized protein OS=Homo sapiens GN=CTSW PE=3 SV=1 - [E9PI30_HUMAN]
B4E358	Uncharacterized protein OS=Homo sapiens GN=CTTN PE=2 SV=1 - [B4E358_HUMAN]
E7EWR0	Uncharacterized protein OS=Homo sapiens GN=CUL1 PE=3 SV=1 - [E7EWR0_HUMAN]
A8MSH7	Uncharacterized protein OS=Homo sapiens GN=CUL4A PE=3 SV=1 - [A8MSH7_HUMAN]
D6RFH4	Uncharacterized protein OS=Homo sapiens GN=CYB5B PE=3 SV=1 - [D6RFH4_HUMAN]
B4DWD7	Uncharacterized protein OS=Homo sapiens GN=CYBRD1 PE=2 SV=1 - [B4DWD7_HUMAN]

C9JFR7	Uncharacterized protein OS=Homo sapiens GN=CYCS PE=3 SV=1 - [C9JFR7_HUMAN]
B4DDP6	Uncharacterized protein OS=Homo sapiens GN=DBNL PE=2 SV=1 - [B4DDP6_HUMAN]
B7Z8C9	Uncharacterized protein OS=Homo sapiens GN=DCAF8 PE=2 SV=1 - [B7Z8C9_HUMAN]
E9PFS5	Uncharacterized protein OS=Homo sapiens GN=DCTN1 PE=4 SV=1 - [E9PFS5_HUMAN]
E5RI97	Uncharacterized protein OS=Homo sapiens GN=DCTN4 PE=4 SV=1 - [E5RI97_HUMAN]
E7EWW8	Uncharacterized protein OS=Homo sapiens GN=DCTN5 PE=4 SV=1 - [E7EWW8_HUMAN]
C9J8R4	Uncharacterized protein OS=Homo sapiens GN=DCUN1D1 PE=4 SV=1 - [C9J8R4_HUMAN]
E7EWT1	Uncharacterized protein OS=Homo sapiens GN=DDOST PE=4 SV=1 - [E7EWT1_HUMAN]
B4DME8	Uncharacterized protein OS=Homo sapiens GN=DDX1 PE=2 SV=1 - [B4DME8_HUMAN]
B4E3E8	Uncharacterized protein OS=Homo sapiens GN=DDX3X PE=2 SV=1 - [B4E3E8_HUMAN]
B4E132	Uncharacterized protein OS=Homo sapiens GN=DDX3Y PE=2 SV=1 - [B4E132_HUMAN]
B7Z6B8	Uncharacterized protein OS=Homo sapiens GN=DECR1 PE=2 SV=1 - [B7Z6B8_HUMAN]
F5H853	Uncharacterized protein OS=Homo sapiens GN=DEF6 PE=4 SV=1 - [F5H853_HUMAN]
F5H0H1	Uncharacterized protein OS=Homo sapiens GN=DHCR24 PE=4 SV=1 - [F5H0H1_HUMAN]
E7ERW8	Uncharacterized protein OS=Homo sapiens GN=DIAPH1 PE=4 SV=1 - [E7ERW8_HUMAN]
E7EMA5	Uncharacterized protein OS=Homo sapiens GN=DIP2A PE=4 SV=2 - [E7EMA5_HUMAN]
F5H7M3	Uncharacterized protein OS=Homo sapiens GN=DLAT PE=3 SV=1 - [F5H7M3_HUMAN]
E7EQD7	Uncharacterized protein OS=Homo sapiens GN=DLG1 PE=4 SV=1 - [E7EQD7_HUMAN]
E7EMZ0	Uncharacterized protein OS=Homo sapiens GN=DMXL1 PE=4 SV=1 - [E7EMZ0_HUMAN]
F5GWV1	Uncharacterized protein OS=Homo sapiens GN=DNAJA4 PE=4 SV=1 - [F5GWV1_HUMAN]
C9JXB9	Uncharacterized protein OS=Homo sapiens GN=DNAJB2 PE=4 SV=1 - [C9JXB9_HUMAN]
E9PEQ4	Uncharacterized protein OS=Homo sapiens GN=DNM2 PE=3 SV=1 - [E9PEQ4_HUMAN]
E3W974	Uncharacterized protein OS=Homo sapiens GN=DNPEP PE=4 SV=2 - [E3W974_HUMAN]
E7EV00	Uncharacterized protein OS=Homo sapiens GN=DOCK10 PE=4 SV=2 - [E7EV00_HUMAN]
C9JDB3	Uncharacterized protein OS=Homo sapiens GN=DOCK4 PE=4 SV=1 - [C9JDB3_HUMAN]
E9PNX5	Uncharacterized protein OS=Homo sapiens GN=DPP3 PE=4 SV=1 - [E9PNX5_HUMAN]
B4DIZ0	Uncharacterized protein OS=Homo sapiens GN=DRP2 PE=2 SV=1 - [B4DIZ0_HUMAN]
E7ERU0	Uncharacterized protein OS=Homo sapiens GN=DST PE=4 SV=1 - [E7ERU0_HUMAN]
E7ETW8	Uncharacterized protein OS=Homo sapiens GN=DYSF PE=4 SV=1 - [E7ETW8_HUMAN]
F1LLU7	Uncharacterized protein OS=Homo sapiens GN=ECI2 PE=4 SV=1 - [F1LLU7_HUMAN]
A6NJH9	Uncharacterized protein OS=Homo sapiens GN=EIF1AY PE=4 SV=1 - [A6NJH9_HUMAN]
F5H6W1	Uncharacterized protein OS=Homo sapiens GN=EIF2B4 PE=3 SV=1 - [F5H6W1_HUMAN]
B4DV79	Uncharacterized protein OS=Homo sapiens GN=EIF3B PE=2 SV=1 - [B4DV79_HUMAN]
B4DVQ5	Uncharacterized protein OS=Homo sapiens GN=EIF3C PE=2 SV=1 - [B4DVQ5_HUMAN]
B4DYY1	Uncharacterized protein OS=Homo sapiens GN=EIF3D PE=2 SV=1 - [B4DYY1_HUMAN]
B7ZAM9	Uncharacterized protein OS=Homo sapiens GN=EIF3K PE=2 SV=1 - [B7ZAM9_HUMAN]
E7EQG2	Uncharacterized protein OS=Homo sapiens GN=EIF4A2 PE=3 SV=1 - [E7EQG2_HUMAN]
E7EPC9	Uncharacterized protein OS=Homo sapiens GN=EIF4B PE=4 SV=1 - [E7EPC9_HUMAN]
E7EX73	Uncharacterized protein OS=Homo sapiens GN=EIF4G1 PE=4 SV=1 - [E7EX73_HUMAN]
E9PCV8	Uncharacterized protein OS=Homo sapiens GN=EIF4G2 PE=4 SV=1 - [E9PCV8_HUMAN]
C9JIW2	Uncharacterized protein OS=Homo sapiens GN=ELMO1 PE=4 SV=1 - [C9JIW2_HUMAN]
D6RBS5	Uncharacterized protein OS=Homo sapiens GN=ELMOD2 PE=4 SV=1 - [D6RBS5_HUMAN]
E9PEW9	Uncharacterized protein OS=Homo sapiens GN=EPB41 PE=4 SV=1 - [E9PEW9_HUMAN]

E9PEX0	Uncharacterized protein OS=Homo sapiens GN=EPB41 PE=4 SV=1 - [E9PEX0_HUMAN]
E9PPD9	Uncharacterized protein OS=Homo sapiens GN=EPB41L2 PE=4 SV=1 - [E9PPD9_HUMAN]
A8K968	Uncharacterized protein OS=Homo sapiens GN=EPB41L3 PE=2 SV=1 - [A8K968_HUMAN]
E9PNT0	Uncharacterized protein OS=Homo sapiens GN=EPS8L2 PE=4 SV=1 - [E9PNT0_HUMAN]
F5GZU8	Uncharacterized protein OS=Homo sapiens GN=ERC1 PE=4 SV=1 - [F5GZU8_HUMAN]
E5RHW4	Uncharacterized protein OS=Homo sapiens GN=ERLIN2 PE=4 SV=1 - [E5RHW4_HUMAN]
B4DT43	Uncharacterized protein OS=Homo sapiens GN=ETFA PE=2 SV=1 - [B4DT43_HUMAN]
C9JT69	Uncharacterized protein OS=Homo sapiens GN=ETFDH PE=4 SV=1 - [C9JT69_HUMAN]
E9PED2	Uncharacterized protein OS=Homo sapiens GN=EXOC4 PE=4 SV=1 - [E9PED2_HUMAN]
B4DEJ6	Uncharacterized protein OS=Homo sapiens GN=FAF1 PE=2 SV=1 - [B4DEJ6_HUMAN]
B4E2M8	Uncharacterized protein OS=Homo sapiens GN=FAF2 PE=2 SV=1 - [B4E2M8_HUMAN]
D6REQ2	Uncharacterized protein OS=Homo sapiens GN=FAM193B PE=4 SV=2 - [D6REQ2_HUMAN]
F5H817	Uncharacterized protein OS=Homo sapiens GN=FAT3 PE=4 SV=1 - [F5H817_HUMAN]
E9PG79	Uncharacterized protein OS=Homo sapiens GN=FCHO2 PE=4 SV=1 - [E9PG79_HUMAN]
B7Z7G2	Uncharacterized protein OS=Homo sapiens GN=FDXR PE=2 SV=1 - [B7Z7G2_HUMAN]
B4DVN5	Uncharacterized protein OS=Homo sapiens GN=FHOD1 PE=2 SV=1 - [B4DVN5_HUMAN]
C9JN41	Uncharacterized protein OS=Homo sapiens GN=FILIP1L PE=4 SV=1 - [C9JN41_HUMAN]
F5H0N4	Uncharacterized protein OS=Homo sapiens GN=FKBP2 PE=3 SV=1 - [F5H0N4_HUMAN]
B7Z7Z8	Uncharacterized protein OS=Homo sapiens GN=FKBP5 PE=2 SV=1 - [B7Z7Z8_HUMAN]
F5H453	Uncharacterized protein OS=Homo sapiens GN=FLII PE=4 SV=1 - [F5H453_HUMAN]
A8MQB8	Uncharacterized protein OS=Homo sapiens GN=FMR1 PE=4 SV=1 - [A8MQB8_HUMAN]
B4DY49	Uncharacterized protein OS=Homo sapiens GN=FRK PE=2 SV=1 - [B4DY49_HUMAN]
B4E0X8	Uncharacterized protein OS=Homo sapiens GN=FUBP1 PE=2 SV=1 - [B4E0X8_HUMAN]
B4DR70	Uncharacterized protein OS=Homo sapiens GN=FUS PE=2 SV=1 - [B4DR70_HUMAN]
B3KPS6	Uncharacterized protein OS=Homo sapiens GN=FYN PE=2 SV=1 - [B3KPS6_HUMAN]
F5H4D6	Uncharacterized protein OS=Homo sapiens GN=G3BP1 PE=4 SV=1 - [F5H4D6_HUMAN]
D6RFJ8	Uncharacterized protein OS=Homo sapiens GN=G3BP2 PE=4 SV=1 - [D6RFJ8_HUMAN]
B7Z8V8	Uncharacterized protein OS=Homo sapiens GN=GALNT2 PE=2 SV=1 - [B7Z8V8_HUMAN]
E9PKU7	Uncharacterized protein OS=Homo sapiens GN=GANAB PE=4 SV=1 - [E9PKU7_HUMAN]
E7EPW1	Uncharacterized protein OS=Homo sapiens GN=GAPDH PE=3 SV=1 - [E7EPW1_HUMAN]
C9JPA8	Uncharacterized protein OS=Homo sapiens GN=GET4 PE=4 SV=1 - [C9JPA8_HUMAN]
F5GYQ4	Uncharacterized protein OS=Homo sapiens GN=GLUD1 PE=4 SV=1 - [F5GYQ4_HUMAN]
F5H7W9	Uncharacterized protein OS=Homo sapiens GN=GOLPH3 PE=4 SV=1 - [F5H7W9_HUMAN]
F5GWK9	Uncharacterized protein OS=Homo sapiens GN=GPR116 PE=4 SV=1 - [F5GWK9_HUMAN]
A8MZ27	Uncharacterized protein OS=Homo sapiens GN=GRID2IP PE=4 SV=2 - [A8MZ27_HUMAN]
E7EPY9	Uncharacterized protein OS=Homo sapiens GN=GRIK1 PE=4 SV=1 - [E7EPY9_HUMAN]
E7EQZ3	Uncharacterized protein OS=Homo sapiens GN=GSPT1 PE=4 SV=1 - [E7EQZ3_HUMAN]
C9J8R8	Uncharacterized protein OS=Homo sapiens GN=GYG1 PE=4 SV=1 - [C9J8R8_HUMAN]
C9IYM4	Uncharacterized protein OS=Homo sapiens GN=GYS1 PE=4 SV=1 - [C9IYM4_HUMAN]
E9PD60	Uncharacterized protein OS=Homo sapiens GN=HARS2 PE=4 SV=1 - [E9PD60_HUMAN]
C9JBZ4	Uncharacterized protein OS=Homo sapiens GN=HAUS8 PE=4 SV=1 - [C9JBZ4_HUMAN]
E9PJ90	Uncharacterized protein OS=Homo sapiens GN=HBS1L PE=4 SV=1 - [E9PJ90_HUMAN]
E9PLX3	Uncharacterized protein OS=Homo sapiens GN=HBXIP PE=4 SV=1 - [E9PLX3_HUMAN]

E7ERH6	Uncharacterized protein OS=Homo sapiens GN=HDAC6 PE=4 SV=1 - [E7ERH6_HUMAN]
C9JJI9	Uncharacterized protein OS=Homo sapiens GN=HERC6 PE=4 SV=1 - [C9JJI9_HUMAN]
F5H5Y5	Uncharacterized protein OS=Homo sapiens GN=HGS PE=4 SV=1 - [F5H5Y5_HUMAN]
E9PL41	Uncharacterized protein OS=Homo sapiens GN=HIF1AN PE=4 SV=1 - [E9PL41_HUMAN]
E9PCK0	Uncharacterized protein OS=Homo sapiens GN=HK1 PE=3 SV=1 - [E9PCK0_HUMAN]
F5H345	Uncharacterized protein OS=Homo sapiens GN=HMBS PE=4 SV=1 - [F5H345_HUMAN]
F2Z2W6	Uncharacterized protein OS=Homo sapiens GN=HMGN1 PE=4 SV=1 - [F2Z2W6_HUMAN]
E9PEK0	Uncharacterized protein OS=Homo sapiens GN=HMGXB3 PE=4 SV=1 - [E9PEK0_HUMAN]
A6NGP5	Uncharacterized protein OS=Homo sapiens GN=HN1L PE=2 SV=2 - [A6NGP5_HUMAN]
D6R9P3	Uncharacterized protein OS=Homo sapiens GN=HNRNPAB PE=4 SV=1 - [D6R9P3_HUMAN]
B2R603	Uncharacterized protein OS=Homo sapiens GN=HNRNPC PE=2 SV=1 - [B2R603_HUMAN]
B4DTC3	Uncharacterized protein OS=Homo sapiens GN=HNRNPD PE=2 SV=1 - [B4DTC3_HUMAN]
E9PCY7	Uncharacterized protein OS=Homo sapiens GN=HNRNPH1 PE=4 SV=1 - [E9PCY7_HUMAN]
B3KX72	Uncharacterized protein OS=Homo sapiens GN=HNRNPU PE=2 SV=1 - [B3KX72_HUMAN]
E9PDE8	Uncharacterized protein OS=Homo sapiens GN=HSPA4L PE=3 SV=1 - [E9PDE8_HUMAN]
B7Z4V2	Uncharacterized protein OS=Homo sapiens GN=HSPA9 PE=2 SV=1 - [B7Z4V2_HUMAN]
B4DY72	Uncharacterized protein OS=Homo sapiens GN=HSPH1 PE=2 SV=1 - [B4DY72_HUMAN]
E9PD66	Uncharacterized protein OS=Homo sapiens GN=HUWE1 PE=4 SV=1 - [E9PD66_HUMAN]
E9PDR5	Uncharacterized protein OS=Homo sapiens GN=IBTK PE=4 SV=1 - [E9PDR5_HUMAN]
B7Z9J8	Uncharacterized protein OS=Homo sapiens GN=IDH3A PE=2 SV=1 - [B7Z9J8_HUMAN]
E9PDD5	Uncharacterized protein OS=Homo sapiens GN=IDH3G PE=3 SV=1 - [E9PDD5_HUMAN]
F5H0L4	Uncharacterized protein OS=Homo sapiens GN=IER5 PE=4 SV=1 - [F5H0L4_HUMAN]
C9JT33	Uncharacterized protein OS=Homo sapiens GN=IGF2BP1 PE=4 SV=1 - [C9JT33_HUMAN]
D6RGV7	Uncharacterized protein OS=Homo sapiens GN=IGSF22 PE=4 SV=1 - [D6RGV7_HUMAN]
C9JZ85	Uncharacterized protein OS=Homo sapiens GN=IKBKG PE=4 SV=1 - [C9JZ85_HUMAN]
B4DFG4	Uncharacterized protein OS=Homo sapiens GN=ILF3 PE=2 SV=1 - [B4DFG4_HUMAN]
C9J406	Uncharacterized protein OS=Homo sapiens GN=IMMT PE=4 SV=1 - [C9J406_HUMAN]
C9JUT0	Uncharacterized protein OS=Homo sapiens GN=ING3 PE=4 SV=1 - [C9JUT0_HUMAN]
E7EMB7	Uncharacterized protein OS=Homo sapiens GN=IPO11 PE=4 SV=1 - [E7EMB7_HUMAN]
C9JL81	Uncharacterized protein OS=Homo sapiens GN=IQCD PE=4 SV=1 - [C9JL81_HUMAN]
F5H7A1	Uncharacterized protein OS=Homo sapiens GN=IQGAP1 PE=4 SV=1 - [F5H7A1_HUMAN]
E7EWZ6	Uncharacterized protein OS=Homo sapiens GN=ITGAV PE=3 SV=1 - [E7EWZ6_HUMAN]
F5GXQ1	Uncharacterized protein OS=Homo sapiens GN=ITIH4 PE=4 SV=1 - [F5GXQ1_HUMAN]
F5H153	Uncharacterized protein OS=Homo sapiens GN=ITSN1 PE=4 SV=1 - [F5H153_HUMAN]
E7EUQ1	Uncharacterized protein OS=Homo sapiens GN=ITSN2 PE=4 SV=1 - [E7EUQ1_HUMAN]
E9PDU1	Uncharacterized protein OS=Homo sapiens GN=KARS PE=3 SV=2 - [E9PDU1_HUMAN]
C9J3I2	Uncharacterized protein OS=Homo sapiens GN=KCMF1 PE=4 SV=1 - [C9J3I2_HUMAN]
Q6ZW95	Uncharacterized protein OS=Homo sapiens GN=KCNK2 PE=2 SV=1 - [Q6ZW95_HUMAN]
F5H275	Uncharacterized protein OS=Homo sapiens GN=KDM3B PE=4 SV=1 - [F5H275_HUMAN]
F5GYH2	Uncharacterized protein OS=Homo sapiens GN=KDM4C PE=4 SV=1 - [F5GYH2_HUMAN]
F5H5V6	Uncharacterized protein OS=Homo sapiens GN=KDM6A PE=4 SV=1 - [F5H5V6_HUMAN]
C9JCC4	Uncharacterized protein OS=Homo sapiens GN=KIAA1199 PE=4 SV=1 - [C9JCC4_HUMAN]
F5H2Q7	Uncharacterized protein OS=Homo sapiens GN=KIAA1715 PE=4 SV=1 - [F5H2Q7_HUMAN]

E9PFS1	Uncharacterized protein OS=Homo sapiens GN=KIAA1755 PE=4 SV=1 - [E9PFS1_HUMAN]
E9PKB8	Uncharacterized protein OS=Homo sapiens GN=KIAA1875 PE=4 SV=1 - [E9PKB8_HUMAN]
C9JKA9	Uncharacterized protein OS=Homo sapiens GN=KIF15 PE=4 SV=1 - [C9JKA9_HUMAN]
B7WPD9	Uncharacterized protein OS=Homo sapiens GN=KIF26B PE=3 SV=1 - [B7WPD9_HUMAN]
F5GYJ7	Uncharacterized protein OS=Homo sapiens GN=KIF3C PE=3 SV=1 - [F5GYJ7_HUMAN]
B4DFZ5	Uncharacterized protein OS=Homo sapiens GN=KLHL2 PE=2 SV=1 - [B4DFZ5_HUMAN]
C9J191	Uncharacterized protein OS=Homo sapiens GN=KLHL22 PE=4 SV=1 - [C9J191_HUMAN]
D6RH21	Uncharacterized protein OS=Homo sapiens GN=KLHL3 PE=4 SV=1 - [D6RH21_HUMAN]
F2Z383	Uncharacterized protein OS=Homo sapiens GN=KLRAQ1 PE=4 SV=1 - [F2Z383_HUMAN]
C9JYI4	Uncharacterized protein OS=Homo sapiens GN=KPNA1 PE=4 SV=1 - [C9JYI4_HUMAN]
E9PCN9	Uncharacterized protein OS=Homo sapiens GN=KPNA3 PE=4 SV=1 - [E9PCN9_HUMAN]
B7Z2R9	Uncharacterized protein OS=Homo sapiens GN=LAMP2 PE=2 SV=1 - [B7Z2R9_HUMAN]
F5H479	Uncharacterized protein OS=Homo sapiens GN=LAMTOR1 PE=4 SV=1 - [F5H479_HUMAN]
E9PHS0	Uncharacterized protein OS=Homo sapiens GN=LANCL1 PE=4 SV=1 - [E9PHS0_HUMAN]
B4DJ10	Uncharacterized protein OS=Homo sapiens GN=LARS PE=2 SV=1 - [B4DJ10_HUMAN]
B4E135	Uncharacterized protein OS=Homo sapiens GN=LIG1 PE=2 SV=1 - [B4E135_HUMAN]
B4DWN1	Uncharacterized protein OS=Homo sapiens GN=LMAN2 PE=2 SV=1 - [B4DWN1_HUMAN]
E9PBF6	Uncharacterized protein OS=Homo sapiens GN=LMNB1 PE=3 SV=1 - [E9PBF6_HUMAN]
F5GZH4	Uncharacterized protein OS=Homo sapiens GN=LNX1 PE=4 SV=1 - [F5GZH4_HUMAN]
B3KXS5	Uncharacterized protein OS=Homo sapiens GN=LONP1 PE=2 SV=1 - [B3KXS5_HUMAN]
B7Z3N3	Uncharacterized protein OS=Homo sapiens GN=LPCAT3 PE=2 SV=1 - [B7Z3N3_HUMAN]
B4DS68	Uncharacterized protein OS=Homo sapiens GN=LRP11 PE=2 SV=1 - [B4DS68_HUMAN]
B4DHL3	Uncharacterized protein OS=Homo sapiens GN=LRRRC28 PE=2 SV=1 - [B4DHL3_HUMAN]
C9JI57	Uncharacterized protein OS=Homo sapiens GN=LRRIQ1 PE=4 SV=1 - [C9JI57_HUMAN]
E9PN53	Uncharacterized protein OS=Homo sapiens GN=LUZP2 PE=4 SV=1 - [E9PN53_HUMAN]
C9J1B9	Uncharacterized protein OS=Homo sapiens GN=LYSMD4 PE=4 SV=1 - [C9J1B9_HUMAN]
F5GXE0	Uncharacterized protein OS=Homo sapiens GN=M6PR PE=4 SV=1 - [F5GXE0_HUMAN]
D6RID6	Uncharacterized protein OS=Homo sapiens GN=MAEA PE=4 SV=1 - [D6RID6_HUMAN]
B4DH53	Uncharacterized protein OS=Homo sapiens GN=MAP1S PE=2 SV=1 - [B4DH53_HUMAN]
B5MEG9	Uncharacterized protein OS=Homo sapiens GN=MAP4 PE=4 SV=2 - [B5MEG9_HUMAN]
E7EX83	Uncharacterized protein OS=Homo sapiens GN=MAP4K4 PE=3 SV=1 - [E7EX83_HUMAN]
E9PFH7	Uncharacterized protein OS=Homo sapiens GN=MAPK8IP3 PE=4 SV=1 - [E9PFH7_HUMAN]
E7ETY4	Uncharacterized protein OS=Homo sapiens GN=MARK2 PE=4 SV=1 - [E7ETY4_HUMAN]
E7ET05	Uncharacterized protein OS=Homo sapiens GN=MARS PE=4 SV=1 - [E7ET05_HUMAN]
F5H6C8	Uncharacterized protein OS=Homo sapiens GN=MAVS PE=4 SV=1 - [F5H6C8_HUMAN]
C9J600	Uncharacterized protein OS=Homo sapiens GN=MCM10 PE=4 SV=1 - [C9J600_HUMAN]
F5H3Y1	Uncharacterized protein OS=Homo sapiens GN=MED12 PE=4 SV=1 - [F5H3Y1_HUMAN]
E7EQ94	Uncharacterized protein OS=Homo sapiens GN=MET PE=4 SV=2 - [E7EQ94_HUMAN]
B4E2X3	Uncharacterized protein OS=Homo sapiens GN=METTL13 PE=2 SV=1 - [B4E2X3_HUMAN]
E9PGF9	Uncharacterized protein OS=Homo sapiens GN=MGEA5 PE=4 SV=1 - [E9PGF9_HUMAN]
F5H4V4	Uncharacterized protein OS=Homo sapiens GN=MKI67 PE=4 SV=1 - [F5H4V4_HUMAN]
C9JI01	Uncharacterized protein OS=Homo sapiens GN=MLPH PE=4 SV=1 - [C9JI01_HUMAN]
F5H079	Uncharacterized protein OS=Homo sapiens GN=MMAB PE=4 SV=1 - [F5H079_HUMAN]

B4DWL9	Uncharacterized protein OS=Homo sapiens GN=MPP7 PE=2 SV=1 - [B4DWL9_HUMAN]
D6RC14	Uncharacterized protein OS=Homo sapiens GN=MRPL3 PE=3 SV=1 - [D6RC14_HUMAN]
F5H2F9	Uncharacterized protein OS=Homo sapiens GN=MSH6 PE=3 SV=1 - [F5H2F9_HUMAN]
E9PNT8	Uncharacterized protein OS=Homo sapiens GN=MTCH2 PE=3 SV=1 - [E9PNT8_HUMAN]
B4DZD5	Uncharacterized protein OS=Homo sapiens GN=MTOR PE=2 SV=1 - [B4DZD5_HUMAN]
C9JNK6	Uncharacterized protein OS=Homo sapiens GN=MTX2 PE=4 SV=1 - [C9JNK6_HUMAN]
B4DRZ1	Uncharacterized protein OS=Homo sapiens GN=MYCN PE=2 SV=1 - [B4DRZ1_HUMAN]
F5H678	Uncharacterized protein OS=Homo sapiens GN=MYO5B PE=4 SV=1 - [F5H678_HUMAN]
F5GWA6	Uncharacterized protein OS=Homo sapiens GN=MYPN PE=4 SV=1 - [F5GWA6_HUMAN]
A8MZ22	Uncharacterized protein OS=Homo sapiens GN=NAP1L4 PE=3 SV=3 - [A8MZ22_HUMAN]
E7EUA9	Uncharacterized protein OS=Homo sapiens GN=NAV3 PE=4 SV=2 - [E7EUA9_HUMAN]
F5GZJ1	Uncharacterized protein OS=Homo sapiens GN=NCAPD2 PE=4 SV=1 - [F5GZJ1_HUMAN]
B4E189	Uncharacterized protein OS=Homo sapiens GN=NCAPH PE=2 SV=1 - [B4E189_HUMAN]
E7EX81	Uncharacterized protein OS=Homo sapiens GN=NCL PE=4 SV=1 - [E7EX81_HUMAN]
E9PC09	Uncharacterized protein OS=Homo sapiens GN=NCSTN PE=4 SV=1 - [E9PC09_HUMAN]
F5H1J6	Uncharacterized protein OS=Homo sapiens GN=NDC80 PE=4 SV=1 - [F5H1J6_HUMAN]
C9IZN5	Uncharacterized protein OS=Homo sapiens GN=NDUFA5 PE=4 SV=1 - [C9IZN5_HUMAN]
F5H7V7	Uncharacterized protein OS=Homo sapiens GN=NDUFA8 PE=4 SV=1 - [F5H7V7_HUMAN]
E9PH64	Uncharacterized protein OS=Homo sapiens GN=NDUFB9 PE=4 SV=1 - [E9PH64_HUMAN]
B4DJ81	Uncharacterized protein OS=Homo sapiens GN=NDUFS1 PE=2 SV=1 - [B4DJ81_HUMAN]
E9PMX3	Uncharacterized protein OS=Homo sapiens GN=NDUFV1 PE=4 SV=1 - [E9PMX3_HUMAN]
E9PGQ8	Uncharacterized protein OS=Homo sapiens GN=NECAP2 PE=4 SV=1 - [E9PGQ8_HUMAN]
D6RBG5	Uncharacterized protein OS=Homo sapiens GN=NEK1 PE=4 SV=1 - [D6RBG5_HUMAN]
C9J5I7	Uncharacterized protein OS=Homo sapiens GN=NFKBIZ PE=4 SV=1 - [C9J5I7_HUMAN]
F2Z2E7	Uncharacterized protein OS=Homo sapiens GN=NFS1 PE=3 SV=1 - [F2Z2E7_HUMAN]
E7ERL0	Uncharacterized protein OS=Homo sapiens GN=NME1 PE=3 SV=1 - [E7ERL0_HUMAN]
B7Z8J4	Uncharacterized protein OS=Homo sapiens GN=NMT1 PE=2 SV=1 - [B7Z8J4_HUMAN]
F5GZD4	Uncharacterized protein OS=Homo sapiens GN=NPC1 PE=4 SV=1 - [F5GZD4_HUMAN]
B4DQV7	Uncharacterized protein OS=Homo sapiens GN=NPC2 PE=2 SV=1 - [B4DQV7_HUMAN]
E9PN47	Uncharacterized protein OS=Homo sapiens GN=NPEPL1 PE=4 SV=1 - [E9PN47_HUMAN]
B4DG89	Uncharacterized protein OS=Homo sapiens GN=NPLOC4 PE=2 SV=1 - [B4DG89_HUMAN]
E9PR11	Uncharacterized protein OS=Homo sapiens GN=NRBP2 PE=4 SV=1 - [E9PR11_HUMAN]
C9JDR0	Uncharacterized protein OS=Homo sapiens GN=NSDHL PE=3 SV=1 - [C9JDR0_HUMAN]
D6RAS3	Uncharacterized protein OS=Homo sapiens GN=NSUN2 PE=4 SV=1 - [D6RAS3_HUMAN]
E9PJP3	Uncharacterized protein OS=Homo sapiens GN=NUCB2 PE=4 SV=1 - [E9PJP3_HUMAN]
F5H2D1	Uncharacterized protein OS=Homo sapiens GN=ODZ2 PE=4 SV=1 - [F5H2D1_HUMAN]
E9PFG7	Uncharacterized protein OS=Homo sapiens GN=OGDH PE=4 SV=1 - [E9PFG7_HUMAN]
E5RFY7	Uncharacterized protein OS=Homo sapiens GN=OSTM1 PE=4 SV=1 - [E5RFY7_HUMAN]
C9JJI8	Uncharacterized protein OS=Homo sapiens GN=PAFAH1B1 PE=4 SV=1 - [C9JJI8_HUMAN]
E9PBS1	Uncharacterized protein OS=Homo sapiens GN=PAICS PE=4 SV=1 - [E9PBS1_HUMAN]
E9PHT6	Uncharacterized protein OS=Homo sapiens GN=PANK4 PE=4 SV=1 - [E9PHT6_HUMAN]
C9J7L3	Uncharacterized protein OS=Homo sapiens GN=PARP15 PE=4 SV=2 - [C9J7L3_HUMAN]
B7Z3X5	Uncharacterized protein OS=Homo sapiens GN=PDHA1 PE=2 SV=1 - [B7Z3X5_HUMAN]

B4DDD7	Uncharacterized protein OS=Homo sapiens GN=PDHB PE=2 SV=1 - [B4DDD7_HUMAN]
B3KQT9	Uncharacterized protein OS=Homo sapiens GN=PDIA3 PE=2 SV=1 - [B3KQT9_HUMAN]
B7Z254	Uncharacterized protein OS=Homo sapiens GN=PDIA6 PE=2 SV=1 - [B7Z254_HUMAN]
D6RF83	Uncharacterized protein OS=Homo sapiens GN=PDLM7 PE=4 SV=1 - [D6RF83_HUMAN]
E7ENB6	Uncharacterized protein OS=Homo sapiens GN=PDZRN3 PE=4 SV=1 - [E7ENB6_HUMAN]
E9PCE8	Uncharacterized protein OS=Homo sapiens GN=PEPD PE=3 SV=1 - [E9PCE8_HUMAN]
E5RGS4	Uncharacterized protein OS=Homo sapiens GN=PFDN1 PE=4 SV=1 - [E5RGS4_HUMAN]
C9JW96	Uncharacterized protein OS=Homo sapiens GN=PHB PE=4 SV=2 - [C9JW96_HUMAN]
F2Z3N8	Uncharacterized protein OS=Homo sapiens GN=PHF16 PE=4 SV=1 - [F2Z3N8_HUMAN]
B4DM11	Uncharacterized protein OS=Homo sapiens GN=PIP4K2C PE=2 SV=1 - [B4DM11_HUMAN]
E9PBQ1	Uncharacterized protein OS=Homo sapiens GN=PIP5K1A PE=4 SV=1 - [E9PBQ1_HUMAN]
F5H0R5	Uncharacterized protein OS=Homo sapiens GN=PIP5K1C PE=4 SV=1 - [F5H0R5_HUMAN]
F5H5E2	Uncharacterized protein OS=Homo sapiens GN=PLBD2 PE=4 SV=1 - [F5H5E2_HUMAN]
C9J228	Uncharacterized protein OS=Homo sapiens GN=PLCH1 PE=4 SV=1 - [C9J228_HUMAN]
B4DTE8	Uncharacterized protein OS=Homo sapiens GN=PLSCR1 PE=2 SV=1 - [B4DTE8_HUMAN]
F5H5K9	Uncharacterized protein OS=Homo sapiens GN=PNPLA6 PE=4 SV=1 - [F5H5K9_HUMAN]
D6R967	Uncharacterized protein OS=Homo sapiens GN=PPA2 PE=4 SV=1 - [D6R967_HUMAN]
F5H158	Uncharacterized protein OS=Homo sapiens GN=PPFIA2 PE=4 SV=1 - [F5H158_HUMAN]
B4DY76	Uncharacterized protein OS=Homo sapiens GN=PPOX PE=2 SV=1 - [B4DY76_HUMAN]
F5H3X9	Uncharacterized protein OS=Homo sapiens GN=PPP2R1A PE=4 SV=1 - [F5H3X9_HUMAN]
F5GYS1	Uncharacterized protein OS=Homo sapiens GN=PPP2R5D PE=4 SV=1 - [F5GYS1_HUMAN]
E9PKF6	Uncharacterized protein OS=Homo sapiens GN=PPP6R3 PE=4 SV=1 - [E9PKF6_HUMAN]
E7EVG6	Uncharacterized protein OS=Homo sapiens GN=PPRC1 PE=4 SV=1 - [E7EVG6_HUMAN]
A6NP52	Uncharacterized protein OS=Homo sapiens GN=PRAF2 PE=4 SV=1 - [A6NP52_HUMAN]
E9PH29	Uncharacterized protein OS=Homo sapiens GN=PRDX3 PE=4 SV=1 - [E9PH29_HUMAN]
E9PEN3	Uncharacterized protein OS=Homo sapiens GN=PRDX4 PE=4 SV=1 - [E9PEN3_HUMAN]
A8K318	Uncharacterized protein OS=Homo sapiens GN=PRKCSH PE=2 SV=1 - [A8K318_HUMAN]
F5H180	Uncharacterized protein OS=Homo sapiens GN=PRKDC PE=4 SV=1 - [F5H180_HUMAN]
E7EPE6	Uncharacterized protein OS=Homo sapiens GN=PRKG2 PE=4 SV=1 - [E7EPE6_HUMAN]
E9PNR9	Uncharacterized protein OS=Homo sapiens GN=PRMT1 PE=4 SV=1 - [E9PNR9_HUMAN]
E5RG77	Uncharacterized protein OS=Homo sapiens GN=PROSC PE=3 SV=1 - [E5RG77_HUMAN]
C9J168	Uncharacterized protein OS=Homo sapiens GN=PRPSAP1 PE=4 SV=1 - [C9J168_HUMAN]
F5GZ16	Uncharacterized protein OS=Homo sapiens GN=PSMD2 PE=4 SV=1 - [F5GZ16_HUMAN]
F5H8K4	Uncharacterized protein OS=Homo sapiens GN=PSMD3 PE=4 SV=1 - [F5H8K4_HUMAN]
B4DZM8	Uncharacterized protein OS=Homo sapiens GN=PSMD5 PE=2 SV=1 - [B4DZM8_HUMAN]
B4DXI8	Uncharacterized protein OS=Homo sapiens GN=PSMD7 PE=2 SV=1 - [B4DXI8_HUMAN]
F5H169	Uncharacterized protein OS=Homo sapiens GN=PSMD9 PE=4 SV=1 - [F5H169_HUMAN]
F5H656	Uncharacterized protein OS=Homo sapiens GN=PSPC1 PE=4 SV=1 - [F5H656_HUMAN]
B8ZZI3	Uncharacterized protein OS=Homo sapiens GN=PTMA PE=4 SV=1 - [B8ZZI3_HUMAN]
B4DSN5	Uncharacterized protein OS=Homo sapiens GN=PTPN1 PE=2 SV=1 - [B4DSN5_HUMAN]
F5H0X4	Uncharacterized protein OS=Homo sapiens GN=PTPRE PE=4 SV=1 - [F5H0X4_HUMAN]
A8MZH8	Uncharacterized protein OS=Homo sapiens GN=PTTG1IP PE=4 SV=1 - [A8MZH8_HUMAN]
B5MD87	Uncharacterized protein OS=Homo sapiens GN=PYCRL PE=4 SV=2 - [B5MD87_HUMAN]
B4DNN3	Uncharacterized protein OS=Homo sapiens GN=QARS PE=2 SV=1 - [B4DNN3_HUMAN]
E9PK89	Uncharacterized protein OS=Homo sapiens GN=RAB3IL1 PE=4 SV=1 - [E9PK89_HUMAN]
B4DJA5	Uncharacterized protein OS=Homo sapiens GN=RAB5A PE=2 SV=1 - [B4DJA5_HUMAN]
F5GZB1	Uncharacterized protein OS=Homo sapiens GN=RAB6A PE=3 SV=1 - [F5GZB1_HUMAN]
F5GZU7	Uncharacterized protein OS=Homo sapiens GN=RABEP1 PE=4 SV=1 - [F5GZU7_HUMAN]
C9JXM3	Uncharacterized protein OS=Homo sapiens GN=RABL3 PE=4 SV=1 - [C9JXM3_HUMAN]
B4DQI8	Uncharacterized protein OS=Homo sapiens GN=RAP1B PE=2 SV=1 - [B4DQI8_HUMAN]
E9PH06	Uncharacterized protein OS=Homo sapiens GN=RAP1GDS1 PE=4 SV=1 - [E9PH06_HUMAN]
Q8N8Y7	Uncharacterized protein OS=Homo sapiens GN=RBMX PE=2 SV=1 - [Q8N8Y7_HUMAN]
B7Z1M1	Uncharacterized protein OS=Homo sapiens GN=RCN1 PE=2 SV=1 - [B7Z1M1_HUMAN]

B7ZKW8	Uncharacterized protein OS=Homo sapiens GN=RCS1 PE=2 SV=1 - [B7ZKW8_HUMAN]
D6RBN9	Uncharacterized protein OS=Homo sapiens GN=RELL1 PE=4 SV=1 - [D6RBN9_HUMAN]
A6NKZ2	Uncharacterized protein OS=Homo sapiens GN=RENBP PE=4 SV=2 - [A6NKZ2_HUMAN]
C9JE27	Uncharacterized protein OS=Homo sapiens GN=RFLL PE=4 SV=1 - [C9JE27_HUMAN]
F5H250	Uncharacterized protein OS=Homo sapiens GN=RHAG PE=4 SV=1 - [F5H250_HUMAN]
E7EWZ5	Uncharacterized protein OS=Homo sapiens GN=RHCE PE=4 SV=1 - [E7EWZ5_HUMAN]
E9PCZ1	Uncharacterized protein OS=Homo sapiens GN=RIMS1 PE=4 SV=1 - [E9PCZ1_HUMAN]
C9J5S3	Uncharacterized protein OS=Homo sapiens GN=RINT1 PE=4 SV=1 - [C9J5S3_HUMAN]
E9PKE9	Uncharacterized protein OS=Homo sapiens GN=RNF121 PE=4 SV=1 - [E9PKE9_HUMAN]
D6RI12	Uncharacterized protein OS=Homo sapiens GN=RNF213 PE=4 SV=1 - [D6RI12_HUMAN]
E9PMA9	Uncharacterized protein OS=Homo sapiens GN=RNH1 PE=4 SV=1 - [E9PMA9_HUMAN]
B4DM74	Uncharacterized protein OS=Homo sapiens GN=RPL18A PE=2 SV=1 - [B4DM74_HUMAN]
C9JXB8	Uncharacterized protein OS=Homo sapiens GN=RPL24 PE=4 SV=1 - [C9JXB8_HUMAN]
E9PLL6	Uncharacterized protein OS=Homo sapiens GN=RPL27A PE=3 SV=1 - [E9PLL6_HUMAN]
C9JU56	Uncharacterized protein OS=Homo sapiens GN=RPL31 PE=4 SV=1 - [C9JU56_HUMAN]
C9K025	Uncharacterized protein OS=Homo sapiens GN=RPL35A PE=4 SV=1 - [C9K025_HUMAN]
E9PEL3	Uncharacterized protein OS=Homo sapiens GN=RPL37A PE=4 SV=1 - [E9PEL3_HUMAN]
E9PIZ3	Uncharacterized protein OS=Homo sapiens GN=RPL8 PE=4 SV=1 - [E9PIZ3_HUMAN]
D6RAN4	Uncharacterized protein OS=Homo sapiens GN=RPL9 PE=4 SV=1 - [D6RAN4_HUMAN]
D6R910	Uncharacterized protein OS=Homo sapiens GN=RPS6KA2 PE=4 SV=1 - [D6R910_HUMAN]
C9J9K3	Uncharacterized protein OS=Homo sapiens GN=RPSA PE=3 SV=1 - [C9J9K3_HUMAN]
B7Z7H6	Uncharacterized protein OS=Homo sapiens GN=RRAS2 PE=2 SV=1 - [B7Z7H6_HUMAN]
F5H774	Uncharacterized protein OS=Homo sapiens GN=RTN3 PE=4 SV=1 - [F5H774_HUMAN]
E5RJE3	Uncharacterized protein OS=Homo sapiens GN=RWDD1 PE=4 SV=1 - [E5RJE3_HUMAN]
C9K0H7	Uncharacterized protein OS=Homo sapiens GN=RYS3 PE=4 SV=2 - [C9K0H7_HUMAN]
F5H3J5	Uncharacterized protein OS=Homo sapiens GN=S100A4 PE=4 SV=1 - [F5H3J5_HUMAN]
E9PGZ4	Uncharacterized protein OS=Homo sapiens GN=SACM1L PE=4 SV=1 - [E9PGZ4_HUMAN]
C9JPX0	Uncharacterized protein OS=Homo sapiens GN=SAFB2 PE=4 SV=2 - [C9JPX0_HUMAN]
F5H353	Uncharacterized protein OS=Homo sapiens GN=SCAMP2 PE=4 SV=1 - [F5H353_HUMAN]
E9PEA8	Uncharacterized protein OS=Homo sapiens GN=SCCPDH PE=4 SV=1 - [E9PEA8_HUMAN]
B7Z5N7	Uncharacterized protein OS=Homo sapiens GN=SCFD1 PE=2 SV=1 - [B7Z5N7_HUMAN]
E7EW05	Uncharacterized protein OS=Homo sapiens GN=SDAD1 PE=4 SV=1 - [E7EW05_HUMAN]
E9PBJ5	Uncharacterized protein OS=Homo sapiens GN=SDHA PE=4 SV=1 - [E9PBJ5_HUMAN]
E7EP00	Uncharacterized protein OS=Homo sapiens GN=SEC24C PE=4 SV=2 - [E7EP00_HUMAN]
B5MCX3	Uncharacterized protein OS=Homo sapiens GN=SEPT2 PE=3 SV=1 - [B5MCX3_HUMAN]
B4DNE4	Uncharacterized protein OS=Homo sapiens GN=SEPT7 PE=2 SV=1 - [B4DNE4_HUMAN]
A6NMH6	Uncharacterized protein OS=Homo sapiens GN=SEPT8 PE=3 SV=1 - [A6NMH6_HUMAN]
E5RIQ9	Uncharacterized protein OS=Homo sapiens GN=SERF1A PE=4 SV=1 - [E5RIQ9_HUMAN]
B4DN87	Uncharacterized protein OS=Homo sapiens GN=SERPINH1 PE=2 SV=1 - [B4DN87_HUMAN]
F2Z2A8	Uncharacterized protein OS=Homo sapiens GN=SGSM2 PE=4 SV=1 - [F2Z2A8_HUMAN]
E7EMJ6	Uncharacterized protein OS=Homo sapiens GN=SGTA PE=4 SV=1 - [E7EMJ6_HUMAN]
B4DQ94	Uncharacterized protein OS=Homo sapiens GN=SH3BP5L PE=2 SV=1 - [B4DQ94_HUMAN]
E5RJR5	Uncharacterized protein OS=Homo sapiens GN=SKP1 PE=4 SV=1 - [E5RJR5_HUMAN]
F5H741	Uncharacterized protein OS=Homo sapiens GN=SLC11A2 PE=4 SV=1 - [F5H741_HUMAN]
E9PR61	Uncharacterized protein OS=Homo sapiens GN=SLC14A1 PE=4 SV=1 - [E9PR61_HUMAN]
F5H843	Uncharacterized protein OS=Homo sapiens GN=SLC16A7 PE=4 SV=1 - [F5H843_HUMAN]
E9PGW3	Uncharacterized protein OS=Homo sapiens GN=SLC17A1 PE=4 SV=1 - [E9PGW3_HUMAN]
F5GX33	Uncharacterized protein OS=Homo sapiens GN=SLC25A13 PE=3 SV=1 - [F5GX33_HUMAN]
B7Z844	Uncharacterized protein OS=Homo sapiens GN=SLC2A14 PE=2 SV=1 - [B7Z844_HUMAN]
E7EMT4	Uncharacterized protein OS=Homo sapiens GN=SLC39A7 PE=4 SV=1 - [E7EMT4_HUMAN]
B4E2Z3	Uncharacterized protein OS=Homo sapiens GN=SLC3A2 PE=2 SV=1 - [B4E2Z3_HUMAN]
B7Z6X9	Uncharacterized protein OS=Homo sapiens GN=SLC43A2 PE=2 SV=1 - [B7Z6X9_HUMAN]
E9PS74	Uncharacterized protein OS=Homo sapiens GN=SLC43A3 PE=4 SV=1 - [E9PS74_HUMAN]
C9JIA1	Uncharacterized protein OS=Homo sapiens GN=SLFN12 PE=4 SV=1 - [C9JIA1_HUMAN]
D6RBH9	Uncharacterized protein OS=Homo sapiens GN=SMAD1 PE=4 SV=1 - [D6RBH9_HUMAN]
B7Z799	Uncharacterized protein OS=Homo sapiens GN=SMARCD1 PE=2 SV=1 - [B7Z799_HUMAN]

E7EPI6	Uncharacterized protein OS=Homo sapiens GN=SMC1A PE=4 SV=1 - [E7EPI6_HUMAN]
E9PD53	Uncharacterized protein OS=Homo sapiens GN=SMC4 PE=4 SV=1 - [E9PD53_HUMAN]
E7EPV7	Uncharacterized protein OS=Homo sapiens GN=SNCA PE=4 SV=1 - [E7EPV7_HUMAN]
E7ESM6	Uncharacterized protein OS=Homo sapiens GN=SND1 PE=4 SV=2 - [E7ESM6_HUMAN]
C9J1U1	Uncharacterized protein OS=Homo sapiens GN=SNRNP48 PE=4 SV=1 - [C9J1U1_HUMAN]
B4DDC7	Uncharacterized protein OS=Homo sapiens GN=SNRPA PE=2 SV=1 - [B4DDC7_HUMAN]
B4DJI7	Uncharacterized protein OS=Homo sapiens GN=SNX11 PE=2 SV=1 - [B4DJI7_HUMAN]
E5RGS8	Uncharacterized protein OS=Homo sapiens GN=SNX16 PE=4 SV=1 - [E5RGS8_HUMAN]
B4DEK4	Uncharacterized protein OS=Homo sapiens GN=SNX2 PE=2 SV=1 - [B4DEK4_HUMAN]
E9PBC7	Uncharacterized protein OS=Homo sapiens GN=SNX5 PE=4 SV=1 - [E9PBC7_HUMAN]
F5H3C5	Uncharacterized protein OS=Homo sapiens GN=SOD2 PE=3 SV=1 - [F5H3C5_HUMAN]
F5H7U6	Uncharacterized protein OS=Homo sapiens GN=SPAG9 PE=4 SV=1 - [F5H7U6_HUMAN]
A8MV89	Uncharacterized protein OS=Homo sapiens GN=SPECC1 PE=4 SV=1 - [A8MV89_HUMAN]
B4DGT1	Uncharacterized protein OS=Homo sapiens GN=SPTAN1 PE=2 SV=1 - [B4DGT1_HUMAN]
B4DHX0	Uncharacterized protein OS=Homo sapiens GN=SQLE PE=2 SV=1 - [B4DHX0_HUMAN]
F5H281	Uncharacterized protein OS=Homo sapiens GN=SRP68 PE=4 SV=1 - [F5H281_HUMAN]
B4DED5	Uncharacterized protein OS=Homo sapiens GN=SRRM1 PE=2 SV=1 - [B4DED5_HUMAN]
E7EMJ8	Uncharacterized protein OS=Homo sapiens GN=SRSF4 PE=4 SV=2 - [E7EMJ8_HUMAN]
A6NNE8	Uncharacterized protein OS=Homo sapiens GN=SRSF7 PE=4 SV=3 - [A6NNE8_HUMAN]
C9K0U8	Uncharacterized protein OS=Homo sapiens GN=SSBP1 PE=4 SV=1 - [C9K0U8_HUMAN]
A6NMU3	Uncharacterized protein OS=Homo sapiens GN=STAM PE=4 SV=1 - [A6NMU3_HUMAN]
B4E2V5	Uncharacterized protein OS=Homo sapiens GN=STOM PE=2 SV=1 - [B4E2V5_HUMAN]
B4E1K7	Uncharacterized protein OS=Homo sapiens GN=STOML2 PE=2 SV=1 - [B4E1K7_HUMAN]
B4DQH7	Uncharacterized protein OS=Homo sapiens GN=STRN4 PE=2 SV=1 - [B4DQH7_HUMAN]
D6RF48	Uncharacterized protein OS=Homo sapiens GN=STX18 PE=4 SV=1 - [D6RF48_HUMAN]
F5GXG3	Uncharacterized protein OS=Homo sapiens GN=STXBP2 PE=4 SV=1 - [F5GXG3_HUMAN]
F5GXC8	Uncharacterized protein OS=Homo sapiens GN=SUCLA2 PE=3 SV=1 - [F5GXC8_HUMAN]
F2Z3D0	Uncharacterized protein OS=Homo sapiens GN=SUMO2 PE=4 SV=1 - [F2Z3D0_HUMAN]
E7ENN3	Uncharacterized protein OS=Homo sapiens GN=SYNE1 PE=4 SV=1 - [E7ENN3_HUMAN]
C9JYN0	Uncharacterized protein OS=Homo sapiens GN=SYPL1 PE=4 SV=1 - [C9JYN0_HUMAN]
B4DIR9	Uncharacterized protein OS=Homo sapiens GN=TAB2 PE=2 SV=1 - [B4DIR9_HUMAN]
F5GWQ7	Uncharacterized protein OS=Homo sapiens GN=TAF15 PE=4 SV=1 - [F5GWQ7_HUMAN]
E7ERI3	Uncharacterized protein OS=Homo sapiens GN=TARS PE=3 SV=1 - [E7ERI3_HUMAN]
F5GXK4	Uncharacterized protein OS=Homo sapiens GN=TBC1D14 PE=4 SV=1 - [F5GXK4_HUMAN]
A8MVD5	Uncharacterized protein OS=Homo sapiens GN=TBCB PE=4 SV=1 - [A8MVD5_HUMAN]
E9PSI4	Uncharacterized protein OS=Homo sapiens GN=TBCEL PE=4 SV=1 - [E9PSI4_HUMAN]
E9PMC5	Uncharacterized protein OS=Homo sapiens GN=TCIRG1 PE=4 SV=1 - [E9PMC5_HUMAN]
C9JJP5	Uncharacterized protein OS=Homo sapiens GN=TFG PE=4 SV=1 - [C9JJP5_HUMAN]
B4DIT7	Uncharacterized protein OS=Homo sapiens GN=TGM2 PE=2 SV=1 - [B4DIT7_HUMAN]
B5MDZ2	Uncharacterized protein OS=Homo sapiens GN=THOC7 PE=4 SV=1 - [B5MDZ2_HUMAN]
E9PQZ7	Uncharacterized protein OS=Homo sapiens GN=TM7SF2 PE=4 SV=1 - [E9PQZ7_HUMAN]
F5H587	Uncharacterized protein OS=Homo sapiens GN=TM7SF3 PE=4 SV=1 - [F5H587_HUMAN]
C9JAP5	Uncharacterized protein OS=Homo sapiens GN=TMBIM1 PE=4 SV=1 - [C9JAP5_HUMAN]
A6NJCO	Uncharacterized protein OS=Homo sapiens GN=TMCC2 PE=4 SV=2 - [A6NJCO_HUMAN]
F5GX39	Uncharacterized protein OS=Homo sapiens GN=TMED2 PE=3 SV=1 - [F5GX39_HUMAN]
C9JUN6	Uncharacterized protein OS=Homo sapiens GN=TMED4 PE=3 SV=1 - [C9JUN6_HUMAN]
E9PI90	Uncharacterized protein OS=Homo sapiens GN=TMEM126A PE=4 SV=1 - [E9PI90_HUMAN]
B4DMG2	Uncharacterized protein OS=Homo sapiens GN=TMEM131 PE=2 SV=1 - [B4DMG2_HUMAN]
F2Z329	Uncharacterized protein OS=Homo sapiens GN=TMEM237 PE=4 SV=1 - [F2Z329_HUMAN]
D6RAA6	Uncharacterized protein OS=Homo sapiens GN=TMEM33 PE=4 SV=1 - [D6RAA6_HUMAN]
F5H4F7	Uncharacterized protein OS=Homo sapiens GN=TMX3 PE=4 SV=1 - [F5H4F7_HUMAN]
E9PNS3	Uncharacterized protein OS=Homo sapiens GN=TOLLIP PE=4 SV=1 - [E9PNS3_HUMAN]
E7EPD0	Uncharacterized protein OS=Homo sapiens GN=TOM1 PE=4 SV=1 - [E7EPD0_HUMAN]
F5GXH5	Uncharacterized protein OS=Homo sapiens GN=TOM1L2 PE=4 SV=1 - [F5GXH5_HUMAN]
B4DZ87	Uncharacterized protein OS=Homo sapiens GN=TOMM70A PE=2 SV=1 - [B4DZ87_HUMAN]
F5H7S3	Uncharacterized protein OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [F5H7S3_HUMAN]

B4DSE2	Uncharacterized protein OS=Homo sapiens GN=TPP1 PE=2 SV=1 - [B4DSE2_HUMAN]
E7EQD1	Uncharacterized protein OS=Homo sapiens GN=TRA2B PE=4 SV=1 - [E7EQD1_HUMAN]
B7WP88	Uncharacterized protein OS=Homo sapiens GN=TRANK1 PE=4 SV=2 - [B7WP88_HUMAN]
F5H897	Uncharacterized protein OS=Homo sapiens GN=TRAP1 PE=3 SV=1 - [F5H897_HUMAN]
F5H012	Uncharacterized protein OS=Homo sapiens GN=TRIM21 PE=4 SV=1 - [F5H012_HUMAN]
C9JNQ0	Uncharacterized protein OS=Homo sapiens GN=TRIM6 PE=4 SV=1 - [C9JNQ0_HUMAN]
C9JFG4	Uncharacterized protein OS=Homo sapiens GN=TRIOBP PE=4 SV=1 - [C9JFG4_HUMAN]
E9PFY7	Uncharacterized protein OS=Homo sapiens GN=TRMT11 PE=4 SV=1 - [E9PFY7_HUMAN]
B7Z897	Uncharacterized protein OS=Homo sapiens GN=TSC1 PE=2 SV=1 - [B7Z897_HUMAN]
E9PGT1	Uncharacterized protein OS=Homo sapiens GN=TSN PE=4 SV=1 - [E9PGT1_HUMAN]
E9PLH9	Uncharacterized protein OS=Homo sapiens GN=TSTA3 PE=4 SV=1 - [E9PLH9_HUMAN]
D6RCE2	Uncharacterized protein OS=Homo sapiens GN=TTC37 PE=4 SV=1 - [D6RCE2_HUMAN]
B5MDV0	Uncharacterized protein OS=Homo sapiens GN=TLL8 PE=4 SV=2 - [B5MDV0_HUMAN]
D6RDV0	Uncharacterized protein OS=Homo sapiens GN=TUSC3 PE=4 SV=1 - [D6RDV0_HUMAN]
B8ZZX4	Uncharacterized protein OS=Homo sapiens GN=TXNDC9 PE=4 SV=1 - [B8ZZX4_HUMAN]
E9PRI8	Uncharacterized protein OS=Homo sapiens GN=TXNRD1 PE=4 SV=1 - [E9PRI8_HUMAN]
F6WBI2	Uncharacterized protein OS=Homo sapiens GN=TXNRD2 PE=4 SV=1 - [F6WBI2_HUMAN]
C9JGI3	Uncharacterized protein OS=Homo sapiens GN=TYMP PE=4 SV=1 - [C9JGI3_HUMAN]
E7EW00	Uncharacterized protein OS=Homo sapiens GN=U2SURP PE=4 SV=1 - [E7EW00_HUMAN]
C9JD99	Uncharacterized protein OS=Homo sapiens GN=UBAP2L PE=4 SV=2 - [C9JD99_HUMAN]
F5H265	Uncharacterized protein OS=Homo sapiens GN=UBC PE=4 SV=1 - [F5H265_HUMAN]
F5H6D6	Uncharacterized protein OS=Homo sapiens GN=UBE3B PE=4 SV=1 - [F5H6D6_HUMAN]
B7Z7P0	Uncharacterized protein OS=Homo sapiens GN=UBE4A PE=2 SV=1 - [B7Z7P0_HUMAN]
F5H2G2	Uncharacterized protein OS=Homo sapiens GN=UBQLN2 PE=4 SV=1 - [F5H2G2_HUMAN]
E7EMW7	Uncharacterized protein OS=Homo sapiens GN=UBR5 PE=4 SV=1 - [E7EMW7_HUMAN]
E9PJ81	Uncharacterized protein OS=Homo sapiens GN=UBXN1 PE=4 SV=1 - [E9PJ81_HUMAN]
E9PGR0	Uncharacterized protein OS=Homo sapiens GN=UCHL5 PE=4 SV=1 - [E9PGR0_HUMAN]
C9JNP9	Uncharacterized protein OS=Homo sapiens GN=UFD1L PE=4 SV=1 - [C9JNP9_HUMAN]
C9JKD6	Uncharacterized protein OS=Homo sapiens GN=UGP2 PE=4 SV=1 - [C9JKD6_HUMAN]
F5H4C5	Uncharacterized protein OS=Homo sapiens GN=UQCC PE=4 SV=1 - [F5H4C5_HUMAN]
E5RHG9	Uncharacterized protein OS=Homo sapiens GN=UQCRB PE=4 SV=1 - [E5RHG9_HUMAN]
F5GZ76	Uncharacterized protein OS=Homo sapiens GN=USO1 PE=4 SV=1 - [F5GZ76_HUMAN]
B7Z7T5	Uncharacterized protein OS=Homo sapiens GN=USP7 PE=2 SV=1 - [B7Z7T5_HUMAN]
B0V043	Uncharacterized protein OS=Homo sapiens GN=VAR5 PE=3 SV=1 - [B0V043_HUMAN]
F5H2A7	Uncharacterized protein OS=Homo sapiens GN=VBP1 PE=4 SV=1 - [F5H2A7_HUMAN]
F5H7T3	Uncharacterized protein OS=Homo sapiens GN=VCL PE=4 SV=1 - [F5H7T3_HUMAN]
F5GZG8	Uncharacterized protein OS=Homo sapiens GN=VPS13C PE=4 SV=1 - [F5GZG8_HUMAN]
E7ETQ7	Uncharacterized protein OS=Homo sapiens GN=VTA1 PE=4 SV=2 - [E7ETQ7_HUMAN]
F5H721	Uncharacterized protein OS=Homo sapiens GN=WBP11 PE=4 SV=1 - [F5H721_HUMAN]
E7EX86	Uncharacterized protein OS=Homo sapiens GN=WBP2 PE=4 SV=1 - [E7EX86_HUMAN]
Q9H8N9	Uncharacterized protein OS=Homo sapiens GN=WDFY1 PE=2 SV=1 - [Q9H8N9_HUMAN]
B4DP38	Uncharacterized protein OS=Homo sapiens GN=WDR77 PE=2 SV=1 - [B4DP38_HUMAN]
B7Z6V3	Uncharacterized protein OS=Homo sapiens GN=WDR81 PE=2 SV=1 - [B7Z6V3_HUMAN]
E9PRI3	Uncharacterized protein OS=Homo sapiens GN=WEE1 PE=4 SV=1 - [E9PRI3_HUMAN]
F5GWT4	Uncharacterized protein OS=Homo sapiens GN=WNK1 PE=4 SV=1 - [F5GWT4_HUMAN]
C9J673	Uncharacterized protein OS=Homo sapiens GN=XPO1 PE=4 SV=1 - [C9J673_HUMAN]
E7ESC6	Uncharacterized protein OS=Homo sapiens GN=XPO7 PE=4 SV=1 - [E7ESC6_HUMAN]
B1AHC7	Uncharacterized protein OS=Homo sapiens GN=XRCC6 PE=4 SV=1 - [B1AHC7_HUMAN]
F5H3P9	Uncharacterized protein OS=Homo sapiens GN=YIPF5 PE=4 SV=1 - [F5H3P9_HUMAN]
F5GYR4	Uncharacterized protein OS=Homo sapiens GN=YME1L1 PE=3 SV=1 - [F5GYR4_HUMAN]
E9PJX1	Uncharacterized protein OS=Homo sapiens GN=ZFPL1 PE=4 SV=1 - [E9PJX1_HUMAN]
Q8N8Y4	Uncharacterized protein OS=Homo sapiens GN=ZNF7 PE=2 SV=1 - [Q8N8Y4_HUMAN]
A1A528	Uncharacterized protein OS=Homo sapiens GN=ZW10 PE=2 SV=1 - [A1A528_HUMAN]
Q6ZRU8	Uncharacterized protein OS=Homo sapiens PE=2 SV=1 - [Q6ZRU8_HUMAN]
Q6ZTQ9	Uncharacterized protein OS=Homo sapiens PE=2 SV=1 - [Q6ZTQ9_HUMAN]
C9IY82	Uncharacterized protein OS=Homo sapiens PE=3 SV=1 - [C9IY82_HUMAN]

F5H167	Uncharacterized protein OS=Homo sapiens PE=3 SV=1 - [F5H167_HUMAN]
A8MZD0	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [A8MZD0_HUMAN]
E9PGW4	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [E9PGW4_HUMAN]
F5GYQ0	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [F5GYQ0_HUMAN]
F5H3W8	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [F5H3W8_HUMAN]
F6WSP1	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [F6WSP1_HUMAN]
C9J8M8	Uncharacterized protein OS=Homo sapiens PE=4 SV=2 - [C9J8M8_HUMAN]
C9JH24	Uncharacterized protein OS=Homo sapiens PE=4 SV=2 - [C9JH24_HUMAN]
C4AM82	Uncharacterized protein OS=Homo sapiens PE=4 SV=3 - [C4AM82_HUMAN]
Q8WUR7	UPF0235 protein C15orf40 OS=Homo sapiens GN=C15orf40 PE=1 SV=1 - [C0040_HUMAN]
Q9H1C7	UPF0467 protein C5orf32 OS=Homo sapiens GN=C5orf32 PE=2 SV=1 - [CE032_HUMAN]
Q969H8	UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1 - [CS010_HUMAN]
Q9Y224	UPF0568 protein C14orf166 OS=Homo sapiens GN=C14orf166 PE=1 SV=1 - [CN166_HUMAN]
Q9HCM3	UPF0606 protein KIAA1549 OS=Homo sapiens GN=KIAA1549 PE=1 SV=4 - [K1549_HUMAN]
Q8IV36	UPF0663 transmembrane protein C17orf28 OS=Homo sapiens GN=C17orf28 PE=1 SV=1 - [CQ028_HUMAN]
Q8WUY1	UPF0670 protein C8orf55 OS=Homo sapiens GN=C8orf55 PE=1 SV=2 - [CH055_HUMAN]
Q9BSF0	UPF0733 protein C2orf88 OS=Homo sapiens GN=C2orf88 PE=2 SV=2 - [CB088_HUMAN]
Q13336	Urea transporter 1 OS=Homo sapiens GN=SLC14A1 PE=2 SV=2 - [UT1_HUMAN]
Q92738	USP6 N-terminal-like protein OS=Homo sapiens GN=USP6NL PE=1 SV=3 - [US6NL_HUMAN]
P54725	UV excision repair protein RAD23 homolog A OS=Homo sapiens GN=RAD23A PE=1 SV=1 - [RD23A_HUMAN]
P54727	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1 - [RD23B_HUMAN]
Q9H270	Vacuolar protein sorting-associated protein 11 homolog OS=Homo sapiens GN=VPS11 PE=1 SV=1 - [VPS11_HUMAN]
Q96RL7	Vacuolar protein sorting-associated protein 13A OS=Homo sapiens GN=VPS13A PE=1 SV=2 - [VP13A_HUMAN]
Q9P253	Vacuolar protein sorting-associated protein 18 homolog OS=Homo sapiens GN=VPS18 PE=1 SV=2 - [VPS18_HUMAN]
Q9UBQ0	Vacuolar protein sorting-associated protein 29 OS=Homo sapiens GN=VPS29 PE=1 SV=1 - [VPS29_HUMAN]
Q96QK1	Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VPS35 PE=1 SV=2 - [VPS35_HUMAN]
Q9BRG1	Vacuolar protein-sorting-associated protein 25 OS=Homo sapiens GN=VPS25 PE=1 SV=1 - [VPS25_HUMAN]
P50552	Vasodilator-stimulated phosphoprotein OS=Homo sapiens GN=VASP PE=1 SV=3 - [VASP_HUMAN]
P49748	Very long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=ACADVL PE=1 SV=1 - [ACADV_HUMAN]
Q12981	Vesicle transport protein SEC20 OS=Homo sapiens GN=BNIP1 PE=1 SV=3 - [SEC20_HUMAN]
Q9NZ43	Vesicle transport protein USE1 OS=Homo sapiens GN=USE1 PE=1 SV=2 - [USE1_HUMAN]
Q9UEU0	Vesicle transport through interaction with t-SNAREs homolog 1B OS=Homo sapiens GN=VTI1B PE=1 SV=3 - [VTI1B_HUMAN]
Q15836	Vesicle-associated membrane protein 3 OS=Homo sapiens GN=VAMP3 PE=1 SV=3 - [VAMP3_HUMAN]
P51809	Vesicle-associated membrane protein 7 OS=Homo sapiens GN=VAMP7 PE=1 SV=3 - [VAMP7_HUMAN]
Q9POL0	Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA PE=1 SV=3 - [VAPA_HUMAN]
Q95292	Vesicle-associated membrane protein-associated protein B/C OS=Homo sapiens GN=VAPB PE=1 SV=3 - [VAPB_HUMAN]
P46459	Vesicle-fusing ATPase OS=Homo sapiens GN=NSF PE=1 SV=3 - [NSF_HUMAN]
O75396	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=4 - [SC22B_HUMAN]
P21796	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2 - [VDAC1_HUMAN]
P45880	Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1 SV=2 - [VDAC2_HUMAN]
Q9Y277	Voltage-dependent anion-selective channel protein 3 OS=Homo sapiens GN=VDAC3 PE=1 SV=1 - [VDAC3_HUMAN]
P21281	V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3 - [VATB2_HUMAN]
P61421	V-type proton ATPase subunit d 1 OS=Homo sapiens GN=ATP6V0D1 PE=1 SV=1 - [VA0D1_HUMAN]
Q9Y5K8	V-type proton ATPase subunit D OS=Homo sapiens GN=ATP6V1D PE=1 SV=1 - [VATD_HUMAN]
Q16864	V-type proton ATPase subunit F OS=Homo sapiens GN=ATP6V1F PE=1 SV=2 - [VATF_HUMAN]
O75348	V-type proton ATPase subunit G 1 OS=Homo sapiens GN=ATP6V1G1 PE=1 SV=3 - [VATG1_HUMAN]
Q9BZH6	WD repeat-containing protein 11 OS=Homo sapiens GN=WDR11 PE=1 SV=1 - [WDR11_HUMAN]
Q6ZMY6	WD repeat-containing protein 88 OS=Homo sapiens GN=WDR88 PE=2 SV=2 - [WDR88_HUMAN]
A4D1P6	WD repeat-containing protein 91 OS=Homo sapiens GN=WDR91 PE=1 SV=2 - [WDR91_HUMAN]
Q2TAY7	WD40 repeat-containing protein SMU1 OS=Homo sapiens GN=SMU1 PE=1 SV=2 - [SMU1_HUMAN]
P13010	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]
Q86Y38	Xylosyltransferase 1 OS=Homo sapiens GN=XYLT1 PE=1 SV=1 - [XYLT1_HUMAN]
Q9P1Z0	Zinc finger and BTB domain-containing protein 4 OS=Homo sapiens GN=ZBTB4 PE=1 SV=3 - [ZBTB4_HUMAN]
Q7Z2W4	Zinc finger CCCH-type antiviral protein 1 OS=Homo sapiens GN=ZC3HAV1 PE=1 SV=3 - [ZCCHV_HUMAN]
Q86UP3	Zinc finger homeobox protein 4 OS=Homo sapiens GN=ZFHX4 PE=1 SV=1 - [ZFHX4_HUMAN]
Q96NC0	Zinc finger matrin-type protein 2 OS=Homo sapiens GN=ZMAT2 PE=1 SV=1 - [ZMAT2_HUMAN]

Q9ULJ3	Zinc finger protein 295 OS=Homo sapiens GN=ZNF295 PE=1 SV=2 - [ZN295_HUMAN]
P15822	Zinc finger protein 40 OS=Homo sapiens GN=HIVEP1 PE=1 SV=3 - [ZEP1_HUMAN]
Q9C0G0	Zinc finger protein 407 OS=Homo sapiens GN=ZNF407 PE=2 SV=2 - [ZN407_HUMAN]
Q09FC8	Zinc finger protein 415 OS=Homo sapiens GN=ZNF415 PE=2 SV=2 - [ZN415_HUMAN]
Q7Z3V5	Zinc finger protein 571 OS=Homo sapiens GN=ZNF571 PE=2 SV=3 - [ZN571_HUMAN]
Q5T619	Zinc finger protein 648 OS=Homo sapiens GN=ZNF648 PE=1 SV=1 - [ZN648_HUMAN]
Q5JPB2	Zinc finger protein 831 OS=Homo sapiens GN=ZNF831 PE=2 SV=4 - [ZN831_HUMAN]
C9JN71	Zinc finger protein 878 OS=Homo sapiens GN=ZNF878 PE=3 SV=2 - [ZN878_HUMAN]
A8MXY4	Zinc finger protein 99 OS=Homo sapiens GN=ZNF99 PE=2 SV=2 - [ZNF99_HUMAN]
Q86YF9	Zinc finger protein DZIP1 OS=Homo sapiens GN=DZIP1 PE=1 SV=1 - [DZIP1_HUMAN]
Q9Y6M5	Zinc transporter 1 OS=Homo sapiens GN=SLC30A1 PE=1 SV=3 - [ZNT1_HUMAN]
Q401N2	Zinc-activated ligand-gated ion channel OS=Homo sapiens GN=ZACN PE=1 SV=2 - [ZACN_HUMAN]

Supplemental Table 2B. Proteins detected in the cytosol fraction of Human RBCs.

Accession	Description
Q9BV57	1,2-dihydroxy-3-keto-5-methylthiopentene dioxygenase OS=Homo sapiens GN=ADI1 PE=1 SV=1 - [MTND_HUMAN]
Q9NRX4	14 kDa phosphohistidine phosphatase OS=Homo sapiens GN=PHPT1 PE=1 SV=1 - [PHP14_HUMAN]
P31946	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3 - [1433B_HUMAN]
P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1 - [1433E_HUMAN]
Q04917	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4 - [1433F_HUMAN]
<i>P61981</i>	<i>14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2 - [1433G_HUMAN]</i>
<i>P27348</i>	<i>14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1 - [1433T_HUMAN]</i>
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1 - [1433Z_HUMAN]
Q9BPX1	17-beta-hydroxysteroid dehydrogenase 14 OS=Homo sapiens GN=HSD17B14 PE=1 SV=1 - [DHB14_HUMAN]
P09543	2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Homo sapiens GN=CNP PE=1 SV=2 - [CN37_HUMAN]
P62333	26S protease regulatory subunit 10B OS=Homo sapiens GN=PSMC6 PE=1 SV=1 - [PRS10_HUMAN]
<i>P62191</i>	<i>26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1 - [PRS4_HUMAN]</i>
P17980	26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=1 SV=3 - [PRS6A_HUMAN]
P43686	26S protease regulatory subunit 6B OS=Homo sapiens GN=PSMC4 PE=1 SV=2 - [PRS6B_HUMAN]
P35998	26S protease regulatory subunit 7 OS=Homo sapiens GN=PSMC2 PE=1 SV=3 - [PRS7_HUMAN]
Q99460	26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1 PE=1 SV=2 - [PSMD1_HUMAN]
O00231	26S proteasome non-ATPase regulatory subunit 11 OS=Homo sapiens GN=PSMD11 PE=1 SV=3 - [PSD11_HUMAN]
<i>O00232</i>	<i>26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV=3 - [PSD12_HUMAN]</i>
Q9UNM6	26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV=2 - [PSD13_HUMAN]
O00487	26S proteasome non-ATPase regulatory subunit 14 OS=Homo sapiens GN=PSMD14 PE=1 SV=1 - [PSDE_HUMAN]
P55036	26S proteasome non-ATPase regulatory subunit 4 OS=Homo sapiens GN=PSMD4 PE=1 SV=1 - [PSMD4_HUMAN]
P48556	26S proteasome non-ATPase regulatory subunit 8 OS=Homo sapiens GN=PSMD8 PE=1 SV=2 - [PSMD8_HUMAN]
O00233	26S proteasome non-ATPase regulatory subunit 9 OS=Homo sapiens GN=PSMD9 PE=1 SV=3 - [PSMD9_HUMAN]
Q13442	28 kDa heat- and acid-stable phosphoprotein OS=Homo sapiens GN=PDAP1 PE=1 SV=1 - [HAP28_HUMAN]
P82664	28S ribosomal protein S10, mitochondrial OS=Homo sapiens GN=MRPS10 PE=1 SV=2 - [RT10_HUMAN]
P25325	3-mercaptopyruvate sulfurtransferase OS=Homo sapiens GN=MPST PE=1 SV=3 - [THTM_HUMAN]
Q96RE3	36 kDa muscleblind protein EXP36 OS=Homo sapiens GN=MBNL1 PE=2 SV=1 - [Q96RE3_HUMAN]
Q9BRJ2	39S ribosomal protein L45, mitochondrial OS=Homo sapiens GN=MRPL45 PE=1 SV=2 - [RM45_HUMAN]
P49189	4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3 - [AL9A1_HUMAN]
P46783	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1 - [RS10_HUMAN]
P62280	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3 - [RS11_HUMAN]
P25398	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3 - [RS12_HUMAN]
P62841	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2 - [RS15_HUMAN]
P62244	40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2 - [RS15A_HUMAN]
P62249	40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2 - [RS16_HUMAN]
P0CW22	40S ribosomal protein S17-like OS=Homo sapiens GN=RPS17L PE=3 SV=1 - [RS17L_HUMAN]
P62269	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3 - [RS18_HUMAN]
P39019	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2 - [RS19_HUMAN]
P60866	40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1 - [RS20_HUMAN]
Q8WVC2	40S ribosomal protein S21 OS=Homo sapiens GN=RPS21 PE=2 SV=1 - [Q8WVC2_HUMAN]
P62851	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1 - [RS25_HUMAN]
P62854	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3 - [RS26_HUMAN]
P62857	40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1 - [RS28_HUMAN]

P62273	40S ribosomal protein S29 OS=Homo sapiens GN=RPS29 PE=1 SV=2 - [RS29_HUMAN]
<i>P23396</i>	<i>40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2 - [RS3_HUMAN]</i>
P62701	40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2 - [RS4X_HUMAN]
P46782	40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4 - [RS5_HUMAN]
<i>P62753</i>	<i>40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1 - [RS6_HUMAN]</i>
P62081	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1 - [RS7_HUMAN]
P46781	40S ribosomal protein S9 OS=Homo sapiens GN=RPS9 PE=1 SV=3 - [RS9_HUMAN]
Q9UPN4	5-azacytidine-induced protein 1 OS=Homo sapiens GN=AZI1 PE=1 SV=3 - [AZI1_HUMAN]
Q13131	5'-AMP-activated protein kinase catalytic subunit alpha-1 OS=Homo sapiens GN=PRKAA1 PE=1 SV=4 - [AAPK1_HUMAN]
Q5TFE4	5'-nucleotidase domain-containing protein 1 OS=Homo sapiens GN=NT5DC1 PE=1 SV=1 - [NT5D1_HUMAN]
Q5JUV3	5'-nucleotidase, cytosolic II (Fragment) OS=Homo sapiens GN=NT5C2 PE=2 SV=1 - [Q5JUV3_HUMAN]
Q8TCD5	5'(3')-deoxyribonucleotidase, cytosolic type OS=Homo sapiens GN=NT5C PE=1 SV=2 - [NT5C_HUMAN]
Q01813	6-phosphofructokinase type C OS=Homo sapiens GN=PFKP PE=1 SV=2 - [K6PP_HUMAN]
P17858	6-phosphofructokinase, liver type OS=Homo sapiens GN=PFKL PE=1 SV=6 - [K6PL_HUMAN]
<i>P08237</i>	<i>6-phosphofructokinase, muscle type OS=Homo sapiens GN=PFKM PE=1 SV=2 - [K6PF_HUMAN]</i>
<i>F5H7U0</i>	<i>6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=3 SV=1 - [F5H7U0_HUMAN]</i>
O95336	6-phosphogluconolactonase OS=Homo sapiens GN=PGLS PE=1 SV=2 - [6PGL_HUMAN]
P05388	60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1 - [RLA0_HUMAN]
P05387	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1 - [RLA2_HUMAN]
P62906	60S ribosomal protein L10a OS=Homo sapiens GN=RPL10A PE=1 SV=2 - [RL10A_HUMAN]
P30050	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1 - [RL12_HUMAN]
P26373	60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4 - [RL13_HUMAN]
P40429	60S ribosomal protein L13a OS=Homo sapiens GN=RPL13A PE=1 SV=2 - [RL13A_HUMAN]
P18621	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3 - [RL17_HUMAN]
P84098	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1 - [RL19_HUMAN]
P46778	60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2 - [RL21_HUMAN]
P35268	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2 - [RL22_HUMAN]
P62829	60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1 - [RL23_HUMAN]
P62750	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1 - [RL23A_HUMAN]
P61353	60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2 - [RL27_HUMAN]
P46779	60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=1 SV=3 - [RL28_HUMAN]
P47914	60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2 - [RL29_HUMAN]
P62888	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2 - [RL30_HUMAN]
P49207	60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3 - [RL34_HUMAN]
P42766	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2 - [RL35_HUMAN]
P18077	60S ribosomal protein L35a OS=Homo sapiens GN=RPL35A PE=1 SV=2 - [RL35A_HUMAN]
Q969Q0	60S ribosomal protein L36a-like OS=Homo sapiens GN=RPL36AL PE=1 SV=3 - [RL36L_HUMAN]
Q02878	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3 - [RL6_HUMAN]
P18124	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1 - [RL7_HUMAN]
P11021	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2 - [GRP78_HUMAN]
Q02952	A-kinase anchor protein 12 OS=Homo sapiens GN=AKAP12 PE=1 SV=4 - [AKA12_HUMAN]
Q2L6I2	ABC50 protein OS=Homo sapiens GN=ABCF1 PE=2 SV=1 - [Q2L6I2_HUMAN]
Q8NG38	ABG-deleted variant of telomerase reverse transcriptase OS=Homo sapiens GN=TERT PE=2 SV=1 - [Q8NG38_HUMAN]

Q8IZT6	Abnormal spindle-like microcephaly-associated protein OS=Homo sapiens GN=ASPM PE=1 SV=2 - [ASPM_HUMAN]
Q9BWD1	Acetyl-CoA acetyltransferase, cytosolic OS=Homo sapiens GN=ACAT2 PE=1 SV=2 - [THIC_HUMAN]
Q92688	Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Homo sapiens GN=ANP32B PE=1 SV=1 - [AN32B_HUMAN]
P61160	Actin-related protein 2 OS=Homo sapiens GN=ACTR2 PE=1 SV=1 - [ARP2_HUMAN]
O15143	Actin-related protein 2/3 complex subunit 1B OS=Homo sapiens GN=ARPC1B PE=1 SV=3 - [ARC1B_HUMAN]
O15144	Actin-related protein 2/3 complex subunit 2 OS=Homo sapiens GN=ARPC2 PE=1 SV=1 - [ARPC2_HUMAN]
B1ALC0	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=3 SV=1 - [B1ALC0_HUMAN]
Q9BPX5	Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5L PE=1 SV=1 - [ARP5L_HUMAN]
P63267	Actin, gamma-enteric smooth muscle OS=Homo sapiens GN=ACTG2 PE=1 SV=1 - [ACTH_HUMAN]
P53999	Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SUB1 PE=1 SV=3 - [TCP4_HUMAN]
O95433	Activator of 90 kDa heat shock protein ATPase homolog 1 OS=Homo sapiens GN=AHSA1 PE=1 SV=1 - [AHSA1_HUMAN]
Q8NER5	Activin receptor type-1C OS=Homo sapiens GN=ACVR1C PE=1 SV=1 - [ACV1C_HUMAN]
P07108	Acyl-CoA-binding protein OS=Homo sapiens GN=DBI PE=1 SV=2 - [ACBP_HUMAN]
Q8N9L9	Acyl-coenzyme A thioesterase 4 OS=Homo sapiens GN=ACOT4 PE=1 SV=2 - [ACOT4_HUMAN]
<i>P13798</i>	<i>Acylamino-acid-releasing enzyme OS=Homo sapiens GN=APEH PE=1 SV=4 - [ACPH_HUMAN]</i>
P07311	Acylphosphatase-1 OS=Homo sapiens GN=ACYP1 PE=1 SV=2 - [ACYP1_HUMAN]
P46108	Adapter molecule crk OS=Homo sapiens GN=CRK PE=1 SV=2 - [CRK_HUMAN]
Q7Z451	Adaptor-related protein complex 2, beta 1 subunit, isoform CRA_f OS=Homo sapiens GN=AP2B1 PE=2 SV=1 - [Q7Z451_HUMAN]
Q96D30	ADD1 protein OS=Homo sapiens GN=ADD1 PE=2 SV=1 - [Q96D30_HUMAN]
P07741	Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2 - [APT_HUMAN]
F5H737	Adenosylhomocysteinase OS=Homo sapiens GN=AHCY PE=3 SV=1 - [F5H737_HUMAN]
<i>P00568</i>	<i>Adenylate kinase isoenzyme 1 OS=Homo sapiens GN=AK1 PE=1 SV=3 - [KAD1_HUMAN]</i>
P30566	Adenylosuccinate lyase OS=Homo sapiens GN=ADSL PE=1 SV=2 - [PUR8_HUMAN]
F5H6Z3	Adenylyl cyclase-associated protein OS=Homo sapiens GN=CAP1 PE=3 SV=1 - [F5H6Z3_HUMAN]
P36405	ADP-ribosylation factor-like protein 3 OS=Homo sapiens GN=ARL3 PE=1 SV=2 - [ARL3_HUMAN]
Q9UUK9	ADP-sugar pyrophosphatase OS=Homo sapiens GN=NUDT5 PE=1 SV=1 - [NUDT5_HUMAN]
P24298	Alanine aminotransferase 1 OS=Homo sapiens GN=GPT PE=1 SV=3 - [ALAT1_HUMAN]
P49588	Alanyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2 - [SYAC_HUMAN]
C9JKR2	Albumin, isoform CRA_k OS=Homo sapiens GN=ALB PE=4 SV=1 - [C9JKR2_HUMAN]
P14550	Alcohol dehydrogenase [NADP+] OS=Homo sapiens GN=AKR1A1 PE=1 SV=3 - [AK1A1_HUMAN]
P11766	Alcohol dehydrogenase class-3 OS=Homo sapiens GN=ADH5 PE=1 SV=4 - [ADHX_HUMAN]
Q06278	Aldehyde oxidase OS=Homo sapiens GN=AOX1 PE=2 SV=2 - [ADO_HUMAN]
P42330	Aldo-keto reductase family 1 member C3 OS=Homo sapiens GN=AKR1C3 PE=1 SV=4 - [AK1C3_HUMAN]
P15121	Aldose reductase OS=Homo sapiens GN=AKR1B1 PE=1 SV=3 - [ALDR_HUMAN]
Q6P4F1	Alpha-(1,3)-fucosyltransferase 10 OS=Homo sapiens GN=FUT10 PE=2 SV=2 - [FUT10_HUMAN]
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3 - [A2MG_HUMAN]
O43768	Alpha-endosulfine OS=Homo sapiens GN=ENSA PE=1 SV=1 - [ENSA_HUMAN]
<i>P06733</i>	<i>Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2 - [ENOA_HUMAN]</i>
<i>Q9NZD4</i>	<i>Alpha-hemoglobin-stabilizing protein OS=Homo sapiens GN=AHSP PE=1 SV=1 - [AHSP_HUMAN]</i>
Q9NTJ4	Alpha-mannosidase 2C1 OS=Homo sapiens GN=MAN2C1 PE=1 SV=1 - [MA2C1_HUMAN]
P54920	Alpha-soluble NSF attachment protein OS=Homo sapiens GN=NAPA PE=1 SV=3 - [SNAA_HUMAN]
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sapiens GN=AIMP1 PE=1 SV=2 - [AIMP1_HUMAN]
A2RU49	Aminoglycoside phosphotransferase domain-containing protein 1 OS=Homo sapiens GN=AGPHD1 PE=2 SV=2 - [AGPD1_HUMAN]

Q6F181	Anamorsin OS=Homo sapiens GN=CIAPIN1 PE=1 SV=2 - [CPIN1_HUMAN]
Q8N302	Angiogenic factor with G patch and FHA domains 1 OS=Homo sapiens GN=AGGF1 PE=1 SV=2 - [AGGF1_HUMAN]
Q9Y5C1	Angiopietin-related protein 3 OS=Homo sapiens GN=ANGPTL3 PE=1 SV=1 - [ANGL3_HUMAN]
Q8N7Z5	Ankyrin repeat domain-containing protein 31 OS=Homo sapiens GN=ANKRD31 PE=5 SV=2 - [ANR31_HUMAN]
A6NCL7	Ankyrin repeat domain-containing protein 33B OS=Homo sapiens GN=ANKRD33B PE=4 SV=1 - [AN33B_HUMAN]
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2 - [ANXA5_HUMAN]
O75843	AP-1 complex subunit gamma-like 2 OS=Homo sapiens GN=AP1G2 PE=1 SV=1 - [AP1G2_HUMAN]
O95782	AP-2 complex subunit alpha-1 OS=Homo sapiens GN=AP2A1 PE=1 SV=3 - [AP2A1_HUMAN]
P53680	AP-2 complex subunit sigma OS=Homo sapiens GN=AP2S1 PE=1 SV=2 - [AP2S1_HUMAN]
Q8NCW5	Apolipoprotein A-I-binding protein OS=Homo sapiens GN=APOA1BP PE=1 SV=2 - [AIBP_HUMAN]
Q1L6K4	Apoptosis-inducing factor short isoform 3 OS=Homo sapiens GN=PDCD8 PE=2 SV=1 - [Q1L6K4_HUMAN]
A6NM98	ARD1 homolog A, N-acetyltransferase (<i>S. cerevisiae</i>), isoform CRA_b OS=Homo sapiens GN=NAA10 PE=4 SV=1 - [A6NM98_HUMAN]
Q15057	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing protein 2 OS=Homo sapiens GN=ACAP2 PE=1 SV=3 - [ACAP2_HUMAN]
Q99490	Arf-GAP with GTPase, ANK repeat and PH domain-containing protein 2 OS=Homo sapiens GN=AGAP2 PE=1 SV=2 - [AGAP2_HUMAN]
Q9ULH1	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 1 OS=Homo sapiens GN=ASAP1 PE=1 SV=4 - [ASAP1_HUMAN]
P05089	Arginase-1 OS=Homo sapiens GN=ARG1 PE=1 SV=2 - [ARG1_HUMAN]
B1AKN3	Arginine-glutamic acid dipeptide (RE) repeats OS=Homo sapiens GN=RERE PE=4 SV=1 - [B1AKN3_HUMAN]
O43776	Asparaginyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=NARS PE=1 SV=1 - [SYNC_HUMAN]
E7ERW2	Aspartate aminotransferase OS=Homo sapiens GN=GOT2 PE=3 SV=1 - [E7ERW2_HUMAN]
<i>P17174</i>	<i>Aspartate aminotransferase, cytoplasmic OS=Homo sapiens GN=GOT1 PE=1 SV=3 - [AATC_HUMAN]</i>
<i>P14868</i>	<i>Aspartyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2 - [SYDC_HUMAN]</i>
Q9NP58	ATP-binding cassette sub-family B member 6, mitochondrial OS=Homo sapiens GN=ABCB6 PE=1 SV=1 - [ABCB6_HUMAN]
P53396	ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3 - [ACLY_HUMAN]
Q08211	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4 - [DHX9_HUMAN]
Q9NUU7	ATP-dependent RNA helicase DDX19A OS=Homo sapiens GN=DDX19A PE=1 SV=1 - [DD19A_HUMAN]
Q7Z478	ATP-dependent RNA helicase DHX29 OS=Homo sapiens GN=DHX29 PE=1 SV=2 - [DHX29_HUMAN]
Q96QE3	ATPase family AAA domain-containing protein 5 OS=Homo sapiens GN=ATAD5 PE=1 SV=4 - [ATAD5_HUMAN]
Q9UII2	ATPase inhibitor, mitochondrial OS=Homo sapiens GN=ATPIF1 PE=1 SV=1 - [ATIF1_HUMAN]
P16066	Atrial natriuretic peptide receptor 1 OS=Homo sapiens GN=NPR1 PE=1 SV=1 - [ANPRA_HUMAN]
Q8WXX7	Autism susceptibility gene 2 protein OS=Homo sapiens GN=AUTS2 PE=1 SV=1 - [AUTS2_HUMAN]
A4D250	B-cell acute lymphoblastic leukemia-expressed protein OS=Homo sapiens GN=BLACE PE=2 SV=1 - [BLACE_HUMAN]
Q9P281	BAH and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=BAHCC1 PE=1 SV=3 - [BAHC1_HUMAN]
P02730	Band 3 anion transport protein OS=Homo sapiens GN=SLC4A1 PE=1 SV=3 - [B3AT_HUMAN]
O43491	Band 4.1-like protein 2 OS=Homo sapiens GN=EPB41L2 PE=1 SV=1 - [E41L2_HUMAN]
Q9H329	Band 4.1-like protein 4B OS=Homo sapiens GN=EPB41L4B PE=2 SV=2 - [E41LB_HUMAN]
O75531	Barrier-to-autointegration factor OS=Homo sapiens GN=BANF1 PE=1 SV=1 - [BAF_HUMAN]
A2RRM7	BBX protein OS=Homo sapiens GN=BBX PE=2 SV=1 - [A2RRM7_HUMAN]
A2A492	BCL2/adenovirus E1B 19kD interacting protein like OS=Homo sapiens GN=BNIPL PE=4 SV=1 - [A2A492_HUMAN]
P07814	Bifunctional aminoacyl-tRNA synthetase OS=Homo sapiens GN=EPRS PE=1 SV=5 - [SYEP_HUMAN]
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cyclizing) OS=Homo sapiens GN=DAK PE=1 SV=2 - [DHAK_HUMAN]
<i>Q13057</i>	<i>Bifunctional coenzyme A synthase OS=Homo sapiens GN=COASY PE=1 SV=4 - [COASY_HUMAN]</i>
<i>P53004</i>	<i>Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2 - [BIEA_HUMAN]</i>
Q6QNY1	Biogenesis of lysosome-related organelles complex 1 subunit 2 OS=Homo sapiens GN=BLOC1S2 PE=1 SV=1 - [BL1S2_HUMAN]

P50583	Bis(5'-nucleosyl)-tetraphosphatase [asymmetrical] OS=Homo sapiens GN=NUDT2 PE=1 SV=3 - [AP4A_HUMAN]
P07738	<i>Bisphosphoglycerate mutase</i> OS=Homo sapiens GN=BPGM PE=1 SV=2 - [PMGE_HUMAN]
Q13867	Bleomycin hydrolase OS=Homo sapiens GN=BLMH PE=1 SV=1 - [BLMH_HUMAN]
Q9H3K6	BolA-like protein 2 OS=Homo sapiens GN=BOLA2 PE=1 SV=1 - [BOLA2_HUMAN]
P13497	Bone morphogenetic protein 1 OS=Homo sapiens GN=BMP1 PE=1 SV=2 - [BMP1_HUMAN]
B7Z5L0	Branched-chain-amino-acid aminotransferase OS=Homo sapiens GN=BCAT1 PE=2 SV=1 - [B7Z5L0_HUMAN]
Q9NWW8	BRISC and BRCA1-A complex member 1 OS=Homo sapiens GN=BABAM1 PE=1 SV=1 - [BABA1_HUMAN]
Q5VW32	BRO1 domain-containing protein BROX OS=Homo sapiens GN=BROX PE=1 SV=1 - [BROX_HUMAN]
Q7L273	BTB/POZ domain-containing protein KCTD9 OS=Homo sapiens GN=KCTD9 PE=2 SV=1 - [KCTD9_HUMAN]
P11586	C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3 - [C1TC_HUMAN]
O15467	C-C motif chemokine 16 OS=Homo sapiens GN=CCL16 PE=1 SV=1 - [CCL16_HUMAN]
P27708	CAD protein OS=Homo sapiens GN=CAD PE=1 SV=3 - [PYR1_HUMAN]
Q9BRF8	Calcineurin-like phosphoesterase domain-containing protein 1 OS=Homo sapiens GN=CPPED1 PE=1 SV=3 - [CPPED_HUMAN]
Q9Y6N3	Calcium-activated chloride channel regulator family member 3 OS=Homo sapiens GN=CLCA3P PE=1 SV=1 - [CLCA3_HUMAN]
Q9Y2V2	Calcium-regulated heat stable protein 1 OS=Homo sapiens GN=CARHSP1 PE=1 SV=2 - [CHSP1_HUMAN]
B2ZWH1	Calcyclin binding protein (Fragment) OS=Homo sapiens GN=CACYBP PE=4 SV=1 - [B2ZWH1_HUMAN]
Q05682	Caldesmon OS=Homo sapiens GN=CALD1 PE=1 SV=3 - [CALD1_HUMAN]
P62158	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2 - [CALM_HUMAN]
P04632	Calpain small subunit 1 OS=Homo sapiens GN=CAPNS1 PE=1 SV=1 - [CPNS1_HUMAN]
P07384	Calpain-1 catalytic subunit OS=Homo sapiens GN=CAPN1 PE=1 SV=1 - [CAN1_HUMAN]
P27797	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1 - [CALR_HUMAN]
P17612	cAMP-dependent protein kinase catalytic subunit alpha OS=Homo sapiens GN=PRKACA PE=1 SV=2 - [KAPCA_HUMAN]
Q92887	Canalicular multispecific organic anion transporter 1 OS=Homo sapiens GN=ABCC2 PE=1 SV=3 - [MRP2_HUMAN]
B1AK87	Capping protein (Actin filament) muscle Z-line, beta OS=Homo sapiens GN=CAPZB PE=4 SV=1 - [B1AK87_HUMAN]
P00915	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2 - [CAH1_HUMAN]
P00918	Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2 - [CAH2_HUMAN]
P35219	Carbonic anhydrase-related protein OS=Homo sapiens GN=CA8 PE=1 SV=3 - [CAH8_HUMAN]
P16152	Carbonyl reductase [NADPH] 1 OS=Homo sapiens GN=CBR1 PE=1 SV=3 - [CBR1_HUMAN]
Q96DG6	Carboxymethylenebutenolidase homolog OS=Homo sapiens GN=CMBL PE=1 SV=1 - [CMBL_HUMAN]
P48052	Carboxypeptidase A2 OS=Homo sapiens GN=CPA2 PE=1 SV=3 - [CBPA2_HUMAN]
A3EY21	Cardiac sodium channel alpha subunit (Fragment) OS=Homo sapiens GN=SCN5A PE=2 SV=1 - [A3EY21_HUMAN]
P19784	Casein kinase II subunit alpha' OS=Homo sapiens GN=CSNK2A2 PE=1 SV=1 - [CSK22_HUMAN]
P67870	Casein kinase II subunit beta OS=Homo sapiens GN=CSNK2B PE=1 SV=1 - [CSK2B_HUMAN]
P04040	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3 - [CATA_HUMAN]
Q05BA3	CCDC19 protein OS=Homo sapiens GN=CCDC19 PE=2 SV=1 - [Q05BA3_HUMAN]
Q9Y5K6	CD2-associated protein OS=Homo sapiens GN=CD2AP PE=1 SV=1 - [CD2AP_HUMAN]
B3KW52	cDNA FLJ42145 fis, clone TESTI4000228, highly similar to Mus musculus ubiquitin family domain containing 1 (Ubf1), mRNA OS=Homo sapiens PE=2 SV=1 - [B3KW52_HUMAN]
B3KXM6	cDNA FLJ45735 fis, clone JCMLC2000273, highly similar to Integrin alpha-M OS=Homo sapiens PE=2 SV=1 - [B3KXM6_HUMAN]
B7Z3E2	cDNA FLJ50777, highly similar to Serine/threonine-protein phosphatase 6 (EC 3.1.3.16) OS=Homo sapiens PE=2 SV=1 - [B7Z3E2_HUMAN]
B4DEW9	cDNA FLJ52478, highly similar to Eukaryotic translation initiation factor 3 subunit 5 OS=Homo sapiens PE=2 SV=1 - [B4DEW9_HUMAN]
B4DUR7	cDNA FLJ52763, highly similar to Adaptin ear-binding coat-associated protein 2 OS=Homo sapiens PE=2 SV=1 - [B4DUR7_HUMAN]
B4DHX4	cDNA FLJ52902, highly similar to Rab GDP dissociation inhibitor alpha OS=Homo sapiens PE=2 SV=1 - [B4DHX4_HUMAN]
B4DS57	cDNA FLJ53176, highly similar to Nuclear autoantigenic sperm protein OS=Homo sapiens PE=2 SV=1 - [B4DS57_HUMAN]

B4DFF1	cDNA FLJ53312, highly similar to Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens PE=2 SV=1 - [B4DFF1_HUMAN]
B4DPJ6	cDNA FLJ53635, highly similar to Homo sapiens tumor protein D52-like 2 (TPD52L2), transcript variant 6, mRNA OS=Homo sapiens PE=2 SV=1 - [B4DPJ6_HUMAN]
B4DH37	cDNA FLJ53760, highly similar to Syntaxin-7 OS=Homo sapiens PE=2 SV=1 - [B4DH37_HUMAN]
B4DW31	cDNA FLJ54186, highly similar to Nuclear receptor-binding protein OS=Homo sapiens PE=2 SV=1 - [B4DW31_HUMAN]
B7Z2C8	cDNA FLJ55681, highly similar to F-box/WD repeat protein 7 OS=Homo sapiens PE=2 SV=1 - [B7Z2C8_HUMAN]
B4E022	cDNA FLJ56274, highly similar to Transketolase (EC 2.2.1.1) OS=Homo sapiens PE=2 SV=1 - [B4E022_HUMAN]
B4DK60	cDNA FLJ57515, highly similar to CDK5 and ABL1 enzyme substrate 1 OS=Homo sapiens PE=2 SV=1 - [B4DK60_HUMAN]
B7Z337	cDNA FLJ57728, highly similar to Ras-related protein Rab-6B OS=Homo sapiens PE=2 SV=1 - [B7Z337_HUMAN]
B4DET8	cDNA FLJ57956 OS=Homo sapiens PE=2 SV=1 - [B4DET8_HUMAN]
B4DKI1	cDNA FLJ58071, highly similar to Annexin A8 OS=Homo sapiens PE=2 SV=1 - [B4DKI1_HUMAN]
B4DZV5	cDNA FLJ58820, highly similar to 55 kDa erythrocyte membrane protein (p55) OS=Homo sapiens PE=2 SV=1 - [B4DZV5_HUMAN]
B4DFA2	cDNA FLJ59316, highly similar to Vesicle-fusing ATPase (EC 3.6.4.6) OS=Homo sapiens PE=2 SV=1 - [B4DFA2_HUMAN]
B4DXR2	cDNA FLJ60945, highly similar to Eukaryotic translation initiation factor 4gamma 3 OS=Homo sapiens PE=2 SV=1 - [B4DXR2_HUMAN]
A8K1R6	cDNA FLJ77026, highly similar to Homo sapiens ataxin 2-like (ATXN2L), transcript variant E, mRNA OS=Homo sapiens PE=2 SV=1 - [A8K1R6_HUMAN]
A8K1B6	cDNA FLJ78533, highly similar to Homo sapiens dynamin 2 (DNM2), transcript variant 4, mRNA OS=Homo sapiens PE=2 SV=1 - [A8K1B6_HUMAN]
Q9H636	CDNA: FLJ22634 fis, clone HSI06565 OS=Homo sapiens GN=GRHPR PE=2 SV=1 - [Q9H636_HUMAN]
Q9ULG6	Cell cycle progression protein 1 OS=Homo sapiens GN=CCPG1 PE=1 SV=3 - [CCPG1_HUMAN]
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2 - [CDC42_HUMAN]
P49454	Centromere protein F OS=Homo sapiens GN=CENPF PE=1 SV=2 - [CENPF_HUMAN]
Q6ZU80	Centrosomal protein of 128 kDa OS=Homo sapiens GN=CEP128 PE=1 SV=2 - [CE128_HUMAN]
Q9BV73	Centrosome-associated protein CEP250 OS=Homo sapiens GN=CEP250 PE=1 SV=2 - [CP250_HUMAN]
Q3B7A2	CEP152 protein (Fragment) OS=Homo sapiens GN=CEP152 PE=2 SV=1 - [Q3B7A2_HUMAN]
B7Z4T9	Chaperonin containing TCP1, subunit 7 (Eta), isoform CRA_c OS=Homo sapiens GN=CCT7 PE=2 SV=1 - [B7Z4T9_HUMAN]
Q7LBR1	Charged multivesicular body protein 1b OS=Homo sapiens GN=CHMP1B PE=1 SV=1 - [CHM1B_HUMAN]
O43633	Charged multivesicular body protein 2a OS=Homo sapiens GN=CHMP2A PE=1 SV=1 - [CHM2A_HUMAN]
Q9H444	Charged multivesicular body protein 4b OS=Homo sapiens GN=CHMP4B PE=1 SV=1 - [CHM4B_HUMAN]
O00299	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4 - [CLIC1_HUMAN]
Q9Y259	Choline/ethanolamine kinase OS=Homo sapiens GN=CHKB PE=1 SV=3 - [CHKB_HUMAN]
Q13185	Chromobox protein homolog 3 OS=Homo sapiens GN=CBX3 PE=1 SV=4 - [CBX3_HUMAN]
P09093	Chymotrypsin-like elastase family member 3A OS=Homo sapiens GN=CELA3A PE=1 SV=3 - [CEL3A_HUMAN]
Q00610	Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5 - [CLH1_HUMAN]
P53675	Clathrin heavy chain 2 OS=Homo sapiens GN=CLTCL1 PE=1 SV=2 - [CLH2_HUMAN]
P09497	Clathrin light chain B OS=Homo sapiens GN=CLTB PE=1 SV=1 - [CLCB_HUMAN]
P53621	Coatomer subunit alpha OS=Homo sapiens GN=COPA PE=1 SV=2 - [COPA_HUMAN]
P48444	Coatomer subunit delta OS=Homo sapiens GN=ARCN1 PE=1 SV=1 - [COPD_HUMAN]
Q9UBF2	Coatomer subunit gamma-2 OS=Homo sapiens GN=COPG2 PE=1 SV=1 - [COPG2_HUMAN]
Q5T0F9	Coiled-coil and C2 domain-containing protein 1B OS=Homo sapiens GN=CC2D1B PE=1 SV=1 - [C2D1B_HUMAN]
Q96CT7	Coiled-coil domain-containing protein 124 OS=Homo sapiens GN=CCDC124 PE=1 SV=1 - [CC124_HUMAN]
Q8IYE1	Coiled-coil domain-containing protein 13 OS=Homo sapiens GN=CCDC13 PE=2 SV=2 - [CCD13_HUMAN]
Q4G0X9	Coiled-coil domain-containing protein 40 OS=Homo sapiens GN=CCDC40 PE=2 SV=2 - [CCD40_HUMAN]
Q96MW1	Coiled-coil domain-containing protein 43 OS=Homo sapiens GN=CCDC43 PE=1 SV=2 - [CCD43_HUMAN]
Q8NEL0	Coiled-coil domain-containing protein 54 OS=Homo sapiens GN=CCDC54 PE=1 SV=2 - [CCD54_HUMAN]
Q16204	Coiled-coil domain-containing protein 6 OS=Homo sapiens GN=CCDC6 PE=1 SV=2 - [CCDC6_HUMAN]

A6NI79	Coiled-coil domain-containing protein 69 OS=Homo sapiens GN=CCDC69 PE=2 SV=1 - [CCD69_HUMAN]
Q6ZRK6	Coiled-coil domain-containing protein 73 OS=Homo sapiens GN=CCDC73 PE=1 SV=2 - [CCD73_HUMAN]
Q7Z6B0	Coiled-coil domain-containing protein 91 OS=Homo sapiens GN=CCDC91 PE=1 SV=2 - [CCD91_HUMAN]
Q8N8F7	Coiled-coil domain-containing transmembrane protein C7orf53 OS=Homo sapiens GN=C7orf53 PE=2 SV=1 - [CG053_HUMAN]
Q02388	Collagen alpha-1(VII) chain OS=Homo sapiens GN=COL7A1 PE=1 SV=2 - [CO7A1_HUMAN]
P29400	Collagen alpha-5(IV) chain OS=Homo sapiens GN=COL4A5 PE=1 SV=2 - [CO4A5_HUMAN]
A6NMZ7	Collagen alpha-6(VI) chain OS=Homo sapiens GN=COL6A6 PE=1 SV=2 - [CO6A6_HUMAN]
A2A2Y8	Collagen, type XVII, alpha 1 OS=Homo sapiens GN=COL17A1 PE=4 SV=1 - [A2A2Y8_HUMAN]
Q9BXJ0	Complement C1q tumor necrosis factor-related protein 5 OS=Homo sapiens GN=C1QTNF5 PE=1 SV=1 - [C1QT5_HUMAN]
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2 - [CO9_HUMAN]
Q15021	Condensin complex subunit 1 OS=Homo sapiens GN=NCAPD2 PE=1 SV=3 - [CND1_HUMAN]
Q9BPX3	Condensin complex subunit 3 OS=Homo sapiens GN=NCAPG PE=1 SV=1 - [CND3_HUMAN]
A8K1H6	COP9 constitutive photomorphogenic homolog subunit 8 (Arabidopsis), isoform CRA_b OS=Homo sapiens GN=COPS8 PE=2 SV=1 - [A8K1H6_HUMAN]
Q13098	COP9 signalosome complex subunit 1 OS=Homo sapiens GN=GPS1 PE=1 SV=4 - [CSN1_HUMAN]
Q9BT78	COP9 signalosome complex subunit 4 OS=Homo sapiens GN=COPS4 PE=1 SV=1 - [CSN4_HUMAN]
Q92905	COP9 signalosome complex subunit 5 OS=Homo sapiens GN=COPS5 PE=1 SV=4 - [CSN5_HUMAN]
Q9UBW8	COP9 signalosome complex subunit 7a OS=Homo sapiens GN=COPS7A PE=1 SV=1 - [CSN7A_HUMAN]
Q9H9Q2	COP9 signalosome complex subunit 7b OS=Homo sapiens GN=COPS7B PE=1 SV=1 - [CSN7B_HUMAN]
Q5JX54	Copine I (Fragment) OS=Homo sapiens GN=CPNE1 PE=2 SV=1 - [Q5JX54_HUMAN]
O75131	Copine-3 OS=Homo sapiens GN=CPNE3 PE=1 SV=1 - [CPNE3_HUMAN]
O14618	Copper chaperone for superoxide dismutase OS=Homo sapiens GN=CCS PE=1 SV=1 - [CCS_HUMAN]
P36551	Coproporphyrinogen-III oxidase, mitochondrial OS=Homo sapiens GN=CPOX PE=1 SV=3 - [HEM6_HUMAN]
B5MDT7	Core-binding factor, runt domain, alpha subunit 2; translocated to, 3, isoform CRA_a OS=Homo sapiens GN=CBFA2T3 PE=4 SV=1 - [B5MDT7_HUMAN]
Q9ULV4	Coronin-1C OS=Homo sapiens GN=CORO1C PE=1 SV=1 - [COR1C_HUMAN]
P57737	Coronin-7 OS=Homo sapiens GN=CORO7 PE=1 SV=2 - [CORO7_HUMAN]
Q96H99	Cortactin OS=Homo sapiens GN=CTTN PE=2 SV=1 - [Q96H99_HUMAN]
Q9UEE9	Craniofacial development protein 1 OS=Homo sapiens GN=CFDP1 PE=1 SV=1 - [CFDP1_HUMAN]
Q49AN0	Cryptochrome-2 OS=Homo sapiens GN=CRY2 PE=1 SV=2 - [CRY2_HUMAN]
P17812	CTP synthase 1 OS=Homo sapiens GN=CTPS PE=1 SV=2 - [PYRG1_HUMAN]
Q9NRF8	CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1 - [PYRG2_HUMAN]
B1AMN7	CTTNBP2 N-terminal like (Fragment) OS=Homo sapiens GN=CTTNBP2NL PE=4 SV=1 - [B1AMN7_HUMAN]
Q13617	Cullin-2 OS=Homo sapiens GN=CUL2 PE=1 SV=2 - [CUL2_HUMAN]
Q13618	Cullin-3 OS=Homo sapiens GN=CUL3 PE=1 SV=2 - [CUL3_HUMAN]
Q86VP6	Cullin-associated NEDD8-dissociated protein 1 OS=Homo sapiens GN=CAND1 PE=1 SV=2 - [CAND1_HUMAN]
O95273	Cyclin-D1-binding protein 1 OS=Homo sapiens GN=CCNDBP1 PE=1 SV=2 - [CCDB1_HUMAN]
P04080	Cystatin-B OS=Homo sapiens GN=CSTB PE=1 SV=2 - [CYTB_HUMAN]
Q9UHD1	Cysteine and histidine-rich domain-containing protein 1 OS=Homo sapiens GN=CHORDC1 PE=1 SV=2 - [CHRD1_HUMAN]
Q9H8E8	Cysteine-rich protein 2-binding protein OS=Homo sapiens GN=CSRP2BP PE=1 SV=3 - [CSR2B_HUMAN]
A8MVQ3	Cysteinyl-tRNA synthetase, isoform CRA_b OS=Homo sapiens GN=CARS PE=2 SV=1 - [A8MVQ3_HUMAN]
P00167	Cytochrome b5 OS=Homo sapiens GN=CYB5A PE=1 SV=2 - [CYB5_HUMAN]
Q8IUI8	Cytokine receptor-like factor 3 OS=Homo sapiens GN=CRLF3 PE=1 SV=2 - [CRLF3_HUMAN]
Q8NCM8	Cytoplasmic dynein 2 heavy chain 1 OS=Homo sapiens GN=DYNC2H1 PE=1 SV=4 - [DYHC2_HUMAN]
P28838	Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3 - [AMPL_HUMAN]

P53384	Cytosolic Fe-S cluster assembly factor NUBP1 OS=Homo sapiens GN=NUBP1 PE=1 SV=2 - [NUBP1_HUMAN]
Q96KP4	Cytosolic non-specific dipeptidase OS=Homo sapiens GN=CNDP2 PE=1 SV=2 - [CNDP2_HUMAN]
P0C869	Cytosolic phospholipase A2 beta OS=Homo sapiens GN=PLA2G4B PE=1 SV=2 - [PA24B_HUMAN]
B7Z522	D-dopachrome tautomerase, isoform CRA_a OS=Homo sapiens GN=DDT PE=4 SV=1 - [B7Z522_HUMAN]
Q9NYF0	Dapper homolog 1 OS=Homo sapiens GN=DACT1 PE=2 SV=2 - [DACT1_HUMAN]
Q8TEB1	DDB1- and CUL4-associated factor 11 OS=Homo sapiens GN=DCAF11 PE=1 SV=1 - [DCA11_HUMAN]
Q3SXM0	DDB1- and CUL4-associated factor 4-like protein 1 OS=Homo sapiens GN=DCAF4L1 PE=2 SV=1 - [DC4L1_HUMAN]
P51397	Death-associated protein 1 OS=Homo sapiens GN=DAP PE=1 SV=3 - [DAP1_HUMAN]
A0PJW8	Death-associated protein-like 1 OS=Homo sapiens GN=DAPL1 PE=1 SV=2 - [DAPL1_HUMAN]
Q9BTC0	Death-inducer obliterator 1 OS=Homo sapiens GN=DIDO1 PE=1 SV=5 - [DIDO1_HUMAN]
Q5JSL3	Dedicator of cytokinesis protein 11 OS=Homo sapiens GN=DOCK11 PE=1 SV=2 - [DOC11_HUMAN]
Q8IZD9	Dedicator of cytokinesis protein 3 OS=Homo sapiens GN=DOCK3 PE=1 SV=1 - [DOCK3_HUMAN]
Q6UWP2	Dehydrogenase/reductase SDR family member 11 OS=Homo sapiens GN=DHRS11 PE=1 SV=1 - [DHR11_HUMAN]
Q9Y238	Deleted in lung and esophageal cancer protein 1 OS=Homo sapiens GN=DLEC1 PE=2 SV=2 - [DLEC1_HUMAN]
B7Z3I9	Delta-aminolevulinic acid dehydratase OS=Homo sapiens PE=2 SV=1 - [B7Z3I9_HUMAN]
Q6ZUT9	DENN domain-containing protein 5B OS=Homo sapiens GN=DENND5B PE=1 SV=2 - [DEN5B_HUMAN]
O43598	Deoxyribonucleoside 5'-monophosphate N-glycosidase OS=Homo sapiens GN=RCL PE=1 SV=1 - [RCL_HUMAN]
P33316	Deoxyuridine 5'-triphosphate nucleotidohydrolase, mitochondrial OS=Homo sapiens GN=DUT PE=1 SV=3 - [DUT_HUMAN]
Q9UL01	Dermatan-sulfate epimerase OS=Homo sapiens GN=DSE PE=1 SV=1 - [DSE_HUMAN]
P81605	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2 - [DCD_HUMAN]
B4DYA6	Destrin (Actin depolymerizing factor), isoform CRA_a OS=Homo sapiens GN=DSTN PE=2 SV=1 - [B4DYA6_HUMAN]
Q86TI2	Dipeptidyl peptidase 9 OS=Homo sapiens GN=DPP9 PE=1 SV=3 - [DPP9_HUMAN]
Q8NFP7	Diphosphoinositol polyphosphate phosphohydrolase 3-alpha OS=Homo sapiens GN=NUDT10 PE=1 SV=1 - [NUD10_HUMAN]
P53602	Diphosphomevalonate decarboxylase OS=Homo sapiens GN=MVD PE=1 SV=1 - [MVD1_HUMAN]
Q16531	DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1 - [DDB1_HUMAN]
O00273	DNA fragmentation factor subunit alpha OS=Homo sapiens GN=DFFA PE=1 SV=1 - [DFFA_HUMAN]
Q92878	DNA repair protein RAD50 OS=Homo sapiens GN=RAD50 PE=1 SV=1 - [RAD50_HUMAN]
P16989	DNA-binding protein A OS=Homo sapiens GN=CSDA PE=1 SV=4 - [DBPA_HUMAN]
O60884	DnaJ homolog subfamily A member 2 OS=Homo sapiens GN=DNAJA2 PE=1 SV=1 - [DNJA2_HUMAN]
Q8WW22	DnaJ homolog subfamily A member 4 OS=Homo sapiens GN=DNAJA4 PE=1 SV=1 - [DNJA4_HUMAN]
P25685	DnaJ homolog subfamily B member 1 OS=Homo sapiens GN=DNAJB1 PE=1 SV=4 - [DNJB1_HUMAN]
Q8WWF6	DnaJ homolog subfamily B member 3 OS=Homo sapiens GN=DNAJB3 PE=1 SV=1 - [DNJB3_HUMAN]
Q96KC8	DnaJ homolog subfamily C member 1 OS=Homo sapiens GN=DNAJC1 PE=1 SV=1 - [DNJC1_HUMAN]
O75937	DnaJ homolog subfamily C member 8 OS=Homo sapiens GN=DNAJC8 PE=1 SV=2 - [DNJC8_HUMAN]
Q8WXX5	DnaJ homolog subfamily C member 9 OS=Homo sapiens GN=DNAJC9 PE=1 SV=1 - [DNJC9_HUMAN]
P49959	Double-strand break repair protein MRE11A OS=Homo sapiens GN=MRE11A PE=1 SV=3 - [MRE11_HUMAN]
O60216	Double-strand-break repair protein rad21 homolog OS=Homo sapiens GN=RAD21 PE=1 SV=2 - [RAD21_HUMAN]
Q9UJU6	Drebrin-like protein OS=Homo sapiens GN=DBNL PE=1 SV=1 - [DBNL_HUMAN]
P36507	Dual specificity mitogen-activated protein kinase kinase 2 OS=Homo sapiens GN=MAP2K2 PE=1 SV=1 - [MP2K2_HUMAN]
P46734	Dual specificity mitogen-activated protein kinase kinase 3 OS=Homo sapiens GN=MAP2K3 PE=1 SV=2 - [MP2K3_HUMAN]
Q4G0W2	Dual specificity phosphatase 28 OS=Homo sapiens GN=DUSP28 PE=2 SV=1 - [DUS28_HUMAN]
Q9NR20	Dual specificity tyrosine-phosphorylation-regulated kinase 4 OS=Homo sapiens GN=DYRK4 PE=1 SV=2 - [DYRK4_HUMAN]
Q8IVF4	Dynein heavy chain 10, axonemal OS=Homo sapiens GN=DNAH10 PE=1 SV=4 - [DYH10_HUMAN]

P63167	Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1 - [DYL1_HUMAN]
Q96FJ2	Dynein light chain 2, cytoplasmic OS=Homo sapiens GN=DYNLL2 PE=1 SV=1 - [DYL2_HUMAN]
Q8WXU2	Dyslexia susceptibility 1 candidate gene 1 protein OS=Homo sapiens GN=DYX1C1 PE=1 SV=2 - [DYXC1_HUMAN]
Q14118	Dystroglycan OS=Homo sapiens GN=DAG1 PE=1 SV=2 - [DAG1_HUMAN]
O95376	E3 ubiquitin-protein ligase ARIH2 OS=Homo sapiens GN=ARIH2 PE=1 SV=1 - [ARI2_HUMAN]
Q9P0J7	E3 ubiquitin-protein ligase KCMF1 OS=Homo sapiens GN=KCMF1 PE=1 SV=2 - [KCMF1_HUMAN]
Q5XPI4	E3 ubiquitin-protein ligase RNF123 OS=Homo sapiens GN=RNF123 PE=1 SV=1 - [RN123_HUMAN]
Q9HAU4	E3 ubiquitin-protein ligase SMURF2 OS=Homo sapiens GN=SMURF2 PE=1 SV=1 - [SMUF2_HUMAN]
Q96F44	E3 ubiquitin-protein ligase TRIM11 OS=Homo sapiens GN=TRIM11 PE=1 SV=2 - [TRI11_HUMAN]
Q5T4S7	E3 ubiquitin-protein ligase UBR4 OS=Homo sapiens GN=UBR4 PE=1 SV=1 - [UBR4_HUMAN]
Q4VBZ6	EEF1D protein OS=Homo sapiens GN=EEF1D PE=1 SV=1 - [Q4VBZ6_HUMAN]
Q9BSW2	EF-hand calcium-binding domain-containing protein 4B OS=Homo sapiens GN=EFCAB4B PE=1 SV=1 - [EFC4B_HUMAN]
Q2Y0W8	Electroneutral sodium bicarbonate exchanger 1 OS=Homo sapiens GN=SLC4A8 PE=1 SV=1 - [S4A8_HUMAN]
P68104	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 - [EF1A1_HUMAN]
P24534	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3 - [EF1B_HUMAN]
P26641	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3 - [EF1G_HUMAN]
P13639	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4 - [EF2_HUMAN]
Q7L9B9	Endonuclease/exonuclease/phosphatase family domain-containing protein 1 OS=Homo sapiens GN=EEPDP1 PE=1 SV=2 - [EEPDP1_HUMAN]
P14625	Endoplasmic OS=Homo sapiens GN=HSP90B1 PE=1 SV=1 - [ENPL_HUMAN]
P24530	Endothelin B receptor OS=Homo sapiens GN=EDNRB PE=1 SV=1 - [EDNRB_HUMAN]
Q96F86	Enhancer of mRNA-decapping protein 3 OS=Homo sapiens GN=EDC3 PE=1 SV=1 - [EDC3_HUMAN]
P84090	Enhancer of rudimentary homolog OS=Homo sapiens GN=ERH PE=1 SV=1 - [ERH_HUMAN]
B7Z2X9	Enolase OS=Homo sapiens GN=ENO2 PE=2 SV=1 - [B7Z2X9_HUMAN]
P98073	Enteropeptidase OS=Homo sapiens GN=TMPRSS15 PE=2 SV=3 - [ENTK_HUMAN]
A7E2D7	EP400 protein (Fragment) OS=Homo sapiens GN=EP400 PE=2 SV=1 - [A7E2D7_HUMAN]
Q96CG1	ETF1 protein OS=Homo sapiens GN=ETF1 PE=2 SV=1 - [Q96CG1_HUMAN]
Q99447	Ethanolamine-phosphate cytidyltransferase OS=Homo sapiens GN=PCYT2 PE=1 SV=1 - [PCY2_HUMAN]
P60842	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1 - [IF4A1_HUMAN]
P38919	Eukaryotic initiation factor 4A-III OS=Homo sapiens GN=EIF4A3 PE=1 SV=4 - [IF4A3_HUMAN]
O14602	Eukaryotic translation initiation factor 1A, Y-chromosomal OS=Homo sapiens GN=EIF1AY PE=1 SV=4 - [IF1AY_HUMAN]
P05198	Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens GN=EIF2S1 PE=1 SV=3 - [IF2A_HUMAN]
P20042	Eukaryotic translation initiation factor 2 subunit 2 OS=Homo sapiens GN=EIF2S2 PE=1 SV=2 - [IF2B_HUMAN]
P41091	Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=3 - [IF2G_HUMAN]
Q9BQI3	Eukaryotic translation initiation factor 2-alpha kinase 1 OS=Homo sapiens GN=EIF2AK1 PE=1 SV=2 - [E2AK1_HUMAN]
O75821	Eukaryotic translation initiation factor 3 subunit G OS=Homo sapiens GN=EIF3G PE=1 SV=2 - [EIF3G_HUMAN]
Q13347	Eukaryotic translation initiation factor 3 subunit I OS=Homo sapiens GN=EIF3I PE=1 SV=1 - [EIF3I_HUMAN]
O75822	Eukaryotic translation initiation factor 3 subunit J OS=Homo sapiens GN=EIF3J PE=1 SV=2 - [EIF3J_HUMAN]
B0QYA6	Eukaryotic translation initiation factor 3, subunit D (Fragment) OS=Homo sapiens GN=EIF3D PE=4 SV=1 - [B0QYA6_HUMAN]
E2QC29	Eukaryotic translation initiation factor 3, subunit E interacting protein (Fragment) OS=Homo sapiens GN=EIF3EIP PE=4 SV=1 - [E2QC29_HUMAN]
P06730	Eukaryotic translation initiation factor 4E OS=Homo sapiens GN=EIF4E PE=1 SV=2 - [IF4E_HUMAN]
Q13541	Eukaryotic translation initiation factor 4E-binding protein 1 OS=Homo sapiens GN=EIF4EBP1 PE=1 SV=3 - [4EBP1_HUMAN]
Q15056	Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H PE=1 SV=5 - [IF4H_HUMAN]
P55010	Eukaryotic translation initiation factor 5 OS=Homo sapiens GN=EIF5 PE=1 SV=2 - [IF5_HUMAN]

P63241	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2 - [IF5A1_HUMAN]
O60841	Eukaryotic translation initiation factor 5B OS=Homo sapiens GN=EIF5B PE=1 SV=4 - [IF2P_HUMAN]
B7ZBH1	Eukaryotic translation initiation factor 6 (Fragment) OS=Homo sapiens GN=EIF6 PE=4 SV=1 - [B7ZBH1_HUMAN]
O00471	Exocyst complex component 5 OS=Homo sapiens GN=EXOC5 PE=1 SV=1 - [EXOC5_HUMAN]
O14980	Exportin-1 OS=Homo sapiens GN=XPO1 PE=1 SV=1 - [XPO1_HUMAN]
A8CZ64	Extracellular signal-regulated kinase-2 splice variant OS=Homo sapiens GN=MAPK1 PE=2 SV=1 - [A8CZ64_HUMAN]
P52907	F-actin-capping protein subunit alpha-1 OS=Homo sapiens GN=CAPZA1 PE=1 SV=3 - [CAZA1_HUMAN]
Q5TI86	F-box protein 7 OS=Homo sapiens GN=FBXO7 PE=2 SV=1 - [Q5TI86_HUMAN]
B1AJQ5	Family with sequence similarity 76, member A (Fragment) OS=Homo sapiens GN=FAM76A PE=4 SV=1 - [B1AJQ5_HUMAN]
Q9Y2M0	Fanconi-associated nuclease 1 OS=Homo sapiens GN=FAN1 PE=1 SV=4 - [FAN1_HUMAN]
Q16658	Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3 - [FSCN1_HUMAN]
P49327	Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3 - [FAS_HUMAN]
Q01469	Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=1 SV=3 - [FABP5_HUMAN]
Q5SYB0	FERM and PDZ domain-containing protein 1 OS=Homo sapiens GN=FRMPD1 PE=1 SV=1 - [FRPD1_HUMAN]
P02792	Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2 - [FRIL_HUMAN]
P11487	Fibroblast growth factor 3 OS=Homo sapiens GN=FGF3 PE=1 SV=1 - [FGF3_HUMAN]
A6NDY9	Filamin A OS=Homo sapiens GN=FLNA PE=2 SV=4 - [A6NDY9_HUMAN]
Q1JUQ5	FK506 binding protein12 OS=Homo sapiens GN=FKBP12-Exp2 PE=2 SV=1 - [Q1JUQ5_HUMAN]
P30043	Flavin reductase OS=Homo sapiens GN=BLVRB PE=1 SV=3 - [BLVRB_HUMAN]
P04075	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2 - [ALDOA_HUMAN]
P09972	Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2 - [ALDOC_HUMAN]
B7ZM02	FSD2 protein OS=Homo sapiens GN=FSD2 PE=2 SV=1 - [B7ZM02_HUMAN]
Q9H9S5	Fukutin-related protein OS=Homo sapiens GN=FKRP PE=1 SV=1 - [FKRP_HUMAN]
P07954	Fumarate hydratase, mitochondrial OS=Homo sapiens GN=FH PE=1 SV=3 - [FUMH_HUMAN]
Q96GK7	Fumarylacetoacetate hydrolase domain-containing protein 2A OS=Homo sapiens GN=FAHD2A PE=1 SV=1 - [FAH2A_HUMAN]
E7ET32	Galactose-1-phosphate uridylyltransferase OS=Homo sapiens GN=GALT PE=3 SV=1 - [E7ET32_HUMAN]
P09382	Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2 - [LEG1_HUMAN]
P17931	Galectin-3 OS=Homo sapiens GN=LGALS3 PE=1 SV=5 - [LEG3_HUMAN]
P60520	Gamma-aminobutyric acid receptor-associated protein-like 2 OS=Homo sapiens GN=GABARAPL2 PE=1 SV=1 - [GBRL2_HUMAN]
Q13630	GDP-L-fucose synthase OS=Homo sapiens GN=TSTA3 PE=1 SV=1 - [FCL_HUMAN]
P53609	Geranylgeranyl transferase type-1 subunit beta OS=Homo sapiens GN=PGGT1B PE=1 SV=2 - [PGTB1_HUMAN]
Q92696	Geranylgeranyl transferase type-2 subunit alpha OS=Homo sapiens GN=RABGGTA PE=1 SV=2 - [PGTA_HUMAN]
Q6PCE3	Glucose 1,6-bisphosphate synthase OS=Homo sapiens GN=PGM2L1 PE=1 SV=3 - [PGM2L_HUMAN]
P11413	Glucose-6-phosphate 1-dehydrogenase OS=Homo sapiens GN=G6PD PE=1 SV=4 - [G6PD_HUMAN]
P06744	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4 - [G6PI_HUMAN]
P48058	Glutamate receptor 4 OS=Homo sapiens GN=GRIA4 PE=2 SV=2 - [GRIA4_HUMAN]
P48506	Glutamate--cysteine ligase catalytic subunit OS=Homo sapiens GN=GCLC PE=1 SV=2 - [GSH1_HUMAN]
P48507	Glutamate--cysteine ligase regulatory subunit OS=Homo sapiens GN=GCLM PE=1 SV=1 - [GSH0_HUMAN]
P15104	Glutamine synthetase OS=Homo sapiens GN=GLUL PE=1 SV=4 - [GLNA_HUMAN]
P35754	Glutaredoxin-1 OS=Homo sapiens GN=GLRX PE=1 SV=2 - [GLRX1_HUMAN]
O76003	Glutaredoxin-3 OS=Homo sapiens GN=GLRX3 PE=1 SV=2 - [GLRX3_HUMAN]
C8KIL8	Glutathion reductase delta8 alternative splicing variant OS=Homo sapiens GN=GSR PE=2 SV=1 - [C8KIL8_HUMAN]
P07203	Glutathione peroxidase 1 OS=Homo sapiens GN=GPX1 PE=1 SV=4 - [GPX1_HUMAN]

P21266	Glutathione S-transferase Mu 3 OS=Homo sapiens GN=GSTM3 PE=1 SV=3 - [GSTM3_HUMAN]
P78417	Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2 - [GSTO1_HUMAN]
P48637	Glutathione synthetase OS=Homo sapiens GN=GSS PE=1 SV=1 - [GSHB_HUMAN]
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3 - [G3P_HUMAN]
P35573	Glycogen debranching enzyme OS=Homo sapiens GN=AGL PE=1 SV=3 - [GDE_HUMAN]
P04921	Glycophorin-C OS=Homo sapiens GN=GYPC PE=1 SV=1 - [GLPC_HUMAN]
P41250	Glycyl-tRNA synthetase OS=Homo sapiens GN=GARS PE=1 SV=3 - [SYG_HUMAN]
Q9UBQ7	Glyoxylate reductase/hydroxypyruvate reductase OS=Homo sapiens GN=GRHPR PE=1 SV=1 - [GRHPR_HUMAN]
P36959	GMP reductase 1 OS=Homo sapiens GN=GMPR PE=1 SV=1 - [GMPR1_HUMAN]
Q86T14	GMP reductase OS=Homo sapiens GN=GMPR2 PE=2 SV=1 - [Q86T14_HUMAN]
Q08378	Golgin subfamily A member 3 OS=Homo sapiens GN=GOLGA3 PE=1 SV=2 - [GOGA3_HUMAN]
Q14789	Golgin subfamily B member 1 OS=Homo sapiens GN=GOLGB1 PE=1 SV=2 - [GOGB1_HUMAN]
B3KUR5	GRAM domain containing 1C, isoform CRA_d OS=Homo sapiens GN=GRAMD1C PE=2 SV=1 - [B3KUR5_HUMAN]
Q5VT94	Growth hormone inducible transmembrane protein OS=Homo sapiens GN=GHITM PE=2 SV=1 - [Q5VT94_HUMAN]
P62826	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3 - [RAN_HUMAN]
O00178	GTP-binding protein 1 OS=Homo sapiens GN=GTPBP1 PE=1 SV=3 - [GTPB1_HUMAN]
O75628	GTP-binding protein REM 1 OS=Homo sapiens GN=REM1 PE=1 SV=2 - [REM1_HUMAN]
Q96F15	GTPase IMAP family member 5 OS=Homo sapiens GN=GIMAP5 PE=2 SV=1 - [GIMA5_HUMAN]
Q8ND71	GTPase IMAP family member 8 OS=Homo sapiens GN=GIMAP8 PE=1 SV=2 - [GIMA8_HUMAN]
Q14C86	GTPase-activating protein and VPS9 domain-containing protein 1 OS=Homo sapiens GN=GAPVD1 PE=1 SV=2 - [GAPD1_HUMAN]
Q8TBN0	Guanine nucleotide exchange factor for Rab-3A OS=Homo sapiens GN=RAB3IL1 PE=1 SV=1 - [R3GEF_HUMAN]
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L1 PE=1 SV=3 - [GBLP_HUMAN]
Q16774	Guanylate kinase OS=Homo sapiens GN=GUK1 PE=1 SV=2 - [KGUA_HUMAN]
Q96PP9	Guanylate-binding protein 4 OS=Homo sapiens GN=GBP4 PE=1 SV=2 - [GBP4_HUMAN]
Q9H0R4	Haloacid dehalogenase-like hydrolase domain-containing protein 2 OS=Homo sapiens GN=HDHD2 PE=1 SV=1 - [HDHD2_HUMAN]
Q9Y450	HBS1-like protein OS=Homo sapiens GN=HBS1L PE=1 SV=1 - [HBS1L_HUMAN]
D3DTT2	HCG1820468, isoform CRA_a OS=Homo sapiens GN=FAM18B2 PE=4 SV=1 - [D3DTT2_HUMAN]
Q2YDC2	HCG1985580, isoform CRA_f OS=Homo sapiens GN=PDCC6 PE=2 SV=1 - [Q2YDC2_HUMAN]
B4DHA4	HCG2031650, isoform CRA_f OS=Homo sapiens GN=NSUN4 PE=2 SV=1 - [B4DHA4_HUMAN]
Q7Z4H3	HD domain-containing protein 2 OS=Homo sapiens GN=HDDC2 PE=1 SV=1 - [HDDC2_HUMAN]
Q5T3Q7	HEAT repeat containing 1 OS=Homo sapiens GN=HEATR1 PE=2 SV=1 - [Q5T3Q7_HUMAN]
B8ZZL8	Heat shock 10kDa protein 1 (Chaperonin 10), isoform CRA_b OS=Homo sapiens GN=HSPE1 PE=3 SV=1 - [B8ZZL8_HUMAN]
P34931	Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2 - [HS71L_HUMAN]
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5 - [HSP71_HUMAN]
P34932	Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4 - [HSP74_HUMAN]
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1 - [HSP7C_HUMAN]
Q92598	Heat shock protein 105 kDa OS=Homo sapiens GN=HSPH1 PE=1 SV=1 - [HS105_HUMAN]
P07900	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5 - [HS90A_HUMAN]
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [HS90B_HUMAN]
Q9NRV9	Heme-binding protein 1 OS=Homo sapiens GN=HEBP1 PE=1 SV=1 - [HEBP1_HUMAN]
A8K466	Hemochromatosis type 2 (Juvenile) OS=Homo sapiens GN=HFE2 PE=2 SV=1 - [A8K466_HUMAN]
Q9BXL5	Hemogen OS=Homo sapiens GN=HEMGN PE=1 SV=1 - [HEMGN_HUMAN]
P69905	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2 - [HBA_HUMAN]

P68871	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2 - [HBB_HUMAN]
P02042	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2 - [HBD_HUMAN]
P02100	Hemoglobin subunit epsilon OS=Homo sapiens GN=HBE1 PE=1 SV=2 - [HBE_HUMAN]
P69891	Hemoglobin subunit gamma-1 OS=Homo sapiens GN=HBG1 PE=1 SV=2 - [HBG1_HUMAN]
P69892	Hemoglobin subunit gamma-2 OS=Homo sapiens GN=HBG2 PE=1 SV=2 - [HBG2_HUMAN]
Q6B0K9	Hemoglobin subunit mu OS=Homo sapiens GN=HBM PE=2 SV=1 - [HBM_HUMAN]
P09105	Hemoglobin subunit theta-1 OS=Homo sapiens GN=HBQ1 PE=1 SV=2 - [HBAT_HUMAN]
P02008	Hemoglobin subunit zeta OS=Homo sapiens GN=HBZ PE=1 SV=2 - [HBAZ_HUMAN]
D9YZU8	Hemoglobin, gamma A OS=Homo sapiens GN=HBG1 PE=3 SV=1 - [D9YZU8_HUMAN]
P51858	Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1 - [HDGF_HUMAN]
B0QZK4	Heterochromatin protein 1, binding protein 3 (Fragment) OS=Homo sapiens GN=HP1BP3 PE=3 SV=1 - [B0QZK4_HUMAN]
P09651	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5 - [ROA1_HUMAN]
O60812	Heterogeneous nuclear ribonucleoprotein C-like 1 OS=Homo sapiens GN=HNRNPCL1 PE=1 SV=1 - [HNRCL_HUMAN]
O14979	Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRPDL PE=1 SV=3 - [HNRDL_HUMAN]
P31942	Heterogeneous nuclear ribonucleoprotein H3 OS=Homo sapiens GN=HNRNPH3 PE=1 SV=2 - [HNRH3_HUMAN]
P52272	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3 - [HNRPM_HUMAN]
O60506	Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2 - [HNRPQ_HUMAN]
P22626	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2 - [ROA2_HUMAN]
P26583	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2 - [HMGB2_HUMAN]
Q5T7C0	High-mobility group box 1 (Fragment) OS=Homo sapiens GN=HMGB1 PE=2 SV=1 - [Q5T7C0_HUMAN]
P49773	Histidine triad nucleotide-binding protein 1 OS=Homo sapiens GN=HINT1 PE=1 SV=2 - [HINT1_HUMAN]
P16403	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]
P10412	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2 - [H14_HUMAN]
P16401	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3 - [H15_HUMAN]
Q96KK5	Histone H2A type 1-H OS=Homo sapiens GN=HIST1H2AH PE=1 SV=3 - [H2A1H_HUMAN]
O60814	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3 - [H2B1K_HUMAN]
P68431	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2 - [H31_HUMAN]
P62805	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]
Q86X55	Histone-arginine methyltransferase CARM1 OS=Homo sapiens GN=CARM1 PE=1 SV=3 - [CARM1_HUMAN]
Q96L73	Histone-lysine N-methyltransferase, H3 lysine-36 and H4 lysine-20 specific OS=Homo sapiens GN=NSD1 PE=1 SV=1 - [NSD1_HUMAN]
B0V2L0	HLA-B associated transcript 1 (Fragment) OS=Homo sapiens GN=BAT1 PE=4 SV=1 - [B0V2L0_HUMAN]
B0UX83	HLA-B associated transcript 3 OS=Homo sapiens GN=BAT3 PE=4 SV=1 - [B0UX83_HUMAN]
Q9UBX0	Homeobox expressed in ES cells 1 OS=Homo sapiens GN=HESX1 PE=1 SV=1 - [HESX1_HUMAN]
P39880	Homeobox protein cut-like 1 OS=Homo sapiens GN=CUX1 PE=1 SV=3 - [CUX1_HUMAN]
P56915	Homeobox protein gooseoid OS=Homo sapiens GN=GSC PE=1 SV=2 - [GSC_HUMAN]
P50502	Hsc70-interacting protein OS=Homo sapiens GN=ST13 PE=1 SV=2 - [F10A1_HUMAN]
Q16543	Hsp90 co-chaperone Cdc37 OS=Homo sapiens GN=CDC37 PE=1 SV=1 - [CDC37_HUMAN]
Q16775	Hydroxyacylglutathione hydrolase, mitochondrial OS=Homo sapiens GN=HAGH PE=1 SV=2 - [GLO2_HUMAN]
P00492	Hypoxanthine-guanine phosphoribosyltransferase OS=Homo sapiens GN=HPRT1 PE=1 SV=2 - [HPRT_HUMAN]
P55899	IgG receptor FcRn large subunit p51 OS=Homo sapiens GN=FCGRT PE=1 SV=1 - [FCGRN_HUMAN]
Q8WZA9	Immunity-related GTPase family Q protein OS=Homo sapiens GN=IRGQ PE=1 SV=1 - [IRGQ_HUMAN]
P78318	Immunoglobulin-binding protein 1 OS=Homo sapiens GN=IGBP1 PE=1 SV=1 - [IGBP1_HUMAN]
O00629	Importin subunit alpha-4 OS=Homo sapiens GN=KPNA4 PE=1 SV=1 - [IMA4_HUMAN]

Q14974	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2 - [IMB1_HUMAN]
O94829	Importin-13 OS=Homo sapiens GN=IPO13 PE=1 SV=3 - [IPO13_HUMAN]
O95373	Importin-7 OS=Homo sapiens GN=IPO7 PE=1 SV=1 - [IPO7_HUMAN]
Q96P70	Importin-9 OS=Homo sapiens GN=IPO9 PE=1 SV=3 - [IPO9_HUMAN]
Q9NQS7	Inner centromere protein OS=Homo sapiens GN=INCENP PE=1 SV=3 - [INCE_HUMAN]
Q15181	Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2 - [IPYR_HUMAN]
Q14571	Inositol 1,4,5-trisphosphate receptor type 2 OS=Homo sapiens GN=ITPR2 PE=1 SV=2 - [ITPR2_HUMAN]
P14735	Insulin-degrading enzyme OS=Homo sapiens GN=IDE PE=1 SV=4 - [IDE_HUMAN]
Q8N201	Integrator complex subunit 1 OS=Homo sapiens GN=INTS1 PE=1 SV=2 - [INT1_HUMAN]
Q9H0H0	Integrator complex subunit 2 OS=Homo sapiens GN=INTS2 PE=1 SV=2 - [INT2_HUMAN]
P80217	Interferon-induced 35 kDa protein OS=Homo sapiens GN=IFI35 PE=1 SV=5 - [IN35_HUMAN]
P32456	Interferon-induced guanylate-binding protein 2 OS=Homo sapiens GN=GBP2 PE=2 SV=3 - [GBP2_HUMAN]
Q5JU85	IQ motif and SEC7 domain-containing protein 2 OS=Homo sapiens GN=IQSEC2 PE=1 SV=1 - [IQEC2_HUMAN]
Q9HBB2	Iron regulatory protein 1 OS=Homo sapiens GN=IRP1 PE=2 SV=1 - [Q9HBB2_HUMAN]
Q9BZI1	Iroquois-class homeodomain protein IRX-2 OS=Homo sapiens GN=IRX2 PE=2 SV=2 - [IRX2_HUMAN]
Q2TAA2	Isoamyl acetate-hydrolyzing esterase 1 homolog OS=Homo sapiens GN=IAH1 PE=1 SV=1 - [IAH1_HUMAN]
Q96CN7	Isochorismatase domain-containing protein 1 OS=Homo sapiens GN=ISOC1 PE=1 SV=3 - [ISOC1_HUMAN]
O75874	Isocitrate dehydrogenase [NADP] cytoplasmic OS=Homo sapiens GN=IDH1 PE=1 SV=2 - [IDHC_HUMAN]
Q13907	Isopentenyl-diphosphate Delta-isomerase 1 OS=Homo sapiens GN=IDI1 PE=1 SV=2 - [IDI1_HUMAN]
Q8IYV9	Izumo sperm-egg fusion protein 1 OS=Homo sapiens GN=IZUMO1 PE=1 SV=2 - [IZUM1_HUMAN]
Q9P2K6	Kelch domain-containing protein 5 OS=Homo sapiens GN=KLHDC5 PE=1 SV=2 - [KLDC5_HUMAN]
Q9P2J3	Kelch-like protein 9 OS=Homo sapiens GN=KLHL9 PE=1 SV=2 - [KLHL9_HUMAN]
Q8N1A0	Keratin-like protein KRT222 OS=Homo sapiens GN=KRT222 PE=2 SV=1 - [KT222_HUMAN]
O76009	Keratin, type I cuticular Ha3-I OS=Homo sapiens GN=KRT33A PE=1 SV=2 - [KT33A_HUMAN]
O76013	Keratin, type I cuticular Ha6 OS=Homo sapiens GN=KRT36 PE=1 SV=1 - [KRT36_HUMAN]
P13645	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 - [K1C10_HUMAN]
P08779	Keratin, type I cytoskeletal 16 OS=Homo sapiens GN=KRT16 PE=1 SV=4 - [K1C16_HUMAN]
P35527	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3 - [K1C9_HUMAN]
Q9NSB4	Keratin, type II cuticular Hb2 OS=Homo sapiens GN=KRT82 PE=1 SV=3 - [KRT82_HUMAN]
P78385	Keratin, type II cuticular Hb3 OS=Homo sapiens GN=KRT83 PE=1 SV=2 - [KRT83_HUMAN]
P04264	Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6 - [K2C1_HUMAN]
P35908	Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2 - [K22E_HUMAN]
P12035	Keratin, type II cytoskeletal 3 OS=Homo sapiens GN=KRT3 PE=1 SV=3 - [K2C3_HUMAN]
Q7RTS7	Keratin, type II cytoskeletal 74 OS=Homo sapiens GN=KRT74 PE=1 SV=2 - [K2C74_HUMAN]
O95678	Keratin, type II cytoskeletal 75 OS=Homo sapiens GN=KRT75 PE=1 SV=2 - [K2C75_HUMAN]
Q6KB66	Keratin, type II cytoskeletal 80 OS=Homo sapiens GN=KRT80 PE=1 SV=2 - [K2C80_HUMAN]
Q9HA64	Ketosamine-3-kinase OS=Homo sapiens GN=FN3KRP PE=1 SV=2 - [KT3K_HUMAN]
A1A5A9	KIAA0999 protein OS=Homo sapiens GN=KIAA0999 PE=2 SV=1 - [A1A5A9_HUMAN]
P33176	Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1 - [KINH_HUMAN]
Q96L93	Kinesin-like protein KIF16B OS=Homo sapiens GN=KIF16B PE=1 SV=2 - [KI16B_HUMAN]
O60333	Kinesin-like protein KIF1B OS=Homo sapiens GN=KIF1B PE=1 SV=5 - [KIF1B_HUMAN]
Q86VH2	Kinesin-like protein KIF27 OS=Homo sapiens GN=KIF27 PE=2 SV=1 - [KIF27_HUMAN]
P50748	Kinetochore-associated protein 1 OS=Homo sapiens GN=KNTC1 PE=1 SV=1 - [KNTC1_HUMAN]

Q9UEF7	Klotho OS=Homo sapiens GN=KL PE=1 SV=2 - [KLOT_HUMAN]
E1B6W7	KLRAQ1 protein OS=Homo sapiens GN=KLRAQ1 PE=2 SV=2 - [E1B6W7_HUMAN]
B0S8I7	L antigen family, member 3 OS=Homo sapiens GN=LAGE3 PE=4 SV=1 - [B0S8I7_HUMAN]
Q6ZMR3	L-lactate dehydrogenase A-like 6A OS=Homo sapiens GN=LDHAL6A PE=2 SV=1 - [LDH6A_HUMAN]
P07195	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2 - [LDHB_HUMAN]
E9PH51	L-lactate dehydrogenase OS=Homo sapiens GN=LDHA PE=3 SV=2 - [E9PH51_HUMAN]
Q7Z4W1	L-xylulose reductase OS=Homo sapiens GN=DCXR PE=1 SV=2 - [DCXR_HUMAN]
Q04760	Lactoylglutathione lyase OS=Homo sapiens GN=GLO1 PE=1 SV=4 - [LGUL_HUMAN]
Q03252	Lamin-B2 OS=Homo sapiens GN=LMNB2 PE=1 SV=3 - [LMNB2_HUMAN]
Q59G12	Lamina-associated polypeptide 2, isoforms beta/gamma variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q59G12_HUMAN]
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4 - [LAMA4_HUMAN]
Q9Y6N6	Laminin subunit gamma-3 OS=Homo sapiens GN=LAMC3 PE=2 SV=3 - [LAMC3_HUMAN]
Q9NS86	LanC-like protein 2 OS=Homo sapiens GN=LANCL2 PE=1 SV=1 - [LANC2_HUMAN]
Q01650	Large neutral amino acids transporter small subunit 1 OS=Homo sapiens GN=SLC7A5 PE=1 SV=2 - [LAT1_HUMAN]
Q9BS40	Latexin OS=Homo sapiens GN=LXN PE=1 SV=2 - [LXN_HUMAN]
Q9Y468	Lethal(3)malignant brain tumor-like protein 1 OS=Homo sapiens GN=L3MBTL1 PE=1 SV=3 - [LMBL1_HUMAN]
Q9UIC8	Leucine carboxyl methyltransferase 1 OS=Homo sapiens GN=LCMT1 PE=1 SV=2 - [LCMT1_HUMAN]
Q9Y608	Leucine-rich repeat flightless-interacting protein 2 OS=Homo sapiens GN=LRRFIP2 PE=1 SV=1 - [LRRF2_HUMAN]
Q8N1G4	Leucine-rich repeat-containing protein 47 OS=Homo sapiens GN=LRRC47 PE=1 SV=1 - [LRC47_HUMAN]
Q96JA1	Leucine-rich repeats and immunoglobulin-like domains protein 1 OS=Homo sapiens GN=LRIG1 PE=1 SV=2 - [LRIG1_HUMAN]
P09960	Leukotriene A-4 hydrolase OS=Homo sapiens GN=LTA4H PE=1 SV=2 - [LKHA4_HUMAN]
Q14847	LIM and SH3 domain protein 1 OS=Homo sapiens GN=LASP1 PE=1 SV=2 - [LASP1_HUMAN]
Q9GZY6	Linker for activation of T-cells family member 2 OS=Homo sapiens GN=LAT2 PE=1 SV=1 - [NTAL_HUMAN]
P24666	Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1 PE=1 SV=3 - [PPAC_HUMAN]
P05455	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2 - [LA_HUMAN]
Q5QPQ1	Lysophospholipase II (Fragment) OS=Homo sapiens GN=LYPLA2 PE=2 SV=1 - [Q5QPQ1_HUMAN]
P14174	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4 - [MIF_HUMAN]
P40925	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4 - [MDHC_HUMAN]
Q6P052	MAN1A1 protein OS=Homo sapiens GN=MAN1A1 PE=2 SV=1 - [Q6P052_HUMAN]
A4D2A2	MCM7 minichromosome maintenance deficient 7 (S. cerevisiae) OS=Homo sapiens GN=MCM7 PE=2 SV=1 - [A4D2A2_HUMAN]
Q9UHV7	Mediator of RNA polymerase II transcription subunit 13 OS=Homo sapiens GN=MED13 PE=1 SV=3 - [MED13_HUMAN]
Q969V1	Melanin-concentrating hormone receptor 2 OS=Homo sapiens GN=MCHR2 PE=1 SV=1 - [MCHR2_HUMAN]
P02795	Metallothionein-2 OS=Homo sapiens GN=MT2A PE=1 SV=1 - [MT2_HUMAN]
P50579	Methionine aminopeptidase 2 OS=Homo sapiens GN=METAP2 PE=1 SV=1 - [AMPM2_HUMAN]
Q9BV20	Methylthioribose-1-phosphate isomerase OS=Homo sapiens GN=MRI1 PE=1 SV=1 - [MTNA_HUMAN]
D3YTD2	MICAL-like 2, isoform CRA_a OS=Homo sapiens GN=MICALL2 PE=4 SV=1 - [D3YTD2_HUMAN]
Q8WYQ5	Microprocessor complex subunit DGCR8 OS=Homo sapiens GN=DGCR8 PE=1 SV=1 - [DGCR8_HUMAN]
P78559	Microtubule-associated protein 1A OS=Homo sapiens GN=MAP1A PE=1 SV=6 - [MAP1A_HUMAN]
Q15691	Microtubule-associated protein RP/EB family member 1 OS=Homo sapiens GN=MAPRE1 PE=1 SV=3 - [MARE1_HUMAN]
Q7Z6P5	Minichromosome maintenance complex component 3 (Fragment) OS=Homo sapiens GN=MCM3 PE=2 SV=1 - [Q7Z6P5_HUMAN]
Q9Y3D6	Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2 - [FIS1_HUMAN]
Q15785	Mitochondrial import receptor subunit TOM34 OS=Homo sapiens GN=TOMM34 PE=1 SV=2 - [TOM34_HUMAN]
Q99797	Mitochondrial intermediate peptidase OS=Homo sapiens GN=MIPEP PE=1 SV=2 - [MIPEP_HUMAN]

Q8TD08	Mitogen-activated protein kinase 15 OS=Homo sapiens GN=MAPK15 PE=1 SV=1 - [MK15_HUMAN]
Q12852	Mitogen-activated protein kinase kinase kinase 12 OS=Homo sapiens GN=MAP3K12 PE=1 SV=2 - [M3K12_HUMAN]
Q9ULH7	MKL/myocardin-like protein 2 OS=Homo sapiens GN=MKL2 PE=1 SV=3 - [MKL2_HUMAN]
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3 - [MOES_HUMAN]
Q2QL34	Mpv17-like protein OS=Homo sapiens GN=MPV17L PE=1 SV=1 - [MP17L_HUMAN]
B5AK60	Multidrug resistance protein 1 OS=Homo sapiens GN=ABCB1 PE=2 SV=1 - [B5AK60_HUMAN]
Q5T3U5	Multidrug resistance-associated protein 7 OS=Homo sapiens GN=ABCC10 PE=1 SV=1 - [MRP7_HUMAN]
Q7Z7M0	Multiple epidermal growth factor-like domains protein 8 OS=Homo sapiens GN=MEGF8 PE=1 SV=2 - [MEGF8_HUMAN]
P60201	Myelin proteolipid protein OS=Homo sapiens GN=PLP1 PE=1 SV=2 - [MYPR_HUMAN]
P41218	Myeloid cell nuclear differentiation antigen OS=Homo sapiens GN=MNDA PE=1 SV=1 - [MNDA_HUMAN]
P12829	Myosin light chain 4 OS=Homo sapiens GN=MYL4 PE=1 SV=3 - [MYL4_HUMAN]
P60660	Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2 - [MYL6_HUMAN]
Q96A32	Myosin regulatory light chain 2, skeletal muscle isoform OS=Homo sapiens GN=MYLPF PE=2 SV=1 - [MLRS_HUMAN]
P35580	Myosin-10 OS=Homo sapiens GN=MYH10 PE=1 SV=3 - [MYH10_HUMAN]
P35579	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4 - [MYH9_HUMAN]
B0I1T2	Myosin-Ig OS=Homo sapiens GN=MYO1G PE=1 SV=2 - [MYO1G_HUMAN]
Q3MIV8	Myosin, heavy chain 11, smooth muscle OS=Homo sapiens GN=MYH11 PE=2 SV=1 - [Q3MIV8_HUMAN]
Q9BUF9	Myosin, light chain 9, regulatory OS=Homo sapiens GN=MYL9 PE=2 SV=1 - [Q9BUF9_HUMAN]
P58546	Myotrophin OS=Homo sapiens GN=MTPN PE=1 SV=2 - [MTPN_HUMAN]
Q76KP1	N-acetyl-beta-glucosaminyl-glycoprotein 4-beta-N-acetylgalactosaminyltransferase 1 OS=Homo sapiens GN=B4GALNT4 PE=1 SV=1 - [B4GN4_HUMAN]
Q9BXD5	N-acetylneuraminase lyase OS=Homo sapiens GN=NPL PE=2 SV=1 - [NPL_HUMAN]
Q8TBE9	N-acetylneuraminase-9-phosphatase OS=Homo sapiens GN=NANP PE=1 SV=1 - [NANP_HUMAN]
Q8TBK2	N-lysine methyltransferase SETD6 OS=Homo sapiens GN=SETD6 PE=1 SV=2 - [SETD6_HUMAN]
Q9NXH9	N(2),N(2)-dimethylguanosine tRNA methyltransferase OS=Homo sapiens GN=TRMT1 PE=1 SV=1 - [TRM1_HUMAN]
O94760	N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 OS=Homo sapiens GN=DDAH1 PE=1 SV=3 - [DDAH1_HUMAN]
A2A2U4	NAD(P)H dehydrogenase, quinone 2 (Fragment) OS=Homo sapiens GN=NQO2 PE=4 SV=1 - [A2A2U4_HUMAN]
Q16795	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitochondrial OS=Homo sapiens GN=NDUFA9 PE=1 SV=2 - [NDUA9_HUMAN]
Q13765	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=1 SV=1 - [NACA_HUMAN]
Q9H0M0	NEDD4-like E3 ubiquitin-protein ligase WWP1 OS=Homo sapiens GN=WWP1 PE=1 SV=1 - [WWP1_HUMAN]
Q15843	NEDD8 OS=Homo sapiens GN=NEDD8 PE=1 SV=1 - [NEDD8_HUMAN]
Q8TBC4	NEDD8-activating enzyme E1 catalytic subunit OS=Homo sapiens GN=UBA3 PE=1 SV=2 - [UBA3_HUMAN]
P61081	NEDD8-conjugating enzyme Ubc12 OS=Homo sapiens GN=UBE2M PE=1 SV=1 - [UBC12_HUMAN]
Q8WXH0	Nesprin-2 OS=Homo sapiens GN=SYNE2 PE=1 SV=3 - [SYNE2_HUMAN]
Q6ZN44	Netrin receptor UNC5A OS=Homo sapiens GN=UNC5A PE=2 SV=3 - [UNC5A_HUMAN]
Q8NEY1	Neuron navigator 1 OS=Homo sapiens GN=NAV1 PE=1 SV=2 - [NAV1_HUMAN]
Q8IVL1	Neuron navigator 2 OS=Homo sapiens GN=NAV2 PE=1 SV=3 - [NAV2_HUMAN]
Q9P2E3	NFX1-type zinc finger-containing protein 1 OS=Homo sapiens GN=ZNF1 PE=1 SV=2 - [ZNF1_HUMAN]
Q8NBF2	NHL repeat-containing protein 2 OS=Homo sapiens GN=NHLRC2 PE=1 SV=1 - [NHLRC2_HUMAN]
O60934	Nibrin OS=Homo sapiens GN=NBN PE=1 SV=1 - [NBN_HUMAN]
C9J8U2	Nicotinate phosphoribosyltransferase domain containing 1, isoform CRA_e OS=Homo sapiens GN=NAPRT1 PE=4 SV=1 - [C9J8U2_HUMAN]
Q9GZT8	NIF3-like protein 1 OS=Homo sapiens GN=NIF3L1 PE=1 SV=2 - [NIF3L_HUMAN]
Q6KC79	Nipped-B-like protein OS=Homo sapiens GN=NIPBL PE=1 SV=2 - [NIPBL_HUMAN]
B1AQP4	Nitrilase 1 OS=Homo sapiens GN=NIT1 PE=4 SV=1 - [B1AQP4_HUMAN]

Q9NPP4	NLR family CARD domain-containing protein 4 OS=Homo sapiens GN=NLRC4 PE=1 SV=2 - [NLRC4_HUMAN]
Q5SYT8	Novel protein similar to Pre-B cell enhancing factor (PBEF) (Fragment) OS=Homo sapiens GN=RP11-92J19.4 PE=1 SV=1 - [Q5SYT8_HUMAN]
Q9UNZ2	NSFL1 cofactor p47 OS=Homo sapiens GN=NSFL1C PE=1 SV=2 - [NSF1C_HUMAN]
Q09161	Nuclear cap-binding protein subunit 1 OS=Homo sapiens GN=NCBP1 PE=1 SV=1 - [NCBP1_HUMAN]
B5B2M6	Nuclear factor of activated T-cells c1 isoform IA-deltaIX OS=Homo sapiens GN=NFATC1 PE=2 SV=1 - [B5B2M6_HUMAN]
Q9Y266	Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1 - [NUDC_HUMAN]
Q8TEM1	Nuclear pore membrane glycoprotein 210 OS=Homo sapiens GN=NUP210 PE=1 SV=3 - [PO210_HUMAN]
P61970	Nuclear transport factor 2 OS=Homo sapiens GN=NUTF2 PE=1 SV=1 - [NUTF2_HUMAN]
Q9H1E3	Nuclear ubiquitous casein and cyclin-dependent kinases substrate OS=Homo sapiens GN=NUCKS1 PE=1 SV=1 - [NUCKS_HUMAN]
P67809	Nuclease-sensitive element-binding protein 1 OS=Homo sapiens GN=YBX1 PE=1 SV=3 - [YBOX1_HUMAN]
Q02818	Nucleobindin-1 OS=Homo sapiens GN=NUCB1 PE=1 SV=4 - [NUCB1_HUMAN]
Q13823	Nucleolar GTP-binding protein 2 OS=Homo sapiens GN=GNL2 PE=1 SV=1 - [NOG2_HUMAN]
P19338	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3 - [NUCL_HUMAN]
P06748	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2 - [NPM_HUMAN]
Q32Q12	Nucleoside diphosphate kinase OS=Homo sapiens GN=NME1-NME2 PE=2 SV=1 - [Q32Q12_HUMAN]
Q96RS6	NudC domain-containing protein 1 OS=Homo sapiens GN=NUDCD1 PE=1 SV=2 - [NUDC1_HUMAN]
Q8IVD9	NudC domain-containing protein 3 OS=Homo sapiens GN=NUDCD3 PE=1 SV=3 - [NUDC3_HUMAN]
Q7Z7N6	NUDT1 protein OS=Homo sapiens GN=NUDT1 PE=2 SV=2 - [Q7Z7N6_HUMAN]
Q9NTK5	Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=1 SV=2 - [OLA1_HUMAN]
O95007	Olfactory receptor 6B1 OS=Homo sapiens GN=OR6B1 PE=2 SV=1 - [OR6B1_HUMAN]
Q9NQR4	Omega-amidase NIT2 OS=Homo sapiens GN=NIT2 PE=1 SV=1 - [NIT2_HUMAN]
Q8WWZ8	Oncoprotein-induced transcript 3 protein OS=Homo sapiens GN=OIT3 PE=1 SV=2 - [OIT3_HUMAN]
Q96CV9	Optineurin OS=Homo sapiens GN=OPTN PE=1 SV=2 - [OPTN_HUMAN]
B5LY72	Orotidine 5'-phosphate decarboxylase OS=Homo sapiens GN=UMPS PE=2 SV=1 - [B5LY72_HUMAN]
Q8TE49	OTU domain-containing protein 7A OS=Homo sapiens GN=OTUD7A PE=1 SV=1 - [OTU7A_HUMAN]
B7WNL4	Oxysterol-binding protein OS=Homo sapiens GN=OSBP PE=3 SV=2 - [B7WNL4_HUMAN]
B4DTR3	Oxysterol-binding protein OS=Homo sapiens GN=OSBP2 PE=2 SV=1 - [B4DTR3_HUMAN]
Q6FIA3	PAC SIN2 protein OS=Homo sapiens GN=PAC SIN2 PE=2 SV=1 - [Q6FIA3_HUMAN]
Q5VU21	PAI-1 mRNA-binding protein variant OS=Homo sapiens GN=SERBP1 PE=2 SV=1 - [Q5VU21_HUMAN]
O75182	Paired amphipathic helix protein Sin3b OS=Homo sapiens GN=SIN3B PE=1 SV=2 - [SIN3B_HUMAN]
Q5T0S4	Palmitoyl-protein thioesterase 1 (Fragment) OS=Homo sapiens GN=PPT1 PE=2 SV=1 - [Q5T0S4_HUMAN]
Q96RD7	Pannexin-1 OS=Homo sapiens GN=PANX1 PE=1 SV=4 - [PANX1_HUMAN]
Q9UL41	Paraneoplastic antigen Ma3 OS=Homo sapiens GN=PNMA3 PE=2 SV=2 - [PNMA3_HUMAN]
P20962	Parathyrosin OS=Homo sapiens GN=PTMS PE=1 SV=2 - [PTMS_HUMAN]
Q9UBV8	Peflin OS=Homo sapiens GN=PEF1 PE=1 SV=1 - [PEF1_HUMAN]
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2 - [PPIA_HUMAN]
Q00688	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1 - [FKBP3_HUMAN]
Q02790	Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3 - [FKBP4_HUMAN]
Q13526	Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1 OS=Homo sapiens GN=PIN1 PE=1 SV=1 - [PIN1_HUMAN]
Q9Y3C6	Peptidyl-prolyl cis-trans isomerase-like 1 OS=Homo sapiens GN=PPIL1 PE=1 SV=1 - [PPIL1_HUMAN]
Q567Q0	Peptidylprolyl isomerase A (Cyclophilin A) OS=Homo sapiens GN=PPIA PE=2 SV=1 - [Q567Q0_HUMAN]
O60664	Perilipin-3 OS=Homo sapiens GN=PLIN3 PE=1 SV=3 - [PLIN3_HUMAN]
Q06830	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1 - [PRDX1_HUMAN]

P32119	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5 - [PRDX2_HUMAN]
P30044	Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=4 - [PRDX5_HUMAN]
<i>P30041</i>	<i>Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3 - [PRDX6_HUMAN]</i>
Q8WW12	PEST proteolytic signal-containing nuclear protein OS=Homo sapiens GN=PCNP PE=1 SV=2 - [PCNP_HUMAN]
Q9NSD9	Phenylalanyl-tRNA synthetase beta chain OS=Homo sapiens GN=FARSB PE=1 SV=3 - [SYFB_HUMAN]
Q92539	Phosphatidate phosphatase LPIN2 OS=Homo sapiens GN=LPIN2 PE=1 SV=1 - [LPIN2_HUMAN]
Q9UKL6	Phosphatidylcholine transfer protein OS=Homo sapiens GN=PCTP PE=1 SV=1 - [PPCT_HUMAN]
P30086	Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3 - [PEBP1_HUMAN]
B3KYB7	Phosphatidylinositol transfer protein, beta (Fragment) OS=Homo sapiens GN=PITPNB PE=4 SV=1 - [B3KYB7_HUMAN]
Q92835	Phosphatidylinositol-3,4,5-trisphosphate 5-phosphatase 1 OS=Homo sapiens GN=INPP5D PE=1 SV=2 - [SHIP1_HUMAN]
Q5T9C9	Phosphatidylinositol-4-phosphate 5-kinase-like protein 1 OS=Homo sapiens GN=PIP5KL1 PE=2 SV=2 - [PI5L1_HUMAN]
P42336	Phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha isoform OS=Homo sapiens GN=PIK3CA PE=1 SV=2 - [PK3CA_HUMAN]
P48426	Phosphatidylinositol-5-phosphate 4-kinase type-2 alpha OS=Homo sapiens GN=PIP4K2A PE=1 SV=2 - [PI42A_HUMAN]
Q96G03	Phosphoglucomutase-2 OS=Homo sapiens GN=PGM2 PE=1 SV=4 - [PGM2_HUMAN]
Q5SZU1	Phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=2 SV=1 - [Q5SZU1_HUMAN]
P00558	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3 - [PGK1_HUMAN]
P07205	Phosphoglycerate kinase 2 OS=Homo sapiens GN=PGK2 PE=1 SV=3 - [PGK2_HUMAN]
P18669	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2 - [PGAM1_HUMAN]
A6NDG6	Phosphoglycolate phosphatase OS=Homo sapiens GN=PGP PE=1 SV=1 - [PGP_HUMAN]
Q5T1Z0	Phospholysine phosphohistidine inorganic pyrophosphate phosphatase OS=Homo sapiens GN=LHPP PE=2 SV=1 - [Q5T1Z0_HUMAN]
Q9HAB8	Phosphopantothenate--cysteine ligase OS=Homo sapiens GN=PPCS PE=1 SV=2 - [PPCS_HUMAN]
O15067	Phosphoribosylformylglycinamide synthase OS=Homo sapiens GN=PFAS PE=1 SV=4 - [PUR4_HUMAN]
P46019	Phosphorylase b kinase regulatory subunit alpha, liver isoform OS=Homo sapiens GN=PHKA2 PE=1 SV=1 - [KPB2_HUMAN]
Q9Y617	Phosphoserine aminotransferase OS=Homo sapiens GN=PSAT1 PE=1 SV=2 - [SERC_HUMAN]
O00625	Pirin OS=Homo sapiens GN=PIR PE=1 SV=1 - [PIR_HUMAN]
Q9GZP4	PITH domain-containing protein 1 OS=Homo sapiens GN=PITHD1 PE=1 SV=1 - [PITH1_HUMAN]
P13796	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6 - [PLSL_HUMAN]
P43034	Platelet-activating factor acetylhydrolase IB subunit alpha OS=Homo sapiens GN=PFAH1B1 PE=1 SV=2 - [LIS1_HUMAN]
P68402	Platelet-activating factor acetylhydrolase IB subunit beta OS=Homo sapiens GN=PFAH1B2 PE=1 SV=1 - [PA1B2_HUMAN]
Q15102	Platelet-activating factor acetylhydrolase IB subunit gamma OS=Homo sapiens GN=PFAH1B3 PE=1 SV=1 - [PA1B3_HUMAN]
Q8IVE3	Pleckstrin homology domain-containing family H member 2 OS=Homo sapiens GN=PLEKHH2 PE=2 SV=2 - [PKHH2_HUMAN]
A0JP02	PLEKHA5 protein OS=Homo sapiens GN=PLEKHA5 PE=2 SV=1 - [A0JP02_HUMAN]
Q68Y55	Poly(RC) binding protein 2 OS=Homo sapiens GN=PCBP2 PE=2 SV=1 - [Q68Y55_HUMAN]
Q15365	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]
P26599	Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1 - [PTBP1_HUMAN]
Q14721	Potassium voltage-gated channel subfamily B member 1 OS=Homo sapiens GN=KCNB1 PE=1 SV=2 - [KCNB1_HUMAN]
Q14CL3	Potassium voltage-gated channel, subfamily H (Eag-related), member 1 OS=Homo sapiens GN=KCNH1 PE=2 SV=1 - [Q14CL3_HUMAN]
Q6S8J7	POTE ankyrin domain family member A OS=Homo sapiens GN=POTEA PE=2 SV=1 - [POTEA_HUMAN]
P57071	PR domain zinc finger protein 15 OS=Homo sapiens GN=PRDM15 PE=2 SV=4 - [PRD15_HUMAN]
O60925	Prefoldin subunit 1 OS=Homo sapiens GN=PFDN1 PE=1 SV=2 - [PFD1_HUMAN]
Q9UHV9	Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1 - [PFD2_HUMAN]
Q9NQP4	Prefoldin subunit 4 OS=Homo sapiens GN=PFDN4 PE=1 SV=1 - [PFD4_HUMAN]
Q99471	Prefoldin subunit 5 OS=Homo sapiens GN=PFDN5 PE=1 SV=2 - [PFD5_HUMAN]

O15212	Prefoldin subunit 6 OS=Homo sapiens GN=PFDN6 PE=1 SV=1 - [PFD6_HUMAN]
Q86TM3	Probable ATP-dependent RNA helicase DDX53 OS=Homo sapiens GN=DDX53 PE=1 SV=3 - [DDX53_HUMAN]
Q9HD20	Probable cation-transporting ATPase 13A1 OS=Homo sapiens GN=ATP13A1 PE=1 SV=2 - [AT131_HUMAN]
Q96GX9	Probable methylthioribulose-1-phosphate dehydratase OS=Homo sapiens GN=APIP PE=1 SV=1 - [MTNB_HUMAN]
Q93008	Probable ubiquitin carboxyl-terminal hydrolase FAF-X OS=Homo sapiens GN=USP9X PE=1 SV=3 - [USP9X_HUMAN]
O00507	Probable ubiquitin carboxyl-terminal hydrolase FAF-Y OS=Homo sapiens GN=USP9Y PE=1 SV=2 - [USP9Y_HUMAN]
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2 - [PROF1_HUMAN]
Q8WUM4	Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1 - [PDC6I_HUMAN]
O14737	Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3 - [PDCD5_HUMAN]
Q9HCU5	Prolactin regulatory element-binding protein OS=Homo sapiens GN=PREB PE=1 SV=2 - [PREB_HUMAN]
Q9UQ80	Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=3 - [PA2G4_HUMAN]
Q96HE9	Proline-rich protein 11 OS=Homo sapiens GN=PRR11 PE=1 SV=1 - [PRR11_HUMAN]
Q8IVL6	Prolyl 3-hydroxylase 3 OS=Homo sapiens GN=LEPREL2 PE=2 SV=1 - [P3H3_HUMAN]
P48147	Prolyl endopeptidase OS=Homo sapiens GN=PREP PE=1 SV=2 - [PPCE_HUMAN]
B4DP21	Prostaglandin E synthase 3 (Cytosolic), isoform CRA_d OS=Homo sapiens GN=PTGES3 PE=2 SV=1 - [B4DP21_HUMAN]
A6XGL3	Protease serine 1 OS=Homo sapiens PE=2 SV=1 - [A6XGL3_HUMAN]
B1AN99	Protease, serine, 3 (Fragment) OS=Homo sapiens GN=PRSS3 PE=3 SV=1 - [B1AN99_HUMAN]
A8K3Z3	Proteasome (Prosome, macropain) 26S subunit, ATPase, 5, isoform CRA_b OS=Homo sapiens GN=PSMC5 PE=2 SV=1 - [A8K3Z3_HUMAN]
B1AJY5	Proteasome (Prosome, macropain) 26S subunit, non-ATPase, 10 OS=Homo sapiens GN=PSMD10 PE=4 SV=1 - [B1AJY5_HUMAN]
Q4VXG9	Proteasome (Prosome, macropain) 26S subunit, non-ATPase, 5 OS=Homo sapiens GN=PSMD5 PE=2 SV=1 - [Q4VXG9_HUMAN]
Q5QPM7	Proteasome (Prosome, macropain) inhibitor subunit 1 (PI31) OS=Homo sapiens GN=PSMF1 PE=2 SV=2 - [Q5QPM7_HUMAN]
Q06323	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1 - [PSME1_HUMAN]
Q9UL46	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=4 - [PSME2_HUMAN]
O95456	Proteasome assembly chaperone 1 OS=Homo sapiens GN=PSMG1 PE=1 SV=1 - [PSMG1_HUMAN]
P25787	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2 - [PSA2_HUMAN]
P25788	Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 PE=1 SV=2 - [PSA3_HUMAN]
P28066	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3 - [PSA5_HUMAN]
P60900	Proteasome subunit alpha type-6 OS=Homo sapiens GN=PSMA6 PE=1 SV=1 - [PSA6_HUMAN]
O14818	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1 - [PSA7_HUMAN]
P20618	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2 - [PSB1_HUMAN]
P49721	Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1 - [PSB2_HUMAN]
P49720	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2 - [PSB3_HUMAN]
P28070	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4 - [PSB4_HUMAN]
P28074	Proteasome subunit beta type-5 OS=Homo sapiens GN=PSMB5 PE=1 SV=3 - [PSB5_HUMAN]
P28072	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4 - [PSB6_HUMAN]
Q99436	Proteasome subunit beta type-7 OS=Homo sapiens GN=PSMB7 PE=1 SV=1 - [PSB7_HUMAN]
Q5VYK3	Proteasome-associated protein ECM29 homolog OS=Homo sapiens GN=ECM29 PE=1 SV=2 - [ECM29_HUMAN]
Q8IVF2	Protein AHNAK2 OS=Homo sapiens GN=AHNAK2 PE=1 SV=2 - [AHNK2_HUMAN]
A8MTP3	Protein arginine methyltransferase 5, isoform CRA_c OS=Homo sapiens GN=PRMT5 PE=4 SV=1 - [A8MTP3_HUMAN]
Q9UKV8	Protein argonaute-2 OS=Homo sapiens GN=EIF2C2 PE=1 SV=3 - [AGO2_HUMAN]
Q8TCG1	Protein CIP2A OS=Homo sapiens GN=KIAA1524 PE=1 SV=2 - [CIP2A_HUMAN]
Q5TDH0	Protein DDI1 homolog 2 OS=Homo sapiens GN=DDI2 PE=1 SV=1 - [DDI2_HUMAN]
Q8N919	Protein deltex-3 OS=Homo sapiens GN=DTX3 PE=2 SV=2 - [DTX3_HUMAN]

Q99497	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2 - [PARK7_HUMAN]
Q96BN8	Protein FAM105B OS=Homo sapiens GN=FAM105B PE=1 SV=3 - [F105B_HUMAN]
Q9NRY5	Protein FAM114A2 OS=Homo sapiens GN=FAM114A2 PE=1 SV=4 - [F1142_HUMAN]
Q96KS9	Protein FAM167A OS=Homo sapiens GN=FAM167A PE=2 SV=1 - [F167A_HUMAN]
P58499	Protein FAM3B OS=Homo sapiens GN=FAM3B PE=1 SV=2 - [FAM3B_HUMAN]
Q5VWP2	Protein FAM46C OS=Homo sapiens GN=FAM46C PE=2 SV=1 - [FA46C_HUMAN]
Q9NUQ9	Protein FAM49B OS=Homo sapiens GN=FAM49B PE=1 SV=1 - [FA49B_HUMAN]
Q6UWF7	Protein FAM55D OS=Homo sapiens GN=FAM55D PE=2 SV=1 - [FA55D_HUMAN]
Q8IWB4	Protein FAM75A7 OS=Homo sapiens GN=FAM75A7 PE=2 SV=4 - [F75A7_HUMAN]
Q9C0D2	Protein KIAA1731 OS=Homo sapiens GN=KIAA1731 PE=2 SV=4 - [K1731_HUMAN]
P05771	Protein kinase C beta type OS=Homo sapiens GN=PRKCB PE=1 SV=4 - [KPCB_HUMAN]
Q96GA3	Protein LTV1 homolog OS=Homo sapiens GN=LTV1 PE=1 SV=1 - [LTV1_HUMAN]
Q8WZA0	Protein LZIC OS=Homo sapiens GN=LZIC PE=1 SV=1 - [LZIC_HUMAN]
Q9P2P1	Protein NYNRIN OS=Homo sapiens GN=NYNRIN PE=1 SV=3 - [NYNRI_HUMAN]
Q15435	Protein phosphatase 1 regulatory subunit 7 OS=Homo sapiens GN=PPP1R7 PE=1 SV=1 - [PP1R7_HUMAN]
P35813	Protein phosphatase 1A OS=Homo sapiens GN=PPM1A PE=1 SV=1 - [PPM1A_HUMAN]
O75688	Protein phosphatase 1B OS=Homo sapiens GN=PPM1B PE=1 SV=1 - [PPM1B_HUMAN]
A6PVN5	Protein phosphatase 2A activator, regulatory subunit 4 OS=Homo sapiens GN=PPP2R4 PE=4 SV=1 - [A6PVN5_HUMAN]
Q9Y570	Protein phosphatase methylesterase 1 OS=Homo sapiens GN=PPME1 PE=1 SV=3 - [PPME1_HUMAN]
Q96M27	Protein PRRC1 OS=Homo sapiens GN=PRRC1 PE=1 SV=1 - [PRRC1_HUMAN]
Q9P258	Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2 - [RCC2_HUMAN]
P31949	Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2 - [S10AB_HUMAN]
P26447	Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1 - [S10A4_HUMAN]
<i>P06703</i>	<i>Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1 - [S10A6_HUMAN]</i>
Q13796	Protein Shroom2 OS=Homo sapiens GN=SHROOM2 PE=1 SV=1 - [SHRM2_HUMAN]
Q9UBV4	Protein Wnt-16 OS=Homo sapiens GN=WNT16 PE=2 SV=1 - [WNT16_HUMAN]
Q9H900	Protein zwilch homolog OS=Homo sapiens GN=ZWILCH PE=1 SV=2 - [ZWILC_HUMAN]
B7Z972	Protein-L-isoaspartate O-methyltransferase OS=Homo sapiens GN=PCMT1 PE=2 SV=1 - [B7Z972_HUMAN]
Q9Y5H8	Protocadherin alpha-3 OS=Homo sapiens GN=PCDHA3 PE=1 SV=1 - [PCDA3_HUMAN]
Q9UN71	Protocadherin gamma-B4 OS=Homo sapiens GN=PCDHGB4 PE=2 SV=1 - [PCDGG_HUMAN]
Q567Q5	PSMA4 protein OS=Homo sapiens GN=PSMA4 PE=2 SV=1 - [Q567Q5_HUMAN]
P00491	Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2 - [PNPH_HUMAN]
A6NGE7	Putative 2-oxo-4-hydroxy-4-carboxy-5-ureidoimidazole decarboxylase OS=Homo sapiens GN=PRHOXNB PE=5 SV=2 - [URAD_HUMAN]
P20848	Putative alpha-1-antitrypsin-related protein OS=Homo sapiens GN=SERPINA2 PE=5 SV=1 - [A1ATR_HUMAN]
A8MPP1	Putative ATP-dependent RNA helicase DDX11-like protein 8 OS=Homo sapiens GN=DDX11L8 PE=1 SV=1 - [D11L8_HUMAN]
A6NDK9	Putative golgin subfamily A member 6C OS=Homo sapiens GN=GOLGA6C PE=5 SV=1 - [GOG6C_HUMAN]
P48741	Putative heat shock 70 kDa protein 7 OS=Homo sapiens GN=HSPA7 PE=5 SV=2 - [HSP77_HUMAN]
Q6GMV3	Putative peptidyl-tRNA hydrolase PTRHD1 OS=Homo sapiens GN=PTRHD1 PE=1 SV=1 - [PTRD1_HUMAN]
Q9C0F0	Putative Polycomb group protein ASXL3 OS=Homo sapiens GN=ASXL3 PE=2 SV=3 - [ASXL3_HUMAN]
A4D2B8	Putative postmeiotic segregation increased 2-like protein 1 OS=Homo sapiens GN=PMS2P1 PE=5 SV=1 - [PM2P1_HUMAN]
P98179	Putative RNA-binding protein 3 OS=Homo sapiens GN=RBM3 PE=1 SV=1 - [RBM3_HUMAN]
Q9P1A2	Putative serine/threonine-protein phosphatase 4 regulatory subunit 1-like OS=Homo sapiens GN=PPP4R1L PE=5 SV=2 - [PP4RL_HUMAN]
Q495Y7	Putative speedy protein E7 OS=Homo sapiens GN=SPDYE7P PE=5 SV=1 - [SPDE7_HUMAN]

Q5QP74	Putative uncharacterized protein C10orf115 OS=Homo sapiens GN=C10orf115 PE=2 SV=2 - [CJ115_HUMAN]
Q8WZ69	Putative uncharacterized protein C11orf40 OS=Homo sapiens GN=C11orf40 PE=2 SV=1 - [CK040_HUMAN]
Q96BT1	Putative uncharacterized protein C3orf49 OS=Homo sapiens GN=C3orf49 PE=2 SV=1 - [CC049_HUMAN]
Q53XA7	Putative uncharacterized protein DKFZp686F13224 OS=Homo sapiens GN=DKFZp686F13224 PE=2 SV=1 - [Q53XA7_HUMAN]
A8MT66	Putative uncharacterized protein ENSP00000383407 OS=Homo sapiens PE=5 SV=1 - [YU005_HUMAN]
Q6ICG7	Putative uncharacterized protein LOC100131404 OS=Homo sapiens PE=2 SV=1 - [YV028_HUMAN]
Q6ZTK2	Putative uncharacterized protein LOC400499 OS=Homo sapiens PE=2 SV=1 - [YP015_HUMAN]
Q9NSJ1	Putative zinc finger protein 355P OS=Homo sapiens GN=ZNF355P PE=5 SV=2 - [Z355P_HUMAN]
Q96GD0	Pyridoxal phosphate phosphatase OS=Homo sapiens GN=PDXP PE=1 SV=2 - [PLPP_HUMAN]
P14618	Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM2 PE=1 SV=4 - [KPYM_HUMAN]
P30613	Pyruvate kinase isozymes R/L OS=Homo sapiens GN=PKLR PE=1 SV=2 - [KPYR_HUMAN]
Q53FA7	Quinone oxidoreductase PIG3 OS=Homo sapiens GN=TP53I3 PE=1 SV=2 - [QORX_HUMAN]
P50395	Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GDI2 PE=1 SV=2 - [GDIB_HUMAN]
Q96NA2	Rab-interacting lysosomal protein OS=Homo sapiens GN=RILP PE=1 SV=1 - [RILP_HUMAN]
Q5HYI8	Rab-like protein 3 OS=Homo sapiens GN=RABL3 PE=1 SV=1 - [RABL3_HUMAN]
Q8WVF3	RAB28 protein OS=Homo sapiens GN=RAB28 PE=2 SV=1 - [Q8WVF3_HUMAN]
B4DKD7	RAB5B, member RAS oncogene family, isoform CRA_b OS=Homo sapiens GN=RAB5B PE=2 SV=1 - [B4DKD7_HUMAN]
P35241	Radixin OS=Homo sapiens GN=RDY PE=1 SV=1 - [RADI_HUMAN]
P46060	Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1 - [RAGP1_HUMAN]
O60518	Ran-binding protein 6 OS=Homo sapiens GN=RANBP6 PE=1 SV=2 - [RNBP6_HUMAN]
P43487	Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1 - [RANG_HUMAN]
Q5T3S7	RAP1 GTPase activating protein OS=Homo sapiens GN=RAP1GAP PE=2 SV=1 - [Q5T3S7_HUMAN]
P63000	Ras-related C3 botulinum toxin substrate 1 OS=Homo sapiens GN=RAC1 PE=1 SV=1 - [RAC1_HUMAN]
P61026	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1 - [RAB10_HUMAN]
P62491	Ras-related protein Rab-11A OS=Homo sapiens GN=RAB11A PE=1 SV=3 - [RB11A_HUMAN]
P61106	Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4 - [RAB14_HUMAN]
P62820	Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=1 SV=3 - [RAB1A_HUMAN]
Q9H0U4	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1 - [RAB1B_HUMAN]
P51148	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2 - [RAB5C_HUMAN]
Q8TDY2	RB1-inducible coiled-coil protein 1 OS=Homo sapiens GN=RB1CC1 PE=1 SV=3 - [RBCC1_HUMAN]
O43665	Regulator of G-protein signaling 10 OS=Homo sapiens GN=RGS10 PE=1 SV=2 - [RGS10_HUMAN]
P00352	<i>Retinal dehydrogenase 1 OS=Homo sapiens GN=ALDH1A1 PE=1 SV=2 - [AL1A1_HUMAN]</i>
Q9Y2V3	Retinal homeobox protein Rx OS=Homo sapiens GN=RAX PE=1 SV=2 - [RX_HUMAN]
P52565	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDI1 PE=1 SV=3 - [GDIR1_HUMAN]
A1A4S6	Rho GTPase-activating protein 10 OS=Homo sapiens GN=ARHGAP10 PE=1 SV=1 - [RHG10_HUMAN]
Q52LW3	Rho GTPase-activating protein 29 OS=Homo sapiens GN=ARHGAP29 PE=1 SV=2 - [RHG29_HUMAN]
Q7Z6I6	Rho GTPase-activating protein 30 OS=Homo sapiens GN=ARHGAP30 PE=1 SV=3 - [RHG30_HUMAN]
O43182	Rho GTPase-activating protein 6 OS=Homo sapiens GN=ARHGAP6 PE=1 SV=3 - [RHG06_HUMAN]
Q92974	Rho guanine nucleotide exchange factor 2 OS=Homo sapiens GN=ARHGEF2 PE=1 SV=4 - [ARHG2_HUMAN]
P13489	Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2 - [RINI_HUMAN]
Q5D1E8	Ribonuclease ZC3H12A OS=Homo sapiens GN=ZC3H12A PE=1 SV=1 - [ZC12A_HUMAN]
P49247	Ribose-5-phosphate isomerase OS=Homo sapiens GN=RPIA PE=1 SV=3 - [RPIA_HUMAN]
P60891	Ribose-phosphate pyrophosphokinase 1 OS=Homo sapiens GN=PRPS1 PE=1 SV=2 - [PRPS1_HUMAN]

P11908	Ribose-phosphate pyrophosphokinase 2 OS=Homo sapiens GN=PRPS2 PE=1 SV=2 - [PRPS2_HUMAN]
Q5VVC8	Ribosomal protein L11 (Fragment) OS=Homo sapiens GN=RPL11 PE=2 SV=1 - [Q5VVC8_HUMAN]
E7EQV9	Ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=3 SV=1 - [E7EQV9_HUMAN]
C9JNW5	Ribosomal protein L24, isoform CRA_e OS=Homo sapiens GN=RPL24 PE=4 SV=1 - [C9JNW5_HUMAN]
D6R9X9	Ribosomal protein L37 OS=Homo sapiens GN=RPL37 PE=3 SV=1 - [D6R9X9_HUMAN]
B3KTM6	Ribosomal protein L5, isoform CRA_b OS=Homo sapiens GN=RPL5 PE=2 SV=1 - [B3KTM6_HUMAN]
Q5T8U3	Ribosomal protein L7a (Fragment) OS=Homo sapiens GN=RPL7A PE=4 SV=1 - [Q5T8U3_HUMAN]
C9J1C5	Ribosomal protein S27-like, isoform CRA_a OS=Homo sapiens GN=RPS27L PE=3 SV=1 - [C9J1C5_HUMAN]
D6R9B6	Ribosomal protein S3A, isoform CRA_e OS=Homo sapiens GN=RPS3A PE=4 SV=1 - [D6R9B6_HUMAN]
Q5SVM6	Ribosomal protein S6 kinase, 90kDa, polypeptide 1 (Fragment) OS=Homo sapiens GN=RPS6KA1 PE=2 SV=1 - [Q5SVM6_HUMAN]
Q9Y3A4	Ribosomal RNA-processing protein 7 homolog A OS=Homo sapiens GN=RRP7A PE=1 SV=2 - [RRP7A_HUMAN]
Q9P2E9	Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4 - [RRBP1_HUMAN]
Q5EBL4	RILP-like protein 1 OS=Homo sapiens GN=RILPL1 PE=1 SV=1 - [RIPL1_HUMAN]
Q9NXI6	RING finger protein 186 OS=Homo sapiens GN=RNF186 PE=2 SV=1 - [RN186_HUMAN]
Q63HN8	RING finger protein 213 OS=Homo sapiens GN=RNF213 PE=1 SV=2 - [RN213_HUMAN]
Q5W0B1	RING finger protein 219 OS=Homo sapiens GN=RNF219 PE=1 SV=1 - [RN219_HUMAN]
O00442	RNA 3'-terminal phosphate cyclase OS=Homo sapiens GN=RTCD1 PE=1 SV=1 - [RTC1_HUMAN]
Q9Y5S9	RNA-binding protein 8A OS=Homo sapiens GN=RBM8A PE=1 SV=1 - [RBM8A_HUMAN]
Q96T51	RUN and FYVE domain-containing protein 1 OS=Homo sapiens GN=RUFY1 PE=1 SV=2 - [RUFY1_HUMAN]
Q9Y265	RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1 - [RUVB1_HUMAN]
Q9Y230	RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3 - [RUVB2_HUMAN]
P31153	S-adenosylmethionine synthase isoform type-2 OS=Homo sapiens GN=MAT2A PE=1 SV=1 - [METK2_HUMAN]
P10768	S-formylglutathione hydrolase OS=Homo sapiens GN=ESD PE=1 SV=2 - [ESTD_HUMAN]
Q13126	S-methyl-5'-thioadenosine phosphorylase OS=Homo sapiens GN=MTAP PE=1 SV=2 - [MTAP_HUMAN]
Q9UDX3	SEC14-like protein 4 OS=Homo sapiens GN=SEC14L4 PE=2 SV=1 - [S14L4_HUMAN]
Q13228	Selenium-binding protein 1 OS=Homo sapiens GN=SELENBP1 PE=1 SV=2 - [SBP1_HUMAN]
Q96T21	Selenocysteine insertion sequence-binding protein 2 OS=Homo sapiens GN=SECISBP2 PE=1 SV=2 - [SEBP2_HUMAN]
Q9UHD8	Septin-9 OS=Homo sapiens GN=SEPT9 PE=1 SV=2 - [SEPT9_HUMAN]
Q96GA7	Serine dehydratase-like OS=Homo sapiens GN=SDSL PE=1 SV=1 - [SDSL_HUMAN]
O43464	Serine protease HTRA2, mitochondrial OS=Homo sapiens GN=HTRA2 PE=1 SV=2 - [HTRA2_HUMAN]
Q07955	Serine/arginine-rich splicing factor 1 OS=Homo sapiens GN=SRSF1 PE=1 SV=2 - [SRSF1_HUMAN]
Q08170	Serine/arginine-rich splicing factor 4 OS=Homo sapiens GN=SRSF4 PE=1 SV=2 - [SRSF4_HUMAN]
Q13242	Serine/arginine-rich splicing factor 9 OS=Homo sapiens GN=SRSF9 PE=1 SV=1 - [SRSF9_HUMAN]
Q9Y5S2	Serine/threonine-protein kinase MRCK beta OS=Homo sapiens GN=CDC42BPB PE=1 SV=2 - [MRCKB_HUMAN]
O95747	Serine/threonine-protein kinase OSR1 OS=Homo sapiens GN=OSXR1 PE=1 SV=1 - [OSXR1_HUMAN]
Q9BRS2	Serine/threonine-protein kinase RIO1 OS=Homo sapiens GN=RIOK1 PE=1 SV=2 - [RIOK1_HUMAN]
Q15173	Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit beta isoform OS=Homo sapiens GN=PPP2R5B PE=1 SV=1 - [2A5B_HUMAN]
P30153	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Homo sapiens GN=PPP2R1A PE=1 SV=4 - [2AAA_HUMAN]
P30154	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A beta isoform OS=Homo sapiens GN=PPP2R1B PE=1 SV=3 - [2AAB_HUMAN]
P67775	Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform OS=Homo sapiens GN=PPP2CA PE=1 SV=1 - [PP2AA_HUMAN]
O15084	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A OS=Homo sapiens GN=ANKRD28 PE=1 SV=5 - [ANR28_HUMAN]
E9PMD7	Serine/threonine-protein phosphatase OS=Homo sapiens GN=PPP1CA PE=3 SV=1 - [E9PMD7_HUMAN]
B4DDZ8	Serine/threonine-protein phosphatase OS=Homo sapiens GN=PPP5C PE=2 SV=1 - [B4DDZ8_HUMAN]

O14830	Serine/threonine-protein phosphatase with EF-hands 2 OS=Homo sapiens GN=PPEF2 PE=1 SV=2 - [PPE2_HUMAN]
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3 - [TRFE_HUMAN]
Q9BXP5	Serrate RNA effector molecule homolog OS=Homo sapiens GN=SRRT PE=1 SV=1 - [SRRT_HUMAN]
O95810	Serum deprivation-response protein OS=Homo sapiens GN=SDPR PE=1 SV=3 - [SDPR_HUMAN]
P49591	Seryl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=SARS PE=1 SV=3 - [SYSC_HUMAN]
Q9C0A6	SET domain-containing protein 5 OS=Homo sapiens GN=SETD5 PE=1 SV=2 - [SETD5_HUMAN]
B2REB8	SET nuclear oncogene OS=Homo sapiens GN=SET PE=3 SV=1 - [B2REB8_HUMAN]
Q8N228	Sex comb on midleg-like protein 4 OS=Homo sapiens GN=SCML4 PE=2 SV=2 - [SCML4_HUMAN]
Q5T123	SH3 domain binding glutamic acid-rich protein like 3 OS=Homo sapiens GN=SH3BGRL3 PE=2 SV=1 - [Q5T123_HUMAN]
O75368	SH3 domain-binding glutamic acid-rich-like protein OS=Homo sapiens GN=SH3BGRL PE=1 SV=1 - [SH3L1_HUMAN]
Q9BQI5	SH3-containing GRB2-like protein 3-interacting protein 1 OS=Homo sapiens GN=SGIP1 PE=1 SV=2 - [SGIP1_HUMAN]
B7ZC39	SH3-domain GRB2-like endophilin B2 OS=Homo sapiens GN=SH3GLB2 PE=4 SV=1 - [B7ZC39_HUMAN]
Q9NR45	Sialic acid synthase OS=Homo sapiens GN=NANS PE=1 SV=2 - [SIAS_HUMAN]
Q6P4A7	Sideroflexin-4 OS=Homo sapiens GN=SFXN4 PE=1 SV=1 - [SFXN4_HUMAN]
P37108	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2 - [SRP14_HUMAN]
O76094	Signal recognition particle 72 kDa protein OS=Homo sapiens GN=SRP72 PE=1 SV=3 - [SRP72_HUMAN]
P49458	Signal recognition particle 9 kDa protein OS=Homo sapiens GN=SRP9 PE=1 SV=2 - [SRP09_HUMAN]
Q1KLZ6	Signal transducer and activator of transcription 5A variant delta5 OS=Homo sapiens GN=STAT5A PE=2 SV=1 - [Q1KLZ6_HUMAN]
O00241	Signal-regulatory protein beta-1 OS=Homo sapiens GN=SIRPB1 PE=1 SV=5 - [SIRB1_HUMAN]
Q8IW23	Similar to glutamate receptor, ionotropic, N-methyl D-aspartate 2C (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q8IW23_HUMAN]
O43805	Sjogren syndrome nuclear autoantigen 1 OS=Homo sapiens GN=SSNA1 PE=1 SV=2 - [SSNA1_HUMAN]
Q96G53	SLC27A4 protein OS=Homo sapiens GN=SLC27A4 PE=2 SV=1 - [Q96G53_HUMAN]
P62306	Small nuclear ribonucleoprotein F OS=Homo sapiens GN=SNRPF PE=1 SV=1 - [RUXF_HUMAN]
Q5SZX1	Smg-5 homolog, nonsense mediated mRNA decay factor (C. elegans) OS=Homo sapiens GN=SMG5 PE=2 SV=1 - [Q5SZX1_HUMAN]
A2RRQ7	SNRNP200 protein OS=Homo sapiens GN=SNRNP200 PE=2 SV=1 - [A2RRQ7_HUMAN]
C0H5W9	SNX6 protein OS=Homo sapiens GN=SNX6 PE=2 SV=1 - [C0H5W9_HUMAN]
P48066	Sodium- and chloride-dependent GABA transporter 3 OS=Homo sapiens GN=SLC6A11 PE=2 SV=1 - [S6A11_HUMAN]
P11166	Solute carrier family 2, facilitated glucose transporter member 1 OS=Homo sapiens GN=SLC2A1 PE=1 SV=2 - [GTR1_HUMAN]
A5X8Z9	Solute carrier family 24 member 5 OS=Homo sapiens GN=SLC24A5 PE=2 SV=1 - [A5X8Z9_HUMAN]
Q00796	Sorbitol dehydrogenase OS=Homo sapiens GN=SORD PE=1 SV=4 - [DHSO_HUMAN]
A6NKH4	Sorting nexin 1, isoform CRA_d OS=Homo sapiens GN=SNX1 PE=4 SV=2 - [A6NKH4_HUMAN]
O60493	Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3 - [SNX3_HUMAN]
Q9Y5X1	Sorting nexin-9 OS=Homo sapiens GN=SNX9 PE=1 SV=1 - [SNX9_HUMAN]
Q8N0X7	Spartin OS=Homo sapiens GN=SPG20 PE=1 SV=1 - [SPG20_HUMAN]
P02549	Spectrin alpha chain, erythrocyte OS=Homo sapiens GN=SPTA1 PE=1 SV=5 - [SPTA1_HUMAN]
O15020	Spectrin beta chain, brain 2 OS=Homo sapiens GN=SPTBN2 PE=1 SV=3 - [SPTN2_HUMAN]
P11277	Spectrin beta chain, erythrocyte OS=Homo sapiens GN=SPTB PE=1 SV=5 - [SPTB1_HUMAN]
Q9C093	Sperm flagellar protein 2 OS=Homo sapiens GN=SPEF2 PE=1 SV=2 - [SPEF2_HUMAN]
Q8NA61	Spermatid-associated protein OS=Homo sapiens GN=SPERT PE=2 SV=1 - [SPERT_HUMAN]
Q9P0W8	Spermatogenesis-associated protein 7 OS=Homo sapiens GN=SPATA7 PE=2 SV=3 - [SPAT7_HUMAN]
P52788	Spermine synthase OS=Homo sapiens GN=SMS PE=1 SV=2 - [SPSY_HUMAN]
P26368	Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2 PE=1 SV=4 - [U2AF2_HUMAN]
B3KUY1	Splicing factor, arginine/serine-rich 2, isoform CRA_d OS=Homo sapiens GN=SFRS2 PE=2 SV=1 - [B3KUY1_HUMAN]

Q9P2P6	StAR-related lipid transfer protein 9 OS=Homo sapiens GN=STARD9 PE=2 SV=2 - [STAR9_HUMAN]
P16949	Stathmin OS=Homo sapiens GN=STMN1 PE=1 SV=3 - [STMN1_HUMAN]
B7Z5G4	Stathmin OS=Homo sapiens GN=STMN3 PE=2 SV=1 - [B7Z5G4_HUMAN]
Q9H2G2	STE20-like serine/threonine-protein kinase OS=Homo sapiens GN=SLK PE=1 SV=1 - [SLK_HUMAN]
Q8IZD0	Sterile alpha motif domain-containing protein 14 OS=Homo sapiens GN=SAMD14 PE=2 SV=2 - [SAM14_HUMAN]
P31948	Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=1 SV=1 - [STIP1_HUMAN]
Q14683	Structural maintenance of chromosomes protein 1A OS=Homo sapiens GN=SMC1A PE=1 SV=2 - [SMC1A_HUMAN]
O95347	Structural maintenance of chromosomes protein 2 OS=Homo sapiens GN=SMC2 PE=1 SV=2 - [SMC2_HUMAN]
Q9BYN0	Sulfiredoxin-1 OS=Homo sapiens GN=SRXN1 PE=1 SV=2 - [SRXN1_HUMAN]
P00441	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2 - [SODC_HUMAN]
O60284	Suppression of tumorigenicity 18 protein OS=Homo sapiens GN=ST18 PE=1 SV=1 - [ST18_HUMAN]
Q9Y2Z0	Suppressor of G2 allele of SKP1 homolog OS=Homo sapiens GN=SUGT1 PE=1 SV=3 - [SUGT1_HUMAN]
O75683	Surfeit locus protein 6 OS=Homo sapiens GN=SURF6 PE=1 SV=3 - [SURF6_HUMAN]
Q59GV3	SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 isoform b variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q59GV3_HUMAN]
O15498	Synaptobrevin homolog YKT6 OS=Homo sapiens GN=YKT6 PE=1 SV=1 - [YKT6_HUMAN]
Q96C24	Synaptotagmin-like protein 4 OS=Homo sapiens GN=SYTL4 PE=1 SV=2 - [SYTL4_HUMAN]
Q9NPQ8	Synembryn-A OS=Homo sapiens GN=RIC8A PE=1 SV=3 - [RIC8A_HUMAN]
Q96SF7	T-box transcription factor TBX15 OS=Homo sapiens GN=TBX15 PE=2 SV=2 - [TBX15_HUMAN]
P17987	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1 - [TCPA_HUMAN]
P78371	T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4 - [TCPB_HUMAN]
P50991	T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4 - [TCPD_HUMAN]
P50990	T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4 - [TCPQ_HUMAN]
P40227	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3 - [TCPZ_HUMAN]
Q92526	T-complex protein 1 subunit zeta-2 OS=Homo sapiens GN=CCT6B PE=1 SV=5 - [TCPW_HUMAN]
Q8N4U5	T-complex protein 11-like protein 2 OS=Homo sapiens GN=TCP11L2 PE=2 SV=1 - [T11L2_HUMAN]
Q5TCU6	Talin 1 OS=Homo sapiens GN=TLN1 PE=2 SV=1 - [Q5TCU6_HUMAN]
Q9Y4G6	Talin-2 OS=Homo sapiens GN=TLN2 PE=1 SV=4 - [TLN2_HUMAN]
B1AKP7	TAR DNA binding protein OS=Homo sapiens GN=TARDBP PE=4 SV=1 - [B1AKP7_HUMAN]
P59537	Taste receptor type 2 member 43 OS=Homo sapiens GN=TAS2R43 PE=2 SV=1 - [T2R43_HUMAN]
Q8TBP0	TBC1 domain family member 16 OS=Homo sapiens GN=TBC1D16 PE=1 SV=1 - [TBC16_HUMAN]
O95759	TBC1 domain family member 8 OS=Homo sapiens GN=TBC1D8 PE=1 SV=3 - [TBCD8_HUMAN]
A8K8E1	TBC1 domain family, member 15, isoform CRA_d OS=Homo sapiens GN=TBC1D15 PE=2 SV=1 - [A8K8E1_HUMAN]
Q9NYB0	Telomeric repeat-binding factor 2-interacting protein 1 OS=Homo sapiens GN=TERF2IP PE=1 SV=1 - [TE2IP_HUMAN]
Q59G71	Tensin variant (Fragment) OS=Homo sapiens PE=2 SV=1 - [Q59G71_HUMAN]
Q99614	Tetratricopeptide repeat protein 1 OS=Homo sapiens GN=TTC1 PE=1 SV=1 - [TTC1_HUMAN]
P52888	Thimet oligopeptidase OS=Homo sapiens GN=THOP1 PE=1 SV=2 - [THOP1_HUMAN]
Q9BRA2	Thioredoxin domain-containing protein 17 OS=Homo sapiens GN=TXNDC17 PE=1 SV=1 - [TXD17_HUMAN]
B1ALW1	Thioredoxin OS=Homo sapiens GN=TXN PE=4 SV=1 - [B1ALW1_HUMAN]
O43396	Thioredoxin-like protein 1 OS=Homo sapiens GN=TXNL1 PE=1 SV=3 - [TXNL1_HUMAN]
P26639	Threonyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=TARS PE=1 SV=3 - [SYTC_HUMAN]
P23919	Thymidylate kinase OS=Homo sapiens GN=DTYMK PE=1 SV=4 - [KTHY_HUMAN]
P0CG34	Thymosin beta-15A OS=Homo sapiens GN=TMSB15A PE=1 SV=1 - [TB15A_HUMAN]
A8MW06	Thymosin beta-4-like protein 3 OS=Homo sapiens GN=TMSL3 PE=2 SV=1 - [TMSL3_HUMAN]

O75663	TIP41-like protein OS=Homo sapiens GN=TIPRL PE=1 SV=2 - [TIPRL_HUMAN]
Q9NYK1	Toll-like receptor 7 OS=Homo sapiens GN=TLR7 PE=1 SV=1 - [TLR7_HUMAN]
Q6LBI8	Topoisomerase IIb (Fragment) OS=Homo sapiens GN=topIIb PE=2 SV=1 - [Q6LBI8_HUMAN]
O14804	Trace amine-associated receptor 5 OS=Homo sapiens GN=TAAR5 PE=2 SV=2 - [TAAR5_HUMAN]
A6NDN0	Trafficking protein particle complex 3, isoform CRA_a OS=Homo sapiens GN=TRAPPC3 PE=4 SV=1 - [A6NDN0_HUMAN]
P37837	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2 - [TALDO_HUMAN]
P23193	Transcription elongation factor A protein 1 OS=Homo sapiens GN=TCEA1 PE=1 SV=2 - [TCEA1_HUMAN]
B8ZZU8	Transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B), isoform CRA_b OS=Homo sapiens GN=TCEB2 PE=4 SV=1 - [B8ZZU8_HUMAN]
Q5VWG9	Transcription initiation factor TFIID subunit 3 OS=Homo sapiens GN=TAF3 PE=1 SV=1 - [TAF3_HUMAN]
Q13263	Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5 - [TIF1B_HUMAN]
Q00577	Transcriptional activator protein Pur-alpha OS=Homo sapiens GN=PURA PE=1 SV=2 - [PURA_HUMAN]
P02786	Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=2 - [TFR1_HUMAN]
B9ZVW9	Transforming growth factor, beta-induced, 68kDa, isoform CRA_a OS=Homo sapiens GN=TGFBI PE=4 SV=1 - [B9ZVW9_HUMAN]
P61586	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1 - [RHOA_HUMAN]
P37802	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3 - [TAGL2_HUMAN]
P55072	Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4 - [TERA_HUMAN]
P13693	Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1 - [TCTP_HUMAN]
Q99598	Translin-associated protein X OS=Homo sapiens GN=TSNAX PE=1 SV=1 - [TSNAX_HUMAN]
Q6IN77	Transportin 2 OS=Homo sapiens GN=TNPO2 PE=2 SV=1 - [Q6IN77_HUMAN]
Q92973	Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2 - [TNPO1_HUMAN]
P22102	Trifunctional purine biosynthetic protein adenosine-3 OS=Homo sapiens GN=GART PE=1 SV=1 - [PUR2_HUMAN]
P60174	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=2 - [TPIS_HUMAN]
Q5SRJ5	Tripartite motif-containing 10 OS=Homo sapiens GN=TRIM10 PE=2 SV=1 - [Q5SRJ5_HUMAN]
Q8NG06	Tripartite motif-containing protein 58 OS=Homo sapiens GN=TRIM58 PE=2 SV=2 - [TRI58_HUMAN]
P29144	Tripeptidyl-peptidase 2 OS=Homo sapiens GN=TPP2 PE=1 SV=4 - [TPP2_HUMAN]
O95620	tRNA-dihydrouridine synthase 4-like OS=Homo sapiens GN=DUS4L PE=2 SV=2 - [DUS4L_HUMAN]
D9YZV5	Tropomyosin 1 (Alpha) isoform 4 OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [D9YZV5_HUMAN]
D9YZV7	Tropomyosin 1 (Alpha) isoform 6 OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [D9YZV7_HUMAN]
Q5VU58	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU58_HUMAN]
Q5VU59	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU59_HUMAN]
Q5VU61	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU61_HUMAN]
Q5VU66	Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1 - [Q5VU66_HUMAN]
P06753	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=1 SV=1 - [TPM3_HUMAN]
Q5LJ98	TROVE domain family, member 2 OS=Homo sapiens GN=TROVE2 PE=2 SV=1 - [Q5LJ98_HUMAN]
P23381	Tryptophanyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=WARS PE=1 SV=2 - [SYWC_HUMAN]
O75157	TSC22 domain family protein 2 OS=Homo sapiens GN=TSC22D2 PE=1 SV=3 - [T22D2_HUMAN]
Q9BQE3	Tubulin alpha-1C chain OS=Homo sapiens GN=TUBA1C PE=1 SV=1 - [TBA1C_HUMAN]
P68371	Tubulin beta-2C chain OS=Homo sapiens GN=TUBB2C PE=1 SV=1 - [TBB2C_HUMAN]
B1AH89	Tubulin tyrosine ligase-like family, member 12 OS=Homo sapiens GN=TTLL12 PE=4 SV=1 - [B1AH89_HUMAN]
Q5QJ74	Tubulin-specific chaperone cofactor E-like protein OS=Homo sapiens GN=TBCEL PE=2 SV=2 - [TBCEL_HUMAN]
Q15813	Tubulin-specific chaperone E OS=Homo sapiens GN=TBCE PE=1 SV=1 - [TBCE_HUMAN]
A8MUB1	Tubulin, alpha 1 (Testis specific), isoform CRA_a OS=Homo sapiens GN=TUBA4A PE=2 SV=1 - [A8MUB1_HUMAN]
Q9UIG0	Tyrosine-protein kinase BAZ1B OS=Homo sapiens GN=BAZ1B PE=1 SV=2 - [BAZ1B_HUMAN]

Q06124	Tyrosine-protein phosphatase non-receptor type 11 OS=Homo sapiens GN=PTPN11 PE=1 SV=2 - [PTN11_HUMAN]
P54577	Tyrosyl-tRNA synthetase, cytoplasmic OS=Homo sapiens GN=YARS PE=1 SV=4 - [SYYC_HUMAN]
Q15696	U2 small nuclear ribonucleoprotein auxiliary factor 35 kDa subunit-related protein 2 OS=Homo sapiens GN=ZRSR2 PE=1 SV=2 - [U2AFM_HUMAN]
O43818	U3 small nucleolar RNA-interacting protein 2 OS=Homo sapiens GN=RRP9 PE=1 SV=1 - [U3IP2_HUMAN]
O15116	U6 snRNA-associated Sm-like protein LSM1 OS=Homo sapiens GN=LSM1 PE=1 SV=1 - [LSM1_HUMAN]
Q9Y333	U6 snRNA-associated Sm-like protein LSM2 OS=Homo sapiens GN=LSM2 PE=1 SV=1 - [LSM2_HUMAN]
Q9Y4Z0	U6 snRNA-associated Sm-like protein LSM4 OS=Homo sapiens GN=LSM4 PE=1 SV=1 - [LSM4_HUMAN]
P62312	U6 snRNA-associated Sm-like protein LSM6 OS=Homo sapiens GN=LSM6 PE=1 SV=1 - [LSM6_HUMAN]
Q9UK45	U6 snRNA-associated Sm-like protein LSM7 OS=Homo sapiens GN=LSM7 PE=1 SV=1 - [LSM7_HUMAN]
Q5TBK7	Ubiquitin carboxyl-terminal esterase L3 (Ubiquitin thioesterase) (Fragment) OS=Homo sapiens GN=UCHL3 PE=2 SV=1 - [Q5TBK7_HUMAN]
A6NCW0	Ubiquitin carboxyl-terminal hydrolase 17-like protein 3 OS=Homo sapiens GN=USP17L3 PE=3 SV=1 - [U17L3_HUMAN]
P45974	Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5 PE=1 SV=2 - [UBP5_HUMAN]
A6NJA2	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP14 PE=3 SV=1 - [A6NJA2_HUMAN]
E9PCQ3	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP15 PE=3 SV=2 - [E9PCQ3_HUMAN]
B7WPF4	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP24 PE=3 SV=2 - [B7WPF4_HUMAN]
E9PM46	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP47 PE=3 SV=1 - [E9PM46_HUMAN]
B4DKA8	Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP8 PE=2 SV=1 - [B4DKA8_HUMAN]
D6RAW0	Ubiquitin carrier protein OS=Homo sapiens GN=UBE2D3 PE=3 SV=1 - [D6RAW0_HUMAN]
C9JZY6	Ubiquitin carrier protein OS=Homo sapiens GN=UBE2H PE=3 SV=1 - [C9JZY6_HUMAN]
F5GX52	Ubiquitin carrier protein OS=Homo sapiens PE=3 SV=1 - [F5GX52_HUMAN]
Q14139	Ubiquitin conjugation factor E4 A OS=Homo sapiens GN=UBE4A PE=1 SV=2 - [UBE4A_HUMAN]
Q5VVQ6	Ubiquitin thioesterase OTU1 OS=Homo sapiens GN=YOD1 PE=1 SV=1 - [OTU1_HUMAN]
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2 - [RS27A_HUMAN]
Q9BSL1	Ubiquitin-associated domain-containing protein 1 OS=Homo sapiens GN=UBAC1 PE=1 SV=1 - [UBAC1_HUMAN]
P61088	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1 - [UBE2N_HUMAN]
Q9C0C9	Ubiquitin-conjugating enzyme E2 O OS=Homo sapiens GN=UBE2O PE=1 SV=3 - [UBE2O_HUMAN]
P49427	Ubiquitin-conjugating enzyme E2 R1 OS=Homo sapiens GN=CDC34 PE=1 SV=2 - [UB2R1_HUMAN]
Q13404	Ubiquitin-conjugating enzyme E2 variant 1 OS=Homo sapiens GN=UBE2V1 PE=1 SV=2 - [UB2V1_HUMAN]
Q15819	Ubiquitin-conjugating enzyme E2 variant 2 OS=Homo sapiens GN=UBE2V2 PE=1 SV=4 - [UB2V2_HUMAN]
P22314	Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3 - [UBA1_HUMAN]
A0AVT1	Ubiquitin-like modifier-activating enzyme 6 OS=Homo sapiens GN=UBA6 PE=1 SV=1 - [UBA6_HUMAN]
P05161	Ubiquitin-like protein ISG15 OS=Homo sapiens GN=ISG15 PE=1 SV=5 - [ISG15_HUMAN]
Q9NT62	Ubiquitin-like-conjugating enzyme ATG3 OS=Homo sapiens GN=ATG3 PE=1 SV=1 - [ATG3_HUMAN]
Q5QPP1	UDP-galactose-4-epimerase (Fragment) OS=Homo sapiens GN=GALE PE=2 SV=1 - [Q5QPP1_HUMAN]
Q9NYU2	UDP-glucose:glycoprotein glucosyltransferase 1 OS=Homo sapiens GN=UGGT1 PE=1 SV=3 - [UGGG1_HUMAN]
Q9Y4X1	UDP-glucuronosyltransferase 2A1 OS=Homo sapiens GN=UGT2A1 PE=1 SV=2 - [UD2A1_HUMAN]
O15294	UDP-N-acetylglucosamine--peptide N-acetylglucosaminyltransferase 110 kDa subunit OS=Homo sapiens GN=OGT PE=1 SV=3 - [OGT1_HUMAN]
P30085	UMP-CMP kinase OS=Homo sapiens GN=CMPK1 PE=1 SV=3 - [KCY_HUMAN]
Q3MIX3	Uncharacterized aarF domain-containing protein kinase 5 OS=Homo sapiens GN=ADCK5 PE=2 SV=2 - [ADCK5_HUMAN]
B4DGV4	Uncharacterized protein (Fragment) OS=Homo sapiens GN=RYSR2 PE=2 SV=1 - [B4DGV4_HUMAN]
Q7Z3E2	Uncharacterized protein C10orf118 OS=Homo sapiens GN=C10orf118 PE=2 SV=2 - [CJ118_HUMAN]
Q96LP6	Uncharacterized protein C12orf42 OS=Homo sapiens GN=C12orf42 PE=2 SV=2 - [CL042_HUMAN]
Q4W4Y0	Uncharacterized protein C14orf28 OS=Homo sapiens GN=C14orf28 PE=2 SV=1 - [CN028_HUMAN]

Q9H972	Uncharacterized protein C14orf93 OS=Homo sapiens GN=C14orf93 PE=2 SV=1 - [CN093_HUMAN]
Q5RHP9	Uncharacterized protein C1orf173 OS=Homo sapiens GN=C1orf173 PE=2 SV=1 - [CA173_HUMAN]
Q6ICL3	Uncharacterized protein C22orf25 OS=Homo sapiens GN=C22orf25 PE=2 SV=1 - [CV025_HUMAN]
Q8NC60	Uncharacterized protein C4orf14 OS=Homo sapiens GN=C4orf14 PE=1 SV=2 - [CD014_HUMAN]
Q9P0K9	Uncharacterized protein C9orf4 OS=Homo sapiens GN=C9orf4 PE=2 SV=2 - [CI004_HUMAN]
Q8IXQ3	Uncharacterized protein C9orf40 OS=Homo sapiens GN=C9orf40 PE=1 SV=1 - [CI040_HUMAN]
Q9UF83	Uncharacterized protein DKFZp434B061 OS=Homo sapiens PE=2 SV=2 - [YM012_HUMAN]
Q6ZSR9	Uncharacterized protein FLJ45252 OS=Homo sapiens PE=1 SV=2 - [YJ005_HUMAN]
Q6ZRH9	Uncharacterized protein FLJ46347 OS=Homo sapiens PE=2 SV=1 - [YB011_HUMAN]
O94964	Uncharacterized protein KIAA0889 OS=Homo sapiens GN=KIAA0889 PE=1 SV=2 - [K0889_HUMAN]
Q8IXQ4	Uncharacterized protein KIAA1704 OS=Homo sapiens GN=KIAA1704 PE=1 SV=1 - [K1704_HUMAN]
Q9C0B2	Uncharacterized protein KIAA1751 OS=Homo sapiens GN=KIAA1751 PE=2 SV=2 - [K1751_HUMAN]
Q0VF49	Uncharacterized protein KIAA2012 OS=Homo sapiens GN=KIAA2012 PE=2 SV=1 - [K2012_HUMAN]
C9JGD3	Uncharacterized protein OS=Homo sapiens GN=A1CF PE=4 SV=2 - [C9JGD3_HUMAN]
E9PG46	Uncharacterized protein OS=Homo sapiens GN=AAK1 PE=4 SV=1 - [E9PG46_HUMAN]
F5GYC1	Uncharacterized protein OS=Homo sapiens GN=ABCD3 PE=3 SV=1 - [F5GYC1_HUMAN]
D6R9I9	Uncharacterized protein OS=Homo sapiens GN=ABCE1 PE=4 SV=1 - [D6R9I9_HUMAN]
B4DNR3	Uncharacterized protein OS=Homo sapiens GN=ABHD14B PE=2 SV=1 - [B4DNR3_HUMAN]
B4DUX0	Uncharacterized protein OS=Homo sapiens GN=ACOT7 PE=2 SV=1 - [B4DUX0_HUMAN]
E9PIS4	Uncharacterized protein OS=Homo sapiens GN=ACOT8 PE=4 SV=1 - [E9PIS4_HUMAN]
F5H4B8	Uncharacterized protein OS=Homo sapiens GN=ACSM2A PE=4 SV=1 - [F5H4B8_HUMAN]
F5H0N0	Uncharacterized protein OS=Homo sapiens GN=ACTG1 PE=3 SV=1 - [F5H0N0_HUMAN]
B4DXP9	Uncharacterized protein OS=Homo sapiens GN=ACTR1A PE=2 SV=1 - [B4DXP9_HUMAN]
B4DTI0	Uncharacterized protein OS=Homo sapiens GN=ACTR3 PE=2 SV=1 - [B4DTI0_HUMAN]
F5GXW0	Uncharacterized protein OS=Homo sapiens GN=ADA PE=4 SV=1 - [F5GXW0_HUMAN]
B7ZAL1	Uncharacterized protein OS=Homo sapiens GN=ADH6 PE=2 SV=1 - [B7ZAL1_HUMAN]
B7Z783	Uncharacterized protein OS=Homo sapiens GN=ADK PE=2 SV=1 - [B7Z783_HUMAN]
E9PEC0	Uncharacterized protein OS=Homo sapiens GN=ADRM1 PE=4 SV=2 - [E9PEC0_HUMAN]
C9J2I0	Uncharacterized protein OS=Homo sapiens GN=AGFG1 PE=4 SV=1 - [C9J2I0_HUMAN]
C9JSL3	Uncharacterized protein OS=Homo sapiens GN=AKR7A2 PE=4 SV=2 - [C9JSL3_HUMAN]
F5H4B6	Uncharacterized protein OS=Homo sapiens GN=ALDH16A1 PE=4 SV=1 - [F5H4B6_HUMAN]
E9PBX3	Uncharacterized protein OS=Homo sapiens GN=ALDH1L1 PE=3 SV=1 - [E9PBX3_HUMAN]
E9PGG4	Uncharacterized protein OS=Homo sapiens GN=ALS2CR11 PE=4 SV=1 - [E9PGG4_HUMAN]
A6NP16	Uncharacterized protein OS=Homo sapiens GN=AMOT PE=4 SV=1 - [A6NP16_HUMAN]
E9PKC5	Uncharacterized protein OS=Homo sapiens GN=AMPD3 PE=4 SV=1 - [E9PKC5_HUMAN]
E7EVE3	Uncharacterized protein OS=Homo sapiens GN=ANK1 PE=4 SV=1 - [E7EVE3_HUMAN]
D6RIY9	Uncharacterized protein OS=Homo sapiens GN=ANK2 PE=4 SV=1 - [D6RIY9_HUMAN]
E7EWS1	Uncharacterized protein OS=Homo sapiens GN=ANKAR PE=4 SV=1 - [E7EWS1_HUMAN]
F5H754	Uncharacterized protein OS=Homo sapiens GN=ANKFY1 PE=4 SV=1 - [F5H754_HUMAN]
B3KUC3	Uncharacterized protein OS=Homo sapiens GN=ANKRD6 PE=2 SV=1 - [B3KUC3_HUMAN]
F2Z3H3	Uncharacterized protein OS=Homo sapiens GN=ANP32A PE=4 SV=2 - [F2Z3H3_HUMAN]
B4DT77	Uncharacterized protein OS=Homo sapiens GN=ANXA7 PE=2 SV=1 - [B4DT77_HUMAN]
F5H0H0	Uncharacterized protein OS=Homo sapiens GN=AP2M1 PE=4 SV=1 - [F5H0H0_HUMAN]

F5H093	Uncharacterized protein OS=Homo sapiens GN=AP3B1 PE=4 SV=1 - [F5H093_HUMAN]
E7EMM2	Uncharacterized protein OS=Homo sapiens GN=AP3D1 PE=4 SV=1 - [E7EMM2_HUMAN]
E9PC21	Uncharacterized protein OS=Homo sapiens GN=AQP1 PE=3 SV=1 - [E9PC21_HUMAN]
<i>B7ZB63</i>	<i>Uncharacterized protein OS=Homo sapiens GN=ARF3 PE=2 SV=1 - [B7ZB63_HUMAN]</i>
E9PNR6	Uncharacterized protein OS=Homo sapiens GN=ARHGAP1 PE=4 SV=1 - [E9PNR6_HUMAN]
C9JVJ3	Uncharacterized protein OS=Homo sapiens GN=ARHGAP42 PE=4 SV=1 - [C9JVJ3_HUMAN]
F5H6Q0	Uncharacterized protein OS=Homo sapiens GN=ARHGDI1 PE=4 SV=1 - [F5H6Q0_HUMAN]
E7EWX4	Uncharacterized protein OS=Homo sapiens GN=ARPC4 PE=4 SV=1 - [E7EWX4_HUMAN]
E9PE48	Uncharacterized protein OS=Homo sapiens GN=ASL PE=3 SV=1 - [E9PE48_HUMAN]
C9JAL9	Uncharacterized protein OS=Homo sapiens GN=ASPSR1 PE=4 SV=1 - [C9JAL9_HUMAN]
E9PBU3	Uncharacterized protein OS=Homo sapiens GN=ATIC PE=4 SV=1 - [E9PBU3_HUMAN]
E5RGN3	Uncharacterized protein OS=Homo sapiens GN=ATOX1 PE=4 SV=1 - [E5RGN3_HUMAN]
D6R9H7	Uncharacterized protein OS=Homo sapiens GN=ATP5G1 PE=3 SV=1 - [D6R9H7_HUMAN]
C9J8H9	Uncharacterized protein OS=Homo sapiens GN=ATP5J2 PE=4 SV=1 - [C9J8H9_HUMAN]
B7Z1R5	Uncharacterized protein OS=Homo sapiens GN=ATP6V1A PE=2 SV=1 - [B7Z1R5_HUMAN]
E7EV59	Uncharacterized protein OS=Homo sapiens GN=ATP6V1C1 PE=4 SV=1 - [E7EV59_HUMAN]
A8MUN4	Uncharacterized protein OS=Homo sapiens GN=ATP6V1E1 PE=2 SV=1 - [A8MUN4_HUMAN]
E9PGE6	Uncharacterized protein OS=Homo sapiens GN=ATP6V1H PE=4 SV=1 - [E9PGE6_HUMAN]
B5MEB7	Uncharacterized protein OS=Homo sapiens GN=ATRIP PE=4 SV=2 - [B5MEB7_HUMAN]
C9J9Y9	Uncharacterized protein OS=Homo sapiens GN=BAG1 PE=4 SV=2 - [C9J9Y9_HUMAN]
F5H239	Uncharacterized protein OS=Homo sapiens GN=BAZ1A PE=4 SV=1 - [F5H239_HUMAN]
E7EX43	Uncharacterized protein OS=Homo sapiens GN=BBS9 PE=4 SV=1 - [E7EX43_HUMAN]
E9PK09	Uncharacterized protein OS=Homo sapiens GN=BCLAF1 PE=4 SV=1 - [E9PK09_HUMAN]
F5H7R6	Uncharacterized protein OS=Homo sapiens GN=BLM PE=4 SV=1 - [F5H7R6_HUMAN]
B4DUS9	Uncharacterized protein OS=Homo sapiens GN=BPNT1 PE=2 SV=1 - [B4DUS9_HUMAN]
C9IZW4	Uncharacterized protein OS=Homo sapiens GN=BRCA1 PE=4 SV=1 - [C9IZW4_HUMAN]
E9PPN5	Uncharacterized protein OS=Homo sapiens GN=BSCL2 PE=4 SV=1 - [E9PPN5_HUMAN]
C9JB13	Uncharacterized protein OS=Homo sapiens GN=BSDC1 PE=4 SV=1 - [C9JB13_HUMAN]
E9PL10	Uncharacterized protein OS=Homo sapiens GN=BTF3L4 PE=4 SV=1 - [E9PL10_HUMAN]
E9PAX7	Uncharacterized protein OS=Homo sapiens GN=C11orf49 PE=4 SV=1 - [E9PAX7_HUMAN]
F5H641	Uncharacterized protein OS=Homo sapiens GN=C12orf10 PE=4 SV=1 - [F5H641_HUMAN]
E9PBZ7	Uncharacterized protein OS=Homo sapiens GN=C13orf38 PE=4 SV=1 - [E9PBZ7_HUMAN]
B5MC47	Uncharacterized protein OS=Homo sapiens GN=C14orf135 PE=4 SV=1 - [B5MC47_HUMAN]
C9IYA4	Uncharacterized protein OS=Homo sapiens GN=C14orf80 PE=4 SV=1 - [C9IYA4_HUMAN]
B8ZZ55	Uncharacterized protein OS=Homo sapiens GN=C2orf70 PE=4 SV=1 - [B8ZZ55_HUMAN]
B4DIY7	Uncharacterized protein OS=Homo sapiens GN=C3orf31 PE=2 SV=1 - [B4DIY7_HUMAN]
C9JF90	Uncharacterized protein OS=Homo sapiens GN=C7orf25 PE=4 SV=1 - [C9JF90_HUMAN]
C9JS17	Uncharacterized protein OS=Homo sapiens GN=C7orf69 PE=4 SV=1 - [C9JS17_HUMAN]
E7ER51	Uncharacterized protein OS=Homo sapiens GN=CA3 PE=4 SV=2 - [E7ER51_HUMAN]
B4DE01	Uncharacterized protein OS=Homo sapiens GN=CAPZA2 PE=2 SV=1 - [B4DE01_HUMAN]
C9JXR7	Uncharacterized protein OS=Homo sapiens GN=CASP3 PE=4 SV=2 - [C9JXR7_HUMAN]
D6RBM3	Uncharacterized protein OS=Homo sapiens GN=CASP6 PE=4 SV=1 - [D6RBM3_HUMAN]
E7EVY3	Uncharacterized protein OS=Homo sapiens GN=CAST PE=4 SV=1 - [E7EVY3_HUMAN]

E9PDE4	Uncharacterized protein OS=Homo sapiens GN=CAST PE=4 SV=1 - [E9PDE4_HUMAN]
B8ZZI4	Uncharacterized protein OS=Homo sapiens GN=CCDC150 PE=4 SV=1 - [B8ZZI4_HUMAN]
C9JGU1	Uncharacterized protein OS=Homo sapiens GN=CCDC66 PE=4 SV=2 - [C9JGU1_HUMAN]
E9PAQ6	Uncharacterized protein OS=Homo sapiens GN=CCT3 PE=3 SV=1 - [E9PAQ6_HUMAN]
E7ENZ3	Uncharacterized protein OS=Homo sapiens GN=CCT5 PE=4 SV=1 - [E7ENZ3_HUMAN]
B4DTR2	Uncharacterized protein OS=Homo sapiens GN=CDH5 PE=2 SV=1 - [B4DTR2_HUMAN]
D6R973	Uncharacterized protein OS=Homo sapiens GN=CDV3 PE=4 SV=1 - [D6R973_HUMAN]
F5GYE8	Uncharacterized protein OS=Homo sapiens GN=CEP112 PE=4 SV=1 - [F5GYE8_HUMAN]
C9J710	Uncharacterized protein OS=Homo sapiens GN=CEP70 PE=4 SV=1 - [C9J710_HUMAN]
E9PP50	Uncharacterized protein OS=Homo sapiens GN=CFL1 PE=4 SV=1 - [E9PP50_HUMAN]
F5H2G1	Uncharacterized protein OS=Homo sapiens GN=CHD4 PE=4 SV=1 - [F5H2G1_HUMAN]
B4DIR6	Uncharacterized protein OS=Homo sapiens GN=CHMP5 PE=2 SV=1 - [B4DIR6_HUMAN]
F5H1F8	Uncharacterized protein OS=Homo sapiens GN=CLIC4 PE=4 SV=1 - [F5H1F8_HUMAN]
E9PMI6	Uncharacterized protein OS=Homo sapiens GN=CLNS1A PE=4 SV=1 - [E9PMI6_HUMAN]
B4DIN1	Uncharacterized protein OS=Homo sapiens GN=CLTA PE=2 SV=1 - [B4DIN1_HUMAN]
B4DP17	Uncharacterized protein OS=Homo sapiens GN=CNBP PE=2 SV=1 - [B4DP17_HUMAN]
F5H8M2	Uncharacterized protein OS=Homo sapiens GN=COG6 PE=4 SV=1 - [F5H8M2_HUMAN]
B4DIH5	Uncharacterized protein OS=Homo sapiens GN=COPS2 PE=2 SV=1 - [B4DIH5_HUMAN]
B4DY81	Uncharacterized protein OS=Homo sapiens GN=COPS3 PE=2 SV=1 - [B4DY81_HUMAN]
E7EM64	Uncharacterized protein OS=Homo sapiens GN=COPS6 PE=4 SV=1 - [E7EM64_HUMAN]
E7EUK5	Uncharacterized protein OS=Homo sapiens GN=CPXM2 PE=4 SV=1 - [E7EUK5_HUMAN]
B4DM67	Uncharacterized protein OS=Homo sapiens GN=CSE1L PE=2 SV=1 - [B4DM67_HUMAN]
E9PDV0	Uncharacterized protein OS=Homo sapiens GN=CTH PE=3 SV=1 - [E9PDV0_HUMAN]
F5GWW0	Uncharacterized protein OS=Homo sapiens GN=CTSB PE=3 SV=1 - [F5GWW0_HUMAN]
C9JH19	Uncharacterized protein OS=Homo sapiens GN=CTSD PE=3 SV=1 - [C9JH19_HUMAN]
E7EWR0	Uncharacterized protein OS=Homo sapiens GN=CUL1 PE=3 SV=1 - [E7EWR0_HUMAN]
A8MSH7	Uncharacterized protein OS=Homo sapiens GN=CUL4A PE=3 SV=1 - [A8MSH7_HUMAN]
E9PP19	Uncharacterized protein OS=Homo sapiens GN=CUL5 PE=4 SV=1 - [E9PP19_HUMAN]
C9JFR7	Uncharacterized protein OS=Homo sapiens GN=CYCS PE=3 SV=1 - [C9JFR7_HUMAN]
E7ERE8	Uncharacterized protein OS=Homo sapiens GN=CYP20A1 PE=4 SV=1 - [E7ERE8_HUMAN]
F2Z305	Uncharacterized protein OS=Homo sapiens GN=DCAF6 PE=4 SV=1 - [F2Z305_HUMAN]
F5H5I7	Uncharacterized protein OS=Homo sapiens GN=DCHS2 PE=4 SV=1 - [F5H5I7_HUMAN]
E9PFS5	Uncharacterized protein OS=Homo sapiens GN=DCTN1 PE=4 SV=1 - [E9PFS5_HUMAN]
F5H223	Uncharacterized protein OS=Homo sapiens GN=DCTN2 PE=4 SV=1 - [F5H223_HUMAN]
E7EWW8	Uncharacterized protein OS=Homo sapiens GN=DCTN5 PE=4 SV=1 - [E7EWW8_HUMAN]
C9J8R4	Uncharacterized protein OS=Homo sapiens GN=DCUN1D1 PE=4 SV=1 - [C9J8R4_HUMAN]
B4DME8	Uncharacterized protein OS=Homo sapiens GN=DDX1 PE=2 SV=1 - [B4DME8_HUMAN]
D6RCM4	Uncharacterized protein OS=Homo sapiens GN=DDX4 PE=3 SV=1 - [D6RCM4_HUMAN]
D6RDA2	Uncharacterized protein OS=Homo sapiens GN=DEK PE=4 SV=1 - [D6RDA2_HUMAN]
E9PML7	Uncharacterized protein OS=Homo sapiens GN=DERA PE=4 SV=1 - [E9PML7_HUMAN]
F5H6K0	Uncharacterized protein OS=Homo sapiens GN=DHX15 PE=4 SV=1 - [F5H6K0_HUMAN]
E7ERW8	Uncharacterized protein OS=Homo sapiens GN=DIAPH1 PE=4 SV=1 - [E7ERW8_HUMAN]
E7EPJ5	Uncharacterized protein OS=Homo sapiens GN=DNAH11 PE=4 SV=1 - [E7EPJ5_HUMAN]

E7EUM8	Uncharacterized protein OS=Homo sapiens GN=DNAH17 PE=4 SV=1 - [E7EUM8_HUMAN]
E7EP17	Uncharacterized protein OS=Homo sapiens GN=DNAH9 PE=4 SV=1 - [E7EP17_HUMAN]
C9JXB9	Uncharacterized protein OS=Homo sapiens GN=DNAJB2 PE=4 SV=1 - [C9JXB9_HUMAN]
C9JUL4	Uncharacterized protein OS=Homo sapiens GN=DNAJB4 PE=4 SV=1 - [C9JUL4_HUMAN]
E7EPX3	Uncharacterized protein OS=Homo sapiens GN=DNPEP PE=3 SV=1 - [E7EPX3_HUMAN]
E7EPS3	Uncharacterized protein OS=Homo sapiens GN=DPH5 PE=4 SV=1 - [E7EPS3_HUMAN]
F5GXD5	Uncharacterized protein OS=Homo sapiens GN=DPP3 PE=4 SV=1 - [F5GXD5_HUMAN]
B4DR31	Uncharacterized protein OS=Homo sapiens GN=DPYSL2 PE=2 SV=1 - [B4DR31_HUMAN]
E7ERU0	Uncharacterized protein OS=Homo sapiens GN=DST PE=4 SV=1 - [E7ERU0_HUMAN]
B4DE23	Uncharacterized protein OS=Homo sapiens GN=EHD2 PE=2 SV=1 - [B4DE23_HUMAN]
B4DF96	Uncharacterized protein OS=Homo sapiens GN=EIF2A PE=2 SV=1 - [B4DF96_HUMAN]
F5H0D0	Uncharacterized protein OS=Homo sapiens GN=EIF2B1 PE=3 SV=1 - [F5H0D0_HUMAN]
F5H6W1	Uncharacterized protein OS=Homo sapiens GN=EIF2B4 PE=3 SV=1 - [F5H6W1_HUMAN]
E9PC74	Uncharacterized protein OS=Homo sapiens GN=EIF2B5 PE=4 SV=1 - [E9PC74_HUMAN]
F5H335	Uncharacterized protein OS=Homo sapiens GN=EIF3A PE=4 SV=1 - [F5H335_HUMAN]
B4DV79	Uncharacterized protein OS=Homo sapiens GN=EIF3B PE=2 SV=1 - [B4DV79_HUMAN]
F5H1I1	Uncharacterized protein OS=Homo sapiens GN=EIF3C PE=4 SV=1 - [F5H1I1_HUMAN]
E5RHC7	Uncharacterized protein OS=Homo sapiens GN=EIF3H PE=4 SV=1 - [E5RHC7_HUMAN]
B7ZAM9	Uncharacterized protein OS=Homo sapiens GN=EIF3K PE=2 SV=1 - [B7ZAM9_HUMAN]
E9PN86	Uncharacterized protein OS=Homo sapiens GN=EIF3M PE=4 SV=1 - [E9PN86_HUMAN]
E7EQG2	Uncharacterized protein OS=Homo sapiens GN=EIF4A2 PE=3 SV=1 - [E7EQG2_HUMAN]
E7EPC9	Uncharacterized protein OS=Homo sapiens GN=EIF4B PE=4 SV=1 - [E7EPC9_HUMAN]
E7EX73	Uncharacterized protein OS=Homo sapiens GN=EIF4G1 PE=4 SV=1 - [E7EX73_HUMAN]
B7Z8R4	Uncharacterized protein OS=Homo sapiens GN=ELF2 PE=2 SV=1 - [B7Z8R4_HUMAN]
D6RA00	Uncharacterized protein OS=Homo sapiens GN=ENOPH1 PE=4 SV=1 - [D6RA00_HUMAN]
F5H1W8	Uncharacterized protein OS=Homo sapiens GN=ENOSF1 PE=4 SV=1 - [F5H1W8_HUMAN]
E9PEX0	Uncharacterized protein OS=Homo sapiens GN=EPB41 PE=4 SV=1 - [E9PEX0_HUMAN]
F5H563	Uncharacterized protein OS=Homo sapiens GN=EPB42 PE=4 SV=1 - [F5H563_HUMAN]
E5RJ56	Uncharacterized protein OS=Homo sapiens GN=EPB49 PE=4 SV=1 - [E5RJ56_HUMAN]
F2Z3A2	Uncharacterized protein OS=Homo sapiens GN=ERGIC1 PE=4 SV=1 - [F2Z3A2_HUMAN]
B4DT43	Uncharacterized protein OS=Homo sapiens GN=ETFA PE=2 SV=1 - [B4DT43_HUMAN]
E7ESD5	Uncharacterized protein OS=Homo sapiens GN=EYA4 PE=4 SV=1 - [E7ESD5_HUMAN]
E7EQR4	Uncharacterized protein OS=Homo sapiens GN=EZR PE=4 SV=2 - [E7EQR4_HUMAN]
B4DEJ6	Uncharacterized protein OS=Homo sapiens GN=FAF1 PE=2 SV=1 - [B4DEJ6_HUMAN]
E9PBZ4	Uncharacterized protein OS=Homo sapiens GN=FAM173B PE=4 SV=1 - [E9PBZ4_HUMAN]
F5GYN0	Uncharacterized protein OS=Homo sapiens GN=FAM186A PE=4 SV=1 - [F5GYN0_HUMAN]
B4DZ15	Uncharacterized protein OS=Homo sapiens GN=FAM186B PE=2 SV=1 - [B4DZ15_HUMAN]
B4E363	Uncharacterized protein OS=Homo sapiens GN=FARSA PE=2 SV=1 - [B4E363_HUMAN]
E9PCI9	Uncharacterized protein OS=Homo sapiens GN=FDPS PE=3 SV=1 - [E9PCI9_HUMAN]
F5H1C6	Uncharacterized protein OS=Homo sapiens GN=FERMT3 PE=4 SV=1 - [F5H1C6_HUMAN]
B7Z7Z8	Uncharacterized protein OS=Homo sapiens GN=FKBP5 PE=2 SV=1 - [B7Z7Z8_HUMAN]
E7EN95	Uncharacterized protein OS=Homo sapiens GN=FLNB PE=4 SV=1 - [E7EN95_HUMAN]
F5H0J2	Uncharacterized protein OS=Homo sapiens GN=FMNL1 PE=4 SV=1 - [F5H0J2_HUMAN]

A8MQB8	Uncharacterized protein OS=Homo sapiens GN=FMR1 PE=4 SV=1 - [A8MQB8_HUMAN]
B3KNR9	Uncharacterized protein OS=Homo sapiens GN=FN3K PE=2 SV=1 - [B3KNR9_HUMAN]
C9JCN8	Uncharacterized protein OS=Homo sapiens GN=FOSL2 PE=3 SV=1 - [C9JCN8_HUMAN]
F5H7C9	Uncharacterized protein OS=Homo sapiens GN=FOXO3 PE=4 SV=1 - [F5H7C9_HUMAN]
B4E0X8	Uncharacterized protein OS=Homo sapiens GN=FUBP1 PE=2 SV=1 - [B4E0X8_HUMAN]
F5H4D6	Uncharacterized protein OS=Homo sapiens GN=G3BP1 PE=4 SV=1 - [F5H4D6_HUMAN]
E9PGM4	Uncharacterized protein OS=Homo sapiens GN=GBE1 PE=4 SV=1 - [E9PGM4_HUMAN]
B4E192	Uncharacterized protein OS=Homo sapiens GN=GFAP PE=2 SV=1 - [B4E192_HUMAN]
B8ZZK2	Uncharacterized protein OS=Homo sapiens GN=GGCT PE=4 SV=1 - [B8ZZK2_HUMAN]
E7ESB6	Uncharacterized protein OS=Homo sapiens GN=GIGYF2 PE=4 SV=1 - [E7ESB6_HUMAN]
E7EMY5	Uncharacterized protein OS=Homo sapiens GN=GLOD4 PE=4 SV=1 - [E7EMY5_HUMAN]
C9JAH0	Uncharacterized protein OS=Homo sapiens GN=GMPPA PE=4 SV=1 - [C9JAH0_HUMAN]
B4DUT7	Uncharacterized protein OS=Homo sapiens GN=GMPS PE=2 SV=1 - [B4DUT7_HUMAN]
E9PHP0	Uncharacterized protein OS=Homo sapiens GN=GNL1 PE=4 SV=1 - [E9PHP0_HUMAN]
D6R917	Uncharacterized protein OS=Homo sapiens GN=GNPDA1 PE=4 SV=1 - [D6R917_HUMAN]
E9PDI1	Uncharacterized protein OS=Homo sapiens GN=GPCPD1 PE=4 SV=1 - [E9PDI1_HUMAN]
F5H504	Uncharacterized protein OS=Homo sapiens GN=GRIPAP1 PE=4 SV=1 - [F5H504_HUMAN]
F5H8M4	Uncharacterized protein OS=Homo sapiens GN=GSN PE=4 SV=1 - [F5H8M4_HUMAN]
E7EQZ3	Uncharacterized protein OS=Homo sapiens GN=GSPT1 PE=4 SV=1 - [E7EQZ3_HUMAN]
E9PLF1	Uncharacterized protein OS=Homo sapiens GN=GSTM2 PE=4 SV=1 - [E9PLF1_HUMAN]
A8MX94	Uncharacterized protein OS=Homo sapiens GN=GSTP1 PE=3 SV=1 - [A8MX94_HUMAN]
C9JA47	Uncharacterized protein OS=Homo sapiens GN=GSTT1 PE=4 SV=1 - [C9JA47_HUMAN]
C9J8R8	Uncharacterized protein OS=Homo sapiens GN=GYG1 PE=4 SV=1 - [C9J8R8_HUMAN]
E9PIZ2	Uncharacterized protein OS=Homo sapiens GN=GYLTL1B PE=4 SV=1 - [E9PIZ2_HUMAN]
B4DF58	Uncharacterized protein OS=Homo sapiens GN=GZF1 PE=2 SV=1 - [B4DF58_HUMAN]
F5H5M6	Uncharacterized protein OS=Homo sapiens GN=HABP2 PE=3 SV=1 - [F5H5M6_HUMAN]
D6RF05	Uncharacterized protein OS=Homo sapiens GN=HARS PE=4 SV=1 - [D6RF05_HUMAN]
E9PC78	Uncharacterized protein OS=Homo sapiens GN=HAS1 PE=4 SV=1 - [E9PC78_HUMAN]
B4DQ69	Uncharacterized protein OS=Homo sapiens GN=HCLS1 PE=2 SV=1 - [B4DQ69_HUMAN]
E5RK19	Uncharacterized protein OS=Homo sapiens GN=HDAC2 PE=4 SV=1 - [E5RK19_HUMAN]
F5GZ06	Uncharacterized protein OS=Homo sapiens GN=HEATR7B2 PE=4 SV=1 - [F5GZ06_HUMAN]
F5H5Y5	Uncharacterized protein OS=Homo sapiens GN=HGS PE=4 SV=1 - [F5H5Y5_HUMAN]
E9PCK0	Uncharacterized protein OS=Homo sapiens GN=HK1 PE=3 SV=1 - [E9PCK0_HUMAN]
F5H345	Uncharacterized protein OS=Homo sapiens GN=HMBS PE=4 SV=1 - [F5H345_HUMAN]
F5H4N1	Uncharacterized protein OS=Homo sapiens GN=HMHA1 PE=4 SV=1 - [F5H4N1_HUMAN]
B8ZZT7	Uncharacterized protein OS=Homo sapiens GN=HN1 PE=4 SV=1 - [B8ZZT7_HUMAN]
B4DL85	Uncharacterized protein OS=Homo sapiens GN=HN1L PE=2 SV=1 - [B4DL85_HUMAN]
E7EWI9	Uncharacterized protein OS=Homo sapiens GN=HNRNPA3 PE=4 SV=1 - [E7EWI9_HUMAN]
D6R9P3	Uncharacterized protein OS=Homo sapiens GN=HNRNPAB PE=4 SV=1 - [D6R9P3_HUMAN]
B4DTC3	Uncharacterized protein OS=Homo sapiens GN=HNRNPD PE=2 SV=1 - [B4DTC3_HUMAN]
B4DKS8	Uncharacterized protein OS=Homo sapiens GN=HNRNPF PE=2 SV=1 - [B4DKS8_HUMAN]
B4DFK9	Uncharacterized protein OS=Homo sapiens GN=HNRNPH2 PE=2 SV=1 - [B4DFK9_HUMAN]
B4DPK8	Uncharacterized protein OS=Homo sapiens GN=HNRNPL PE=2 SV=1 - [B4DPK8_HUMAN]

B3KX72	Uncharacterized protein OS=Homo sapiens GN=HNRNPU PE=2 SV=1 - [B3KX72_HUMAN]
E9PAZ1	Uncharacterized protein OS=Homo sapiens GN=HOMER2 PE=4 SV=1 - [E9PAZ1_HUMAN]
E5RIP8	Uncharacterized protein OS=Homo sapiens GN=HRSP12 PE=4 SV=1 - [E5RIP8_HUMAN]
E9PDE8	Uncharacterized protein OS=Homo sapiens GN=HSPA4L PE=3 SV=1 - [E9PDE8_HUMAN]
B4DL87	Uncharacterized protein OS=Homo sapiens GN=HSPB1 PE=2 SV=1 - [B4DL87_HUMAN]
E7EXB4	Uncharacterized protein OS=Homo sapiens GN=HSPD1 PE=3 SV=1 - [E7EXB4_HUMAN]
E9PD66	Uncharacterized protein OS=Homo sapiens GN=HUWE1 PE=4 SV=1 - [E9PD66_HUMAN]
E9PDR5	Uncharacterized protein OS=Homo sapiens GN=IBTK PE=4 SV=1 - [E9PDR5_HUMAN]
E7EP25	Uncharacterized protein OS=Homo sapiens GN=IFT172 PE=4 SV=1 - [E7EP25_HUMAN]
D6RGV7	Uncharacterized protein OS=Homo sapiens GN=IGSF22 PE=4 SV=1 - [D6RGV7_HUMAN]
B4DFG4	Uncharacterized protein OS=Homo sapiens GN=ILF3 PE=2 SV=1 - [B4DFG4_HUMAN]
C9J406	Uncharacterized protein OS=Homo sapiens GN=IMMT PE=4 SV=1 - [C9J406_HUMAN]
E7EMB7	Uncharacterized protein OS=Homo sapiens GN=IPO11 PE=4 SV=1 - [E7EMB7_HUMAN]
B4E0R6	Uncharacterized protein OS=Homo sapiens GN=IPO5 PE=2 SV=1 - [B4E0R6_HUMAN]
C9JL81	Uncharacterized protein OS=Homo sapiens GN=IQCD PE=4 SV=1 - [C9JL81_HUMAN]
E7EWZ6	Uncharacterized protein OS=Homo sapiens GN=ITGAV PE=3 SV=1 - [E7EWZ6_HUMAN]
Q6ZW95	Uncharacterized protein OS=Homo sapiens GN=KCNK2 PE=2 SV=1 - [Q6ZW95_HUMAN]
F5GX28	Uncharacterized protein OS=Homo sapiens GN=KDM4B PE=4 SV=1 - [F5GX28_HUMAN]
E7ET98	Uncharacterized protein OS=Homo sapiens GN=KHDRBS1 PE=4 SV=2 - [E7ET98_HUMAN]
E7EP96	Uncharacterized protein OS=Homo sapiens GN=KHSRP PE=4 SV=1 - [E7EP96_HUMAN]
F5H6Z7	Uncharacterized protein OS=Homo sapiens GN=KIAA0174 PE=4 SV=1 - [F5H6Z7_HUMAN]
E7EQI7	Uncharacterized protein OS=Homo sapiens GN=KIAA0196 PE=4 SV=1 - [E7EQI7_HUMAN]
E9PEZ5	Uncharacterized protein OS=Homo sapiens GN=KIAA1107 PE=4 SV=1 - [E9PEZ5_HUMAN]
C9J352	Uncharacterized protein OS=Homo sapiens GN=KPNA1 PE=4 SV=1 - [C9J352_HUMAN]
E9PCN9	Uncharacterized protein OS=Homo sapiens GN=KPNA3 PE=4 SV=1 - [E9PCN9_HUMAN]
F5H4G7	Uncharacterized protein OS=Homo sapiens GN=KPNA6 PE=4 SV=1 - [F5H4G7_HUMAN]
F5H6G5	Uncharacterized protein OS=Homo sapiens GN=KRT6B PE=3 SV=1 - [F5H6G5_HUMAN]
C9JVN9	Uncharacterized protein OS=Homo sapiens GN=L2HGDH PE=4 SV=1 - [C9JVN9_HUMAN]
A6NGS5	Uncharacterized protein OS=Homo sapiens GN=LAMA2 PE=4 SV=2 - [A6NGS5_HUMAN]
E9PHS0	Uncharacterized protein OS=Homo sapiens GN=LANCL1 PE=4 SV=1 - [E9PHS0_HUMAN]
B4DJ10	Uncharacterized protein OS=Homo sapiens GN=LARS PE=2 SV=1 - [B4DJ10_HUMAN]
F5GY62	Uncharacterized protein OS=Homo sapiens GN=LIMCH1 PE=4 SV=1 - [F5GY62_HUMAN]
F5GX84	Uncharacterized protein OS=Homo sapiens GN=LMCD1 PE=4 SV=1 - [F5GX84_HUMAN]
E9PBF6	Uncharacterized protein OS=Homo sapiens GN=LMNB1 PE=3 SV=1 - [E9PBF6_HUMAN]
F5H2T1	Uncharacterized protein OS=Homo sapiens GN=LRCH2 PE=4 SV=1 - [F5H2T1_HUMAN]
B4DS68	Uncharacterized protein OS=Homo sapiens GN=LRP11 PE=2 SV=1 - [B4DS68_HUMAN]
E5RJ48	Uncharacterized protein OS=Homo sapiens GN=LYPLA1 PE=4 SV=1 - [E5RJ48_HUMAN]
F5GWL8	Uncharacterized protein OS=Homo sapiens GN=LZTFL1 PE=4 SV=1 - [F5GWL8_HUMAN]
F5GXU8	Uncharacterized protein OS=Homo sapiens GN=LZTR1 PE=4 SV=1 - [F5GXU8_HUMAN]
C9JIR0	Uncharacterized protein OS=Homo sapiens GN=MAD1L1 PE=4 SV=1 - [C9JIR0_HUMAN]
F5H3U9	Uncharacterized protein OS=Homo sapiens GN=MAGOHB PE=4 SV=1 - [F5H3U9_HUMAN]
B4DH44	Uncharacterized protein OS=Homo sapiens GN=MAP2K4 PE=2 SV=1 - [B4DH44_HUMAN]
B7Z895	Uncharacterized protein OS=Homo sapiens GN=MAP7D2 PE=2 SV=1 - [B7Z895_HUMAN]

D6REN1	Uncharacterized protein OS=Homo sapiens GN=MARCH1 PE=4 SV=1 - [D6REN1_HUMAN]
A8MUU4	Uncharacterized protein OS=Homo sapiens GN=MBP PE=4 SV=2 - [A8MUU4_HUMAN]
D6REY2	Uncharacterized protein OS=Homo sapiens GN=MCC PE=4 SV=1 - [D6REY2_HUMAN]
C9J600	Uncharacterized protein OS=Homo sapiens GN=MCM10 PE=4 SV=1 - [C9J600_HUMAN]
E5RK04	Uncharacterized protein OS=Homo sapiens GN=MCM4 PE=4 SV=1 - [E5RK04_HUMAN]
E5RJR1	Uncharacterized protein OS=Homo sapiens GN=MCTP1 PE=4 SV=1 - [E5RJR1_HUMAN]
F5H3Y1	Uncharacterized protein OS=Homo sapiens GN=MED12 PE=4 SV=1 - [F5H3Y1_HUMAN]
E7EQ94	Uncharacterized protein OS=Homo sapiens GN=MET PE=4 SV=2 - [E7EQ94_HUMAN]
E9PGF9	Uncharacterized protein OS=Homo sapiens GN=MGEA5 PE=4 SV=1 - [E9PGF9_HUMAN]
C9JLN4	Uncharacterized protein OS=Homo sapiens GN=MICAL1 PE=4 SV=1 - [C9JLN4_HUMAN]
B7Z849	Uncharacterized protein OS=Homo sapiens GN=MICAL2 PE=2 SV=1 - [B7Z849_HUMAN]
F5H4V4	Uncharacterized protein OS=Homo sapiens GN=MKI67 PE=4 SV=1 - [F5H4V4_HUMAN]
D6RCK3	Uncharacterized protein OS=Homo sapiens GN=MOBK1A PE=4 SV=1 - [D6RCK3_HUMAN]
F5H8H9	Uncharacterized protein OS=Homo sapiens GN=MPHOSPH8 PE=4 SV=1 - [F5H8H9_HUMAN]
F5H2F9	Uncharacterized protein OS=Homo sapiens GN=MSH6 PE=3 SV=1 - [F5H2F9_HUMAN]
E9PMQ0	Uncharacterized protein OS=Homo sapiens GN=MYB PE=4 SV=1 - [E9PMQ0_HUMAN]
B4DRZ1	Uncharacterized protein OS=Homo sapiens GN=MYCN PE=2 SV=1 - [B4DRZ1_HUMAN]
F5GWY8	Uncharacterized protein OS=Homo sapiens GN=MYO1A PE=4 SV=1 - [F5GWY8_HUMAN]
F5H7I9	Uncharacterized protein OS=Homo sapiens GN=MYPN PE=4 SV=1 - [F5H7I9_HUMAN]
F2Z2Y6	Uncharacterized protein OS=Homo sapiens GN=NAA38 PE=4 SV=1 - [F2Z2Y6_HUMAN]
E7EQ69	Uncharacterized protein OS=Homo sapiens GN=NAA50 PE=4 SV=1 - [E7EQ69_HUMAN]
A8MU28	Uncharacterized protein OS=Homo sapiens GN=NAE1 PE=4 SV=1 - [A8MU28_HUMAN]
F5H1X5	Uncharacterized protein OS=Homo sapiens GN=NAP1L1 PE=3 SV=1 - [F5H1X5_HUMAN]
C9JZI7	Uncharacterized protein OS=Homo sapiens GN=NAP1L4 PE=3 SV=1 - [C9JZI7_HUMAN]
E7EPF5	Uncharacterized protein OS=Homo sapiens GN=NCK2 PE=4 SV=1 - [E7EPF5_HUMAN]
Q6ZS38	Uncharacterized protein OS=Homo sapiens GN=NDUFS7 PE=2 SV=1 - [Q6ZS38_HUMAN]
D6RBG5	Uncharacterized protein OS=Homo sapiens GN=NEK1 PE=4 SV=1 - [D6RBG5_HUMAN]
B4DZU0	Uncharacterized protein OS=Homo sapiens GN=NEK2 PE=2 SV=1 - [B4DZU0_HUMAN]
C9JFL6	Uncharacterized protein OS=Homo sapiens GN=NFE2L2 PE=4 SV=1 - [C9JFL6_HUMAN]
D6RF93	Uncharacterized protein OS=Homo sapiens GN=NFKB1 PE=4 SV=1 - [D6RF93_HUMAN]
B5MDB6	Uncharacterized protein OS=Homo sapiens GN=NLRC3 PE=4 SV=2 - [B5MDB6_HUMAN]
A6NJK7	Uncharacterized protein OS=Homo sapiens GN=NOL8 PE=4 SV=2 - [A6NJK7_HUMAN]
E9PN47	Uncharacterized protein OS=Homo sapiens GN=NPEPL1 PE=4 SV=1 - [E9PN47_HUMAN]
B7Z463	Uncharacterized protein OS=Homo sapiens GN=NPEPPS PE=2 SV=1 - [B7Z463_HUMAN]
B4DG89	Uncharacterized protein OS=Homo sapiens GN=NPLOC4 PE=2 SV=1 - [B4DG89_HUMAN]
B4E2P3	Uncharacterized protein OS=Homo sapiens GN=NR5A2 PE=2 SV=1 - [B4E2P3_HUMAN]
E9PR11	Uncharacterized protein OS=Homo sapiens GN=NRBP2 PE=4 SV=1 - [E9PR11_HUMAN]
B9A035	Uncharacterized protein OS=Homo sapiens GN=NT5C3 PE=4 SV=1 - [B9A035_HUMAN]
E5RHQ3	Uncharacterized protein OS=Homo sapiens GN=NUDCD1 PE=4 SV=1 - [E5RHQ3_HUMAN]
E5RFP0	Uncharacterized protein OS=Homo sapiens GN=NUDCD2 PE=4 SV=1 - [E5RFP0_HUMAN]
F5H2E3	Uncharacterized protein OS=Homo sapiens GN=OPHN1 PE=4 SV=1 - [F5H2E3_HUMAN]
F5GYN4	Uncharacterized protein OS=Homo sapiens GN=OTUB1 PE=4 SV=1 - [F5GYN4_HUMAN]
B4DUA5	Uncharacterized protein OS=Homo sapiens GN=P4HB PE=2 SV=1 - [B4DUA5_HUMAN]

E7EQV3	Uncharacterized protein OS=Homo sapiens GN=PABPC1 PE=4 SV=1 - [E7EQV3_HUMAN]
E9PBS1	Uncharacterized protein OS=Homo sapiens GN=PAICS PE=4 SV=1 - [E9PBS1_HUMAN]
E9PHT6	Uncharacterized protein OS=Homo sapiens GN=PANK4 PE=4 SV=1 - [E9PHT6_HUMAN]
B4DV82	Uncharacterized protein OS=Homo sapiens GN=PARP2 PE=2 SV=1 - [B4DV82_HUMAN]
E5RGG7	Uncharacterized protein OS=Homo sapiens GN=PCMTD1 PE=3 SV=1 - [E5RGG7_HUMAN]
E9PQD8	Uncharacterized protein OS=Homo sapiens GN=PDDC1 PE=4 SV=1 - [E9PQD8_HUMAN]
F6T1Q0	Uncharacterized protein OS=Homo sapiens GN=PDE12 PE=4 SV=1 - [F6T1Q0_HUMAN]
E9PJ64	Uncharacterized protein OS=Homo sapiens GN=PDE4DIP PE=4 SV=1 - [E9PJ64_HUMAN]
B3KQU2	Uncharacterized protein OS=Homo sapiens GN=PDIA2 PE=2 SV=1 - [B3KQU2_HUMAN]
B3KQT9	Uncharacterized protein OS=Homo sapiens GN=PDIA3 PE=2 SV=1 - [B3KQT9_HUMAN]
F2Z2Y4	Uncharacterized protein OS=Homo sapiens GN=PD XK PE=4 SV=1 - [F2Z2Y4_HUMAN]
E9PCE8	Uncharacterized protein OS=Homo sapiens GN=PEPD PE=3 SV=1 - [E9PCE8_HUMAN]
E9PN05	Uncharacterized protein OS=Homo sapiens GN=PICALM PE=4 SV=1 - [E9PN05_HUMAN]
F2Z3N1	Uncharacterized protein OS=Homo sapiens GN=PIP5K1B PE=4 SV=2 - [F2Z3N1_HUMAN]
F5GWE5	Uncharacterized protein OS=Homo sapiens GN=PITPNA PE=4 SV=1 - [F5GWE5_HUMAN]
E7EU42	Uncharacterized protein OS=Homo sapiens GN=PLA2G4A PE=4 SV=1 - [E7EU42_HUMAN]
E5RIM3	Uncharacterized protein OS=Homo sapiens GN=PLAA PE=4 SV=1 - [E5RIM3_HUMAN]
B7Z6M1	Uncharacterized protein OS=Homo sapiens GN=PLS3 PE=2 SV=1 - [B7Z6M1_HUMAN]
B4E0V0	Uncharacterized protein OS=Homo sapiens GN=PNPO PE=2 SV=1 - [B4E0V0_HUMAN]
F5H696	Uncharacterized protein OS=Homo sapiens GN=POLQ PE=4 SV=1 - [F5H696_HUMAN]
A6NM32	Uncharacterized protein OS=Homo sapiens GN=PPIH PE=3 SV=1 - [A6NM32_HUMAN]
B8ZZ77	Uncharacterized protein OS=Homo sapiens GN=PPIL3 PE=3 SV=1 - [B8ZZ77_HUMAN]
A8MYH2	Uncharacterized protein OS=Homo sapiens GN=PPIP5K1 PE=4 SV=1 - [A8MYH2_HUMAN]
D6RFZ8	Uncharacterized protein OS=Homo sapiens GN=PPIP5K2 PE=4 SV=1 - [D6RFZ8_HUMAN]
F5GYS1	Uncharacterized protein OS=Homo sapiens GN=PPP2R5D PE=4 SV=1 - [F5GYS1_HUMAN]
B4DLT9	Uncharacterized protein OS=Homo sapiens GN=PPP6R1 PE=2 SV=1 - [B4DLT9_HUMAN]
E9PH29	Uncharacterized protein OS=Homo sapiens GN=PRDX3 PE=4 SV=1 - [E9PH29_HUMAN]
E9PBE1	Uncharacterized protein OS=Homo sapiens GN=PRKCZ PE=4 SV=1 - [E9PBE1_HUMAN]
E7EPE6	Uncharacterized protein OS=Homo sapiens GN=PRKG2 PE=4 SV=1 - [E7EPE6_HUMAN]
E9PNE8	Uncharacterized protein OS=Homo sapiens GN=PRPH PE=3 SV=1 - [E9PNE8_HUMAN]
C9J168	Uncharacterized protein OS=Homo sapiens GN=PRPSAP1 PE=4 SV=1 - [C9J168_HUMAN]
E7EPA1	Uncharacterized protein OS=Homo sapiens GN=PRPSAP2 PE=4 SV=1 - [E7EPA1_HUMAN]
F5GX11	Uncharacterized protein OS=Homo sapiens GN=PSMA1 PE=4 SV=1 - [F5GX11_HUMAN]
F5GZ16	Uncharacterized protein OS=Homo sapiens GN=PSMD2 PE=4 SV=1 - [F5GZ16_HUMAN]
<i>F5H8K4</i>	<i>Uncharacterized protein OS=Homo sapiens GN=PSMD3 PE=4 SV=1 - [F5H8K4_HUMAN]</i>
E9PHI9	Uncharacterized protein OS=Homo sapiens GN=PSMD6 PE=4 SV=1 - [E9PHI9_HUMAN]
<i>B4DXI8</i>	<i>Uncharacterized protein OS=Homo sapiens GN=PSMD7 PE=2 SV=1 - [B4DXI8_HUMAN]</i>
F5H4W8	Uncharacterized protein OS=Homo sapiens GN=PSMG2 PE=4 SV=1 - [F5H4W8_HUMAN]
B8ZZQ6	Uncharacterized protein OS=Homo sapiens GN=PTMA PE=4 SV=1 - [B8ZZQ6_HUMAN]
B4DX06	Uncharacterized protein OS=Homo sapiens GN=PTPN7 PE=2 SV=1 - [B4DX06_HUMAN]
B4DNN3	Uncharacterized protein OS=Homo sapiens GN=QARS PE=2 SV=1 - [B4DNN3_HUMAN]
E7EX40	Uncharacterized protein OS=Homo sapiens GN=RAB11FIP1 PE=4 SV=1 - [E7EX40_HUMAN]
C9J7D1	Uncharacterized protein OS=Homo sapiens GN=RAB7A PE=4 SV=1 - [C9J7D1_HUMAN]

B4DHR0	Uncharacterized protein OS=Homo sapiens GN=RABEP2 PE=2 SV=1 - [B4DHR0_HUMAN]
F5H4C2	Uncharacterized protein OS=Homo sapiens GN=RANBP3 PE=4 SV=1 - [F5H4C2_HUMAN]
B4DW94	Uncharacterized protein OS=Homo sapiens GN=RAP1B PE=2 SV=1 - [B4DW94_HUMAN]
E9PGY3	Uncharacterized protein OS=Homo sapiens GN=RAPGEF5 PE=4 SV=1 - [E9PGY3_HUMAN]
F5H3T8	Uncharacterized protein OS=Homo sapiens GN=RARS PE=4 SV=1 - [F5H3T8_HUMAN]
E9PC52	Uncharacterized protein OS=Homo sapiens GN=RBBP7 PE=4 SV=1 - [E9PC52_HUMAN]
E9PSF1	Uncharacterized protein OS=Homo sapiens GN=RBM38 PE=4 SV=1 - [E9PSF1_HUMAN]
B4DNY1	Uncharacterized protein OS=Homo sapiens GN=RBM6 PE=2 SV=1 - [B4DNY1_HUMAN]
F5GX07	Uncharacterized protein OS=Homo sapiens GN=REXO2 PE=4 SV=1 - [F5GX07_HUMAN]
D6RJG8	Uncharacterized protein OS=Homo sapiens GN=RFESD PE=4 SV=1 - [D6RJG8_HUMAN]
E7ETT0	Uncharacterized protein OS=Homo sapiens GN=RICTOR PE=4 SV=1 - [E7ETT0_HUMAN]
E9PCZ1	Uncharacterized protein OS=Homo sapiens GN=RIMS1 PE=4 SV=1 - [E9PCZ1_HUMAN]
F6V054	Uncharacterized protein OS=Homo sapiens GN=RIN3 PE=4 SV=1 - [F6V054_HUMAN]
E7EP22	Uncharacterized protein OS=Homo sapiens GN=RNFB14 PE=4 SV=1 - [E7EP22_HUMAN]
F6SYU7	Uncharacterized protein OS=Homo sapiens GN=RNFB40 PE=4 SV=1 - [F6SYU7_HUMAN]
A6NKB8	Uncharacterized protein OS=Homo sapiens GN=RNPEP PE=4 SV=1 - [A6NKB8_HUMAN]
E9PF63	Uncharacterized protein OS=Homo sapiens GN=ROCK2 PE=4 SV=1 - [E9PF63_HUMAN]
C9J6A7	Uncharacterized protein OS=Homo sapiens GN=RPE PE=4 SV=1 - [C9J6A7_HUMAN]
E7EPB3	Uncharacterized protein OS=Homo sapiens GN=RPL14 PE=4 SV=1 - [E7EPB3_HUMAN]
B4DM74	Uncharacterized protein OS=Homo sapiens GN=RPL18A PE=2 SV=1 - [B4DM74_HUMAN]
E5RIT6	Uncharacterized protein OS=Homo sapiens GN=RPL26L1 PE=3 SV=1 - [E5RIT6_HUMAN]
F2Z3A5	Uncharacterized protein OS=Homo sapiens GN=RPL3 PE=3 SV=1 - [F2Z3A5_HUMAN]
B8ZZK4	Uncharacterized protein OS=Homo sapiens GN=RPL31 PE=4 SV=1 - [B8ZZK4_HUMAN]
E7EWF1	Uncharacterized protein OS=Homo sapiens GN=RPL4 PE=4 SV=1 - [E7EWF1_HUMAN]
E9PKZ0	Uncharacterized protein OS=Homo sapiens GN=RPL8 PE=4 SV=1 - [E9PKZ0_HUMAN]
B4E1M5	Uncharacterized protein OS=Homo sapiens GN=RPL9 PE=2 SV=1 - [B4E1M5_HUMAN]
E9PS50	Uncharacterized protein OS=Homo sapiens GN=RPS13 PE=4 SV=1 - [E9PS50_HUMAN]
E5RH77	Uncharacterized protein OS=Homo sapiens GN=RPS14 PE=4 SV=1 - [E5RH77_HUMAN]
E9PM36	Uncharacterized protein OS=Homo sapiens GN=RPS2 PE=3 SV=1 - [E9PM36_HUMAN]
D6RDJ2	Uncharacterized protein OS=Homo sapiens GN=RPS23 PE=3 SV=1 - [D6RDJ2_HUMAN]
C9J9K3	Uncharacterized protein OS=Homo sapiens GN=RPSA PE=3 SV=1 - [C9J9K3_HUMAN]
D6RA96	Uncharacterized protein OS=Homo sapiens GN=RTEL1 PE=4 SV=1 - [D6RA96_HUMAN]
E5RJE3	Uncharacterized protein OS=Homo sapiens GN=RWDD1 PE=4 SV=1 - [E5RJE3_HUMAN]
F5H2Z8	Uncharacterized protein OS=Homo sapiens GN=SAGE1 PE=4 SV=1 - [F5H2Z8_HUMAN]
B7Z5N7	Uncharacterized protein OS=Homo sapiens GN=SCFD1 PE=2 SV=1 - [B7Z5N7_HUMAN]
B4DH22	Uncharacterized protein OS=Homo sapiens GN=SCRN1 PE=2 SV=1 - [B4DH22_HUMAN]
F5H3U4	Uncharacterized protein OS=Homo sapiens GN=SEC14L2 PE=4 SV=1 - [F5H3U4_HUMAN]
B4DNE4	Uncharacterized protein OS=Homo sapiens GN=SEPT7 PE=2 SV=1 - [B4DNE4_HUMAN]
E5RIQ9	Uncharacterized protein OS=Homo sapiens GN=SERF1A PE=4 SV=1 - [E5RIQ9_HUMAN]
B4E3A8	Uncharacterized protein OS=Homo sapiens GN=SERPINB1 PE=2 SV=1 - [B4E3A8_HUMAN]
C9JN98	Uncharacterized protein OS=Homo sapiens GN=SERPINE2 PE=3 SV=1 - [C9JN98_HUMAN]
E9PJ04	Uncharacterized protein OS=Homo sapiens GN=SF3B2 PE=4 SV=1 - [E9PJ04_HUMAN]
E7EMJ6	Uncharacterized protein OS=Homo sapiens GN=SGTA PE=4 SV=1 - [E7EMJ6_HUMAN]

F5H229	Uncharacterized protein OS=Homo sapiens GN=SH3GL2 PE=4 SV=1 - [F5H229_HUMAN]
F5H7E2	Uncharacterized protein OS=Homo sapiens GN=SKIV2L2 PE=4 SV=1 - [F5H7E2_HUMAN]
E5RI56	Uncharacterized protein OS=Homo sapiens GN=SKP1 PE=4 SV=1 - [E5RI56_HUMAN]
E5RJR5	Uncharacterized protein OS=Homo sapiens GN=SKP1 PE=4 SV=1 - [E5RJR5_HUMAN]
C9JK45	Uncharacterized protein OS=Homo sapiens GN=SLC35E1 PE=4 SV=1 - [C9JK45_HUMAN]
B4DDK0	Uncharacterized protein OS=Homo sapiens GN=SLC39A9 PE=2 SV=1 - [B4DDK0_HUMAN]
F5H6G3	Uncharacterized protein OS=Homo sapiens GN=SLFN12L PE=4 SV=1 - [F5H6G3_HUMAN]
D6RBH9	Uncharacterized protein OS=Homo sapiens GN=SMAD1 PE=4 SV=1 - [D6RBH9_HUMAN]
F5H0U7	Uncharacterized protein OS=Homo sapiens GN=SMC4 PE=4 SV=1 - [F5H0U7_HUMAN]
E7EPV7	Uncharacterized protein OS=Homo sapiens GN=SNCA PE=4 SV=1 - [E7EPV7_HUMAN]
E7ESM6	Uncharacterized protein OS=Homo sapiens GN=SND1 PE=4 SV=2 - [E7ESM6_HUMAN]
B4DDC7	Uncharacterized protein OS=Homo sapiens GN=SNRPA PE=2 SV=1 - [B4DDC7_HUMAN]
E9PLV3	Uncharacterized protein OS=Homo sapiens GN=SNX19 PE=4 SV=1 - [E9PLV3_HUMAN]
B4DEK4	Uncharacterized protein OS=Homo sapiens GN=SNX2 PE=2 SV=1 - [B4DEK4_HUMAN]
Q9H5Q8	Uncharacterized protein OS=Homo sapiens GN=SNX25 PE=2 SV=1 - [Q9H5Q8_HUMAN]
E9PBC7	Uncharacterized protein OS=Homo sapiens GN=SNX5 PE=4 SV=1 - [E9PBC7_HUMAN]
B4DGT1	Uncharacterized protein OS=Homo sapiens GN=SPTAN1 PE=2 SV=1 - [B4DGT1_HUMAN]
C9J0K6	Uncharacterized protein OS=Homo sapiens GN=SRI PE=4 SV=1 - [C9J0K6_HUMAN]
E7ETK6	Uncharacterized protein OS=Homo sapiens GN=SRP68 PE=4 SV=1 - [E7ETK6_HUMAN]
B4DED5	Uncharacterized protein OS=Homo sapiens GN=SRRM1 PE=2 SV=1 - [B4DED5_HUMAN]
A6NMU3	Uncharacterized protein OS=Homo sapiens GN=STAM PE=4 SV=1 - [A6NMU3_HUMAN]
C9JZ93	Uncharacterized protein OS=Homo sapiens GN=STAMBP PE=4 SV=1 - [C9JZ93_HUMAN]
C9JQV3	Uncharacterized protein OS=Homo sapiens GN=STK11IP PE=4 SV=1 - [C9JQV3_HUMAN]
B4E2V5	Uncharacterized protein OS=Homo sapiens GN=STOM PE=2 SV=1 - [B4E2V5_HUMAN]
B0AZV0	Uncharacterized protein OS=Homo sapiens GN=STRAP PE=2 SV=1 - [B0AZV0_HUMAN]
B5MCT9	Uncharacterized protein OS=Homo sapiens GN=STXBP4 PE=4 SV=1 - [B5MCT9_HUMAN]
F2Z3D0	Uncharacterized protein OS=Homo sapiens GN=SUMO2 PE=4 SV=1 - [F2Z3D0_HUMAN]
C9JIE4	Uncharacterized protein OS=Homo sapiens GN=SYNM PE=3 SV=1 - [C9JIE4_HUMAN]
A6NCR3	Uncharacterized protein OS=Homo sapiens GN=SYNPO2L PE=4 SV=1 - [A6NCR3_HUMAN]
F5GWQ7	Uncharacterized protein OS=Homo sapiens GN=TAF15 PE=4 SV=1 - [F5GWQ7_HUMAN]
B8ZZD4	Uncharacterized protein OS=Homo sapiens GN=TAX1BP1 PE=4 SV=1 - [B8ZZD4_HUMAN]
C9JF30	Uncharacterized protein OS=Homo sapiens GN=TBC1D26 PE=4 SV=1 - [C9JF30_HUMAN]
F5GWQ9	Uncharacterized protein OS=Homo sapiens GN=TBC1D3H PE=4 SV=1 - [F5GWQ9_HUMAN]
E5RIW3	Uncharacterized protein OS=Homo sapiens GN=TBCA PE=4 SV=1 - [E5RIW3_HUMAN]
A8MVD5	Uncharacterized protein OS=Homo sapiens GN=TBCB PE=4 SV=1 - [A8MVD5_HUMAN]
C9J1U4	Uncharacterized protein OS=Homo sapiens GN=TBCD PE=4 SV=1 - [C9J1U4_HUMAN]
D6RB05	Uncharacterized protein OS=Homo sapiens GN=TEC PE=4 SV=1 - [D6RB05_HUMAN]
E7EVC1	Uncharacterized protein OS=Homo sapiens GN=TEKT4 PE=4 SV=1 - [E7EVC1_HUMAN]
C9JVR1	Uncharacterized protein OS=Homo sapiens GN=TENC1 PE=4 SV=1 - [C9JVR1_HUMAN]
E7EQE3	Uncharacterized protein OS=Homo sapiens GN=TEP1 PE=4 SV=1 - [E7EQE3_HUMAN]
C9JJP5	Uncharacterized protein OS=Homo sapiens GN=TFG PE=4 SV=1 - [C9JJP5_HUMAN]
B4DIT7	Uncharacterized protein OS=Homo sapiens GN=TGM2 PE=2 SV=1 - [B4DIT7_HUMAN]
E5RIQ8	Uncharacterized protein OS=Homo sapiens GN=THG1L PE=4 SV=1 - [E5RIQ8_HUMAN]

E7EUG5	Uncharacterized protein OS=Homo sapiens GN=THUMPD2 PE=4 SV=1 - [E7EUG5_HUMAN]
A6NJC0	Uncharacterized protein OS=Homo sapiens GN=TMCC2 PE=4 SV=2 - [A6NJC0_HUMAN]
B7Z588	Uncharacterized protein OS=Homo sapiens GN=TNFRSF10B PE=2 SV=1 - [B7Z588_HUMAN]
E9PFH4	Uncharacterized protein OS=Homo sapiens GN=TNPO3 PE=4 SV=1 - [E9PFH4_HUMAN]
A8MY3	Uncharacterized protein OS=Homo sapiens GN=TNRC6B PE=4 SV=3 - [A8MY3_HUMAN]
F5H7S3	Uncharacterized protein OS=Homo sapiens GN=TPM1 PE=3 SV=1 - [F5H7S3_HUMAN]
B4DSE2	Uncharacterized protein OS=Homo sapiens GN=TPP1 PE=2 SV=1 - [B4DSE2_HUMAN]
E7EQD1	Uncharacterized protein OS=Homo sapiens GN=TRA2B PE=4 SV=1 - [E7EQD1_HUMAN]
B7WP88	Uncharacterized protein OS=Homo sapiens GN=TRANK1 PE=4 SV=2 - [B7WP88_HUMAN]
F5H012	Uncharacterized protein OS=Homo sapiens GN=TRIM21 PE=4 SV=1 - [F5H012_HUMAN]
C9JNQ0	Uncharacterized protein OS=Homo sapiens GN=TRIM6 PE=4 SV=1 - [C9JNQ0_HUMAN]
F5H1Z0	Uncharacterized protein OS=Homo sapiens GN=TRIP11 PE=4 SV=1 - [F5H1Z0_HUMAN]
E9PGT1	Uncharacterized protein OS=Homo sapiens GN=TSN PE=4 SV=1 - [E9PGT1_HUMAN]
E7ES78	Uncharacterized protein OS=Homo sapiens GN=TTC26 PE=4 SV=1 - [E7ES78_HUMAN]
D6RCE2	Uncharacterized protein OS=Homo sapiens GN=TTC37 PE=4 SV=1 - [D6RCE2_HUMAN]
A6NKB1	Uncharacterized protein OS=Homo sapiens GN=TTN PE=4 SV=4 - [A6NKB1_HUMAN]
D6RDV0	Uncharacterized protein OS=Homo sapiens GN=TUSC3 PE=4 SV=1 - [D6RDV0_HUMAN]
E9PIR7	Uncharacterized protein OS=Homo sapiens GN=TXNRD1 PE=3 SV=1 - [E9PIR7_HUMAN]
E7EWE1	Uncharacterized protein OS=Homo sapiens GN=UBA5 PE=4 SV=1 - [E7EWE1_HUMAN]
F2Z363	Uncharacterized protein OS=Homo sapiens GN=UBE2F PE=4 SV=1 - [F2Z363_HUMAN]
D6RDM7	Uncharacterized protein OS=Homo sapiens GN=UBE2K PE=4 SV=1 - [D6RDM7_HUMAN]
E9PJ81	Uncharacterized protein OS=Homo sapiens GN=UBXN1 PE=4 SV=1 - [E9PJ81_HUMAN]
B4DW59	Uncharacterized protein OS=Homo sapiens GN=UCL5 PE=2 SV=1 - [B4DW59_HUMAN]
A6NJ11	Uncharacterized protein OS=Homo sapiens GN=UFD1L PE=4 SV=1 - [A6NJ11_HUMAN]
C9JKD6	Uncharacterized protein OS=Homo sapiens GN=UGP2 PE=4 SV=1 - [C9JKD6_HUMAN]
C9J1U3	Uncharacterized protein OS=Homo sapiens GN=UNC80 PE=4 SV=1 - [C9J1U3_HUMAN]
E9PG85	Uncharacterized protein OS=Homo sapiens GN=UROS PE=4 SV=1 - [E9PG85_HUMAN]
B7Z7T5	Uncharacterized protein OS=Homo sapiens GN=USP7 PE=2 SV=1 - [B7Z7T5_HUMAN]
B0V043	Uncharacterized protein OS=Homo sapiens GN=VAR5 PE=3 SV=1 - [B0V043_HUMAN]
F5H2A7	Uncharacterized protein OS=Homo sapiens GN=VBP1 PE=4 SV=1 - [F5H2A7_HUMAN]
E5RHE1	Uncharacterized protein OS=Homo sapiens GN=VDAC3 PE=4 SV=1 - [E5RHE1_HUMAN]
F5H4L7	Uncharacterized protein OS=Homo sapiens GN=VPS26A PE=4 SV=1 - [F5H4L7_HUMAN]
F5GYF5	Uncharacterized protein OS=Homo sapiens GN=VPS35 PE=4 SV=1 - [F5GYF5_HUMAN]
A6NG10	Uncharacterized protein OS=Homo sapiens GN=WBP2 PE=2 SV=2 - [A6NG10_HUMAN]
C9JJ54	Uncharacterized protein OS=Homo sapiens GN=WDFY1 PE=4 SV=1 - [C9JJ54_HUMAN]
F2Z372	Uncharacterized protein OS=Homo sapiens GN=WDFY4 PE=4 SV=1 - [F2Z372_HUMAN]
B4DP38	Uncharacterized protein OS=Homo sapiens GN=WDR77 PE=2 SV=1 - [B4DP38_HUMAN]
C9JY66	Uncharacterized protein OS=Homo sapiens GN=WHSC1L1 PE=3 SV=1 - [C9JY66_HUMAN]
E7EVF6	Uncharacterized protein OS=Homo sapiens GN=WIP1 PE=4 SV=1 - [E7EVF6_HUMAN]
F5GWT4	Uncharacterized protein OS=Homo sapiens GN=WNK1 PE=4 SV=1 - [F5GWT4_HUMAN]
C9K044	Uncharacterized protein OS=Homo sapiens GN=XAF1 PE=4 SV=1 - [C9K044_HUMAN]
Q5T6H7	Uncharacterized protein OS=Homo sapiens GN=XPNPEP1 PE=2 SV=1 - [Q5T6H7_HUMAN]
F2Z2X4	Uncharacterized protein OS=Homo sapiens GN=XPO4 PE=4 SV=1 - [F2Z2X4_HUMAN]

E7ESC6	Uncharacterized protein OS=Homo sapiens GN=XPO7 PE=4 SV=1 - [E7ESC6_HUMAN]
F5H1I8	Uncharacterized protein OS=Homo sapiens GN=XRCC6 PE=4 SV=1 - [F5H1I8_HUMAN]
E9PKX1	Uncharacterized protein OS=Homo sapiens GN=ZCCHC11 PE=4 SV=1 - [E9PKX1_HUMAN]
F5GWB2	Uncharacterized protein OS=Homo sapiens GN=ZNF225 PE=4 SV=1 - [F5GWB2_HUMAN]
F5H6H3	Uncharacterized protein OS=Homo sapiens GN=ZNF268 PE=4 SV=1 - [F5H6H3_HUMAN]
Q96I98	Uncharacterized protein OS=Homo sapiens GN=ZNF324 PE=2 SV=1 - [Q96I98_HUMAN]
C9JLT3	Uncharacterized protein OS=Homo sapiens GN=ZNF502 PE=4 SV=1 - [C9JLT3_HUMAN]
C9JZC2	Uncharacterized protein OS=Homo sapiens GN=ZNF621 PE=4 SV=1 - [C9JZC2_HUMAN]
E5RH94	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [E5RH94_HUMAN]
E9PQI5	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [E9PQI5_HUMAN]
F5GWM9	Uncharacterized protein OS=Homo sapiens PE=4 SV=1 - [F5GWM9_HUMAN]
C9J8M8	Uncharacterized protein OS=Homo sapiens PE=4 SV=2 - [C9J8M8_HUMAN]
C9JH24	Uncharacterized protein OS=Homo sapiens PE=4 SV=2 - [C9JH24_HUMAN]
E9PDI5	Uncharacterized protein OS=Homo sapiens PE=4 SV=2 - [E9PDI5_HUMAN]
Q9H993	UPF0364 protein C6orf211 OS=Homo sapiens GN=C6orf211 PE=1 SV=1 - [CF211_HUMAN]
Q9H7C9	UPF0366 protein C11orf67 OS=Homo sapiens GN=C11orf67 PE=1 SV=1 - [CK067_HUMAN]
Q14667	UPF0378 protein KIAA0100 OS=Homo sapiens GN=KIAA0100 PE=1 SV=3 - [K0100_HUMAN]
Q3KRA6	UPF0538 protein C2orf76 OS=Homo sapiens GN=C2orf76 PE=2 SV=3 - [CB076_HUMAN]
Q5T6V5	UPF0553 protein C9orf64 OS=Homo sapiens GN=C9orf64 PE=1 SV=1 - [CI064_HUMAN]
Q8N8R5	UPF0565 protein C2orf69 OS=Homo sapiens GN=C2orf69 PE=2 SV=1 - [CB069_HUMAN]
Q9Y224	UPF0568 protein C14orf166 OS=Homo sapiens GN=C14orf166 PE=1 SV=1 - [CN166_HUMAN]
Q9NWW4	UPF0587 protein C1orf123 OS=Homo sapiens GN=C1orf123 PE=1 SV=1 - [CA123_HUMAN]
A6NDU8	UPF0600 protein C5orf51 OS=Homo sapiens GN=C5orf51 PE=1 SV=1 - [CE051_HUMAN]
P06132	Uroporphyrinogen decarboxylase OS=Homo sapiens GN=UROD PE=1 SV=2 - [DCUP_HUMAN]
P54725	UV excision repair protein RAD23 homolog A OS=Homo sapiens GN=RAD23A PE=1 SV=1 - [RD23A_HUMAN]
P54727	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1 - [RD23B_HUMAN]
P21281	V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3 - [VATB2_HUMAN]
Q9Y5K8	V-type proton ATPase subunit D OS=Homo sapiens GN=ATP6V1D PE=1 SV=1 - [VATD_HUMAN]
O75348	V-type proton ATPase subunit G 1 OS=Homo sapiens GN=ATP6V1G1 PE=1 SV=3 - [VATG1_HUMAN]
Q9UBQ0	Vacuolar protein sorting-associated protein 29 OS=Homo sapiens GN=VPS29 PE=1 SV=1 - [VPS29_HUMAN]
Q9UN37	Vacuolar protein sorting-associated protein 4A OS=Homo sapiens GN=VPS4A PE=1 SV=1 - [VPS4A_HUMAN]
Q9NP79	Vacuolar protein sorting-associated protein VTA1 homolog OS=Homo sapiens GN=VTA1 PE=1 SV=1 - [VTA1_HUMAN]
P35968	Vascular endothelial growth factor receptor 2 OS=Homo sapiens GN=KDR PE=1 SV=2 - [VGFR2_HUMAN]
O15195	Villin-like protein OS=Homo sapiens GN=VILL PE=2 SV=3 - [VILL_HUMAN]
P18206	Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4 - [VINC_HUMAN]
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4 - [VWF_HUMAN]
Q5MNZ9	WD repeat domain phosphoinositide-interacting protein 1 OS=Homo sapiens GN=WIFI1 PE=1 SV=3 - [WIFI1_HUMAN]
O75083	WD repeat-containing protein 1 OS=Homo sapiens GN=WDR1 PE=1 SV=4 - [WDR1_HUMAN]
Q9BZH6	WD repeat-containing protein 11 OS=Homo sapiens GN=WDR11 PE=1 SV=1 - [WDR11_HUMAN]
Q6ZQ6	WD repeat-containing protein 87 OS=Homo sapiens GN=WDR87 PE=2 SV=3 - [WDR87_HUMAN]
P13010	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]
A4UGR9	Xin actin-binding repeat-containing protein 2 OS=Homo sapiens GN=XIRP2 PE=1 SV=2 - [XIRP2_HUMAN]
Q86Y38	Xylosyltransferase 1 OS=Homo sapiens GN=XYLT1 PE=1 SV=1 - [XYLT1_HUMAN]

Q7Z2W4	Zinc finger CCCH-type antiviral protein 1 OS=Homo sapiens GN=ZC3HAV1 PE=1 SV=3 - [ZCCHV_HUMAN]
Q9UJW7	Zinc finger protein 229 OS=Homo sapiens GN=ZNF229 PE=2 SV=3 - [ZN229_HUMAN]
Q3ZCT1	Zinc finger protein 260 OS=Homo sapiens GN=ZNF260 PE=2 SV=3 - [ZN260_HUMAN]
Q53GI3	Zinc finger protein 394 OS=Homo sapiens GN=ZNF394 PE=2 SV=2 - [ZN394_HUMAN]
Q6AHZ1	Zinc finger protein 518A OS=Homo sapiens GN=ZNF518A PE=1 SV=2 - [Z518A_HUMAN]
Q96C55	Zinc finger protein 524 OS=Homo sapiens GN=ZNF524 PE=2 SV=1 - [ZN524_HUMAN]
Q08AN1	Zinc finger protein 616 OS=Homo sapiens GN=ZNF616 PE=2 SV=2 - [ZN616_HUMAN]
Q5T619	Zinc finger protein 648 OS=Homo sapiens GN=ZNF648 PE=1 SV=1 - [ZN648_HUMAN]
Q6PK81	Zinc finger protein 773 OS=Homo sapiens GN=ZNF773 PE=2 SV=1 - [ZN773_HUMAN]
Q7Z570	Zinc finger protein 804A OS=Homo sapiens GN=ZNF804A PE=1 SV=3 - [Z804A_HUMAN]
B7ZM62	ZMYND8 protein OS=Homo sapiens GN=ZMYND8 PE=2 SV=1 - [B7ZM62_HUMAN]